Abstract: The present invention relates to novel compositions of active agents and methods for the treatment of HIV infection and AIDS. In particular, the present invention relates to novel methods for treatment of HIV infection and prevention of AIDS.
FIELD OF THE INVENTION

The present invention relates to novel methods in the treatment of HIV infections and AIDS. In particular, the present invention relates to specific methods with treatment of HIV-specific vaccine peptides administered in a dosis regimen, optionally together with a reservoir purging agent and/or immunomodulatory compounds, wherein progress or effect of the immunization phase is monitored or followed by measurement of HIV DNA levels.

BACKGROUND OF THE INVENTION

HIV-1 infection is today perceived as an incurable chronic viral infection in which lifelong combination antiretroviral therapy (cART) is needed to avoid disease. Very early during acute HIV infection a latent reservoir is established and despite effective cART, HIV-1 persists in latently infected cells. Upon treatment interruption, the virus quickly replicates, and viremia rebounds to pre-treatment levels. In the inactive, resting state latently infected cells are unrecognizable to the immune system and unresponsive to antiretroviral drugs. The size of the reservoir likely varies between individuals and may be influenced by a number of different factors such as host immune constitution, time from diagnosis to initiation, level of persistent immune activation, antiretroviral treatment regimens used and individual response to treatment. Earlier studies employing viral outgrowth assays indicated that the number of latent CD4 T cells harboring replication-competent virus was approximately 1 per 10^6 cells.

A broad range of bioanalytical assays have been used in the attempt to quantify the reservoir but it is currently unclear which assay(s) should be used to monitor HIV-1 reservoirs in clinical studies of eradication strategies. Upon activation, resting T cells carrying replication competent integrated proviral DNA are capable of resuming HIV transcription. One of the proposed ways of curing HIV-1 is to activate and kill latently infected cells in the presence of antiretroviral therapy. Epigenetic modulation of the molecular mechanisms that block transcription of integrated HIV DNA can reactivate HIV-1 expression in resting infected memory CD4 T cells and disrupt latency. Histone deacetylase inhibitors (HDACI) turn on genes by promoting acetylation of lysine residues on histones. This induces chromatin relaxation and transcriptional activation. The HDACi romidepsin (Celgene) potently activates HIV-1 expression in latently infected cell lines and primary T cells.

Vacc-4x is a peptide-based HIV-1 therapeutic vaccine that aims to improve immune responses to p24Gag since this has been associated with slower disease progression and improved virus control. The primary objective of Vacc-4x immunization is to strengthen the immune system’s response to HIV p24. The enhanced immune response to HIV-1 following
immunization with Vacc-4x could improve the host immune system as part of an HIV functional cure treatment strategy.

In one of the largest randomized, placebo controlled HIV therapeutic vaccine trials conducted to date (study CT-BI/Vacc-4x/2007/1), Vacc-4x and rhuGM-CSF (Leukine®) as adjuvant showed a significant reduction in viral load (VL) set point in the Vacc-4x group as compared to placebo and a significant reduction in VL set point from historic preART values, despite higher preART values being present in the Vacc-4x group as compared to placebo. Additionally Vacc-4x was shown to be immunogenic, inducing proliferative responses in both CD4 and CD8 T-cells.

New HIV p24 peptides are described in WO91/13360, wherein the peptides are used in a method of discriminating between a false and true diagnosed HIV-positive serum sample. Johnson R.P., et al., The Journal of Immunology, Vol.147, p.1512-1521, No.5, September 1, 1991 describe an analysis of the fine specificity of gag-specific CTL-responses in three HIV-1 seropositive individuals, the gag-specific CTL-responses were found to be mediated by CD3+CD8+ lymphocytes which are HLA class I restricted.

EP-A-0 356 007 discloses antigenic determinants, in particular it relates to synthetic polypeptide sequences which are related to proteins present in the HIV-1 and which can be used as a basis for a potential vaccine against AIDS.

Rosenberg E.S. et al., Science, Vol.278, 21 November 1997, p.1447-1450 describe that virus specific CD4+ T helper lymphocytes are critical to the maintenance of effective immunity in a number of chronic viral infections, but are characteristically undetectable in chronic human immunodeficiency virus-type 1 (HIV-1) infection. HIV-1-specific proliferative responses to p24 were inversely related to viral load. They conclude that the HIV-1-specific helper cells are likely to be important in immunotherapeutic interventions and vaccine development.

EP 0 230 222, EP 0 270 114, DE 37 11 016 and GB 2 188 639 all in the name of F. Hoffmann-La Roche & Co. Aktiengesellschaft concern recombinant expression and purification of an HTLVIII Gag/Env gene protein or fusion proteins. The proteins consisting of native sequences can be purified to homogeneity and used as a basis for diagnostic tests for detection of antibodies against viruses associated with AIDS. The gag/env protein may also be formulated for use as a vaccine for protection against AIDS through prophylactic immunization.

International Patent Application WO00/52040 discloses methods for treating HIV infections by administering e.g. HIV specific peptides based on conserved regions of HIV gag p24.

There is a need to provide improved methods in the treatment of HIV infections and AIDS.
OBJECT OF THE INVENTION

It is an object of embodiments of the invention to provide effective methods, which can be used in the treatment of HIV infection and prevention of AIDS.

The present invention is based on the finding that HIV-specific vaccine peptides may be used in specific dosage regimens, wherein HIV-1 viral DNA is monitored subsequently or simultaneously as a measure of effect of the vaccine, optionally together with specific reservoir purging agents. This may provide an effective method in the treatment and/or eradication of HIV infection and AIDS.

SUMMARY OF THE INVENTION

It has been found that the effect of a treatment with HIV-specific vaccine peptides administered in a specific dosage regimen may be monitored by measuring HIV-1 viral DNA.

So, in a first aspect of the present invention is provided a method for reducing and/or delaying pathological effects of human immunodeficiency virus I (HIV) or for reducing the risk of developing acquired immunodeficiency syndrome (AIDS) in a human subject infected with HIV, the method comprising the steps of:

a) a therapeutic HIV-1 immunization phase consisting of the administering in one or more doses of an effective amount of one or more HIV-specific peptide selected from the list consisting of the amino acid sequence shown in SEQ ID NO: 18 (Vacc-10), SEQ ID NO: 11 (Vacc-11), SEQ ID NO: 6 (Vacc-12), and SEQ ID NO: 3 (Vacc-13) over a period of 1-12 weeks;

b) one or more subsequent or simultaneous measurements of HIV-1 DNA levels in said human subject infected with HIV; and optionally

c) a subsequent viral reactivation phase consisting of the administering of an effective amount of a reservoir purging agent.

In some embodiments the methods described in the present invention do not comprise the administering of a reservoir purging agent, such as a histone deacetylase (HDAC) inhibitor, and/or an immunomodulatory compound.

In some embodiments the methods described in the present invention do not comprise the administering of an immunomodulatory compound.

In some embodiments the methods described in the present invention do not comprise the administering of a reservoir purging agent, such as a histone deacetylase (HDAC) inhibitor.

In a second aspect of the present invention is provided a method for monitoring the effect of a therapeutic HIV-1 immunization phase consisting of the administering in one or more doses of an effective amount of one or more HIV-specific peptide selected from the list consisting of the amino acid sequence shown in SEQ ID NO: 18 (Vacc-10), SEQ ID NO: 11 (Vacc-11), SEQ ID NO: 6 (Vacc-12), and SEQ ID NO: 3 (Vacc-13) over a period of 1-12 weeks; in reducing and/or delaying pathological effects of human immunodeficiency virus I
(HIV) or in reducing the risk of developing acquired immunodeficiency syndrome (AIDS) in a human subject infected with HIV, the method comprising the step of

a) One or more measurements of HIV-1 DNA levels in said human subject infected with HIV-1 subsequent or simultaneous to said immunization phase.

In a third aspect of the present invention there is provided an effective amount of one or more HIV-specific peptides selected from the list consisting of the amino acid sequence shown in SEQ ID NO: 18 (Vacc-10), SEQ ID NO: 11 (Vacc-11), SEQ ID NO: 6 (Vacc-12) for use in a method for reducing and/or delaying pathological effects of human immunodeficiency virus I (HIV) or for reducing the risk of developing acquired immunodeficiency syndrome (AIDS) in a human infected with HIV, the method comprising the steps of:

a) a therapeutic HIV-1 immunization phase consisting of the administering in one or more doses of an effective amount of one or more HIV-specific peptide selected from the list consisting of the amino acid sequence shown in SEQ ID NO: 18 (Vacc-10), SEQ ID NO: 11 (Vacc-11), SEQ ID NO: 6 (Vacc-12), and SEQ ID NO: 3 (Vacc-13) over a period of 1-12 weeks;

b) a subsequent or simultaneous measurement of HIV-1 DNA levels in said human subject infected with HIV; and optionally

c) a subsequent viral reactivation phase consisting of the administering of an effective amount of a reservoir purging agent.

In some aspects of the present invention the therapeutic HIV-1 immunization phase consist of the administering in one or more doses of an effective amount of one or more HIV-specific peptide selected from the group of amino acid sequences:

Xaa₁ Xaa₂ Xaa₃ Xaa₄ Xaa₅ Xaa₆ Ala Xaa₇ Xaa₈ Gln Thr Pro Trp Xaa₁₄ Xaa₁₅ Xaa₁₆ Xaa₁₇ Xaa₁₈ Val Xaa₂₀ (SEQ ID NO: 1);

wherein Xaa in position 1 is Lys or Arg,
Xaa in position 2 is Ala, Gly, Ser or Arg,
Xaa in position 3 is Leu or Met,
Xaa in position 4 is Gly or Arg,
Xaa in position 5 is Pro, Thr, Val, Ser, Gln or Ala,
Xaa in position 6 is Gly, Ala, Lys, Arg, Gln or Glu,
Xaa in position 8 is Thr or Ser,
Xaa in position 9 is Leu or Ile,
Xaa in position 14 is Thr, Ser or Val,
Xaa in position 15 is Ala or Ser,
Xaa in position 16 is Cys or Ser,
Xaa in position 17 is Gln or Leu,
Xaa in position 18 is Gly, Glu or Arg, and
Xaa in position 20 is Gly or Arg;

Xaa₂ Xaa₃ Xaa₄ Xaa₅ Gly Leu Asn Pro Leu Val [Gly]ₙ Xaa₁₂ Xaa₁₃ Tyr Xaa₁₅ Pro Xaa₁₇ Xaa₁₈
5  Ile Leu Xaa₂₁ Xaa₂₂ (SEQ ID NO: 4);
wherein Xaa in position 1 is Arg, Lys, Asp or none,
Xaa in position 2 is Trp, Gly, Lys or Arg,
Xaa in position 3 is Ile, Leu, Val or Met,
Xaa in position 4 is Ile, Val or Leu,

10  Xaa in position 5 Leu, Met, Val or Pro,
Xaa in position 12 is Arg or Lys,
Xaa in position 13 is Met or Leu,
Xaa in position 15 is Ser, Cys or Gln,
Xaa in position 17 is Thr, Val, Ile, Ser or Ala,

15  Xaa in position 18 is Ser, Gly or Thr,
Xaa in position 21 is Asp, Glu, Cys or Gly,
Xaa in position 22 is Gly or none, and

n = 0, 1, 2 or 3;

20  Xaa₂ Xaa₃ Xaa₄ Pro Ile Pro Xaa₇ Xaa₉ Xaa₁₀ Xaa₁₁ Xaa₁₂ [Gly]ₙ Xaa₁₃ Xaa₁₄ Xaa₁₅ Xaa₁₆
Xaa₁₇ Xaa₁₈ Xaa₁₉ Xaa₂₀ Xaa₂₁ Xaa₂₂ Xaa₂₃ Xaa₂₄ (SEQ ID NO: 9);
wherein Xaa in position 1 is Asn, Ser, Gly, His, Ala, Pro, Arg or none,
Xaa in position 2 is Asn, Ala or Lys,
Xaa in position 3 is Pro, Gln, Gly, Ile or Leu,

25  Xaa in position 7 is Val or Ala,
Xaa in position 8 is Gly or Lys,
Xaa in position 9 is Glu, Asp, Lys, Phe or Thr,
Xaa in position 10 is Ile, Met, Val or Leu,
Xaa in position 11 is Tyr, Leu or none,

30  Xaa in position 12 is Ser or none,
Xaa in position 13 is Arg or none,
Xaa in position 14 is Asp, Arg, Trp, Ala or none,
Xaa in position 15 is Ile or none,
Xaa in position 16 is Tyr or none,

35  Xaa in position 17 is Lys or Arg,
Xaa in position 18 is Arg, Lys or Asp,
Xaa in position 19 is Trp or Gly,
Xaa in position 20 is Ile, Met, Val, Gln or Ala,
Xaa in position 21 is Ile, Val or Ala,
Xaa in position 22 is Leu, Met or Val,
Xaa in position 23 is Gly or Cys,
Xaa in position 24 is Leu or none,
n = 1, 2 or 3; and

\[Xaa_1 \ Xaa_2 \ Ile \ Ile \ Xaa_5 \ Xaa_6 \ Xaa_7 \ Xaa_8 \ Xaa_9 \ Xaa_{11} \ [Gly]_n \ [Arg]_m \ Xaa_{12} \ Xaa_{13} \ Xaa_{14} \ Xaa_{15} \ Xaa_{16} \ Xaa_{17} \ Xaa_{18} \ Xaa_{19} \ Xaa_{20} \ Xaa_{21} \ Xaa_{22} \ Xaa_{23} \ Xaa_{24} \ Xaa_{25}\] (SEQ ID NO: 15);
wherein Xaa in position 1 is Pro, Lys, Arg or none,
Xaa in position 2 is Glu, Arg, Phe or Lys,
Xaa in position 5 is Pro or Thr,
Xaa in position 6 is Met, Thr or Nleu,
Xaa in position 7 is Phe or Leu,
Xaa in position 8 is Ser, Thr, Ala or Met,
Xaa in position 9 is Ala, Glu or Leu,
Xaa in position 11 is Ser or none,
Xaa in position 12 is Ala, Arg or none,
Xaa in position 13 is Ile, Leu or none,
Xaa in position 14 is Ser, Ala, Leu or none,
Xaa in position 15 is Tyr, Glu or Asp,
Xaa in position 16 is Gly or Asp,
Xaa in position 17 is Ala or Leu,
Xaa in position 18 is Thr, Ile, Val, Leu or Asn,
Xaa in position 19 is Pro, Thr or Ser,
Xaa in position 20 is Tyr, Phe, Nleu, His or Gln,
Xaa in position 21 is Asp, Asn, Leu or Ala,
Xaa in position 22 is Leu, Ile, Val or Asn,
Xaa in position 23 is Asn, Tyr, Cys or Gly,
Xaa in position 24 is Thr, Met, Ile, Ala, Val or none,
Xaa in position 25 is Gly or none,
n = 1, 2 or 3 and m = 0, 1, 2 or 3 independent of each other;

wherein the terminal ends of each HIV specific peptide may be free carboxyl- or amino-groups, amides, acyls or acetyls; and wherein each peptide optionally is in the form of an acetate salt; over a period of 1-12 weeks; and optionally a subsequent viral reactivation phase consisting of the administering of an effective amount of a reservoir purging agent.

In some embodiments the one or more HIV-specific peptide is selected from the group of amino acid sequences of SEQ ID NOs: 1, 4, 9 and 15; wherein the terminal ends of
each HIV specific peptide may be free carboxyl- or amino- groups, amides, acyls or acetyl; and wherein each peptide is in the form of an acetate salt.

In some embodiments the peptide consisting of the amino acid sequence shown in SEQ ID NO: 18 (Vacc-10) is in the form of an acetate salt.

In some embodiments the peptide consisting of the amino acid sequence shown in SEQ ID NO: 11 (Vacc-11) is in the form of an acetate salt.

In some embodiments the peptide consisting of the amino acid sequence shown in SEQ ID NO: 6 (Vacc-12) is in the form of an acetate salt.

In some embodiments the peptide consisting of the amino acid sequence shown in SEQ ID NO: 3 (Vacc-13) is in the form of an acetate salt.

In some embodiments one, two, three or four peptide acetate salts is/are used in the methods according to the invention.

LEGENDS TO THE FIGURE

15 Fig. 1 illustrates various embodiments of the method according to the present invention in a flow diagram.

DETAILED DISCLOSURE OF THE INVENTION

The present invention is based on the finding that significant reductions in the HIV-1 reservoir size due to increased levels and responsiveness of HIV-1-specific cytotoxic T lymphocytes in Vacc-4x immunized subjects can be observed, and that the progress of treatment may be monitored by measurements of HIV-1 DNA levels. This may be used to follow the success of the treatment, as a guidance in its development, and for the selection of patients which benefit from treatment with the vaccine.

25 Definitions

When terms such as “one”, “a” or “an” are used in this disclosure they mean “at least one”, or “one or more” unless otherwise indicated. Further, the term “comprising” is intended to mean “including” and thus allows for the presence of other constituents, features, conditions, or steps than those explicitly recited.

“HIV” unless otherwise indicated generally denotes human immunodeficiency virus I. "HIV disease" is composed of several stages including the acute HIV infection which often manifests itself as a flu-like infection and the early and medium stage symptomatic disease, which has several non-characteristic symptoms such as skin rashes, fatigue, night sweats, slight weight loss, mouth ulcers, and fungal skin and nail infections. Most HIV infected will experience mild symptoms such as these before developing more serious
illnesses. It is generally believed that it takes five to seven years for the first mild symptoms to appear. As HIV disease progresses, some individuals may become quite ill even if they have not yet been diagnosed with AIDS (see below), the late stage of HIV disease. Typical problems include chronic oral or vaginal thrush (a fungal rash or spots), recurrent herpes blisters on the mouth (cold sores) or genitals, ongoing fevers, persistent diarrhea, and significant weight loss. "AIDS" is the late stage HIV disease and is a condition which progressively reduces the effectiveness of the immune system and leaves individuals susceptible to opportunistic infections and tumors.

"Reducing and/or delaying pathological effect of HIV" is in the present context meant to denote that use of the methods of the invention provides for a statistically significant reduction and/or delay in morbidity seen in individual infected with HIV which are treated according to the present invention. That is, the time of onset of manifest disease symptoms characterizing AIDS is later compared to non-treated controls and/or the number of pathological manifestations is reduced to controls not receiving the treatment of the present invention.

The term "peptide" is in the present context intended to mean both short peptides of from 2 to 10 amino acid residues, oligopeptides of from 11 to 100 amino acid residues, and polypeptides of more than 100 amino acid residues. When referring to amino acids in peptides, it is intended that the amino acids are L-amino acids, unless other information is provided.

A "variant" or "analogue" of a peptide refers to a peptide having an amino acid sequence that is substantially identical to a reference peptide, typically a native or "parent" polypeptide. The peptide variant may possess one or more amino acid substitutions, deletions, and/or insertions at certain positions within the native amino acid sequence.

