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(54) Title: COMBINATION OF A COMPOUND HAVING THE ABILITY TO REARRANGE A LYSOSOMAL ENZYME AND AMBROXOL AND/OR A DERIVATIVE OF AMBROXOL

(57) Abstract: It is herewith declared that the information stored on the carrier which is submitted together with the international patent application is identical to the written sequence listing of the international patent application.



WO 2013/091897 A1

Combination of a compound having the ability to rearrange a lysosomal enzyme and Ambroxol and/or a derivative of Ambroxol

The present invention is related to a combination comprising a first constituent and a second constituent, use of the combination, a pharmaceutical preparation comprising a first constituent, a second constituent, optionally a further constituent, use of a pharmaceutical preparation for the manufacture of a medicament, a method for the treatment of a disease and a method for increasing activity of a lysosomal protein having reduced activity.

Lysosomal storage diseases, also referred to herein as lysosomal storage disorders or LSDs, are a group of rare inherited metabolic disorders that result from defects in lysosomal function (Winchester B, et al. (2000), Biochem. Soc. Trans. 28 (2): 150–4). AN LSD results when a specific organelle in the body's cells – the lysosome – malfunctions. The lysosome processes unwanted material via the activity of proteins, such as enzymes, into substances that the cell can utilize. Some of the more prominent LSDs are Pompe's disease and Fabry disease.

LSDs are caused by lysosomal dysfunction usually as a consequence of reduced or absent activity of a single protein, such as a lysosomal enzyme, which is required for the metabolism of lipids, glycoproteins or mucopolysaccharides.

Like other genetic diseases, individuals typically inherit an LSD from their parents. Individually, an LSD occurs with frequencies of less than about 1:100,000 to 1:250,000, however, as a group the incidence is about 1:5,000 - 1:10,000. The overall majority of these disorders are autosomal recessively inherited; only a few are X-linked inherited, such as Fabry disease and Hunter syndrome.

Although each disorder results from different gene mutations that translate into a deficiency or reduction of a lysosomal protein's activity they all share a common biochemical characteristic – nearly all lysosomal disorders originate from an abnormal accumulation of substances inside the lysosome which is due to a deficiency or reduction of the particular lysosomal protein's activity.

LSDs affect mostly children and they often die at a young and unpredictable age, many within a few months or years of birth. Many children die of this disease following years of suffering from various symptoms of their particular disorder. The symptoms of LSDs vary, depending on the particular disorder and other variables like the age of onset, and can be mild to severe. They can include developmental delay, movement disorders, seizures, dementia, deafness and/or blindness. Some people with an LSD have enlarged livers (hepatomegaly) and enlarged spleens (splenomegaly), pulmonary and cardiac problems, and bones that develop abnormally.

There are up to now no causative cures for LSDs available. Therapies which have been tried with different success at least in animal models comprise enzyme replacement therapy, bone marrow transplantation, substrate reduction therapy, gene therapy and pharmacological chaperone therapy. Although the experimental technique of gene therapy may offer such cures in the future, therapies available at present are mainly symptomatic.

In enzyme replacement therapy, also referred to herein as ERT, recombinantly expressed or synthesized enzyme is administered intravenously which compensates for the deficiency of the affected enzyme. Cell receptors mediate the uptake of the recombinant enzyme. The constant need for treatment, the costs associated therewith and particularly the dependence on cell receptor mediated uptake are disadvantageous in ERT.

Bone marrow transplantation also referred to herein as BMT, has been supposed as a therapy for the treatment of LSDs. However, the efficacy and effectiveness of this therapy is in doubt or may be insufficient, as it is still not known whether the number of cells transferred by BMT will provide enough of the missing enzyme to successfully treat patients suffering from an LSD in general.

In substrate reduction therapy, also referred to herein as SRT, upstream inhibition of biological pathways results in reduction of substrate produced for the enzyme to process. In SRT adverse effects may result from an influence of the applied substances on other cellular processes.

Pharmacological chaperone therapy, also referred to herein as PCT, is a technique used to stabilize the lysosomal protein having reduced activity produced by patients and was examined for certain LSDs.

PCT is based on the use of chaperone molecules that assist the folding of mutated enzymes and improve their stability and lysosomal trafficking.

In Pompe disease it was shown for deoxynojirimycin, an iminosugar also referred to herein as DNJ, to increase acid α -glucosidase activity, the mutated enzyme in Pompe disease (Parenti G. et al., Mol Ther. 2007 Mar;15(3):508-14. Epub 2007 Jan 9).

In Fabry disease, which is caused by mutations of lysosomal alpha-galactosidase A, also referred to herein as GLA or α -Gal A, 1-Deoxygalactonojirimycin, also referred to herein as DGJ, AT1001 or Migalastat, an inhibitor of GLA serves as a pharmacological chaperone and enhances the activity of the mutated enzyme (Asano N. et al., Eur J Biochem. 2000 Jul; 267(13):4179-86).

Iminosugars such as DGJ or DNJ may thus be selectively used as pharmacological chaperones to assist rearrangement of lysosomal proteins having reduced activity, such as mutant lysosomal proteins, and allow the treatment of patients with an LSD such as Fabry disease or Pompe disease. Pharmacological Chaperone administered in an effective dose can cause an increase of mutation reduced enzyme activity. Nevertheless, a person skilled in the art will acknowledge that the administration of pharmacological chaperones, such as iminosugars, such as for example Isfagomine applied in Gaucher's Disease, have to be exactly dosed, for example in a scheme where dosing is followed by a withdrawal period that enables the cell and enzyme to temporarily get rid of the inhibitor to display beneficial effect (Khanna et al; FEBS Journal 277 (2010) 1618–1638). It will be also acknowledged that to distinguish between inhibitory and subinhibitory concentrations is difficult and needs to be evaluated in in vivo studies. While treating Fabry's disease, i.e. mutant GLA in a cell-free environment, dosages far below 1 μ M are already inhibiting (Asano et al., supra). Asano and colleagues demonstrated a decrease of GLA activity in patient lymphoblasts while exceeding concentration of 100 μ M DGJ. In a cell culture based GLA overexpression system concentrations up to 1mM (Wu et al., Hum Mutat, 2011 Aug; 32(8):965-77) are tolerated without any detectable inhibition of the tested mutant enzyme. However, it is important to

note that a washout period is also included in the therein described experimental setup, i.e. two hours prior to the conduction of the assay the medium containing the Pharmacological Chaperone is removed, which is indicative for a disturbance of the assay by high concentrations of iminosugars, i.e. inhibition. In accordance therewith it will be also acknowledged that a therapy using higher doses is hard to establish in patients and therefore needs to be considered carefully in clinical trials where 10 μ M are acknowledged as to be a clinically achievable concentration.

Pharmacological chaperone therapy aims to restore the activity of a lysosomal protein having reduced activity. Nevertheless the development of pharmacological chaperon treatment of LSDs is still at an early stage. Its strong dependence on the particular patient which is to be treated, more specifically the particular LSD and its particular cause, for example, a particular mutation of the specific lysosomal protein, affect efficacy and effectiveness of said therapy.

Furthermore the successful administration of pharmacological chaperones against LSDs is hampered by the inhibitory effect and potential toxicity when high doses of pharmacological chaperones are administered (Wu et al., supra; Asano et al., supra and Khanna et al., supra).

The problem underlying the present invention is to provide a means for the treatment of LSDs, and methods for the treatment of LSDs.

It is a further problem underlying the present invention to enhance protein activity of a lysosomal protein having reduced activity.

It is a still further problem of the present invention to enhance the efficacy and effectiveness of a pharmacological chaperone therapy in the treatment of LSDs.

These and other problems underlying the instant invention are solved by the subject matter of the attached independent claims. Preferred embodiments may be taken from the attached dependent claims.

The underlying problems are also solved in a first aspect of the invention by a combination comprising a first constituent and a second constituent, wherein the first constituent is a

compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, and wherein the second constituent is Ambroxol and/or a derivative of Ambroxol.

The underlying problems are also solved in a second aspect of the invention by a combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to the first aspect of the invention; wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is alpha-allo-homonojirimycin or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a third aspect of the invention by a combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to the first aspect of the invention; wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is alpha-galacto-homonojirimycin or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a fourth aspect of the invention by a combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to the first aspect of the invention; wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is alpha-allo-homonojirimycin or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a fifth aspect of the invention by a combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to the first aspect of the invention; wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is beta-1-C-butyl-deoxygalactonojirimycin

or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a sixth aspect of the invention by a combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to the first aspect of the invention; wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,

wherein the first constituent is galactose, preferably D-galactose or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a seventh aspect of the invention by a combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to the first aspect of the invention; wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is N-acetyl-glucosamine-thiazoline (NGT) or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in an eighth aspect of the invention by a combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to the first aspect of the invention; wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 6-acetamido-6-deoxycastanospermine (ACAS) or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a ninth aspect of the invention by a combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to the first aspect of the invention; wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,

wherein the first constituent is bisnaphthalimide nitro-indan-1-one or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a tenth aspect of the invention by a combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to the first aspect of the invention; wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,

wherein the first constituent is pyrrolo[3,4-d]pyridazin-1-one or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in an eleventh aspect of the invention by a combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to the first aspect of the invention; wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is pyrimethamine (PYR) or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a twelfth aspect of the invention by a combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to the first aspect of the invention; wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is N-octyl-4-epi-beta-valienamine (NOEV) or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a 13th aspect of the invention by a combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to the first aspect of the invention; wherein the first constituent is a

compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,

wherein the first constituent is N-butyl-DNJ or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a 14th aspect of the invention by a combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to the first aspect of the invention; wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,

wherein the first constituent is deoxynojirimycin or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a 15th aspect of the invention by the use of a combination of any one of the first, second, third, fourth, fifth, sixth, seventh, eighth, ninth, tenth, eleventh, twelfth, 13th and 14th aspect of the invention, for the manufacture of a medicament for the treatment or prevention of a disease.

The underlying problems are also solved in a 16th aspect of the invention by the use of a compound having the ability to rearrange a lysosomal protein in or for the manufacture of a medicament for the treatment of a disease, wherein the medicament comprises a combination comprising a first constituent and a second constituent, wherein the first constituent is a or the compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, and wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a 17th aspect of the invention by the use Ambroxol and/or a derivative of Ambroxol in or for the manufacture of a medicament for the treatment of a disease, wherein the medicament comprises a combination comprising a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity and wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in an 18th aspect of the invention by a pharmaceutical preparation comprising a first constituent, a second constituent optionally a further constituent,

wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the second constituent is Ambroxol and/or a derivative of Ambroxol and wherein the further constituent is selected from the group comprising a pharmaceutically acceptable excipients and a pharmaceutically active agents, and wherein preferably the pharmaceutical preparation increases the reduced activity of the lysosomal protein, and wherein the lysosomal protein has a reduced activity.

The underlying problems are also solved in a 19th aspect of the invention by pharmaceutical preparation comprising a first constituent and a second constituent, wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to the 18th aspect of the invention; wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 1-deoxygalactonojirimycin or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a 20th aspect of the invention by a pharmaceutical preparation comprising a first constituent and a second constituent, wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to the 18th aspect of the invention; wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is alpha-galacto-homonojirimycin or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a 21st aspect of the invention by a pharmaceutical preparation comprising a first constituent and a second constituent, wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to the 18th

aspect of the invention; wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is alpha-allo-homonojirimycin or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a 22nd aspect of the invention by a pharmaceutical preparation comprising a first constituent and a second constituent, wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to the 18th aspect of the invention; wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is beta-1-C-butyl-deoxygalactonojirimycin or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a 23rd aspect of the invention by a pharmaceutical preparation comprising a first constituent and a second constituent, wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to the 18th aspect of the invention; wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is galactose, preferably D-galactose or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a 24th aspect of the invention by a pharmaceutical preparation comprising a first constituent and a second constituent, wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to the first aspect of the invention; wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is N-acetyl-glucosamine-thiazoline (NGT) or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a 25th aspect of the invention by a pharmaceutical preparation comprising a first constituent and a second constituent, wherein the

pharmaceutical preparation is preferably a pharmaceutical preparation according to the 18th aspect of the invention;

wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 6-acetamido-6-deoxycastanospermine (ACAS) or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a 26th aspect of the invention by a pharmaceutical preparation comprising a first constituent and a second constituent, wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to the 18th aspect of the invention;

wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is bisnaphthalimide nitro-indan-1-one or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a 27th aspect of the invention by a pharmaceutical preparation comprising a first constituent and a second constituent, wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to the 18th aspect of the invention;

wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is pyrrolo[3,4-d]pyridazin-1-one or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a 28th aspect of the invention by a pharmaceutical preparation comprising a first constituent and a second constituent, wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to the 18th

aspect of the invention; wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is pyrimethamine (PYR) or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a 29th aspect of the invention by a pharmaceutical preparation comprising a first constituent and a second constituent, wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to the 18th aspect of the invention;

wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is N-octyl-4-epi-beta-valienamine (NOEV) or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a 30th aspect of the invention by a pharmaceutical preparation comprising a first constituent and a second constituent, wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to the 18th aspect of the invention;

wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is N-butyl-DNJ or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a 31st aspect of the invention by a pharmaceutical preparation comprising a first constituent and a second constituent, wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to the 18th aspect of the invention;

wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is Deoxynojirimycin or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a 32nd aspect of the invention by a method for preparing a pharmaceutical preparation, preferably a pharmaceutical preparation according to any one of the 18th, 19th, 20th, 21st, 22nd, 23rd, 24th, 25th, 26th, 27th, 28th, 29th, 30th and 31st aspect of the invention, comprising the steps of formulating a first constituent as defined in any one of the aspects of the invention and a second constituent as defined in any one of the aspects of the invention into a single dosage form or into two separate dosage forms, wherein in case of two separate dosage forms a first of the two separate dosage forms contains the first constituent and a second of the two separate dosage forms contains the second constituent.

The underlying problems are also solved in a 33rd aspect of the invention by the use of a pharmaceutical preparation for the manufacture of a medicament, wherein the pharmaceutical preparation is a pharmaceutical preparation according to any one of the 18th, 19th, 20th, 21st, 22nd, 23rd, 24th, 25th, 26th, 27th, 28th, 29th, 30th and 31st aspect of the invention, and wherein the pharmaceutical preparation is for the treatment and/or prevention of a disease.

The underlying problems are also solved in a 34th aspect of the invention by a method for the treatment of a disease, wherein the method comprises administering to a subject a first constituent as defined in any one of the first, second, third, fourth, fifth, sixth, seventh, eighth, ninth, tenth, eleventh, twelfth, 13th, 14th, 15th, 16th, 17th, 18th, 19th, 20th, 21st, 22nd, 23rd, 24th, 25th, 26th, 27th, 28th, 29th, 30th and 31st aspect of the invention, prior to, concomitantly with or after a second constituent as defined in any one of the first, second, third, fourth, fifth, sixth, seventh, eighth, ninth, tenth, eleventh, twelfth, 13th, 14th, 15th, 16th, 17th, 18th, 19th, 20th, 21st, 22nd, 23rd, 24th, 25th, 26th, 27th, 28th, 29th, 30th and 31st aspect of the invention, and/or a combination according to any one of the first, second, third, fourth, fifth, sixth, seventh, eighth, ninth, tenth, eleventh, twelfth, 13th and 14th aspect of the invention and/or a pharmaceutical preparation according to any one of the 18th, 19th, 20th, 21st, 22nd, 23rd, 24th, 25th, 26th, 27th, 28th, 29th, 30th and 31st aspect of the invention.

The underlying problems are also solved in a 35th aspect of the invention by a method for increasing activity of a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the method comprises administering to a cell a compound having the ability to rearrange the lysosomal protein, and Ambroxol and/or a derivative thereof.

The underlying problems are also solved in a 36th aspect of the invention by a method for increasing activity of a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the method comprises administering to a subject a compound having the ability to rearrange the lysosomal protein, and Ambroxol and/or a derivative thereof, wherein the compound having the ability to rearrange the lysosomal protein increases the activity of the lysosomal protein.

The underlying problems are also solved in a 37th aspect of the invention by a method for increasing activity of a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the method comprises, administering to a subject a first constituent as defined in any one first, second, third, fourth, fifth, sixth, seventh, eighth, ninth, tenth, eleventh, twelfth, 13th, 14th, 15th, 16th, 17th, 18th, 19th, 20th, 21st, 22nd, 23rd, 24th, 25th, 26th, 27th, 28th, 29th, 30th and 31st aspect of the invention and, prior to, concomitantly with or after a second constituent as defined in any one the first, second, third, fourth, fifth, sixth, seventh, eighth, ninth, tenth, eleventh, twelfth, 13th, 14th, 15th, 16th, 17th, 18th, 19th, 20th, 21st, 22nd, 23rd, 24th, 25th, 26th, 27th, 28th, 29th, 30th and 31st aspect of the invention, and/or a combination according to any one of the first, second, third, fourth, fifth, sixth, seventh, eighth, ninth, tenth, eleventh, twelfth, 13th and 14th aspect of the invention and/or a pharmaceutical preparation according to any one of the 18th, 19th, 20th, 21st, 22nd, 23rd, 24th, 25th, 26th, 27th, 28th, 29th, 30th and 31st aspect of the invention, wherein the combination and/or the pharmaceutical preparation increases the activity of the lysosomal protein.

The underlying problems are also solved in a 38th aspect of the invention by a combination comprising a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity wherein the reduced activity is reduced due to a mutation of the lysosomal protein, and wherein the second constituent is Ambroxol and/or a derivative of Ambroxol, wherein the combination is preferably a combination according to any one of the

first, second, third, fourth, fifth, sixth, seventh, eighth, ninth, tenth, eleventh, twelfth, 13th and 14th aspect of the invention, for use in a method of personalized therapeutic treatment of a subject, wherein the method comprises the following steps:

step a): determining whether in a sample of the subject the lysosomal protein has a reduced activity, preferably such reduced activity results from one or more mutation of the lysosomal protein compared to the wild type lysosomal protein;

step b): identifying a compound having the ability to rearrange the lysosomal protein having reduced activity, and wherein the compound is suitable for or is increasing the reduced activity of the lysosomal protein; and

step c): administering to the subject the first constituent prior to, concomitantly with or after the second constituent.

The underlying problems are also solved in a 39th aspect of the invention by pharmaceutical preparation comprising a first constituent, a second constituent and optionally a further constituent,

wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the reduced activity is reduced due to a mutation of the lysosomal protein, and wherein the second constituent is Ambroxol and/or a derivative of Ambroxol, wherein the further constituent is selected from the group comprising pharmaceutically acceptable excipients and pharmaceutically active agents, wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of the 18th, 19th, 20th, 21st, 22nd, 23rd, 24th, 25th, 26th, 27th, 28th, 29th, 30th and 31st aspect of the invention, for use in a method of personalized therapeutic treatment of a subject, wherein the method comprises the following steps:

step a): determining whether in a sample of the subject the lysosomal protein has a reduced activity, preferably such reduced activity results from one or more mutation of the lysosomal protein compared to the wild type lysosomal protein;

step b): identifying a compound having the ability to rearrange the lysosomal protein having reduced activity, and wherein the compound is suitable for or is increasing the reduced activity of the lysosomal protein; and

step c): administering to the subject the first constituent prior to, concomitantly with or after the second constituent.

Other aspects and embodiments of the invention are disclosed in the following. For avoidance of doubt, any embodiment referred to herein is preferably an embodiment of the invention.

Embodiment 1: A combination comprising a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, and wherein the second constituent is Ambroxol and/or a derivative of Ambroxol.

Embodiment 2: The combination according to embodiment 1, wherein the lysosomal protein is affected in a disease.

Embodiment 3: The combination according to any one of embodiments 1 to 2, wherein the lysosomal protein is selected from the group comprising an enzyme, a trans-membrane protein or a soluble non-enzymatic protein.

Embodiment 4: The combination according to embodiment 3, wherein the lysosomal protein is an enzyme, wherein the enzyme has a reduced activity, preferably a reduced activity due to a mutation of the enzyme.

Embodiment 5: The combination according to embodiment 4, wherein the enzyme is selected from the group comprising a lysosomal hydrolase and a phosphotransferase.

Embodiment 6: The combination according to any one of embodiments 1 to 5, wherein the compound having the ability to rearrange a lysosomal protein is a chaperon.

Embodiment 7: The combination according to embodiment 6, wherein the chaperon is a pharmacological chaperon or a pharmaceutically acceptable salt, solvate or derivative thereof.

Embodiment 8: The combination according to embodiment 7, wherein the pharmacological chaperon is a sugar and/or an imino sugar or a pharmaceutically acceptable salt, solvate or derivative of the sugar and/or the imino sugar.

Embodiment 9: The combination according to embodiment 9, wherein the sugar is galactose, preferably D-galactose.

Embodiment 10: The combination according to embodiment 8, wherein the imino sugar is selected from the group comprising 1-deoxygalactonojirimycin (DGJ), alpha-galactohomonojirimycin, alpha-allo-homonojirimycin, beta-1-C-butyl-deoxygalactonojirimycin, beta-1-C-butyl-deoxynojirimycin, N-nonyl-deoxynojirimycin (NN-DNJ), N-octyl-2,5-anhydro-2,5-imino-D-glucitol, N-octyl-isofagomine, N-octyl-beta-valienamine (NOV), Isofagomine (IFG), calystegine A3, calystegine B1, calystegine B2, calystegine C1, 1,5-dideoxy-1,5-iminoxylitol (DIX), alpha-1-C-nonyl-DIX, alpha-1-C-octyl-1-DNJ, N-acetylglucosamine-thiazoline (NGT), 6-acetamido-6-deoxycastanospermine (ACAS), bisnaphthalimide nitro-indan-1-one, pyrrolo[3,4-d]pyridazin-1-one, pyrimethamine (PYR), N-octyl-4-epi-beta-valienamine (NOEV), N-butyl-DNJ, Deoxynojirimycin (DNJ), N-Acetylgalactosamine (GalNAc), 2-Acetamido-1,2-dideoxynojirimycin (AdDNJ), N-dodecyl-DNJ, 6-nonyl-isofagomine, N-methyl calystegine A3, calystegine B2, 4-epi-isofagomine, 1-deoxynojirimycin, alpha-homonojirimycin, castanospermine, 1-deoxymannonojirimycin, Swainsonine, Mannostatin A, 2-hydroxy-isofagomine, 1-deoxyfuconojirimycin, beta-homofuconojirimycin, 2,5-imino-1,2,5-trideoxy-L-glucitol, 2,5-dideoxy-2,5-imino-D-fucitol, 2,5-imino-1,2,5-trideoxy-D-altritol, 1,2-dideoxy-2-N-acetamido-nojirimycin, 1,2-dideoxy-2-N-acetamido-galaconojirimycin, 2-N-acetylamino-isofagomine, 1,2-dideoxy-2-acetamido-nojirimycin, nagastain, 2-N-acetamido-isofagomine, 1,2-dideoxy-2-acetamido-nojirimycin, 1-deoxyiduronojirimycin, 2-carboxy-3,4,5-trideoxypiperidine, 6-carboxy-isofagomine, 2,6-dideoxy-2,6-imino-sialic acid, Siastin B and Castanospermine (CAS), and derivatives thereof; and pharmaceutically acceptable salts thereof.

Embodiment 11: The combination according to any one of embodiments 1 to 10, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 12: The combination according to any one of embodiments 1 to 11, wherein at least one of the first constituent and the second constituent is/are present as a solvate or a pharmaceutically acceptable salt thereof.

Embodiment 13: The combination according to embodiment 12, wherein the second constituent is Ambroxol, wherein Ambroxol is present as a pharmaceutically acceptable salt of Ambroxol, and wherein, preferably, the pharmaceutically acceptable salt of Ambroxol is Ambroxol hydrochloride.

Embodiment 14: The combination according to embodiment 12, wherein the second constituent is a derivative of Ambroxol, wherein the derivative of Ambroxol is bromhexine, and wherein preferably the pharmaceutically acceptable salt of bromhexine is bromhexine hydrochloride.

Embodiment 15: The combination according to any one of embodiments 1 to 14, wherein the combination is suitable for or is for use in the treatment and/or prevention of a disease.

Embodiment 16: The combination according to any one of embodiments 1 to 15, wherein the combination is a pharmaceutical combination.

Embodiment 17: The combination according to any one of embodiments 1 to 16, wherein the combination is suitable for or is for use in a method for the treatment of a subject comprising the administration of the combination to the subject.

Embodiment 18: The combination according to any one of embodiments 15 to 17, wherein the disease is a lysosomal storage disease.

Embodiment 19: The combination according to embodiment 18, wherein the lysosomal storage disease is a lysosomal storage disease having a defective degradation of a sphingolipid component.

Embodiment 20: The combination according to embodiment 19, wherein the lysosomal storage disease having a defective degradation of a sphingolipid component is selected from the group comprising Fabry disease, Gaucher disease type I, Gaucher disease type II, Gaucher disease type III, acid sphingomyelinase deficiency, Farber disease, GM1 gangliosidosis type I, GM1 gangliosidosis type II, GM1 gangliosidosis type III, GM2 gangliosidosis, Krabbe disease, Metachromatic leukodystrophy type I, Metachromatic leukodystrophy type II and Metachromatic leukodystrophy type III.

Embodiment 21: The combination according to embodiment 19, wherein the lysosomal storage disease having a defective degradation of a sphingolipid component is selected from the group consisting of Fabry disease, acid sphingomyelinase deficiency, Farber disease, GM1 gangliosidosis type I, GM1 gangliosidosis type II, GM1 gangliosidosis type III, GM2 gangliosidosis, Krabbe disease, Metachromatic leukodystrophy type I, Metachromatic leukodystrophy type II and Metachromatic leukodystrophy type III.

Embodiment 22: The combination according to embodiment 20 or embodiment 21, wherein the lysosomal storage disease is Fabry disease.

Embodiment 23: The combination according to embodiment 20 or embodiment 21, wherein the acid sphingomyelinase deficiency is selected from the group comprising Niemann-Pick type A and type Niemann-Pick B.

Embodiment 24: The combination according to embodiment 20 or embodiment 21, wherein the GM2 gangliosidosis is selected from the group comprising Tay-Sachs type I, Tay-Sachs type II, Tay-Sachs type III and Sandhoff.

Embodiment 25: The combination according to embodiment 24, wherein the lysosomal storage disease is Tay-Sachs type I, Tay-Sachs type II or Tay-Sachs type III.

Embodiment 26: The combination according to embodiment 24, wherein the lysosomal storage disease is Sandhoff.

Embodiment 27: The combination according to embodiment 20 or embodiment 21, wherein the lysosomal storage disease is GM1 gangliosidosis type I, GM1 gangliosidosis type II or GM1 gangliosidosis type III

Embodiment 28: The combination according to embodiment 18, wherein the lysosomal storage disease is a lysosomal storage disease having a defective metabolism of glycosaminoglycans.

Embodiment 29: The combination according to embodiment 28, wherein the lysosomal storage disease having a defective metabolism of glycosaminoglycans is selected from the group comprising MPS I, MPS II, MPS III, MPS IV, MPS VI, MPS VI and MPS IX.

Embodiment 30: The combination according to embodiment 29, wherein MPS I is selected from the group comprising Hurler disease, Hurler-Scheie syndrome and Scheie syndrome.

Embodiment 31: The combination according to embodiment 29, wherein MPS II is selected from the group comprising Hunter syndrome.

Embodiment 32: The combination according to embodiment 29, wherein MPS III is selected from the group comprising Sanfilippo syndrome Type A, Sanfilippo syndrome Type B, Sanfilippo syndrome Type C and Sanfilippo syndrome Type D.

Embodiment 33: The combination according to embodiment 29, wherein MPS IV is selected from the group comprising Morquio type A and Morquio type B.

Embodiment 34: The combination according to embodiment 29, wherein MPS VI is selected from the group comprising Maroteaux-Lamy.

Embodiment 35: The combination according to embodiment 29, wherein MPS VII is selected from the group comprising Sly.

Embodiment 36: The combination according to embodiment 29, wherein MPS IX is selected from the group comprising Hyaluronidase deficiency and Multiple Sulfatase deficiency.

Embodiment 37: The combination according to embodiment 18, wherein the lysosomal storage disease is a lysosomal storage disease having a defective degradation of a glycan portion of a glycoprotein.

Embodiment 38: The combination according to embodiment 37, wherein the lysosomal storage disease having a defective degradation of a glycan portion of a glycoprotein is selected from the group comprising Aspartylglucosaminuria, Fucosidosis type I, Fucosidosis type II, Mannosidosis, Sialidosis type I and Sialidosis type II.

Embodiment 39: The combination according to embodiment 18, wherein the lysosomal storage disease is a lysosomal storage disease having a defective degradation of glycogen.

Embodiment 40: The combination according to embodiment 39, wherein the lysosomal storage disease having a defective degradation of glycogen is selected from the group comprising Pompe's disease.

Embodiment 41: The combination according to embodiment 40, wherein the lysosomal storage disease is Pompe's disease.

Embodiment 42: The combination according to embodiment 18, wherein the lysosomal storage disease is a lysosomal storage disease having a defective degradation of polypeptides.

Embodiment 43: The combination according to embodiment 42, wherein the lysosomal storage disease having a defective degradation of polypeptides is selected from the group comprising pycnodysostosis.

Embodiment 44: The combination according to embodiment 18, wherein the lysosomal storage disease is a lysosomal storage disease having a defective degradation or transport of cholesterol, cholesterol esters and/or other complex lipids.

Embodiment 45: The combination according to embodiment 44, wherein the lysosomal storage disease having a defective degradation or transport of cholesterol, cholesterol esters and/or other complex lipids is selected from the group comprising Neuronal Ceroid Lipofuscinosis type I, Neuronal Ceroid Lipofuscinosis type II, Neuronal Ceroid Lipofuscinosis type III and Neuronal Ceroid Lipofuscinosis type IV.

Embodiment 46: The combination according to embodiment 18, wherein the lysosomal storage disease is a lysosomal storage disease having multiple deficiencies of lysosomal enzymes.

Embodiment 47: The combination according to embodiment 46, wherein the lysosomal storage disease having multiple deficiencies of lysosomal enzymes is selected from the group comprising Galactosialidosis, Danon Disease, Pyknodysostosis, multiple sulfatase deficiency, GM2 Gangliosidosis, Mucopolidosis type II and Mucopolidosis type III.

Embodiment 48: The combination according to embodiment 47, wherein the lysosomal storage disease is Mucopolidosis type II or Mucopolidosis type III.

Embodiment 49: The combination according to embodiment 18, wherein the lysosomal storage disease is a lysosomal storage disease having transport and trafficking defects.

Embodiment 50: The combination according to embodiment 49, wherein the lysosomal storage disease having transport and trafficking defects is selected from the group comprising Cystinosis, Danon disease, Mucopolidosis type IV, Infantile sialic acid storage disease and Salla disease.

Embodiment 51: The combination according to any one of embodiments 5 to 50, wherein the lysosomal hydrolase is selected from the group comprising alpha-galactosidase A, alpha-glucosidase, sphingomyelinase, glucocerebrosidase, acid beta-glucosidase, N-acetylglucosamine-1-phosphotransferase, β -Hexosaminidase A, β -Hexosaminidase B, α -L-Iduronidase, Iduronatsulfatsulfatase, Heparansulfatsulfamidase, α -N-Acetylglukoseamidase, α -Glukosaminid-N-Acetyltransferase, N-Acetyl-glukosamin-6-sulfatsulfatase, β -

Galactosidase, N-Acetylgalactosamin-4-sulfat-Sulfatase, β -Glucuronidase, Arylsulfatase A, Arylsulfatase B, Neuraminidase, α -Fucosidase, acid Ceramidase, Aspartylglukosaminidase, α -Mannosidase, β -Mannosidase, acid Lipase, acid β -Galactosidase, galactocerebrosidase, acid α -Mannosidase, acid β -Mannosidase, acid α -L-fucosidase, α -N-Acetylgalactosaminidase, α -N-Acetylglucosaminidase, β -glucuronidase, sialidase and galactosylceramidase.

Embodiment 52: The combination according to any one of embodiments 5 to 51, wherein the lysosomal hydrolase is selected from the group consisting of α -galactosidase A, α -glucosidase, sphingomyelinase, N-acetylglucosamine-1-phosphotransferase, β -Hexosaminidase A, Hexosaminidase B, α -L-Iduronidase, Iduronatsulfat sulfatase, Heparansulfatsulfamidase, α -N-Acetylglukoseamidase, α -Glukosaminid-N-Acetyltransferase, N-Acetyl-glukosamin-6-sulfatsulfatase, β -Galactosidase, N-Acetylgalactosamin-4-sulfat-Sulfatase, β -Glucuronidase, Arylsulfatase A, Arylsulfatase B, Neuraminidase, α -Fucosidase, acid Ceramidase, Aspartylglukosaminidase, α -Mannosidase, β -Mannosidase, acid Lipase, acid β -Galactosidase, galactocerebrosidase, acid α -Mannosidase, acid β -Mannosidase, acid α -L-fucosidase, α -N-Acetylglucosaminidase, α -N-Acetylgalactosaminidase, β -glucuronidase, sialidase and galactosylceramidase.

Embodiment 53: The combination according to any one of embodiments 3 to 4 and 6 to 17, wherein the lysosomal protein is a trans-membrane protein, wherein the trans-membrane protein has a reduced activity, preferably a reduced activity due to a mutation of the trans-membrane protein.

Embodiment 54: The combination according to any one of embodiments 3 to 4, 6 to 17 and 53, preferably embodiment 53, wherein the trans-membrane protein is a trans-membrane protein affected in a lysosomal storage disease.

Embodiment 55: The combination according to any one of embodiments 3 to 4, 6 to 17, 53 and 54, preferably embodiment 54, wherein the trans-membrane protein is selected from the group comprising NPC1, Acetyl-CoA α -Glukosaminid-N-Acetyltransferase, LAMP2, lysosomal free sialic acid transporter SLC17A5.

Embodiment 56: The combination according to embodiment 55, wherein the trans-membrane protein is NPC1.

Embodiment 57: The combination according to embodiment 56, wherein the lysosomal storage disease is Niemann-Pick type C1.

Embodiment 58: The combination according to embodiment 55, wherein the trans-membrane protein is Acetyl-CoA α -Glukosaminid-N-Acetyltransferase.

Embodiment 59: The combination according to embodiment 58, wherein the lysosomal storage disease is Sanfillipo syndrome Typ C.

Embodiment 60: The combination according to embodiment 55, wherein the trans-membrane protein is LAMP2.

Embodiment 61: The combination according to embodiment 60, wherein the lysosomal storage disease is Danon Disease.

Embodiment 62: The combination according to embodiment 60, wherein the lysosomal storage disease is Cystinosis.

Embodiment 63: The combination according to embodiment 55, wherein the trans-membrane protein is lysosomal free sialic acid transporter SLC17A5.

Embodiment 64: The combination according to embodiment 63, wherein the lysosomal storage disease is free sialic acid storage disease.

Embodiment 65: The combination according to any one of embodiments 3 to 4 and 6 to 17, wherein the lysosomal protein is a soluble non-enzymatic protein, wherein the soluble non-enzymatic protein has a reduced activity, preferably a reduced activity due to a mutation of the soluble non-enzymatic protein.

Embodiment 66: The combination according to any one of embodiments 3 to 4, 6 to 17 and 65, wherein the soluble non-enzymatic protein is a soluble non-enzymatic protein affected in a lysosomal storage disease.

Embodiment 67: The combination according to any one of embodiments 3 to 4, 6 to 17, 65 and 66, preferably embodiment 66, wherein the soluble non-enzymatic protein is NPC2.

Embodiment 68: The combination according to any one of embodiments 65 to 67, wherein the lysosomal storage disease is Niemann-Pick type C2.

Embodiment 69: The combination according to any one of embodiments 15 to 17, wherein the disease is different from a lysosomal storage disease.

Embodiment 70: The combination according to embodiment 69, wherein the disease different from a lysosomal storage disease is Parkinson disease.

Embodiment 71: The combination according to any one of embodiments 1 to 70, wherein the first constituent is 1-deoxygalactonojirimycin or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 72: The combination according to any one of embodiments 1 to 70, wherein the first constituent is alpha-galacto-homonojirimycin or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 73: The combination according to any one of embodiments 1 to 70, wherein the first constituent is alpha-allo-homonojirimycin or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 74: The combination according to any one of embodiments 1 to 70,

wherein the first constituent is beta-1-C-butyl-deoxygalactonojirimycin or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 75: The combination according to any one of embodiments 1 to 70, wherein the first constituent is galactose, preferably D-galactose, or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 76: The combination according to any one of embodiments 71 to 75,

wherein the lysosomal protein is affected in a disease;

wherein the lysosomal protein is α -galactosidase A; and/or

wherein the disease preferably is Fabry disease.

Embodiment 77: The combination according to any one of embodiments 1 to 70,

wherein the first constituent is N-butyl-DNJ or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 78: The combination according to any one of embodiments 1 to 70,

wherein the first constituent is Deoxynojirimycin or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 79: The combination according to any one of embodiments 77 to 78,

wherein the lysosomal protein is affected in a disease;

wherein the lysosomal protein is alpha-glucosidase; and/or

wherein the disease preferably is Pompe disease.

Embodiment 80: The combination according to any one of embodiments 1 to 70,

wherein the first constituent is N-acetyl-glucosamine-thiazoline (NGT) or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 81: The combination according to any one of embodiments 1 to 70, wherein the first constituent is 6-acetamido-6-deoxycastanospermine (ACAS) or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 82: The combination according to any one of embodiments 1 to 70, wherein the first constituent is bisnaphthalimide nitro-indan-1-one or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 83: The combination according to any one of embodiments 1 to 70, wherein the first constituent is pyrrolo[3,4-d]pyridazin-1-one or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 84: The combination according to any one of embodiments 1 to 70, wherein the first constituent is pyrimethamine (PYR) or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 85: The combination according to any one of embodiments 80 to 84, wherein the lysosomal protein is affected in a disease; wherein the lysosomal protein is beta-hexosaminidase A/B; and/or wherein the disease preferably is Tay-Sachs and/or Sandhoff disease.

Embodiment 86: The combination according to any one of embodiments 1 to 70, wherein the first constituent is N-octyl-4-epi-beta-valienamine (NOEV) or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 87: The combination according embodiment 86, wherein the lysosomal protein is affected in a disease; wherein the lysosomal protein is beta-galactosidase; and/or

wherein the disease preferably is Gm1-Gangliosidosis.

Embodiment 88: The combination according to any one of embodiments 1 to 70, wherein the lysosomal protein is affected in a disease; wherein the lysosomal protein is acid sphingomyelinase; and/or wherein the disease preferably is Niemann-Pick disease type A or Niemann-Pick disease type B.

