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(54) Titre : VECTEURS RETROVIRAUX A SPECIFICITE CELLULAIRE DOTES DE DOMAINES DE RECONNAISSANCE
D'ANTICORPS ET PROCEDES DE FABRICATION DESDITS VECTEURS POUR LE TRANSFERT SELECTIF DE
GENES

(54) Title: CELL-SPECIFIC RETROVIRAL VECTORS WITH ANTIBODY DOMAINS AND METHOD FOR THE
PRODUCTION THEREOF FOR SELECTIVE GENE TRANSFER

(57) **Abrégé/Abstract:**

The invention relates to cell-specific retroviral vectors with antibody recognition domains (scFv), which are suitable for cell-specific transduction of a selected mammal cell type (cell targeting), to methods for the production of the cell-specific retroviral vectors and to the use thereof in gene transfers into selected cells. The invention further relates to retroviral packaging cells to obtain cell-specific retroviral vectors according to the present invention.



Abstract

The invention relates to cell-specific retroviral vectors with antibody recognition domains (scFv), which are suitable for cell-specific transduction of a selected mammal cell type (cell targeting), to methods for the production of the cell-specific retroviral vectors and to the use thereof in gene transfers into selected cells. The invention further relates to retroviral packaging cells to obtain cell-specific retroviral vectors according to the present invention.

CELL-SPECIFIC RETROVIRAL VECTORS WITH ANTIBODY DOMAINS AND METHOD FOR THE PRODUCTION THEREOF FOR SELECTIVE GENE TRANSFER

The present invention relates to cell-specific retroviral vectors with antibody recognition domains (scFv) which are suitable for cell-specific transduction of a selected mammalian cell type (cell targeting), methods for the preparation of the cell-specific retroviral vectors and their use for the gene transfer into selected cells. The invention further relates to retroviral packaging cells to obtain the cell-specific retroviral vectors of the present invention.

The majority of retroviral vectors which are presently used in gene therapeutic research are derived from the amphotropic murine leukemia virus (MLV). The host cell range of the amphotropic MLV is determined by the surface envelope protein (SU) encoded by the env gene. The protein products of the env gene form the outer envelope of the retroviral vector. The SU proteins interact with, i.e. bind to a specific protein (receptor) on the surface of the host cell. The env gene products of the amphotropic MLV enable gene transfer into a great number of different mammalian cells. However, a selective gene transfer in particular cell or tissue types of humans or other mammals is not possible, since the receptor for the MLV envelope proteins on the surface of mammalian cells which mediates the entry of amphotropic MLV vectors and gene transfer, is found on nearly all of these cells. Accordingly, the host cell range of the amphotropic MLV is not specific.

A host cell specificity, however, is advantageous e.g. for gene therapeutic use, since in a gene therapy outside of the organism (*ex vivo*) (Anderson et al., Science 256 (1992) 808-813; Yu et al., H. Gene Therapy 8 (1997) 1065-1072) laborious purifications of cells are avoided. It is desired for therapeutic, diagnostic or vaccination use *in vivo*, that retroviral vectors are targeted specifically to the desired host cells and subsequently transfer the therapeutic gene. By modification of the surface envelope protein a restriction of the host cell range of the amphotropic MLV could be achieved. A modification of the surface capsid protein was done by fusion with a hormone domain. A transduction of the cells carrying the specific hormone receptor occurred (Kasahara et al., Science 266 (1994) 1373-1375). Further, the surface envelope protein was modified by fusion with a single chain antibody fragment (single chain variable fragment, in the following referred to as "scFv"). The fragment represented the antigen binding domain of an antibody and is a fusion protein composed of the variable domains Vh and Vl of a monoclonal antibody. Both domains are linked via a glycine and serine oligopeptide [-(ser-gly₄)₃-gly-] which enables the correct folding of the fusion protein (Huston et al., Methods Enzymol. 203 (1991) 46-88, Whitlow et al., Methods: A companion to Methods Enzymol. 2 (1991) 97-105). All modifications of the MLV surface capsid protein using a scFv carried out so far showed that while binding of the

vectors to the host target cell occurred, however, there was no entry into the cell (Russel et al., *Nucleic Acid Res.* 21 (1993) 1081-1085). Furthermore it is known that the surface envelope protein of the MLV generally does not enable for extensive modifications (Cosset et al., *J. Virol.* 69 (1995) 6314-632). Modifications in which a portion of the binding domain of the MLV-SU protein has been replaced led to an incorrect processing and, thus, to a defective transport of the SU protein to the cell surface (Weiss et al., In J.A. Levy (ed.) *The Retroviridae* 2 (1993) 1-108; Morgan et al., *J. Virol.* 67 (1993) 4712-4721; Russel et al., *Nucleic Acid Res.* 21 (1993) 1081-1085). Accordingly, the development of cell-specific retroviral vectors on the basis of MLV with modified surface envelope proteins is only little promising.

Retroviral vectors on the basis of spleen necrosis virus SNV are more suitable for a targeted gene transfer into e.g. human cells, since the surface envelope protein of SNV enables for extensive modifications after which still correct processing occurs (Martinez and Dornburg, *Virol.* 208 (1995) 234-241; Chu et al., *Gene Therapy* 1 (1994) 292-299; Chu and Dornburg *J. Virol.* 69 (1995) 2659-2663). At least two components are required for the preparation of such vectors. On the one hand a so-called expression construct is to be prepared enabling packaging in and transfer through a retrovirus. The expression construct comprises a coding DNA fragment of the desired gene product, e.g. a gene for gene therapy or as a vaccine. The expression construct has to comprise a nucleotide sequence which is referred to as packaging signal ψ and controls the efficient packaging of mRNA in retroviral particles. Furthermore a packaging or helper cell is required which provides the gag, pol and env gene products of SNV, without packaging of the gag, pol and env genes into a retrovirus. The gag, pol and env genes which are present in the packaging cell have to be ψ -negative. Following transduction of the expression construct by means of transfection of the respective plasmid DNA into the packaging cells, retroviral particles are released into the cell culture supernatant, which particles contain the expression construct and are only able to transduce said construct but not the gag, pol and env genes into the target cell. Said vectors are replication incompetent and only pass one cycle of replication. The general method for the preparation of replication incompetent retroviral vectors is known in the prior art (Weiss et al., In J.A. Levy (ed.) *The Retroviridae* 2 (1993) 1-108; Morgan et al., *J. Virol.* 67 (1993) 4712-4721; Russel et al., *Nucleic Acid Res.* 21 (1993) 1081-1085; Cosset et al., *J. Virol.* 69 (1995); Martinez and Dornburg, *Virol.* 208 (1995) 234-241; Chu et al., *Gene Therapy* 1 (1994) 292-299; Chu and Dornburg, *J. Virol.* 69 (1995) 2659-2663).