"Conservative" amino acid substitutions are those in which an amino acid residue is replaced with an amino acid residue having a side chain with similar physicochemical properties. Families of amino acid residues having similar side chains are known in the art, and include amino acids with basic side chains (e.g., lysine, arginine, histidine), acidic side chains (e.g., aspartic acid, glutamic acid), uncharged polar side chains (e.g., glycine, asparagine, glutamine, serine, threonine, tyrosine, cysteine, tryptophan), nonpolar side chains (e.g., alanine, valine, leucine, isoleucine, proline, phenylalanine, methionine), beta-branched side chains (e.g., threonine, valine, isoleucine) and aromatic side chains (e.g., tyrosine, phenylalanine, tryptophan, histidine). A particular form of conservative amino acid substitutions include those with amino acids, which are not among the normal 20 amino acids encoded by the genetic code. Since preferred embodiments of the present invention entail use of synthetic peptides, it is unproblematic to provide such "non-naturally occurring" amino acid residues in the peptides disclosed herein, and thereby it is possible to exchange the natural saturated carbon chains in the side chains of amino acid residues with shorter or longer saturated carbon chains – for instance, lysine may be substituted with an amino acid
having an the side chain -(CH₂)ₙNH₂, where n is different from 4, and arginine may be
substituted with an amino acid having the side chain -(CH₂)ₙNHC(=NH₂)NH₂, where n is
different from 3, etc. Similarly, the acidic amino acids aspartic acid and glutamic acid may be
substituted with amino acid residues having the side chains -(CH₂)ₙCOOH, where n>2.

A "retro form" of a peptide is a form of a peptide where the order of the amino acids
in N- to C-terminal direction has been inverted. For instance, the retro form of ALDFR is the
peptide RFDLA.

The term "substantially identical" in the context of two amino acid sequences means
that the sequences, when optimally aligned, such as by the programs GAP or BESTFIT using
default gap weights, share at least about 50, at least about 60, at least about 70, at least
about 80, at least about 90, at least about 95, at least about 98, or at least about 99 percent
sequence identity. In one embodiment, residue positions that are not identical differ by
conservative amino acid substitutions. Sequence identity is typically measured using
sequence analysis software. Protein analysis software matches similar sequences using
measures of similarity assigned to various substitutions, deletions and other modifications,
including conservative amino acid substitutions. For instance, the publicly available GCG
software contains programs such as "Gap" and "BestFit" which can be used with default
parameters to determine sequence homology or sequence identity between closely related
polypeptides, such as homologous polypeptides from different species of organisms or
between a wild-type protein and a mutein thereof. See, e.g., GCG Version 6.1. Polypeptide
sequences can also be compared using FASTA or ClustalW, applying default or recommended
parameters. A program in GCG Version 6.1., FASTA (e.g., FASTA2 and FASTA3) provides
alignments and percent sequence identity of the regions of the best overlap between the
query and search sequences (Pearson, Methods Enzymol. 1990;183:63-98; Pearson, Methods
Mol. Biol. 2000;132:185-219). Another preferred algorithm when comparing a sequence to a
database containing a large number of sequences from various organisms, or when deducing
the is the computer program BLAST, especially blastp, using default parameters. See, e.g.,
1997;25:3389-402 (1997); each herein incorporated by reference. "Corresponding" amino
acid positions in two substantially identical amino acid sequences are those aligned by any of
the protein analysis software mentioned herein, typically using default parameters.

A nucleic acid is "operably linked" when it is placed into a functional relationship with
another nucleic acid sequence. For example, DNA for a presequence or secretory leader is
operably linked to DNA for a polypeptide if it is expressed as a preprotein that participates in
the secretion of the polypeptide; a promoter or enhancer is operably linked to a coding
sequence if it affects the transcription of the sequence; or a ribosome-binding site is operably
linked to a coding sequence if it is positioned so as to facilitate translation. Generally,
"operably linked" means that the DNA sequences being linked are contiguous, and, in the
case of a secretory leader, contiguous and in reading phase. However, enhancers do not have
to be contiguous. Linking is accomplished by ligation at convenient restriction sites. If such sites do not exist, the synthetic oligonucleotide adaptors or linkers are used in accordance with conventional practice.

An "isolated" molecule is a molecule that is the predominant species in the composition wherein it is found with respect to the class of molecules to which it belongs (i.e., it makes up at least about 50% of the type of molecule in the composition and typically will make up at least about 70%, at least about 80%, at least about 85%, at least about 90%, at least about 95%, or more of the species of molecule, e.g., peptide, in the composition). Commonly, a composition of an antibody molecule will exhibit 98% - 99% homogeneity for antibody molecules in the context of all present peptide species in the composition or at least with respect to substantially active peptide species in the context of proposed use.

In the context of the present invention, "treatment" or "treating" refers to preventing, alleviating, managing, curing or reducing one or more symptoms or clinically relevant manifestations of a disease or disorder, unless contradicted by context. For example, "treatment" of a patient in whom no symptoms or clinically relevant manifestations of a disease or disorder have been identified is preventive or prophylactic therapy, whereas "treatment" of a patient in whom symptoms or clinically relevant manifestations of a disease or disorder have been identified generally does not constitute preventive or prophylactic therapy.

The term antigen denotes a substance of matter which is recognized by the immune system's specifically recognizing components (antibodies, T-cells).

The term "immunogen" is in the present context intended to denote a substance of matter, which is capable of inducing an adaptive immune response in an individual, where said adaptive immune response targets the immunogen. In other words, an immunogen is an antigen, which is capable of inducing immunity.

The terms "epitope", "antigenic determinant" and "antigenic site" are used interchangeably herein and denotes the region in an antigen or immunogen which is recognized by antibodies (in the case of antibody binding epitopes, also known as "B-cell epitopes") or by T-cell receptors when the epitope is complexed to an MHC molecule (in the case of T-cell receptor binding epitopes, i.e. "T-cell epitopes").

The term "immunogenically effective amount" has its usual meaning in the art, i.e. an amount of an immunogen, which is capable of inducing an immune response, which significantly engages pathogenic agents, which share immunological features with the immunogen.

The term "vaccine" is used for a composition comprising an immunogen and which is capable of inducing an immune response which is either capable of reducing the risk of developing a pathological condition or capable of inducing a therapeutically effective immune
response which may aid in the cure of (or at least alleviate the symptoms of) a pathological condition.

The term “pharmaceutically acceptable” has its usual meaning in the art, i.e. it is used for a substance that can be accepted as part of a medicament for human use when treating the disease in question and thus the term effectively excludes the use of highly toxic substances that would worsen rather than improve the treated subject’s condition.

A “T helper lymphocyte epitope” (a Th epitope) is peptide, which binds an MHC Class II molecule and can be presented on the surface of an antigen presenting cell (APC) bound to the MHC Class II molecule. An “immunological carrier” is generally a substance of matter which includes one or many Th epitopes, and which increase the immune response against an antigen to which it is coupled by ensuring that T-helper lymphocytes are activated and proliferate. Examples of known immunological carriers are the tetanus and diphtheria toxoids and keyhole limpet hemocyanin (KLH).

The term “adjuvant” has its usual meaning in the art of vaccine technology, i.e. a substance or a composition of matter which is 1) not in itself capable of mounting a specific immune response against the immunogen of the vaccine, but which is 2) nevertheless capable of enhancing the immune response against the immunogen. Or, in other words, vaccination with the adjuvant alone does not provide an immune response against the immunogen, vaccination with the immunogen may or may not give rise to an immune response against the immunogen, but the combined vaccination with immunogen and adjuvant induces an immune response against the immunogen which is stronger than that induced by the immunogen alone.

The term “HIV-1 DNA levels” as used herein refers to the total amount of copies of measurable cellular human immunodeficiency virus-1 (HIV-1) DNA in non-integrated, circular and well as integrated forms in copies per $10^6$ CD4+ T cells of peripheral blood obtained from patients infected with HIV-1.

Specific aspects and embodiments of the invention

One aspect of the present invention relates to the use of one or more HIV-specific peptide as defined above.

In certain embodiments, peptides comprise an N- or C-terminal modification, such as an amidation, acylation, or acetylation. When the C-terminal end of a peptide is an amide, suitable amides included those having the formula -C(O)-NR=R', wherein R and R' are independently selected from hydrogen and C₁-₆ alkyl, which alkyl group may be substituted with one of more fluoro atoms, for example -CH₃, -CH₂CH₃ and -CF₃, a particular amide group which may be mentioned is -C(O)NH₂. When the N-terminal end of the peptide is acetylated, suitable acetylated N-terminal ends include those of formula -NH-C(O)R₂, wherein R₂ is hydrogen, C₁-₆ alkyl, which alkyl group may be substituted with one of more fluoro atoms, for example -CH₃, -CH₂CH₃ and -CF₃, or phenyl.
Since the peptides are contemplated as vaccine agents, they are in certain embodiments coupled to a carrier molecule, such as an immunogenic carrier. The peptides may thus be linked to other molecules either as recombinant fusions (e.g. via CLIP technology) or through chemical linkages in an oriented (e.g. using heterobifunctional cross-linkers) or nonoriented fashion. Linking to carrier molecules such as for example diphtheria toxin, polylysine constructs etc, are all possible according to the invention.

The immunogenic carrier is conveniently selected from carrier proteins such as those conventionally used in the art (e.g. diphtheria or tetanus toxoid, KLH etc.), but it is also possible to use shorter peptides (T-helper epitopes) which can induce T-cell immunity in larger proportions of a population. Details about such T-helper epitopes can e.g. be found in WO 00/20027, which is hereby incorporated by reference herein – all immunologic carriers and “promiscuous” (i.e. universal) T-helper epitopes discussed therein are useful as immunogenic carriers in the present invention.

In certain embodiments, the carrier is a virus like particle, i.e. a particle sharing properties with virions without being infectious. Such virus-like particles may be provided chemically (e.g. Jennings and Bachmann Ann. Rev. Pharmacol. Toxicol. 2009. 49:303-26 Immunodrugs: Therapeutic VLP-based vaccines for chronic diseases) or using cloning techniques to generate fusion proteins (e.g. Peabody et al. J. Mol. Biol. 2008; 380: 252-63. Immunogenic display of diverse peptides on virus-like particles of RNA phage MS2). Another example is “Remune”, an HIV vaccine originally made by Immune Response Corporation, which consists of formalin inactivated HIV that has been irradiated to destroy the viral genome. The company was started by Jonas Salk who used the same technique to generate the killed polio vaccine in widespread use today.

One aspect of the present invention relates to the use of an immunogenic composition (such as a vaccine composition) comprising a composition of at least one HIV-specific peptides, in combination with an effective amount of a reservoir purging agent, optionally together with a pharmaceutically acceptable diluent or vehicle and optionally one or more immunological adjuvant.

In common for aspects of the invention is that they all include embodiments where the at least one HIV-specific peptide is selected from the group of amino acid sequences of SEQ ID NOs: 1, 4, 9 and 15, as defined above; wherein the terminal ends of each HIV specific peptide may be free carboxyl- or amino- groups, amides, acyls or acetyl; and in the form of an acetate salt.

In some embodiments two or more of the Cys residues of said HIV-specific peptide may form part of an intrachain- or interchain disulphide binding, a -S-(CH₂)ₚ-S-, or a -(CH₂)ₚ-bridge wherein p = 1-8 optionally intervened by one or more heteroatoms such as O, N and S and/or the said peptide sequences are immobilized to a solid support.

In some embodiments the amino acid sequence of SEQ ID NO: 1 is selected from the group of SEQ ID NO: 2 and SEQ ID NO: 3.
In some embodiments the amino acid sequence of SEQ ID NO: 4 is selected from the group of SEQ ID NO: 5, SEQ ID NO: 6, SEQ ID NO: 7 and SEQ ID NO: 8.

In some embodiments the amino acid sequence of SEQ ID NO: 9 is selected from the group of SEQ ID NO: 10, SEQ ID NO: 11, SEQ ID NO: 12, SEQ ID NO: 13 and SEQ ID NO: 14.

In some embodiments the amino acid sequence of SEQ ID NO: 15 is selected from the group of SEQ ID NO: 16, SEQ ID NO: 17, SEQ ID NO: 18, SEQ ID NO: 19 and SEQ ID NO: 20.

In some embodiments the at least one HIV-specific peptide comprises at least, two, three, or four peptides selected from each of the groups of SEQ ID NO: 1, SEQ ID NO: 4, SEQ ID NO: 9 and SEQ ID NO: 15.

In some embodiments the at least one HIV-specific peptide consists of or comprises the peptides of SEQ ID NO: 3, SEQ ID NO: 6, SEQ ID NO: 11 and SEQ ID NO: 18.

Preparation of immunogenic compositions includes the use of state-of-the-art constituents such as immunological adjuvants. Apart from these adjuvants, which are detailed, by way of example, below, immunogenic compositions are prepared as generally taught in the art:

The preparation of vaccines which contain peptide sequences as active ingredients is generally well understood in the art, as exemplified by U.S. Patents 4,608,251; 4,601,903; 4,599,231; 4,599,230; 4,596,792; and 4,578,770, all incorporated herein by reference. Typically, such vaccines are prepared as injectable either as liquid solutions or suspensions; solid forms suitable for solution in, or suspension in, liquid prior to injection may also be prepared. The preparation may also be emulsified. The active immunogenic ingredient is often mixed with excipients which are pharmaceutically acceptable and compatible with the active ingredient. Suitable excipients are, for example, water, saline, dextrose, glycerol, ethanol, or the like, and combinations thereof. In addition, if desired, the vaccine may contain minor amounts of auxiliary substances such as wetting or emulsifying agents, pH buffering agents, or adjuvants which enhance the effectiveness of the vaccines; cf. the detailed discussion of adjuvants below.

The vaccines are conventionally administered parenterally, by injection, for example, either subcutaneously, intracutaneously, intradermally, subdermally or intramuscularly. Additional formulations which are suitable for other modes of administration include suppositories and, in some cases, oral, nasal, buccal, sublingual, intraperitoneal, intravaginal, anal, epidural, spinal, and intracranial formulations. For suppositories, traditional binders and carriers may include, for example, polyalkalene glycols or triglycerides; such suppositories may be formed from mixtures containing the active ingredient in the range of 0.5% to 10% (w/w), preferably 1-2% (w/w). Oral formulations include such normally employed excipients as, for example, pharmaceutical grades of mannitol, lactose, starch, magnesium stearate, sodium saccharine, cellulose, magnesium carbonate, and the like. These compositions take
the form of solutions, suspensions, tablets, pills, capsules, sustained release formulations or powders and may contain 10-95% (w/w) of active ingredient, preferably 25-70% (w/w).

The peptides may be formulated into a vaccine as neutral or salt forms. Pharmaceutically acceptable salts include acid addition salts (formed with the free amino groups of the peptide) and which are formed with inorganic acids such as, for example, hydrochloric or phosphoric acids, or such organic acids as acetic, oxalic, tartaric, mandelic, and the like. Salts formed with the free carboxyl groups may also be derived from inorganic bases such as, for example, sodium, potassium, ammonium, calcium, or ferric hydroxides, and organic bases such as isopropylamine, trimethylamine, 2-ethylamino ethanol, histidine, procaine, and the like.

The vaccines are administered in a manner compatible with the dosage formulation, and in such amount as will be therapeutically effective and immunogenic. The quantity to be administered depends on the subject to be treated, including, e.g., the capacity of the individual's immune system to mount an immune response, and the degree of immunity desired. Suitable dosage ranges are of the order of several hundred micrograms of active ingredient per vaccination with a preferred range from about 0.1 μg to 2,000 μg (even though higher amounts in the 1-10 mg range are contemplated), such as in the range from about 0.5 μg to 1,800 μg, preferably in the range from 1 μg to 1,500 μg and especially in the range from about 100 μg to 1200 μg. Suitable regimens for initial administration and booster shots are also variable but are typified by an initial administration followed by subsequent inoculations or other administrations.

Some of the peptides are sufficiently immunogenic in a vaccine, but for some of the others the immune response will be enhanced if the vaccine further comprises an adjuvant substance. The immunogenic molecules described herein can therefore be formulated with adjuvants:

The adjuvants to be combined are known to induce humoral responses and include: i) Salt suspensions (e.g. varieties of salts containing aluminum ions or calcium ions), ii) Oil-in-water emulsions (e.g. varieties of squalane-based or squalene-based emulsions), iii) Water-in-oil emulsions (e.g. Montanide ISA51 or ISA720), iv) Neutral liposomes, v) Cationic liposomes, vi) Microspheres, vii) Immunostimulating complexes (e.g. ISCOMs or ISCOMATRIX), viii) Pattern-recognition receptor agonists (e.g. agonists for C-type lectin receptors (CLRs), NOD-like receptors (NLRs), RIG-like helicases (RLHs), Triggering receptor expressed on myeloid cells (TREM) and Toll-like receptors (TLRs)), ix) Saponins (i.e. Any saponin derived from Quillaja saponaria or Platycodon grandiflorum), x) Virosomes/Virus-like particles, xi) Enterotoxins (i.e. Cholera toxin, CTA1-DD or Escherichia coli heat-labile enterotoxin), and combinations thereof.

For a further enhancement of the vaccine antigenic properties, they could be combined with a well-known adjuvant with an oral immune modulant or adjuvant such as a Cox-2 inhibitor or an immunomodulating compound.
A further aspect of the invention is the use of the vaccine combined with adjuvant, with one or more further therapeutic agents, such as an (oral) immunomodulating agent and/or a second reservoir purging agent.

The terms “therapeutic agent”, such as “immunomodulating agent” or virus reservoir purging agent as used herein, includes but is not limited to cytokines, such as interferons, monoclonal antibodies, such as anti-PD1 antibodies, cyclophosphamide, Thalidomide, Levamisole, and Lenalidomide.

“A virus reservoir purging agent”, includes but is not limited to auranofin, IL-7, prostratin, bryostatin, HDAC inhibitors, such as vorinostat, Disulfiram and any suitable agent disclosed in any one of WO2013050422, WO2012051492 A3 and in Barton et al., Clinical Pharmacology & Therapeutics (2013); 93 1, 46–56, including but not limited to a NF-kappa-B-inducer selected from the group comprising: PMA, prostratin, bryostatin and TNF-alpha, and/or b) a histone deacetylase inhibitor selected from the different families (hydroxamates, cyclic peptides, aliphatic acids, and benzamides) including: TSA, SAHA, MS-275, aminosuberoyl hydroxamic acids, M-Carboxycinnamic acid bishydroxamate, LAQ-824, LBH-589, belinostat (PXD-101 ), Panobinostat (LBH-589), a cinnamic hydroxamic acid analogue of M-carboxycinnamic acid bishydroxamate, IF2357, aryloxyalkanoic acid hydroxamides, depsipeptide, apicidin, cyclic hydroxamic acid- containing peptide group of molecules, FK-228, red FK, cyclic peptide mimic linked by an aliphatic chain to a hydroxamic acid, butyrate, phenylbutyrate, sodium butyrate, valproic acid, pivaloyloxymethyl butyrate, 5 NOX-275, and MGCD0103. Any of the above virus reservoir purging agents may be used alone or in combination with any one other suitable virus reservoir purging agent, such as with another class of HIV inducers.

DNA methylation, probably together with repressive histone modifications, may also contribute to a "lock" in a silent state of the provirus and makes its return to an active state difficult. These observations suggest that HDAC or HMT or DNA methylation inhibitors together with efficient cART constitute good anti-latency drug candidates aimed at reducing/eliminating the pool of latent reservoirs to a level bearable by the host immune system.

Accordingly suitable immunomodulatory compounds or (reservoir) purging agents may be DNA methylation inhibitors selected from the two classes (non-nucleoside and nucleoside demethylating agents) including: 5-azacytidine (azacitidine), Sinefungin, 5-aza-2'-deoxycytidine (5-aza-CdR, decitabine), 1-3-Darabinofuranosyl-5-azacytosine (fazarabine) and dihydro-5-azacytidine (DHAC), 5-fluorodeoxycytidine (FdC), oligodeoxynucleotide duplexes containing 2-H pyrimidinone, zebularine, antisense oligodeoxynucleotides (ODNs), MG98, (-)-epigallocatechin-3-gallate, hydralazine, procaine and procarbamide.