Embodiment 89: The combination according to any one of embodiments 1 to 70, wherein the lysosomal protein is affected in a disease; wherein the lysosomal protein is N-acetylglucosamine-1-phosphotransferase; and/or wherein the disease preferably is mucopolidosis type II or mucopolidosis type IIIA.

Embodiment 90: The combination according to any one of embodiments 1 to 70, wherein the first constituent is N-nonyl-deoxynojirimycin (NN-DNJ) or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 91: The combination according to any one of embodiments 1 to 70, wherein the first constituent is N-octyl-2,5-anhydro-2,5-imino-D-glucitol or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 92: The combination according to any one of embodiments 1 to 70, wherein the first constituent is N-octyl-isofagomine or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 93: The combination according to any one of embodiments 1 to 70, wherein the first constituent is N-octyl-beta-valienamine (NOV) or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 94: The combination according to any one of embodiments 1 to 70, wherein the first constituent is Isofagomine (IFG) or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 95: The combination according to any one of embodiments 1 to 70, wherein the first constituent is calystegine A3 or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 96: The combination according to any one of embodiments 1 to 70, wherein the first constituent is calystegine B1 or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 97: The combination according to any one of embodiments 1 to 70, wherein the first constituent is calystegine B2 or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 98: The combination according to any one of embodiments 1 to 70, wherein the first constituent is calystegine C1 or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 99: The combination according to any one of embodiments 1 to 70, wherein the first constituent is 1,5-dideoxy-1,5-iminoxylitol (DIX) or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 100: The combination according to any one of embodiments 1 to 70, wherein the first constituent is alpha-1-C-nonyl-DIX or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 101: The combination according to any one of embodiments 1 to 70, wherein the first constituent is alpha-1-C-octyl-1-DNJ or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 102: The combination according to any one of embodiments 1 to 70, wherein the first constituent is N-butyl-DNJ or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

Embodiment 103: The combination according to any one of embodiments 90 to 102, wherein the lysosomal protein is affected in a disease; and/or wherein the lysosomal protein is glucocerebrosidase; and/or wherein the disease preferably is Parkinson disease.

Embodiment 104: A combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to any one of embodiments 1 to 70; wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 1-deoxygalactonojirimycin or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 105: The combination according to embodiment 104, wherein the lysosomal protein is α -galactosidase A.

Embodiment 106: The combination according to any one of embodiments 104 to 105, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 107: The combination according to any one of embodiments 104 to 106, wherein the combination is for use in the treatment or prevention of Fabry's disease.

Embodiment 108: A combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to any one of embodiments 1 to 70;

wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,

wherein the first constituent is alpha-galacto-homonojirimycin or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 109: The combination according to embodiment 108, wherein the lysosomal protein is α -galactosidase A.

Embodiment 110: The combination according to any one of embodiments 108 to 109, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 111: The combination according to any one of embodiments 108 to 110, wherein the combination is for use in the treatment or prevention of Fabry's disease.

Embodiment 112: A combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to any one of embodiments 1 to 70;

wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,

wherein the first constituent is alpha-allo-homonojirimycin or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 113: The combination according to embodiment 112, wherein the lysosomal protein is α -galactosidase A.

Embodiment 114: The combination according to any one of embodiments 112 to 113, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 115: The combination according to any one of embodiments 112 to 114, wherein the combination is for use in the treatment or prevention of Fabry's disease.

Embodiment 116: A combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to any one of embodiments 1 to 70;

wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,

wherein the first constituent is beta-1-C-butyl-deoxygalactonojirimycin or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 117: The combination according to embodiment 116, wherein the lysosomal protein is α -galactosidase A.

Embodiment 118: The combination according to any one of embodiments 116 to 117, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 119: The combination according to any one of embodiments 116 to 118, wherein the combination is for use in the treatment or prevention of Fabry's disease.

Embodiment 120: A combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to any one of embodiments 1 to 70;

wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,

wherein the first constituent is galactose, preferably D-galactose or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 121: The combination according to embodiment 120, wherein the lysosomal protein is α -galactosidase A.

Embodiment 122: The combination according to any one of embodiments 120 to 121, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 123: The combination according to any one of embodiments 120 to 122, wherein the combination is for use in the treatment or prevention of Fabry's disease.

Embodiment 124: A combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to any one of embodiments 1 to 70;

wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,

wherein the first constituent is N-acetyl-glucosamine-thiazoline (NGT) or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 125: The combination according to embodiment 124, wherein the lysosomal protein is beta-hexosaminidase A/B.

Embodiment 126: The combination according to any one of embodiments 124 to 125, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 127: The combination according to any one of embodiments 124 to 126, wherein the combination is for use in the treatment or prevention of Tay-Sachs or Sandhoff.

Embodiment 128: A combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to any one of embodiments 1 to 70;

wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,

wherein the first constituent is 6-acetamido-6-deoxycastanospermine (ACAS) or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 129: The combination according to embodiment 128, wherein the lysosomal protein is beta-hexosaminidase A/B.

Embodiment 130: The combination according to any one of embodiments 128 to 129, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 131: The combination according to any one of embodiments 128 to 130, wherein the combination is for use in the treatment or prevention of Tay-Sachs or Sandhoff.

Embodiment 132: A combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to any one of embodiments 1 to 70;

wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,

wherein the first constituent is bisnaphthalimide nitro-indan-1-one or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 133: The combination according to embodiment 132, wherein the lysosomal protein is beta-hexosaminidase A/B.

Embodiment 134: The combination according to any one of embodiments 132 to 133, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 135: The combination according to any one of embodiments 132 to 134, wherein the combination is for use in the treatment or prevention of Tay-Sachs or Sandhoff.

Embodiment 136: A combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to any one of embodiments 1 to 70;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is pyrrolo[3,4-d]pyridazin-1-one or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 137: The combination according to embodiment 136, wherein the lysosomal protein is beta-hexosaminidase A/B.

Embodiment 138: The combination according to any one of embodiments 136 to 137, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 139: The combination according to any one of embodiments 136 to 138, wherein the combination is for use in the treatment or prevention of Tay-Sachs or Sandhoff.

Embodiment 140: A combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to any one of embodiments 1 to 70;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is pyrimethamine (PYR) or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 141: The combination according to embodiment 140, wherein the lysosomal protein is beta-hexosaminidase A/B.

Embodiment 142: The combination according to any one of embodiments 140 to 141, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 143: The combination according to any one of embodiments 140 to 142, wherein the combination is for use in the treatment or prevention of Tay-Sachs or Sandhoff.

Embodiment 144: A combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to any one of embodiments 1 to 70;

wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,

wherein the first constituent is N-octyl-4-epi-beta-valienamine (NOEV) or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 145: The combination according to embodiment 144, wherein the lysosomal protein is beta-galactosidase.

Embodiment 146: The combination according to any one of embodiments 144 to 145, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 147: The combination according to any one of embodiments 144 to 146, wherein the combination is for use in the treatment or prevention of GM1-Gangliosidosis.

Embodiment 148: A combination comprising a first constituent and a second constituent,

wherein the combination is preferably a combination according to any one of embodiments 1 to 70;

wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,

wherein the first constituent is N-butyl-DNJ or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 149: The combination according to embodiment 148, wherein the lysosomal protein is alpha-glucosidase.

Embodiment 150: The combination according to any one of embodiments 148 to 149, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 151: The combination according to any one of embodiments 148 to 150, wherein the combination is for use in the treatment or prevention of Pompe's disease.

Embodiment 152: A combination comprising a first constituent and a second constituent, wherein the combination is preferably a combination according to any one of embodiments 1 to 70;

wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,

wherein the first constituent is deoxynojirimycin or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 153: The combination according to embodiment 152, wherein the lysosomal protein is alpha-glucosidase.

Embodiment 154: The combination according to any one of embodiments 152 to 153, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 155: The combination according to any one of embodiments 152 to 154, wherein the combination is for use in the treatment or prevention of Pompe's disease.

Embodiment 156: Use of a combination as defined in any one of embodiments 1 to 155, for the manufacture of a medicament for the treatment or prevention of a disease.

Embodiment 157: Use according to embodiment 156, wherein the disease is a lysosomal storage disease.

Embodiment 158: Use according to embodiment 157, wherein the lysosomal storage disease is a lysosomal storage disease having a defective degradation of a sphingolipid component.

Embodiment 159: Use according to embodiment 158, wherein the lysosomal storage disease having a defective degradation of a sphingolipid component is selected from the group comprising Fabry disease, Gaucher disease type I, Gaucher disease type II, Gaucher disease type III, acid sphingomyelinase deficiency, Farber disease, GM1 gangliosidosis type I, GM1 gangliosidosis type II, GM1 gangliosidosis type III, GM2 gangliosidosis, Krabbe disease, Metachromatic leukodystrophy type I, Metachromatic leukodystrophy type II and Metachromatic leukodystrophy type III.

Embodiment 160: Use according to embodiment 158, wherein the lysosomal storage disease having a defective degradation of a sphingolipid component is selected from the group consisting of Fabry disease, acid sphingomyelinase deficiency, Farber disease, GM1 gangliosidosis type I, GM1 gangliosidosis type II, GM1 gangliosidosis type III, GM2 gangliosidosis, Krabbe disease, Metachromatic leukodystrophy type I, Metachromatic leukodystrophy type II and Metachromatic leukodystrophy type III.

Embodiment 161: Use according to any one of embodiments 159 to 160, wherein the lysosomal storage disease having a defective degradation of sphingolipid components is Fabry disease.

Embodiment 162: Use according to any one of embodiments 159 to 160, wherein the acid sphingomyelinase deficiency is selected from the group comprising Niemann-Pick type A and Niemann-Pick type A and B.

Embodiment 163: Use according to any one of embodiments 159 to 160, wherein the GM2 gangliosidosis is selected from the group comprising Tay-Sachs type I, Tay-Sachs type II, Tay-Sachs type III and Sandhoff.

Embodiment 164: Use according to embodiment 163, wherein the lysosomal storage disease is Tay-Sachs type I, Tay-Sachs type II or Tay-Sachs type III.

Embodiment 165: Use according to embodiment 163, wherein the lysosomal storage disease is Sandhoff.

Embodiment 166: Use according to any one of embodiments 159 to 160, wherein the lysosomal storage disease is GM1 gangliosidosis type I, GM1 gangliosidosis type II or GM1 gangliosidosis type III.

Embodiment 167: Use according to embodiment 157, wherein the lysosomal storage disease is a lysosomal storage disease having a defective metabolism of glycosaminoglycans.

Embodiment 168: Use according to embodiment 167, wherein the lysosomal storage disease having a defective metabolism of glycosaminoglycans is selected from the group comprising MPS I, MPS II, MPS III, MPS IV, MPS VI, MPS VI and MPS IX.

Embodiment 169: Use according to embodiment 168, wherein MPS I is selected from the group comprising Hurler disease, Hurler-Scheie syndrome and Scheie syndrome.

Embodiment 170: Use according to embodiment 168, wherein MPS II is selected from the group comprising Hunter syndrome.

Embodiment 171: Use according to embodiment 168, wherein MPS III is selected from the group comprising Sanfilippo syndrome type A, Sanfilippo syndrome type B, Sanfilippo syndrome type C and Sanfilippo syndrome type D.

Embodiment 172: Use according to embodiment 168, wherein MPS IV is selected from the group comprising Morquio type A and Morquio type B.

Embodiment 173: Use according to embodiment 168, wherein MPS VI is selected from the group comprising Maroteaux-Lamy.

Embodiment 174: Use according to embodiment 168, wherein MPS VII is selected from the group comprising Sly.

Embodiment 175: Use according to embodiment 168, wherein MPS IX is selected from the group comprising Hyaluronidase deficiency and Multiple Sulfatase deficiency.

Embodiment 176: Use according to embodiment 157, wherein the lysosomal storage disease is a lysosomal storage disease having a defective degradation of a glycan portion of a glycoprotein.

Embodiment 177: Use according to embodiment 176, wherein the lysosomal storage disease having a defective degradation of a glycan portion of a glycoprotein is selected from the group comprising Aspartylglucosaminuria, Fucosidosis type I, Fucosidosis type II, Mannosidosis, Sialidosis type I and Sialidosis type II.

Embodiment 178: Use according to embodiment 157, wherein the lysosomal storage disease is a lysosomal storage disease having a defective degradation of glycogen.

Embodiment 179: Use according to embodiment 178, wherein the lysosomal storage disease having a defective degradation of glycogen is selected from the group comprising Pompe's disease.

Embodiment 180: Use according to embodiment 179, wherein the lysosomal storage disease is Pompe's disease.

Embodiment 181: Use according to embodiment 157, wherein the lysosomal storage disease is a lysosomal storage disease having a defective degradation of polypeptides.

Embodiment 182: Use according to embodiment 181, wherein the lysosomal storage disease having a defective degradation of polypeptides is selected from the group comprising pycnodysostosis.

Embodiment 183: Use according to embodiment 157, wherein the lysosomal storage disease is a lysosomal storage disease having a defective degradation or transport of cholesterol, cholesterol esters and/or other complex lipids.

Embodiment 184: Use according to embodiment 183, wherein the lysosomal storage disease having a defective degradation or transport of cholesterol, cholesterol esters and/or other complex lipids is selected from the group comprising Neuronal Ceroid Lipofuscinosis type I, Lipofuscinosis type II, Lipofuscinosis type III and Lipofuscinosis type IV.

Embodiment 185: Use according to embodiment 157, wherein the lysosomal storage disease is a lysosomal storage disease having multiple deficiencies of lysosomal enzymes.

Embodiment 186: Use according to embodiment 185, wherein the lysosomal storage disease having multiple deficiencies of lysosomal enzymes is selected from the group comprising Galactosialidosis, Danon Disease, Pycnodysostosis, multiple sulfatase deficiency, GM2 Gangliosidosis, Mucopolidosis type II and Mucopolidosis type III.

Embodiment 187: Use according to embodiment 186, wherein the lysosomal storage disease is Mucopolidosis type II or Mucopolidosis type III.

Embodiment 188: Use according to embodiment 157, wherein the lysosomal storage disease is a lysosomal storage disease having transport and trafficking defects.

Embodiment 189: Use according to embodiment 188, wherein the lysosomal storage disease having transport and trafficking defects is selected from the group comprising Cystinosis, Danon disease, Mucopolidosis type IV, Infantile sialic acid storage disease and Salla disease.

Embodiment 190: Use according to embodiment 157, wherein the lysosomal storage disease is Niemann -Pick type C1.

Embodiment 191: Use according to embodiment 157, wherein the lysosomal storage disease is San filipo syndrome type C.

Embodiment 192: Use according to embodiment 157, wherein the lysosomal storage disease is Danon disease.

Embodiment 193: Use according to embodiment 157, wherein the lysosomal storage disease is free sialic acid storage disease.

Embodiment 194: Use according to embodiment 157, wherein the lysosomal storage disease is Niemann -Pick type C2.

Embodiment 195: Use according to embodiment 156, wherein the disease is different from a lysosomal storage disease.

Embodiment 196: Use according to embodiment 195, wherein the disease different from a lysosomal storage disease is Parkinson disease.

Embodiment 197: Use of a compound having the ability to rearrange a lysosomal protein in or for the manufacture of a medicament for the treatment of a disease, wherein the medicament comprises a combination comprising a first constituent and a second constituent, wherein the first constituent is a or the compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, and wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 198: Use according to embodiment 197, wherein the combination is a combination according to any one of embodiments 1 to 155, preferably according to any one of embodiments 104 to 155.

Embodiment 199: Use of Ambroxol and/or a derivative of Ambroxol in or for the manufacture of a medicament for the treatment of a disease, wherein the medicament comprises a combination comprising a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity and wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 200: Use according to embodiment 199, wherein the combination is a combination according to any one of embodiments 1 to 155, preferably according to any one of embodiments 104 to 155.

Embodiment 201: Use according to any one of embodiments 197 to 200, wherein the disease is a lysosomal storage disease.

Embodiment 202: Use according to any one of embodiments 197 to 200, wherein the disease is different from a lysosomal storage disease.

Embodiment 203: Use according to embodiment 202, wherein the disease different from a lysosomal storage disease is Parkinson disease.

Embodiment 204: A pharmaceutical preparation comprising a first constituent, a second constituent optionally a further constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the second constituent is Ambroxol and/or a derivative of Ambroxol and wherein the further constituent is selected from the group comprising a pharmaceutically acceptable excipient and a pharmaceutically active agent, and wherein preferably the pharmaceutical preparation increases the reduced activity of the lysosomal protein, and wherein the lysosomal protein has a reduced activity.

Embodiment 205: The pharmaceutical preparation according to embodiment 204, wherein the lysosomal protein is affected in a disease.

Embodiment 206: The pharmaceutical preparation according to any one of embodiments 204 to 205, wherein the lysosomal protein is selected from the group comprising an enzyme, a trans-membrane protein and a soluble non-enzymatic protein.

Embodiment 207: The pharmaceutical preparation according to embodiment 206, wherein the lysosomal protein is an enzyme, wherein the enzyme has a reduced activity, preferably a reduced activity due to a mutation of the enzyme.

Embodiment 208: The pharmaceutical preparation according to embodiment 207, wherein the enzyme is a lysosomal hydrolase.

Embodiment 209: The pharmaceutical preparation according to any one of embodiments 204 to 208, wherein the compound having the ability to rearrange a lysosomal protein is a chaperon.

Embodiment 210: The pharmaceutical preparation according to embodiment 209, wherein the chaperon is a pharmacological chaperon or a pharmaceutically acceptable salt, solvate or derivative thereof.

Embodiment 211: The pharmaceutical preparation according to embodiment 210, wherein the pharmacological chaperon is a sugar and/or an imino sugar or a pharmaceutically acceptable salt, solvate or derivative of the sugar and/or the imino sugar.

Embodiment 212: The pharmaceutical preparation according to embodiment 211, wherein the sugar is galactose, preferably D-galactose.

Embodiment 213: The pharmaceutical preparation according to embodiment 211, wherein the imino sugar is selected from the group comprising 1-deoxygalactonojirimycin (DGJ), alpha-galacto-homonojirimycin, alpha-allo-homonojirimycin, beta-1-C-butyl-

deoxygalactonojirimycin, N-nonyl-deoxynojirimycin (NN-DNJ), N-octyl-2,5-anhydro-2,5-imino-D-glucitol, N-octyl-isofagomine, N-octyl-beta-valienamine (NOV), Isofagomine (IFG), calystegines A3 B1, B2, C1, 1,5-dideoxy-1,5-iminoxylitol (DIX), alpha-1-C-nonyl-DIX, alpha-1-C-octyl-1-DNJ, N-acetyl-glucosamine-thiazoline (NGT), 6-acetamido-6-deoxycastanospermine (ACAS), bisnaphthalimide nitro-indan-1-one, pyrrolo[3,4-d]pyridazin-1-one, pyrimethamine (PYR), N-actyl-4-epi-beta-valienamine (NOEV), N-butyl-DNJ, Deoxynojirimycin (DNJ), N-Acetyl-galactosamine (GalNAc), 2-Acetamido-1,2-dideoxynojirimycin (AdDNJ) and Castanospermine (CAS), and derivatives thereof; and pharmaceutically acceptable salts thereof.

Embodiment 214: The pharmaceutical preparation according to any one of embodiments 204 to 213, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 215: The pharmaceutical preparation according to any one of embodiments 204 to 214, wherein at least one of the first constituent and the second constituent is/are present as a solvate or a pharmaceutically acceptable salt thereof.

Embodiment 216: The pharmaceutical preparation according to embodiment 215, wherein the second constituent is present as a pharmaceutically acceptable salt thereof, wherein the second constituent is Ambroxol, wherein preferably the pharmaceutically acceptable salt of Ambroxol is Ambroxol hydrochloride.

Embodiment 217: The pharmaceutical preparation according to embodiment 215, wherein the second constituent is present as a pharmaceutically acceptable salt thereof, wherein the second constituent is a derivative of Ambroxol, wherein the derivative of Ambroxol is bromhexine, and wherein preferably the pharmaceutically acceptable salt of bromhexine is bromhexine hydrochloride.

Embodiment 218: The pharmaceutical preparation according to any one of embodiments 204 to 217, wherein the pharmaceutical preparation is suitable for or is for use in the treatment and/or prevention of a disease.

Embodiment 219: The pharmaceutical preparation according to any one of embodiments 204 to 218, wherein the first constituent and the second constituent are formulated into separate dosage forms, wherein preferably each dosage form is independently selected from the group comprising tablets, capsules, powder, mixture, effervescence tablets and solutions.

Embodiment 220: The pharmaceutical preparation according to any one of embodiments 204 to 219, wherein the first constituent and the second constituent are each and individually administered subcutaneously, intranasally, intravenously, sublingually, transmucosally and/or intracranially.

Embodiment 221: The pharmaceutical preparation according to any one of embodiments 204 to 220, wherein the pharmaceutical preparation is suitable for or is for use in a method for the treatment of a subject wherein the method comprises administering the pharmaceutical preparation to the subject.

Embodiment 222: The pharmaceutical preparation according to any one of embodiments 205 to 221, wherein the disease is a lysosomal storage disease.

Embodiment 223: The pharmaceutical preparation according to any one of embodiments 205 to 221, wherein the disease is different from a lysosomal storage disease.

Embodiment 224: The pharmaceutical preparation according to embodiment 223, wherein the disease different from a lysosomal storage disease is Parkinson disease.

Embodiment 225: A pharmaceutical preparation comprising a first constituent and a second constituent,
wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of embodiments 204 to 224;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is 1-deoxygalactonojirimycin or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 226: The pharmaceutical preparation according to embodiment 225, wherein the lysosomal protein is α -galactosidase A.

Embodiment 227: The pharmaceutical preparation according to any one of embodiments 225 to 226, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 228: The pharmaceutical preparation according to any one of embodiments 225 to 227, wherein the pharmaceutical preparation is for use in the treatment or prevention of Fabry's disease.

Embodiment 229: A pharmaceutical preparation comprising a first constituent and a second constituent,
wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of embodiments 204 to 224;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is alpha-galacto-homonojirimycin or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 230: The pharmaceutical preparation according to embodiment 229, wherein the lysosomal protein is α -galactosidase A.

Embodiment 231: The pharmaceutical preparation according to any one of embodiments 229 to 230, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 232: The pharmaceutical preparation according to any one of embodiments 229 to 231, wherein the pharmaceutical preparation is for use in the treatment or prevention of Fabry's disease.

Embodiment 233: A pharmaceutical preparation comprising a first constituent and a second constituent,
wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of embodiments 204 to 224;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is alpha-allo-homonojirimycin or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 234: The pharmaceutical preparation according to embodiment 233, wherein the lysosomal protein is α -galactosidase A.

Embodiment 235: The pharmaceutical preparation according to any one of embodiments 233 to 234, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 236: The pharmaceutical preparation according to any one of embodiments 233 to 235, wherein the pharmaceutical preparation is for use in the treatment or prevention of Fabry's disease.

Embodiment 237: A pharmaceutical preparation comprising a first constituent and a second constituent,
wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of embodiments 204 to 224;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is beta-1-C-butyl-deoxygalactonojirimycin or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 238: The pharmaceutical preparation according to embodiment 237, wherein the lysosomal protein is α -galactosidase A.

Embodiment 239: The pharmaceutical preparation according to any one of embodiments 237 to 238, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 240: The pharmaceutical preparation according to any one of embodiments 237 to 239, wherein the pharmaceutical preparation is for use in the treatment or prevention of Fabry's disease.

Embodiment 241: A pharmaceutical preparation comprising a first constituent and a second constituent,
wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of embodiments 204 to 224;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is galactose, preferably D-galactose or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 242: The pharmaceutical preparation according to embodiment 241, wherein the lysosomal protein is α -galactosidase A.

Embodiment 243: The pharmaceutical preparation according to any one of embodiments 241 to 242, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 244: The pharmaceutical preparation according to any one of embodiments 241 to 243, wherein the pharmaceutical preparation is for use in the treatment or prevention of Fabry's disease.

Embodiment 245: A pharmaceutical preparation comprising a first constituent and a second constituent,

wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of embodiments 204 to 224;

wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,

wherein the first constituent is N-acetyl-glucosamine-thiazoline (NGT) or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 246: The pharmaceutical preparation according to embodiment 245, wherein the lysosomal protein is beta-hexosaminidase A/B.

Embodiment 247: The pharmaceutical preparation according to any one of embodiments 245 to 246, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 248: The pharmaceutical preparation according to any one of embodiments 245 to 247, wherein the pharmaceutical preparation is for use in the treatment or prevention of Tay-Sachs or Sandhoff.

Embodiment 249: A pharmaceutical preparation comprising a first constituent and a second constituent,

wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of embodiments 204 to 224;

wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,

wherein the first constituent is 6-acetamido-6-deoxycastanospermine (ACAS) or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 250: The pharmaceutical preparation according to embodiment 249, wherein the lysosomal protein is beta-hexosaminidase A/B.

Embodiment 251: The pharmaceutical preparation according to any one of embodiments 249 to 250, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 252: The pharmaceutical preparation according to any one of embodiments 249 to 251, wherein the pharmaceutical preparation is for use in the treatment or prevention of Tay-Sachs or Sandhoff.

Embodiment 253: A pharmaceutical preparation comprising a first constituent and a second constituent,
wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of embodiments 204 to 224;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is bisnaphthalimide nitro-indan-1-one or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 254: The pharmaceutical preparation according to embodiment 253, wherein the lysosomal protein is beta-hexosaminidase A/B.

Embodiment 255: The pharmaceutical preparation according to any one of embodiments 253 to 254, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 256: The pharmaceutical preparation according to any one of embodiments 253 to 255, wherein the pharmaceutical preparation is for use in the treatment or prevention of Tay-Sachs or Sandhoff.

Embodiment 257: A pharmaceutical preparation comprising a first constituent and a second constituent,
wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of embodiments 204 to 224;

wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is pyrrolo[3,4-d]pyridazin-1-one or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 258: The pharmaceutical preparation according to embodiment 257, wherein the lysosomal protein is beta-hexosaminidase A/B.

Embodiment 259: The pharmaceutical preparation according to any one of embodiments 257 to 257, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 260: The pharmaceutical preparation according to any one of embodiments 257 to 259, wherein the pharmaceutical preparation is for use in the treatment or prevention of Tay-Sachs or Sandhoff.

Embodiment 261: A pharmaceutical preparation comprising a first constituent and a second constituent,
wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of embodiments 204 to 224;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is pyrimethamine (PYR) or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 262: The pharmaceutical preparation according to embodiment 261, wherein the lysosomal protein is beta-hexosaminidase A/B.

Embodiment 263: The pharmaceutical preparation according to any one of embodiments 261 to 262, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 264: The pharmaceutical preparation according to any one of embodiments 261 to 263, wherein the pharmaceutical preparation is for use in the treatment or prevention of Tay-Sachs or Sandhoff.

Embodiment 265: A pharmaceutical preparation comprising a first constituent and a second constituent,
wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of embodiments 204 to 224;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is N-octyl-4-epi-beta-valienamine (NOEV) or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 266: The pharmaceutical preparation according to embodiment 265, wherein the lysosomal protein is beta-galactosidase.

Embodiment 267: The pharmaceutical preparation according to any one of embodiments 265 to 266, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 268: The pharmaceutical preparation according to any one of embodiments 265 to 267, wherein the pharmaceutical preparation is for use in the treatment or prevention of GM1-Gangliosidosis.

Embodiment 269: A pharmaceutical preparation comprising a first constituent and a second constituent,
wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of embodiments 204 to 224;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is N-butyl-DNJ or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 270: The pharmaceutical preparation according to embodiment 269, wherein the lysosomal protein is alpha-glucosidase.

Embodiment 271: The pharmaceutical preparation according to any one of embodiments 269 to 270, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 272: The pharmaceutical preparation according to any one of embodiments 269 to 271, wherein the pharmaceutical preparation is for use in the treatment or prevention of Pompe's disease.

Embodiment 273: A pharmaceutical preparation comprising a first constituent and a second constituent,
wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of embodiments 204 to 224;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is Deoxynojirimycin or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.

Embodiment 274: The pharmaceutical preparation according to embodiment 273, wherein the lysosomal protein is alpha-glucosidase.

Embodiment 275: The pharmaceutical preparation according to any one of embodiments 273 to 274, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

Embodiment 276: The pharmaceutical preparation according to any one of embodiments 273 to 275, wherein the pharmaceutical preparation is for use in the treatment or prevention of Pompe's disease.

Embodiment 277: A method for preparing a pharmaceutical preparation, preferably a pharmaceutical preparation according to any one of embodiments 204 to 276, comprising the steps of

formulating a first constituent as defined in any one of the preceding embodiments and a second constituent as defined in any one of the preceding embodiments into a single dosage form or into two separate dosage forms, wherein in case of two separate dosage forms a first of the two separate dosage forms contains the first constituent and a second of the two separate dosage forms contains the second constituent.

Embodiment 278: Use of a pharmaceutical preparation for the manufacture of a medicament, wherein the pharmaceutical preparation is a pharmaceutical preparation according to any one of embodiments 204 to 276, and wherein the pharmaceutical preparation is for the treatment and/or prevention of a disease.

Embodiment 279: Use according to embodiment 278, wherein the disease is a lysosomal storage disease.

Embodiment 280: Use according to embodiment 278, wherein the disease is different from a lysosomal storage disease.

Embodiment 281: Use according to embodiment 280, wherein the disease different from a lysosomal storage disease is Parkinson disease.

Embodiment 282: A method for the treatment of a disease, wherein the method comprises administering to a subject a first constituent as defined in any one of embodiments 1 to 276 and, prior to, concomitantly with or after a second constituent as defined in any one of embodiments 1 to 276, and/or a combination according to any one of embodiments 1 to 155 and/or a pharmaceutical preparation according to any one of embodiments 204 to 276.

Embodiment 283: The method according to embodiment 282, wherein the disease is a lysosomal storage disease.

Embodiment 284: The method according to embodiment 282, wherein the disease is different from a lysosomal storage disease.

Embodiment 285: The method according to embodiment 284, wherein the disease different from a lysosomal storage disease is Parkinson disease.

Embodiment 286: A method for increasing activity of a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the method comprises administering to a cell a compound having the ability to rearrange the lysosomal protein, and Ambroxol and/or a derivative thereof.

Embodiment 287: A method for increasing activity of a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the method comprises administering to a subject a compound having the ability to rearrange the lysosomal protein, and Ambroxol and/or a derivative thereof, wherein the compound having the ability to rearrange the lysosomal protein increases the activity of the lysosomal protein.

Embodiment 288: A method for increasing activity of a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the method comprises administering to a subject a first constituent as defined in any one of embodiments 1 to 276 and, prior to, concomitantly with or after a second constituent as defined in any one of embodiments 1 to 276, and/or a combination according to any one of embodiments 1 to 155 and/or a pharmaceutical preparation according to any one of embodiments 204 to 276, wherein the combination and/or the pharmaceutical preparation increases the activity of the lysosomal protein.

Embodiment 289: A combination comprising a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity wherein the reduced activity is reduced due to a mutation of the lysosomal protein, and wherein the second constituent is Ambroxol and/or a derivative of Ambroxol, wherein the combination is preferably a combination according to any one of embodiments 1 to 155,

for use in a method of personalized therapeutic treatment of a subject, wherein the method comprises the following steps:

step a): determining whether in a sample of the subject the lysosomal protein has a reduced activity, preferably such reduced activity results from one or more mutation of the lysosomal protein compared to the wild type lysosomal protein;

step b): identifying a compound having the ability to rearrange the lysosomal protein having reduced activity, and wherein the compound is suitable for or is increasing the reduced activity of the lysosomal protein; and

step c): administering to the subject the first constituent prior to, concomitantly with or after the second constituent.

Embodiment 290: A pharmaceutical preparation comprising a first constituent, a second constituent and optionally a further constituent,

wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the reduced activity is reduced due to a mutation of the lysosomal protein, and

wherein the second constituent is Ambroxol and/or a derivative of Ambroxol,

wherein the further constituent is selected from the group comprising pharmaceutically acceptable excipients and pharmaceutically active agents,

wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of embodiments 204 to 276,

for use in a method of personalized therapeutic treatment of a subject, wherein the method comprises the following steps:

step a): determining whether in a sample of the subject the lysosomal protein has a reduced activity, preferably such reduced activity results from one or more mutation of the lysosomal protein compared to the wild type lysosomal protein;

step b): identifying a compound having the ability to rearrange the lysosomal protein having reduced activity, and wherein the compound is suitable for or is increasing the reduced activity of the lysosomal protein; and

step c): administering to the subject the first constituent prior to, concomitantly with or after the second constituent.

The present inventors have surprisingly found that the combined use of a compound having the ability to rearrange a lysosomal protein having reduced activity and Ambroxol or a derivative of Ambroxol is suitable for the treatment of LSDs such as Fabry disease and Pompe disease. In such combination, preferably the compound having the ability to rearrange a lysosomal protein having reduced activity is a chaperone, preferably a pharmacological chaperone.

More specifically, the present inventors when conducting experiments in a cell culture system which is an acknowledged model for LSDs and Fabry disease in particular (Ishii et al.; *Biochem. J.* (2007) 406, 285–295; Shin et al.; *Pharmacogenet Genomics*. Sep 2008; 18(9): 773–780; Parenti G et al., *Mol Ther.* 2007; 15:508-14; Okumiya T et al., *Mol Genet Metab* 2007 Jan;90(1):49-57) have found that when DGJ of Galactose is administered together with Ambroxol the activity of mutant alpha-galactosidase A is surprisingly and unexpectedly increased compared to the effect arising from administering DGJ or Galactose only.

When conducting further studies the present inventors also found that when NB-DNJ is administered together with Ambroxol the activity of mutant acid α -Glucosidase is surprisingly and unexpectedly increased compared to the effect arising from administering NB-DNJ only.

In connection therewith it is particularly noteworthy that, as also found by the instant inventors, administering Ambroxol alone, i.e. without DGJ, does not exhibit any enhancing effect on alpha-galactosidase A activity in alpha-galactosidase A mutants frequently resulting in Fabry disease, and that administering Ambroxol alone, i.e. without NB-DNJ does not exhibit any enhancing effect on acid α -Glucosidase activity in acid α -Glucosidase mutants frequently resulting in Pompe disease.

It is the merit of the present inventors having found that administering a combination according to the present invention is advantageous over the enhancement of the activity of the lysosomal protein having a reduced activity with a compound having the ability to rearrange a lysosomal protein alone. More particularly, administering Ambroxol and/or a derivative of Ambroxol in combination with said compound having the ability to rearrange a lysosomal protein further enhances the activity of the lysosomal protein having reduced protein activity compared to administering said compound having the ability to rearrange a lysosomal protein alone. Thus administration of a combination of the present invention is advantageous over the administration of the compound having the ability to rearrange a lysosomal protein alone in that further enhancement of the activity of the protein having reduced activity is achieved and/or a significantly lower amount of the compound having the ability to rearrange a lysosomal protein may be administered to a patient still resulting in the same or similar beneficial effect of treatment as if the compound having the ability to rearrange a lysosomal proteins is administered alone at a higher concentration or in a higher amount.

More particularly, it is the merit of the present inventors having found that the combined use of a pharmacological chaperone and Ambroxol and/or a derivative of Ambroxol is suitable for increasing the activity of a mutant protein in the treatment of an LSD such as Fabry disease compared to the application of said pharmacological chaperone alone. The amount of pharmacological chaperone administered to a subject can be reduced if the pharmacological chaperone is administered as the first constituent of the combination according to the present invention and will result in the same or similar level of activity or a higher level of activity of the lysosomal protein as if the pharmacological chaperone is administered alone in a higher amount.

Furthermore, it will be acknowledged by a person skilled in the art that any feature, advantage, embodiment of and any statement made herein in relation to a combination of the present invention equally applies to a pharmaceutical preparation according to the present invention, a method according to the present invention, a use according to the present invention, and vice versa.

Furthermore, it will be acknowledged by a person skilled in the art that any feature, advantage, embodiment of and any statement made herein in relation to any aspect of the present invention equally applies to each and any of the other aspects of the present invention.

The term “reduced activity” of a lysosomal protein as used herein preferably means that the activity of a lysosomal protein, e.g. a lysosomal protein being a lysosomal protein mutant resulting from a mutation, is reduced compared to a control, such as a control protein, for example said protein without such mutation. In an embodiment such reduced activity is the activity of an enzyme which is reduced due to a mutation of said enzyme compared to an activity of said enzyme without such mutation and wherein, preferably, said reduced enzyme activity results in a disease, e.g. an LSD, such as Fabry’s disease.

It is a further advantage of the combination according to the present invention that the activity of a lysosomal protein having a reduced activity, wherein treatment with a pharmacological chaperone alone does not result in an increase of activity of the lysosomal protein mutant sufficient to treat a subject, can be elevated to a level sufficient for the treatment, preferably therapeutic treatment of the subject. In other words, in particular lysosomal protein mutants wherein the treatment with a particular pharmacological chaperone results in an increase of activity of the lysosomal protein mutants the increase is too low to treat the subject. If the pharmacological chaperone is administered as a first constituent of the combination according to the present invention the increase of activity of the lysosomal protein mutant can be sufficient to treat the subject.

A sufficient treatment as used herein preferably is a treatment which results in elevating and/or increasing the activity of a lysosomal protein having a reduced activity due to a mutation, i.e. a lysosomal protein mutant, to a level of 20%, preferably of 25%, of the activity of the lysosomal protein not having the mutation and preferably determined in the absence of treatment. In an embodiment, such sufficient treatment results in a therapeutic effect including, but not limited to, treatment of the LSD from which the subject suffers or at least amelioration of the symptoms of the subject suffering from the LSD.

It will be immediately acknowledged by a skilled person that various methods for determining the activity of a protein, such as the activity of an enzyme, and thus determining whether said

activity is reduced compared to a control such as a protein without mutation(s) which result(s) in such reduced activity, are known to a person skilled in the art and particularly depend on the protein the activity of which is determined and, preferably, compared to a control. For example, determining the activity of α -galactosidase A and/or α -glucosidase is described as enzymatic measurement of α -galactosidase A and α -glucosidase in Example 1 herein, respectively. In order to determine the activity of β -Hexosaminidase A/B and/or whether said activity is reduced preferably the assay as described in Wood et al. is used (Wood, S and Macdougall, BG Am J Hum Genet 28:489-495, 1976). In order to determine the activity of beta-galactosidase and/or whether said activity is reduced preferably the assay as described in Yang et al. is used (Yang et al., Journal of Biomedical Science 2010, 17:79).