The tropism (host cell specificity) of spleen necrosis virus is determined by the surface envelope protein (SU protein) encoded by the env gene of SNV. The wild type SNV surface envelope protein does not allow for a selective gene transfer into particular human cells or tissues since the specific acceptor protein (receptor) is not present on the surface of human cells (Dornburg, *Gene Therapy* 2 (1995) 1-10). Therefore, a method has been

developed to substitute the SU protein of SNV by the antigen recognizing domains of antibodies. Said [SNV-scFv-Env] vectors with the two different scFv known heretofore were able to transmit the psi positive reporter gene, bacterial β -galactosidase, into the selected human target cells. Said scFv are directed against the hapten dinitrophenol (DNP) or against an unknown surface molecule on colon CA cells and other cancer cells, respectively. (Chu et al., *Gene Therapy* 1 (1994) 292-299, Chu et al., *BioTechniques* 18 (1995) 890-899; Chu and Dornburg, *J. Virol.* 71 (1997) 720-725). A packaging cell line (DSH-CXL) has been developed, containing the psi-negative SNV genes gag, pol and env as well as the psi-positive reporter gene expression construct (pCXL). Following transfection of the packaging cell using plasmid DNA of another env expression gene (pTC53), in which the entire surface envelope protein was substituted by a single chain antibody fragment (scFv), retroviral vectors were released into the cell supernatant which in addition to the wild type surface envelope protein also carried the chimeric [scFv-Env] surface protein on their surface. By means of said vectors the reporter gene could be transferred into the target cells specific for scFv, canine osteosarcoma cells (D17), which were conjugated with DNP, or HeLA cells (human cervical carcinoma cells), respectively. However, this method described for the preparation of cell-specific retroviral vectors has the disadvantage that only already known and cloned scFv may be used. Further, it has been found by us that not every scFv is suitable as a portion of a [SNV-scFv-Env] vector for cell transduction (transfer of the desired gene to the target cell).

Generally, the gene transfer into mammalian cells by means of retroviruses has the following benefits:

- Normally, one copy of the desired gene is transferred into the mammalian cell.
- Generally, the desired gene is transferred without mutation or rearrangements.
- Stable introduction of the desired gene into the genome of the target cell occurs.

Furthermore, it is desired that the retroviral vector has a particular cell-specificity by which e.g. the therapeutic gene may be introduced into a selected cell population.

Therefore, it is an object of the present invention to provide cell-specific retroviral vectors having antibody recognition domains for selective gene transfer into mammalian cells as well as a universal method for the preparation thereof. It is possible to improve the gene transfer by means of said vectors. A further object of the invention is to provide retroviral packaging cells for obtaining the vectors according to the present invention. The solution of said objects is obvious from the claims, the following description and the figures.

The object is solved by the method of the present invention for the preparation of cell-specific retroviral vectors comprising the following steps of: a) Immunizing a mammal with one or more cell population(s), b) Isolation of RNA from the mammal immunized, comprising the B cell RNA, c) preparation of cDNA portions of the variable regions of the immunoglobulin heavy and light chains from the isolated RNA by means of RT-PCR using

primers for the immunoglobulin heavy and light chain wherein the primers comprise the nucleic acid sequence for an oligopeptide linker, d) ligation of the cDNA portions to scFv-cDNAs, e) ligation of the scFv-cDNAs into a phagemid vector and transformation of a host bacterium with said phagemid vector, f) isolation of phages which bind to the cell population(s) used in step a) by means of selection, g) isolation of cell-specific phages from the phages obtained in step f) which only bind to the cell population(s) used in step a) by means of selection, h) excising the scFv encoding DNA fragments from the cell-specific phages obtained in step g) and ligation into a psi-negative retroviral Env expression vector, i) transformation of the resulting Env-scFv expression vector into a packaging cell, and j) isolation of retroviral vectors secreted from the packaging cell.

Optionally, the method according to the present invention comprises further isolation of cell-specific phages obtained in step g). Optionally, the method of the invention further comprises the step: k) isolation of retroviral vectors secreted by the packaging cell, which transduce the cells of the cell population(s) by means of selection. Furthermore, the steps f) and/or g) may be repeated at least once.

A method is preferred in which the immunized mammal is selected from the group consisting of mouse, rat, guinea pig, rabbit, goat or sheep. Also preferred is a method in which the cell population(s) is/are selected from the group consisting of man, mouse, rat, sheep, cattle or pig. Particularly preferred is a method, in which the cell population(s) is/are selected from the group consisting of T cells, epithelial cells, muscle cells, stem cells, neural cells, hematopoietic cells, carcinoma cells or liver cells. A method is preferred in which the env gene is derived from spleen necrosis virus (SNV). Particularly preferred is a method in which the expression vector is the vector with the designation pTC53.

The cell-specific retroviral vectors obtainable by means of the method of the invention may be used as medicaments. Preferred is the use for the preparation of a medicament for gene therapy, vaccination therapy or diagnostics. Particularly preferred is the therapy of cystic fibrosis, ADA deficiency, HIV infections, leukemia, chronic granulomatosis.

Furthermore, the invention is solved by the provision of retroviral packaging cells for obtaining retroviral vectors of the invention, transformed both with one or more psi-negative expression construct(s) expressing gag, pol and/or env gene products and with a psi-negative Env-scFv expression construct according to claim 1h). A packaging cell is preferred further comprising a psi-positive expression construct, comprising a nucleic acid fragment, which is to be introduced into the cell to be transduced by the retroviral vector. Particularly preferred is a packaging cell wherein the nucleic acid fragment comprises a therapeutic gene or its DNA fragment and/or a reporter gene. Particularly preferred is a packaging cell wherein the therapeutic gene or its nucleic acid fragment comprises CFTR, phox91, ADA, IL-16, p53 or revM10 gene or vaccination genes, e.g. recombinant gp120 and IL-16. Further particularly

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preferred is a packaging cell wherein the reporter gene comprises β -galactosidase, green fluorescent protein, luciferase or neomycin.

5 According to one aspect of the present invention, there is provided a method for the production of cell-specific retroviral vectors comprising the following steps: a) immunizing a mammal with one or more cell population(s), b) isolating RNA from the immunized mammal, comprising B cell
10 RNA, c) producing cDNA regions of the variable regions of an immunoglobulin heavy and light chain from the isolated RNA by means of RT-PCR with primers for the immunoglobulin heavy and light chain wherein the primers comprise a nucleic acid sequence which encodes an oligopeptide linker, d) ligating
15 the cDNA regions to form scFv-cDNAs, e) ligating the scFv-cDNAs into a phagemid-vector and transformation of a host bacterium with the phagemid vector, f) isolating phages binding to the cell population(s) used in step a) by means of selection, g) isolating cell-specific phages from the
20 phages obtained in step f) which only bind to the cell population(s) used in step a) by means of selection, h) excising the scFv-encoding DNA fragments from the cell-specific phages obtained in step g) and ligating the scFv-encoding DNA fragments into a psi-negative retroviral
25 Env expression vector to form an Env-scFv expression vector comprising a chimeric Env-scFv surface protein encoding DNA sequence, i) transforming said Env-scFv expression vector into a packaging cell, wherein the packaging cell comprises a psi-negative env, gag, pol expression construct; and j)
30 isolating the retroviral vectors secreted by the packaging cell.

The figures are meant to illustrate the invention.

Fig. 1 schematically shows the ENV-scFv expression construct pTC53, pT-scFv and pT/zeo, the transfection thereof into the packaging cell DSH-CXL which secretes the vectors of the present invention.

Fig. 2 schematically shows the preparation, isolation and selection of the vectors of the invention.

Fig. 3 is a schematic illustration of an immunoglobulin and the scFv resulting therefrom. Furthermore, scFv display phages and SNV-scFv-Env vectors are schematically depicted.

Fig. 4 shows the nucleic acid sequence of pTC53.