Other suitable immunomodulatory compounds or (reservoir) purging agents to be used according to the present invention includes histone deacetylase inhibitor selected from the different families of HDACI (hydroxamates, cyclic peptides, aliphatic acids, and
benzamides) including TSA, SAHA, MS-275, aminosuberoyl hydroxamic acids, M-Carboxycinnamic acid bishydroxamate, LAQ-824, LBH-589, belinostat (PXD-101 ), Panobinostat (LBH-589), a cinnamic hydroxamic acid analogue of M-carboxycinnamic acid bishydroxamate, IF2357, aryloxyalkanoic acid hydroxamides, depsipeptide, apicidin, cyclic hydroxamic acid-containing peptide group of molecules, FK-228, red FK, cyclic peptide mimic linked by an aliphatic chain to a hydroxamic acid, butyrate, phenylbutyrate, sodium butyrate, valproic acid, pivaloyloxymethyl butyrate, 5 NOX-275, MGCD0103, BET family protein inhibitors/antagonists, such as JQ1, I-BET, I-Bet151, MS417, and GW841819X (Nicodeme et al. (2010) Nature 468:1119-1123; Filippakopoulos et al. (2010) Nature 468:1067-1073), and thienotriazolodiazepine compounds, such as those described in U.S. Patent Application Publication No. 2010/0286127.

Other suitable immunomodulatory compounds or (reservoir) purging agents to be used according to the present invention includes histone methyltransferase inhibitors (chaetocin and BIX-01294); Inhibitors of Enhances of Zeste 2 (EZH2) – such as 3-deazaneplanocin A (DZNep) used alone or in combination with other classes of immunomodulatory compounds or (reservoir) purging agents.

Other suitable adjuvants include response-selective C5a agonists, such as EP54 and EP67 described in Hung CY et al. An agonist of human complement fragment C5a enhances vaccine immunity against Coccidioides infection. Vaccine (2012) and Kollessery G et al.


Various methods of achieving adjuvant effect for the vaccine are thus known.


In the methods and compositions of the invention the at least one HIV-specific peptide and the reservoir purging agent, may be administered in combination with one or more further therapeutically active agents, such as agents for the treatment of HIV and/or AIDS.

The failure of antiretroviral therapy (ART) to eradicate HIV-1 infection lies in the observation that HIV-1 remains quiescent in latent reservoirs. Latently infected resting CD4+ cells (either naive or long lived memory cells) carry transcriptionally silent HIV-1 and represent the predominant reservoir of HIV-1 infection. Other cells may also act as reservoirs (Reviewed in Alexaki et al., 2008, Curr. HIV Res. 6:388-400), such as macrophages, dendritic cells and astrocytes (where HIV-1 infection occurs via a CD4-independent mechanism). It is these latent reservoirs that represent the major challenge to eradication of HIV-1 infection. Approaches towards eradication include attempts to purge reservoirs by selective activation of latently infected cells (such as memory cells) in the presence of ART such that released virus may not infect and replicate in neighbouring cells (Richman et al., 2009, Science 323:1304-1307). Agents include histone deacetylase inhibitors, cytokines, such as IL-2 and IL-7, as well as bryostatin, the protein kinase C activator (Kovochich et al., 2011, PLoS ONE 6 (4):e18270). Therapeutic vaccines have the advantage of being able to penetrate sanctuary sites less well accessed by ART such as lymphoid tissue (Pantaleo et al., 1991, Proc. Natl. Acad. Sci. USA 88:9838–42; Fox et al., 1991, J. Infect. Dis. 164:1051–57) and the central nervous system (Alexaki et al., 2008, Curr. HIV Res. 6:388-400), that represent regions for viral persistence. This relates to therapeutic interventions targeting both the virus itself as well as HIV-associated immune activation.

A number of studies have been conducted with the aim of providing compounds that can safely and effectively be used to treat diseases associated with abnormal production of TNF-α. See, e.g., Marriott, J.B., et al, Expert Opin. Biol. Ther. (4): 1-8 (2001); G.W. Muller, et al, Journal of Medicinal Chemistry, 39(17): 3238-3240 (1996); and G.W. Muller, et al, Bioorganic & Medicinal Chemistry Letters, 8: 2669-2674 (1998). Some studies have focused on a group of compounds selected for their capacity to potently inhibit TNF-α production by LPS stimulated PBMC. L.G. Corral, et al, Ann. Rheum. Dis., 58 (suppl 1): 1107-1113 (1999). These compounds, often referred to as immunomodulatory compounds, show not only potent inhibition of TNF-α, but also marked inhibition of LPS induced monocyte IL1β and IL12 production. LPS induced IL6 is also inhibited by immunomodulatory compounds, albeit partially. These compounds are potent stimulators of LPS induced IL10. Particular examples include, but are not limited to, the substituted 2-(2,6-dioxopiperidin-3-yl)phthalimides and substituted 2-(2,6-dioxopiperidin-3-yl)-1-oxoisindoles as described in US 6281230 and US
Monocyte/macrophage function is part of the Innate Immune System that serves as a first line of defense against an infection. By modulating the host's monocytes and macrophages, immunomodulatory compounds can change the dynamics of the response to a viral infection, such as influenza.

Histone deacetylases (HDAC) are a class of enzymes that remove acetyl groups from N-acetylated lysines amino acid on histone proteins. Currently 18 HDACs have been identified in mammals. They have been divided into four classes based on cellular localization, function, and sequence similarity. Class I includes HDACs 1, 2, 3, and 8 which are found primarily in the nucleus. Class II HDACs (HDACs 4, 5, 6, 7, 9, and 10) are found primarily in the cytoplasm but may be able to shuttle between the nucleus and the cytoplasm; class IIa comprises four HDACs (HDACs 4, 5, 7 and 9) while class IIb comprises two HDACs (HDACs 6 and 10) which are expressed only in the cytoplasm. HDAC11, which is ubiquitously expressed, shares sequence similarities with both class I and class II HDACs and represents Class IV. Class III (also called "sirtuin family") groups NAD+-dependent proteins which do not act primarily on histones.

In the methods of the invention the at least one HIV-specific peptide, may optionally be administered with a reservoir purging agent, and optionally together with another immunomodulatory compound and/or a second reservoir purging agent, such as another histone deacetylase (HDAC) inhibitor.

The immunomodulatory compounds may be selected from anti-PD1 antibodies, such as MDX-1106 (Merck), THALOMID® (thalidomide), anti-PD1 antibodies, cyclophosphamide, Levamisole, lenalidomide, CC-4047 (pomalidomide), CC-11006 (Celgene), and CC-10015 (Celgene), and immunomodulatory compound described in any one of WO2007028047, WO2002059106, and WO2002094180. The immunomodulatory compound may be selected from 4-((amino)-2-(2,6-dioxo(3-piperidyl))-isoindoline-1,3-dione and 3-(4-amino-1-oxo-1,3-dihydro-isoindol-2-yl)-piperidine-2,6-dione. In particular the immunomodulatory compound is lenalidomide. The immunomodulatory compound may be enantiomerically pure. The second reservoir purging agent, such as a histone deacetylase (HDAC) inhibitor, may be selected from M344 (4-(dimethylamino)-N-[7-(hydroxyamino)-7-oxoheptyl]benzamide), chidamide (CS055/HBI-800), 4SC-202, (4SC), Resminostat (4SC), hydroxamic acids such as vorinostat (SAHA), belinostat (PXD101), LAQ824, trichostatin A and panobinostat (LBH589); benzamides such as entinostat (MS-275), CI994, and mocetinostat (MGCD0103), cyclic tetrapeptides (such as trapoxin, such as trapoxin B), and the depsipeptides, such as romidepsin (Istodax® (Celgene)), electrophilic ketones, and the aliphatic acid compounds such as phenylbutyrate, valproic acid, Oxamflatin, ITF2357 (generic givinostat), Apicidin, MC1293, CG05, and CG06; compounds that activate transcription factors including NF-KappaB, Prostratin, auranofin, bryostatin, a nontumorigenic phorbol ester, DPP (12-deoxyphorbol-13-phenylacetate), PMA, and Phorbol 12-myristate 13-acetate (PMA); Compounds that activate HIV mRNA elongation including P-TEF-b kinase and
hexamethylbisacetamide (HMBA); IL-7; T-cell stimulating factors including anti-CD3/CD28 - T-cell stimulating Ab's; Kinase inhibitors including Tyrphostin A, Tyrphostin B, and Tyrphostin C; PTEN (phosphatase and tensin homologue) gene inhibitors including SF1670 (Echelon Bioscience), Disulfiram (DSF), an inhibitor of acetaldehyde dehydrogenase, Protein Tyrosine Phosphatase Inhibitors including bpV(HOpic), bpV(phen), and bpV(pic) (Calbiochem; EMD Millipore), Toll-like receptors agonists including Toll-like receptor-9 (TLR9) and Toll-like receptor-7 (TLR9) agonists, quercetin, lipoic acid, sodium butyrate, TNF-alpha, PHA, Tat.

In the methods of the invention the components of the at least one HIV-specific peptide and/or the one or more further therapeutically active agents, may be administered simultaneously, sequentially or separately in any order.

Thus the invention provides a pharmaceutical composition comprising one, two or more components of the at least one HIV-specific peptide and/or the one or more further therapeutically active agents optionally in combination with one or more pharmaceutically acceptable adjuvants, diluents or carriers.

Similarly, the invention also provides a combination product comprising of components of the at least one HIV-specific peptide and/or the one or more further therapeutically active agents, wherein each of component is formulated in admixture with a pharmaceutically-acceptable adjuvant, diluent or carrier. In this aspect of the invention, the combination product may be either a single (combination) pharmaceutical formulation or a kit-of-parts. In a kit-of-parts some or all of the components may be formulated separately and may each be provided in a form that is suitable for administration in conjunction with the other(s).

The component(s) may also be provided for use, e.g. with instructions for use, in combination with one or more further component(s) as defined above.

The peptides for use in the invention may be produced synthetically using art recognised methods. Further details for the synthetic production of such peptides are found in the Examples. Alternatively peptides may be produced recombinantly, if possible. When recombinantly producing peptides for use in the invention by means of transformed cells, it is convenient, although far from essential, that the expression product is either exported out into the culture medium or carried on the surface of the transformed cell.

When an effective producer cell has been identified it is preferred, on the basis thereof, to establish a stable cell line which carries the vector of the invention and which expresses the nucleic acid fragment of the invention. Preferably, this stable cell line secretes or carries the peptide expression product, thereby facilitating purification thereof.

In general, plasmid vectors containing replicon and control sequences which are derived from species compatible with the host cell are used in connection with the hosts. The vector ordinarily carries a replication site, as well as marking sequences which are capable of providing phenotypic selection in transformed cells. For example, *E. coli* is typically transformed using pBR322, a plasmid derived from an *E. coli* species (see, e.g., Bollivar *et al.*,}
The pBR322 plasmid contains genes for ampicillin and tetracycline resistance and thus provides easy means for identifying transformed cells. The pBR plasmid, or other microbial plasmid or phage must also contain, or be modified to contain, promoters which can be used by the prokaryotic microorganism for expression.

Those promoters most commonly used in recombinant DNA construction include the β-lactamase (penicillinase) and lactose promoter systems (Chang et al., 1978; Itakura et al., 1977; Goeddel et al., 1979) and a tryptophan (trp) promoter system (Goeddel et al., 1979; EP-A-0 036 776). While these are the most commonly used, other microbial promoters have been discovered and utilized, and details concerning their nucleotide sequences have been published.

In addition to prokaryotes, eukaryotic microbes, such as yeast cultures may also be used, and also here the promoter should be capable of driving expression. Saccharomyces cerevisiae, or common baker’s yeast is the most commonly used among eukaryotic microorganisms, although a number of other strains are commonly available. For expression in Saccharomyces, the plasmid YRp7, for example, is commonly used (Stinchcomb et al., 1979; Kingsman et al., 1979; Tschemper et al., 1980).

Suitable promoting sequences in yeast vectors include the promoters for 3-phosphoglycerate kinase (Hitzman et al., 1980) or other glycolytic enzymes (Hess et al., 1968; Holland et al., 1978), such as enolase, glyceraldehyde-3-phosphate dehydrogenase, hexokinase, pyruvate decarboxylase, phosphofructokinase, glucose-6-phosphate isomerase, 3-phosphoglycerate mutase, pyruvate kinase, triosephosphate isomerase, phosphoglucone isomerase, and glucokinase. In constructing suitable expression plasmids, the termination sequences associated with these genes are also incorporated into the expression vector 3’ of the sequence desired to be expressed to provide polyadenylation of the mRNA and termination.

Other promoters, which have the additional advantage of transcription controlled by growth conditions are the promoter region for alcohol dehydrogenase 2, isocitrochrome C, acid phosphatase, degradative enzymes associated with nitrogen metabolism, and the aforementioned glyceraldehyde-3-phosphate dehydrogenase, and enzymes responsible for maltose and galactose utilization. Any plasmid vector containing a yeast-compatible promoter, origin of replication and termination sequences is suitable.

In addition to microorganisms, cultures of cells derived from multicellular organisms may also be used as hosts. In principle, any such cell culture is workable, whether from vertebrate or invertebrate culture. Examples of such useful host cell lines are VERO and HeLa cells, Chinese hamster ovary (CHO) cell lines, and W138, Per.C6, BHK, COS-7 293, Spodoptera frugiperda (SF) cells, Drosophila melanogaster cell lines (such as Schneider 2 (S2)), and MDCK cell lines.

Expression vectors for such cells ordinarily include (if necessary) an origin of replication, a promoter located in front of the gene to be expressed, along with any
necessary ribosome binding sites, RNA splice sites, polyadenylation site, and transcriptional terminator sequences.

For use in mammalian cells, the control functions on the expression vectors are often provided by viral material. For example, commonly used promoters are derived from polyoma, Adenovirus 2, and most frequently Simian Virus 40 (SV40). The early and late promoters of SV40 virus are particularly useful because both are obtained easily from the virus as a fragment which also contains the SV40 viral origin of replication (Fiers et al., 1978). Smaller or larger SV40 fragments may also be used, provided there is included the approximately 250 bp sequence extending from the HindIII site toward the BglII site located in the viral origin of replication. Further, it is also possible, and often desirable, to utilize promoter or control sequences normally associated with the desired gene sequence, provided such control sequences are compatible with the host cell systems.

An origin of replication may be provided either by construction of the vector to include an exogenous origin, such as may be derived from SV40 or other viral (e.g., other Polyoma viruses, Adeno, VSV, BPV) or may be provided by the host cell chromosomal replication mechanism. If the vector is integrated into the host cell chromosome, the latter is often sufficient.

As for routes of administration and administration schemes of polypeptide based vaccines which have been detailed above, these are also applicable for the nucleic acid vaccines of the invention and all discussions above pertaining to routes of administration and administration schemes for polypeptides apply mutatis mutandis to nucleic acids. To this should be added that nucleic acid vaccines can also be administered intravenously and intraarterially. Furthermore, it is well-known in the art that nucleic acid vaccines can be administered by use of a so-called gene gun and/or by use of electroporation, and hence also these and equivalent modes of administration are regarded as part of the present invention.

Under normal circumstances, the nucleic acid fragment is introduced in the form of a vector wherein expression is under control of a viral promoter. For more detailed discussions of vectors according to the invention, cf. the discussion above. Also, detailed disclosures relating to the formulation and use of nucleic acid vaccines are available, cf. Donnelly JJ et al., 1997, Annu. Rev. Immunol. 15: 617-648 and Donnelly JJ et al., 1997, Life Sciences 60: 163-172. Both of these references are incorporated by reference herein.

An alternative of using peptide immunogens or nucleic acid immunogens is the use of live immunogen technology. This entails administering a non-pathogenic microorganism which has been transformed with a nucleic acid fragment or a vector of the present invention. The non-pathogenic microorganism can be any suitable attenuated bacterial strain (attenuated by means of passaging or by means of removal of pathogenic expression products by recombinant DNA technology), e.g. Mycobacterium bovis BCG., non-pathogenic Streptococcus spp., E. coli, Salmonella spp., Vibrio cholerae, Shigella, etc. Reviews dealing with preparation of state-of-the-art live vaccines can e.g. be found in Saliou P, 1995, Rev.
Prat. 45: 1492-1496 and Walker PD, 1992, Vaccine 10: 977-990, both incorporated by reference herein. For details about the nucleic acid fragments and vectors used in such live vaccines, cf. the discussion below.

As an alternative to bacterial live immunogens, the nucleic acid fragment of the invention can be incorporated in a non-virulent viral vaccine vector such as a vaccinia strain or any other suitable poxvirus.

Normally, the non-pathogenic microorganism or virus is administered only once to a subject, but in certain cases it may be necessary to administer the microorganism/virus more than once in a lifetime in order to maintain protective immunity. It is even contemplated that immunization schemes as those detailed above for polypeptide vaccination will be useful when using live or virus vaccines.

Alternatively, live or virus immunization is combined with previous or subsequent polypeptide and/or nucleic acid immunization. For instance, it is possible to effect primary immunization with a live or virus vaccine followed by subsequent booster immunizations using the polypeptide or nucleic acid approach.

PREAMBLE TO EXAMPLES

20 HIV-SPECIFIC PEPTIDES FOR USE ACCORDING TO THE INVENTION

The present invention involves the use of HIV-specific peptides based on conserved regions of HIV gag p24, antigens in free or carrier-bound form comprising at least one of the said peptides.

The HIV-specific peptides for use according to the invention originate from the four different conserved areas of the HIV-1 core protein p24, having the properties of maintaining the uniqueness (sensitivity and specificity) of the HIV-1-epitope. Further these peptides possess no recognized cytotoxic T lymphocyte (CTL) antagonistic effect and have at least one potential CTL epitope.

The HIV-specific peptides, for use according to the invention, which have met the above criteria are selected from the group of amino acid sequences of SEQ ID NOs: 1, 4, 9 and 15, as defined above; wherein the terminal ends of each HIV specific peptide may be free carboxyl- or amino- groups, amides, acyls or acetyl; or acetate salts of any of the HIV specific peptides.

The HIV-specific peptide sequences have the potential to serve as a particularly good antigen wherein the antigen comprises at least one peptide selected from the group of sequences of SEQ ID NO: 1, SEQ ID NO: 4, SEQ ID NO: 9 or SEQ ID NO: 15. The antigenicity may be adapted through adjusting the ratio or concentration of different peptides or size of the peptides by for instance dimerisation or polymerisation and/or immobilisation to a solid
phase. The antigen may comprise two or more polypeptide sequences which are either linked
by a bridge for instance a disulphide bridge between the Cys residues of the chains or bridges
like C₁₋₆ alkylene possibly intervened by one or more heteroatoms like O, S, or N or
preferably they are unlinked. The chains may be immobilized to a solid phase in monomeric,
dimeric or oligomeric forms. Further amino acids may be added to the ends in order to
achieve an «arm» to facilitate immobilization.

All amino acids in the HIV-specific peptides of the invention can be in both D- or L-
form, although the naturally occurring L-form is preferred.

The C- and N-terminal ends of the HIV-specific peptide sequences could deviate
from the natural sequences by modification of the terminal NH₂-group and/or COOH-group,
they may for instance be acylated, acetylated, amidated or salts thereof; or modified to
provide a binding site for a carrier or another molecule. When the C-terminal end of a peptide
is an amide, suitable amides included those having the formula -C(O)-NR'R', wherein R' and
R' are independently selected from hydrogen and C₁₋₆ alkyl, which alkyl group may be
substituted with one of more fluoro atoms, for example -CH₃, -CH₂CH₃ and -CF₃, a particular
amide group which may be mentioned is -C(O)NH₂. When the N-terminal end of the peptide
is acetylated, suitable acetylated N-terminal ends include those of formula -NH-C(O)R',
wherein R' is hydrogen, C₁₋₆ alkyl, which alkyl group may be substituted with one of more
fluoro atoms, for example -CH₃, -CH₂CH₃ and -CF₃, or phenyl.