It will be acknowledged by a person skilled in the art that in certain diseases resulting from reduced activity of a lysosomal protein typically resulting from a mutation of said lysosomal protein, the administration of Ambroxol alone results in an enhancement of the activity of a/the mutant lysosomal protein. It is thus important to understand that the present invention is based on the finding that the effect of a compound having the ability to rearrange a protein having reduced activity, such as a pharmacological chaperone, is enhanced when administered together with Ambroxol and/or a derivative of Ambroxol, wherein the administration of Ambroxol and/or a derivative of Ambroxol is not enhancing the activity of the protein having reduced activity when administered alone. The administration of said combination is thus advantageous over the administration of said compound having the ability to rearrange a protein having reduced activity alone in that Ambroxol enhances the activity of said compound.

In connection therewith it will be acknowledged by a skilled person that any derivative of Ambroxol and/or bromhexine enhancing the activity of a/the protein having reduced activity when administered in combination with said compound having the ability to rearrange a/the protein having reduced activity is suitable for the practicing of the instant invention and is, accordingly, encompassed by the present invention.

A compound having the ability to rearrange a lysosomal protein having reduced activity as used herein preferably means a small molecule which reversibly binds to the lysosomal protein having reduced activity, said binding resulting in an increase in the reduced activity,

i.e. said binding resulting in an activity which is higher compared to the activity of the lysosomal protein without the compound having the ability to rearrange a/the lysosomal protein having reduced activity. Preferably, said binding corrects a folding defect of the protein having reduced activity through a stabilization effect. More preferably, said binding prevents degradation of said protein. A small molecule as used herein preferably is a small molecule according to Lipinski's rule of five. In an embodiment of the combination according to the present invention the compound having the ability to rearrange a lysosomal protein having reduced activity preferably is a chaperone and more preferably is a pharmacological chaperone. In a preferred embodiment of the combination according to the present invention a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, is capable of increasing and/or elevating the reduced activity. In a further embodiment a compound having the ability to rearrange a lysosomal protein is a compound which is capable of increasing and/or elevating the reduced activity irrespective of whether or not said compound has the ability to rearrange a lysosomal protein, i.e. the activity of the lysosomal protein is higher when the compound is present compared to the activity of the lysosomal protein when the compound is not present.

A person skilled in the art will acknowledge that a pharmacological chaperon, such as an iminosugar, e.g. DGJ or DNJ, interacts with a specific enzyme and/or a specific enzyme having a mutation. The use of pharmacological chaperones in the treatment of LSDs is summarized in, e.g., Parenti et al., (Parenti et al., 2009 EMBO Mol Med 1, 268–279).

For example, NB-DNJ but not NB-DGJ effects folding of beta-Glucocerebrosidase (Butters TD et al., Glycobiology 2005, 15:43R-52R). However, sphingolipid synthesis can be inhibited by both drugs. Furthermore, DNJ is often used as negative control for testing putative drugs and it has no effect on enhancing protein activity in Fabry (Fan et al. 2003, Methods in Enzymology, Vol. 363, p. 412-420) or Gaucher. Interestingly, Parenti et al. showed an effect of DNJ in enhancing protein activity in Pompe's disease (Parenti et al. 2007 Mol Ther. 2007 Mar;15(3):508-14. Epub 2007 Jan 9).

Further examples for the specific interaction of pharmacological chaperons are described in Butters et al. (2005, supra) and Parenti et al. (2009, supra).

The present inventors assume, without wishing to be bound by any theory, that the specific interaction and non-interaction, respectively, of pharmacological chaperons with lysosomal protein mutants is due to a stereochemical hindrance, for example with L-sugars or with DNJ, which prevents access to the active site of the lysosomal protein. Accordingly, rearrangement of the lysosomal protein mutant is or is not possible and thus enhancement of transport and/or activity is or is not prevented. The instant inventors further assume that due to interaction of the pharmacological chaperone with the active site of the lysosomal protein a rearrangement of the lysosomal protein is induced leading to a stabilization of the lysosomal protein mutant, similar to wild type. Thus the mutant lysosomal protein being no longer misfolded or less misfolded is no longer recognized by the cellular degradation machinery and accordingly and preferably escapes early degradation. As a result transport into the lysosome is enhanced. Applying different assays and methods Yam et al. (Yam et al., *supra*) and Ishii et al. (Ishii et al., *supra*) both showed accumulation of the enzyme after treatment in the lysosome in Fabry.

In addition, it is important to understand that a disease which results from a lysosomal protein having a reduced activity and wherein the reduced activity is due to a mutation of the lysosomal protein may actually be the result of different mutations, i.e. different subjects may suffer from the same disease having different mutations of the lysosomal proteins. By way of example, more than 400 mutations which result in Fabry's disease are known to date. In connection therewith a person skilled in the art will immediately acknowledge that for successful treatment of such disease the reduced activity of the lysosomal protein having reduced activity has to be increased. More particularly, the particular lysosomal protein mutant has to be a mutant responsive for the treatment with a compound having the ability to rearrange a lysosomal protein having a reduced activity, such as a pharmacological chaperone. A "responsive mutant" as used herein preferably means a lysosomal protein having a particular mutation, wherein the lysosomal protein has a reduced activity due to the mutation and wherein the compound having the ability to rearrange a lysosomal protein having a reduced activity is suitable for or is increasing and/or enhancing the reduced activity so that the activity of the lysosomal protein is increased.

In an embodiment of the combination and/or pharmaceutical preparation according to the present invention the activity of the lysosomal protein is reduced due to a mutation. In a further embodiment of the present invention the mutation is a mutation responsive to a/the

compound having the ability to rearrange the lysosomal protein, wherein the lysosomal protein has a reduced activity. In an embodiment of the combination and/or the pharmaceutical preparation according to the present invention the compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, is suitable for and/or is capable of increasing the activity of the lysosomal protein.

A person skilled in the art will thus also acknowledge that a compound having the ability to rearrange a protein having reduced activity may be specifically effective in enhancing the activity of a specific protein in a specific disease. In other words a certain compound having the ability to rearrange a protein having reduced activity, such as a certain pharmacological chaperone, may be administered in a certain disease as said pharmacological chaperone enhances the activity of a certain protein having reduced activity in said certain disease.

In an embodiment of the compound according to the present invention a compound having the ability to rearrange a protein having reduced activity is an active site-specific chaperone.

The following table 1 is taken from Fan et al. (Fan JQ et al., Biol Chem. 2008 Jan;389(1):1-11.) and comprises a list of exemplary active site-specific chaperones, also referred to herein as ASSC, and the respective LSD, wherein the activity of the respective affected protein having reduced activity may be enhanced upon treatment with the indicated ASSC.

Table 1: Compounds having the ability to rearrange a protein having reduced activity useful in treatment of the respective LSD by enhancing the activity of the respective protein having reduced activity. Additionally, table 1 shows competitive inhibitors of enzymes which result in an LSD if mutated.

Disorder	Deficient enzyme	Competitive inhibitors	References
Gaucher disease	Acid β -glucosidase or glucocerebrosidase	isofagomine N-dodecyl-DNJ calystegines A ₁ , B ₁ , B ₂ , C ₁ 6-nonyl-isofagomine N-octyl- β -valienamine (NOV)	Ogawa et al., 1998; Zhu et al., 2005; Chang et al., 2006
Fabry disease	α -Galactosidase A	1-deoxygalactonojirimycin α -allo-homonojirimycin α -galacto-homonojirimycin β -1-C-butyl-deoxynojirimycin calystegines A ₁ , B ₂ N-methyl calystegines A ₁ , B ₂	Fan et al., 1999; Asano et al., 2000
G _{M1} -gangliosidosis	Acid β -galactosidase	4-epi-isofagomine 1-deoxygalactonojirimycin	Fan and Ishii, 2003b
Krabbe disease	Galactocerebrosidase	4-epi-isofagomine 1-deoxygalactonojirimycin	Asano et al., 1994; Ichikawa et al., 1998
Pompe disease	α -glucosidase	1-deoxynojirimycin (DNJ) α -homonojirimycin castanospermine	Asano et al., 1994; Pili et al., 1995; Asano et al., 1997
Morquio disease B	Acid β -galactosidase	4-epi-isofagomine 1-deoxygalactonojirimycin	Asano et al., 1994; Ichikawa et al., 1998
α -Mannosidosis	Acid α -mannosidase	1-deoxymannonojirimycin Swainsonine Mannostatin A	Dorling et al., 1980; Aoyagi et al., 1989; Ichikawa et al., 1998
β -Mannosidosis	Acid β -mannosidase	2-hydroxy-isofagomine	Ichikawa et al., 1998
Fucosidosis	Acid α -L-fucosidase	1-deoxyfucono-jirimycin β -homofucono-jirimycin 2,5-imino-1,2,5-trideoxy-L-glucitol 2,5-dideoxy-2,5-imino-D-fucitol 2,5-imino-1,2,5-trideoxy-O-altritol	Asano et al., 2001
Sanfilippo disease B	α -N-Acetylglucosaminidase	1,2-dideoxy-2-N-acetamido-nojirimycin	Ichikawa et al., 1998
Schindler-Kanzaki disease	α -N-Acetylgalactosaminidase	1,2-dideoxy-2-N-acetamido-galactonojirimycin	Ichikawa et al., 1998
Tay-Sachs disease	β -Hexosaminidase A	2-N-acetylamino-isofagomine 1,2-dideoxy-2-acetamido-nojirimycin nagstain and its derivatives	Aoyagi et al., 1992; Tatsuta et al., 1995; Ichikawa et al., 1998; Asano et al., 2001
Sandhoff disease	β -Hexosaminidase B	2-N-acetamido-isofagomine 1,2-dideoxy-2-acetamido-nojirimycin nagstain and its derivatives	Aoyagi et al., 1992; Tatsuta et al., 1995; Ichikawa et al., 1998; Asano et al., 2001
Hurler-Scheie disease	α -L-Iduronidase	1-deoxyiduronojirimycin 2-carboxy-3,4,5-trideoxypiperidine	Cenci di Bello et al., 1984; Ichikawa et al., 1998
Sly disease	β -Glucuronidase	6-carboxy-isofagomine 2-carboxy-3,4,5-trideoxypiperidine	Cenci di Bello et al., 1984; Ichikawa et al., 1998
Sialidosis	Sialidase	2,6-dideoxy-2,6-imino-sialic acid Siestatin B	Umezawa et al., 1974; Ichikawa et al., 1998

Additionally, Sawkar et al. (Sawkar, A. R. et al., Proc. Natl. Acad. Sci. U. S. A. 2002, 99, 15428-15433) showed that NB-DNJ and NN-DNJ can function as pharmacological chaperones in Gaucher's disease.

A person skilled in the art will acknowledge that in an embodiment an LSD is characterized by a lysosomal dysfunction which results from genetic mutation(s) usually leading to a reduced activity, including deficiency of protein activity, of a protein such as an enzyme, a membrane transport protein, a trans-membrane protein or a soluble non-enzymatic protein preferably required for or involved in the metabolism of lipids, glycoproteins or mucopolysaccharides. Said protein having reduced activity preferably is a lysosomal protein and the reduced activity thereof more preferably results from misfolding of the protein. Correcting said misfolding of said protein by rearrangement which comprises stabilization of said lysosomal protein having reduced activity, results in an enhancement of said reduced protein's activity and is, in an embodiment of the invention, thus suitable for the treatment of a disease, preferably an LSD.

In molecular biology, chaperones are proteins that assist folding or unfolding and assembly or disassembly of other macromolecular structures, such as proteins, but do not occur in these structures when the latter are performing their normal biological functions. Nevertheless, chaperones are not strictly concerned with protein folding. Moreover, a chaperon may assist in the assembly of, e.g., nucleosomes from folded histones and DNA, and such assembly chaperones, especially in the nucleus, are concerned with the assembly of folded subunits into oligomeric structures.

Chaperones as preferably used herein do not necessarily convey steric information required for proteins to fold. One major function of chaperones is to prevent both newly synthesized polypeptide chains and assembled subunits from aggregating into nonfunctional structures. It is for this reason that many chaperones, but by no means all, are also heat shock proteins because the tendency to aggregate increases as proteins are denatured by stress. Many chaperones are heat shock proteins, that is, proteins expressed in response to elevated temperatures or other cellular stresses. The reason for this behavior is that protein folding is severely affected by heat and, therefore, some chaperones act to repair the potential damage caused by misfolding.

The term “affected” as used herein preferably means “to change” or “being changed” and more preferably refers to a changed behavior, structure, activity or condition. For example, an affected protein may be affected in such that its activity is changed, i.e. increased or decreased, or as a further example an affected protein may be affected in such that its structure or sequence is changed compared to the protein’s structure or sequence in an unaffected stage. The term “to affect” refers to herein preferably to an influence which a change has on something else. Thus said protein being affected in a disease means that said changed activity and/or structure or sequence is changed compared to the activity and/or structure or sequence of said protein in a healthy patient. In a preferred embodiment thereof, the change goes along with a change in the condition of a subject the metabolism of lysosomal protein of which is affected; more preferably such change goes along with a disease or a predisposition of a disease.

Other chaperones are involved in folding newly made proteins as they are extruded from the ribosome. Although most newly synthesized proteins can fold in the absence of chaperones, a minority strictly requires them.

Other types of chaperones are involved in transport across membranes, for example membranes of the mitochondria and endoplasmic reticulum (ER) in eukaryotes. Bacterial translocation-specific chaperones maintain newly synthesized precursor polypeptide chains in a translocation-competent (generally unfolded) state and guide them to the translocon.

Molecular chaperones as used herein are preferably involved in protein folding and provide other functions in addition to stabilization. Molecular chaperones recognize unfolded protein elements such as exposed hydrophobic patches and bind to unfolded proteins and prevent aggregation or provide stabilization during folding, after which the folded protein dissociates. Molecular chaperones also direct proteins that do not fold correctly to degradation pathways.

Chemical chaperones, as used herein, preferably induce stabilizing effects by non-specific binding. Chemical chaperones include but are not limited to glycerol.

Chaperon functions also comprise among others assistance in protein degradation, bacterial adhesion activity and responding to diseases linked to protein aggregation.

Protein misfolding is recognized as an important pathophysiological cause of protein deficiency or reduced activity in many genetic disorders, especially in LSDs. Inherited mutations can disrupt native protein folding, thereby producing proteins with misfolded conformations. These misfolded proteins are consequently retained and degraded by endoplasmic reticulum-associated degradation, although they would otherwise be catalytically fully or partially active. Lack of protein function or reduced activity of lysosomal proteins results in substrate accumulation, impaired transport of ions or molecules, and disruption of biosynthetic pathways.

Recently, it was found that potent competitive inhibitors for enzymes associated with LSDs enhance the activity of such enzymes in cells when administered at concentrations lower than that normally required to inhibit the intracellular enzyme activity. The effect is particularly significant on certain defective or mutant enzymes, but also occurs in cells containing the normal enzyme type, preferably the non-defective enzyme and/or the non-mutant enzyme. The use of such pharmacological chaperones, such as DGJ, in PCT of an LSD preferably results from stabilizing the lysosomal mutant protein having reduced activity by the selective binding of the pharmacological chaperone to said mutant protein in its misfolded state. More preferably, said binding results in a more stable state of the protein and increased stability leads to decrease in degradation by endoplasmic reticulum-associated degradation. A pharmacological chaperone is preferably an active site-directed competitive inhibitor or a specific-site chaperone. Such active site-specific chaperones preferably act as reversible inhibitors that efficiently bind the mutant protein in the endoplasmic reticulum and facilitate folding and/or stabilization of the mutant protein resulting in more protein being processed and transported to the correct location, for example the lysosome. The pharmacological chaperone thereby preferably acts as a reversible binder and dissociates in the correct location where a higher level of residual protein activity is achieved. Nevertheless, a person skilled in the art will acknowledge that high concentrations of pharmacological chaperones will result in an inhibitory effect on the activity of the mutant protein (Asano et al., *supra*; Khanna et al., *supra*).

Accordingly, pharmacological chaperones are preferably applied in a sub-inhibitory concentration, wherein the stabilization of the mutant protein after rescue from the endoplasmatic reticulum preferably also results in a longer half-life of the mutant protein. Specific-site chaperones in contrast to active site chaperones do not bind the active site of the mutant protein. Therefore specific-site chaperones may remain bound to the mutant protein as long as the function of said protein is not inhibited.

Pharmacological chaperones used in the combination of the present invention preferably have at least one pharmaceutical characteristics selected from the group comprising low toxicity, good solubility and brain-barrier penetration.

Pharmacological chaperones as used herein are preferably selected from the group comprising sugars and iminosugar and derivatives of the sugars and iminosugar as well as pharmaceutical acceptable salts thereof.

The following table 2 is taken from Toprak et al. (Toprak et al., 2004, The Journal of biological chemistry, Vol. 279, No. 14, Issue of April 2, pp. 13478–13487, 2004) and shows illustrative but non-limiting examples of imino sugars useful as pharmacological chaperones in the combination according to the present invention, preferably wherein the lysosomal protein is hexosaminidase A.

Table 2: Imino sugars – pharmacological chaperones

Structure	Name	Ki / IC ₅₀
	<i>N</i> -Acetyl-galactosamine (GalNAc)	Ki = 1.9 mM ¹
	Deoxynojirimycin (DNJ)	N/A
	2-Acetamido-1,2-dideoxynojirimycin (AdDNJ)	Ki = 700 nM ²
	2-Acetamido-2-deoxynojirimycin (ADNJ)	Ki = 5 nM ²
	Castanospermine (CAS)	N/A
	6-Acetamido-6-deoxycastanospermine (ACAS)	IC ₅₀ = 500 nM ¹
	<i>N</i> -Acetyl-glucosamine-thiazoline (NGT)	Ki = 280 nM ² Ki = 300 nM ¹

Pharmacological chaperones as used herein are preferably selected from the group comprising 1-deoxygalactonojirimycin (DGJ), alpha-galacto-homonojirimycin, alpha-allo-homonojirimycin, beta-1-C-butyl-deoxygalactonojirimycin, beta-1-C-butyl-deoxynojirimycin, N-nonyl-deoxynojirimycin (NN-DNJ), N-octyl-2,5-anhydro-2,5-imino-D-glucitol, N-octyl-isofagomine, N-octyl-beta-valienamine (NOV), Isfagomine (IFG), calystegine A3, calystegine B1, calystegine B2, calystegine C1, 1,5-dideoxy-1,5-iminoxylitol (DIX), alpha-1-

C-nonyl-DIX, alpha-1-C-octyl-1-DNJ, N-acetyl-glucosamine-thiazoline (NGT), 6-acetamido-6-deoxycastanospermine (ACAS), bisnaphthalimide nitro-indan-1-one, pyrrolo[3,4-d]pyridazin-1-one, pyrimethamine (PYR), N-actyl-4-epi-beta-valienamine (NOEV), N-butyl-DNJ, Deoxynojirimycin (DNJ), N-Acetyl-galactosamine (GalNAc), 2-Acetamido-1,2-dideoxynojirimycin (AdDNJ), N-dodecyl-DNJ, 6-nonyl-isofagomine, N-methyl calystegine A3, calystegine B2, 4-epi-isofagomine, 1-deoxynojirimycin, alpha-homonojirimycin, castanospermine, 1-deoxymannojirimycin, Swainsonine, Mannostatin A, 2-hydroxy-isofagomine, 1-deoxyfuconojirimycin, beta-homofuconojirimycin, 2,5-imino-1,2,5-trideoxy-L-glucitol, 2,5-dideoxy-2,5-imino-D-fucitol, 2,5-imino-1,2,5-trideoxy-D-altritol, 1,2-dideoxy-2-N-acetamido-nojirimycin, 1,2-dideoxy-2-N-acetamido-galaconojirimycin, 2-N-acetylamino-isofagomine, 1,2-dideoxy-2-acetamido-nojirimycin, nagastain, 2-N-acetamido-isofagomine, 1,2-dideoxy-2-acetamido-nojirimycin, 1-deoxyduronojirimycin, 2-carboxy-3,4,5-trideoxypiperidine, 6-carboxy-isofagomine, 2,6-dideoxy-2,6-imino-sialic acid, Siastin B and Castanospermine (CAS), and derivatives thereof; and pharmaceutically acceptable salts thereof.

A person skilled in the art will know other pharmacological chaperones which may be applied for pharmacological chaperone therapy of LSDs in the combination according to the present invention.

Preferably a pharmacological chaperone has the ability to rearrange a lysosomal protein having reduced activity, said reduced activity preferably resulting from a mutation and being the or a cause of an LSD.

In an embodiment of the combination according to the present invention, the combination comprises a pharmaceutical acceptable salt of a pharmacological chaperon. In an embodiment of the combination according to the present invention the pharmaceutically acceptable salt is selected from the group comprising an organic acid and an inorganic salt. In an embodiment of the combination according to the present invention an organic acid is selected from the group comprising citrate, acetate, benzoate and tartrate, preferably isofagomine tartrate. In an embodiment of the combination according to the present invention an inorganic salt is selected from the group comprising hydrochloride and hydrobromide, i.e. HCl and HBr, as for example DGJ-HCl.

A person skilled in the art will acknowledge that the symptoms of LSDs vary, depending on the particular disorder or disease, and other variables like the age of onset, and can be mild to severe. They can include developmental delay, movement disorders, seizures, dementia, deafness and/or blindness. Some people with an LSD have enlarged livers (hepatomegaly) and enlarged spleens (splenomegaly), pulmonary and cardiac problems, and bones that grow abnormally.

For the diagnosis of an LSD the majority of patients are initially screened by an enzyme assay, which is the most efficient method to arrive at a definitive diagnosis. In other cases, for example, in some families where the disease-causing mutation(s) is/are known and in certain genetic isolates, mutation analysis may be performed. In addition, after a diagnosis is made by biochemical means, mutation analysis may be performed for certain disorders.

Representative, not-limiting examples of LSDs are described in the following.

Fabry disease, also referred to herein as Fabry's disease, Anderson-Fabry disease, angiokeratoma corporis diffusum and alpha-galactosidase A deficiency, is a rare X-linked inherited LSD, which can cause a wide range of systemic symptoms (James, William D.; Berger, Timothy G.; et al. (2006). *Andrews' Diseases of the Skin: clinical Dermatology*. Saunders Elsevier. p. 538, ISBN0-7216-2921-0). A deficiency of the enzyme alpha galactosidase A, also referred to herein as a-GAL A, GAL, GLA or alpha-Gal A, due to mutation causes a glycolipid known as globotriaosylceramide, also referred to herein as Gb3, GL-3, or ceramide trihexoside to accumulate within the blood vessels, other tissues, and organs (Karen JK et al., 2005 Dermatol. Online J. 11 (4): 8). This accumulation leads to an impairment of their proper functioning. The DNA mutations which cause the disease are X-linked. The condition affects hemizygous males, as well as homozygous, and potentially heterozygous females, so called carriers. Whilst males typically experience severe symptoms, women can range from being asymptomatic to having severe symptoms. This variability is thought to be due to X-inactivation patterns during embryonic development of the female (James, William D. supra).

Symptoms are typically first experienced in early childhood and can be very difficult to understand; the rarity of Fabry disease to many clinicians sometimes leads to misdiagnoses. Manifestations of the disease usually increase in number and severity while at individual ages. Kidney complications are a common and serious effect of the disease; renal insufficiency and renal failure may worsen throughout life. Proteinuria which causes foamy urine, is often the first sign of kidney involvement. End stage renal failure in males can typically occur in the third decade of life, and is a common cause of death due to the disease.

Fabry disease is indicated when associated symptoms are present, and can be diagnosed by a blood test to measure the level of alpha-galactosidase activity. Chromosomal analysis of the alpha-galactosidase A gene is the most accurate method of diagnosis, and many mutations which cause the disease have been noted.

Naturally, a-Gal A is likely to be present only at very low levels in the blood, particularly in males. In females, owing to X-inactivation patterns, levels are commonly normal even if the patient is not asymptomatic.

Glycogen storage disease type II, also referred to herein as Pompe disease, Pompe's disease or acid maltase deficiency is an autosomal recessive metabolic disorder which damages muscle and nerve cells throughout the body. It is the only glycogen storage disease with a defect in lysosomal metabolism.

Pompe disease is caused by an accumulation of glycogen in the lysosome due to deficiency of the lysosomal acid alpha-glucosidase enzyme, which is a lysosomal hydrolase. The enzyme degrades alpha -1,4 and alpha -1,6 linkages in glycogen, maltose and isomaltose and is required for the degradation of 1-3% of cellular glycogen. The deficiency of this enzyme results in the accumulation of structurally normal glycogen in lysosomes and cytoplasm in affected individuals. The build-up of glycogen causes progressive muscle weakness (myopathy) throughout the body and affects various body tissues, particularly in the heart, skeletal muscles, liver and nervous system. Excessive glycogen storage within lysosomes may interrupt normal functioning of other organelles and lead to cellular injury.

The enzyme deficiency is caused by a mutation in a gene, namely acid alpha-glucosidase, also referred to herein as acid maltase, on long arm of chromosome 17 at 17q25.2-q25.3 (base pairs 75,689,876 to 75,708,272). The number of mutations described in 2010 was 289 with 67 being non-pathogenic mutations and 197 being pathogenic mutations. The remainder is still being evaluated for their association with disease. Most cases appear to be due to three mutations. A transversion (T → G) mutation is the most common among adults with this disorder. This mutation interrupts a splicing site.

Pompe disease associated symptoms include severe lack of muscle tone, weakness, an enlarged liver (hepatomegaly), and an enlarged heart (cardiomegaly). Mental function is not affected. Development appears normal for the first weeks or months but slowly declines as the disease progresses. Swallowing may become difficult and the tongue may protrude and become enlarged. Most children die from respiratory or cardiac complications before 2 years of age.

Juvenile onset symptoms appear in early to late childhood and include progressive weakness of respiratory muscles in the trunk, diaphragm and lower limbs, as well as exercise intolerance. Intelligence is normal.

Adult onset symptoms also involve generalized muscle weakness and wasting of respiratory muscles in the trunk, lower limbs, and diaphragm. Many patients report respiratory distress, headache at night or upon waking, diminished deep tendon reflexes, and proximal muscle weakness, such as difficulty in climbing stairs. Intellect is not affected. A small number of adult patients live without major symptoms or limitations.

Pompe's disease is one of the infiltrative causes of restrictive cardiomyopathy. The infantile form usually comes to medical attention within the first few months of life. The usual presenting features are cardiomegaly (92%), hypotonia (88%), cardiomyopathy (88%), respiratory distress (78%), muscle weakness (63%), feeding difficulties (57%) and failure to thrive (53%). The main clinical findings include floppy baby appearance, delayed motor milestones and feeding difficulties. Moderate hepatomegaly may be present. Facial features include macroglossia, open mouth, wide open eyes, nasal flaring (due to respiratory distress), and poor facial muscle tone. Cardiopulmonary involvement is manifest by increased

respiratory rate, use of accessory muscles for respiration, recurrent chest infections, decreased air entry in the left lower zone (due to cardiomegaly), arrhythmias and evidence of heart failure. Median age at death in untreated cases is 8.7 months and is usually due to cardiorespiratory failure.

The usual initial investigations of this form of the disease include chest X ray, electrocardiogram and echocardiography. Typical findings are those of an enlarged heart with nonspecific conduction defects. Biochemical investigations include serum creatine kinase (typically increased 10-fold) with lesser elevations of the serum aldolase, aspartate transaminase, alanine transaminase and lactic dehydrogenase. Diagnosis is made by estimating the acid alpha glucoside activity in either skin biopsy (fibroblasts), muscle biopsy (muscle cells) or in white blood cells. The choice of sample depends on the facilities available at the diagnostic laboratory.

In various embodiments of the combination according to the present invention the disease is different from a lysosomal storage disease. In an embodiment of the combination according to the present invention the disease different from a lysosomal storage disease is a disease associated with an LSD. In an embodiment of the combination according to the present invention the disease different from a lysosomal storage disease is a neurodegenerative disorder. In an embodiment of the combination according to the present invention the neurodegenerative disorder is selected from the group comprising Parkinson's disease. In connection therewith it has to be understood that it has been recently discovered that there is a link between mutations in lysosomal enzymes and neurological disorders different from a lysosomal storage disease.

As one example there is a well-established link between mutations in the glucocerebrosidase gene and Parkinson's disease. More specifically, although Parkinson's disease results from accumulation of synuclein, mutations of the glucocerebrosidase gene occur in Parkinson patients with a frequency that is 6-fold increased (Sidransky et al., NEJM, 2009). Accordingly, Parkinson's disease is a disease associated with an LSD, more particularly with Gaucher's disease. More particularly, it has been shown that beta glucocerebrosidase co-localizes with alpha synuclein, which is the plaque forming material in Parkinson's disease. (Sidransky et al., NEJM, 2009). The present inventors thus currently assume that the

misfolding of mutant glucocerebrosidase in Gaucher may enhance accumulation of synuclein. Accordingly, the combination according to the present invention wherein the first constituent is a compound having the ability to rearrange mutant glucocerebrosidase, wherein the mutant glucocerebrosidase has a reduced activity, and wherein the second constituent is Ambroxol and/or a derivative of Ambroxol is suitable for or is for the treatment of Parkinson's disease. In an embodiment of the present invention the combination according to the present invention wherein the lysosomal protein is affected in a disease, the disease is a disease different from an LSD, preferably the disease is a neurodegenerative disorder, more preferably the disease is a disease associated with an LSD. In an embodiment of the combination according to the present invention the lysosomal protein is glucocerebrosidase and the disease preferably is Parkinson disease.

Parkinson's disease also referred to herein as Parkinson disease or PD, is a degenerative disorder of the central nervous system that often impairs the sufferer's motor skills, speech, and other functions (Jankovic J et al., April 2008, *J. Neurol. Neurosurg. Psychiatr.* **79** (4): 368–76).

Parkinson's disease also referred to herein preferably as PD, belongs to a group of conditions called movement disorders.

Parkinson's disease is characterized by muscle rigidity, tremor, a slowing of physical movement (bradykinesia) and a loss of physical movement (akinesia) in extreme cases. The primary symptoms are the results of decreased stimulation of the motor cortex by the basal ganglia, normally caused by the insufficient formation and action of dopamine, which is produced in the dopaminergic neurons of the brain, specifically the substantia nigra. Secondary symptoms may include high level cognitive dysfunction and subtle language problems. PD is both chronic and progressive. PD is the most common cause of chronic progressive Parkinsonism, a term which refers to the syndrome of tremor, rigidity, bradykinesia and postural instability. While many forms of Parkinsonism are idiopathic, "secondary" cases may result from toxicity most notably of drugs, head trauma, or other medical disorders.

The term Parkinsonism is used for symptoms of tremor, stiffness, and slowing of movement caused by loss of dopamine. "Parkinson's disease" is the synonym of "primary Parkinsonism", i.e., isolated Parkinsonism due to a neurodegenerative process without any secondary systemic cause. In some cases, it would be inaccurate to say that the cause is "unknown", because a small proportion is caused by genetic mutations. It is possible for a patient to be initially diagnosed with Parkinson's disease but then to develop additional features, requiring revision of the diagnosis (National Institute for Health and Clinical Excellence. Clinical guideline 35: Parkinson's disease. London, June 2006).

The usual anti-Parkinson's medications are typically either less effective or completely ineffective in controlling symptoms; patients may be exquisitely sensitive to neuroleptic medications like haloperidol, so correct differential diagnosis is important. Essential tremor may be mistaken for Parkinson's disease, but lacks all other features besides tremor, and has particular characteristics distinguishing it from Parkinson's disease, such as improvement with beta blockers and alcoholic beverages. PD is not considered to be a fatal disease by itself, but it progresses with time. The average life expectancy of a PD patient is generally lower than for people who do not have the disease. In the late stages of the disease, PD may cause complications such as choking, pneumonia, and falls that can lead to death. The progression of symptoms in PD may take 20 years or more. In some people, however, the disease progresses more quickly. There is no way to predict what course the disease will take for an individual person. With appropriate treatment, most people with PD can live productive lives for many years after diagnosis. There are some indications that PD acquires resistance to drug treatment by evolving into a Parkinson-plus disorder, usually Lewy body dementia, although transitions to progressive supranuclear palsy or multiple system atrophy are not unknown (Apaydin H. et al., January 2002, *Archives of Neurology* 59 (1): 102–12; Spanaki C. et al., October 2006, *Neurology* 67 (8): 1518–9.).

Niemann-Pick diseases are genetic diseases which are classified in a subgroup of LSDs called sphingolipidoses or lipid storage disorders in which harmful quantities of fatty compounds, or lipids, accumulate in the spleen, liver, lungs, bone marrow, and brain. In the classic infantile type A variant, a missense mutation causes deficiency of sphingomyelinase. Sphingomyelin is a component of cell membrane including the organellar membrane and so the enzyme deficiency blocks degradation of lipid, resulting in the accumulation of sphingomyelin within

lysosomes in the macrophage-monocyte phagocyte lineage. Affected cells become enlarged, sometimes up to 90 microns in diameter, secondary to the distention of lysosomes with sphingomyelin and cholesterol.

Symptoms are related to the organs in which the fatty compounds or lipids accumulate. Enlargement of the liver and spleen (hepatosplenomegaly) may cause reduced appetite, abdominal distension and pain as well as thrombocytopenia secondary to splenomegaly. Sphingomyelin accumulation in the central nervous system, including the cerebellum, results in unsteady gait (ataxia), slurring of speech (dysarthria) and disordered swallowing (dysphagia). Basal ganglia dysfunction causes abnormal posturing of the limbs, trunk and face (dystonia) and upper brainstem disease results in impaired voluntary rapid eye movements (supranuclear gaze palsy). More widespread disease involving the cerebral cortex and subcortical structures is responsible for gradual loss of intellectual abilities causing dementia and seizures. Sleep related disorders are also seen, including gelastic cataplexy, which means a sudden loss of muscle tone associated with laughter, and sleep inversion, which means sleepiness during the day and wakefulness at night.

Treatments for Niemann-Pick disease are limited with care being mostly supportive. Anecdotally, organ transplant has been attempted with limited success. Future prospects include enzyme replacement and gene therapy. Bone marrow transplant has been attempted for Type B. Supportive care through nutrition, medication, physical therapy and being followed by specialists can help with quality of life.

The drug Zavesca comprising Miglustat as an active ingredient, provided by Actelion, has been approved at least in the European Union for the treatment of progressive neurological manifestations in adult patients and pediatric patients with Niemann-Pick type C disease, also referred to herein as NPC.

In an embodiment of the lysosomal protein having reduced activity, said lysosomal protein is affected in an LSD.

A person skilled in the art will know other LSDs which can be treated with PCT and to which the combination according to the present invention may be applied.

AN LSD, as preferably used herein in connection with the various aspects of the present invention such as a combination according to the present invention, a use according to the present invention, a pharmaceutical preparation according to the present invention and/or a method according to the present invention may be selected from the group comprising LSDs having and/or being characterized in at least one of the following:

- a) a defective metabolism of glycosaminoglycans such as in, e.g., mucopolysaccharidosis, also referred to herein preferably as MPS, comprising among others MPS I, such as Hurler, Hurler-Scheie, Scheie; MPS II, such as Hunter; MPS III such as Sanfilippo syndrome type A, Sanfilippo syndrome type B, Sanfilippo syndrome type C and Sanfilippo syndrome type D; MPS IV such as Morquio type A and B; MPS VI such as Maroteaux-Lamy; MPS VII such as Sly; MPS IX such as hyaluronidase deficiency, multiple sulfatase deficiency;
- b) a defective degradation of glycan portion of glycoprotein such as in, among others, aspartylglucosaminuria; fucosidosis, type I and II; mannosidosis; sialidosis, type I and II;
- c) a defective degradation of glycogen such as in, e.g., Pompe's disease;
- d) a defective degradation of sphingolipid components such as in, among others, acid sphingomyelinase deficiency such as Niemann-Pick A & B; Fabry disease; Farber disease; Gaucher disease type I, II and III, GM1 gangliosidosis, type I, II and III; GM2 gangliosidosis such as Tay-Sachs type I, II, III and Sandhoff; Krabbe disease; metachromatic leukodystrophy, type I, II and III;
- e) a defective degradation of polypeptides such as in, e.g., pycnodysostosis;
- f) a defective degradation or transport of cholesterol, cholesterol esters, or other complex lipids such as in, e.g. neuronal ceroid lipofuscinosis, type I, II, III and IV;
- g) multiple deficiencies of lysosomal enzymes such as in, e.g., galactosialidosis, Danon Disease, GM2 gangliosidosis, mucopolipidosis type II and mucopolipidosis III; and

h) transport and trafficking defects such as in, e.g., cystinosis; Danon disease; mucopolidosis type IV; Niemann-Pick type C; infantile sialic acid storage disease; Salla disease.

In an embodiment of the present invention, wherein an LSD is an LSD having a defective degradation of a sphingolipid component, it will be understood by a person skilled in the art that wherein a degradation of a sphingolipid component is defective, preferably also the degeneration of several sphingolipid components may be defective.

In a preferred embodiment the lysosomal protein is selected from the group comprising an enzyme, a membrane transport protein, a trans-membrane protein or a soluble non-enzymatic protein. Preferably, the reduced activity is reduced as a result of at least one mutation within the nucleotide sequence of a gene coding for the lysosomal protein. The gene product of said mutant gene is more preferably the protein having reduced activity. Most preferably said reduced activity results in an LSD and/or the reduced activity is reduced due to a mutation of the amino acid sequence of the lysosomal protein.

In a still further preferred embodiment of the lysosomal protein being an enzyme said lysosomal enzyme is selected from the group comprising a lysosomal hydrolase and a phosphotransferase.

In an embodiment of the combination according to the present invention the phosphotransferase is selected from the group comprising N-acetylglucosamine-1- phosphate transferase.

In an embodiment of the combination according to the present invention the lysosomal storage disease is a lysosomal storage disease having a defective degradation of a glycan portion of a glycoprotein. In connection therewith it will be acknowledged that N-glycosylated proteins, in principle, can be affected by a lack of or a reduction of degradation resulting in lysosomal storage. In case of aspartylglucosaminuria, for example, Asn-GlcNAc units accumulate in the lysosome, whereas in case of manosidase the oligosaccharides $\text{Man}(\alpha 1 \rightarrow 3)\text{Man}(\beta 1 \rightarrow 4)\text{GlcNAc}$, $\text{Man}(\alpha 1 \rightarrow 2)\text{Man}(\alpha 1 \rightarrow 3)\text{Man}(\beta 1 \rightarrow 4)\text{GlcNAc}$ and $\text{Man}(\alpha 1 \rightarrow 2)\text{Man}(\alpha 1 \rightarrow 2)\text{Man}(\alpha 1 \rightarrow 3)\text{Man}(\beta 1 \rightarrow 4)\text{GlcNAc}$ are formed (DeGasperi R et

al., J Biol Chem 1991, 266:16556-16563.). It will be understood that in the latter case high mannosylated proteins are affected with preference. As to fucosidosis, fucose containing sugars are degraded. It will be understood that fucose is a sugar which is capable of forming the last moiety of a chain or is capable of linking two different sugars.