The term amphotropic virus used herein means infection and replication in murine and human cells, in contrast to an ecotropic virus which only replicates in murine cells. The term retroviral vector used herein means replication deficient retroviral virus particle which instead of retroviral mRNA may transmit a foreign introduced RNA of a gene, e.g. of a therapeutic gene or a fragment thereof or of a reporter gene. The term antibody recognition domain (scFv) used herein means an antigen binding site of an antibody comprising V_H and V_L chain. The term SNV used herein represents spleen necrosis virus with its strains and substrains. SNV belongs to the avian reticulo endotheliosis viruses (REV), type D retrovirus.

To provide the cell-specific antibody recognition domains (scFv) a new combinatory phage cDNA library of the variable domains of the light and heavy chains of the immunoglobulins is prepared. For this purpose, a mammal, e.g. a mouse, rat, rabbit, guinea pig, goat or sheep is immunized with a sufficient titer of one or more cell population(s) in a usual manner. The cell population is the cell type forming a surface receptor to which the retroviral vectors of the present invention specifically bind. The cells may be derived from a mammal which is different from the mammal to be immunized, e.g. from human, mouse, rat, sheep, cattle or pig. The cells may be such cells in which for example a somatic gene therapy, a vaccination therapy or diagnostics is to be carried out. Typical examples of such cells are T-cells, liver cells, muscle cells, neural cells, fibroblasts, epithelial cells, stem cells or hematopoietic cells. For immunization one or more cell population(s) may be simultaneously administered to the mammal depending on the cell population(s) for which the retroviral vector of the present invention has to be specific.

For the preparation of a cDNA library B cell RNA of the immunized mammal is first isolated in a known manner. The mRNA sequences of the regions of the heavy and light chain (V_H and V_L) of the immunoglobulins responsible for antigen recognition, are transcribed into cDNA and amplified in a usual manner by means of reverse transcription

and subsequent polymerase chain amplification. The primer pairs and their sequences for V_H and V_L regions are known to the person skilled in the art. For example, they are contained in the kit commercially available from Pharmacia company, or may be obtained from known data bases (EMBL), respectively. It is known to the skilled artisan that he has to use different primer sequences for every immunized mammalian species. The sequences are also contained in the known data bases. The cDNA fragments of the V_H and V_L regions are then linked to scFv-cDNAs by means of a ligase reaction in a usual manner. It is obvious to the skilled artisan that during ligation different combinations of cDNA fragments are prepared. The resulting scFv-cDNAs may then be cloned into a phagemid vector, e.g. pCANTA 5E phagemid, Pharmacia company. Subsequently, host bacteria, e.g. *E. coli* TG1 are transformed with the phagemid vector.

The recombinant phages produced by the bacteria are then isolated in a usual manner and selected for the presence of cell-specific scFv peptides. The phages are then contacted in a usual manner with the cell population(s) which have been used for immunization. Phages not binding to cells do not carry a specific scFv peptide and are removed by means of washing steps in a usual manner. Phages binding to the cells present the desired scFv peptide on their surface and are eluted in a usual manner. Phages presenting the desired scFv peptide are amplified by allowing them to infect host bacteria in a usual manner. This selection step may be repeated once or several times to enrich the binding phages. This procedure is referred to as panning. The phages are subjected to a further selection after panning or directly after the first selection step. For this purpose the phages are contacted with one or more other cell population(s) which are different from the cells used for immunization. Phages not binding to said cells present a cell-specific scFv peptide. They are isolated from the cell supernatant in a usual manner and are used for a host bacteria infection for amplification. Also this selection step may be repeated once or several times (Marks et al., *BioTechnology* 10 (1992) 779; Clackson et al., *Nature* 352 (1991) 624; Marks et al., *J. Mol. Biol.* 222 (1991) 581; Chaudhary et al., *Proc. Natl. Acad. Sci USA* 87 (1990) 1066; Chiswell et al., *TIBTECH* 10 (1992) 80; McCafferty et al., *Nature* 348 (1990) 552; Huston et al., *Proc. Natl. Acad. Sci. USA* 85 (1988) 5879).

For the preparation of the cell-specific retroviral vectors of the present invention the above described phage cDNA library is used as a starting material. The scFv-cDNAs of the phages remaining after the second selection step and the panning method which has optionally been carried out are excised in a usual manner from the phage DNA and inserted into a retroviral Env expression gene. The retroviral env expression gene may be derived from SNV. A typical example for a SNV-scFv-env expression construct is pTC53. The sequence is shown in Figure 4B. A typical example for a wild type (wt) SNV-env expression construct is pIM29.

The construction of the expression plasmids encoding wt-SNV-ENV proteins, e.g. pIM29 and the chimeric SNV-scFv-ENV proteins has been previously described by Chu et al. (*J. Virol.* 71 (1997) 720-725). The expression of the DNA coding for the wt-env gene is controlled by an MLV promoter. The env-cDNA was excised from a plasmid encoding the complete SNV virus via restriction sites *SacII* and *AvrII* and introduced by insertion into a linker (L). To enable a correct processing of the protein pIM29 contains the polyadenylation site of Simian Virus 40 (SV 40). Thus, from this plasmid the expression of the wt-env gene may occur, so that following proteolytic cleavage of a precursor protein the outer glycoprotein (SU) and the transmembrane protein (TM) are obtained. However, other plasmids, promoters, linkers, polyadenylation signals known to the skilled artisan and further DNA elements required for a correct processing may be used.

For the expression of SNV-scFv-ENV proteins scFv obtained in a known manner are introduced into a SNV-ENV expression construct, e.g. pTC53, in a usual manner. The restriction recognition sites for the enzymes *SfiI* and *NotI* present in pTC53 enable the molecular cloning of e.g. scFv between the SNV-env- leader sequence and the DNA region encoding the transmembrane protein (TM protein). The protease cleavage site between SU and TM present in wt-ENV is deleted in pTC53, so that a fusion protein is expressed which consists of a single chain antibody fragment at the N-terminus and of the SNV-TM at the C-terminus. The regulatory elements, such as MLV promoter and SV40 polyadenylation signal are identical to those of the pIM29 vector. For enhancing the expression of a chimeric env gene an adenoviral leader sequence, e.g. *Avtl* (Sheay et al., *BioTechniques* 15 (1993) 856-861) is inserted into the expression plasmid pTC53. A zoezin cassette (pSV2zeo; Invitrogen Comp., The Netherlands) functions to select stably transfected cells, so that single cell clones may be established.

The psi-negative SNV-scFv-env expression construct may be introduced into packaging cells by means of electroporation or other known methods. A typical packaging cell is for example DSH-CXL. Further, packaging cells have psi-negative env, gag, pol expression constructs and for the desired gene transfer into the specific target cells a further psi-positive expression construct comprising for example a gene or DNA fragment for gene therapy, vaccination therapy or a reporter gene for diagnostics. Following transfection of the packaging cells a transient expression and release of retroviral vectors which present in addition to natural SU proteins recombinant SU-scFv proteins into the cell culture supernatant occurs. The retroviral vectors may then be used for transduction of the target cell, i.e. the cell population used for immunization in a usual manner. Optionally, said step may be a further selection step. Only the retroviral vectors of the invention which transduce the target cell in a sufficient manner are further employed. These vectors may be subjected to a further selection step. The vectors may be used in a conventional manner for transduction of cell populations different from the cell population used for immunizing said mammal.