The HIV-specific peptides for use according to the invention consist of 6 to 50 amino
acids, preferably between 10 and 30 amino acids. They cover all natural variation of amino
acids in the identified positions.

The polypeptide antigen for use according to the invention is either in a free or in a
carrier-bound form. The carrier or solid phase to which the peptide is optionally bound can be
selected from a wide variety of known carriers. It should be selected with regard to the
intended use of the immobilized polypeptide as an immunizing component in a vaccine.

In a preferred embodiment the HIV specific peptides for use according to the
present invention comprises antigens containing the peptides of the SEQ ID NOs: 1, 4, 9 and
15, more preferably the peptides occur in the ratio 1:1:1:1 w/w.

In a further preferred embodiment the HIV specific peptides for use according to the
invention comprise the following:

RALGPAATLQTPWTASLGVG (SEQ ID NO: 3)
RWLLLGLNLVGGGRLYSPTSILG (SEQ ID NO: 6)
RAIPIPAGTLLSGGGRAIYKRTAILG (SEQ ID NO: 11)
and
RFIIIPNIFTALSGGRALLYGATPYAIG (SEQ ID NO: 18) (NI in position 6 is Norleucine)
or salts thereof, particularly acetate salts.

In some embodiments the HIV specific peptides for use according to the invention are modified at the C-terminus as follows:

5  RALGPAATLQTPWATASLGV-GNH₂ (SEQ ID NO: 3)
   RWLLLGLNLVGGGRLYSPTLSILG-NH₂ (SEQ ID NO: 6)
   RAIPIPAGTLLSGGGRAIYKRTAILG-NH₂ (SEQ ID NO: 11)
   and
   RFIIPNFTALSGGGRRALLYGATPYAIG-NH₂ (SEQ ID NO: 18)

10  or salts thereof, particularly acetate salts. (In this application also referred to in the examples as Vacc-4x).

One of the sequences contains a B-cell epitope and will activate the humoral immune system, whereas the other sequences contribute with CTL-epitopes and the amino acid changes implemented within the frame of the CTL-epitope are designed to achieve enhanced binding. Other amino acid changes have been conducted in order to facilitate the synthesis of the peptide and/or increase the solubility of the peptide.

As described above some aspects of the present invention relates to methods for reducing and/or delaying pathological effects of human immunodeficiency virus I (HIV) or for reducing the risk of developing acquired immunodeficiency syndrome (AIDS) in a human subject infected with HIV, the method comprising the steps of:

a) a therapeutic HIV-1 immunization phase consisting of the administering in one or more doses of an effective amount of one or more HIV-specific peptide selected from the list consisting of the amino acid sequence shown in SEQ ID NO: 18 (Vacc-10), SEQ ID NO: 11 (Vacc-11), SEQ ID NO: 6 (Vacc-12), and SEQ ID NO: 3 (Vacc-13) over a period of 1-12 weeks;

b) one or more subsequent or simultaneous measurements of HIV-1 DNA levels in said human subject infected with HIV; and optionally

c) a subsequent viral reactivation phase consisting of the administering of an effective amount of a reservoir purging agent.

Another aspect relates to a method for monitoring the effect of a therapeutic HIV-1 immunization phase consisting of the administering in one or more doses of an effective amount of one or more HIV-specific peptide selected from the list consisting of the amino acid sequence shown in SEQ ID NO: 18 (Vacc-10), SEQ ID NO: 11 (Vacc-11), SEQ ID NO: 6 (Vacc-12), and SEQ ID NO: 3 (Vacc-13) over a period of 1-12 weeks; in reducing and/or delaying pathological effects of human immunodeficiency virus I (HIV) or in reducing the risk of developing acquired immunodeficiency syndrome (AIDS) in a human subject infected with HIV, the method comprising the step of one or more measurements of HIV-1 DNA levels in
said human subject infected with HIV-1 subsequent or simultaneous to said immunization phase.

In some embodiments, the subjects are being treated with a combination antiretroviral therapy (cART) prior to and/or during, and/or after said immunization phase, and/or said viral reactivation phase.

In some embodiments, the method further comprises a step b2) subsequent to step b) of selecting human subjects, wherein the level of HIV-1 DNA in said subjects is at least 1, such as at least 2, 3, 4, 5, 6, 7, 8, 9, 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 100, 200, 300, 400, 500, 600, 700, 800, 900, 1000, 2000, 3000, 4000, 5000, 6000, 7000, 8000, 9000, or 10000 HIV-1 DNA copy per million cell over a period of 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 100, 110, 120, 130, 140, 150, 160, 170, 180, 190, 200, 210, 220, 230, 240, 250, 260, 270, 280, 290, 300, 310, 320, 330, 340, 350, 360, or 370 weeks after the therapeutic HIV-1 immunization phase consisting of the administering in one or more doses under step a); and repeating step a) and/or b) and/or optionally step c) for said selected subjects.

In some embodiments, the method further comprises a step b2) subsequent to step b) of selecting human subjects, wherein the level of HIV-1 DNA in said subjects is at least 10%, such as at least 15%, 20%, 25%, 30%, 35%, 40%, 45%, 50%, 55%, 60%, 65%, 70%, 75%, 80%, 85%, 90%, or 95% of said level prior to said immunization phase as measured over a period of 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 100, 110, 120, 130, 140, 150, 160, 170, 180, 190, 200, 210, 220, 230, 240, 250, 260, 270, 280, 290, 300, 310, 320, 330, 340, 350, 360, or 370 weeks after said therapeutic HIV-1 immunization phase consisting of the administering in one or more doses under step a); and repeating step a) and/or b) and/or optionally step c) for said selected subjects.

In some embodiments, the method further comprises a step b2) subsequent to step b) of selecting human subjects, wherein the level of HIV-1 DNA in said subjects is less than 10000, such as less than 9000, 8000, 7000, 6000, 5000, 4000, 3000, 2000, 1000, 900, 800, 700, 600, 500, 400, 300, 200, 100, 95, 90, 85, 80, 75, 70, 65, 60, 55, 50, 45, 40, 35, 30, 25, 20, 15, 10, 5, 4, 3, 2, or 1 HIV-1 DNA copy per million cell over a period of 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 100, 110, 120, 130, 140, 150, 160, 170, 180, 190, 200, 210, 220, 230, 240, 250, 260, 270, 280, 290, 300, 310, 320, 330, 340, 350, 360, or 370 weeks after the therapeutic HIV-1 immunization phase consisting of the administering in one or more doses under step a); and treating said selected subjects under step c).

In some embodiments, the method further comprises a step b2) subsequent to step b) of selecting human subjects, wherein the level of HIV-1 DNA in said subjects is less than
95%, such as less than 90%, 85%, 80%, 75%, 70%, 65%, 60%, 55%, 50%, 45%, 40%, 35%, 30%, 25%, 20%, 15%, 10%, 5% of said level prior to said immunization phase as measured over a period of 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 100, 110, 120, 130, 140, 150, 160, 170, 180, 190, 200, 210, 220, 230, 240, 250, 260, 270, 280, 290, 300, 310, 320, 330, 340, 350, 360, or 370 weeks after said therapeutic HIV-1 immunization phase consisting of the administering in one or more doses under step a); and treating said selected subjects under step c).

In some embodiments, the method further comprises a step b2) subsequent to step

b) of selecting human subjects, wherein the level of HIV-1 DNA in said subjects is less than 10000, such as less than 9000, 8000, 7000, 6000, 5000, 4000, 3000, 2000, 1000, 900, 800, 700, 600, 500, 400, 300, 200, 100, 95, 90, 85, 80, 75, 70, 65, 60, 55, 50, 45, 40, 35, 30, 25, 20, 15, 10, 5, 4, 3, 2, or 1 HIV-1 DNA copy per million cell over a period of 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 100, 110, 120, 130, 140, 150, 160, 170, 180, 190, 200, 210, 220, 230, 240, 250, 260, 270, 280, 290, 300, 310, 320, 330, 340, 350, 360, or 370 weeks after the therapeutic HIV-1 immunization phase consisting of the administering in one or more doses under step a); and repeating step a) and/or b) and/or optionally step c) for said selected subjects.

In some embodiments, the method further comprises a step b2) subsequent to step

b) of selecting human subjects, wherein the level of HIV-1 DNA in said subjects is less than 95%, such as less than 90%, 85%, 80%, 75%, 70%, 65%, 60%, 55%, 50%, 45%, 40%, 35%, 30%, 25%, 20%, 15%, 10%, 5% of said level prior to said immunization phase as measured over a period of 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 100, 110, 120, 130, 140, 150, 160, 170, 180, 190, 200, 210, 220, 230, 240, 250, 260, 270, 280, 290, 300, 310, 320, 330, 340, 350, 360, or 370 weeks after said therapeutic HIV-1 immunization phase consisting of the administering in one or more doses under step a); and repeating step a) and/or b) and/or optionally step c) for said selected subjects.

In some embodiments, the method further comprises a step b2) subsequent to step

b) of selecting human subjects, wherein the level of HIV-1 DNA in said subjects decreases by more than 10%, such as at least 15%, 20%, 25%, 30%, 35%, 40%, 45%, 50%, 55%, 60%, 65%, 70%, 75%, 80%, 85%, 90%, or 95% of said level prior to said immunization phase as measured over a period of 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 100, 110, 120, 130, 140, 150, 160, 170, 180, 190, 200, 210, 220, 230, 240, 250, 260, 270, 280, 290, 300, 310, 320, 330, 340, 350, 360, or 370 weeks after said therapeutic HIV-1 immunization phase consisting of the administering in one or more doses under step a); and repeating step a) and/or b) and/or optionally step c) for said selected subjects.
In some embodiments, the method further comprises a step b) subsequent to step b) of selecting human subjects, wherein the level of HIV-1 DNA in said subjects decreases less than 10%, such as less than 9%, 8%, 7%, 6%, 5%, 4%, 3%, 2%, 1% of said level prior to said immunization phase as measured over a period of 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 100, 110, 120, 130, 140, 150, 160, 170, 180, 190, 200, 210, 220, 230, 240, 250, 260, 270, 280, 290, 300, 310, 320, 330, 340, 350, 360, or 370 weeks after said therapeutic HIV-1 immunization phase consisting of the administering in one or more doses under step a); and treating said selected subjects under step c).

In some embodiments, the method further comprises a step a-1) preceding step a) of measurement of HIV-1 DNA levels in said human subject infected with HIV.

In some embodiments, the method comprises in step a) the administering of two, three, four, five or more doses of an effective amount of one or more HIV-specific peptide selected from the list consisting of the amino acid sequence shown in SEQ ID NO: 18 (Vacc-10), SEQ ID NO: 11 (Vacc-11), SEQ ID NO: 6 (Vacc-12), and SEQ ID NO: 3 (Vacc-13) over a period of 1-12 weeks.

In some embodiments, an adjuvant, such as recombinant human granulocyte-macrophage colony-stimulating factor (rhuGM-CSF) or a water-in-oil adjuvant such as ISA51 or ISA720, or an oil-in-water adjuvant such as Provax, is administered in conjunction to, prior to or simultaneously with said therapeutic HIV-1 immunization.

In some embodiments, the reservoir purging agent is administered over a period of 1, 2, 3, or 4 consecutive weeks at least about 1, 2, 3, or 4 weeks after said therapeutic HIV-1 immunization phase.

In some embodiments, the viral reactivation phase includes the administration of 1-10 doses, such as 2-10 doses, such as 3-10, such as 4-10, such as 5-10, such as 6-10, such as 7-10, such as 8-10, such as 9-10, such as 10 doses, or 1-9 doses, such as 1-8 doses, such as 1-7, such as 1-6, such as 1-5, such as 1-4, such as 1-3, such as 3 doses of an effective amount of a reservoir purging agent.

In some embodiments, step a) and/or b) are independently repeated 1, 2, 3, 4, 5, 6, 7, 8, 9, or 10 times in any order.

In some embodiments, the reservoir purging agent is an HDAC inhibitor, such as romidepsin or panobinostat.

In some embodiments, the reservoir purging agent is romidepsin administered by infusions at a dosing of up to 2.5 mg/m2, such as up to 5 mg/m2, such as up to 7.5 mg/m2, such as up to 10 mg/m2, such as up to 12 mg/m2, such as up to 12.5 mg/m2, such as up to 14 mg/m2, such as between 2.5 mg/m2 and 7.5 mg/m2, such as around 5 mg/m2.

In some embodiments, the effect on the HIV-1 latent reservoir is in HIV-infected patients virologically suppressed on cART.
In some embodiments, each peptide is given in a dose of 0.1 mg-10 mg per administration, such as 0.1-10 mg per administration, such as 0.1-9 mg per administration, such as 0.1-8 mg per administration, such as 0.1-7 mg per administration, such as 0.1-6 mg per administration, such as 0.1-5 mg per administration, such as 0.1-4 mg per administration, such as 0.1-3 mg per administration, such as 0.1-2 mg per administration, such as 0.1-1.2 mg per administration, such as 0.1-0.9 mg per administration, such as 0.1-0.6 mg per administration, such as 0.1-0.4 mg per administration.

In some embodiments, the therapeutic HIV-1 immunization phase is over a period of 1-12 weeks, such as over a period of 2-12 weeks, such as over a period of 3-12 weeks, such as over a period of 4-12 weeks, such as over a period of 5-12 weeks, such as over a period of 6-12 weeks, such as over a period of 7-12 weeks, such as over a period of 8-12 weeks.

In some embodiments, the therapeutic HIV-1 immunization phase includes the administration of 1-10 doses, such as 2-10 doses, such as 3-10, such as 4-10, such as 5-10, such as 6-10, such as 7-10, such as 8-10, such as 9-10, such as 10 doses.

In some embodiments, the one or more peptide is in the form of an acetate salt.

In some embodiments, the acetate content of the salt is between 4% and 18%, such as between 5% and 17%, such as between 6% and 16%, such as between 7% and 15%, such as between 8% and 14%, such as between 9% and 14%, such as between 9% and 13%, such as between 10% and 14%, such as between 11% and 14%, or between 5% and 16%, such as between 5% and 15%, such as between 5% and 14%, such as between 6% and 14%, such as between 6% and 13%, such as between 7% and 12%, such as between 7% and 11%, such as between 8% and 11%, such as between 9% and 11%, or between 3% and 18%, such as between 3% and 17%, such as between 3% and 16%, such as between 3% and 15%, such as between 3% and 14%, such as between 3% and 13%, such as between 3% and 11%, such as between 3% and 10%, such as between 4% and 10%, such as between 4% and 9%, such as between 4% and 8%, such as between 4% and 7%, such as between 4% and 6%, such as between 4% and 5%.

In some embodiments, one, two, three or four peptides are used in the therapeutic HIV-1 immunization phase.

In some embodiments, all four peptide as acetate salts are used in the therapeutic HIV-1 immunization phase.

In some embodiments, the peptides have amide C-terminal ends of formula -C(O)NH2, or acetate salts thereof.

In some embodiments, all four peptide are used in the ratio of 1:1:1:1 w/w.

In some embodiments, the one, two, three or four peptides are in a dissolved liquid state.

In some embodiments, the liquid is water.

In some embodiments, the method further comprises the administering of one or more further therapeutically active agent selected from an immunomodulatory compound
and a second reservoir purging agent, such as a histone deacetylase (HDAC) inhibitor, or BET family protein inhibitors/antagonists, such as JQ1, I-BET, I-Bet151, MS417, GW841819X, and thienotriazolodiazepine compounds, such as those described in U.S. Patent Application Publication No. 2010/0286127.

In some embodiments, the immunomodulatory compound is selected from anti-PD1 antibodies, such as MDX-1106 (Merck), THALOMID® (thalidomide), anti-PD1 antibodies, cyclophosphamide, Levamisole, lenalidomide, CC-4047 (pomalidomide), CC-11006 (Celgene), and CC-10015 (Celgene), and immunomodulatory compounds described in any one ofWO2007028047, WO2002059106, and WO2002094180.

In some embodiments, the immunomodulatory compound is lenalidomide.

In some embodiments, the reservoir purging agent is selected from M344 (4-(dimethylamino)-N-[7-(hydroxyamino)-7-oxoheptyl]benzamide), chidamide (CS055/HBI-800), 4SC-202, (4SC), Resminostat (4SC), hydroxamic acids such as vorinostat (SAHA), belinostat (PXD101), LAQ824, trichostatin A and panobinostat (LBH589); benzamides such as entinostat (MS-275), CI994, and mocetinostat (MGCD0103), cyclic tetrapeptides (such as trapoxin, such as trapoxin B), and the depsipeptides, such as romidepsin (ISTODAX), electrophilic ketones, and the aliphatic acid compounds such as phenylbutyrate, valproic acid, Oxamflatin, ITF2357 (generic givinostat), Apicidin, MC1293, CG05, and CG06; compounds that activate transcription factors including NF-KappaB, Prostratin, auranofin, bryostatin, a nontumorogenic phorbolester, DPP (12-deoxyphorbol-13-phenylacetate), PMA, and Phorbol 12-myristate 13-acetate (PMA); Compounds that activate HIV mRNA elongation including P-TEF-b kinase and hexamethylbisacetamide (HMBA); IL-7; T-cell stimulating factors including anti-CD3/CD28 - T-cell stimulating Ab's; Kinase inhibitors including Tyroptostin A, Tyrophostin B, and Tyrophostin C;PTEN (phosphatase and tensin homologue) gene inhibitors including SF1670 (Echelon Bioscience), Disulfiram (DSF), an inhibitor of acetaldehyde dehydrogenase, Protein Tyrosine Phosphatase Inhibitors including bpV(HOpic), bpV(phen), and bpV(pic) (Calbiochem; EMD Millipore), Toll-like receptors agonists including Toll-like receptor-9 (TLR9) and Toll-like receptor-7 (TLR9) agonists, quercetin, lipoic acid, sodium butyrate, TNF-alpha, PHA and Tat.

DESCRIPTION OF THE PREPARATION OF THE PEPTIDES

The peptides of the invention can be produced by any known method of producing a linear amino acid sequence, such as recombinant DNA techniques. A nucleic acid sequence which encodes a peptide of the invention or a multimer of the said peptides, is introduced into an expression vector. Suitable expression vectors are for instance plasmids, cosmids, viruses and YAC (yeast artificial chromosome) which comprise necessary control regions for

In case a linked or cyclic peptide is desired, the amino acid sequence is subjected to a chemical oxidation step in order to cyclize or link the two cysteine residues within one or between two peptide sequences, when the appropriate linear amino acid sequences are synthesized, see Akaji et al., Tetrahedron Letter, 33, 8, p.1073-1076, 1992.

GENERAL DESCRIPTION OF SYNTHESIS
The amino acid derivatives were supplied by Bachem AG, Switzerland.
The peptides described herein preferably have a free amino group at the N-terminus and an amidated C-terminus. The counter ion of all peptides described herein is acetate which is bound in ionic form to charged functional groups (i.e. guanidino side chains arginine and the ε-amino groups of lysine [Vacc-11] and the side chains of arginine [Vacc-10, Vacc-12 and Vacc-13]). All amino acid residues except the achiral glycine are in the L-configuration.
The peptides described herein were assembled on tricyclic amide linker resins utilising an 9-fluorenylmethoxyxycarbonyl (Fmoc) strategy.