A person skilled in the art will acknowledge that, for example, lysosomal proteins, secretory proteins and membrane proteins, such as adhesion proteins and/or transporter proteins, can be glycosylated.

In an embodiment of the combination according to the present invention the lysosomal storage disease is a lysosomal storage disease having a defective degradation of polypeptides. In connection therewith it will be understood that Cathepsin K is a cysteine protease. Accordingly, all proteins comprising a cysteine residue and a nearby amino acid having a basic side chain are candidates for degradation by Cathepsin K. It will also be acknowledged that Cysteine proteases carry out various tasks in different physiological processes of the cell.

In an embodiment of the combination according to the present invention the lysosomal storage disease is a lysosomal storage disease having a defective degradation or transport of cholesterol, cholesterol esters and/or other complex lipids. A complex lipid as used herein preferably refers to lipids comprising lipofuscin. In connection therewith it will be understood that lipofuscin accumulates in NCLs and comprise a protein moiety, a lipid moiety, preferably a sugar and preferably metal ion, wherein the lipid moiety results from oxidation of unsaturated fatty acids.

In an embodiment of the combination according to the present invention the lysosomal storage disease is a lysosomal storage disease having multiple deficiencies of lysosomal enzymes. In an embodiment the lysosomal storage disease having multiple deficiencies of lysosomal enzymes is selected from the group comprising galactosialidosis, Danon Disease (LAMP2), pycnodysostosis (Cathepsin K), multiple sulfatase deficiency (Fgly-Generating Enzyme), GM2 gangliosidosis (GM2-Activator), mucopolipidosis type II and mucopolipidosis type III (N-acetylglucosaminyl-1-phosphotransferase). In connection therewith „multiple deficiencies“ as used herein preferably means at least two deficiencies of at least two different enzyme activities in the lysosome, preferably of at least two different enzymes. As an

example, the gene CTSA is mutated in galactosialidosis. CTSA encodes Cathepsin A (CatA), which is a factor forming complex(es) with other lysosomal hydrolases (beta-galactosidase). In other words, CatA is important for the functioning of different hydrolases. If not present various degradation processes will not function properly. Accordingly, multiple deficiencies are caused. In case of mucopolipidosis the same principle applies: In mucopolipidosis II the enzyme N-acetylglucosaminyl-1-phosphotransferase is mutated, which is responsible for marking lysosomal enzymes for their transport into the lysosome. The mutation in said gene leading to a reduced activity of the protein/gene product results in multiple deficiencies of enzyme activities which occur in the lysosome. In other words, the mutated genes the mutation of which results in multiple deficiencies assist other enzymes in correct processing and assist said enzymes in their catalytic function, respectively.

In an embodiment of the combination according to the present invention the lysosomal protein is an enzyme, more preferably, the enzyme is selected from the group comprising mutant α -galactosidase A preferably resulting in Fabry disease, mutant acid α -glucosidase preferably resulting in Pompe disease, mutant acid sphingomyelinase preferably resulting in Niemann-Pick diseases type A or B, mutant N-acetylglucosamine-1-phosphotransferase preferably resulting in mucopolipidosis type II and IIIA, mutant beta-hexosaminidase A/B preferably resulting in Tay-Sachs' or Sandhoff disease, mutant beta galactosidase preferably resulting in GM1 gangliosidosis, mutant acid Lipase preferably resulting in Wolman Disease and/or CESD, mutant galactosylceramidase preferably resulting in Krabbe Disease, mutant arylsulfatase B preferably resulting in Maroteaux-Lamy Syndrome, i.e. mucopolysaccharidosis VI, mutant alpha-L iduronidase preferably resulting in Hurler Syndrome, i.e. mucopolysaccharidosis I, mutant alpha-mannosidase preferably resulting in alpha-mannosidosis, mutant beta-glucuronidase preferably resulting in Sly Syndrome, mutant arylsulfatase A preferably resulting in metachromatic leukodystrophy, mutant NPC1 protein preferably resulting in Niemann-Pick disease type C, mutant α -L-iduronidase preferably resulting in Hurler-Scheie syndrome, mutant α -L-iduronidase preferably resulting in Scheie syndrome, mutant iduronatsulfatsilfatase preferably resulting in Hunter syndrome, mutant α -N-acetylglukoseamidase preferably resulting in Sanfilippo syndrome type B, mutant acetyl-CoA: α -glukosaminid-N-acetyltransferase preferably resulting in Sanfilippo syndrome type C, mutant N-acetyl-glukosamin-6-sulfatsulfatase preferably resulting in Sanfilippo syndrome type D, mutant N-acetyl-glukosamin-6-sulfatsulfatase preferably resulting in Morquio type A,

mutant β -galactosidase preferably resulting in Morquio type B, mutant N-acetylgalactosamin-4-sulfat-sulfatase preferably resulting in Maroteaux-Lamy, mutant β -glucuronidase preferably resulting in Sly, mutant hyaluronidase preferably resulting in hyaluronidase deficiency; mutant N-acetylglucosamine-1-phosphatetransferase preferably resulting in N-acetylglucosamine-1-phosphate transferase mucopolidose III gamma and mutation of multiple sulfatase enzymes preferably resulting in multiple sulfatase deficiency.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is castegine A3 or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is α -galactosidase A, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Fabry's disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is castegine B2 or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is α -galactosidase A, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Fabry's disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a

compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is N-methyl calystegine A3 or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is α -galactosidase A, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Fabry's disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is N-methyl calystegine B2 or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is α -galactosidase A, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Fabry's disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 2-N-acetylamino-isofagomine or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is beta-hexosaminidase A preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis,

Fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Tay-Sachs disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 1,2-dideoxy-2-acetamido-nojirimycin or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is beta-hexosaminidase A preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, Fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Tay-Sachs disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is nagastain or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is beta-hexosaminidase A preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Tay-Sachs disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein

has a reduced activity, wherein the first constituent is nagastain or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is beta-hexosaminidase B, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Sandhoff disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 2-N-acetylamino-isofagomine or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is beta-hexosaminidase B, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, Fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Sandhoff disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 1,2-dideoxy-2-acetamido-nojirimycin or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is beta-hexosaminidase B, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs

disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Sandhoff disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 4-epi-isofagomine or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is acid beta-galactosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is GM1-gangliosidosis.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 4-epi-isofagomine or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is acid beta-galactosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Morquio disease B.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 4-epi-isofagomine or a

pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is galactocerebrosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Krabbe disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 1-deoxygalactonojirimycin or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is acid beta-galactosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is GM1-gangliosidosis.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 1-deoxygalactonojirimycin or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is acid beta-galactosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff

disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Morquio disease B.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 1-deoxygalactonojirimycin or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is galactocerebrosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Krabbe disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 1-deoxymannojirimycin or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is acid alpha-manosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is alpha-manosidosis.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is swainsonine or a pharmaceutically

acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is acid alpha-mannosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is alpha-mannosidosis.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is mannosatin A or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is acid alpha-mannosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is alpha-mannosidosis.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 2-hydroxy-isofagomine or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is acid beta-mannosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff

disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is beta-mannosidosis.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 1-deoxyfuconojirimycin or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is acid alpha-fucosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is fucosidosis.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is beta-homofuconojirimycin or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is acid alpha-fucosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is fucosidosis.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 2,5-imino-1,2,5-trideoxy-L-

glucitol or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is acid alpha-fucosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is fucosidosis.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 2,5-dideoxy-2,5-imino-D-fucitol or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is acid alpha-fucosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is fucosidosis.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 2,5-imino-1,2,5-trideoxy-D-altritol or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is acid alpha-fucosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff

disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is fucosidosis.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 1,2-dideoxy-2-N-acetamido-nojirimycin or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is alpha-N-Acetylglucosaminidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Sanfilippo disease B.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 1,2-dideoxy-2-N-acetamido-galactonojirimycin or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is alpha-N-Acetylgalactosaminidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Schindler-Kanzaki disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 1-deoxyduronojirimycin or a

pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is alpha-L-iduronidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Hurler-Scheie disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 2-carboxy-3,4,5-trideoxypiperidine or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is alpha-L-Iduronidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Hurler-Scheie disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 6-carboxy-isofagomine or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is beta-glucuronidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-

Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Sly disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 2-carboxy-3,4,5-trideoxypiperidine or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is beta-glucuronidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Sly disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 2,6-dideoxy-2,6-imino-sialic acid or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is sialidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is sialidosis.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is siastin B or a pharmaceutically

acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is sialidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is sialidosis.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is N-nonyl-deoxynojirimycin (NN-DNJ) or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is glucocerebrosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Parkinson disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is N-octyl-2,5-anhydro-2,5-imino-D-glucitol or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is glucocerebrosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Parkinson disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is N-octyl-isofagomine or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is glucocerebrosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Parkinson disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is N-octyl-beta-valienamine (NOV) or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is glucocerebrosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Parkinson disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is isofagomine (IFG) or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is glucocerebrosidase, preferably

the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Parkinson disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is calystegine A3 or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is glucocerebrosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Parkinson disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is calystegine B1 or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is glucocerebrosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Parkinson disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is calystegine B2 or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is glucocerebrosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Parkinson disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is calystegine C1 or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is glucocerebrosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Parkinson disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is 1,5-dideoxy-1,5-iminoxylitol (DIX) or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is glucocerebrosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is

selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Parkinson disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is alpha-1-C-nonyl-DIX or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is glucocerebrosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Parkinson disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is alpha-1-C-octyl-1-DNJ or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is glucocerebrosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Parkinson disease.

In an embodiment of the combination according to the present invention the combination comprises a first constituent and a second constituent, wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the first constituent is N-butyl-DNJ or a pharmaceutically acceptable salt thereof; and wherein the second constituent is Ambroxol and/or a derivative thereof, preferably the lysosomal protein is glucocerebrosidase, preferably the combination is for use in the treatment or prevention of a disease, whereby the disease is selected from the group comprising Fabry disease, GM1-gangliosidosis, Pompe disease, Krabbe disease, Morquio disease B, alpha-mannosidosis, beta-mannosidosis, fucosidosis, San filippo disease B, Schindler-Kanzaki-disease, Tay-Sachs disease, Sandhoff disease, Hurler-Scheie disease, Sly disease, sialidosis and Parkinson disease, more preferably the disease is Parkinson disease.

It will be understood that a person skilled in the art will be aware of other proteins having reduced activity in respective LSDs, which may be treated with the combination according to the present invention.

In a still further preferred embodiment of the lysosomal protein being a trans-membrane protein, said lysosomal trans-membrane protein is selected from the group comprising mutant NPC1 preferably resulting in Nieman-Pick disease type C1.

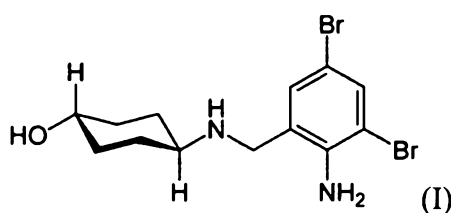
In a still further preferred embodiment of the lysosomal protein being a soluble non-enzymatic protein said lysosomal non-enzymatic protein is selected from the group comprising NPC2, preferably resulting in Niemann-Pick disease type C2.

It will be understood that a person skilled in the art will be aware of other proteins having reduced activity in respective LSDs, which may be treated with the combination according to the present invention.

Ambroxol, also referred to herein as 2-amino-3,5-dibromo-N-[trans-4-hydroxy cyclohexyl]benzylamine, is a drug from the expectorant class widely used for increasing surfactant secretion in lung and for decreasing mucus viscosity (Disse, B.G. et al., 1987, Eur. J. Respir. Dis. 153, 255-262). Ambroxol has been reported to protect alpha-1-proteinase

inhibitor from oxidative inactivation and to inhibit the generation of reactive oxygen species from activated phagocytes in vitro (Rozniecki, J. and Nowak, D., 1987, *Lu Resp.* 4, 14-15; Winsel, K. and Becher, G., 1922, *Eur. Resp. J.* 5, 289). Furthermore, pulse radiolysis experiments showed that Ambroxol is a good scavenger of the primary water radical species, particularly e^-_{aq} and $\cdot OH$ radicals (Tamba, M. and Torreffianni, A. 2001, *Rad. Phys. And Chem.* 60, 43-52).

Ambroxol as used herein, preferably has the formula according to formula (I):



Ambroxol is the active ingredient of Mucosolvan, Ambrosan or Mucoangin. The compound is a mucoactive drug with several properties including secretolytic and secretomotoric actions that restore physiological clearance mechanisms of the respiratory tract which play an important role in the body's natural defense mechanisms. It stimulates synthesis and release of surfactant by type II pneumocytes. Surfactants act as anti-glue factors by reducing the adhesion of mucus to the bronchial wall, in improving its transport and in providing protection against infection and irritating agents (Sanderson RJ et al. *Respir Phys* 1976; 27: 379-392.; Kido H et al. *Biol Chem* 2004; 385: 1029-1034). Furthermore, Ambroxol is indicated as secretolytic therapy in bronchopulmonary diseases associated with abnormal mucus secretion and impaired mucus transport. It promotes mucus clearance, facilitates expectoration and eases productive cough, allowing patients to breathe freely and deeply (Malerba and Ragnoli, *Expert Opin Drug Metab Toxicol* 2008; 4(8): 1119-1129). There are many different formulations developed since the first marketing authorization in 1978. A major product is the syrup with two concentrations of the compound, 30 mg/ml and 315 mg/ml, which can be given in adults and children from the age of 1 year on and even from infant in the latter concentration. Other formulations are tablets containing 30 mg or 60 mg, and a pastille to be sucked with 15 mg Ambroxol. There is also a sustained release form with 75 mg to be given once a day. Ambroxol is also available as dry powder sachets, inhalation solution, drops and ampoules as well as effervescent tablets. Ambroxol also provides pain

relief in acute sore throat. Pain in sore throat is the hallmark of acute pharyngitis (de Mey et al., *Arzneimittelforschung* 2008; 58(11): 557 - 568).

Ambroxol is a very potent inhibitor of the neuronal Na⁺ channels (Weiser T., *Neurosci Lett.* 2006; 395: 179 – 184). This property led to the development of a lozenge containing 20 mg of Ambroxol. Many state-of-the-art clinical studies (de Mey et al., *supra*) have demonstrated the efficacy of Ambroxol in relieving pain in acute sore throat, with a fast onset of action and a long duration of effect of at least 3 hours. Additional anti-inflammatory properties of Ambroxol are of clinical relevance since treatment lead to a marked reduction of redness of the patient's sore throat.

Lysosomal protein having reduced activity, such as mutant lysosomal hydrolases, such as alpha-galactosidase A affected in Fabry disease or acid α -glucosidase affected in Pompe disease form a heterogeneous group of enzymes and moreover pharmacologic chaperones such as DGJ or NB-DNJ, specifically act only in relation to a certain LSD, i.e. in relation to a particularly specific enzyme as has already been outlined above.

Moreover, DGJ treatment does also not address all mutations of alpha-galactosidase A (alpha-Gal A) related to Fabry disease, which inhibits the broad application of the compound in clinical use. This was shown when Fabry patient-derived T-cells were grown in the presence of DGJ. Alpha-Gal A activity increased to more than 50% of normal in several mutations but was unaffected in others. (Shin SH et al., *Pharmacogenet Genomics.* 2008 Sep; 18(9):773-80.).

It is therefore a common believe in the field of LSDtherapy that the response of a certain mutant enzyme affected in an LSD to a certain pharmacological chaperone is dependent on the affected enzyme, on (a)certain mutation(s) of the enzyme and thus on the certain patients (Wu et al., *supra*).

It is thus the merit of the present inventors having found that the combined use of a pharmacological chaperone and Ambroxol is suitable for increasing the activity of a mutant protein in the treatment of an LSD such as Fabry disease compared to the application of said pharmacological chaperone alone.

At least insofar the present invention turns away from the teaching of prior art in that administering Ambroxol according to the present invention is useful independent from the particular LSD and the particular pharmacological chaperone in that Ambroxol is administered in addition to a pharmacological chaperone specific for the particular treated disease causing an increase of the activity of a mutant enzyme compared to the administering of said pharmacological chaperone alone, although, preferably, Ambroxol is not causing an increase of the activity of said mutant enzyme when administered alone.

A person skilled in the art will immediately acknowledge that the combination according to the present invention is able to extend the use of a compound having the ability to rearrange a lysosomal protein having reduced activity, such as a pharmacological chaperone, e.g. an iminosugar such as DGJ, to a wider range of indications as if administered alone in that the treatment with the combination according to the present invention allows the treatment of the protein having reduced activity due to (a) particular mutation(s) which may be only insufficiently treated with the compound having the ability to rearrange a lysosomal protein having reduced activity, such as DGJ, alone. More specifically, the additional application of Ambroxol, Bromhexine and/or a derivative thereof comprised in the combination of the present invention may enhance the activity of the mutant protein having reduced activity to a level sufficient to prevent the patient from having symptoms, from the onset of the disease or at least for suffering from less severe symptoms. In contrast, the responsiveness of a mutant protein having said particular mutation to the treatment with DGJ alone may not be sufficient to induce a therapeutic effect.

Valenzano et al. (Valenzano et al., *Assay Drug Dev Technol.* 2011 Jun;9(3):213-35), for example, describe that in most LSDs, 1% to 6% of normal activity has been estimated to be sufficient to delay or prevent disease onset or to yield a more mild form of the disease. For instance, 1%–2% of normal α -iduronidase activity has been reported in mild cases of MPS I (Hopwood and Muller, *Clin Sci (Lond)*. 1979 Sep;57(3):265-72) ; <1% of normal GCase activity has been reported in mild Gaucher disease (Desnick RJ and Fan JQ, In: Futerman AH, editor; Zimran A, editor. *Gaucher Disease*. CRC Press; Boca Raton, FL: 2006. p. 544); 6% of normal β -galactosidase activity has been reported in late-onset GM1 gangliosidosis (Suzuki Y et al., *Perspect Med Chem.* 2009;3:7–19); 2–4% of normal β -hexosaminidase

activity has been reported in the adult form of GM2 gangliosidosis (Conzelmann et al. *Am J Hum Genet.* 1983 Sep;35(5):900-13); and 10% of normal enzyme activity seems to prevent GM1 and GM2 gangliosidoses altogether (Leinekugel P. et al., *Hum Genet.* 1992;88:513–523; Mahuran DJ., *Biochim Biophys Acta.* 1999;1455:105–138).

It will thus be acknowledged by a person skilled in the art that 10%, preferably 20% and more preferably 25% of protein activity compared to wild type protein activity is considered to be sufficient for a therapeutically effective treatment, i.e. a treatment sufficient to induce a therapeutic effect, and/or an enhancement of the activity of the mutant protein having reduced activity to a level sufficient to prevent the patient from having symptoms, from the onset of the disease or at least for suffering from less severe symptoms.

For example, enhancing the activity of a mutant protein compared to wild type protein activity to 5% by administering DGJ alone, the activity of the mutant protein being 1% without any treatment, will not be sufficient to prevent the onset of disease, i.e. the treatment will not be therapeutically effective. By contrast, the application of the combination according to the present invention, going along with additional application of Ambroxol, bromhexine and/or a derivative thereof may further enhance the mutant protein's activity to 20%, thus being therapeutically effective and prevent the patient from suffering from symptoms, or at least may cause less severe symptoms.

In other words, the present inventors found that applying DGJ and Ambroxol in combination according to the present invention is able to increase the activity of mutant protein measured in accordance with the cell-culture system as described herein to a level which would prompt a skilled clinician to evaluate the patient's health status as "healthy", i.e. having an activity of mutant protein which usually does not result in symptoms otherwise caused by the mutant protein's reduced activity, whereas the treatment with DGJ alone results in protein activity of the mutant protein which would prompt the skilled clinician to evaluate the patient's health status as being „unclear“ or as “diseased”.

The present inventors also surprisingly observed an increase in the activity of a mutant protein upon treatment with the combination according to the present invention compared to the

activity of the mutant protein upon treatment with the compound having the ability to rearrange a lysosomal protein having reduced activity alone.

It is commonly known in the art that a certain protein having reduced activity due to a certain mutation will respond better to the treatment with a compound having the ability to rearrange a lysosomal protein having reduced activity than other proteins having other mutations. The effectiveness of treatment may thus allow distinguishing between responsive mutations and non-responsive mutations of the protein having reduced activity.

The present inventors observed that more responsive mutations, i.e. proteins having mutations where the enhancement of protein activity due to administration of the compound having the ability to rearrange a lysosomal protein having reduced activity was higher, also exhibited a stronger effect in further enhancing the activity of said protein having said certain mutation by applying the combination according to the present invention. Insofar, restoration of the activity of a lysosomal protein the activity of which is reduced in a disease and more specifically a lysosomal storage disease to an activity of at least 10 %, preferably at least 20 % and more preferably at least 25 % of the lysosomal protein of wild type and/or of the lysosomal protein in a healthy subject, is preferably regarded as a treatment, preferably as a treatment of the disease.

For example, and as may also be taken from Fig. 3 and Example 3 of the instant application, a stronger effect in further enhancing the activity of certain GLA mutations, which exhibit a strong responsiveness to treatment with DGJ alone (see also table 3), could be seen when applying the combination according to the present invention. More specifically, said certain GLA mutations are, for example, E59K (14.2-fold), A156V (3.9- fold) and R301Q (5.6- fold) whereas the effect in enhancing the activity of mutant proteins was less strong with regard to proteins less responsive to DGJ treatment. R356W (3.7-fold) and R363H (1.8-fold) additionally show a relatively strong residual activity, i.e. high protein activity without treatment.

In connection therewith it is important to note that other studies, e.g. related to the treatment of Gaucher's disease with ryanodine receptors (RyRs), already considered an increase of 1.3-

fold compared to the residual activity of the mutant enzyme as being clinically relevant (Wang et al., ACS Chemical Biology, 2011).

Furthermore, Wu et al. (Wu et al., *supra*) showed that a protein having a particular mutation and being responsive to the treatment with a pharmacological chaperone in vitro was not responsive when tested in a patient having said mutant form of the protein, whereby the patient was treated with 10 μ M DGJ, and whereas in all other patients tested, the responsiveness of the mutations of said patients tested in the HEK cell culture system represented the responsiveness in clinical application.

It is thus a preferred embodiment of the combination of the present invention that the activity of the protein having reduced activity is reduced due to a mutation of said protein, wherein the mutation is responsive to the treatment with a compound having the ability to rearrange the lysosomal protein having reduced activity when administered alone, i.e. the activity of the protein having reduced activity is enhanced due to the treatment with a compound having the ability to rearrange the lysosomal protein.

In a more preferred embodiment of the combination of the present invention the protein having reduced activity is responsive to the treatment with a compound having the ability to rearrange the lysosomal protein having reduced activity in a patient, wherein the enhancement of the activity of said protein results in a protein activity of preferably 5% and most preferably 10% compared to the wild type protein activity.

As Ambroxol is widely used in connection with different diseases and symptoms including acute pain, such as toothache, postoperative pain or herpes zoster induced pain, as well as cranial infection, stroke, burns, pancreatitis or trauma, colic and headache, a person skilled in the art will acknowledge that a dosage applied in the treatment of a patient suffering from or being at risk of suffering from a lysosomal disease, such as Fabry disease will take into consideration the already approved dosages to be administered in connection with said diseases for which Ambroxol has been approved by health authorities. Accordingly, dosages of Ambroxol preferably range from 30 to 4000 mg per day as described in US patent application 11/856280 using e.g. Ambroxol tablets comprising 150-1200 mg as essentially described in US patent application 10/888362 filed July 9, 2004 by Boehringer Ingelheim

International GmbH or a lozenge as described in US patent 6,663,889 filed April 23, 2002 by Boehringer Ingelheim International GmbH. A person skilled in the art will also acknowledge that up to 20g of Ambroxol per patient is applied for detoxification of patients suffering from liver failure.

In the cell culture system preferably used in connection with the present invention an amount of Ambroxol that results in an increase of alpha-galactosidase A activity is preferably in a range of 20 to 60 μ M, preferably 40 μ M, in order to achieve a stable effect, and may also be taken from Fig. 2 and Example 3 as described herein.

A person skilled in the art will further acknowledge that a dosage to be administered to a patient in order to treat a particular LSD, whereby said patient is suffering from or is at risk of developing the same, will depend on the particular LSD to be treated as well as on the particular combination according to the present invention and more particularly on the particular compound having the ability to rearrange a lysosomal protein having reduced activity such as a particular chaperone, and on the particular derivative of Ambroxol such as bromhexine or Ambroxol contained in said composition, and on the particular condition of the patient to be treated. Accordingly, a person skilled in the art will apply a dosage of Ambroxol in the light of what has been outlined above, starting from a low dosage of Ambroxol and varying said dosage depending on the success of treatment, whereby the success of treatment may be monitored and controlled by regularly assessing the activity of the lysosomal protein having reduced activity affected in the particular LSD.

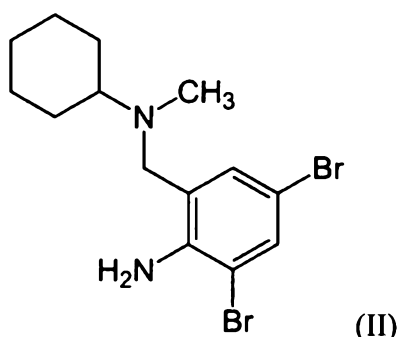
When conducting experiments in the cell culture system as outlined herein the present inventors have also found that when DGJ is administered together with bromhexine instead of Ambroxol, the activity of alpha-galactosidase A is also increased compared to the effect arising from DGJ alone.

In connection therewith it is important to note that Leet et al. (Lee et al. Pharmacol Res. 2004 Jan;49(1):93-8) showed that a maximum concentration of Ambroxol in human plasma as high as 61.5 ng/ml could be achieved which corresponds to 0.15 μ M. With regard to Bromhexine Bechgaard et al. (Bechgaard et al. Biopharm Drug Dispos. 1982 Oct-Dec;3(4):337-44) disclose a concentration of bromhexine of 0.08 μ M. A person skilled in the art will

acknowledge that when administering a 32 mg tablet (77.6 μ mol) an intestinal absorption of 22-27% occurs. Referring to the blood volume and after subtraction of loss by first pass metabolism this should result in approximately 0.7 μ M, i.e. which refers to a 10-fold increase. In clinical trials 10 μ M DGJ have been acknowledged as to be a clinically achievable concentration (Wu et al., 2011, Hum Mutat).

A person skilled in the art will acknowledge that Ambroxol is a metabolite of bromhexine.

Bromhexine, also referred to herein as 2,4-dibromo-6-[[cyclohexyl(methyl)amino]methyl]aniline is a mucolytic agent used in the treatment of respiratory disorders associated with viscid or excessive mucus. Bromhexine as referred to herein preferably has the formula:



In addition, bromhexine has antioxidant properties. Bromhexine thus supports the body's own natural mechanisms for clearing mucus from the respiratory tract. It is secretolytic, i.e. it increases the production of serous mucus in the respiratory tract and makes the phlegm thinner and less sticky. This contributes to a secretomotoric effect, i.e. it helps the cilia - tiny hairs that line the respiratory tract - to transport the phlegm out of the lungs. For this reason it is often added to some antitussive (cough) syrups.

Bromhexine is a synthetic derivative of the herbal active ingredient vasicine. It has been shown to increase the proportion of serous bronchial secretion, making it more easily expectorated. Bromhexine also enhances mucus transport by reducing mucus viscosity and by activating the ciliated epithelium. In clinical studies, Bromhexine showed secretolytic and secretomotoric effects in the bronchial tract area which facilitates expectoration and eases cough. It is indicated as "secretolytic therapy in bronchopulmonary diseases associated with

abnormal mucus secretion and impaired mucus transport". Bromhexine is contained in various formulations, high and low strength syrups 8mg/5ml, 4 mg/5ml, tablets and soluble tablets (both with 8 mg bromhexine) and solution for oral use 10 mg/ 5 ml. The posology varies with the age, but there are products for all age groups from infant on. Bromhexine is a well-established and well tolerated product in its known indication. A skilled person will agree that a dosage schedule of 8-16 mg administered three times per day will serve as a good starting point for a therapy, wherein the combination of the present invention will be applied.

It is particularly noteworthy that the use of Ambroxol and/or a derivative thereof, such as bromhexine in connection with the present invention is advantageous for the clinical application of the combination of the present invention in that Ambroxol and bromhexine constitute well tolerated drugs approved for other clinical uses.

Application of high concentrations of pharmacological chaperones is known to result in an inhibitory effect on the mutant protein's activity and may also exhibit toxicity. As will also be described herein in connection with Figures 5A and 5B the combined use of a dose of a pharmacological chaperone, preferably a sub-inhibitory dose, and Ambroxol and/or a derivative such as bromhexine is of further advantage as substantially lower doses of the pharmacological chaperone may be used in order to achieve the same effect. In other words, the treatment with a combination of the present invention is able to substitute the application of a high dose of the pharmacological chaperone in order to achieve the same effect, i.e. the same increase in mutant enzyme activity.

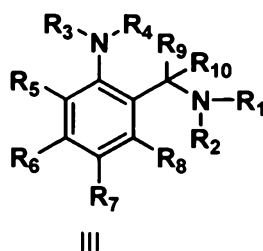
In a preferred embodiment of the present invention the combination of the present invention comprises a compound having the ability to rearrange a lysosomal protein having reduced activity and Ambroxol and/or a derivative of Ambroxol. In a further preferred embodiment said derivative of Ambroxol is bromhexine. It is thus also an embodiment of the present invention that a combination according to the invention comprises bromhexine and Ambroxol; Ambroxol; bromhexine; Ambroxol and bromhexine and optionally at least one derivative of bromhexine other than Ambroxol; Ambroxol and/or at least one derivative of bromhexine other than Ambroxol and/or bromhexine; and/or at least one derivative of bromhexine other than Ambroxol and/or bromhexine.

In an embodiment of the combination of the present invention the salt of Ambroxol or bromhexine is derived from an organic or inorganic acid. In an embodiment of the combination of the present invention the salt of Ambroxol or bromhexine is derived from an organic or inorganic acid selected from the group comprising hydrohalides such HCl, HBr, H₃PO₄, as well as organic acids such as acetic acid, benzoic acid, tartaric acid and citric acid.

A derivative as used herein preferably means any pharmaceutically acceptable salt, solvate, ester or amide, or salt or solvate of such ester or amide, of a compound comprised in a combination of the present invention or any other compound which upon administration to the recipient is capable of providing, directly or indirectly, the compound or an active metabolite or residue thereof, e.g. a prodrug. Preferred pharmaceutically acceptable derivatives according to the invention are any pharmaceutically acceptable salts, solvates or prodrugs.

A person skilled in the art will immediately acknowledge that Ambroxol is a metabolite of bromhexine and, accordingly, that Ambroxol is a derivative of bromhexine, as well as vice versa, preferably bromhexine is a derivative of Ambroxol. In a preferred embodiment of the composition according to the present invention a derivative of bromhexine and/or Ambroxol, or the compound having the ability to rearrange a lysosomal protein having reduced activity is a pharmaceutically active and pharmaceutically acceptable derivative. In other words, the derivative is preferably less toxic, more preferably not toxic, and is enhancing the activity of the protein having reduced activity when applied in the combination according to the present invention.

In an embodiment of the composition according to the present invention a derivative of bromhexine and/or Ambroxol is a compound of formula (III):



or a pharmaceutically acceptable salt thereof, wherein

each and any of R1, R2, R3, and R4 is individually and independently selected from hydrogen atom, an aliphatic carbon group with 1 to 12 carbon atoms, whereby this aliphatic carbon group can be either linear or cyclic, optionally with one or more O, N, S, P and/or halide substituents on the aliphatic carbon group, and an aromatic or heteroaromatic group, whereby this aromatic or heteroaromatic group constitutes of 2 to 18 carbon atoms, optionally with one or more additional N, O, S or halide atoms;

each and any of R5, R6, R7 and R8 is individually and independently selected from hydrogen atom, halide substituents such as I, Br, Cl, F, oxygen substituents such as OH, Oalkyl, OAryl, alkyl carboxylic acid derivatives such as CN, CO₂alkyl, CO₂aryl, CONH₂, CONHalkyl, CONHaryl, CON(alkyl)₂, CON(aryl)₂, acyl substituents such as C(O)alkyl, C(O)aryl, sulfur substituents such as SH, Salkyl, Saryl, SOalkyl, SOaryl, SO₂alkyl, SO₂aryl, SO₃alkyl, SO₃aryl, nitrogen substituents such as NH₂, NHalkyl, NHaryl, Nalkyl₂, Naryl₂, N(alkyl)aryl, NHSO₂alkyl, NHSO₂aryl, N(alkyl)SO₂alkyl, N(alkyl)SO₂aryl, N(aryl)SO₂alkyl, N(aryl)SO₂aryl, whereby alkyl stands for a linear or cyclic aliphatic carbon group from C1 to C6 with additional O, N, S, P and/or halide substituents on the aliphatic carbon group, and whereby aryl stands for an aromatic or heteroaromatic group with 2 to 18 carbon atoms, optionally with one or more additional N, O, S or halide atoms; and

each and any of R9 and R10 is individually and independently selected from hydrogen atom and an aliphatic carbon group from C1-C4, optionally with one or more additional O, N and/or halide substituents.

In an embodiment of the composition according to the present invention a derivative of bromhexine and/or Ambroxol is a compound of formula (III) or a pharmaceutically acceptable salt thereof, wherein

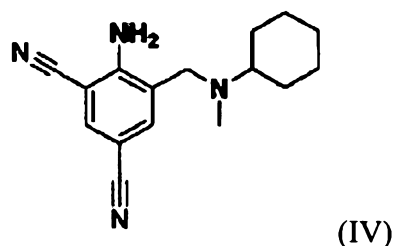
each and any of R1, R2, R3 and R4 is individually and independently selected from hydrogen atom, an aliphatic carbon group with 1 to 6 carbon atoms, whereby the aliphatic carbon group can be either linear or cyclic, optionally with one or more O, N, and/or halide substituents on the aliphatic carbon group, and an aromatic or heteroaromatic group, whereby the aromatic or

heteroaromatic group constitutes of 3 to 9 carbon atoms, optionally with one or more additional N, O, or halide atoms;

each and any of R5, R6, R7 and R8 is individually and independently selected from hydrogen atom, halide substituents such as I, Br, Cl, F, oxygen substituents such as OH, Oalkyl, Oaryl, alkyl, carboxylic acid derivatives such as CN, CO₂alkyl, CONH₂, CONHalkyl, CONHaryl, acyl substituents such as C(O)aryl, nitrogen substituents such as NH₂, NHalkyl, NHaryl, NHSO₂alkyl, NHSO₂aryl, whereby alkyl stands for a linear or cyclic aliphatic carbon group from C1 to C4, optionally with one or more O, N, and/or halide substituents on the alkyl group, and whereby aryl stands for an aromatic or heteroaromatic group with 3 to 9 carbon atoms, optionally with one or more additional N, O or halide atoms; and

each and any of R9 and R10 is individually and independently selected from hydrogen atom and an aliphatic carbon group from C1-C4, optionally with one or more O, N and/or halide substituents.

In an embodiment of the composition according to the present invention a derivative of bromhexine and/or Ambroxol is a compound of the formula (IV)

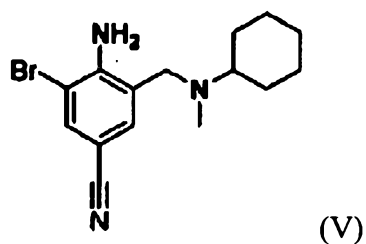


or a pharmaceutically acceptable salt thereof.

A compound of the formula (IV) is also referred to herein preferably as SF-54B.

In an embodiment of the composition according to the present invention a derivative of bromhexine and/or Ambroxol is a compound of the formula (V)

114

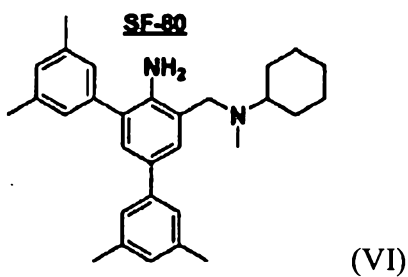


(V)

or a pharmaceutically acceptable salt thereof.

A compound of the formula (V) is also referred to herein preferably as SF-55C.

In an embodiment of the composition according to the present invention a derivative of bromhexine and/or Ambroxol is a compound of formula (VI)

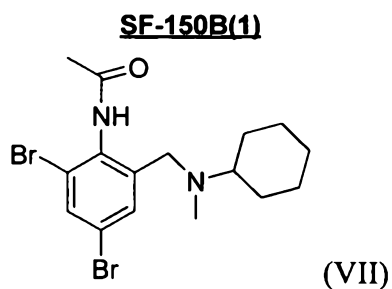


(VI)

or a pharmaceutically acceptable salt thereof.

A compound of the formula (VI) is also referred to herein preferably as SF-80.

In an embodiment of the composition according to the present invention a derivative of bromhexine and/or Ambroxol is a compound of formula (VII)



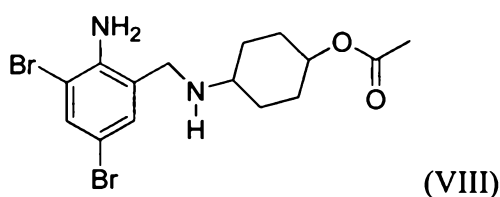
(VII)

or a pharmaceutically acceptable salt thereof.

A compound of formula (VII) is also referred to herein preferably as SF-150B or SF-150B(1).

In an embodiment of the composition according to the present invention a derivative of bromhexine and/or Ambroxol is a compound of formula (VIII)

SF-153B

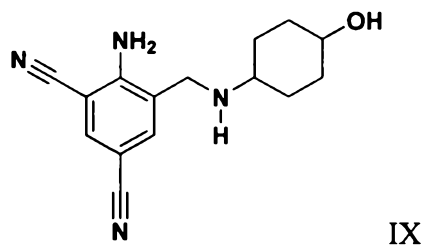


or a pharmaceutically acceptable salt thereof.

A compound of formula (VIII) is also referred to herein preferably as SF-153B.

In an embodiment of the composition according to the present invention a derivative of bromhexine and/or Ambroxol is a compound of formula (IX)

SF-124B



or a pharmaceutically acceptable salt thereof.