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Thus, vectors not transducing the other cells but only the target cells may be obtained in a double selection step.

For establishing of stable packaging cell lines which constitutively release the retroviral vectors of the invention, a selection marker, e.g. the zeocin resistance gene (Invitrogen), may be inserted in a usual manner into the scFv expression construct, e.g. pTC53. The scFv expression constructs provided with the zeocin resistance gene are transferred into the packaging cells e.g. by means of the liposome technique (lipofectamine, Gibco BRL). After a selection of about two weeks of the transfectants in zeocin containing culture medium cell clones may be established which transduced target cell populations with a titre of about 10^4 - 10^6 retroviral vectors pro ml depending on the scFv-cDNA fragment.

The gene transduced with the retroviral vectors of the invention into the target cell population or populations may for example be the RNA of a therapeutic gene or a fragment thereof. Therapeutic genes may for example be the CFTR gene, ADA gene, LDL receptor, β -globin, factor VIII or factor IX, dystrophin gene. The target cells in the case of the CFTR gene would be e.g. lung epithelial cells, in the case of the ADA gene the stem cells of bone marrow or T lymphocytes, for LDL receptor the liver cells, for dystrophin gene skeletal muscle cells, for β -globin gene hematopoietic stem cells, for factor VIII or factor IX fibroblasts and liver cells. It is obvious to the skilled artisan that this listing represents only a selection of therapeutic genes and other genes may also be used for a gene therapy. The DNA fragments of a therapeutic gene comprise for example antisense nucleic acids or ribozymes. Further, DNA fragments may comprise portions of a gene

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containing the trinucleotide repeats of e.g. the fragile X gene.

Further, the RNA of a reporter gene, e.g. β -galactosidase, GFP, luciferase or neomycin, may be introduced into the retroviral vectors of the present invention. The reporter genes enable the determination whether the target cells have been transduced with the retroviral vectors.

Further, the RNA of a gene or a fragment thereof may be transduced into the target cell for vaccination purposes. A typical vaccination gene is for example the recombinant gp120 or gp160 of HIV. The transduction of immune cells with these genes or fragments stimulates the antibody formation against viral gene products.

The vectors of the present invention may for example be applied by i.v. or i.m. injections. The packaging cells of the present invention may however be enclosed into e.g. organoids (Teflon* bags), which are then implanted into the organism and secrete the vectors according to the present invention into the blood stream or tissue. Further application forms are obvious to the person skilled in the art.

The retroviral packaging cell of the invention for obtaining the pseudotyped retroviral vectors of the present invention is provided by transfecting a cell line e.g. a human cell line with psi-negative expression construct expressing the gag and pol gene products of SNV and

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with the psi-negative SNV-Env expression construct and/or psi-negative SNV-scFv-env expression construct in a conventional manner.

Furthermore, packaging cells may be used, which already contain the psi-negative expression constructs for gag and pol gene products. Into such packaging cells only the psi-negative expression construct for the virus envelope and the psi-positive expression construct for the nucleic acid sequence to be transduced into the target cell have to be transfected. Methods for transfection of the expression constructs are known to the skilled artisan. By the packaging cells of the invention, retroviral vector particles are released into the cell supernatant which contain the expression construct, but not the constructs encoding GAG, POL and ENV proteins. Thus, only the desired e.g. therapeutic gene or reporter gene is transferred into the target cell.

The illustrated invention opens up following possibilities:

- Genes, gene fragments or other nucleic acid sequences may be transferred into selected mammalian cells.
- further enhancement of the efficiency of the nucleic acid transfer may be achieved by improvement of the env gene constructs.
- gene therapy, labeling and vaccination strategies may be developed, for which a selective nucleic acid transfer into selected mammalian cells is desirable.

The following examples illustrate the invention and are not intended to be limiting:

1. Isolation and cloning of cell-specific scFv

For the preparation, isolation and selection of cell-specific scFv a mouse was immunized with the human T cell line T-C8166 (Clapham et al., *Virology* 158 (1987) 44-51) in a conventional manner, the spleen removed and RNA was isolated. Cloning of the scFv-cDNAs was carried out with the commercially available kit of Pharmacia company according to the manufacturer's instructions. The resulting phages were examined in a conventional manner with respect to their binding characteristics to target cells. There were isolated 150 phages which specifically bound to the target cells. The 150 thus obtained cell-specific scFv were used to prepare the SNV-scFv vectors according to the present invention.

2. Cloning of specific scFv-cDNA fragments into Env expression constructs

The scFv-cDNAs of 150 cell-specific scFv were excised in a usual manner from the phagemid DNA and each of the DNAs was ligated into the expression construct pTC53. pTC53 was obtained by modification of the universal eukaryotic vector pRD114 (Chu et al., *J. Virol.* 71 (1997) 720-725; Sheay et al. *BioTechniques* 15 (1993) 856-861; Chu et al., *BioTechniques* 18 (1995) 890 - 895). In this vector the SNV-wt-env gene was deleted except for the leader sequence and the transmembrane-protein encoding cDNA. An additionally

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inserted spacer enables the insertion of a foreign DNA (here scFv-cDNA) following the ENV-leader sequence via the restriction recognition site NaeI. The sequence of pTC53 is shown in figure 4. For the insertion of the scFv-cDNA the Env-expression construct pTC53 was modified so that Sfi I and Not I specific restriction endonuclease recognition sites are inserted between the SNV-leader sequence and SNV-transmembrane sequence (TM) in a usual manner. For this purpose a recombinant PCR is carried out in a usual manner starting from the DNA of the plasmid PKA1558 (Scov. H. & Andersen, K.B., 1993) and the DNA coding for the anti-transferrin receptor scFv so that via Nru I (5' and 3') an insertion of the amplified fragment into the Nae I restricted pTC53 is possible. The thus inserted fragment contains the multiple Sfi I/Not I cloning site since the primers used further include a neighboring Sfi I or Not I recognition site, respectively, in addition to the terminal Nru I recognition site. For recombinant PCR following primers were used:

20 PKATFNNRu+:

5'-GGGCCCTCGCGAGCGGCCCGGCCGACATCAAGATGACCCAGTCTCCA-3'

Nru I Sfi I

PKATFNNRu-:

5'-GGGCCCTCGCGATGCGGCCGCTGAGGAGACTGTGAGAGTGGTGCC-3'

25 Nru I Not I

The PCR conditions were: 94°C/3 min, 94°C/1 min, 59°C/1 min, 72°C/1 min., 25 cycles, 72°C/10 min and then cooling to 4°C. The PCR fragment was gel-electrophorized, extracted from the gel matrix (Qiaex, Qiagen Comp.) and

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ligated in a conventional manner with the plasmid pTC53 opened with Nae I.

The scFv-cDNAs from the phagemid (pCANTA 5E) were excised by means of the restriction endonucleases Sfi I and Not I. For this purpose the phagemid-plasmid DNA was prepared by means of known methods, and in each case 8 μ g of plasmid DNA were digested with 60 U each of the restriction endonucleases Sfi I and Not I at 50°C for 1.5 h and subsequently at 37°C for 1.5 h. The reaction batch occurred in a volume of 200 μ l which was supplemented with 20 μ l BSA (10 x conc.) and 20 μ l reaction puffer 3 (10 x conc.). Upon completion of the reaction period the batch was electrophorized on a 1% agarose gel. Following separation the scFv-cDNA specific band (about 750 bp) was purified from the agarose gel by means of known methods.