In brief the tricyclic amide linker resin is transferred into a solid phase peptide synthesis (SPPS)-reactor with a stirrer. Synthesis is then started with a 9-fluorenylmethoxycarbonyl (Fmoc)-deprotection of the resin according to the general description given below, followed by a coupling procedure with Fmoc-Gly-OH. This step is again followed by an Fmoc-deprotection and subsequent coupling of the amino acid derivates, peptides or dipeptides according to the sequence. The last coupling step is performed with side-chain protected Fmoc-Arg-OH. After final Fmoc-deprotection, the peptide resin is dried in a desiccator under reduced pressure.

Fmoc-deprotecting procedure:

Step 1: Washing;
Step 2: Fmoc-deprotection;
Steps 3-9: Washing.
Each step consists of addition of solvents/reagents, stirring at room temperature and filtration.

The peptide resin is treated with cold TFA in the presence of deionised water and 1, 2-Ethanediithiol (EDT), (Vacc-10 and Vacc-13) or triisopropylsilane (TIS) (Vacc-11 and Vacc-12) for approximately two to three hours at room temperature. After filtering off and washing the resin with TFA, the peptide is precipitated in diisopropyl ether (IPE). It is then filtered off, washed with IPE and dried in a desiccator under reduced pressure.

The material obtained in the previous stage is purified by preparative HPLC on reversed phase columns with acetonitrile (ACN) gradient elution and ultraviolet (UV) detection at $\lambda = 220$ nano-metres (nm) using a TEAP and/or TFA system. Vacc-10 is only purified using the TFA system.

For Vacc-13, a perchlorate system for preparative HPLC purification prior to using TEAP and TFA system has been introduced. Sodium perchlorate is listed as a raw material.

The last stage of manufacture of Vacc-4x acetate is the exchange from the TFA salt, obtained in stage three, into the acetate salt by ion exchange. The lyophilised material from one or several combined preparative HPLC runs is dissolved in varying concentrations of acetic acid or in purified water according to the properties of the individual peptides. The dissolved peptide is loaded onto the ion exchange resin (acetate form) and equilibrated with 5% acetic acid (or 20% purified water for Vacc-13). The elution is performed with 5% acetic acid (or purified water for Vacc-13), checked by thin-layer chromatography (TLC), filtered through a 0.2 µm membrane filter and lyophilised to yield the final product as a white to off-white powder.

Although the Vacc-4x formulation does not contain any ionic excipients, the peptides and their counter ions (acetate) account for a certain osmolality. The range of 10 - 100 mOsm/kg was defined based on the result obtained for the technical sample. Potential variability due to the four peptides is taken into account. For the drug product, approximately 1 mg of each of the four Vacc-4x peptides was used. The lyophilisate is reconstituted with 0.30 mL of WFI. Taking the acetic acid contents of the peptides listed in table 1 into account, the acetic acid content of Vacc-4x is approximately 0.40 mg in 0.30 mL of solution. The theoretical osmolality is approximately 23 mOsmol/L by calculation, which correlates well with the values determined in the Vacc-4x batches (20-23 mOsmol/kg).
Table 1  Acetic acid contents of the four peptides (GMP grade material, two batches each)

<table>
<thead>
<tr>
<th>Active substance</th>
<th>Peptide batch used for Vacc-4x batches 1011584 and 1012951</th>
<th>Acetic acid content [%]</th>
<th>Peptide batch used for Vacc-4x batch 1018724</th>
<th>Acetic acid content [%]</th>
</tr>
</thead>
<tbody>
<tr>
<td>Vacc-10 Acetate</td>
<td>1008290</td>
<td>11.3</td>
<td>1015501</td>
<td>12.2</td>
</tr>
<tr>
<td>Vacc-11 Acetate</td>
<td>1009945</td>
<td>17.2</td>
<td>1015502</td>
<td>14.8</td>
</tr>
<tr>
<td>Vacc-12 Acetate</td>
<td>1008294</td>
<td>9.9</td>
<td>1015503</td>
<td>10.0</td>
</tr>
<tr>
<td>Vacc-13 Acetate</td>
<td>1008296</td>
<td>4.6</td>
<td>1015504</td>
<td>5.1</td>
</tr>
</tbody>
</table>

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EXAMPLE 1
Preparation of K A L G P G A T L Q T P W T A C Q G V G - NH₂ (SEQ ID NO: 2).

The peptide was synthesized in amide form, from corresponding starting materials according to the general description of synthesis. The purity was determined by HPLC analysis and the structure was confirmed by amino acid analysis and mass spectrometry (LDI-MS).

Preparation of R A L G P A A T L Q T P W T A S L G V G (SEQ ID NO: 3).

The peptide was synthesized in amide form, from corresponding starting materials according to the general description of synthesis. The purity was determined by HPLC analysis and the structure was confirmed by amino acid analysis and mass spectrometry (LDI-MS).

Molecular formula: \( C_{88}H_{144}O_{25}N_{26} \)


The peptide was synthesized in amide form, from the corresponding starting materials according to the general description of synthesis. The purity was determined by HPLC analysis and the structure was confirmed by amino acid analysis and mass spectrometry (LDI-MS).

Mass spectral analysis: Theoretical molecular weight: 2454.9
Experimental molecular weight: 2454.8 ES+

The peptide was synthesized in amide form, from the corresponding starting materials according to the general description of synthesis. The purity was determined by HPLC analysis and the structure was confirmed by amino acid analysis and mass spectrometry (LDI-MS).

Molecular weight (free base) : 2552
Molecular formula : \( C_{119}H_{195}O_{29}N_{33} \)

Preparation of \( \text{KILLLGLNPLLGVGGGRLYSPTSILG} \) (SEQ ID NO: 7), \( \text{RLLLGLNPLLGVGGGRLYSPTTILG} \) (SEQ ID NO: 8) and \( \text{NIPIPVGDYGGGDYKRWQALCL} \) (SEQ ID NO: 21).

The peptides are synthesized in amide form, from the corresponding starting materials according to the general description of synthesis. The purity are determined by HPLC analysis and the structures are confirmed by amino acid analysis and mass spectrometry (LDI-MS).

Preparation of \( \text{RNIPVGDYGGGDYKRWQALCL} \) (SEQ ID NO: 10).

The peptide was synthesized in amide form, from the corresponding starting materials according to the general description of synthesis. The purity was determined by HPLC analysis and the structure was confirmed by amino acid analysis and mass spectrometry (LDI-MS).

Mass spectral analysis : Theoretical molecular weight : 2817.3
Experimental molecular weight : 2813.7 ES+

Preparation of \( \text{RAIPAGTLLSGGGRAIYKRWAILG} \) (SEQ ID NO: 11).

The peptide was synthesized in amide form, from the corresponding starting materials according to the general description of synthesis. The purity was determined by HPLC analysis and the structure was confirmed by amino acid analysis and mass spectrometry (LDI-MS).

Molecular weight (free base) : 2707
Molecular formula : \( C_{123}H_{206}O_{29}N_{36} \)

Preparation of \( \text{ALPIPAFGIYGGGRYKRWQALG} \) (SEQ ID NO: 12), \( \text{KIPIPVGF} \)

\( \text{IGGGWYKRWAILE} \) (SEQ ID NO: 13) and \( \text{KIPIPVGTLLSGGGRIYKRA} \)

\( \text{ILG} \) (SEQ ID NO : 14). The peptides are synthesized in amide form, from the corresponding starting materials according to the general description of synthesis. The purity are determined by HPLC analysis and the structures are confirmed by amino acid analysis and mass spectrometry (LDI-MS).

Preparation of \( \text{KFIFPNIFSALGGAIASYDLN} \)

\( \text{INCN} \) (SEQ ID NO : 16).
The peptide was synthesized in amide form, from the corresponding starting materials according to the general description of synthesis. Nl in the sequence is Norleucine. The purity was determined by HPLC analysis and the structure was confirmed by amino acid analysis and mass spectrometry (LDI-MS).

Mass spectral analysis : Theoretical molecular weight : 2783.3
Experimental molecular weight : 2783.3 ES+


The peptide was synthesized in amide form, from the corresponding starting materials according to the general description of synthesis. Nl in the sequence is Norleucine. The purity was determined by HPLC analysis and the structure was confirmed by amino acid analysis and mass spectrometry (LDI-MS).

Mass spectral analysis : Theoretical molecular weight : 2932.4
Experimental molecular weight : 2931.8 ES+


The peptide was synthesized in amide form, from the corresponding starting materials according to the general description of synthesis. Nl in the sequence is Norleucine. The purity was determined by HPLC analysis and the structure was confirmed by amino acid analysis and mass spectrometry (LDI-MS).

Molecular weight (free base) : 2894
Molecular formula : C_{137}H_{217}O_{32}N_{57}

Preparation of K I I P N I F S A L G G G R L L Y G A T P Y A I G (SEQ ID NO: 19), R I I P N I F T A L S G G G R L L Y G A T P Y A I G (SEQ ID NO: 20) and W I I P N I F S A L G G A I S Y D L N T N I L N C I (SEQ ID NO : 22). The peptides are synthesized in amide form, from the corresponding starting materials according to the general description of synthesis. The purity are determined by HPLC analysis and the structures are confirmed by amino acid analysis and mass spectrometry (LDI-MS).

EXAMPLE 2

A vaccine comprising the peptides of the SEQ ID NOs: 3, 6, 11 and 18 was prepared (also refered to herein as Vacc-4x). The freeze-dried peptides were dissolved in sterile water at a final concentration of 4 mg/ml. The final salt concentration was 0.9 %. A preparation of a granulocyte-macrophage-colony stimulating factor (GM-CSF) was also prepared, according to the manufacturer’s directions for use, to a final concentration of 0.3 mg/ml. The two solutions are administered intracutaneously. A typical injection dose is 100 μl.

EXAMPLE 3
An antigen solution or suspension is mixed with equal parts of Freund's adjuvant of Behring, complete or incomplete, and is then finely emulsified by being drawn up into, and vigorously pressed out of, an injection syringe, or with a homogenisator. The emulsion should remain stable for at least 30 minutes. The antigen-adjuvant emulsion is best injected subcutaneously as a depot.

EXAMPLE 4

Toxicity studies were performed in mice and rats on the peptide composition of the vaccine in Example 2. The mouse was selected for the study to provide comparative data from a second commonly used rodent species. The test substance was a mixture of four peptides supplied as one vial containing lyophilised material for reconstitution with physiological saline, and dose levels were expressed in terms of total peptide load. The individual peptides were present in ratio 1:1:1:1 w/w giving dose levels of each peptide of 0.0075 mg/kg body weight, 0.075 mg/kg body weight and 0.75 mg/kg body weight, which are up to 500 fold the intended human dose. The test animals were divided into four groups of ten animals each (five males and five females); a saline control group and groups for low, intermediate and high doses. The test composition was administered once, by intravenous infusion into a tail vein at a dose rate of 3 ml/minute. The animals were killed at day 15 and 16 by intraperitoneal injection of sodium pentobarbitone.

The results of these studies indicated that the dose levels administered to the mice and rats elicited no adverse reactions and that the no effect level was in excess of 3 mg/kg.

EXAMPLE 5

Immunoassay for detection of antibodies induced by HIV-1.

The magnetic particle reagents are to be prepared according to the manufacturers recommended protocol. Dynal AS, is the manufacturer of the Dynabeads, which are employed. The magnetic particles coated with ligand are called Reagent 1. A peptide according to the invention is covalently coupled to the pre-activated surface of the magnetic particles. It is also possible to physically absorb the peptide to the surface of the magnetic particles. The concentration of particles in Reagent 1 is within the range from 1 mg/ml to 15 mg/ml. The particle size varies between 0.2 μm to 15 μm. The concentration of peptides is within the range from 0.01 mg/mg particle to 1 mg/mg particle.

The anti-human Ig Alkaline Phosphatase (AP) conjugated antibody reagent is prepared according to the recommended protocol of Dako AS. This protocol is a standard procedure in this field. This reagent is called Reagent 2.

The substrate solution phenolphthalein-monophosphate is to be prepared according to the recommended protocol of Fluka AG. This protocol is a standard procedure in this field. The substrate solution is called Reagent 3.
The washing and incubation buffer which is used is standard 0.05M tris-base buffer with the following additional compounds: Tween 20 (0.01% to 0.1%), glycerol (0.1% to 10%) and sodium chloride (0.2% to 0.1%).

The assay procedure comprises an incubation step wherein 1 drop of Reagent 1 is mixed with 2 drops of washing buffer in each well. After mixing, 30 µl of sample is added and the solution is incubated for 5 minutes. The magnetic particles can be trapped by a magnet and the liquid removed, before the magnet is separated. Then the wells are washed twice in 4 drops of washing solution, before incubation with Reagent 2. 1 drop of Reagent 2 is added with 2 drops of washing buffer and the solution is incubated for 5 minutes. The magnetic particles can be trapped by a magnet and the liquid removed, before the magnet is separated. Then the washing step is repeated before incubation with Reagent 3. 2 drops of Reagent 3 is added to each well and the solution is incubated for 3 minutes. The results can be read against a white background. Positive results are red (3+ = strong red) whereas negative results are clearly light yellow/brown solutions as obtained in the negative control.

The immunoassay kit could be used in detection of antibodies, induced either by HIV virus or HIV-specific peptides or proteins, for instance the peptides of the present invention.

The above Examples are only meant as illustrating the invention. It must be understood that a person skilled in the art can modify the peptides, antigens and vaccines herein described without deviating from the concept and scope of this invention as set forth in the claims.

The polypeptides of the invention can be used in a combination of at least one peptide selected from each group of sequences, SEQ ID NOs: 1, 4, 9 and 15 to form antigens and the active principle of a prophylactic or therapeutic vaccine intended to provide protection against the human immunodeficiency virus type 1 (HIV-1). The vaccine may include compounds having beneficial effects in protecting or stimulating the host’s immune system (human being or vertebrate animal) for instance interleukins, interferons, granulocyte macrophage growth factors, haematopoietic growth factors or similar. Preferably the vaccine composition further contain an adjuvant or vehicle, more preferably the adjuvant or vehicle is Monophosphoryl Lipid A (MPL ©) possibly with alum, Freund’s adjuvant (complete or incomplete) or aluminum hydroxide. The optimal amount of adjuvant/vehicle will depend on the type(s) which is chosen.

The peptide or vaccine formulation can be freeze-dried prior to storage. The vaccine may be stored preferably at low temperature, in ampoules containing one or more dosage units, ready for use. Persons skilled in the art will appreciate that a suitable dose may depend on the body weight of the patient, the type of disease, severity of condition, administration route and several other factors. The vaccine might be administered up to twelve times and through injection, typically it will be administered about six times. In preparation of an injection solution the peptides are dissolved in sterile water or sodium chloride solution at a final concentration of 1-3 mg/ml per peptide and 0-0.9% sodium chloride. Typically an
injection volume is 100 µl to 200 µl (2 x 100 µl). The peptide is preferably co-administered with a suitable adjuvant and/or a granulocyte-macrophage growth factor for instance Leucomax® «Shering Plough». Suitable administration may be intracutane, subcutane, intravenous, peroral, intramuscular, intranasal, mucosal or any other suitable route. Booster administrations may be required in order to maintain protection.

EXAMPLE 6

The anti-HIV p24 immune response resulting from Vacc-4x immunization could in combination with ART potentially improve immune reconstitution in patients who have not fully regained a healthy CD4 level (>600x10^6/L). Potential benefits of Vacc-4x in subjects with incomplete immune reconstitution include a possible sustained improvement in the immune response to p24 and HIV.

Potential risks include the discomfort and inconvenience associated with the immunizations and the risk of known or unknown side effects of exposure to Vacc-4x and Leukine (rhu-GM-CSF) including, most commonly, local reactions at the site of injections and fatigue (likelihood not yet determined).

The results of non-clinical single-dose studies in mice and rats indicate that the dose levels of intravenous Vacc-4x elicited no adverse reactions and that the no effect level was in excess of 3 mg/kg, which constitutes a 500 fold safety margin over the planned human dose level.

In a rabbit study the effect of Vacc-4x was evaluated in the presence of concomitant GM-CSF, the adjuvant used in the clinical program. Local intradermal reactions such as erythema and edema were noted, however, similar effects were noted in control animals both macroscopically and histological. These local reactions were slightly more pronounced in the Vacc-4x treated animals. There were no systemic reactions in this study. These data indicate that Vacc-4x has no limiting toxicology in a model that is relevant to the proposed clinical study.

The therapeutic vaccine candidate Vacc-4x, has been studies in a Phase I and three Phase II clinical studies. The Phase I study enrolled 11 HIV-positive subjects, including nine subjects on ART. Subjects were maintained on ART (if entered on ART); all subjects were treated with 12 immunizations of Vacc-4x at a dose of 0.4 mg/injection over a period of 26 weeks. Immunizations were performed following injection of rhu-GM-CSF (Leucomax® [molgramostim]) as adjuvant. All subjects experienced one or more adverse events (AEs); nine subjects experienced events judged related to treatment. The adverse reactions reported were mild or moderate in severity except for severe local reactions in one subject. No subjects were withdrawn due to treatment-related AEs or toxicological reactions; no serious adverse events (SAEs) occurred. Treatment related events observed in more than one subject included painful injection (seven subjects), fatigue-vertigo (four subjects), influenza-like symptoms (two subjects), and irritated skin at injection site (two subjects).
All subjects experienced a cell-mediated immune response, measured by delayed-type hypersensitivity (DTH) skin reaction. Some cell-mediated immune response, measured by γ IFN release using enzyme-linked immunosorbent spot assay (ELISPOT), was reported for 45% of the subjects; no antibody response to Vacc-4x peptides was observed.

The Phase II dose-finding study (CTN B-HIV 2/2001) enrolled 40 HIV positive subjects, of which 38 completed the trial. Subjects were maintained on ART and treated with 10 immunizations at a dose of 0.4 mg (20 subjects) or 1.2 mg (20 subjects) per Vacc 4x injection, over a period of 26 weeks. Immunizations with Vacc 4x were performed following injection of rhu-GM-CSF (Leucorex [molgramostim]) as a local adjuvant. ART was interrupted from Week 26 to Week 30 to allow exposure to the subject’s own virus (autologous immunization). ART was resumed from Week 30 to Week 38 to allow maturation of immune responses to the Vacc 4x peptides and to the subject’s own virus. ART was discontinued from Week 38 to Week 52 when the study was formally concluded. Treatment-related AEs were observed in 20 subjects (8 subjects in the 0.4mg group and 12 subjects in the 1.2mg group). No SAEs were reported during the period of immunization. One subject experienced a transient vasovagal reaction in conjunction with immunization and the DTH test at Week 26 and Week 38. A second subject experienced a vasovagal reaction in conjunction with the DTH test at Week 52. For the laboratory parameters, vital signs, and performance status, no changes attributable to immunization were observed. Changes in HIV RNA, CD4 cell counts, and CD8 cell counts showed no safety concerns related to immunization.

Immunological responses reported as DTH positive reactions were observed for all subjects. Overall, positive responses both for induration and erythema were statistically significantly higher in the high dose (HD, 1.2mg Vacc-4x) group compared to the low dose (LD, 0.4 mg Vacc 4x) group. The dose-dependent differences in DTH reactions were maintained throughout the study. T-cell proliferation appeared stable after Week 12 and demonstrated an HD advantage, consistent with the DTH results. ART was interrupted at Week 38 with planned restart when CD4 counts fell to less than 200/μL or when AIDS- or HIV related events were observed (i.e. clinical practice). DTH responses to Vacc-4x (high versus low response determined at Week 38) were associated with reduced viral loads and correspondingly improved CD4 counts at the end of the study (Week 52).