A compound of formula (IX) is also referred to herein preferably as SF-124B.

Suitable pharmaceutically acceptable salts of the compounds comprised in a combination of the present invention, more particularly, of Ambroxol, of a derivative of Ambroxol, such as bromhexine or of a compound having the ability to rearrange a lysosomal protein having reduced activity, include acid salts, for example sodium, potassium, calcium, magnesium and

tetraalkylammonium and the like, or mono- or di- basic salts with the appropriate acid, for example organic carboxylic acids such as acetic, lactic, tartaric, malic, isethionic, lactobionic and succinic acids; organic sulfonic acids such as methanesulfonic, ethanesulfonic, benzenesulfonic and p-toluenesulfonic acids and inorganic acids such as hydrochloric, sulfuric, phosphoric and sulfamic acids and the like. Some of the compounds of this invention may be crystallized or recrystallized from solvents such as aqueous and organic solvents. In such cases solvates may be formed. The present invention encompasses within its scope stoichiometric solvates including hydrates as well as compounds containing variable amounts of water that may be produced by processes such as lyophilisation.

In an embodiment of the method for preparing a pharmaceutical preparation of the present invention the method comprises the step of formulating as a first constituent a compound having the ability to rearrange a lysosomal protein, and as a second constituent Ambroxol and/or a derivative of Ambroxol either into a single dosage form or into two separate dosage forms. In an embodiment thereof a first of the two separate dosage forms contains the first constituent and a second of the two separate dosage forms contains the second constituent. In a further embodiment thereof the first constituent and the second constituent are formulated into one single dosage form.

In an embodiment of the pharmaceutical preparation according to the present invention a dosage form is selected from the group comprising tablets, capsules, powder, mixture, effervescence tablets and solutions.

It will be immediately understood by a person skilled in the art that in an embodiment of the method for preparing a pharmaceutical preparation of the present invention, wherein the first constituent and the second constituent are formulated into a single dosage form both the first constituent and the second constituent are comprised, for example, in a single tablet, a single capsule, a single powder, a single mixture, a single effervescence tablet or a single solution. Accordingly, in an embodiment of the method for preparing a pharmaceutical preparation, wherein a first of the two separate dosage forms contains the first constituent and a second of the two separate dosage forms contains the second constituent the first constituent is formulated into a first dosage form and the second constituent is formulated into a second dosage form, wherein preferably the first dosage form and the second dosage form each and

individually are selected from the group comprising tablets, capsules, powder, mixture, effervescence tablets and solutions. Accordingly, the first constituent is, for example, formulated into a first dosage form, wherein the dosage form is a tablet and the second constituent is formulated into a second dosage form, wherein the second dosage form is a tablet, a capsule, a powder, a mixture, an effervescence tablet or a solution. It is within the present invention that the first constituent and the second constituent of the composition of the invention may be contained in the same dosage form, whereby the first constituent and the second constituent are contained in the same physical entity of such dosage form, for example in the same tablet or the same solution; in such embodiment the first and the second constituent can be admixed with each other or can be physically or chemically separated from each other within such physical entity.

A subject as referred to herein preferably means an individual, preferably a human, to which the combination according to the present invention and/or pharmaceutical preparation according to the present invention and/or compound having the ability to rearrange the lysosomal protein according to the present invention and/or Ambroxol and/or a derivative thereof according to the present invention is to be administered and/or is administered and/or has been administered independent from the health status of the individual and/or the genetic status of the individual. In an embodiment of the various aspects of the present invention a subject is a patient. Patient as used herein preferably means a subject, wherein the subject is suffering from or is at risk of suffering from a disease.

In an embodiment of the combination according to the present invention the combination is for use in a method of personalized therapeutic treatment of a subject. In connection therewith it has to be understood that a skilled clinician will categorize a patient's health status, such as the status of a human male, to be "ill" if, for example, a value of GLA activity is determined in a sample from said patient which is lower than a cut-off value of 5 nmol MU/mg total protein/h. It will be acknowledged by a skilled person that the decision whether or whether not the combination according to the present invention will be administered to a patient is preferably solely dependent on the genetic status of the patient, i.e. the particular mutation of the respective lysosomal protein. More particularly such method of personalized therapeutic treatment of a subject preferably comprises the following steps:

step a): determining whether in a sample of the subject the lysosomal protein has a reduced activity, preferably such reduced activity results from one or more mutation being contained in the lysosomal protein compared to the wild type lysosomal protein;

step b): identifying a compound having the ability to rearrange the lysosomal protein having reduced activity, and wherein the compound is suitable for or is increasing the reduced activity of the lysosomal protein; and

step c): administering to the subject the pharmaceutical preparation, wherein the first constituent is the compound suitable for or increasing the reduced activity of the lysosomal protein identified in step b) and wherein the second constituent is Ambroxol and/or a derivative of Ambroxol.

Accordingly, if a patient shows less than 5 nmol MU/mg/hr GLA activity said patient has to have a “responsive mutation” such as one disclosed in table 3 herein, whereas the health status of a patient having a value of GLA activity of more than 5 nmol MU/mg total protein but less than 10 nmol MU/mg total protein/h will be categorized to be “unclear”.

If, according to the invention a protein, polypeptide or peptide, including but not limited to an enzyme, is said to have a mutation, this preferably means that the amino acid sequence of the protein, polypeptide or peptide is changed, more preferably changed compared to the amino acid of the wild type amino acid sequence and/or the amino acid sequence of a healthy subject.

It is within the present invention that each and any compound disclosed herein constitutes another aspect of the present invention.

In an embodiment, the term “wherein the lysosomal protein has a reduced activity” means “wherein the lysosomal protein has reduced activity”.

As preferably used herein, reference to a range defined by an upper limited and a lower limit discloses each and any integer contained within the upper limit and the lower limit including the upper limit value/figure and the lower value/figure. For example, 1 to 6 carbon atoms

means in accordance therewith, 1 carbon atom, 2 carbon atoms, 3 carbon atoms, 4 carbon atoms, 5 carbon atoms and 6 carbon atoms.

The present invention is now further illustrated by the following figures and examples from which further features, embodiments and advantages may be taken.

More specifically:

Fig. 1 is a diagram showing GLA activity of GLA mutants as present in Fabry disease and of wild type GLA with and without administration of 40 μ M Ambroxol. The y-axis indicates the activity of GLA normalized to one hundred percent wild type (WT) GLA activity in the absence of treatment with Ambroxol or pharmacological chaperone. On the x-axis the mutation of the respective analyzed GLA mutants is indicated, the activity of which was determined with (grey bars) and without (white bars) the addition of 40 μ M Ambroxol. The underlying experiment was carried out at least 3 times as described in Example 1; statistical analysis was carried out using two-sided student's T-test. Error bars are indicated as standard error of the mean. p-values are *= $p \leq 0.05$; **= $p \leq 0.01$; ***= $p \leq 0.005$.

Fig. 2 is a diagram showing a dose-response relationship of GLA activity as a function of Ambroxol concentration. The y-axis indicates the percentage of wild type GLA activity normalized to the activity of wild-type GLA in the absence of treatment with Ambroxol or a pharmacological chaperone. The x-axis indicates the concentration of Ambroxol which was added to the respective sample. The activity of wild type GLA was determined upon addition of the depicted concentrations of Ambroxol. The underlying experiment was carried out at least 3 times as described in Example 1. Error bars are indicated as standard error of the mean.

Fig. 3A is a diagram showing GLA activity of alpha-galactosidase (GLA) mutants upon no addition of either Ambroxol or DGJ, upon addition of 20 μ M DGJ and upon addition of 20 μ M DGJ and 40 μ M Ambroxol (indicated from left to right for each mutant). The y-axis indicates the activity of GLA normalized to one hundred percent wild-type (WT) GLA activity in the absence or presence of treatment with a pharmacological chaperone or the combination of a pharmacological chaperone (DGJ) and Ambroxol. On the x-axis the mutation of the respective analyzed GLA mutants is indicated, the activity of which was

determined upon addition of 20 μ M DGJ (grey bars), addition of 20 μ M DGJ and 40 μ M Ambroxol and without addition of either Ambroxol or DGJ (white bars). The underlying experiment was carried out at least 3 times as described in Example 1. Statistical analysis was carried out using two-sided student's T-test. Error bars are indicated as standard error of the mean. p-values are *= $p \leq 0.05$; **= $p \leq 0.01$; ***= $p \leq 0.005$.

Fig. 3B is a diagram showing GLA activity of alpha-galactosidase (GLA) mutants upon no addition of either galactose or Ambroxol, upon addition of galactose and upon addition of both galactose and Ambroxol (indicated from left to right for each mutant). The y-axis indicates the activity of GLA normalized to one hundred percent wild-type (WT) GLA activity in the absence or presence of treatment with a pharmacological chaperone or the combination of a pharmacological chaperone and Ambroxol, whereby the pharmacological chaperone is the sugar galactose. On the x-axis the mutation of the respective analyzed GLA mutants is indicated, the activity of which was determined upon addition of 100 mM galactose (light grey bars), the addition of 100 mM galactose and 40 μ M Ambroxol (dark grey bars) and without the addition of either Ambroxol or galactose (white bars). The experiment has been conducted 2 times, the values are therefore given as mean $\pm$ SD.

Fig. 4 is the result of a Western Blot analysis showing protein levels of GLA in cells transfected with the indicated GLA mutants, whereby the cells were not treated (indicated by "-"), treated with DGJ (indicated by "DGJ") or treated with both DGJ and Ambroxol (indicated by "DGJ/ABX"). The Experiment was carried out as described in Example 1.

Fig. 5A is a diagram showing a dose-response relationship of the activity of mutant GLA having mutation A156V, also referred to herein as GLA[A156V], as a function of DGJ concentration in the absence of Ambroxol and at 40 μ M Ambroxol, respectively. The y-axis indicates the activity of GLA[A156V] as fold increase of GLA[A156V] activity without treatment with/addition of DGJ which was set as 1.0. On the x-axis the concentration of DGJ applied is indicated. The underlying experiment was carried out four times as described in Example 1, with and without application of 40 μ M Ambroxol, as indicated. Error bars are indicated as standard error of the mean. p-values are *= $p \leq 0.05$; **= $p \leq 0.01$; ***= $p \leq 0.005$.

Fig. 5B is a diagram showing a dose-response relationship of the activity of mutant GLA having mutation R301G, also referred to herein as GLA[R301G], as a function of DGJ concentration in the absence of Ambroxol and at 40 μ M Ambroxol, respectively. The y-axis indicates the activity of GLA[R301G] as fold increase of GLA [R301G] activity without treatment with/addition of DGJ which was set as 1.0. On the x-axis the concentration of DGJ applied is indicated. The underlying experiment was carried out four times as described in Example 1, with and without application of 40 μ M Ambroxol, as indicated. Error bars are indicated as standard error of the mean. p-values are *= $p \leq 0.05$; **= $p \leq 0.01$; ***= $p \leq 0.005$.

Fig. 6 is a diagram showing alpha-galactosidase (GLA) activity of various GLA mutants upon addition of (from left to right for each mutation) neither DGJ nor Ambroxol or Bromhexine, Bromhexine alone, DGJ alone, DGJ and Ambroxol, and DGJ and Bromhexine. The y-axis indicates the activity of GLA normalized to one hundred percent wild type (WT) GLA activity in the absence or presence of treatment with Bromhexine, pharmacological chaperone or the combination of pharmacological chaperone and Ambroxol or Bromhexine, respectively, with the pharmacological chaperone being DGJ. On the x-axis the mutation of the respective analyzed GLA mutants is indicated, the activity of which was determined (from left to right for each mutation) upon addition of neither Ambroxol nor DGJ (white bars), upon addition of 40 μ M Bromhexine (light grey bars), upon addition of 20 μ M DGJ (dark grey bars), upon addition of 20 μ M DGJ and 40 μ M Ambroxol (dark grey bars with vertical lines), and upon addition of 20 μ M DGJ and 40 μ M Bromhexine (white bars with diagonal lines). The experiment was carried out at least 3 times as described in Example 1, statistical analysis was carried out using two-sided student's T-test. Error bars are indicated as standard error of the mean. p-values are *= $p \leq 0.05$; **= $p \leq 0.01$; ***= $p \leq 0.005$. The indicated stars show the significances of treatment with DGJ alone compared to the treatment with a combination of Bromhexine and DGJ.

Fig. 7A is a diagram showing acid α -glucosidase (GAA) activity of various GAA mutants as found in Pompe disease upon addition of (from left to right for each mutation) neither Ambroxol nor NB-DNJ, Ambroxol, NB-DNJ, and NB-DNJ and Ambroxol. The y-axis indicates the activity of acid α -glucosidase normalized to one hundred percent wild type (WT) acid α -glucosidase activity in the absence or presence of treatment with Ambroxol, pharmacological chaperone or the combination of pharmacological chaperone and Ambroxol

with the pharmacological chaperone being NB-DNJ. On the x-axis the mutation of the respective analyzed acid α -glucosidase mutants is indicated, the activity of which was determined (from left to right for each mutation) upon addition of neither Ambroxol nor NB-DNJ (white bars), upon addition of 40 μ M Ambroxol (light grey bars), upon addition of 20 μ M NB-DNJ (dark grey bars), and upon addition of 20 μ M NB-DNJ and 40 μ M Ambroxol (black bars). The underlying experiment was carried out at least 3 times as described in Example 1. Error bars are indicated as standard error of the mean.

Fig. 7B is a diagram showing acid α -glucosidase activity of various acid α -glucosidase mutants upon addition of (from left to right for each mutation) neither DNJ nor Ambroxol (white bars), upon addition of 20 μ M DNJ (light grey bars), and upon addition of 20 μ M DNJ and 40 μ M Ambroxol (dark grey bars). The design is similar to Fig 7A, the iminosugar used in this experiment is DNJ instead of NB-DNJ.

Fig. 8 is a diagram showing the activity of α -galactosidase A as a function of the concentration of DGJ and as a function of the concentration of DGJ with a constant concentration of Ambroxol of 40 μ M. The y-axis indicates the activity of α -galactosidase A normalized to one hundred percent of wild-type (WT) α -galactosidase A activity in the absence of treatment with Ambroxol, pharmacological chaperone or the combination of pharmacological chaperone and Ambroxol with the pharmacological chaperone being DGJ. On the x-axis the concentration of DGJ is indicated. The Experiment was carried out at least 3 times as described in Example 1. Error bars are indicated as standard error of the mean (Microsoft, Redmont, Washington, USA). The insert in Fig. 8 shows the activity of α -galactosidase A as a function of the concentration of Ambroxol at a pH of 4.5 and at a pH of 6.7, whereby the activity is normalized activity of untreated α -galactosidase A.

Figs. 9A and 9B are diagrams showing the activity of acid α -galactosidase A in the presence of DGJ alone, Ambroxole, Bromhexine and various compounds, whereby in case of Ambroxole, Bromhexine and the various compounds, DGJ was added such that the concentration of DGJ was 20 μ M, while the concentration of Ambroxol, Bromhexine and the various compounds was (from left to right for any of Ambroxol, Bromhexine and the various compounds) 100 nM (white bars), 1 μ M (grey bars), 10 μ M (dark grey bars) and 40 μ M (light grey bars). The y-axis indicates the activity of α -galactosidase A normalized to one hundred

percent wild-type (WT) α -galactosidase A activity in the absence or presence of Ambroxol (ABX), Bromhexine (BHX) or the compounds indicated on the x-axis in combination with the pharmacological chaperone DGJ. The various compounds are derivatives of Ambroxol and Bromhexine, respectively.

Fig. 10 is a panel of diagrams showing the activity of α -galactosidase A mutant [A156V], also referred to as GLA[A156V] as a function of the concentration of Ambroxol referred to as ABX (A), Bromhexine referred to as BHX (B), compound SF-54B (C), compound SF-55C (D), compound SF-150 B (E) and compound SF-153B (F) with DGJ being added to each reaction as pharmacological chaperone. The y-axis indicates the activity of α -galactosidase A normalized to one hundred percent of wild-type (WT) α -galactosidase A activity in the absence of treatment with the pharmacological chaperone DGJ. On the x-axis the concentration of the Ambroxol, Bromhexine and the other compounds is indicated. The experiment was carried out at least 3 times (except for E and F as indicated in the figure where n was 2) as described in Example 1. Error bars are indicated as standard error of the mean (or standard deviation when $N < 3$). p-values are $*$ = $p \leq 0,05$; $**$ = $p \leq 0,01$; $***$ = $p \leq 0,005$. The indicated stars show the significances of treatment with DGJ alone compared to the treatment with a combination of ABX, BHX or the named compound and DGJ. BHX (B) and especially SF-55C (D) showed a toxic effect on the cells at a concentration of 80 μ M.

Fig. 11A is a diagram indicating acid α -glucosidase activity of acid α -glucosidase mutant Y455F as present in Pompe disease upon addition of (from left to right) neither DNJ nor any one of Bromhexine (BHX), compound SF-54C and SF-55C, upon addition of DNJ, upon addition of DNJ and Bromhexine (BHX), upon addition of DNJ and compound SF-54B, and upon addition of DNJ and compound SF-55C. The imino sugar used as a chaperone is DNJ. The design of the underlying experiment was similar to the one underlying Fig. 7A. Concentration of DNJ was 20 μ M; concentration of each and any one Bromhexine, compound SF-54B and compound SF-55C was 40 μ M. The underlying experiments were carried out at least 3 times (unless indicated otherwise). Error bars are indicated as standard error of the mean (or standard deviation when $N < 3$). p-values are $*$ = $p \leq 0,05$; $**$ = $p \leq 0,01$; $***$ = $p \leq 0,005$.

Fig. 11B is a diagram indicating acid α -glucosidase activity of acid α -glucosidase mutants Y455F and L552P as present in Pompe disease upon addition of (from left to right)

neither DNJ nor any one of compound SF-124B, compound SF-150B and SF-153B, upon addition of DNJ, upon addition of DNJ and compound SF-124B, upon addition of DNJ and compound SF-150B, and upon addition of DNJ and compound SF-153B. The imino sugar used as a chaperone is DNJ. The design of the underlying experiment was similar to the one underlying Fig. 7B. Concentration of DNJ was 20 μ M; concentration of each and any one Bromhexine, compound SF-54B and compound SF-55C was 40 μ M. The underlying experiments were carried out at least 3 times (unless indicated otherwise). Error bars are indicated as standard error of the mean (or standard deviation when $N < 3$). p-values are *= $p \leq 0,05$; **= $p \leq 0,01$; ***= $p \leq 0,005$.

Example 1: Materials and Methods

Cloning of α -galactosidase A in an overexpression vector

A bacterial clone containing the full length cDNA of α -galactosidase A (IRAU969H0320D, aligned to accession no. NM_000169.2 (GeneBank)) was received from ImaGenes GmbH, Berlin, Germany. Amplification was performed with the primers 5'-AGGTCGGATCCG ACAATGCAGCTGAGGAACC-3' (SEQ ID NO:1) and 5'-GGTGTTCTGAATTAAAGTAAGTCTTTTAATGACATCTGCA-3' (SEQ ID NO:2) introducing unique restriction sites for BamHI and BstBI. The PCR was taken out using cloned Pfu DNA polymerase (Stratagene). The cDNA was subcloned into pGEM-T Easy vector (Promega). Prior to ligation, the cDNA insert was incubated with Taq polymerase (Qiagen) to add a poly (A) overhang at the 3'- end for TA -cloning. For expression, the α -Gal A cDNA was ligated into target vector pcDNA3.1/V5-His6 (Invitrogen).

Cloning of acid α -glucosidase in an overexpression vector

A bacterial clone containing the full length cDNA of α -galactosidase A (IRATp970C0971D, aligned to accession no. NM_000152.3, GeneBank) was received from ImaGenes GmbH, Berlin, Germany. Amplification was performed with the primers 5'-TAG GAG CTG TCC AGG CCA TC-3' (SEQ ID NO:3) and 5'-GAG AGA CTA ACA CAC TCC GC -3' (SEQ ID NO:4). The PCR was carried out using iProof DNA Polymerase (BioRad). The cDNA was subcloned into pCRII TA TOPO vector (Invitrogen). Prior to ligation, the cDNA insert was incubated with Taq polymerase (Qiagen) to add a poly (A) overhang at the 3'- end for TA-

cloning. For expression, the GAA cDNA was ligated into target vector pcDNA3.1/V5-His6 (Invitrogen) using sticky end ligation utilizing unique restriction sites (HindIII, XhoI) offered by both vectors multiple cloning sites.

Site-directed mutagenesis of α -galactosidase A

Expression vectors containing α -Gal A and acid glucosidase mutations were generated by site-directed PCR mutagenesis (Andreotti et al., *Orphanet Journal of Rare Diseases* 2011, 6:66) using the QuikChange® II XL Site-Directed Mutagenesis Kit (Stratagene). Nucleotide exchanges, deletions or insertions were individually introduced by PCR amplification using PfuUltra DNA polymerase, the pcDNA3.1/GLA and pcDNA3.1/GAA plasmid vector containing the wild type sequence of the named genes were used as template and a 27-33-mer primer set, with sense and antisense primers harbouring one of the respective sequence modifications in the middle of their sequence. Each obtained mutant plasmid was subjected to sequencing analysis on a 3130 xl Genetic Analyzer (Applied Biosystems)

Cell culture

HEK293H cells were maintained in DMEM (Dulbecco's Modified Eagle Medium, GIBCO - 31966) supplemented with 10% FBS (fetal bovine serum; PAA - A11-151) and 1% penicillin/streptomycin (GIBCO - 15140). All cells were incubated in a water-jacket incubator (Binder, Tuttlingen, GERMANY) at 37 °C under 5% CO₂. DGJ, Ambroxol and Bromhexine were received from Sigma Aldrich (Munich, GERMANY) and added to the culture medium from an aqueous stock solution (10mM). Bromhexine was obtained from Sigma Aldrich (Bromhexine Hydrochloride 17343 - $\geq 98.0\%$).

Transient expression of mutant enzymes in HEK293H cells

1.5×10^5 cells were seeded 24 hours before transfection in each well of a 24 well culture plate using 500 μ l DMEM medium (GIBCO) supplemented with 10 % Fetal bovine serum (PAA). Transient expression of mutant enzymes in HEK293H cells was carried out using Lipofectamine 2000 transfection reagent (Invitrogen) according to the manufacturer's protocol. Typically, prior to transfection, a mixture of plasmid DNA (0.8 μ g) and Lipofectamine 2000 transfection reagent (2 μ l) in 100 μ l of serum-free DMEM or Opti-MEM medium (GIBCO) was incubated at room temperature (20-25 °C) for 20 min and applied to the cells thereafter. The cells were subsequently incubated for 6 hours at 37 °C, the medium

containing the transfection reagent was removed and 500 μ l fresh DMEM was added. In this step, the compounds were added where intended. The cells were incubated for another 48 hours and harvested

Enzymatic measurement of α -galactosidase A

Typically, 48 hrs after transfection with plasmid vectors containing the GLA cDNA carrying the respective mutation, cells were homogenized in 200 μ l bidistilled water and subjected to 5 freeze-thaw cycles using liquid nitrogen. The supernatant collected after centrifugation of the homogenate at 10000 x g for 5 min was used in enzyme assays. Protein concentration was measured with the BCA protein assay kit (Thermo Scientific) according to the manufacturer's manual. 10 μ l of the cell lysates with a concentration of 50 μ g/ml were assayed with 20 μ l of 4MU- α -D-galactopyranoside (4MU- α -Gal, 2 mM) in 0.06 M phosphate citrate buffer (pH 4.7) with some adaptations from the original method described by Desnick et al. (J Lab. Clin Med 81: 157, 1973). Enzyme reactions were terminated after 1 hr of incubation at 37°C by the addition of 0.2 ml of 1.0 M glycine buffer (pH 10.5), prepared by adjusting the pH using 1.0 M NaOH. The released 4MU was determined by fluorescence measurement at 360 and 465 nm as the excitation and emission wavelengths respectively in a micro plate fluorescence reader (Tecan, Männedorf, SWITZERLAND). The measured enzyme activity was first calculated as nm MU/mg protein/hr and normalized to one hundred percent wild-type activity. Experiments were conducted at least 3 times, statistical analysis was carried out using two-sided student's T-test. Error bars are indicated as standard error of the mean using the Excel Software (Microsoft, Redmont, Washington, USA). p-values are *= $p \leq 0.05$; **= $p \leq 0.01$; ***= $p \leq 0.005$

Enzymatic measurement of α -glucosidase

Sample preparation was carried out analogously to the α -galactosidase A recordings. As substrate, a 1.3mg 4-Methylumbelliferyl- α -D-Glucopyranoside/ml solution (in sodium acetate buffer, pH 4.0) was used and 5 μ g sample/reaction were added. The reaction was incubated for 60 minutes at 37°C and stopped with 1M Glycine-NaOH, pH 10.5. The released 4MU was determined by fluorescence measurement at 360 and 465 nm as the excitation and emission wavelengths respectively in a micro plate fluorescence reader (Tecan, Männedorf, SWITZERLAND). The measured enzyme activity was first calculated as nm MU/mg protein/hr and normalized to one hundred percent wild-type activity. Experiments were

conducted at least 3 times, statistical analysis was carried out using two-sided student's T-test. Error bars are indicated as standard error of the mean using the Excel Software (Microsoft, Redmont, Washington, USA). p-values are *= $p \leq 0.05$; **= $p \leq 0.01$; ***= $p \leq 0.005$

Western blot analysis

Western blot analysis for the detection of α -Gal A protein was performed using a commercially available rabbit anti- α -Gal A polyclonal antibody ([H-104] Santa Cruz) Furthermore, a mouse GAPDH monoclonal antibody ([6C5] abcam) was used as an internal loading control.

HEK293H cell lysates were generated by aspirating the media off the 24 well culture plates, washing the cells once with 1 x PBS (Biochrom AG) and applying 200 μ l ice cold RIPA buffer supplemented with protease inhibitor cocktail tablets (Roche) to the cells prior to a 20 minute incubation on ice. The cells were then rinsed off from the wells, transferred to micro centrifuge tubes and spun at 14000 x g for 10 minutes at 4°C to pellet the cell debris. The supernatant was used for the analysis. After the determination of protein concentration, 50 μ g protein were mixed with a suitable volume of 5 x Laemmli loading buffer, boiled for 5 minutes on a thermo shaker (Eppendorf), centrifuged at 14000 x g for 10 minutes at 4°C and loaded on a Criterion precast 4-15% Tris-HCl gel (Bio-Rad). Proteins were transferred electrophoretically to a Nitrocellulose (Amersham Hybond ECL) membrane (GE Healthcare). The membrane was blocked with 5% (w/v) non-fat dried skimmed milk in TBS-Tween 20 [10 mM Tris/HCl (pH 7.5) with 150 mM NaCl and 0.1% Tween 20] at room temperature for 1 hour, and then treated with a primary antibody against GAPDH diluted 1:10000 in a milk/blot solution [3% (w/v) non-fat dried skimmed milk in TBS-Tween 20] at 4°C overnight. After the blot was washed three times with excess TBS-Tween 20, the membrane was treated for 1 h at room temperature with a primary antibody against α -Gal A diluted 1:500 in the 3% milk/blot solution. After another wash procedure, a secondary antibody mix of an Alexa Fluor labeled 680 anti-rabbit IgG antibody produced in goat (Molecular Probes) and an IRDye800 conjugated anti-mouse IgG antibody produced in goat (Rockland) both diluted 1:10000 in the 3% milk/blot solution was applied to the membrane. Following extensive washing with TBS-Tween 20, protein bands were visualized with an Odyssey Infrared Imager (Li-Cor) and quantified using the Odyssey software.

Inhibition of agalsidase alfa (Replagal[®], Shire Pharmaceuticals)

Replagal was diluted to 10nM in citrate-phosphate-buffer (0.06 M) adjusted to different pH-values (4.5; 6.7). Subsequently, different concentrations of the compounds (DGJ, Ambroxol) were added ranging from 0-1mM. Finally, the substrate 4-methylumbelliferyl- α -D-galaktopyranosid (1 mM) was added to start a 15- minute incubation in a water bath at 37°C. The reaction was stopped by the addition of 200 μ l glycine-NaOH-buffer (1M, pH 10.5). Fluorescence measurement was carried out in a micro plate fluorescence reader (Tecan, Männedorf, SWITZERLAND) according to the method above.

Example 2: Effect of Ambroxol on GLA enzyme activity

Various GLA mutants indicated in Fig. 1 were prepared as described in Example 1. To the enzyme preparation for the individual mutant 40 μ M Ambroxol was given. The results are indicated in Fig. 1.

Fig. 1 is a diagram indicating GLA enzymatic activity for wild type GLA and indicated mutant GLA forms with and without Ambroxol. The activity was normalized with the activity observed for wild type GLA having been set as 100 %.

The experiment was conducted as described in Example 1 at least 3 times; statistical analysis was carried out using two-sided student's T-test. Error bars are indicated as standard error of the mean using the Excel Software (Microsoft, Redmont, Washington, USA). p-values are *= $p \leq 0.05$; **= $p \leq 0.01$; ***= $p \leq 0.005$

As may be taken from Fig. 1, the addition of Ambroxol resulted in an increase in enzyme activity of the GLA of the wild type. In contrast thereto, Ambroxol had no effect on the mutant forms of GLA.

The present inventors used the system established by Asano et al. (Asano et al., 2000 supra) and could reproduce the results with regard to the enhancing effect of DGJ when administered alone. More importantly, the present inventors also found that GLA is not inhibited by Ambroxol in said cell free assay.

The following table 3 lists the responsiveness of overexpressed proteins having mutations resulting in Fabry's disease with regard to therapy with DGJ alone, as well as the additional effect in terms of enhancing enzyme activity when applying Ambroxol in accordance with the present invention.

Table 3: List of DGJ responsive GLA mutants.

No.	mutation	Reference for positive testing of chaperone (DGJ) effect	Tested in-house	Non-responder (own data)	additive Ambroxol effect demonstrated
1	A20P	Ishii et al., 2007			
2	N34S	Benjamin et al., 2008			
3	P40S	Benjamin et al., 2008			
4	T41I	Benjamin et al., 2008, Shin et al., 2008			
5	H46P		X		
6	R49C	Shin et al., 2008	X	X	
7	R49L				
8	M51K	Benjamin et al., 2008, Shin et al., 2008	X		(*)
9	M51I		X		**
10	E59K	Ishii et al., 2007, Benjamin et al., 2008	X		****
11	S65I		X		(*)
12	E66Q	Ishii et al., 2007, Benjamin et al., 2008			
13	M72V	Ishii et al., 2007			
14	A73V		X		(*)
15	I91T	Ishii et al., 2007, Benjamin et al., 2008			

16	W95S	Benjamin et al., 2008			
17	A97V	Ishii et al., 2007, Benjamin et al., 2008, Shin et al., 2007, 2008			
18	R100K	Benjamin et al., 2008			
19	R112C	Benjamin et al., 2008, Shin et al., 2007, 2008	X	X	
20	R112H	Ishii et al., 2007, Benjamin et al., 2008, Shin et al., 2007, 2008	X		(*)
21	F113L	Ishii et al., 2007, Benjamin et al., 2008			
22	L120V		X		
23	S126G		X		
24	D136E		X		
25	N139S		X		
26	A143T	Benjamin et al., 2008, Shin et al., 2007, 2008	X		
27	G144V	Benjamin et al., 2008			
28	P146S	Ishii et al., 2007			
29	S148N	Benjamin et al., 2008			
30	A156V	Ishii et al., 2007	X		****
31	D165H		X		
32	L166V	Ishii et al., 2007			
33	D170V	Benjamin et al., 2008			
34	C172Y	Benjamin et al., 2008	X	X	
35	D175N		X		
36	G183D	Benjamin et al., 2008			
37	G183V		X		
38	S201F	Shin et al. 2008			
39	P205R	Benjamin et al., 2008			
40	P205T	Benjamin et al., 2008,			

		Shin et al., 2008			
41	Y207C	Benjamin et al., 2008			
42	Y207S	Benjamin et al., 2008, Shin et al., 2008			
43	N215S	Ishii et al., 2007, Benjamin et al., 2008, Shin et al., 2008	X		
44	I219T		X		
45	R220Q		X		
46	N224S		X		
47	H225D		X		
48	H225R		X		**
49	I232T		X		(*)
50	S235C	Benjamin et al., 2008			
51	S238N		X		
52	I242N		X		
53	D244N	Benjamin et al., 2008			
54	P259R	Shin et al., 2008			
55	N263S	Benjamin et al., 2008			
56	D264Y		X		
57	L268S		X		
58	V269M		X		
59	V269A		X		(*)
60	S276G	Benjamin et al., 2008, Shin et al. 2008			
61	Q279E	Ishii et al., 2007, Benjamin et al., 2008			
62	T282I		X		
63	W287C	Benjamin et al., 2008			
64	A288P	Benjamin et al., 2008			
65	I289F	Benjamin et al., 2008			
66	M290I		X		

67	L294S		X		
68	F295C	Shin et al., 2008			
69	M296I	Benjamin et al., 2008			
70	M296V	Ishii et al., 2007, Benjamin et al., 2008			
71	S297C		X		
72	L300P	Benjamin et al., 2008, Shin et al., 2007, 2008			
73	R301G		X		
74	R301Q	Ishii et al., 2007, Benjamin et al., 2008, Shin et al., 2008	X		****
75	R301P		X		
76	L310F		X		
77	L311V		X		
78	D313Y		X		
79	V316E	Benjamin et al., 2008			
80	V316I		X		
81	I319T		X		
82	N320I		X		
83	N320Y	Benjamin et al., 2008			
84	Q3121H		X		
85	G325D	Benjamin et al., 2008			
86	G325S		X		
87	Q327E		X		
88	G328A	Benjamin et al., 2008, Shin et al., 2008	X		*
89	R342Q	Benjamin et al., 2008	X	X	
90	S345P		X		
91	R356W	Ishii et al., 2007, Benjamin et al., 2008, Shin et al., 2007 (non-	X		****

		responder)			
92	E358A	Benjamin et al., 2008			
93	E358K	Benjamin et al., 2008			
94	G360C		X		
95	G361R		X		
96	R363C	Benjamin et al., 2008			
97	R363H	Benjamin et al., 2008	X		***
98	G373D	Ishii et al, 2007			
99	G373S	Ishii et al, 2007			
100	E398A		X		
101	P409A	Benjamin et al., 2008			
102	T410I		X		*
103	L415F		X		
104	E418G		X		

All mutants were tested with regard to DGJ-responsiveness as referenced in table 3. No reference indicates the mutation is exclusively tested in the hands of the present inventors. An additional effect of the administration of the combination according to the present invention, i.e. Ambroxol in combination with DGJ, was considered significant when * $p \leq 0.1$, ** $p \leq 0.05$, *** $p \leq 0.01$, **** $p \leq 0.005$. For some mutations the additive effect of Ambroxol has only been repeated twice resulting only in a trend of increased enzyme activity indicated by the symbol (*). In table 3 an „X“ in the column „non-responder, own data“ is indicative for mutations where the results of the experiments conducted by the instant inventors is different to at least one of the indicated references with regard to the application of DGJ alone.

Example 3: Dose-dependent efficacy of Ambroxol

In order to determine the dose-dependent efficacy of Ambroxol on the enzyme activity of wild type GLA, wild type GLA was exposed to various titers of Ambroxol.

The results are indicated in Fig. 2 which is a diagram indicating the relative activity of wild type GLA at various concentrations of Ambroxol with the activity of GLA without Ambroxol being set as 100 %.

As may be taken from Fig. 2, the enzyme activity of wild type GLA increases with increasing titers of Ambroxol. Furthermore, it is important to note that Ambroxol is not inhibiting wild type GLA. There seems to be a local maximum or potentially saturation at or beyond an Ambroxol titer of $> 60 \mu\text{M}$. The EC_{50} was determined to be $21.2 \mu\text{M}$.

In connection therewith it is important to note that in contrast to mutant GLA the activity of wild type GLA may be increased upon addition of Ambroxol which may be taken from Fig. 1 and Fig. 2.

Example 4: Synergistic effect of combined use of DGJ and Ambroxol on enzyme activity of GLA mutants

The enzyme activity of various mutant forms of GLA was determined upon exposure of the mutant forms of GLA to (a) none of DGJ and Ambroxol, (b) $20 \mu\text{M}$ DGJ, or (c) $20 \mu\text{M}$ DGJ and $40 \mu\text{M}$ Ambroxol.

The result is indicated in Fig. 3.

As may be taken from Fig. 3, all mutants of GLA which showed a beneficial effect on DGJ treatment also responded to Ambroxol when both DGJ and Ambroxol were added to the growth medium together.

A person skilled in the art will acknowledge that the assay described herein may be used to identify particularly well responding mutants. Because of this, the use of Ambroxol in combination with DGJ will be therapeutically effective/advantageous for patients responding to DGJ treatment. More precisely, for all patients DGJ treatment has been approved to enhance Gal A activity, additional treatment with Ambroxol can only be of an even higher benefit.

In connection therewith it is important to note that for all mutant enzymes tested an increased activity was measured upon application of Ambroxol in addition to the pharmacological chaperone compared to the application of the pharmacological chaperone alone. This proves the existence of a clear tendency which underlines the advantage of the combination according to the present invention compared to the application of the pharmacological chaperone alone. A person skilled in the art will thus immediately acknowledge that an increase in mutant enzyme activity upon administration of the combination according to the present invention has not to be necessarily significantly elevated compared to the administration of the pharmacological chaperone alone as, on the one hand, in clinical application the enhancement of mutant enzyme activity to only a low degree may already result in preventing symptoms or disease as has been outlined above. On the other hand, a person skilled in the art will also acknowledge that even if the application of the combination of the present invention is not leading to significantly higher activities of mutant enzymes, the combination of the present invention may still be advantageous over applying the pharmacological chaperone in that the significant costs usually related to pharmacological chaperone therapy may be lowered if the additional application of Ambroxol according to the present invention can substitute for a certain amount of pharmacological chaperone applied, still achieving the same or similar clinical benefit for the patient treated.

A Western Blot analysis was performed as described in Example 1 and the result is indicated in Fig. 4. As may be taken therefrom, the stabilizing effect of DGJ is enhanced/improved by Ambroxol. Ambroxol contributes to the higher amount of a Gal A in the cell. While being stabilized the enzyme is no longer depleted from the cells by the endoplasmic reticulum degradation machinery. The higher amount of protein/enzyme accompanies/correlates with the higher activity measured.