The purified fragment was ligated with the Env expression construct pTC53 which has also been opened with the restriction endonucleases Sfi I and Not I. For this purpose

equimolar amounts of the scFv-cDNA fragment and pTC53 fragment were supplemented with 200 U T4 ligase and 1.5 μ l 10x ligase buffer in a 15 μ l volume. The batch was incubated at 4°C over night. To enable an efficient transformation of bacteria the bacterial strains TOP10F' and JS5 were made competent by means of a modified method according to Hanahan (1983). Following inoculation of 100 ml LB-medium with 500 μ l of an overnight culture, the bacterial suspension was incubated at 37°C up to a density (OD_{550}) of 0.6. Subsequently, the bacteria were chilled on ice, pelleted at 6000 rpm and 4°C (Minifuge RF, Heraeus, Hanau) and resuspended in 40 ml TFB1 buffer (30 mM KOAc, 100 mM RbCl₂, 10 mM CaCl₂, 15% glycerol, pH 5.8, adjusted with acetic acid, thereafter filter sterilized). After an incubation period of 15 min on ice and centrifugation at 6000 rpm and 4°C the bacterial pellet was resuspended in 4 ml of TFB2 buffer (10 mM MOPS, 75 mM CaCl₂, 10 mM RbCl₂, 15% glycerol, pH-Wert 6.5, adjusted with KOH solution, thereafter filter sterilized). The bacterial suspension was then divided into aliquots of 100 μ l each and then shock frozen on dry ice. The storage was carried out at -70°C. For transformation, 100 μ l of competent bacteria were thawed on ice and following addition of 1-2 μ l of the respective ligation batch incubated on ice for 30 min. After a subsequent temperature shock (45 s at 42°C, thereafter 2 min on ice) the bacteria were added with 500 μ l SOC medium (GIBCO/BRL, Eggenstein) and cultivated for 1 h at 37°C for expression of antibiotic resistance in a bacterial shaker. The bacterial suspension was streaked out on LB agar plates supplemented with the antibiotic ampicillin and incubated at 37°C over night.

The preparation of plasmids from bacteria (*E. coli* TopF10) was done with the QIAGEN plasmid kits of QIAGEN company, Hilden. For the preparation of a low amount of plasmid DNA the bacteria of a 15 ml overnight culture (LB medium with 50 μ g/ml ampicillin) were lysed with the solutions provided by the manufacturer and purified via an anion exchange column (tip 20). For the preparation of large amounts of plasmid DNA (maxi preparation) 400 ml overnight cultures were prepared.

3. Selection of retroviral vectors

Transient transfection of the scFv-pTC53 expression constructs into the packaging cell DSH-CXL by means of electroporation: for each electroporation 2×10^6 DSH-CXL cells were resuspended in 480 μ l PBS and added to a Gene-Pulser cuvette (0.4 cm electrode, gap 50, Biorad, Munich). Thereafter 20 μ g of recombinant plasmid DNA were added to the cell solution. The content of the cuvette was subjected to an electric pulse in an electroporator (Gene-Pulser Apparatus, Biorad, Munich) at 270 V and 960 μ F. After 10 min of incubation of the cuvette on ice the cells were added into 20 ml of fresh culture medium in a medium sized cell culture flask (Nunc, Wiesbaden). The next day the DSH-CXL cells were added with fresh medium and cultivated.

The virus containing supernatant of the transfectants was used for transduction of the target cells. The day before transduction the C8166 target cells were transferred to fresh medium in a ratio of 1:2. The supernatants were filter sterilized with a 0.45 μ m filter (Sartorius). 7 ml of the supernatant were directly employed for transduction of 2×10^5 C8166 cells. To stabilize the junction of the retroviral vectors to the cell surface, 40 μ g/ml Polybrene were added. Following 2 h of incubation at 37°C the cells were washed with PBS and transferred into fresh culture medium.

Detection of β -galactosidase activity (X-Gal assay): For examination of a successful transduction an X-Gal assay was carried out after 72 hours according to a modified method of Sanes et al. (1986). The cell culture supernatant was removed and the cells washed with PBS without (Ca^{2+} and Mg^{2+}). Subsequently, the cells were overlaid with a fixation solution (2% formaldehyde, 0.2% glutaraldehyde in PBS) for 5 min and washed with PBS. Thereafter, the cells were resuspended in 3 ml of X-Gal reaction mix solution (1 mg/ml, 5 mM K-Ferricyanide, 5 mM K-Ferrocyanide, 2 mM MgCl_2). After an approx. 4 h period of incubation of the batch at 37° blue staining of the transduced cells occurred.

6 of the 150 tested scFv-pTC53 expression constructs were cell-specific (M8, K6, 7A5, 7E10, 6C3, 7B4). That means, that per cell-specific construct 10-20 blue-stained C8166 cells could be recognized. Compared to non-cell specific scFv-clones, this result is significant. Stable packaging cell lines were generated from 6 cell-specific scFv expression constructs.

4. Establishment of stable packaging cell lines.

Preparation of zeocin resistance gene by means of PCR starting from DNA of the plasmid pSCV Zeo (Invitrogen Comp.): To select packaging cells after a stable transfection with the pTC53-zeo-scFv plasmid for a stable expression of the resistance gene, a zeocin cassette was integrated. For this purpose, a zeocin cassette was amplified by means of recombinant PCR from the vector pZeoSV2 (+/-) of Invitrogen Company (NV Leek, The Netherlands) and provided with the restriction sites NdeI 5' and 3' so that the cassette subsequently could be inserted into the NdeI restricted portion of the pUC19 backbone of pTC53. The PCR-batch (100 μ l) contained: 1 x PCR buffer (Taq: 10 mM Tris/HCl, pH 8.8, 50 mM KCl, 1.5 mM MgCl_2 , 0.01% gelatin, 10 μ M (+)- and 10 μ M (-)-primer, 200 μ M of each deoxynucleotide, 2.5 units of Taq polymerase and 100 ng of plasmid DNA. Following oligonucleotides have been used:

ZEO2184+NDE:

5'-GGAAATTCCATATGGAATTCCTACATAACTTACGGTAAATGGC-3'

Nde I

ZEO3258-NDE:

5'-GGAATTCCATATGGAATTCCTCAGTCCTGCTCCTCGGCC-3'

Nde I

The PCR-conditions were: 94°C/3 min, [94°C/1 min, 60°C/1 min, 72°C/1,5 min.] 30 cycles, 72°C/10 min and 4°C final temperature.

Insertion of a zeocin resistance gene into the scFv pTC53 Env expression constructs positive in the transient test.

Transfection by means of Lipofectamin™ (GIBCO/BRL, Life Technologies, Eggenstein).

Lipofectamin™: N-[2-({2,5bis[-(3-aminopropyl)amino]-1-oxypentyl}amino)ethyl]-N,N-dimethyl-2,3-bis (9-octadecenyloxy)-1-propanaminium trifluoroacetate)/Dioleoyl-phosphatidylethanolamine; 3 : 1 (w/w)

One day before transfection 1×10^6 cells were seeded in a 60 mm cell culture dish (Greiner, Nuertingen). For transfection 1 - 5 µg (depending on the experimental assay) of recombinant plasmid DNA were resuspended in 200 µl serum free medium. Simultaneously 8-25 µl (depending on the experimental assay) Lipofectamin™ were diluted in 200 µl serumfree medium. After combining both solutions a 45 min incubation at room temperature followed. The DNA-liposome mixture was filled up to a final volume of 2 ml and supplied to the cells washed with serumfree medium. Thereafter, the cells were incubated for 5 hours at 37°C. Subsequently, 2 ml of fresh medium containing the double concentration of FCS were added. The next day medium was changed.