During the immunization period, CD4 counts were stable or increased. Interruption of ART resulted in reduction of CD4 counts. However, 14 weeks after the last interruption of ART (Week 52), the mean CD4 counts were still above 200 x 106 cells/L. No difference between the LD and the HD groups was observed. The majority of subjects remained off ART following completion of the study (Week 52); permission was given to follow the subjects until they resumed ART. The duration of treatment interruption was linked to immune responsiveness to the peptides. When subjects were compared to similar subjects in the Netherlands that had stopped treatment without Vacc-4x administration, a significantly
slower decline in CD4 cells was noted for the Vacc-4x subjects. The median treatment interruption achieved for all the subjects that participated in the Vacc-4x Phase II clinical study was 31 months.

CTN BI Vacc-4x/2009/1 was an open-label follow-up of study CTN B-HIV-2/2001 to determine whether a re-boost with Vacc-4x could reactivate or increase the immune response obtained during the immunization performed in the CTN B-HIV-2/2001 study. The secondary objectives were to evaluate: the in vivo immunogenicity of Vacc-4x by evaluation of DTH and to compare the DTH response to DTH in the initial study; the effect of Vacc-4x on CD4 counts, CD8 counts and HIV viral RNA; and the safety and tolerability of Vacc-4x. All 26 subjects included in the study received two booster administrations of Vacc-4x.

A total of 74 AEs were reported by 23 subjects. Most adverse events (n=60) were scored as possibly/probably related to the study treatment. The majority (98%) of the related adverse events were mild. Two adverse events related to study treatment, one headache and one injection site indurations, were scored as moderate intensity. Itching (injection site pruritus) was the most frequent reported adverse event related to the study treatment. Nineteen patients (73%) reported this adverse event at least once. Ten of these patients reported itching related to both immunizations, while for the other nine patients it was only reported once. Five patients reported swelling related to the immunization. For three of these patients swelling was reported after both immunizations. No patient died during the study. No patient reported serious adverse events and no clinically relevant changes were recorded.

The study demonstrated that Vacc-4x peptides induced T cell responses lasting up to seven years. By re-boosting it was possible to increase killing markers, this again indicates that T cells had increased their potential to kill HIV-infected cells. Before re-boosting, all the patients had returned to CD4, CD8 and viral load levels that were similar to those before ART was stopped in the main study. Re-boosting had no negative effect on the CD4, CD8 and viral load of the patients. No safety concern was reported as a result of the re-boost of these patients.

The Phase II Study CT-BI Vacc-4x 2007/1 (EudraCT Number 2007-006302-13) was performed in US and Europe (UK, Germany, Spain and Italy). The study was a randomized, double-blind, multicenter, immunogenicity study of Vacc-4x versus placebo in patients infected with HIV-1 who have maintained an adequate response to ART. The primary objective was to evaluate the effect of Vacc-4x immunizations versus placebo on CD4 counts, T-cell function (ELISPOT, T-cell proliferative responses and intracellular cytokine staining) and the response to interruption of ART. The necessity to resume ART between the interruption of ART at Week 28 and the end of the study at Week 52, due to decreased CD4 count or increased viral loads, was monitored as one of the primary efficacy endpoints.

In the ITT analysis population, it was concluded that Vacc-4x did not reduce the proportion of subjects requiring resumption of ART after ART cessation at Week 28 in
comparison with placebo. There was also no effect compared with placebo on the percentage change in CD4 count between Week 28 and the last CD4 assessment before resumption of ART. The time to restarting ART was similar in Vacc-4x and placebo-treated subjects.

The viral load results after ART cessation varied between subjects with evidence of favourable effects of Vacc-4x immunization over placebo. There were no significant differences in the repeated measures ANOVA for viral load over Weeks 4 to 52 when data included all evaluable subjects, irrespective of whether they were or were not taking ART. In the subgroup of subjects who remained off ART until Week 52, the average viral load was lower in the Vacc-4x-treated subjects than the placebo group. A post-hoc analysis showed the Week 52 (Last Observation Carried Forward [LOCF]) viral load to be statistically significantly lower in the Vacc-4x group than the placebo group.

The analysis of change in HIV-1 RNA from Week 28 through to Week 52 revealed a statistically significant difference between groups in favour of Vacc-4x. The AUC in those who remained off ART at Week 52 was lower in the Vacc-4x group than in the placebo group. A post-hoc analysis showed this difference in AUC to be statistically significant.

No safety concern was raised during this study. The study was supervised by a Data Safety Monitoring Board (DSMB).

EXAMPLE 7

Test of peptides together with IMiDs for increased proliferation, polyfunctionality, IL-2 secretion and IFN-γ production.

Expansion of polyfunctional HIV-specific T-cells upon stimulation with Dendritic Cells, pre-incubated with peptides to be used according to the invention, may be studied by methods described by Keersmaecker et al. (J. Virol., 2012 86:9351-9360) and referenced therein, HIV proteins Gag or Nef, they are incubated with peptides to be used according to the invention, before they are used to stimulate T-cells in a co-culture.

Keersmaecker et al. found that the presence of IMiDs (Lenalidomide (IMiD3; CC-5013) and pomalidomide (IMiD1; CC-4047) during in vitro T-cell stimulation with dendritic cells presenting Gag- or Nef-specific peptides, resulted in a number of improvements in the function of the T-cells. Among these were; polyfunctional HIV specific CD8+ T cells with enhanced lytic capacity, more Gag antigen epitopes recognized and at lower antigen peptide concentrations, reduced proliferation of CD4+ T cells with increased number of polyfunctional CD4+ T-cells, increased IL-2 production by CD8+ T-cells, detectable IFN-γ production by CD8+ T-cells and CD4 T-cells after antigen stimulation.

"Expansion of Polyfunctional HIV-Specific T Cells upon Stimulation with mRNA Electroporated Dendritic Cells in the Presence of Immunomodulatory Drugs"

Brenda De Keersmaecker, Sabine D. Allard, Patrick Lacor, Rik Schots, Kris Thielemans, and Joeri L. Aerts

EXAMPLE 8

Suggested clinical study protocol for the test of Peptide composition comprising 4 peptides in combination with Lenalidomide and HDAC inhibitor

Immunizations (four primary immunizations and two booster immunizations) at Weeks 1, 2, 3 and 4, and booster immunizations at Weeks 12 and 13 with either:

1) Peptide composition with GM-CSF as adjuvant and Lenalidomide (CC-5013), or

2) Peptide composition with GM-CSF as adjuvant and Placebo for Lenalidomide (CC-5013).

3) Placebo

Suggested doses:

Peptide composition: 0.6, 0.9, 1.2 and 1.5 mg (Equimolar amount of each peptide)
Lenalidomide: 5,10, and 25 mg.
Subjects randomized to the Lenalidomide (CC-5013) arm will take a single oral dose of Lenalidomide (CC-5013) daily the two preceding days before immunization with the Peptide composition and on the day of each immunization.

The Peptide composition used according to this clinical trial setup consists of SEQ ID NO:3, SEQ ID NO:6, SEQ ID NO:11, and SEQ ID NO:18.

At week 20 subjects in all study arms will receive 20 mg panobinostat (LBH589) orally on days 1, 3, and 5 (i.e. 3 times a week) every other week for a period of 8 weeks (up to week 28) while maintaining background ART. This will be followed by a 24 week follow up period (up to week 52). Upon completion of the study, subjects may be invited to participate in an additional observational study in which ART will be interrupted to evaluate the effect of study treatment on virological control. Enrolment into this part of the study will be optional and determined by the effect of study treatments on the latent HIV-1 reservoir. (Maximum duration of treatment interruption: 16 weeks).

In summary:

Study arm 1: Peptide composition + IMiD + HDAC (panobinostat)
Study arm 2: Peptide composition + HDAC (panobinostat)
Study arm 3: HDAC (panobinostat)

Depletion of the viral reservoir as a result of the combination treatments according to the present invention may be quantified by for instance following the procedures set forth in Lehrman et al. (The Lancet (366), 2005, pp. 549-555) and references there in. In brief, this includes measuring in samples of patient blood obtained before, during and after
treatment; p24 expression from stimulated latently infected cells, plasma HIV RNA concentration (viral load), and integrated HIV DNA by realtime PCR analysis.

EXAMPLE 9

5 DC/ T-CELL PROLIFERATION ASSAY

Dendritic cells (DC) were generated from monocytes isolated from buffy coat preparations from healthy blood donors. Briefly, peripheral blood mononuclear cells were separated by a density gradient centrifugation and the monocytes were then negatively isolated using the Dynabeads Untouched Human Monocytes (Invitrogen, Carlsbad, CA) following the manufacturer’s instructions. The monocytes were cultured with IL-4 (20 ng/ml; Immunotools, Friesoythe; Germany) and GM-CSF (100 ng/ml; Immunotools) in X-VIVO15 medium (Lonza, Basel, Switzerland) for 5–6 days to generate immature DC. Cytokines were replenished every 2–3 days. The maturation of the cells was performed for 24 hours with IFN-γ (1000 IU/ml), TNF-α (50 ng/ml), IL-1β (25 ng/ml) IFN-α (3000 IU/ml). After maturation, the DC were pulsed for 2 hours at 37°C with peptides at 10μg/ml, before extensive washing and co-culture with Peripheral blood mononuclear cells (PBMC) labelled with a fluorescent dye (VPD450, BD biosciences, Sam Jose, CA). Various ratios with DC:T cell were tested alongside with appropriate controls. IL-2 (50U/ml) and IL-7 (50ng/ml) (Both, Immunotools) and wells with or without IMiDs were added at the start of co-culture. At day 6-10, the level of T cell proliferation was analysed by flow cytometry. The supernatants from the co-culture wells were investigated with Luminex technology to establish any suppressor activity.

EXAMPLE 10

The peptides according to the invention used in the following examples were synthesized by Schafer-N as c-terminal amides using the Fmoc-strategy of Sheppard, (1978) J.Chem.Soc., Chem. Commun., 539.

Cell penetration assay

Intracellular staining for biotinylated peptides

96-well U-bottom polystyrene plates (NUNC, cat no: 163320) were used for staining of human PBMCs. Briefly, 8ul of N- or C-terminally biotinylated peptides according to the invention (i.e. 5mM, 2.5mM & 1.25mM tested for each peptide) were incubated at 37°C for 2h with 40ul of PBMC (12.5 x 106 cells/ml) from blood donors. Cells were then washed 3x with 150ul of Cellwash (BD, cat no: 349524), followed by resuspension of each cell pellet with 100ul of Trypsin-EDTA (Sigma, cat no: T4424), then incubated at 37°C for 5 min. Trypsinated cells were then washed 3x with 150ul of Cellwash (BD, cat no: 349524), followed by resuspension with BD Cytofix/Cytoperm™ plus (BD, cat no: 554715), then incubated at 4°C for 20 min according to manufacturer. Cells were then washed 2x with 150ul PermWash
Cells were then stained with Streptavidin-APC (BD, cat no: 554067) & Anti-hCD11c (eBioscience, cat no: 12-0116) according to manufacturer at 4°C for 30 min aiming to visualize biotinylated peptides & dendritic cells, respectively. Cells were then washed 3x with 150ul PermWash, followed by resuspension in staining buffer (BD, cat no: 554656) before flow cytometry. Dendritic cells were gated as CD11c+ events outside lymphocyte region (i.e. higher FSC & SSC signals than lymphocytes). 200 000 total cells were acquired on a FACSCanto II flow cytometer with HTS loader, and histograms for both total cells & dendritic cells with respect to peptide-fluorescence (i.e. GeoMean) were prepared.

Extracellular staining for biotinylated peptides

96-well U-bottom polystyrene plates (NUNC, cat no: 163320) were used for staining of human PBMCs. Briefly, 8ul of N- or C-terminally biotinylated peptides according to table 1 (i.e. 5mM, 2.5mM & 1.25mM tested for each peptide; all peptides manufactured by solid phase synthesis by commercial suppliers) were incubated at 37°C for 2h with 40ul of PBMC (12.5 x 106 cells/ml) from blood donors. Cells were then washed 3x with 150ul of Cellwash (BD, cat no: 349524), then stained with Streptavidin-APC (BD, cat no: 554067) & Anti-hCD11c (eBioscience, cat no: 12-0116) according to manufacturer at 4°C for 30 min aiming to visualize biotinylated peptides & dendritic cells, respectively. Cells were then washed 3x with 150ul of Cellwash (BD, cat no: 349524), followed by resuspension in staining buffer (BD, cat no: 554656) before flow cytometry. Dendritic cells were gated as CD11c+ events outside lymphocyte region (i.e. higher FSC & SSC signals than lymphocytes). 200 000 total cells were acquired on a FACSCanto II flow cytometer with HTS loader, and histograms for both total cells & dendritic cells with respect to peptide-fluorescence (i.e. GeoMean) were prepared.

It was clearly seen that the CMI peptides according to the invention had improved ability to enter the cell compared to its native counterparts

The data are geomean-value of each tested peptide, as calculated by the FACS Duva software. The Geomean values by trypsinizing/Cytofix/Cytoperm.: 

EXAMPLE 11

Positive CTL response may alternatively be assayed by ELISPOT assay.

Human IFN-gamma cytotoxic T-cell (CTL) response by ELISPOT assay

Briefly, at day 1, PBMC samples from HCV patients were incubated in flasks (430 000 PBMCs/cm2) for 2h at 37°C, 5% CO2 in covering amount of culture media (RPMI 1640 Fisher Scientific; Cat No. PAAE15-039 supplemented with L- Glutamine, (MedProbe Cat. No. 13E17-605E, 10% Foetal Bovine serum (FBS), Fisher Scientific Cat. No. A15-101) and Penicillin/Streptomycin, (Fisher Acientific Cat. No. P11-010) in order to allow adherence of monocytes. Non-adherent cells were isolated, washed, and frozen in 10% V/V DMSO in FBS until further usage. Adherent cells were carefully washed with culture media, followed by incubation at 37°C until day 3 in culture media containing 2μg/ml final concentration of hrGM-CSF (Xiamen amoytop biotech co, cat no: 3004.9090.90) & 1μg/ml hrIL-4 (Invitrogen,
Cat no: PHC0043) and optionally an immunomodulation agent (IMiD), and this procedure was then repeated at day 6. At day 7, cultured dendritic cells (5 000-10 000 per well) were added to ELISPOT (Millipore multiscreeen HTS) plates coated with 0.5μg/well anti-human γ Interferon together with thawed autologous non-adherent cells (200 000 per well), antigen samples (1-8μg/ml final concentration for peptide antigens; 5μg/ml final concentration for Concanavalin A (Sigma, Cat no: C7275) or PHA (Sigma, Cat no: L2769)) & anti-Anergy antibodies (0.03-0.05μg/ml final concentration for both anti-PD-1 (eBioscience, cat no: 16-9989-82) & anti-PD-L1 (eBioscience, cat no: 16-5983-82)). Plates were incubated overnight and spots were developed according to manufacturer. Spots were read on ELISPOT reader (CTL-ImmunoSpot® S5 UV Analyzer).

EXAMPLE 12

ELISPOT assay

At day one, PBMC samples from blood donors were thawed, washed with warm medium and incubated in flasks (250000PBMCs/cm2) for 24 hours at 37°C, 5% CO2 in covering amount of culture media (RPMI 1640 with ultra-glutamine, Lonza, BE12-702F701; 10% Foetal Bovine serum (FBS), Fisher Scientific Cat. No. A15-101; Penicillin/Streptomycin, Fisher Scientific Cat. No. P11-010) to allow the cells to recover after thawing. At day two, the cells were added to a Falcon Microtest Tissue Culture plate, 96well flat bottom, at 500 000 cells per well in a volume of 200μl total medium. Parallel wells were added the indicated stimuli in duplicate and optionally an immunomodulation agent (IMiD), or left with medium as a control for 6 days at 37°C, 5% CO2. After the six days of incubation, 100μl of the cell suspension were transferred to an ELISPOT (Millipore multiscreeen HTS) plate coated with 1μg/ml native influenza M2e protein. After a 24 hour incubation, the plate was washed four times with PBS + 0,05% Tween20, and a fifth time with PBS, 200μl/well. A mouse Anti-human IgG or IgM biotin (Southern Biotech 9040-08 and 9020-08) was diluted in PBS with 0.5% FBS and incubated for 90 minutes at 37°C. The washing was repeated as described, before 80μl Streptavidin-Alkaline-Phosphatase (Sigma Aldrich, S289) was added each well and incubated at 60 minutes in the dark, at room temperature. The wells were then washed 2 times with PBS + 0.05% Tween20 and 4 times with PBS, 200μl/well, before the substrate, Vector Blue Alkaline Phosphatase Substrate kit III (Vector Blue, SK-5300 ) was added and let to develop for 7 minutes at room temperature. The reaction was stopped with running water, the plates let dry and the sport enumerated by an ELISPOT reader (CTL-ImmunoSpot® S5 UV Analyzer).

ELISA

100 μl of antigen as indicated (pre-incubated in Coating buffer - 0.05M Na2CO3 pH9.6; denoted CB - in cold at 8μg/ml 1-3 days) or just CB (background control) was used for coating wells in microtiter plates at 4°C. The microtiter plates are then washed 3x with washing buffer (PBS + 1% v/v Triton-X100; denoted WB), followed by 2h blocking at room
temperature (RT) with 200 µl/well of blocking buffer (PBS + 1% w/v BSA). Plates are then washed 3x with WB, followed by 1h incubation at 37°C with 50-70 µl/well of added human (or rabbit or sheep) sera (serial dilutions ranging from 1:5 – 1:250 in dilution buffer (PBS + 1% v/v Triton-X100 + 1% w/v BSA; denoted DB)). Plates are then washed 6x with WB, followed by 1h incubation at RT with 70 µl/well of Alkaline Phosphatase-conjugated Protein G (3µg/ml in DB; Calbiochem 539305) or goat anti-mouse IgG biotin (1µg/ml, Southern Biotech, 1030-08. In case of the goat anti-mouse IgG biotin, the plates were washed one extra step as described, before addition of 100µl Streptavidin-Alkaline-Phosphatase (1µg/ml, Sigma Aldrich, S289) and incubated 1 hour at RT. Plates are then washed 6x with WB, followed by 10-60 min incubation at room temperature with 100 µl/well of 0.3% w/v of Phenolphthalein monophosphate (Sigma P-5758). Plates are finally quenched by adding 100 µl/well of Quench solution (0.1M TRIS + 0.1M EDTA + 0.5M NaOH + 0.01% w/v NaN3; pH14), followed by a measurement with an ELISA reader (ASYS UVM 340) at 550 nm. The strength of the sera, i.e. the magnitude of the humoral immune response, is then reported as the dilution of sera that result in the described Optical Density (OD) value, or the OD value at the indicated dilution of sera.

EXAMPLE 13

CLINICAL TRIAL PROTOCOL - Phase I/IIa Study to Evaluate the Effect of Therapeutic HIV-1 Immunization using Vacc-4x + rhuGM-CSF, and HIV-1 Reactivation using Romidepsin, on the Viral Reservoir in Virologically Suppressed HIV-1 Infected Adults on cART.

The primary objective is to measure the effect of treatment with Vacc-4x + rhuGM-CSF and cyclic romidepsin treatment on the HIV-1 latent reservoir in HIV-infected patients virologically suppressed on cART.