It can be taken therefrom that the application of the combination according to the present invention compared to application of DGJ alone leads to a higher amount of Gal A in the respective cell. It will be acknowledged by a person skilled in the art that the mutation of Gal A such as the ones of Fig. 4 do not lead to reduced expression of GalA. Rather, the mutant Gal A is subject to a more rapid degradation, preferably due to misfolding of the protein. It is assumed by the present inventors that application of DGJ and/or Ambroxol has no effect on the level of protein expression of the lysosomal protein. More particularly, the combination

according to the present invention seems to counteract, like a chaperone, preferably a pharmacological chaperone, the more rapid degradation, more preferably such counteracting occurs through stabilizing the mutant protein and thus leading to a less rapid degradation of the protein. It can be taken from Figure 4 and more particularly from the elevated amount of protein level which can be detected that Ambroxol applied in combination with DGJ results in a stabilization of the mutant protein and/or results in less rapid degradation

Example 5: A combinational treatment of endogenous GLA mutants can substitute for high DGJ doses

The enzyme activity of mutant forms of GLA was determined upon exposure of the mutant forms of GLA to (a) different concentrations of DGJ and to 40 μ M Ambroxol or (b) different concentrations of DGJ and no Ambroxol as described in Example 1.

The results are indicated in Fig. 5.

More particularly, Fig. 5A shows a diagram indicating a dose-response relationship of GLA activity of GLA mutant form A156V; and Fig.5B shows a diagram indicating a dose-response relationship of GLA activity of GLA mutant form R301G.

As may be taken from Figs. 5A and B, treatment with 40 μ M Ambroxol in addition to treatment with DGJ alone enhances GLA activity.

It can immediately be seen, that an increase of 3.8-fold of GLA[A156V] activity is achieved by application of 8 μ M DGJ and 40 μ M Ambroxol, whereas treatment with as much as 20 μ M DGJ has to be applied in order to reach the same increase in mutant enzyme activity without the additional treatment of Ambroxol, which is emphasized by the horizontal and vertical dotted line.

In other words, approximately half of the concentration of DGJ, namely 8.9 μ M, has to be applied, if 40 μ M Ambroxol are applied in addition, to reach the same effect as if 20 μ M DGJ

is applied alone. The absolute measures of said assay are approximately 300 nmol MU/mg total protein/h.

That there is a synergistic effect which arises from the combined use of both DGJ and Ambroxol is evident from the fact that alpha-galactosidase A activity is increased up to 8-fold in A156V mutant (Fig. 5A) when both 20 μ M DGJ and 40 μ M Ambroxol are administered, whereas said activity is only increased to 4-fold of untreated GLA[A156V] activity in case of 20 μ M DGJ administration only.

It can be immediately seen, that a 4.1-fold increase of GLA[R301G] activity is achieved by application of 20 μ M DGJ and 40 μ M Ambroxol, whereas treatment with 20 μ M DGJ without additional treatment with Ambroxol results only in an increase of GLA activity of 2.8-fold which is emphasized by the horizontal and vertical dotted line.

In connection therewith it is important to note that the absolute value of GLA activity was determined to be 3.7 ± 0.4 nmol MU/mg total protein/h. A skilled clinician will categorize a patient's health status, such as the status of a human male, to be "ill" if a value of GLA activity is determined in a sample from said patient which is lower than a cut-off value of 5 nmol MU/mg total protein/h. It will be acknowledged by a skilled person that the decision whether or whether not the combination according to the present invention such as ABX/DGJ will be administered to a patient is preferably solely dependent on the genetic status of the patient, i.e. the particular mutation of the respective lysosomal protein. More particularly, method of personalized therapeutic treatment of a subject would comprise the following steps:

step a): determining whether in a sample of the subject the lysosomal protein has a reduced activity, preferably such reduced activity results from one or more mutation being contained in the lysosomal protein compared to the wild type lysosomal protein;

step b): identifying a compound having the ability to rearrange the lysosomal protein having reduced activity, and wherein the compound is suitable for or is increasing the reduced activity of the lysosomal protein; and

step c): administering to the subject the pharmaceutical preparation, wherein the first constituent is the compound suitable for or increasing the reduced activity of the lysosomal protein identified in step b) and wherein the second constituent is Ambroxol and/or a derivative of Ambroxol.

Accordingly, if a patient shows less than 5 nmol MU/mg/hr GLA activity said patient has to have a “responsive mutation” such as one disclosed in table 3 herein, whereas the health status of a patient having a value of GLA activity of more than 5 nmol MU/mg total protein but less than 10 nmol MU/mg total protein/h will be categorized to be “unclear”.

In other words, the present data demonstrate that applying DGJ and Ambroxol in combination according to the present invention is able to raise the activity of mutant protein, as measured in the cell-culture system used herein, to a level which would prompt a skilled clinician to evaluate the patient’s health status as “healthy”, i.e. having an activity of mutant protein which usually does not result in symptoms otherwise caused by the mutant protein’s reduced activity, whereas the treatment with DGJ alone results in protein activity of the mutant protein which would prompt the skilled clinician to evaluate the patient’s health status as being „unclear“.

It is important to note that in the herein described cell-culture system, a concentration of DGJ as low as 4.72µM if applied in combination with 40µM Ambroxol is sufficient to elevate the mutant protein activity to a level which is only reached if as much as 20µM DGJ is applied, if DGJ is applied alone. In other words, this means that approximately only 25% of the concentration of DGJ has to be applied if applied in combination with 40µM Ambroxol compared to the concentration of DGJ which has to be applied to reach the same increase of protein activity if DGJ is applied alone.

From this follows that the combination of the present invention is particularly useful in the treatment of those LSDs and more specifically those forms of Fabry disease involving mutant forms of GLA where the chaperone and more specifically DGJ alone is not sufficiently effective.

Example 6: Similar effects for Ambroxol and Ambroxol derivative Bromhexine

In this example the effect on enzyme activity of mutant forms of GLA of Ambroxol and its derivative Bromhexine when combined with DGJ was compared.

The experiment was conducted as described in Example 1.

The results are shown in Fig. 6.

As is evident from Fig. 6 Ambroxol and Bromhexine perform similarly on the mutant GLAs. In accordance with the results obtained by treatment with Ambroxol alone, see Example 2 herein, no effect could be seen by a treatment using bromhexine alone; and treatment with the chaperone DGJ alone increased alpha-Galactosidase A activity. However, upon treatment with a combination of chaperone DGJ and Ambroxol or a combination of chaperone DGJ and Bromhexine, the enzyme activity of mutant forms of GLA could be significantly increased.

As may be taken from Fig. 6 the treatment with DGJ alone always lead to a highly significant increase in protein, i.e. enzyme activity compared to “no treatment”. There was no significant difference between protein activity measured after application of a combination of Ambroxol and DGJ compared to protein activity measured after application of a combination of Bromhexine and DGJ. Protein activity measured after application of a combination of Ambroxol and DGJ compared to protein activity measured after application of DGJ alone was significant for the mutant proteins having mutations E59K, A156V (see also Fig. 3) and G328A.

Example 7: Similar effects for Ambroxol and NB-DNJ in Pompe disease

In this example the effect on enzyme activity of mutant forms of acid α -glucosidase of Ambroxol and its derivative Ambroxol when combined with NB-DNJ were compared.

The experiment was conducted as described in Example 1.

The results are indicated in Fig. 7.

As is evident from Fig. 7 and analogous to what has been outlined above in connection with the combination according to the present invention and Fabry's disease (see Figs. 1 to 6) the application of the combination according to the present invention enhances the activity of the respective mutant protein in Pompe's disease compared to the application of the pharmacological chaperone alone. More specifically, the application of 20 μ M NB-DNJ in combination with the application of 40 μ M Ambroxol enhanced the activity of acid α -glucosidase compared to the application of 20 μ M NB-DNJ alone.

More particularly, the enzyme activity of mutations Y455F, P545L and L552P was significantly increased by NB-DNJ and the combined treatment according to the present invention using both NB-DNL and Ambroxol. Said mutations showed a tendency for higher activities when additionally treated with Ambroxol. However, only L552P showed significance after 4 experiments.

Most importantly, Y575S showed a clear tendency to be responsive only towards the combination of the present invention.

It is important to note that Ambroxol administered alone is not leading to an enhancement of the activity of mutant acid α -glucosidase.

Example 8: Inhibitory effect of DGJ is not affected by Ambroxol

Wild type α -galactosidase A (Replagal[®], agalsidase alfa) was incubated with different concentrations of DGJ and with either no or a constant concentration of 40 μ M Ambroxol. The result is shown in Figure 8.

More particularly, Fig. 8 shows that the inhibitory effect of DGJ on enzyme activity remains unchanged by Ambroxol.

It can be taken therefrom that Ambroxol has no effect on the inhibition of WT α -galactosidase A by DGJ. The IC_{50} was determined to be unchanged in both applications (0.07 μ M). Inlay diagram (upper right) shows the effect of addition of Ambroxol only at two different pH values (pH 4.5 and pH 6.7). No reduction of the enzyme activity could be detected.

It may be taken therefrom that Ambroxol alone displays neither an inhibitory nor a stimulating effect on the wild type enzyme.

Example 9: Effect of candidate compounds on the activity of α -galactosidase A mutant A156V

Different compounds which are also referred to herein as candidate compounds or derivatives of Ambroxol and Bromhexine, respectively, were tested in combination with DGJ as to their effect on α -galactosidase A mutant A156V activity.

The results are shown in Fig. 9A, Fig. 9B and Fig. 10.

More particularly, Figs. 9A and 9B are diagrams indicating the activity of acid α -galactosidase A mutant A156V; the y-axis indicated the activity of α -galactosidase A mutant A156V normalized to one hundred percent wild-type (WT) α -galactosidase A activity in the absence or presence of treatment with Ambroxol (ABX), bromhexine (BHX) or the various compounds as indicated on the x-axis in combination with 20 μ M the pharmacological chaperone DGJ.

It can be taken therefrom that particularly compounds SF-54B and SF-55C show a high increase in mutant enzyme activity when applied in combination with DGJ, whereby the concentration of SF-54B and SF-55C is 10 μ M and 40 μ M, respectively. Most important, application of compound SF-80 when applied at a concentration as low as 1 μ M in combination with DGJ resulted in an increase in mutant enzyme activity as high as if Ambroxol was applied at a concentration of up to 10 μ M in combination with DGJ.

Fig. 10 is a panel of diagrams showing the activity of α -galactosidase A mutant A156V. The y-axis indicates the activity of α -galactosidase A normalized to one hundred percent of wild-type (WT) α -galactosidase A activity in the absence of treatment with pharmacological chaperone DGJ and increasing concentrations of Ambroxol (ABX), Bromhexine (BHX) or the compounds indicated on the x-axis in combination with the pharmacological chaperone DGJ. On the x-axis the concentration of the compound is indicated. The experiment was carried out at least 3 times as described in Example 1 (except for Figs. 10E and 10F, as indicated in the figure). Error bars are indicated as standard error of the mean (or standard deviation when $N < 3$) using the Excel Software (Microsoft, Redmont, Washington, USA). p-values are $*$ = $p \leq 0,05$; $**$ = $p \leq 0,01$; $***$ = $p \leq 0,005$. The indicated stars show the significances of treatment with DGJ alone compared to the treatment with a combination of ABX, BHX or the named compound and DGJ. BHX (B) and especially SF-55C (D) showed a toxic effect on the cells in a concentration of $80 \mu\text{M}$.

It can be taken therefrom that compounds SF-54B, SF-55C, SF-150B and SF153B show a high increase in mutant enzyme activity when applied in combination with DGJ.

Example 10: Effect of candidate compounds effect on the activity of α -galactosidase A mutants Y455F and L552P

Different compounds which are also referred to herein as candidate compounds or derivatives of Ambroxol and Bromhexine, respectively, were tested in combination with DNJ for the effect on the activity of α -galactosidase A mutants Y455F and L552P, respectively.

The results are shown in Fig. 11A and Fig. 11B.

Figs. 11A and 11B are diagrams indicating the activity of acid α -glucosidase. The design is similar to Fig 7A and 7B, the iminosugar used in these experiments is DNJ. The diagram in figure 11A shows the effect on mutant acid α -glucosidase activity after addition of $20 \mu\text{M}$ DNJ alone and $20 \mu\text{M}$ DNJ with $40 \mu\text{M}$ Bromhexine (BHX) or a derivative compound thereof. The diagram in Fig. 1 11B shows the effect on mutant acid α -glucosidase after addition of $20 \mu\text{M}$ DNJ alone and $20 \mu\text{M}$ DNJ with $40 \mu\text{M}$ of three Ambroxol (ABX)-like compounds. The experiments were carried out at least 3 times (unless otherwise indicated). Error bars are

indicated as standard error of the mean (or standard deviation when $N < 3$) using the Excel Software (Microsoft, Redmont, Washington, USA). p-values are $*$ = $p \leq 0,05$; $**$ = $p \leq 0,01$; $***$ = $p \leq 0,005$.

It can be taken therefrom that the increase in the activity of acid α -glucosidase of mutant Y455F is comparable when DNJ is applied alone or when Bromehexine, compounds SF-54B, SF-55C and SF-124B, respectively, are applied in combination with DNJ. Most importantly, the increase in the activity of mutant Y455F is higher when DNJ is applied in combination with SF -150B and SF-153B compared to the effect arising from DNJ only.

It can also be taken therefrom that the increase in the activity of mutant L552P is higher when compounds SF-124B, SF-150B and SF-153B, respectively are applied in combination with DNJ compared to the effect arising from DNJ only.

The features of the present invention disclosed in the specification, the claims, the sequence listing and/or the drawings may both separately and in any combination thereof be material for realizing the invention in various forms thereof.

Claims

1. A combination comprising a first constituent and a second constituent,
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, and
wherein the second constituent is Ambroxol and/or a derivative of Ambroxol.
2. The combination according to claim 1, wherein the lysosomal protein is affected in a disease.
3. The combination according to any one of claims 1 to 2, wherein the lysosomal protein is selected from the group comprising an enzyme, a trans-membrane protein or a soluble non-enzymatic protein.
4. The combination according to claim 3, wherein the lysosomal protein is an enzyme, wherein the enzyme has a reduced activity, preferably a reduced activity due to a mutation of the enzyme.
5. The combination according to claim 4, wherein the enzyme is selected from the group comprising a lysosomal hydrolase and a phosphotransferase.
6. The combination according to any one of claims 1 to 5, wherein the compound having the ability to rearrange a lysosomal protein is a chaperon.
7. The combination according to claim 6, wherein the chaperon is a pharmacological chaperon or a pharmaceutically acceptable salt, solvate or derivative thereof.
8. The combination according to claim 7, wherein the pharmacological chaperon is a sugar and/or an imino sugar or a pharmaceutically acceptable salt, solvate or derivative of the sugar and/or the imino sugar.

9. The combination according to claim 9, wherein the sugar is galactose, preferably D-galactose.
10. The combination according to claim 8, wherein the imino sugar is selected from the group comprising 1-deoxygalactonojirimycin (DGJ), alpha-galactohomonojirimycin, alpha-allo-homonojirimycin, beta-1-C-butyl-deoxygalactonojirimycin, beta-1-C-butyl-deoxynojirimycin, N-nonyl-deoxynojirimycin (NN-DNJ), N-octyl-2,5-anhydro-2,5-imino-D-glucitol, N-octyl-isofagomine, N-octyl-beta-valienamine (NOV), Isofagomine (IFG), calystegine A3, calystegine B1, calystegine B2, calystegine C1, 1,5-dideoxy-1,5-iminoxylitol (DIX), alpha-1-C-nonyl-DIX, alpha-1-C-octyl-1-DNJ, N-acetyl-glucosamine-thiazoline (NGT), 6-acetamido-6-deoxycastanospermine (ACAS), bisnaphthalimide nitro-indan-1-one, pyrrolo[3,4-d]pyridazin-1-one, pyrimethamine (PYR), N-acetyl-4-epi-beta-valienamine (NOEV), N-butyl-DNJ, Deoxynojirimycin (DNJ), N-Acetyl-galactosamine (GalNAc), 2-Acetamido-1,2-dideoxynojirimycin (AdDNJ), N-dodecyl-DNJ, 6-nonyl-isofagomine, N-methyl calystegine A₃, calystegine B₂, 4-epi-isofagomine, , 1-deoxynojirimycin, alpha-homonojirimycin, castanospermine, 1-deoxymannojirimycin, Swainsonine, Mannostatin A, 2-hydroxy-isofagomine, 1-deoxyfuconojirimycin, beta-homofuconojirimycin, 2,5-imino-1,2,5-trideoxy-L-glucitol, 2,5-dideoxy-2,5-imino-D-fucitol, 2,5-imino-1,2,5-trideoxy-D-altritol, 1,2-dideoxy-2-N-acetamido-nojirimycin, 1,2-dideoxy-2-N-acetamidogalactonojirimycin, 2-N-acetylamino-isofagomine, 1,2-dideoxy-2-acetamidonojirimycin, nagastain, 2-N-acetamido-isofagomine, 1,2-dideoxy-2-acetamidonojirimycin, 1-deoxyduronojirimycin, 2-carboxy-3,4,5-trideoxypiperidine, 6-carboxy-isofagomine, 2,6-dideoxy-2,6-imino-sialic acid, Siastin B and Castanospermine (CAS), and derivatives thereof; and pharmaceutically acceptable salts thereof.
11. The combination according to any one of claims 1 to 10, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

12. The combination according to any one of claims 1 to 11, wherein at least one of the first constituent and the second constituent is/are present as a solvate or a pharmaceutically acceptable salt thereof.
13. The combination according to claim 12, wherein the second constituent is Ambroxol, wherein Ambroxol is present as a pharmaceutically acceptable salt of Ambroxol, and wherein, preferably, the pharmaceutically acceptable salt of Ambroxol is Ambroxol hydrochloride.
14. The combination according to claim 12, wherein the second constituent is a derivative of Ambroxol, wherein the derivative of Ambroxol is bromhexine, and wherein preferably the pharmaceutically acceptable salt of bromhexine is bromhexine hydrochloride.
15. The combination according to any one of claims 1 to 14, wherein the combination is suitable for or is for use in the treatment and/or prevention of a disease.
16. The combination according to any one of claims 1 to 15, wherein the combination is a pharmaceutical combination.
17. The combination according to any one of claims 1 to 16, wherein the combination is suitable for or is for use in a method for the treatment of a subject comprising the administration of the combination to the subject.
18. The combination according to any one of claims 15 to 17, wherein the disease is a lysosomal storage disease.
19. The combination according to claim 18, wherein the lysosomal storage disease is a lysosomal storage disease having a defective degradation of a sphingolipid component.
20. The combination according to claim 19, wherein the lysosomal storage disease having a defective degradation of a sphingolipid component is selected from the

group comprising Fabry disease, Gaucher disease type I, Gaucher disease type II, Gaucher disease type III, acid sphingomyelinase deficiency, Farber disease, GM1 gangliosidosis type I, GM1 gangliosidosis type II, GM1 gangliosidosis type III, GM2 gangliosidosis, Krabbe disease, Metachromatic leukodystrophy type I, Metachromatic leukodystrophy type II and Metachromatic leukodystrophy type III.

21. The combination according to claim 19, wherein the lysosomal storage disease having a defective degradation of a sphingolipid component is selected from the group consisting of Fabry disease, acid sphingomyelinase deficiency, Farber disease, GM1 gangliosidosis type I, GM1 gangliosidosis type II, GM1 gangliosidosis type III, GM2 gangliosidosis, Krabbe disease, Metachromatic leukodystrophy type I, Metachromatic leukodystrophy type II and Metachromatic leukodystrophy type III.
22. The combination according to claim 20 or claim 21, wherein the lysosomal storage disease is Fabry disease.
23. The combination according to claim 20 or claim 21, wherein the acid sphingomyelinase deficiency is selected from the group comprising Niemann-Pick type A and type Niemann-Pick B.
24. The combination according to claim 20 or claim 21, wherein the GM2 gangliosidosis is selected from the group comprising Tay-Sachs type I, Tay-Sachs type II, Tay-Sachs type III and Sandhoff.
25. The combination according to claim 24, wherein the lysosomal storage disease is Tay-Sachs type I, Tay-Sachs type II or Tay-Sachs type III.
26. The combination according to claim 24, wherein the lysosomal storage disease is Sandhoff.
27. The combination according to claim 20 or claim 21, wherein the lysosomal storage disease is GM1 gangliosidosis type I, GM1 gangliosidosis type II or GM1 gangliosidosis type III

28. The combination according to claim 18, wherein the lysosomal storage disease is a lysosomal storage disease having a defective metabolism of glycosaminoglycans.
29. The combination according to claim 28, wherein the lysosomal storage disease having a defective metabolism of glycosaminoglycans is selected from the group comprising MPS I, MPS II, MPS III, MPS IV, MPS VI, MPS VI and MPS IX.
30. The combination according to claim 29, wherein MPS I is selected from the group comprising Hurler disease, Hurler-Scheie syndrome and Scheie syndrome.
31. The combination according to claim 29, wherein MPS II is selected from the group comprising Hunter syndrome.
32. The combination according to claim 29, wherein MPS III is selected from the group comprising Sanfilippo syndrome Type A, Sanfilippo syndrome Type B, Sanfilippo syndrome Type C and Sanfilippo syndrome Type D.
33. The combination according to claim 29, wherein MPS IV is selected from the group comprising Morquio type A and Morquio type B.
34. The combination according to claim 29, wherein MPS VI is selected from the group comprising Maroteaux-Lamy.
35. The combination according to claim 29, wherein MPS VII is selected from the group comprising Sly.
36. The combination according to claim 29, wherein MPS IX is selected from the group comprising Hyaluronidase deficiency and Multiple Sulfatase deficiency.
37. The combination according to claim 18, wherein the lysosomal storage disease is a lysosomal storage disease having a defective degradation of a glycan portion of a glycoprotein.

38. The combination according to claim 37, wherein the lysosomal storage disease having a defective degradation of a glycan portion of a glycoprotein is selected from the group comprising Aspartylglucosaminuria, Fucosidosis type I, Fucosidosis type II, Mannosidosis, Sialidosis type I and Sialidosis type II.
39. The combination according to claim 18, wherein the lysosomal storage disease is a lysosomal storage disease having a defective degradation of glycogen.
40. The combination according to claim 39, wherein the lysosomal storage disease having a defective degradation of glycogen is selected from the group comprising Pompe's disease.
41. The combination according to claim 40, wherein the lysosomal storage disease is Pompe's disease.
42. The combination according to claim 18, wherein the lysosomal storage disease is a lysosomal storage disease having a defective degradation of polypeptides.
43. The combination according to claim 42, wherein the lysosomal storage disease having a defective degradation of polypeptides is selected from the group comprising pycnodysostosis.
44. The combination according to claim 18, wherein the lysosomal storage disease is a lysosomal storage disease having a defective degradation or transport of cholesterol, cholesterol esters and/or other complex lipids.
45. The combination according to claim 44, wherein the lysosomal storage disease having a defective degradation or transport of cholesterol, cholesterol esters and/or other complex lipids is selected from the group comprising Neuronal Ceroid Lipofuscinosis type I, Neuronal Ceroid Lipofuscinosis type II, Neuronal Ceroid Lipofuscinosis type III and Neuronal Ceroid Lipofuscinosis type IV.

46. The combination according to claim 18, wherein the lysosomal storage disease is a lysosomal storage disease having multiple deficiencies of lysosomal enzymes.
47. The combination according to claim 46, wherein the lysosomal storage disease having multiple deficiencies of lysosomal enzymes is selected from the group comprising Galactosialidosis, Danon Disease, Pyknodysostosis, multiple sulfatase deficiency, GM2 Gangliosidosis, Mucopolidosis type II and Mucopolidosis type III.
48. The combination according to claim 47, wherein the lysosomal storage disease is Mucopolidosis type II or Mucopolidosis type III.
49. The combination according to claim 18, wherein the lysosomal storage disease is a lysosomal storage disease having transport and trafficking defects.
50. The combination according to claim 49, wherein the lysosomal storage disease having transport and trafficking defects is selected from the group comprising Cystinosis, Danon disease, Mucopolidosis type IV, Infantile sialic acid storage disease and Salla disease.
51. The combination according to any one of claims 5 to 50, wherein the lysosomal hydrolase is selected from the group comprising alpha-galactosidase A, alpha-glucosidase, sphingomyelinase, glucocerebrosidase, acid beta-glucosidase, N-acetylglucosamine-1-phosphotransferase, β -Hexosaminidase A, β -Hexosaminidase B, α -L-Iduronidase, Iduronatsulfatsulfatase, Heparansulfatsulfamidase, α -N-Acetylglukoseamidase, α -Glukosaminid-N-Acetyltransferase, N-Acetyl-glukosamin-6-sulfatsulfatase, β -Galactosidase, N-Acetylgalactosamin-4-sulfat-Sulfatase, β -Glucuronidase, Arylsulfatase A, Arylsulfatase B, Neuraminidase, α -Fucosidase, acid Ceramidase, Aspartylglukosaminidase, α -Mannosidase, β -Mannosidase, acid Lipase, acid β -Galactosidase, galactocerebrosidase, acid α -Mannosidase, acid β -Mannosidase, acid alpha-L-fucosidase, alpha-N-Acetylgalactosaminidase, alpha-N-Acetylglukosaminidase, beta-glucuronidase, sialidase and galactosylceramidase.

52. The combination according to any one of claims 5 to 51, wherein the lysosomal hydrolase is selected from the group consisting of alpha-galactosidase A, alpha-glucosidase, sphingomyelinase, N-acetylglucosamine-1-phosphotransferase, β -Hexosaminidase A, Hexosaminidase B, α -L-Iduronidase, Iduronatsulfat sulfatase, Heparansulfatsulfamidase, α -N-Acetylglukoseamidase, α -Glukosaminid-N-Acetyltransferase, N-Acetyl-glukosamin-6-sulfatsulfatase, β -Galactosidase, N-Acetylgalactosamin-4-sulfat-Sulfatase, β -Glucuronidase, Arylsulfatase A, Arylsulfatase B, Neuraminidase, α -Fucosidase, acid Ceramidase, Aspartylglukosaminidase, α -Mannosidase, β -Mannosidase, acid Lipase, acid β -Galactosidase, galactocerebrosidase, acid α -Mannosidase, acid β -Mannosidase, acid alpha-L-fucosidase, alpha-N-Acetylglucosaminidase, alpha-N-Acetylgalactosaminidase, beta-glucuronidase, sialidase and galactosylceramidase.
53. The combination according to any one of claims 3 to 4 and 6 to 17, wherein the lysosomal protein is a trans-membrane protein, wherein the trans-membrane protein has a reduced activity, preferably a reduced activity due to a mutation of the trans-membrane protein.
54. The combination according to any one of claims 3 to 4, 6 to 17 and 53, preferably claim 53, wherein the trans-membrane protein is a trans-membrane protein affected in a lysosomal storage disease.
55. The combination according to any one of claims 3 to 4, 6 to 17, 53 and 54, preferably claim 54, wherein the trans-membrane protein is selected from the group comprising NPC1, Acetyl-CoA α -Glukosaminid-N-Acetyltransferase, LAMP2, lysosomal free sialic acid transporter SLC17A5.
56. The combination according to claim 55, wherein the trans-membrane protein is NPC1.
57. The combination according to claim 56, wherein the lysosomal storage disease is Niemann-Pick type C1.

58. The combination according to claim 55, wherein the trans-membrane protein is Acetyl-CoA α -Glukosaminid-N-Acetyltransferase.
59. The combination according to claim 58, wherein the lysosomal storage disease is Sanfillipo syndrome Typ C.
60. The combination according to claim 55, wherein the trans-membrane protein is LAMP2.
61. The combination according to claim 60, wherein the lysosomal storage disease is Danon Disease.
62. The combination according to claim 60, wherein the lysosomal storage disease is Cystinosis.
63. The combination according to claim 55, wherein the trans-membrane protein is lysosomal free sialic acid transporter SLC17A5.
64. The combination according to claim 63, wherein the lysosomal storage disease is free sialic acid storage disease.
65. The combination according to any one of claims 3 to 4 and 6 to 17, wherein the lysosomal protein is a soluble non-enzymatic protein, wherein the soluble non-enzymatic protein has a reduced activity, preferably a reduced activity due to a mutation of the soluble non-enzymatic protein.
66. The combination according to any one of claims 3 to 4, 6 to 17 and 65, wherein the soluble non-enzymatic protein is a soluble non-enzymatic protein affected in a lysosomal storage disease.
67. The combination according to any one of claims 3 to 4, 6 to 17, 65 and 66, preferably claim 66, wherein the soluble non-enzymatic protein is NPC2.

68. The combination according to any one of claims 65 to 67, wherein the lysosomal storage disease is Niemann-Pick type C2.
69. The combination according to any one of claims 15 to 17, wherein the disease is different from a lysosomal storage disease.
70. The combination according to claim 69, wherein the disease different from a lysosomal storage disease is Parkinson disease.
71. The combination according to any one of claims 1 to 70,
wherein the first constituent is 1-deoxygalactonojirimycin or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.
72. The combination according to any one of claims 1 to 70,
wherein the first constituent is alpha-galacto-homonojirimycin or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.
73. The combination according to any one of claims 1 to 70,
wherein the first constituent is alpha-allo-homonojirimycin or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.
74. The combination according to any one of claims 1 to 70,
wherein the first constituent is beta-1-C-butyl-deoxygalactonojirimycin or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

75. The combination according to any one of claims 1 to 70,
wherein the first constituent is galactose, preferably D-galactose, or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.
76. The combination according to any one of claims 71 to 75,
wherein the lysosomal protein is affected in a disease;
wherein the lysosomal protein is α -galactosidase A; and/or
wherein the disease preferably is Fabry disease.
77. The combination according to any one of claims 1 to 70,
wherein the first constituent is N-butyl-DNJ or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.
78. The combination according to any one of claims 1 to 70,
wherein the first constituent is Deoxynojirimycin or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.
79. The combination according to any one of claims 77 to 78,
wherein the lysosomal protein is affected in a disease;
wherein the lysosomal protein is alpha-glucosidase; and/or
wherein the disease preferably is Pompe disease.
80. The combination according to any one of claims 1 to 70,
wherein the first constituent is N-acetyl-glucosamine-thiazoline (NGT) or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

81. The combination according to any one of claims 1 to 70,
wherein the first constituent is 6-acetamido-6-deoxycastanospermine (ACAS)
or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable
salt thereof.
82. The combination according to any one of claims 1 to 70,
wherein the first constituent is bisnaphthalimide nitro-indan-1-one or a
pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable
salt thereof.
83. The combination according to any one of claims 1 to 70,
wherein the first constituent is pyrrolo[3,4-d]pyridazin-1-one or a
pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable
salt thereof.
84. The combination according to any one of claims 1 to 70,
wherein the first constituent is pyrimethamine (PYR) or a pharmaceutically
acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable
salt thereof.
85. The combination according to any one of claims 80 to 84,
wherein the lysosomal protein is affected in a disease;
wherein the lysosomal protein is beta-hexosaminidase A/B; and/or
wherein the disease preferably is Tay-Sachs and/or Sandhoff disease.
86. The combination according to any one of claims 1 to 70,
wherein the first constituent is N-octyl-4-epi-beta-valienamine (NOEV) or a
pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

87. The combination according claim 86,
wherein the lysosomal protein is affected in a disease;
wherein the lysosomal protein is beta-galactosidase; and/or
wherein the disease preferably is Gm1-Gangliosidosis.
88. The combination according to any one of claims 1 to 70,
wherein the lysosomal protein is affected in a disease;
wherein the lysosomal protein is acid sphingomyelinase; and/or
wherein the disease preferably is Niemann-Pick disease type A or Niemann-Pick disease type B.
89. The combination according to any one of claims 1 to 70,
wherein the lysosomal protein is affected in a disease;
wherein the lysosomal protein is N-acetylglucosamine-1-phosphotransferase;
and/or
wherein the disease preferably is mucopolipidosis type II or mucopolipidosis type IIIA.
90. The combination according to any one of claims 1 to 70,
wherein the first constituent is N-nonyl-deoxynojirimycin (NN-DNJ) or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.
91. The combination according to any one of claims 1 to 70,
wherein the first constituent is N-octyl-2,5-anhydro-2,5-imino-D-glucitol or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

92. The combination according to any one of claims 1 to 70,
wherein the first constituent is N-octyl-isofagomine or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.
93. The combination according to any one of claims 1 to 70,
wherein the first constituent is N-octyl-beta-valienamine (NOV) or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.
94. The combination according to any one of claims 1 to 70,
wherein the first constituent is Isofagomine (IFG) or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.
95. The combination according to any one of claims 1 to 70,
wherein the first constituent is calystegine A3 or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.
96. The combination according to any one of claims 1 to 70,
wherein the first constituent is calystegine B1 or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.
97. The combination according to any one of claims 1 to 70,
wherein the first constituent is calystegine B2 or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

98. The combination according to any one of claims 1 to 70,
wherein the first constituent is calystegine C1 or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.
99. The combination according to any one of claims 1 to 70,
wherein the first constituent is 1,5-dideoxy-1,5-iminoxylitol (DIX) or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.
100. The combination according to any one of claims 1 to 70,
wherein the first constituent is alpha-1-C-nonyl-DIX or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.
101. The combination according to any one of claims 1 to 70,
wherein the first constituent is alpha-1-C-octyl-1-DNJ or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.
102. The combination according to any one of claims 1 to 70,
wherein the first constituent is N-butyl-DNJ or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol or a pharmaceutically acceptable salt thereof.

103. The combination according to any one of claims 90 to 102,
wherein the lysosomal protein is affected in a disease; and/or
wherein the lysosomal protein is glucocerebrosidase; and/or
wherein the disease preferably is Parkinson disease.
104. A combination comprising a first constituent and a second constituent,
wherein the combination is preferably a combination according to any one of
claims 1 to 70;
wherein the first constituent is a compound having the ability to rearrange a
lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is 1-deoxygalactonojirimycin or a
pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.
105. The combination according to claim 104, wherein the lysosomal protein is α -
galactosidase A.
106. The combination according to any one of claims 104 to 105, wherein the derivative
of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.
107. The combination according to any one of claims 104 to 106, wherein the
combination is for use in the treatment or prevention of Fabry's disease.
108. A combination comprising a first constituent and a second constituent,
wherein the combination is preferably a combination according to any one of
claims 1 to 70;
wherein the first constituent is a compound having the ability to rearrange a
lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is alpha-galacto-homonojirimycin or a
pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.

109. The combination according to claim 108, wherein the lysosomal protein is α -galactosidase A.
110. The combination according to any one of claims 108 to 109, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.
111. The combination according to any one of claims 108 to 110, wherein the combination is for use in the treatment or prevention of Fabry's disease.
112. A combination comprising a first constituent and a second constituent,
wherein the combination is preferably a combination according to any one of claims 1 to 70;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is alpha-allo-homonojirimycin or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.
113. The combination according to claim 112, wherein the lysosomal protein is α -galactosidase A.
114. The combination according to any one of claims 112 to 113, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.
115. The combination according to any one of claims 112 to 114, wherein the combination is for use in the treatment or prevention of Fabry's disease.
116. A combination comprising a first constituent and a second constituent,
wherein the combination is preferably a combination according to any one of claims 1 to 70;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,

wherein the first constituent is beta-1-C-butyl-deoxygalactonojirimycin or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol and/or a derivative thereof.

117. The combination according to claim 116, wherein the lysosomal protein is α -galactosidase A.
118. The combination according to any one of claims 116 to 117, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.
119. The combination according to any one of claims 116 to 118, wherein the combination is for use in the treatment or prevention of Fabry's disease.
120. A combination comprising a first constituent and a second constituent,
wherein the combination is preferably a combination according to any one of claims 1 to 70;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is galactose, preferably D-galactose or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.
121. The combination according to claim 120, wherein the lysosomal protein is α -galactosidase A.
122. The combination according to any one of claims 120 to 121, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.
123. The combination according to any one of claims 120 to 122, wherein the combination is for use in the treatment or prevention of Fabry's disease.
124. A combination comprising a first constituent and a second constituent,

wherein the combination is preferably a combination according to any one of claims 1 to 70;

wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,

wherein the first constituent is N-acetyl-glucosamine-thiazoline (NGT) or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol and/or a derivative thereof.

125. The combination according to claim 124, wherein the lysosomal protein is beta-hexosaminidase A/B.
126. The combination according to any one of claims 124 to 125, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.
127. The combination according to any one of claims 124 to 126, wherein the combination is for use in the treatment or prevention of Tay-Sachs or Sandhoff.
128. A combination comprising a first constituent and a second constituent,
wherein the combination is preferably a combination according to any one of claims 1 to 70;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is 6-acetamido-6-deoxycastanospermine (ACAS) or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.
129. The combination according to claim 128, wherein the lysosomal protein is beta-hexosaminidase A/B.
130. The combination according to any one of claims 128 to 129, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

131. The combination according to any one of claims 128 to 130, wherein the combination is for use in the treatment or prevention of Tay-Sachs or Sandhoff.
132. A combination comprising a first constituent and a second constituent,
wherein the combination is preferably a combination according to any one of claims 1 to 70;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is bisnaphthalimide nitro-indan-1-one or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.
133. The combination according to claim 132, wherein the lysosomal protein is beta-hexosaminidase A/B.
134. The combination according to any one of claims 132 to 133, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.
135. The combination according to any one of claims 132 to 134, wherein the combination is for use in the treatment or prevention of Tay-Sachs or Sandhoff.
136. A combination comprising a first constituent and a second constituent,
wherein the combination is preferably a combination according to any one of claims 1 to 70;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is pyrrolo[3,4-d]pyridazin-1-one or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.
137. The combination according to claim 136, wherein the lysosomal protein is beta-hexosaminidase A/B.

138. The combination according to any one of claims 136 to 137, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.
139. The combination according to any one of claims 136 to 138, wherein the combination is for use in the treatment or prevention of Tay-Sachs or Sandhoff.
140. A combination comprising a first constituent and a second constituent,
wherein the combination is preferably a combination according to any one of claims 1 to 70;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is pyrimethamine (PYR) or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.
141. The combination according to claim 140, wherein the lysosomal protein is beta-hexosaminidase A/B.
142. The combination according to any one of claims 140 to 141, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.
143. The combination according to any one of claims 140 to 142, wherein the combination is for use in the treatment or prevention of Tay-Sachs or Sandhoff.
144. A combination comprising a first constituent and a second constituent,
wherein the combination is preferably a combination according to any one of claims 1 to 70;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is N-octyl-4-epi-beta-valienamine (NOEV) or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.