For establishing of stable packaging cell clones the cells were overlaid with a selection medium 24 hours after transfection. The Zeocin™ resistance gene (*Streptoalloteichus hinductanus* bleomycin gene) was used as a selection marker. The selection was carried out in DMEM Medium supplemented with 525 µg/ml Zeocin™ (phleomycin from *Streptomyces verticillius*; Invitrogen BV, The Netherlands). The cells were added with fresh selection medium twice a week. After about 4 weeks cell foci representing cell clones could be identified. These colonies were removed individually and transferred into a 24 well plate (flat bottom, Nunc, Wiesbaden) in cell culture medium without antibiotic

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supplementation. The medium was changed twice a week. When the cells reached a confluence of about 90%, they were expanded into larger cell culture vials.

SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANT: BUNDESREPUBLIK DEUTSCHLAND, LETZTVERTRETEN DURCH DEN
PRÄSIDENTEN DES PAUL-ENRLICH-INSTITUTS
- (ii) TITLE OF INVENTION: CELL-SPECIFIC RETROVIRAL VECTORS WITH ANTIBODY
DOMAINS AND METHOD FOR THE PRODUCTION THEROF
- (iii) NUMBER OF SEQUENCES: 31
- (iv) CORRESPONDENCE ADDRESS:
- (A) ADDRESSEE: SMART & BIGGAR
 - (B) STREET: P.O. BOX 2999, STATION D
 - (C) CITY: OTTAWA
 - (D) STATE: ONT
 - (E) COUNTRY: CANADA
 - (F) ZIP: K1P 5Y6
- (v) COMPUTER READABLE FORM:
- (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: ASCII (text)
- (vi) CURRENT APPLICATION DATA:
- (A) APPLICATION NUMBER: CA 2,311,602
 - (B) FILING DATE: 27-NOV-1998
 - (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
- (A) APPLICATION NUMBER: DE 197 25 854.6
 - (B) FILING DATE: 28-NOV-1997
- (viii) ATTORNEY/AGENT INFORMATION:
- (A) NAME: SMART & BIGGAR
 - (B) REGISTRATION NUMBER:
 - (C) REFERENCE/DOCKET NUMBER: 76260-10
- (ix) TELECOMMUNICATION INFORMATION:
- (A) TELEPHONE: (613)-232-2486
 - (B) TELEFAX: (613)-232-8440

(2) INFORMATION FOR SEQ ID NO: 1:

- (i) SEQUENCE CHARACTERISTICS:
- (A) LENGTH: 4776 Base pairs
 - (B) TYPE: Nucleotide
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: other nucleic acid
- (iii) HYPOTHETICAL: NO
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
- | | | | | | | |
|------------|------------|------------|------------|------------|------------|-----|
| GAATTCCCGT | ACGAGCCATA | GATAAAATAA | AAGATTTTAT | TTAGTCTCCA | GAAAAAGGGG | 60 |
| GGAATGAAAG | ACCCACCTG | TAGGTTTGGC | AAGCTAGCTT | AAGTAACGCC | ATTTTGCAAG | 120 |
| GCATGGAAAA | ATACATAACT | GAGAATAGAG | AAGTTCAGAT | CAAGGTCAGG | AACAGATGGA | 180 |
| ACAGCTGAAT | ATGGGCCAAA | CAGGATATCT | GTGGTAAGCA | GTTCTGCCC | CGGCTCAGGG | 240 |
| CCAAGAACAG | ATGGAACAGC | TGAATATGGG | CCAAACAGGA | TATCTGTGGT | AAGCAGTTCC | 300 |
| TGCCCCGGCT | CAGGGCCAAG | AACAGATGGT | CCCCAGATGC | GGTCCAGCCC | TCAGCAGTTT | 360 |
| CTAGAGAACC | ATCAGATGTT | TCCAGGGTGC | CCCAAGGACC | TGAAATGACC | CTGTGCCTTA | 420 |
| TTTGAATAA | CCAATCAGTT | CGCTTCTCGC | TTCTGTTCGC | GCGCTTCTGC | TCCCCGAGCT | 480 |
| CAATAAAAGA | GCCCACAACC | CCTCACTCGG | GGCGCCAGTC | CTCCGATTGA | CTGAGTCGCC | 540 |
| CGGGTGGGGG | AGCTCGCTGT | TGGGCTCGCG | GTTGAGGACA | AACTCTTCGC | GGTCTTTCCA | 600 |
| GTACTCTTGG | ATCGGAAACC | CGTCGGCCTC | CGAACGGTAC | TCCGCCACCG | AGGGACCTGA | 660 |
| GCGAGTCCGC | ATCGACCGGA | TCGGAAAACC | TCTCGAGAAA | GGCGTCTAAC | CAGTCACAGT | 720 |
| CGCAAGGTAG | GCTGAGCACC | GTGGCCGGGC | GGCACGGGTG | GCGGTCGGGG | TTGTTTCTGG | 780 |
| CGGAGGTGCT | GCTGATGATG | TAATTAAGTA | GGCGGTCTTG | AGACGGCGAT | GGTCGAGGTG | 840 |

	AGGTGTGGCA	GGCTTGAGAT	CTGGCCATAC	ACTTGAGTGA	CAATGACATC	CACTTTGCCT	900
	TTCTCTCCAC	AGGTGTCCAC	TCCCAGGTCC	AACCGGATCC	GAGCTCCACC	GCGGTAAAGG	960
	TCGCTGGGAA	GACCCCGTGG	ATCCACCACT	CTCGACTCAA	GAAAGCTCCT	GACAACCAAG	1020
	AAGAATGGAC	TGTCTCACCA	ACCTCCGATC	CGCTGAGGGT	AAAGTTGACC	AGGCGAGCAA	1080
	AATCCTAATT	CTCCTTGTGG	CTTGGTGGGG	GTTTGGGACC	ACTGCCGAAG	TTTCGACTGC	1140
	CGGCTCCGGG	GGCGGTGGTT	CTGGTGGTGG	TTCTGGTGGT	GGTGGTTCTG	GTGGTGGTGG	1200
	TTCTGGCGCC	AGCCCAGTCC	AGTTTATCCC	CCTGCTTGTG	GGTCTAGGGA	TTTCAGGGGC	1260
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	TGACTCCCTG	GCTGAGGTTG	TCTTACAAAA	TAGAAGAGGG	TTAGACCTAT	TGACTGCCGA	1440
	ACAAGGAGGA	ATATGTCTCG	CACTCCAGGA	GAAGTGTTGT	TTTTACGCTA	ACAAGTCGGG	1500
	TATCGTACGT	GACAAGATCC	GAAAACCTCA	AGAGGACCTT	ATCGAGAGAA	AACGTGCACT	1560
	GTACGACAAC	CCCCTGTGGA	GCGGCTTGAA	CGGCTTCCTT	CCATATTTGC	TACCCTTGTT	1620
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	AAATTTGTGA	TGCTATTGCT	TTATTTGTAA	CCATTATAAG	CTGCAATAAA	CAAGTTAACA	2040
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CTCGCGCGTT TCGGTGATGA CGGTGAAAAC CTCTGACACA TGCAGCTCCC GGAGACGGTC 4440
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 CATTCGCCAT TCAGGCTGCG CAACTGTTGG GAAGGGCGAT CGGTGCGGGC CTCTTCGCTA 4680
 TTACGCCAGC TGGCGAAAGG GGGATGTGCT GCAAGGCGAT TAAGTTGGGT AACGCCAGGG 4740
 TTTTCCCAGT CACGACGTTG TAAAACGACG GCCAGT 4776