Endpoints:
Primary Endpoints:

1) Safety and tolerability evaluation as measured by adverse events (AE), adverse reactions (AR), serious adverse events (SAE), serious adverse reactions (SAR), serious unexpected adverse reactions (SUSAR)
2) Latent reservoir size measured in CD4+ T cells by:
a) HIV-1 viral outgrowth assay (HIV-1 RNA per 106 in resting memory CD4+ T cells (RUPM))
b) Integrated HIV-1 DNA (copies per 106 CD4+ T cells)
c) Total HIV-1 DNA (copies per 106 CD4+ T cells)

Secondary Endpoints PART B
1) Time to re-initiation of cART
2) Time to detectable viremia during cessation of cART
3) HIV transcription measured as cell associated unspliced HIV-1 RNA (copies per 10⁶ CD4+ T cells)
4) HIV-specific T-cell responses as measured by ELISpot, proliferation and/or intracellular cytokine staining
5) Plasma HIV-1 viral load
6) Histone H3 acetylation as measured in lymphocytes
7) T cell count and phenotype
8) Antibody titer to Vacc-4x peptides and to p24 as measured by ELISA.

10) An Open Phase I/IIa Study to Evaluate the Effect of Therapeutic HIV-1 Immunization using Vacc-4x + rhuGM-CSF, and HIV-1 Reactivation using Romidepsin, on the Viral Reservoir in Virologically Suppressed HIV-1 Infected Adults on cART. The study is conducted to evaluate the safety/tolerability of Vacc-4x + rhuGM-CSF as adjunctive therapy to romidepsin and to assess the impact on the latent HIV reservoir and the ability to control viral load during an Analytical Treatment Interruption (n=20, ie. 20 patients).

Target Population: Virologically suppressed (pVL < 50copies/mL) HIV-1 infected adults currently on cART.

Study Procedures/frequency:
1. A pre-treatment phase of 4 weeks (visit 1 to visit 2) to confirm the stability of the latent HIV-1 reservoir and determine baseline HIV-1 T lymphocyte specific immunity.
2. A therapeutic HIV-1 immunization phase of 12 weeks (from visit 2 to visit 7) in which Vacc-4x will be administered together with rhuGM-CSF at visit 2, 3, 4, 5, 6 and 7 follow by a follow-up period of 2 weeks (visit 8-9).
3. A viral reactivation phase of 3 weeks (visit 10-visit 12) consisting of one cycle of romidepsin infusions at a dosing of 5 mg/m².
   A post-treatment observation phase of ~8 weeks (visit 13-visit 14) to assess the effect of the investigational treatment on the size of the latent HIV-1 reservoir.
4. An Analytical Treatment Interruption phase of 16 weeks (from after visit 15-34).

Investigational Medicinal Products:
Vacc-4x: 1.2mg administered intradermally at day 0, 7, 14, 21, 77 and 84 (visit 2, 3, 4, 5, 6 and 7)
rhuGM-CSF: Leukine® (Sanofi) 0.06mg administered intradermally, 10 min prior to Vacc-4x administration, at day 0, 7, 14, 21, 77 and 84 (visit 2, 3, 4, 5, 6 and 7)
Romidepsin: Istodax® (Celgene) 5mg/m² administered by 3 intravenous infusion in three consecutive weeks (day 105, 112 and 119) (visit 10, 11b and 12) (corresponding to one 28 day cycle).
Trial Design:
1. A pre-treatment phase of 4 weeks (visit 1 to visit 2) to confirm the stability of the latent HIV-1 reservoir and determine baseline HIV-1 T lymphocyte specific immunity.
2. A therapeutic HIV-1 immunization phase of 12 weeks (2 to visit 7) in which Vacc-4x will be administered together with rhuGM-CSF at visit 2, 3, 4, 5, 6 and 7 followed by a follow-up period of 2 weeks (visit 8 to visit 9).
3. A viral reactivation phase of 3 weeks (visit 10 to visit 12) consisting of one cycle of romidepsin infusions at a dosing of 5 mg/m2.
4. A post-treatment observation phase of ~8 weeks (visit 13 to visit 14) to assess the effect of the romidepsin on the size of the latent HIV-1 reservoir.
5. An Analytical Treatment Interruption phase of 16 weeks (visit 15-34).

Treatment

Vacc-4x

Vacc-4x, consists of four synthetic peptides (Vacc-10 acetate, Vacc-11 acetate, Vacc-12 acetate, and Vacc-13 acetate), each corresponding to conserved domains on the HIV-1 p24 capsid protein representing the native Gag regions with residues 166-185, 252-269, 264-284, and 335-354, respectively.

Vacc-4x is manufactured in accordance with Good Manufacturing Practice (GMP) and is supplied as sterile vials of freeze-dried white powder. There is no additional ingredient in the product.

RhuGM-CSF (sargramostim, Leukine®, Sanofi)

Leukine® is manufactured by Sanofi and supplied by Genzyme. It is a glycoprotein of 127 amino acids characterized by three primary molecular species having molecular masses of 19,500, 16,800 and 15,500 daltons. The liquid Leukine® presentation is formulated as a sterile, preserved (1.1% benzyl alcohol), injectable solution (500mcg/mL) in a vial. Lyophilized Leukine® is a sterile, white, preservative-free powder (250mcg) that requires reconstitution with 1 mL Sterile Water for Injection, USP or 1 mL Bacteriostatic Water for Injection, USP. Liquid Leukine® has a pH range of 6.7 – 7.7 and lyophilized Leukine® has a pH range of 7.1 – 7.7.

For further information refer to IB (Leukine® prescribing information).

Romidepsin (Istodax®, Celgene)

Istodax® is manufactured by Celgene Corporation. This histone deacetylase (HDAC) inhibitor is a bicyclic depsipeptide. At room temperature, romidepsin is a white powder and is described chemically as (1S,4S,7Z,10S,16E,21R)-7-ethylidene-4,21-bis(1-methylethyl)-2-oxa-12,13-dithia-5,8,20,23-tetraazabicyclo[8.7.6] tricos-16-ene-3,6,9,19,22-pentone. The
empirical formula is C24H36N4O6S2. IstoDEX® is supplied as a kit containing two vials.

IstoDEX® (romidepsin) for injection is a sterile lyophilized white powder and is supplied in a single-use vial containing 10mg romidepsin and 20mg povidone, USP. Diluent for IstoDEX® is a sterile clear solution and is supplied in a single-use vial containing a 2-mL deliverable volume. Diluent for IstoDEX® contains 80% (v/v) propylene glycol, USP and 20% (v/v) dehydrated alcohol, USP.

For further information refer to IB for romidepsin.

Vacc-4x

Each dose of Vacc-4x (0.1mL of a 12mg/mL solution), will be administered by intradermal injections following the intradermal administration of rhuGM-CSF (Leukine®) as adjuvant. A total of 6 Vacc-4x/rhuGM-CSF immunizations (visit 3, 4, 5, 6, 7 and 8) are planned in the HIV-1 therapeutic vaccination phase.

Approximately 10 minutes before each administration of Vacc-4x, rhuGM-CSF will be administered intradermally as an adjuvant. Vacc-4x must be administered intradermally at the same site as rhuGM-CSF, superficial to the deltoid muscle and in the same arm during the course of the study.

When administering the intradermal injection, utmost care must be taken so that no material is injected subcutaneously. If administered correctly, after puncture of the skin a small bleb should appear following the injection of only a small amount of product. An injection that is too superficial should be avoided as this will result in loss of the sample volume from the injection site during injection or after withdrawal of the needle.

RhuGM-CSF

Each dose of rhuGM-CSF (0.1mL of 0.60mg/mL solution) will be administered as an adjuvant by intradermal injection 10 minutes prior to the intradermal administration of Vacc-4x immunizations (visit 3, 4, 5, 6, 7 and 8) during the HIV-1 therapeutic vaccination phase. rhuGM-CSF must be administered intradermally at the same site as Vacc-4x, superficial to the deltoid muscle and in the same arm during the entire course of the study.

When administering the intradermal injection, utmost care must be taken so that no material is injected subcutaneously. If administered correctly, after puncture of the skin a small bleb should appear following the injection of only a small amount of product. An injection that is too superficial should be avoided as this will result in loss of the sample volume from the injection site during injection or after withdrawal of the needle.

Romidepsin

The dose is 5mg/m2 administered intravenously over a 4 hour period on Days 1, 8, and 15 of a 28-day cycle (visit 10, 11 and 12).
Trial Assessment:
Laboratory Assessment
Biochemistry:
Routine biochemistry includes haematology parameters (haemoglobin, total and differential leukocyte count, platelet count), ALAT, bilirubin, alkaline phosphatase, creatinine, sodium, potassium, phosphorus, magnesium, calcium, urea, albumin and CRP.
HIV Virology:
HIV-1 viral outgrowth (HIV-1 RNA per 10^6 resting memory CD4+ T cells (RUPM)): The gold standard assay used to measure the frequency of resting CD4+ T cells carrying latent but replication competent virus is based on co-culture of highly purified resting CD4+ T cells from the patient together with PBMCs from an HIV-negative donor and is measured as infectious units per million cells (IUPM) [Finzi 1999, Chun 2007].
Integrated HIV-1 DNA (copies per 10^6 CD4+ T cells): Within infected cells, HIV DNA can exist as linear non-integrated forms, circular forms and as an integrated provirus. In patients receiving effective cART, the majority of HIV DNA is integrated in resting latently infected CD4+T cells. The most widely used technique to quantify the number of cells that contain integrated virus is the Alu-LTR PCR assay [Sonza 1996].
Total HIV-1 DNA (copies per 10^6 CD4+ T cells): Total HIV DNA quantifies integrated and non-integrated DNA as well as latent and defective virus. There is a strong correlation between total HIV DNA and integrated HIV DNA in patients on cART and therefore cell-associated HIV DNA is likely to be a good surrogate marker of the total number of latently infected cells [Koelsch 2008].
Unspliced HIV-1 RNA (copies per 10^6 CD4+ T cells): HIV transcription is measured as copies of cell-associated unspliced HIV-1 RNA/106 CD4+ T cells using digital droplet PCR
Plasma HIV-1 RNA detection by NAT screen: Measured by a transcription mediated amplification (TMA)-based methodology, usually referred to as a nucleic acid test (NAT)-screen (PROCLEYX ULTRIO Plus, Genprobe).
Plasma HIV RNA, quantitative viral load: Measured by Roche VL (routine clinical assay)
Histone H3 acetylation: Measured in lymphocytes using flow cytometry with intracellular cytokine stain on fresh isolated PBMCs.
T Cell count (CD4 and CD8)
Phylogenetic analysis

Immunology:
HIV-specific T cell response as measured by ELISpot, proliferation and/or intracellular cytokine staining

EXAMPLE 14
Overall Study Design

The study CT-BI Vacc-4x 2012/1 (EudraCT Number: 2012-002281-12) is an open label, multicenter, follow-up, re-boosting study of subjects who previously completed the immunization regimen with Vacc-4x active and stopped ART (at Week 28) in the CT-BI Vacc-4x 2007/1 Study (EudraCT Number 2007-006302-13). No restart of ART is required. The subjects will be re-boosted with two immunizations (Visit 2 and Visit 3) of Vacc-4x, 1.2 mg peptides (12 mg/mL), with a 2 week interval between immunizations.

At Visit 5 ART medication (if being used) will be stopped in subjects whose CD4 count ≥ 350x106 /L, for a 16 week period. At Week 29 ART will be restarted, according to the Investigator and subject’s decision, and the subjects will be followed for an additional 8 weeks (End of Study; Visit 10).

DTH will be measured at Visit 2, and at Visit 4 (3 weeks after the second re-boosting immunization).

Viral load, CD4 and CD8 counts will be measured at all study visits. T-cell responses will be measured at Visits 2, 4, 6, 9 and 10 (End of Study) both by ELISPOT, T-cell proliferation assay and intracellular cytokine staining.

Vital signs and clinical laboratory tests will be done at all study visits.

Monitoring of AEs and concomitant medications will be done continuously from the time of signing ICF through the End of Study.

The prior study (CT-BI Vacc-4X 2007/1) was completed by 88 subjects who received active Vacc-4x and stopped ART at Week 28. It was estimated that approximately 30-40 of these subjects would be eligible for this follow-up re-boost study.

All 33 subjects enrolled into this re-boost study were included in the Safety analysis set. Three subjects were excluded from the Intention To Treat (ITT) analysis set in each case this was because ART was not received from Screening to Week 12 and so it was not possible to discontinue ART at Week 12 as required by the protocol. The same three subjects were excluded from the PP analysis set plus a further three subjects who did not discontinue ART at Week 12. For the enrolled patients the mean time between the last immunization in the prior study (CT-BI Vacc-4X 2007/1) and the first immunization in this study was about 197 weeks.
Proviral DNA

HIV-1 DNA levels will be determined by real-time PCR (Taqman) assay targeting HIV-1 gag gene. Briefly, DNA for each subject at visit 2, 4, 6, 9 and 10 will be extracted from total PBMCs (1-4 million cells), eluted in appropriate storage buffer, quantified and stored at -20°C until used. Equal amounts of DNA (≈300ng) will be used to quantify gag and albumin genes in order to determine the number of copies of HIV-DNA per million of cells.

There was a decrease in proviral DNA levels of approximately 50% after immunization while on ART, based on geometric means. This could suggest immune-based killing of infected cells following re-boosting with Vacc-4x. The per protocol (PP) analysis set consists of all subjects who received two re-boosting Vacc-4x immunizations, who discontinued ART at Week 12 (as planned), and who did not incur a major protocol deviation (violation) that would challenge the validity of their data.

<table>
<thead>
<tr>
<th>Table 2</th>
<th>Proviral DNA Over Time</th>
</tr>
</thead>
<tbody>
<tr>
<td>copies/mL</td>
<td>PP (N = 27)</td>
</tr>
<tr>
<td><strong>Baseline</strong></td>
<td></td>
</tr>
<tr>
<td>n</td>
<td>26</td>
</tr>
<tr>
<td>Mean</td>
<td>98.4</td>
</tr>
<tr>
<td>Geometric mean</td>
<td>22.3</td>
</tr>
<tr>
<td>SD</td>
<td>146.69</td>
</tr>
<tr>
<td>Median</td>
<td>57.0</td>
</tr>
<tr>
<td>Q1 to Q3</td>
<td>0.0 to 114.0</td>
</tr>
<tr>
<td>Min to max</td>
<td>0 to 598</td>
</tr>
<tr>
<td><strong>Week 4</strong></td>
<td></td>
</tr>
<tr>
<td>n</td>
<td>27</td>
</tr>
<tr>
<td>Mean</td>
<td>100.8</td>
</tr>
<tr>
<td>Geometric mean</td>
<td>12.9</td>
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<tr>
<td>SD</td>
<td>176.64</td>
</tr>
<tr>
<td>Median</td>
<td>45.0</td>
</tr>
<tr>
<td>Q1 to Q3</td>
<td>0.0 to 90.0</td>
</tr>
<tr>
<td>Min to max</td>
<td>0 to 769</td>
</tr>
</tbody>
</table>

Geometric mean calculated as anti-log of (mean log_{10} VL values + 1) - 1

The comparison of proviral DNA at Baseline and Week 4 in the 2012/1 study is summarized in Table 3. There was a decrease from Baseline to Week 4 of 50% in the PP analysis set, which was statistically significant.
<table>
<thead>
<tr>
<th></th>
<th>Copies/mL</th>
<th>PP</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Baseline</strong></td>
<td></td>
<td></td>
</tr>
<tr>
<td>n</td>
<td></td>
<td>26</td>
</tr>
<tr>
<td>Mean</td>
<td></td>
<td>98.4</td>
</tr>
<tr>
<td>Geometric mean</td>
<td></td>
<td>22.3</td>
</tr>
<tr>
<td>SD</td>
<td></td>
<td>146.69</td>
</tr>
<tr>
<td>Median</td>
<td></td>
<td>57.0</td>
</tr>
<tr>
<td>Q1 to Q3</td>
<td></td>
<td>0.0 to 114.0</td>
</tr>
<tr>
<td>Min to max</td>
<td></td>
<td>0 to 598</td>
</tr>
<tr>
<td><strong>Week 4</strong></td>
<td></td>
<td></td>
</tr>
<tr>
<td>n</td>
<td></td>
<td>26</td>
</tr>
<tr>
<td>Mean</td>
<td></td>
<td>93.9</td>
</tr>
<tr>
<td>Geometric mean</td>
<td></td>
<td>11.4</td>
</tr>
<tr>
<td>SD</td>
<td></td>
<td>176.36</td>
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<tr>
<td>Median</td>
<td></td>
<td>34.5</td>
</tr>
<tr>
<td>Q1 to Q3</td>
<td></td>
<td>0.0 to 89.0</td>
</tr>
<tr>
<td>Min to max</td>
<td></td>
<td>0 to 769</td>
</tr>
<tr>
<td>Geometric mean ratio (Week 4/Baseline) [a]</td>
<td>0.5</td>
<td>0.31 to 0.93</td>
</tr>
<tr>
<td>95% CI</td>
<td>0.030</td>
<td></td>
</tr>
<tr>
<td>p value [b]</td>
<td>26</td>
<td></td>
</tr>
</tbody>
</table>

Geometric means are calculated as the antilog of the (mean log10 VL values + 1) -1

[a] Calculated as anti-log \{mean[(log10 Week 4 + 1) - (log10 Baseline + 1)]\}

[b] Non-parametric Wilcoxon signed-rank test

Assays for measurement of HIV-1 DNA levels:

Integrated HIV-1 DNA (copies per 10^6 CD4+ T cells): Within infected cells, HIV DNA can exist as linear non-integrated forms, circular forms and as an integrated provirus. In patients receiving effective cART, the majority of HIV DNA is integrated in resting latently infected CD4+T cells. The most widely used technique to quantify the number of cells that contain integrated virus is the Alu-LTR PCR assay [Sonzia; J Virol. Jun 1996; 70(6): 3863-3869][Liszewski; Methods April 2009; 47(4): 254-260]. Alternative methods are described by Graaf, Deeks & CO for integrated PCR: "repetitive-sampling Alu-gag PCR", doi:10.1016/j.ymeth.2009.01.002. In some embodiments HIV-1 DNA levels is measured as described in Graf, E. H., A. M. Mexas, et al. (2011). "Elite Suppressors Harbor Low Levels of Integrated HIV DNA and High Levels of 2-LTR Circular HIV DNA Compared to HIV+ Patients On and Off HAART." PLoS Pathog 7(2).

Total HIV-1 DNA (copies per 10^6 CD4+ T cells): Total HIV DNA quantifies integrated and non-integrated DNA as well as latent and defective virus. There is a strong correlation between total HIV DNA and integrated HIV DNA in patients on cART and therefore cell-associated HIV
DNA is likely to be a good surrogate marker of the total number of latently infected cells [Koelsch; J Infect Dis. 2008 Feb 1;197(3):411-9. doi: 10.1086/525283].

In one example total HIV-1 DNA levels may be measured as follows:

5 HIV-1 DNA levels may be determined by real-time PCR (Taqman) assay targeting HIV-1 gag gene. Briefly, DNA for each subject is extracted from total PBMCs (1-4 million cells), eluted in appropriate storage buffer, quantified and stored at -20°C until used. Equal amounts of DNA (≈300ng) will be used to quantify gag and albumin genes in order to determine the number of copies of HIV-1 DNA per million of cells. Throughout the specification and the claims which follow, unless the context requires otherwise, the word ‘comprise’, and variations such as ‘comprises’ and ‘comprising’, will be understood to imply the inclusion of a stated integer, step, group of integers or group of steps but not to the exclusion of any other integer, step, group of integers or group of steps.