145. The combination according to claim 144, wherein the lysosomal protein is beta-galactosidase.
146. The combination according to any one of claims 144 to 145, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.
147. The combination according to any one of claims 144 to 146, wherein the combination is for use in the treatment or prevention of GM1-Gangliosidosis.
148. A combination comprising a first constituent and a second constituent,
wherein the combination is preferably a combination according to any one of claims 1 to 70;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is N-butyl-DNJ or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.
149. The combination according to claim 148, wherein the lysosomal protein is alpha-glucosidase.
150. The combination according to any one of claims 148 to 149, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.
151. The combination according to any one of claims 148 to 150, wherein the combination is for use in the treatment or prevention of Pompe's disease.
152. A combination comprising a first constituent and a second constituent,
wherein the combination is preferably a combination according to any one of claims 1 to 70;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,

wherein the first constituent is deoxynojirimycin or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol and/or a derivative thereof.

153. The combination according to claim 152, wherein the lysosomal protein is alpha-glucosidase.
154. The combination according to any one of claims 152 to 153, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.
155. The combination according to any one of claims 152 to 154, wherein the combination is for use in the treatment or prevention of Pompe's disease.
156. Use of a combination as defined in any one of claims 1 to 155, for the manufacture of a medicament for the treatment or prevention of a disease.
157. Use according to claim 156, wherein the disease is a lysosomal storage disease.
158. Use according to claim 157, wherein the lysosomal storage disease is a lysosomal storage disease having a defective degradation of a sphingolipid component.
159. Use according to claim 158, wherein the lysosomal storage disease having a defective degradation of a sphingolipid component is selected from the group comprising Fabry disease, Gaucher disease type I, Gaucher disease type II, Gaucher disease type III, acid sphingomyelinase deficiency, Farber disease, GM1 gangliosidosis type I, GM1 gangliosidosis type II, GM1 gangliosidosis type III, GM2 gangliosidosis, Krabbe disease, Metachromatic leukodystrophy type I, Metachromatic leukodystrophy type II and Metachromatic leukodystrophy type III.
160. Use according to claim 158, wherein the lysosomal storage disease having a defective degradation of a sphingolipid component is selected from the group consisting of Fabry disease, acid sphingomyelinase deficiency, Farber disease, GM1 gangliosidosis type I, GM1 gangliosidosis type II, GM1 gangliosidosis type III,

GM2 gangliosidosis, Krabbe disease, Metachromatic leukodystrophy type I, Metachromatic leukodystrophy type II and Metachromatic leukodystrophy type III.

161. Use according to any one of claims 159 to 160, wherein the lysosomal storage disease having a defective degradation of sphingolipid components is Fabry disease.
162. Use according to any one of claims 159 to 160, wherein the acid sphingomyelinase deficiency is selected from the group comprising Niemann-Pick type A and Niemann-Pick type A and B.
163. Use according to any one of claims 159 to 160, wherein the GM2 gangliosidosis is selected from the group comprising Tay-Sachs type I, Tay-Sachs type II, Tay-Sachs type III and Sandhoff.
164. Use according to claim 163, wherein the lysosomal storage disease is Tay-Sachs type I, Tay-Sachs type II or Tay-Sachs type III.
165. Use according to claim 163, wherein the lysosomal storage disease is Sandhoff.
166. Use according to any one of claims 159 to 160, wherein the lysosomal storage disease is GM1 gangliosidosis type I, GM1 gangliosidosis type II or GM1 gangliosidosis type III.
167. Use according to claim 157, wherein the lysosomal storage disease is a lysosomal storage disease having a defective metabolism of glycosaminoglycans.
168. Use according to claim 167, wherein the lysosomal storage disease having a defective metabolism of glycosaminoglycans is selected from the group comprising MPS I, MPS II, MPS III, MPS IV, MPS VI, MPS VI and MPS IX.
169. Use according to claim 168, wherein MPS I is selected from the group comprising Hurler disease, Hurler-Scheie syndrome and Scheie syndrome.

170. Use according to claim 168, wherein MPS II is selected from the group comprising Hunter syndrome.
171. Use according to claim 168, wherein MPS III is selected from the group comprising Sanfilippo syndrome type A, Sanfilippo syndrome type B, Sanfilippo syndrome type C and Sanfilippo syndrome type D.
172. Use according to claim 168, wherein MPS IV is selected from the group comprising Morquio type A and Morquio type B.
173. Use according to claim 168, wherein MPS VI is selected from the group comprising Maroteaux-Lamy.
174. Use according to claim 168, wherein MPS VII is selected from the group comprising Sly.
175. Use according to claim 168, wherein MPS IX is selected from the group comprising Hyaluronidase deficiency and Multiple Sulfatase deficiency.
176. Use according to claim 157, wherein the lysosomal storage disease is a lysosomal storage disease having a defective degradation of a glycan portion of a glycoprotein.
177. Use according to claim 176, wherein the lysosomal storage disease having a defective degradation of a glycan portion of a glycoprotein is selected from the group comprising Aspartylglucosaminuria, Fucosidosis type I, Fucosidosis type II, Mannosidosis, Sialidosis type I and Sialidosis type II.
178. Use according to claim 157, wherein the lysosomal storage disease is a lysosomal storage disease having a defective degradation of glycogen.
179. Use according to claim 178, wherein the lysosomal storage disease having a defective degradation of glycogen is selected from the group comprising Pompe's disease.

180. Use according to claim 179, wherein the lysosomal storage disease is Pompe's disease.
181. Use according to claim 157, wherein the lysosomal storage disease is a lysosomal storage disease having a defective degradation of polypeptides.
182. Use according to claim 181, wherein the lysosomal storage disease having a defective degradation of polypeptides is selected from the group comprising pycnodysostosis.
183. Use according to claim 157, wherein the lysosomal storage disease is a lysosomal storage disease having a defective degradation or transport of cholesterol, cholesterol esters and/or other complex lipids.
184. Use according to claim 183, wherein the lysosomal storage disease having a defective degradation or transport of cholesterol, cholesterol esters and/or other complex lipids is selected from the group comprising Neuronal Ceroid Lipofuscinosis type I, Lipofuscinosis type II, Lipofuscinosis type III and Lipofuscinosis type IV.
185. Use according to claim 157, wherein the lysosomal storage disease is a lysosomal storage disease having multiple deficiencies of lysosomal enzymes.
186. Use according to claim 185, wherein the lysosomal storage disease having multiple deficiencies of lysosomal enzymes is selected from the group comprising Galactosialidosis, Danon Disease, Pycnodysostosis, multiple sulfatase deficiency, GM2 Gangliosidosis, Mucopolidosis type II and Mucopolidosis type III.
187. Use according to claim 186, wherein the lysosomal storage disease is Mucopolidosis type II or Mucopolidosis type III.

188. Use according to claim 157, wherein the lysosomal storage disease is a lysosomal storage disease having transport and trafficking defects.
189. Use according to claim 188, wherein the lysosomal storage disease having transport and trafficking defects is selected from the group comprising Cystinosis, Danon disease, Mucopolidosis type IV, Infantile sialic acid storage disease and Salla disease.
190. Use according to claim 157, wherein the lysosomal storage disease is Niemann -Pick type C1.
191. Use according to claim 157, wherein the lysosomal storage disease is San filipo syndrome type C.
192. Use according to claim 157, wherein the lysosomal storage disease is Danon disease.
193. Use according to claim 157, wherein the lysosomal storage disease is free sialic acid storage disease.
194. Use according to claim 157, wherein the lysosomal storage disease is Niemann -Pick type C2.
195. Use according to claim 156, wherein the disease is different from a lysosomal storage disease.
196. Use according to claim 195, wherein the disease different from a lysosomal storage disease is Parkinson disease.
197. Use of a compound having the ability to rearrange a lysosomal protein in or for the manufacture of a medicament for the treatment of a disease, wherein the medicament comprises a combination comprising a first constituent and a second constituent, wherein the first constituent is a or the compound having the ability to

rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, and wherein the second constituent is Ambroxol and/or a derivative thereof.

198. Use according to claim 197, wherein the combination is a combination according to any one of claims 1 to 155, preferably according to any one of claims 104 to 155.
199. Use of Ambroxol and/or a derivative of Ambroxol in or for the manufacture of a medicament for the treatment of a disease, wherein the medicament comprises a combination comprising a first constituent and a second constituent,
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity and
wherein the second constituent is Ambroxol and/or a derivative thereof.
200. Use according to claim 199, wherein the combination is a combination according to any one of claims 1 to 155, preferably according to any one of claims 104 to 155.
201. Use according to any one of claims 197 to 200, wherein the disease is a lysosomal storage disease.
202. Use according to any one of claims 197 to 200, wherein the disease is different from a lysosomal storage disease.
203. Use according to claim 202, wherein the disease different from a lysosomal storage disease is Parkinson disease.
204. A pharmaceutical preparation comprising a first constituent, a second constituent optionally a further constituent,
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the second constituent is Ambroxol and/or a derivative of Ambroxol and
wherein the further constituent is selected from the group comprising a pharmaceutically acceptable excipient and a pharmaceutically active agent, and

wherein preferably the pharmaceutical preparation increases the reduced activity of the lysosomal protein, and wherein the lysosomal protein has a reduced activity.

205. The pharmaceutical preparation according to claim 204, wherein the lysosomal protein is affected in a disease.
206. The pharmaceutical preparation according to any one of claims 204 to 205, wherein the lysosomal protein is selected from the group comprising an enzyme, a trans-membrane protein and a soluble non-enzymatic protein.
207. The pharmaceutical preparation according to claim 206, wherein the lysosomal protein is an enzyme, wherein the enzyme has a reduced activity, preferably a reduced activity due to a mutation of the enzyme.
208. The pharmaceutical preparation according to claim 207, wherein the enzyme is a lysosomal hydrolase.
209. The pharmaceutical preparation according to any one of claims 204 to 208, wherein the compound having the ability to rearrange a lysosomal protein is a chaperon.
210. The pharmaceutical preparation according to claim 209, wherein the chaperon is a pharmacological chaperon or a pharmaceutically acceptable salt, solvate or derivative thereof.
211. The pharmaceutical preparation according to claim 210, wherein the pharmacological chaperon is a sugar and/or an imino sugar or a pharmaceutically acceptable salt, solvate or derivative of the sugar and/or the imino sugar.
212. The pharmaceutical preparation according to claim 211, wherein the sugar is galactose, preferably D-galactose.

213. The pharmaceutical preparation according to claim 211, wherein the imino sugar is selected from the group comprising 1-deoxygalactonojirimycin (DGJ), alpha-galacto-homonojirimycin, alpha-allo-homonojirimycin, beta-1-C-butyl-deoxygalactonojirimycin, N-nonyl-deoxynojoirrimycin (NN-DNJ), N-octyl-2,5-anhydro-2,5-imino-D-glucitol, N-octyl-isofagomine, N-octyl-beta-valienamine (NOV), Isofagomine (IFG), calystegines A3 B1, B2, C1, 1,5-dideoxy-1,5-iminoxylitol (DIX), alpha-1-C-nonyl-DIX, alpha-1-C-octyl-1-DNJ, N-acetyl-glucosamine-thiazoline (NGT), 6-acetamido-6-deoxycastanospermine (ACAS), bisnaphthalimide nitro-indan-1-one, pyrrolo[3,4-d]pyridazin-1-one, pyrimethamine (PYR), N-actyl-4-epi-beta-valienamine (NOEV), N-butyl-DNJ, Deoxynojoirrimycin (DNJ), N-Acetyl-galactosamine (GalNAc), 2-Acetamido-1,2-dideoxynojoirrimycin (AddDNJ) and Castanospermine (CAS), and derivatives thereof; and pharmaceutically acceptable salts thereof.
214. The pharmaceutical preparation according to any one of claims 204 to 213, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.
215. The pharmaceutical preparation according to any one of claims 204 to 214, wherein at least one of the first constituent and the second constituent is/are present as a solvate or a pharmaceutically acceptable salt thereof.
216. The pharmaceutical preparation according to claim 215, wherein the second constituent is present as a pharmaceutically acceptable salt thereof, wherein the second constituent is Ambroxol, wherein preferably the pharmaceutically acceptable salt of Ambroxol is Ambroxol hydrochloride.
217. The pharmaceutical preparation according to claim 215, wherein the second constituent is present as a pharmaceutically acceptable salt thereof, wherein the second constituent is a derivative of Ambroxol, wherein the derivative of Ambroxol is bromhexine, and wherein preferably the pharmaceutically acceptable salt of bromhexine is bromhexine hydrochloride.

218. The pharmaceutical preparation according to any one of claims 204 to 217, wherein the pharmaceutical preparation is suitable for or is for use in the treatment and/or prevention of a disease.
219. The pharmaceutical preparation according to any one of claims 204 to 218, wherein the first constituent and the second constituent are formulated into separate dosage forms, wherein preferably each dosage form is independently selected from the group comprising tablets, capsules, powder, mixture, effervescence tablets and solutions.
220. The pharmaceutical preparation according to any one of claims 204 to 219, wherein the first constituent and the second constituent are each and individually administered subcutaneously, intranasally, intravenously, sublingually, transmucosally and/or intracranially.
221. The pharmaceutical preparation according to any one of claims 204 to 220, wherein the pharmaceutical preparation is suitable for or is for use in a method for the treatment of a subject wherein the method comprises administering the pharmaceutical preparation to the subject.
222. The pharmaceutical preparation according to any one of claims 205 to 221, wherein the disease is a lysosomal storage disease.
223. The pharmaceutical preparation according to any one of claims 205 to 221, wherein the disease is different from a lysosomal storage disease.
224. The pharmaceutical preparation according to claim 223, wherein the disease different from a lysosomal storage disease is Parkinson disease.
225. A pharmaceutical preparation comprising a first constituent and a second constituent, wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of claims 204 to 224;

wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,

wherein the first constituent is 1-deoxygalactonojirimycin or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol and/or a derivative thereof.

226. The pharmaceutical preparation according to claim 225, wherein the lysosomal protein is α -galactosidase A.
227. The pharmaceutical preparation according to any one of claims 225 to 226, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.
228. The pharmaceutical preparation according to any one of claims 225 to 227, wherein the pharmaceutical preparation is for use in the treatment or prevention of Fabry's disease.
229. A pharmaceutical preparation comprising a first constituent and a second constituent,
wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of claims 204 to 224;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is alpha-galacto-homonojirimycin or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.
230. The pharmaceutical preparation according to claim 229, wherein the lysosomal protein is α -galactosidase A.
231. The pharmaceutical preparation according to any one of claims 229 to 230, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

232. The pharmaceutical preparation according to any one of claims 229 to 231, wherein the pharmaceutical preparation is for use in the treatment or prevention of Fabry's disease.
233. A pharmaceutical preparation comprising a first constituent and a second constituent,
wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of claims 204 to 224;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is alpha-allo-homonojirimycin or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.
234. The pharmaceutical preparation according to claim 233, wherein the lysosomal protein is α -galactosidase A.
235. The pharmaceutical preparation according to any one of claims 233 to 234, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.
236. The pharmaceutical preparation according to any one of claims 233 to 235, wherein the pharmaceutical preparation is for use in the treatment or prevention of Fabry's disease.
237. A pharmaceutical preparation comprising a first constituent and a second constituent,
wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of claims 204 to 224;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,

wherein the first constituent is beta-1-C-butyl-deoxygalactonojirimycin or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol and/or a derivative thereof.

238. The pharmaceutical preparation according to claim 237, wherein the lysosomal protein is α -galactosidase A.
239. The pharmaceutical preparation according to any one of claims 237 to 238, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.
240. The pharmaceutical preparation according to any one of claims 237 to 239, wherein the pharmaceutical preparation is for use in the treatment or prevention of Fabry's disease.
241. A pharmaceutical preparation comprising a first constituent and a second constituent,
wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of claims 204 to 224;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is galactose, preferably D-galactose or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.
242. The pharmaceutical preparation according to claim 241, wherein the lysosomal protein is α -galactosidase A.
243. The pharmaceutical preparation according to any one of claims 241 to 242, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.

244. The pharmaceutical preparation according to any one of claims 241 to 243, wherein the pharmaceutical preparation is for use in the treatment or prevention of Fabry's disease.
245. A pharmaceutical preparation comprising a first constituent and a second constituent,
wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of claims 204 to 224;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is N-acetyl-glucosamine-thiazoline (NGT) or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.
246. The pharmaceutical preparation according to claim 245, wherein the lysosomal protein is beta-hexosaminidase A/B.
247. The pharmaceutical preparation according to any one of claims 245 to 246, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.
248. The pharmaceutical preparation according to any one of claims 245 to 247, wherein the pharmaceutical preparation is for use in the treatment or prevention of Tay-Sachs or Sandhoff.
249. A pharmaceutical preparation comprising a first constituent and a second constituent,
wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of claims 204 to 224;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is 6-acetamido-6-deoxycastanospermine (ACAS) or a pharmaceutically acceptable salt thereof; and

wherein the second constituent is Ambroxol and/or a derivative thereof.

250. The pharmaceutical preparation according to claim 249, wherein the lysosomal protein is beta-hexosaminidase A/B.
251. The pharmaceutical preparation according to any one of claims 249 to 250, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.
252. The pharmaceutical preparation according to any one of claims 249 to 251, wherein the pharmaceutical preparation is for use in the treatment or prevention of Tay-Sachs or Sandhoff.
253. A pharmaceutical preparation comprising a first constituent and a second constituent,
wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of claims 204 to 224;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is bisnaphthalimide nitro-indan-1-one or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.
254. The pharmaceutical preparation according to claim 253, wherein the lysosomal protein is beta-hexosaminidase A/B.
255. The pharmaceutical preparation according to any one of claims 253 to 254, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.
256. The pharmaceutical preparation according to any one of claims 253 to 255, wherein the pharmaceutical preparation is for use in the treatment or prevention of Tay-Sachs or Sandhoff.

257. A pharmaceutical preparation comprising a first constituent and a second constituent,
wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of claims 204 to 224;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is pyrrolo[3,4-d]pyridazin-1-one or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.
258. The pharmaceutical preparation according to claim 257, wherein the lysosomal protein is beta-hexosaminidase A/B.
259. The pharmaceutical preparation according to any one of claims 257 to 257, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.
260. The pharmaceutical preparation according to any one of claims 257 to 259, wherein the pharmaceutical preparation is for use in the treatment or prevention of Tay-Sachs or Sandhoff.
261. A pharmaceutical preparation comprising a first constituent and a second constituent,
wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of claims 204 to 224;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is pyrimethamine (PYR) or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.

262. The pharmaceutical preparation according to claim 261, wherein the lysosomal protein is beta-hexosaminidase A/B.
263. The pharmaceutical preparation according to any one of claims 261 to 262, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.
264. The pharmaceutical preparation according to any one of claims 261 to 263, wherein the pharmaceutical preparation is for use in the treatment or prevention of Tay-Sachs or Sandhoff.
265. A pharmaceutical preparation comprising a first constituent and a second constituent,
wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of claims 204 to 224;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is N-octyl-4-epi-beta-valienamine (NOEV) or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.
266. The pharmaceutical preparation according to claim 265, wherein the lysosomal protein is beta-galactosidase.
267. The pharmaceutical preparation according to any one of claims 265 to 266, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.
268. The pharmaceutical preparation according to any one of claims 265 to 267, wherein the pharmaceutical preparation is for use in the treatment or prevention of GM1-Gangliosidosis.

269. A pharmaceutical preparation comprising a first constituent and a second constituent,
wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of claims 204 to 224;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is N-butyl-DNJ or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.
270. The pharmaceutical preparation according to claim 269, wherein the lysosomal protein is alpha-glucosidase.
271. The pharmaceutical preparation according to any one of claims 269 to 270, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.
272. The pharmaceutical preparation according to any one of claims 269 to 271, wherein the pharmaceutical preparation is for use in the treatment or prevention of Pompe's disease.
273. A pharmaceutical preparation comprising a first constituent and a second constituent,
wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of claims 204 to 224;
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity,
wherein the first constituent is Deoxynojirimycin or a pharmaceutically acceptable salt thereof; and
wherein the second constituent is Ambroxol and/or a derivative thereof.
274. The pharmaceutical preparation according to claim 273, wherein the lysosomal protein is alpha-glucosidase.

275. The pharmaceutical preparation according to any one of claims 273 to 274, wherein the derivative of Ambroxol is bromhexine or a pharmaceutically acceptable salt thereof.
276. The pharmaceutical preparation according to any one of claims 273 to 275, wherein the pharmaceutical preparation is for use in the treatment or prevention of Pompe's disease.
277. A method for preparing a pharmaceutical preparation, preferably a pharmaceutical preparation according to any one of claims 204 to 276, comprising the steps of
formulating a first constituent as defined in any one of the preceding claims and a second constituent as defined in any one of the preceding claims into a single dosage form or into two separate dosage forms, wherein in case of two separate dosage forms a first of the two separate dosage forms contains the first constituent and a second of the two separate dosage forms contains the second constituent.
278. Use of a pharmaceutical preparation for the manufacture of a medicament, wherein the pharmaceutical preparation is a pharmaceutical preparation according to any one of claims 204 to 276, and wherein the pharmaceutical preparation is for the treatment and/or prevention of a disease.
279. Use according to claim 278, wherein the disease is a lysosomal storage disease.
280. Use according to claim 278, wherein the disease is different from a lysosomal storage disease.
281. Use according to claim 280, wherein the disease different from a lysosomal storage disease is Parkinson disease.
282. A method for the treatment of a disease, wherein the method comprises administering to a subject a first constituent as defined in any one of claims 1 to 276 and, prior to, concomitantly with or after a second constituent as defined in any one

of claims 1 to 276, and/or a combination according to any one of claims 1 to 155 and/or a pharmaceutical preparation according to any one of claims 204 to 276.

- 283. The method according to claim 282, wherein the disease is a lysosomal storage disease.
- 284. The method according to claim 282, wherein the disease is different from a lysosomal storage disease.
- 285. The method according to claim 284, wherein the disease different from a lysosomal storage disease is Parkinson disease.
- 286. A method for increasing activity of a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the method comprises administering to a cell a compound having the ability to rearrange the lysosomal protein, and Ambroxol and/or a derivative thereof.
- 287. A method for increasing activity of a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the method comprises administering to a subject a compound having the ability to rearrange the lysosomal protein, and Ambroxol and/or a derivative thereof, wherein the compound having the ability to rearrange the lysosomal protein increases the activity of the lysosomal protein.
- 288. A method for increasing activity of a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the method comprises administering to a subject a first constituent as defined in any one of claims 1 to 276 and, prior to, concomitantly with or after a second constituent as defined in any one of claims 1 to 276, and/or a combination according to any one of claims 1 to 155 and/or a pharmaceutical preparation according to any one of claims 204 to 276, wherein the combination and/or the pharmaceutical preparation increases the activity of the lysosomal protein.

289. A combination comprising a first constituent and a second constituent,
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity wherein the reduced activity is reduced due to a mutation of the lysosomal protein, and
wherein the second constituent is Ambroxol and/or a derivative of Ambroxol,
wherein the combination is preferably a combination according to any one of claims 1 to 155,

for use in a method of personalized therapeutic treatment of a subject, wherein the method comprises the following steps:

- step a): determining whether in a sample of the subject the lysosomal protein has a reduced activity, preferably such reduced activity results from one or more mutation of the lysosomal protein compared to the wild type lysosomal protein;
- step b): identifying a compound having the ability to rearrange the lysosomal protein having reduced activity, and wherein the compound is suitable for or is increasing the reduced activity of the lysosomal protein; and
- step c): administering to the subject the first constituent prior to, concomitantly with or after the second constituent.

290. A pharmaceutical preparation comprising a first constituent, a second constituent and optionally a further constituent,
wherein the first constituent is a compound having the ability to rearrange a lysosomal protein, wherein the lysosomal protein has a reduced activity, wherein the reduced activity is reduced due to a mutation of the lysosomal protein, and
wherein the second constituent is Ambroxol and/or a derivative of Ambroxol,
wherein the further constituent is selected from the group comprising pharmaceutically acceptable excipients and pharmaceutically active agents,

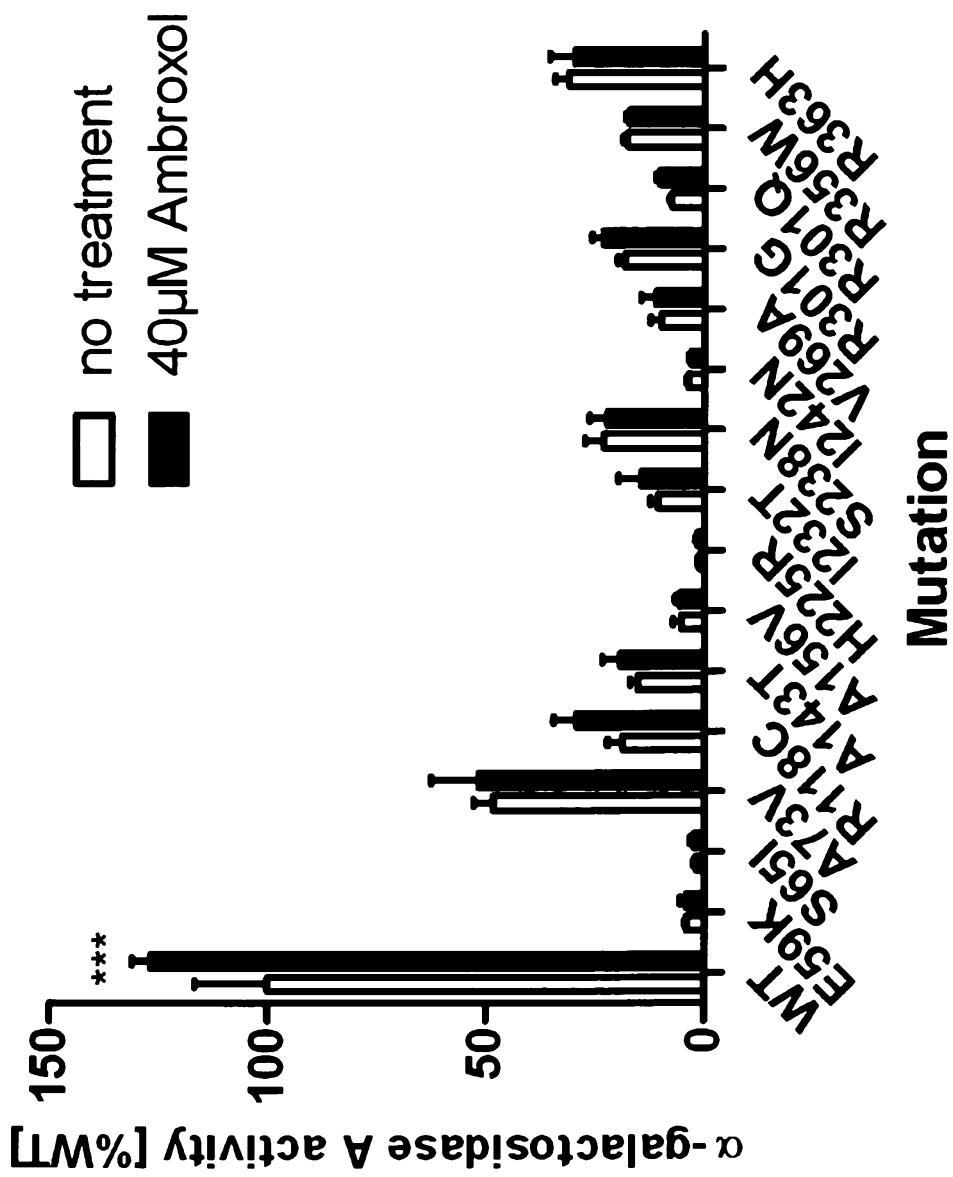
wherein the pharmaceutical preparation is preferably a pharmaceutical preparation according to any one of claims 204 to 276,

for use in a method of personalized therapeutic treatment of a subject, wherein the method comprises the following steps:

- step a): determining whether in a sample of the subject the lysosomal protein has a reduced activity, preferably such reduced activity results from one or more mutation of the lysosomal protein compared to the wild type lysosomal protein;
- step b): identifying a compound having the ability to rearrange the lysosomal protein having reduced activity, and wherein the compound is suitable for or is increasing the reduced activity of the lysosomal protein; and
- step c): administering to the subject the first constituent prior to, concomitantly with or after the second constituent..

Figure 1:

Ambroxol alone has no effect on Fabry mutants



2/16

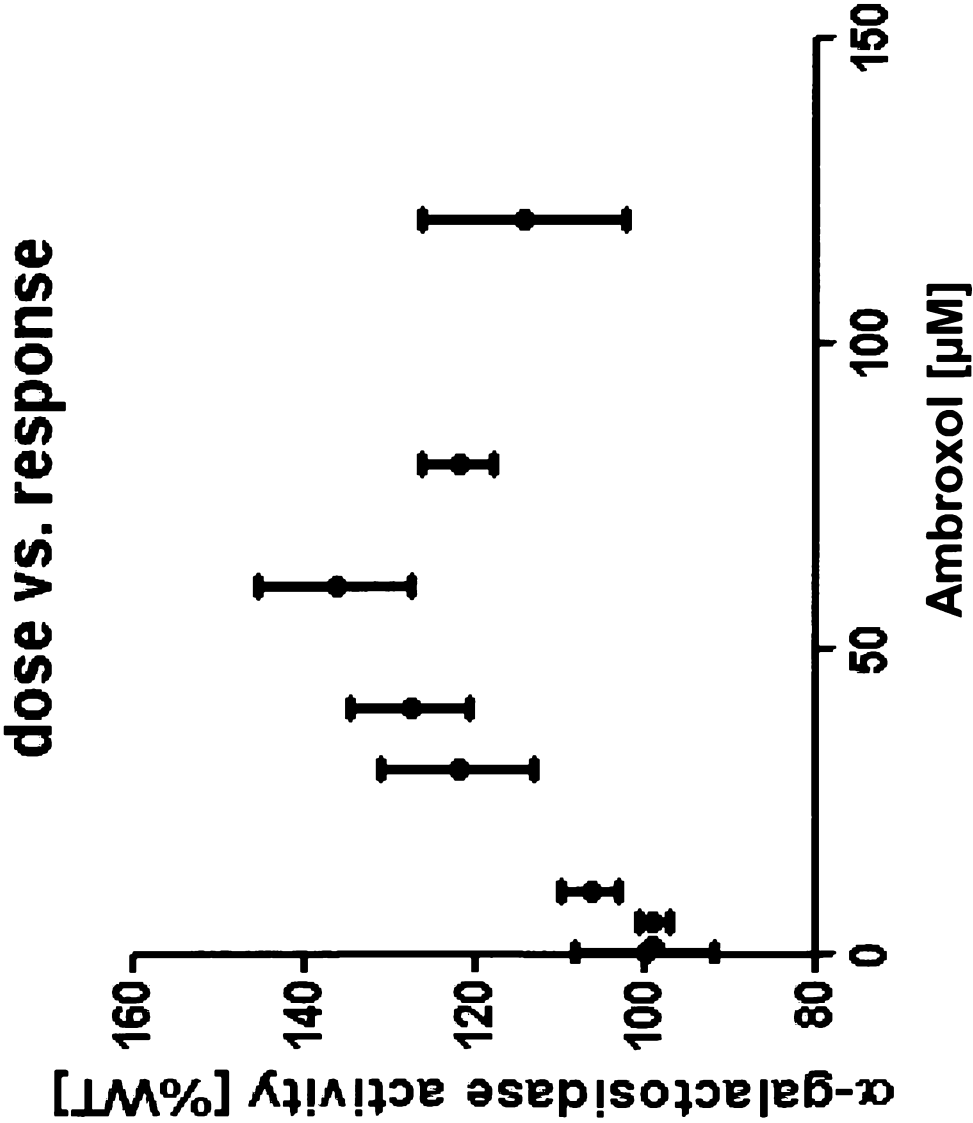


Figure 2:

Figure 3A:

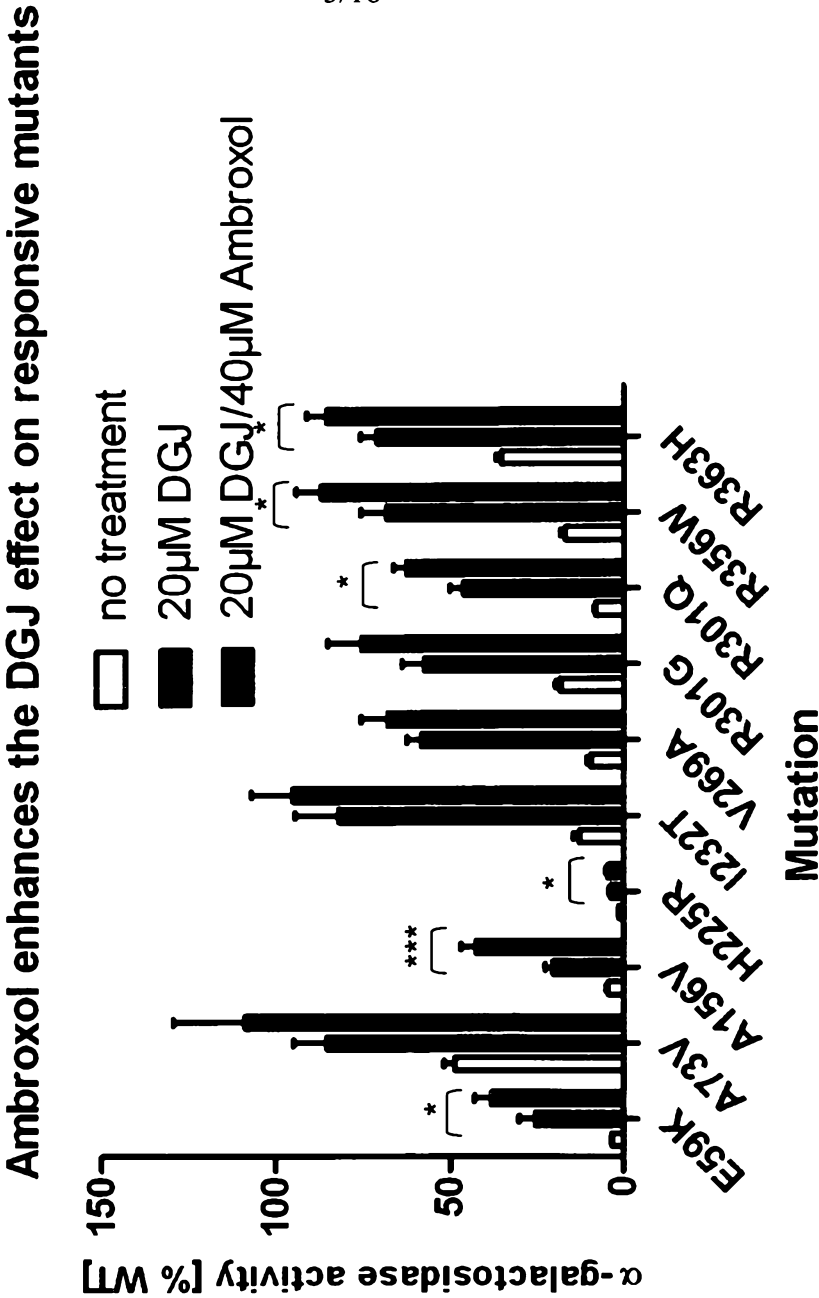


Fig. 3B: Also non-iminosugar Galactose and ABX show the combinatoric effect on Fabry mutants [N=2, values are mean $\pm$ SD]

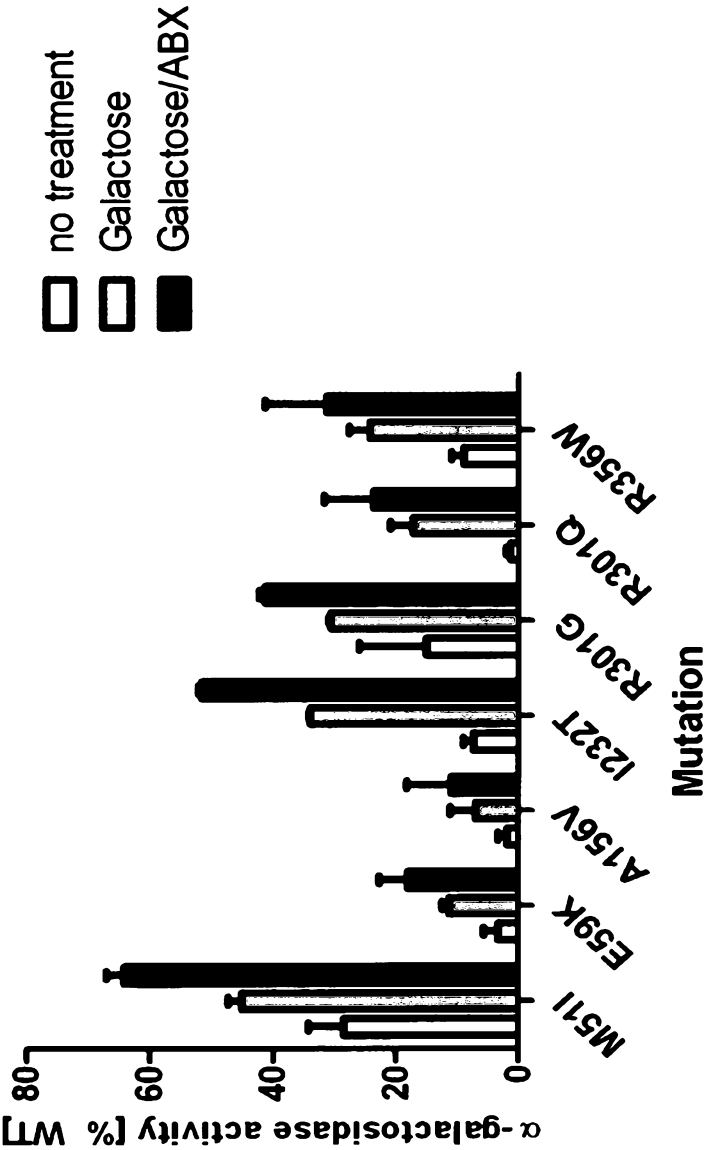


Figure 4:

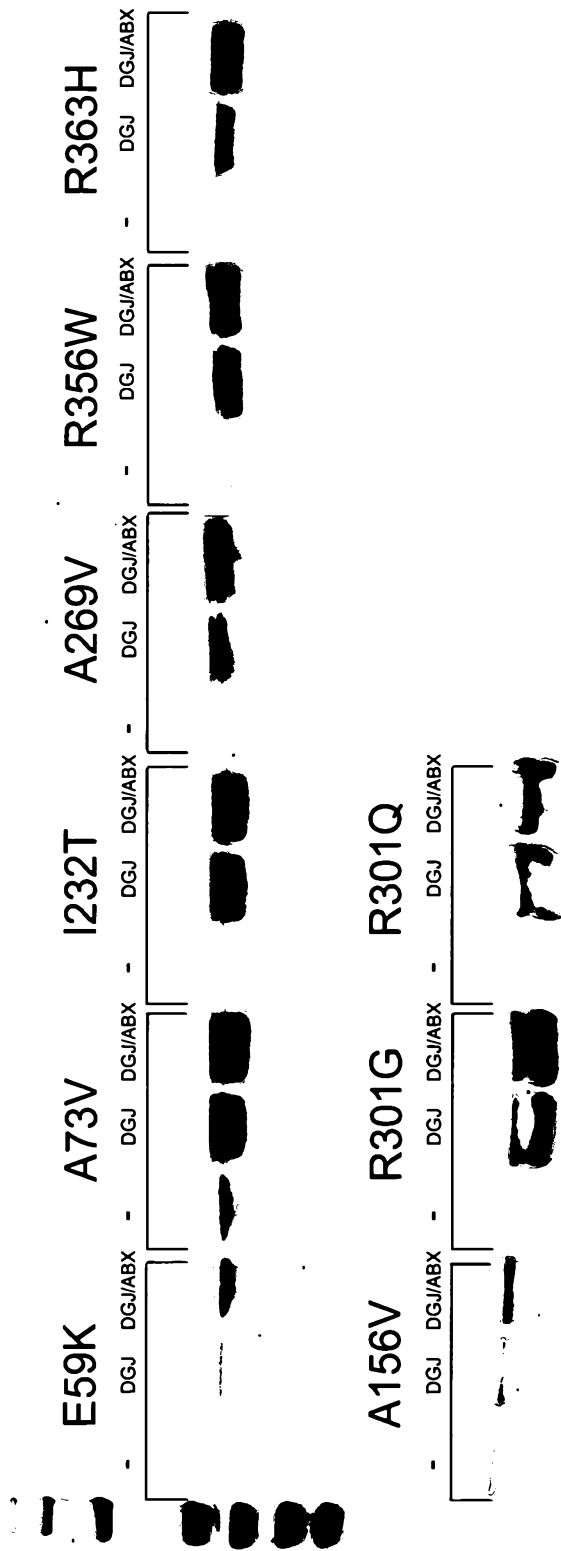
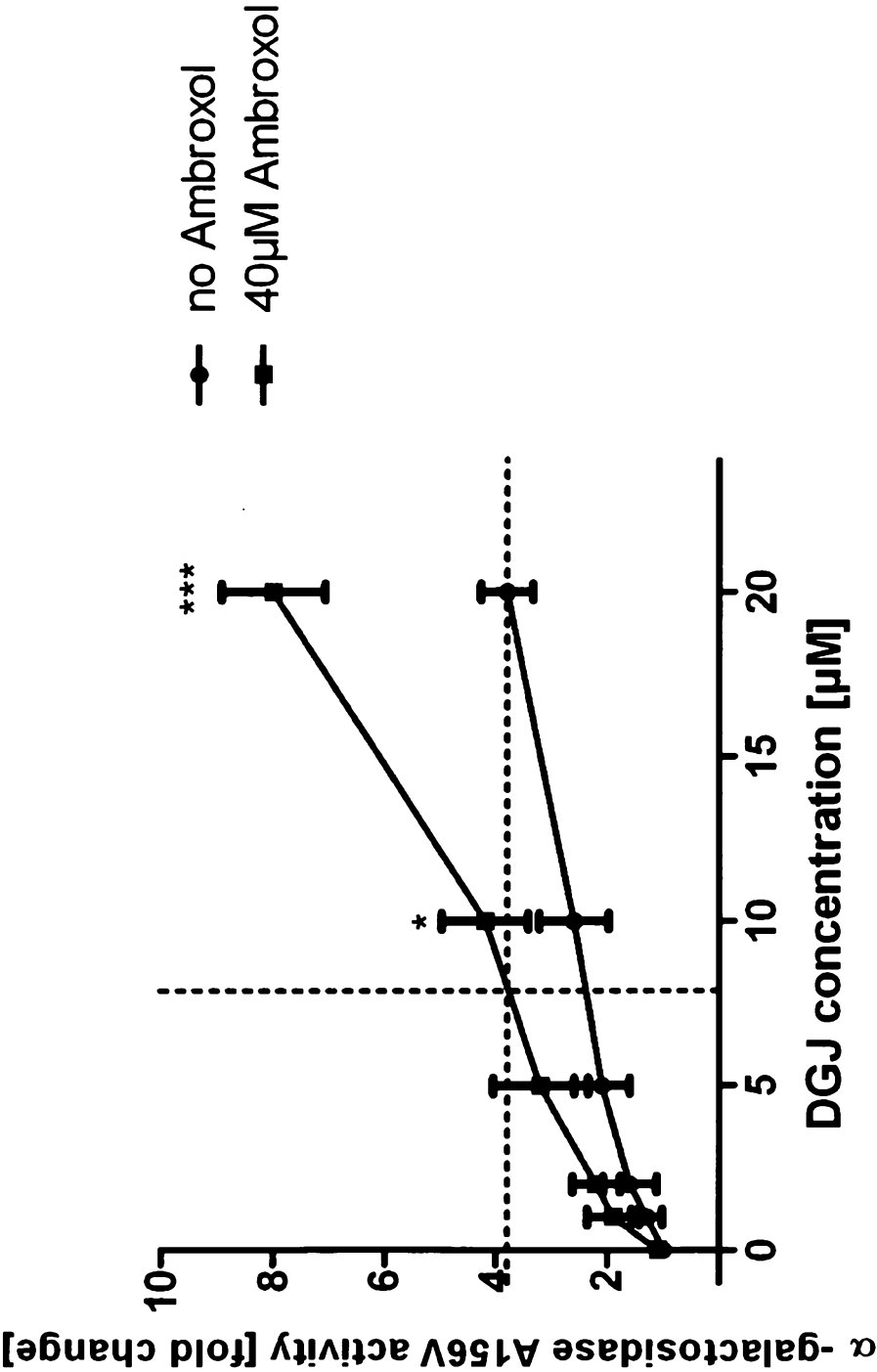


Figure 5A: A combinational treatment of Gla[A156V] can substitute for high DGJ dosages



7/16

Figure 5B: A combinational treatment of endogenous Gla[R301G] can substitute for high DGJ dosages

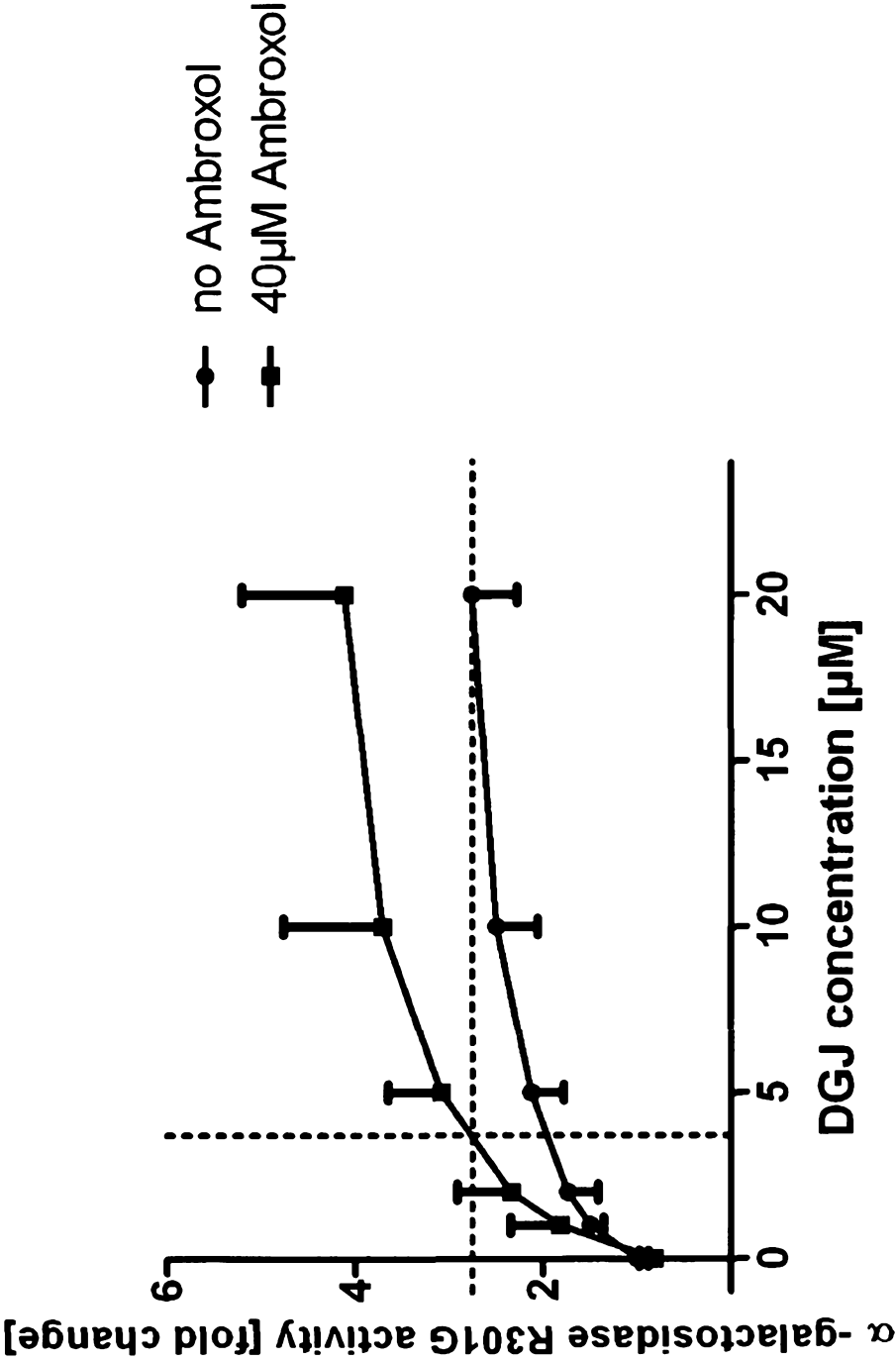


Figure 6:

Bromhexin shows a similar effect to structure analogue Ambroxol

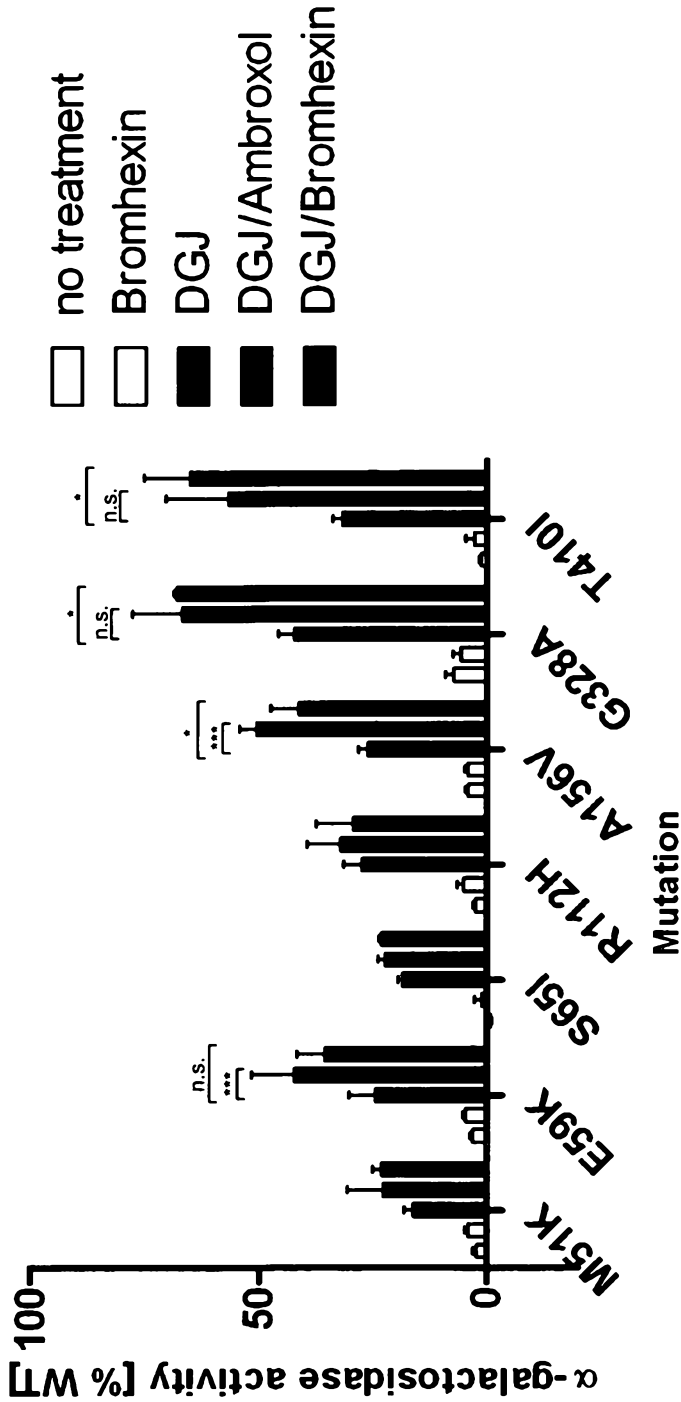


Figure 7A: Effect of NB-DNJ and NB-DNJ/Ambroxol combinational treatment on Pompe Disease mutations

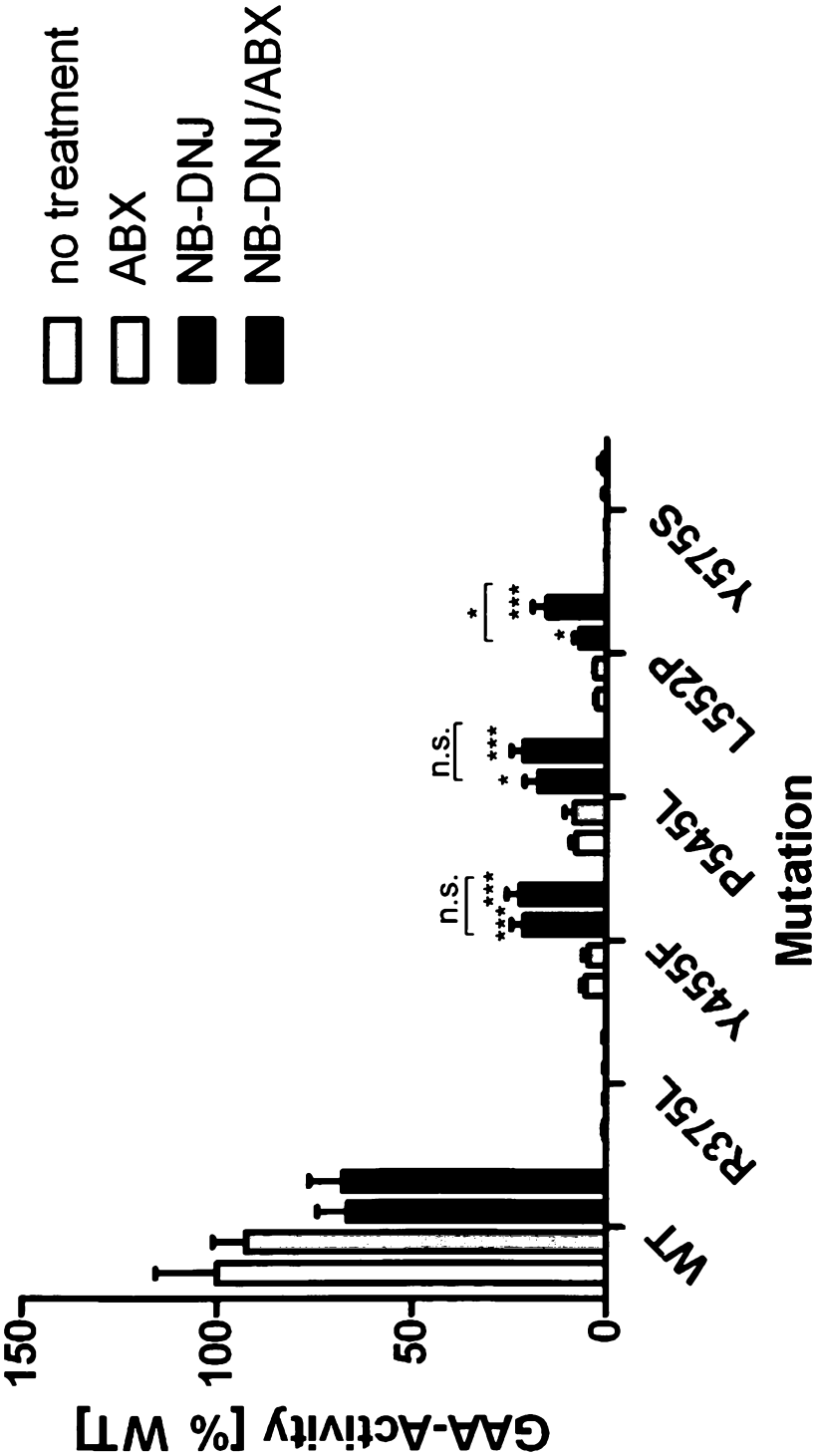


Figure 7B: Effect of DNJ and DNJ/Ambroxol combinational treatment on iminosugar-responsive Pompe Disease mutants [values are mean $\pm$ SEM]

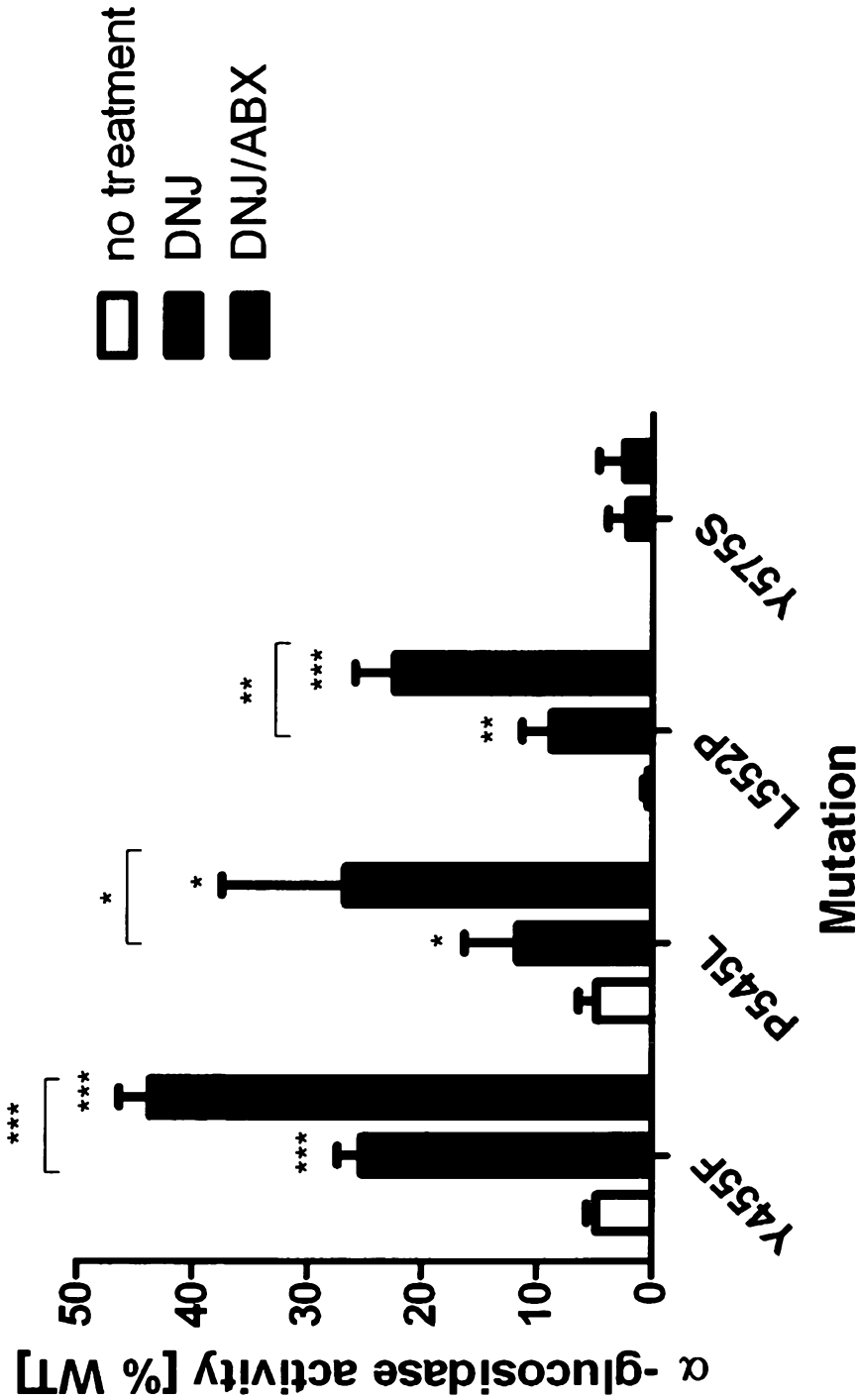
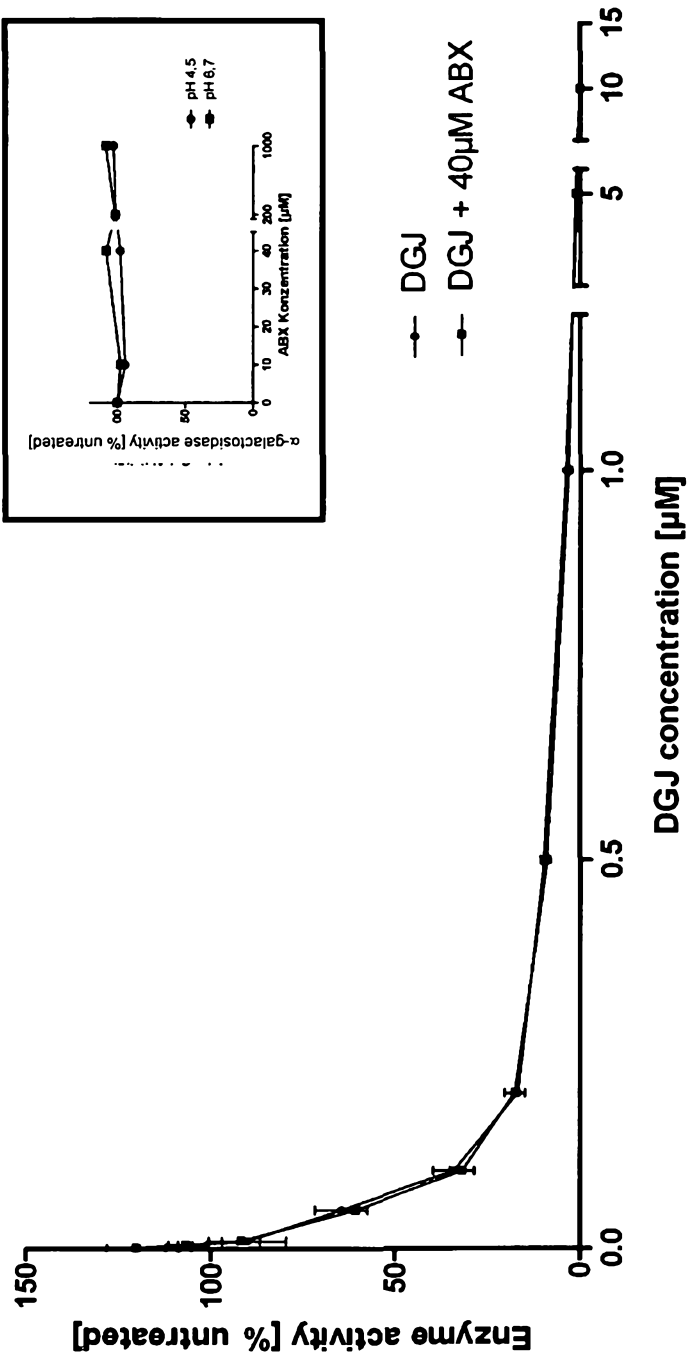


Figure 8: Inhibitory effect of DGJ is not affected by Ambroxol



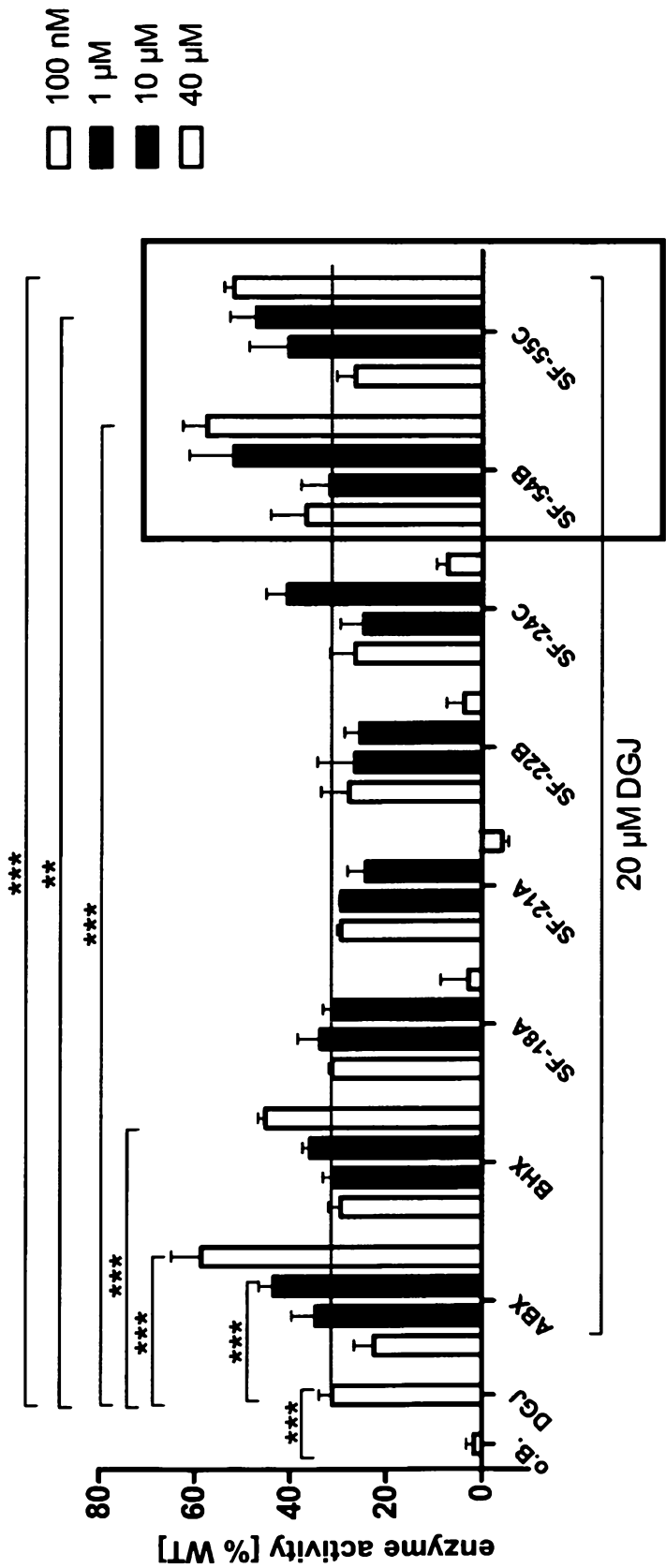


Figure 9A:

Figure 9B:

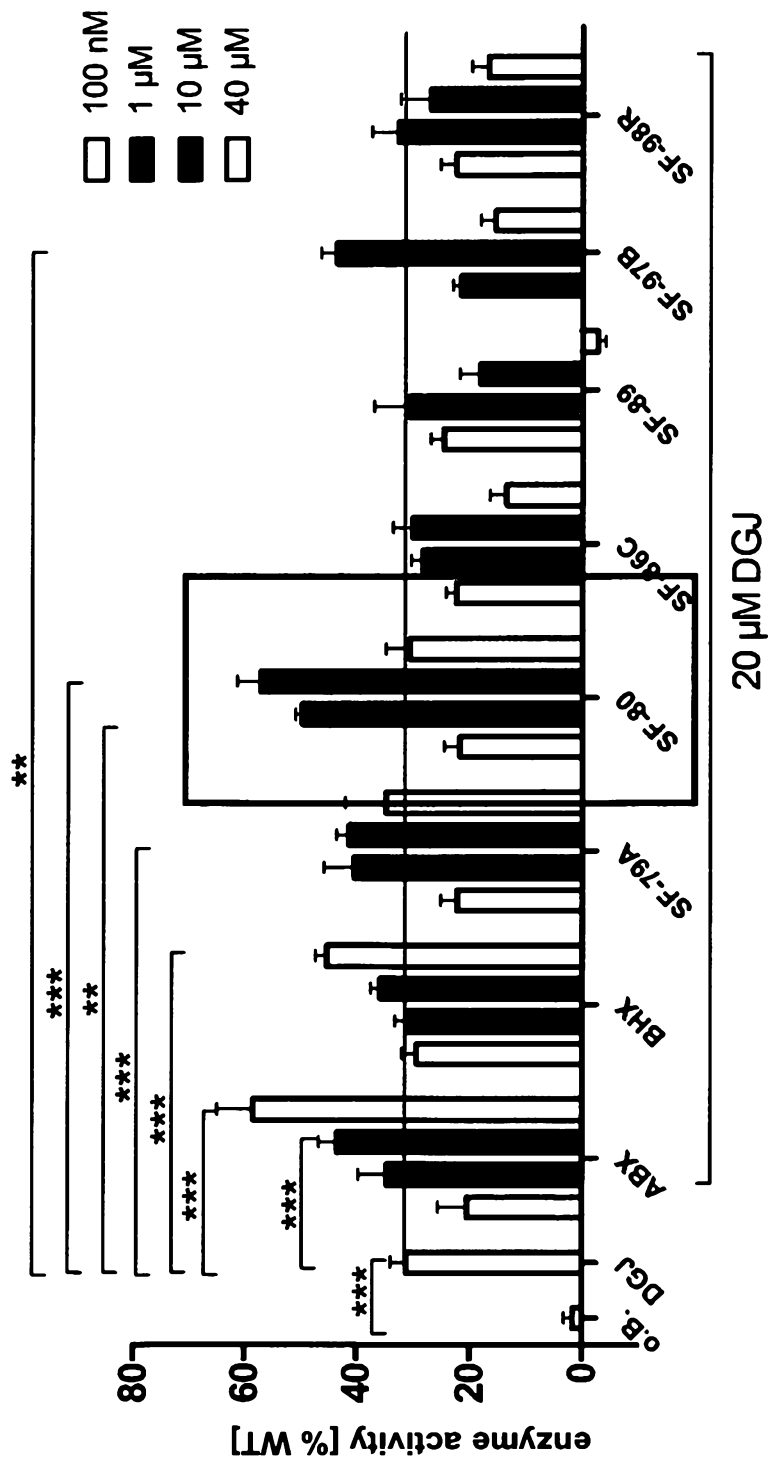


Figure 10: Ambroxol and derivatives in combination with DGJ increase GLA[A156V] activity concentration-dependent

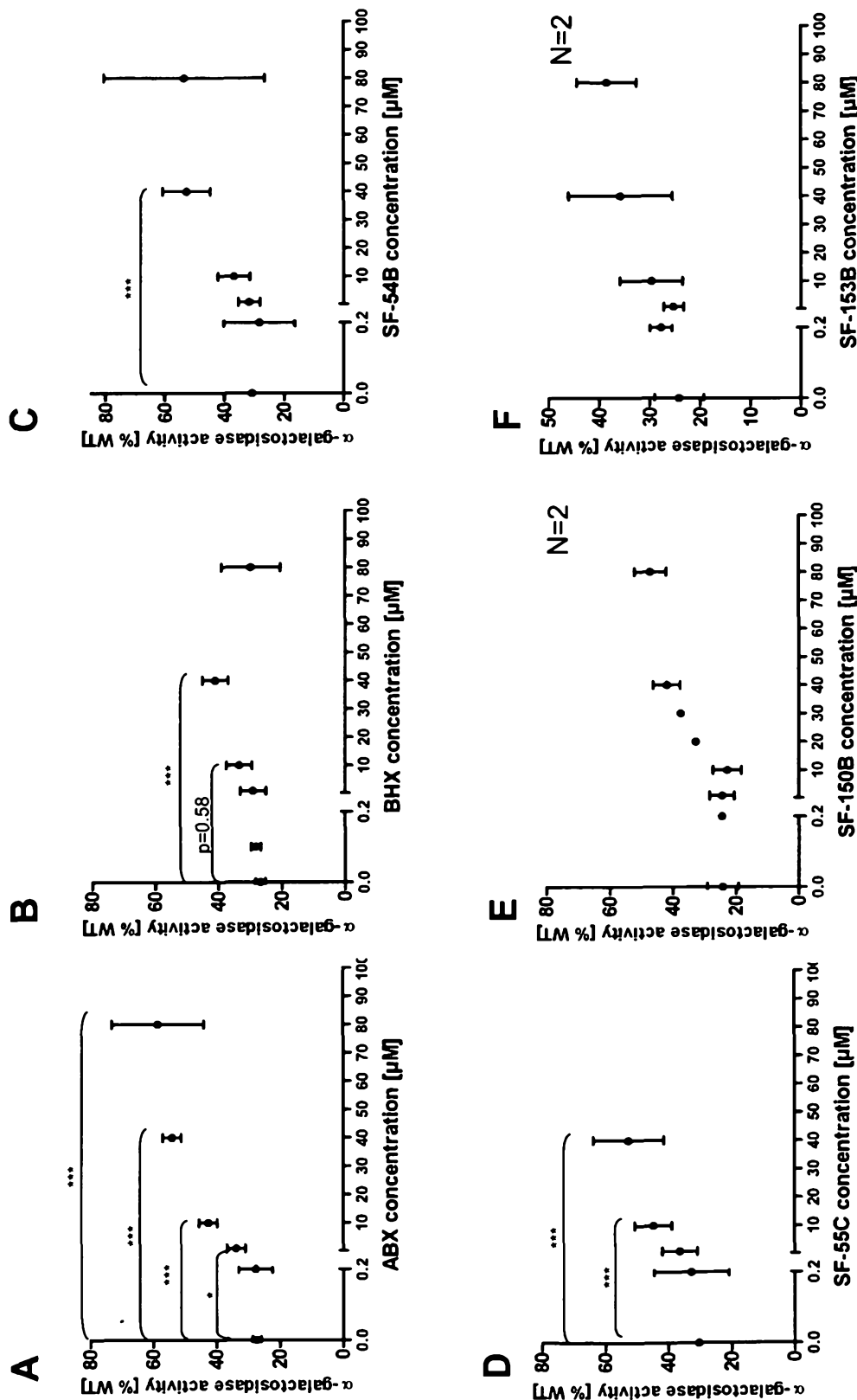
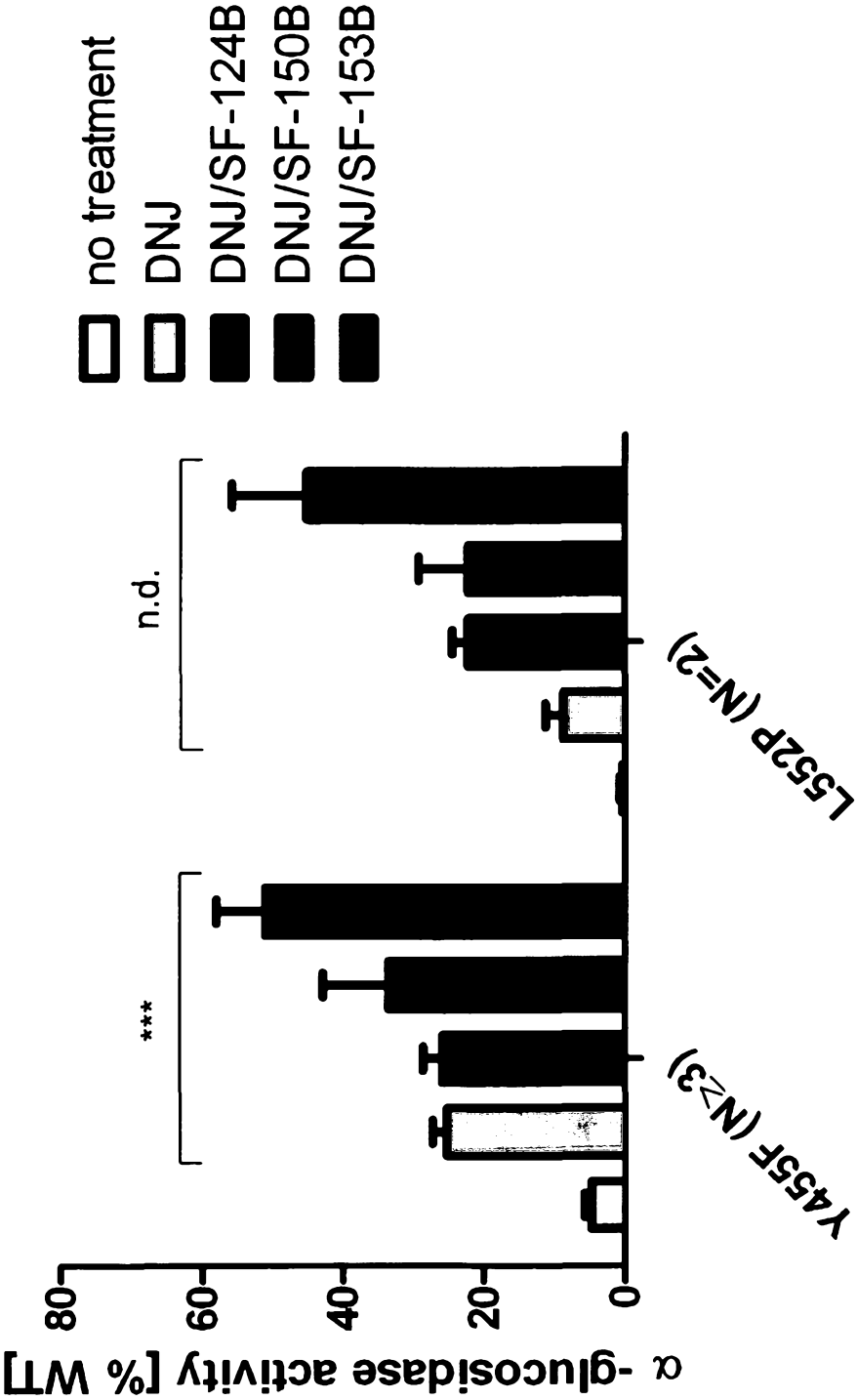


Figure 11A: Pompe mutation Y455F is not responding to Bromhexine-like structures



Figure 11B: Pompe mutations Y455F and L552P are responding to Ambroxol-like structures



SEQUENCE LISTING

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