10

(2) INFORMATION FOR SEQ ID NO: 2:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 12 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

Met Lys Asp Pro Thr Cys Arg Phe Gly Lys Leu Ala
 1 5 10

20

(2) INFORMATION FOR SEQ ID NO: 3:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 21 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:

Met Glu Lys Tyr Ile Thr Glu Asn Arg Glu Val Gln Ile Lys Val Arg
 1 5 10 15

30

Asn Arg Trp Asn Ser
 20

(2) INFORMATION FOR SEQ ID NO: 4:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 8 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:

Met Gly Gln Thr Gly Tyr Leu Trp
 1 5

40

(2) INFORMATION FOR SEQ ID NO: 5:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 13 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:

Met Glu Gln Leu Asn Met Gly Gln Thr Gly Tyr Leu Trp
 1 5 10

50

(2) INFORMATION FOR SEQ ID NO: 6:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 12 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:

Met Val Pro Arg Cys Gly Pro Ala Leu Ser Ser Phe
 1 5 10

60

(2) INFORMATION FOR SEQ ID NO: 7:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 10 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:

10 Met Phe Pro Gly Cys Pro Lys Asp Leu Lys
 1 5 10

(2) INFORMATION FOR SEQ ID NO: 8:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 15 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:

20 Met Val Glu Val Arg Cys Gly Arg Leu Glu Ile Trp Pro Tyr Thr
 1 5 10 15

(2) INFORMATION FOR SEQ ID NO: 9:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 24 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:

30 Met Thr Ser Thr Leu Pro Phe Ser Pro Gln Val Ser Thr Pro Arg Ser
 1 5 10 15

Asn Arg Ile Arg Ala Pro Pro Arg
 20

(2) INFORMATION FOR SEQ ID NO: 10:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 232 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:

40 Met Asp Cys Leu Thr Asn Leu Arg Ser Ala Glu Gly Lys Val Asp Gln
 1 5 10 15

Ala Ser Lys Ile Leu Ile Leu Leu Val Ala Trp Trp Gly Phe Gly Thr
 20 25 30

50 Thr Ala Glu Val Ser Thr Ala Gly Ser Gly Gly Gly Gly Ser Gly Gly
 35 40 45

Gly Ser Gly Gly Gly Gly Ser Gly Gly Gly Gly Ser Gly Ala Ser Pro
 50 55 60

Val Gln Phe Ile Pro Leu Leu Val Gly Leu Gly Ile Ser Gly Ala Thr
 65 70 75 80

Leu Ala Gly Gly Thr Gly Leu Gly Val Ser Val His Thr Tyr His Lys
 85 90 95

60 Leu Ser Asn Gln Leu Ile Glu Asp Val Gln Ala Leu Ser Gly Thr Ile
 100 105 110

Asn Asp Leu Gln Asp Gln Ile Asp Ser Leu Ala Glu Val Val Leu Gln
115 120 125

Asn Arg Arg Gly Leu Asp Leu Leu Thr Ala Glu Gln Gly Gly Ile Cys
130 135 140

10 Leu Ala Leu Gln Glu Lys Cys Cys Phe Tyr Ala Asn Lys Ser Gly Ile
145 150 155 160

Val Arg Asp Lys Ile Arg Lys Leu Gln Glu Asp Leu Ile Glu Arg Lys
165 170 175

Arg Ala Leu Tyr Asp Asn Pro Leu Trp Ser Gly Leu Asn Gly Phe Leu
180 185 190

20 Pro Tyr Leu Leu Pro Leu Leu Gly Pro Leu Phe Gly Leu Ile Leu Phe
195 200 205

Leu Thr Leu Gly Pro Cys Ile Met Lys Thr Leu Thr Arg Ile Ile His
210 215 220

Asp Lys Ile Gln Ala Val Lys Ser
225 230

(2) INFORMATION FOR SEQ ID NO: 11:

(i) SEQUENCE CHARACTERISTICS:

30 (A) LENGTH: 14 Amino acids
(B) TYPE: Amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:

Met Asp Thr Leu Gly Val Arg Trp Ser Lys Asn Ser Arg Val
1 5 10

(2) INFORMATION FOR SEQ ID NO: 12:

(i) SEQUENCE CHARACTERISTICS:

40 (A) LENGTH: 15 Amino acids
(B) TYPE: Amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:

Met Tyr Lys Ser Ser Gly Glu Cys Gly Arg Gly Leu Arg Arg Pro
1 5 10 15

(2) INFORMATION FOR SEQ ID NO: 13:

(i) SEQUENCE CHARACTERISTICS:

50 (A) LENGTH: 16 Amino acids
(B) TYPE: Amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 13:

Met Ile Arg Tyr Ile Asp Glu Phe Gly Gln Thr Thr Thr Arg Met Gln
1 5 10 15

(2) INFORMATION FOR SEQ ID NO:14:

(i) SEQUENCE CHARACTERISTICS:

60 (A) LENGTH: 4 Amino acids
(B) TYPE: Amino acid
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 14:

Met Leu Tyr Leu

1

(2) INFORMATION FOR SEQ ID NO: 15:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 6 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 15:

Met Leu Leu Leu Tyr Leu

1

5

(2) INFORMATION FOR SEQ ID NO: 16:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 12 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 16:

Met Phe Gln Val Gln Gly Glu Val Trp Glu Val Phe

1

5

10

(2) INFORMATION FOR SEQ ID NO: 17:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 26 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 17:

Met Val Ile Ala Val Ser Cys Val Lys Leu Leu Ser Ala His Asn Ser

1

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15

Thr Gln His Thr Ser Arg Lys His Lys Val

20

25

(2) INFORMATION FOR SEQ ID NO: 18:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 49 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 18:

Met Ser Glu Leu Thr His Ile Asn Cys Val Ala Leu Thr Ala Arg Phe

1

5

10

15

Pro Val Gly Lys Pro Val Val Pro Ala Ala Leu Met Asn Arg Pro Thr

20

25

30

Arg Gly Glu Arg Arg Phe Ala Tyr Trp Ala Leu Phe Arg Phe Leu Ala

35

40

45

His

(2) INFORMATION FOR SEQ ID NO: 19:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 4 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 19:

Met Leu Thr Leu

1

(2) INFORMATION FOR SEQ ID NO: 20:

(i) SEQUENCE CHARACTERISTICS:

10 (A) LENGTH: 9 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 20:

Met Arg Leu Ser Lys Arg Ile Phe Thr

1

5

(2) INFORMATION FOR SEQ ID NO: 21:

(i) SEQUENCE CHARACTERISTICS:

20 (A) LENGTH: 11 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 21:

Met Ser Lys Leu Gly Leu Thr Val Thr Asn Ala

1

5

10

(2) INFORMATION FOR SEQ ID NO:22:

(i) SEQUENCE CHARACTERISTICS:

30 (A) LENGTH: 70 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 22:

Met Arg Cys Glu Ile Pro His Arg Cys Val Arg Arg Lys Tyr Arg Ile

1

5

10

15

Arg Arg His Ser Pro Phe Arg Leu Arg Asn Cys Trp Glu Gly Arg Ser

20

25

30

40

Val Arg Ala Ser Ser Leu Leu Arg Gln Leu Ala Lys Gly Gly Cys Ala

35

40

45

Ala Arg Arg Leu Ser Trp Val Thr Pro Gly Phe Ser Gln Ser Arg Arg

50

55

60

Cys Lys Thr Thr Ala Ser

65

70

50 (2) INFORMATION FOR SEQ ID NO: 23:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 88 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 23:

Met Ile Pro Arg Asp Pro Arg Ser Pro Ala Pro Asp Leu Ser Ala Ile

1

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60 Asn Gln Pro Ala Gly Arg Ala Glu Arg Arg Ser Gly Pro Ala Thr Leu

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22

Ser Ala Ser Ile Gln Ser Ile Asn Cys Cys Arg Glu Ala Arg Val Ser
 35 40 45

Ser Ser Pro Val Asn Ser Leu Arg Asn Val Val Ala Ile Ala Thr Gly
 50 55 60

10 Ile Val Val Ser Arg Ser Ser Phe Gly Met Ala Ser Phe Ser Ser Gly
 65 70 75 80

Ser Gln Arg Ser Arg Arg Val Thr
 85

(2) INFORMATION FOR SEQ ID NO: 24:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 56 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

20 (ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 24:

Met Leu Cys Lys Lys Ala Val Ser Ser Phe Gly Pro Pro Ile Val Val
 1 5 10 15

Arg Ser Lys Leu Ala Ala Val Leu Ser Leu Met Val Met Ala Ala Leu
 20 25 30

His Asn Ser Leu Thr Val Met Pro Ser Val Arg Cys Phe Ser Val Thr
 35 40 45

30 Gly Glu Tyr Ser Thr Lys Ser Phe
 50 55

(2) INFORMATION FOR SEQ ID NO: 25:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 49 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

40 (ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 25:

Met Arg Arg Pro Ser Cys Ser Cys Pro Ala Ser Ile Arg Asp Asn Thr
 1 5 10 15

Ala Pro His Ser Arg Thr Leu Lys Val Leu Ile Ile Gly Lys Arg Ser
 20 25 30

Ser Gly Arg Lys Leu Ser Arg Ile Leu Pro Leu Leu Arg Ser Ser Ser
 35 40 45

50 Met

(2) INFORMATION FOR SEQ ID NO: 26:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 6 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 26:

Met Pro Gln Lys Arg Glu
 1 5

60

(2) INFORMATION FOR SEQ ID NO: 27:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 27 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 27:

10 Met Leu Asn Thr His Thr Leu Pro Phe Ser Ile Leu Leu Lys His Leu
 1 5 10 15

Ser Gly Leu Leu Ser His Glu Arg Ile His Ile
 20 25

(2) INFORMATION FOR SEQ ID NO: 28:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 5 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 28:

20 Met Tyr Leu Glu Lys
 1 5

(2) INFORMATION FOR SEQ ID NO: 29:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 26 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 29:

30 Met Thr Leu Thr Tyr Lys Asn Arg Arg Ile Thr Arg Pro Phe Arg Leu
 1 5 10 15

Ala Arg Phe Gly Asp Asp Gly Glu Asn Leu
 20 25

(2) INFORMATION FOR SEQ ID NO: 30:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 11 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 30:

40 Met Gln Leu Pro Glu Thr Val Thr Ala Cys Leu
 1 5 10

(2) INFORMATION FOR SEQ ID NO: 31:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 31 Amino acids

(B) TYPE: Amino acid

(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: Protein

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 31:

50 Met Pro Gly Ala Asp Lys Pro Val Arg Ala Arg Gln Arg Val Leu Ala
 1 5 10 15

Gly Val Gly Ala Gly Leu Thr Met Arg His Gln Ser Arg Leu Tyr
 20 25 30

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CLAIMS:

1. A method for the production of cell-specific retroviral vectors comprising the following steps:

5 a) immunizing a mammal with one or more cell population(s),

b) isolating RNA from the immunized mammal, comprising B cell RNA,

10 c) producing cDNA regions of the variable regions of an immunoglobulin heavy and light chain from the isolated RNA by means of RT-PCR with primers for the immunoglobulin heavy and light chain wherein the primers comprise a nucleic acid sequence which encodes an oligopeptide linker,

d) ligating the cDNA regions to form scFv-cDNAs,

15 e) ligating the scFv-cDNAs into a phagemid-vector and transformation of a host bacterium with the phagemid vector,

f) isolating phages binding to the cell population(s) used in step a) by means of selection,

20 g) isolating cell-specific phages from the phages obtained in step f) which only bind to the cell population(s) used in step a) by means of selection,

25 h) excising the scFv-encoding DNA fragments from the cell-specific phages obtained in step g) and ligating the scFv-encoding DNA fragments into a psi-negative retroviral Env expression vector to form an Env-scFv expression vector comprising a chimeric Env-scFv surface protein encoding DNA sequence,

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i) transforming said Env-scFv expression vector into a packaging cell, wherein the packaging cell comprises a psi-negative env, gag, pol expression construct; and

j) isolating the retroviral vectors secreted by
5 the packaging cell.

2. The method according to claim 1 wherein said isolating step g) comprises isolating individual cell-specific phages.

3. The method according to claim 1 or 2 wherein the
10 steps f) and/or g) are repeated at least once.

4. The method according to any one of claims 1 to 3 further comprising the step of:

k) isolating, by means of selection, from the retroviral vectors isolated in step j), those retroviral
15 vectors with the capability to transduce the cells of the cell population(s).

5. The method according to any one of claims 1 to 4 wherein the mammal is selected from the group consisting of mouse, rat, guinea pig, rabbit, goat and sheep.

20 6. The method according to any one of claims 1 to 4 wherein the cell population(s) is/are selected from the group consisting of human cells, mouse cells, rat cells, sheep cells, cattle cells and pig cells.

7. The method according to claim 6 wherein the cell
25 population(s) is/are selected from the group consisting of T cells, epithelial cells, muscle cells, hematopoietic cells, stem cells, neural cells, carcinoma cells and liver cells.

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8. The method according to any one of claims 1 to 7 wherein the env gene of the psi-negative retroviral Env expression vector is derived from spleen necrosis virus (SNV).

5 9. The method according to claim 8 wherein the expression vector is the vector having the designation pTC53.

SMART & BIGGAR

OTTAWA, CANADA

PATENT AGENTS

Application number / numéro de demande: 2311602

Figures: 1, 2 + 3

Pages: _____

Unscannable items
received with this application
(Request original documents in File Prep. Section on the 10th floor)

Documents reçu avec cette demande ne pouvant être balayés
(Commander les documents originaux dans la section de préparation des dossiers au
10^{ème} étage)

DNA sequence 4776 b.p. GATTCCGGTAC ... ACCACGGCCAGT linear

[illegible]

Fig. 4A

Fig. 4B