15 All patents and patent applications referred to herein are incorporated by reference in their entirety.

The application of which this description and claims forms part may be used as a basis for priority in respect of any subsequent application. The claims of such subsequent application may be directed to any feature or combination of features described herein. They may take the form of product, composition, process, or use claims and may include, by way of example and without limitation, the claims.
CLAMS

1. A method for reducing and/or delaying pathological effects of human immunodeficiency virus I (HIV) or for reducing the risk of developing acquired immunodeficiency syndrome (AIDS) in a human subject infected with HIV, the method comprising the steps of:
   
   a) a therapeutic HIV-1 immunization phase consisting of the administering in one or more doses of an effective amount of one or more HIV-specific peptide selected from the list consisting of the amino acid sequence shown in SEQ ID NO: 18 (Vacc-10), SEQ ID NO: 11 (Vacc-11), SEQ ID NO: 6 (Vacc-12), and SEQ ID NO: 3 (Vacc-13) over a period of 1-12 weeks;
   
   b) one or more subsequent or simultaneous measurements of HIV-1 DNA levels in said human subject infected with HIV; and optionally
   
   c) a subsequent viral reactivation phase consisting of the administering of an effective amount of a reservoir purging agent.

2. A method for monitoring the effect of a therapeutic HIV-1 immunization phase consisting of the administering in one or more doses of an effective amount of one or more HIV-specific peptide selected from the list consisting of the amino acid sequence shown in SEQ ID NO: 18 (Vacc-10), SEQ ID NO: 11 (Vacc-11), SEQ ID NO: 6 (Vacc-12), and SEQ ID NO: 3 (Vacc-13) over a period of 1-12 weeks; in reducing and/or delaying pathological effects of human immunodeficiency virus I (HIV) or in reducing the risk of developing acquired immunodeficiency syndrome (AIDS) in a human subject infected with HIV, the method comprising the step of
   
   a) One or more measurements of HIV-1 DNA levels in said human subject infected with HIV subsequent or simultaneous to said immunization phase.

3. The method or effective amount according to any one of claims 1-2, wherein said subjects are being treated with a combination antiretroviral therapy (cART) prior to and/or during, and/or after said immunization phase, and/or said viral reactivation phase.

4. The method according to any one of claims 1-3, which method further comprises a step b2) subsequent to step b) of selecting human subjects, wherein the level of HIV-1 DNA in said subjects is at least 1, such as at least 2, 3, 4, 5, 6, 7, 8, 9, 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 100, 200, 300, 400, 500, 600, 700, 800, 900,
1000, 2000, 3000, 4000, 5000, 6000, 7000, 8000, 9000, or 10000 HIV-1 DNA copy per million cell over a period of 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 100, 110, 120, 130, 140, 150, 160, 170, 180, 190, 200, 210, 220, 230, 240, 250, 260, 270, 280, 290, 300, 310, 320, 330, 340, 350, 360, or 370 weeks after the therapeutic HIV-1 immunization phase consisting of the administering in one or more doses under step a); and repeating step a) and/or b) and/or optionally step c) for said selected subjects.

5. The method according to any one of claims 1-4, which method further comprises a step b2) subsequent to step b) of selecting human subjects, wherein the level of HIV-1 DNA in said subjects is at least 10%, such as at least 15%, 20%, 25%, 30%, 35%, 40%, 45%, 50%, 55%, 60%, 65%, 70%, 75%, 80%, 85%, 90%, or 95% of said level prior to said immunization phase as measured over a period of 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 100, 110, 120, 130, 140, 150, 160, 170, 180, 190, 200, 210, 220, 230, 240, 250, 260, 270, 280, 290, 300, 310, 320, 330, 340, 350, 360, or 370 weeks after said therapeutic HIV-1 immunization phase consisting of the administering in one or more doses under step a); and repeating step a) and/or b) and/or optionally step c) for said selected subjects.

6. The method according to any one of claims 1-5, which method further comprises a step b2) subsequent to step b) of selecting human subjects, wherein the level of HIV-1 DNA in said subjects is less than 10000, such as less than 9000, 8000, 7000, 6000, 5000, 4000, 3000, 2000, 1000, 900, 800, 700, 600, 500, 400, 300, 200, 100, 95, 90, 85, 80, 75, 70, 65, 60, 55, 50, 45, 40, 35, 30, 25, 20, 15, 10, 5, 4, 3, 2, or 1 HIV-1 DNA copy per million cell over a period of 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 100, 110, 120, 130, 140, 150, 160, 170, 180, 190, 200, 210, 220, 230, 240, 250, 260, 270, 280, 290, 300, 310, 320, 330, 340, 350, 360, or 370 weeks after the therapeutic HIV-1 immunization phase consisting of the administering in one or more doses under step a); and treating said selected subjects under step c).

7. The method according to any one of claims 1-6, which method further comprises a step b2) subsequent to step b) of selecting human subjects, wherein the level of HIV-1 DNA in said subjects is less than 95%, such as less than 90%, 85%, 80%, 75%, 70%, 65%, 60%, 55%, 50%, 45%, 40%, 35%, 30%, 25%, 20%, 15%, 10%, 5%, 4%, 3%, 2%, or 1% of said level prior to said immunization phase as measured over a period of 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 100, 110, 120, 130, 140, 150, 160, 170, 180, 190, 200, 210, 220, 230, 240, 250, 260, 270, 280, 290, 300, 310, 320, 330, 340, 350, 360, or 370 weeks after said therapeutic HIV-1 immunization phase
consisting of the administering in one or more doses under step a); and treating said selected subjects under step c).

8. The method according to any one of claims 1-7, which method further comprises a step b2) subsequent to step b) of selecting human subjects, wherein the level of HIV-1 DNA in said subjects is less than 100, such as less than 95, 90, 85, 80, 75, 70, 65, 60, 55, 50, 45, 40, 35, 30, 25, 20, 15, 10, 5, 4, 3, 2, 1 HIV-1 DNA copy per million cell over a period of 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 100, 110, 120, 130, 140, 150, 160, 170, 180, 190, 200, 210, 220, 230, 240, 250, 260, 270, 280, 290, 300, 310, 320, 330, 340, 350, 360, or 370 weeks after the therapeutic HIV-1 immunization phase consisting of the administering in one or more doses under step a); and repeating step a) and/or b) and/or optionally step c) for said selected subjects.

9. The method according to any one of claims 1-8, which method further comprises a step b2) subsequent to step b) of selecting human subjects, wherein the level of HIV-1 DNA in said subjects is less than 95%, such as less than 90%, 85%, 80%, 75%, 70%, 65%, 60%, 55%, 50%, 45%, 40%, 35%, 30%, 25%, 20%, 15%, 10%, 5% of said level prior to said immunization phase as measured over a period of 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 100, 110, 120, 130, 140, 150, 160, 170, 180, 190, 200, 210, 220, 230, 240, 250, 260, 270, 280, 290, 300, 310, 320, 330, 340, 350, 360, or 370 weeks after said therapeutic HIV-1 immunization phase consisting of the administering in one or more doses under step a); and repeating step a) and/or b) and/or optionally step c) for said selected subjects.

10. The method according to any one of claims 1-9, which method further comprises a step b2) subsequent to step b) of selecting human subjects, wherein the level of HIV-1 DNA in said subjects decreases by more than 10%, such as at least 15%, 20%, 25%, 30%, 35%, 40%, 45%, 50%, 55%, 60%, 65%, 70%, 75%, 80%, 85%, 90%, or 95% of said level prior to said immunization phase as measured over a period of 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 100, 110, 120, 130, 140, 150, 160, 170, 180, 190, 200, 210, 220, 230, 240, 250, 260, 270, 280, 290, 300, 310, 320, 330, 340, 350, 360, or 370 weeks after said therapeutic HIV-1 immunization phase consisting of the administering in one or more doses under step a); and repeating step a) and/or b) and/or optionally step c) for said selected subjects.

11. The method according to any one of claims 1-10, which method further comprises a step b2) subsequent to step b) of selecting human subjects, wherein the level of HIV-1 DNA in said subjects decreases less than 10%, such as less than 9%, 8%, 7%, 6%, 5%, 4%, 3%,
2%, 1% of said level prior to said immunization phase as measured over a period of 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, 100, 110, 120, 130, 140, 150, 160, 170, 180, 190, 200, 210, 220, 230, 240, 250, 260, 270, 280, 290, 300, 310, 320, 330, 340, 350, 360, or 370 weeks after said therapeutic HIV-1 immunization phase consisting of the administering in one or more doses under step a); and treating said selected subjects under step c).

12. The method according to any one of claims 1-11, which method further comprises a step a-1) preceding step a) of measurement of HIV-1 DNA levels in said human subject infected with HIV.

13. The method according to any one of claims 1-12, which method comprises in step a) the administering of two, three, four, five or more doses of an effective amount of one or more HIV-specific peptide selected from the list consisting of the amino acid sequence shown in SEQ ID NO: 18 (Vacc-10), SEQ ID NO: 11 (Vacc-11), SEQ ID NO: 6 (Vacc-12), and SEQ ID NO: 3 (Vacc-13) over a period of 1-12 weeks.

14. The method according to any one of claims 1-13, wherein an adjuvant, such as recombinant human granulocyte-macrophage colony-stimulating factor (rhuGM-CSF) or a water-in-oil adjuvant, such as ISA51 or ISA720, or an oil-in-water adjuvant such as Provax, is administered in conjunction to, prior to or simultaneously with said therapeutic HIV-1 immunization.

15. The method according to any one of claims 1-14, wherein the reservoir purging agent is administered over a period of 1, 2, 3, or 4 consecutive weeks at least about 1, 2, 3, or 4 weeks after said therapeutic HIV-1 immunization phase.

16. The method according to any one of claims 1-15, wherein the viral reactivation phase includes the administration of 1-10 doses, such as 2-10 doses, such as 3-10, such as 4-10, such as 5-10, such as 6-10, such as 7-10, such as 8-10, such as 9-10, such as 10 doses, or 1-9 doses, such as 1-8 doses, such as 1-7, such as 1-6, such as 1-5, such as 1-4, such as 1-3, such as 3 doses of an effective amount of a reservoir purging agent.

17. The method according to claim 1-16, wherein step a) and/or b) are independently repeated 1, 2, 3, 4, 5, 6, 7, 8, 9, or 10 times in any order.

18. The method according to any one of claims 1-17, wherein the reservoir purging agent is an HDAC inhibitor, such as romidepsin or panobinostat.
19. The method according to claim 18, wherein the reservoir purging agent is romidepsin administered by infusions at a dosing of up to 2.5 mg/m², such as up to 5 mg/m², such as up to 7.5 mg/m², such as up to 10 mg/m², such as up to 12 mg/m², such as up to 12.5 mg/m², such as up to 14 mg/m², such as between 2.5 mg/m² and 7.5 mg/m², such as around 5 mg/m².

20. The method according to any one of claims 1-19, wherein the effect on the HIV-1 latent reservoir is in HIV-infected patients virologically suppressed on cART.

21. The method according to any one of claims 1-20, wherein each peptide is given in a dose of 0.1 mg-10 mg per administration, such as 0.1-10 mg per administration, such as 0.1-9 mg per administration, such as 0.1-8 mg per administration, such as 0.1-7 mg per administration, such as 0.1-6 mg per administration, such as 0.1-5 mg per administration, such as 0.1-4 mg per administration, such as 0.1-3 mg per administration, such as 0.1-2 mg per administration, such as 0.1-1.2 mg per administration, such as 0.1-0.6 mg per administration, such as 0.1-0.4 mg per administration.

22. The method according to any one of claims 1-21, wherein the therapeutic HIV-1 immunization phase is over a period of 1-12 weeks, such as over a period of 2-12 weeks, such as over a period of 3-12 weeks, such as over a period of 4-12 weeks, such as over a period of 5-12 weeks, such as over a period of 6-12 weeks, such as over a period of 7-12 weeks, such as over a period of 8-12 weeks

23. The method according to any one of claims 1-22, wherein the therapeutic HIV-1 immunization phase includes the administration of 1-10 doses, such as 2-10 doses, such as 3-10, such as 4-10, such as 5-10, such as 6-10, such as 7-10, such as 8-10, such as 9-10, such as 10 doses.

24. The method according to any one of claims 1-23, wherein said one or more peptide is in the form of an acetate salt.

25. The method according to claim 24, wherein the acetate content of the salt is between 4% and 18%, such as between 5% and 17%, such as between 6% and 16%, such as between 7% and 15%, such as between 8% and 14%, such as between 9% and 14%, such as between 9% and 13%, such as between 10% and 14%, such as between 11% and 14%, or between 5% and 16%, such as between 5% and 15%, such as between 5% and 14%, such as between 6% and 14%, such as between 6% and 13%, such as between 7% and 12%, such as between 7% and 11%, such as between 8% and 11%, such as between 9%
and 11%, or between 3% and 18%, such as between 3% and 17%, such as between 3% and 16%, such as between 3% and 15%, such as between 3% and 14%, such as between 3% and 13%, such as between 3% and 11%, such as between 3% and 10%, such as between 4% and 10%, such as between 4% and 9%, such as between 4% and 8%, such as between 4% and 7%, such as between 4% and 6%, such as between 4% and 5%.

26. The method according to any one of claims 1-25, wherein one, two, three or four peptides are used in the therapeutic HIV-1 immunization phase.

27. The method according to any one of claims 1-26, wherein all four peptide as acetate salts are used in the therapeutic HIV-1 immunization phase.

28. The method according to any one of claims 1-27, wherein the peptides have amide C-terminal ends of formula -C(O)NH2, or acetate salts thereof.

29. The method according to any one of claims 1-28, wherein all four peptide are used in the ratio of 1:1:1:1 w/w.

30. The method according to any one of claims 1-29, wherein said one, two, three or four peptides are in a dissolved liquid state.

31. The method according to claim 30, wherein said liquid is water.

32. The method according to any one of claims 1-31, which method further comprises the administering of one or more further therapeutically active agent selected from an immunomodulatory compound and a second reservoir purging agent, such as a histone deacetylase (HDAC) inhibitor, or a BET family protein inhibitors/antagonist.

33. The method according to claim 32, wherein the immunomodulatory compound is selected from anti-PD1 antibodies, such as MDX-1106 (Merck), THALOMID® (thalidomide), anti-PD1 antibodies, cyclophosphamide, Levamisole, lenalidomide, CC-4047 (pomalidomide), CC-11006 (Celgene), and CC-10015 (Celgene), and immunomodulatory compounds described in any one of WO2007028047, WO2002059106, and WO2002094180.

34. The method according to claim 33, wherein the immunomodulatory compound is lenalidomide.
35. The method according to claims 33 or 34, wherein the reservoir purging agent is selected from M344 (4-(dimethylamino)-N-[7-(hydroxyamino)-7-oxoheptyl]benzamide), chidamide (CS055/HBI-800), 4SC-202, (4SC), Resminostat (4SC), hydroxamic acids such as vorinostat (SAHA), belinostat (PXD101), LAQ824, trichostatin A and panobinostat (LBH589); benzamides such as entinostat (MS-275), CI994, and mocetinostat (MGCD0103), cyclic tetrapeptides (such as trapoxin, such as trapoxin B), and the depsipeptides, such as romidepsin (ISTODAX), electrophilic ketones, and the aliphatic acid compounds such as phenylbutyrate, valproic acid, Oxamflatin, ITF2357 (generic givinostat), Apicidin, MC1293, CG05, and CG06; compounds that activate transcription factors including NF-KappaB, Prostratin, auranofin, bryostatin, a nontumorigenic phorbol ester, DPP (12-deoxyphorbol-13-phenylacetate), PMA, and Phorbol 12-myristate 13-acetate (PMA); Compounds that activate HIV mRNA elongation including P-TEF-b kinase and hexamethylyisacetamide (HMBA); IL-7; T-cell stimulating factors including anti-CD3/CD28 - T-cell stimulating Ab's; Kinase inhibitors including Tyrophostin A, Tyrophostin B, and Tyrophostin C; PTEN (phosphatase and tensin homologue) gene inhibitors including SF1670 (Echelon Bioscience), Disulfiram (DSF), an inhibitor of acetaldehyde dehydrogenase, Protein Tyrosine Phosphatase Inhibitors including bpV(HOpic), bpV(phen), and bpV(pic) (Calbiochem; EMD Millipore), Toll-like receptors agonists including Toll-like receptor-9 (TLR9) and Toll-like receptor-7 (TLR9) agonists, quercetin, lipoic acid, sodium butyrate, TNF-alpha, PHA, Tat, BET family protein inhibitors/antagonists, such as JQ1, I-BET, I-Bet151, MS417, GW841819X, and thienotriazolodiazepine compounds, such as those described in U.S. Patent Application Publication No. 2010/0286127.

36. An effective amount of one or more HIV-specific peptides selected from the list consisting of the amino acid sequence shown in SEQ ID NO: 18 (Vacc-10), SEQ ID NO: 11 (Vacc-11), SEQ ID NO: 6 (Vacc-12) for use in a method for reducing and/or delaying pathological effects of human immunodeficiency virus I (HIV) or for reducing the risk of developing acquired immunodeficiency syndrome (AIDS) in a human infected with HIV, the method comprising the steps of:

a) a therapeutic HIV-1 immunization phase consisting of the administering in one or more doses of an effective amount of one or more HIV-specific peptide selected from the list consisting of the amino acid sequence shown in SEQ ID NO: 18 (Vacc-10), SEQ ID NO: 11 (Vacc-11), SEQ ID NO: 6 (Vacc-12), and SEQ ID NO: 3 (Vacc-13) over a period of 1-12 weeks;

b) a subsequent or simultaneous measurement of HIV-1 DNA levels in said human subject infected with HIV; and optionally

c) a subsequent viral reactivation phase consisting of the administering of an effective amount of a reservoir purging agent.
37. The effective amount according to claim 36, wherein the method is as defined in any one of claims 2-35.
Fig. 1

Vaccinate → Measure DNA

Below Cutoff? Yes → Reactivate

No → Go back to Vaccinate and Measure DNA

Repeats

Vaccinate → Measure DNA

Going Down? Yes → Reactivate

No → Go back to Vaccinate and Measure DNA

Repeats

Vaccinate → Measure DNA

Going Down? Yes → Reactivate

Below Cutoff? Yes → Reactivate

No → Stop
A. CLASSIFICATION OF SUBJECT MATTER
INV. A61K39/21
ADD.

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)
A61K

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)
EPO-Internal, WPI Data, BIOSIS, EMBASE

C. DOCUMENTS CONSIDERED TO BE RELEVANT

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T later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention

X document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone

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& document member of the same patent family

Date of the actual completion of the international search: 2 September 2015

Date of mailing of the international search report: 25/09/2015

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Rojo Romeo, Elena
The query sequence SEQ ID NO:18 has 100.00 % identity (100.00 % similarity) over 27 positions in a common overlap (range (q:s): 1-27:1-27) with subject GSP:AAB18710 (length: 27) from WO200052040-A1 published on 2000-09-08.

The query sequence SEQ ID NO:11 has 100.00 % identity (100.00 % similarity) over 26 positions in a common overlap (range (q:s): 1-26:1-26) with subject GSP:AAB18703 (length: 26) from WO200052040-A1 published on 2000-09-08.

The query sequence SEQ ID NO:6 has 100.00 % identity (100.00 % similarity) over 21 positions in a common overlap (range (q:s): 4-24:3-23) with subject GSP:AAB18699 (length: 23) from WO200052040-A1 published on 2000-09-08.

The query sequence SEQ ID NO:3 has 100.00 % identity (100.00 % similarity) over 20 positions in a common overlap (range (q:s): 1-20:1-20) with subject GSP:AAB18695 (length: 20) from WO200052040-A1 published on 2000-09-08.

see examples, in particular, examples 2, 4, 13, 16,


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