

NATIVE LIGATION PROCESS

FIELD OF THE INVENTION

5 The present invention relates to a process for producing a polypeptide comprising at least one step of native ligation using a peptide functionalized by a selenium-containing group. The invention also relates to peptides and selenium-based compounds. The invention also relates to a synthesis kit for the synthesis of peptides.

TECHNOLOGICAL BACKGROUND

10 The synthesis of polypeptides by conventional solid-phase methods, amino acid by amino acid, is limited by low yields when the polypeptides synthesized are large. In order to overcome this limitation, it is known to assemble two polypeptides by chemical ligation in order to produce a longer polypeptide.

15 Total polypeptide synthesis is increasingly useful for preparing proteins with well-defined structures or bearing natural modifications, such as post-translational modifications, or unnatural modifications. Chemical ligation methods respond to this need; however they prove limited in their use and their industrial application.

20 Generally, in these methods, it is desirable for the link between the polypeptides assembled by ligation to be native, i.e. to correspond to the natural structure of the polypeptides.

The main native ligation method currently in existence is that of Kent and Dawson, described for example in international applications WO 96/34878 and WO 98/28434. This method is based on a chemoselective reaction between a (C-terminal) peptide thioester and a cysteinyl-peptide. Nevertheless, the main drawback of this 25 method is the production of the peptide thioesters which require complex chemical processes. These methods of the prior art are unsatisfactory because they inevitably lead to mixtures that may be difficult to separate, thus affecting the purity of the end product obtained, and to inevitable losses in yield.

30 An alternative method is the ligation known as the Staudinger reaction, described in international applications WO 01/68565 and WO 01/87920. This comprises the reaction of a phosphinothioester with an azide and the hydrolysis of the combined reagents in order to form an amide bond. However this method is difficult to apply on an industrial scale.

35 Another method, described in international application WO 2007/037812, is based on the reaction of an α -ketoacid with an N-alkoxyamine in a decarboxylative condensation reaction. However, the ketoacids are molecules which are difficult to produce and incorporate into peptides. Also, this third method is difficult to apply in

peptide synthesis laboratories that do not have the means to carry out complex organic syntheses.

The document WO 2011/051906 describes a native ligation process involving a thiol compound. Although this method is highly satisfactory, the reaction kinetics of this thiol compound are sometimes slow, in particular when the amino acids surrounding the bond between the two peptides are hindered.

A need therefore remained for a native ligation method provided with rapid reaction kinetics so as to facilitate the ligation of peptides by the formation of a bond between two hindered amino acids.

SUMMARY OF THE INVENTION

The invention is based on the development of a ligation method based on the use of a particular amide peptide, the SeEA peptide, and a cysteinyl peptide. When these two peptides are placed in contact in aqueous solution, a native peptide bond forms between the two fragments. The ligation is very efficient and takes place with excellent yields. It operates over a wide pH range.

This ligation has similarities with SEA ligation, described in the application FR 09 57 639, although there are significant differences between the properties of these two ligations and of the SeEA and SEA segments.

In fact, the kinetics of SeEA ligation are *very* significantly superior to the kinetics of SEA ligation, obtained with the sulphur-containing analogue SEA, or to the kinetics of Native Chemical Ligation (NCL) based on the use of peptide thioesters.

The kinetic superiority of SeEA does not in any way reduce the usefulness of SEA ligation. On the contrary, it complements it by allowing the formation of junctions that are difficult to produce by the other techniques of SEA ligation or NCL (thioester chemistry).

In the present application, the SEA group represents an $-N(CH_2-CH_2-S-G_4)_2$ group, with G_4 representing H or a sulphur protective group or a disulphide bond.

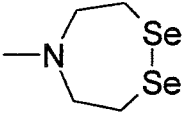
A first subject of the present invention relates to a functionalized peptide selected from:

a) the peptide of formula (I): $X_1-N(CH_2CH_2SeH)_2$; or

b) the peptide of formula (I'):

$$X_1-N \begin{array}{c} \diagup \quad \diagdown \\ \text{Se} \quad \text{Se} \end{array} \quad ;$$

in which X_1 represents a peptide fragment and the $-N(CH_2CH_2SeH)_2$ group or



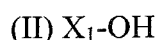
forms an amide bond with the C=O termination

of the amino acid residue of the peptide fragment X_1 which is in the C-terminal position.

According to an embodiment, X_1 comprises from 2 to 300 amino acid residues, preferably from 5 to 100 amino acid residues, more particularly preferably from 8 to 50 amino acid residues.

Another subject of the present invention relates to a process for obtaining a functionalized peptide (I) or (I') according to the invention, comprising the following steps a1 and b1:

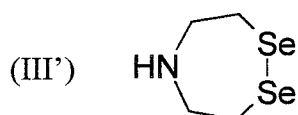
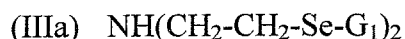
a1) peptide synthesis for providing a polypeptide of formula (II):



preferably comprising protective groups on its amine and carboxylic acid functions, with the exception of its C-terminal carboxylic acid function,

b1) functionalization of the polypeptide obtained in step a) comprising:

- the reaction of the polypeptide of formula (II) with an amine compound selected from:



in which G_1 represents a selenium protective group or a hydrogen atom,

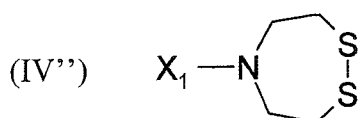
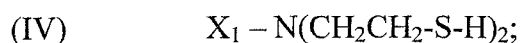
in liquid phase, in order to form the polypeptide of formula (Ia) $X_1-N(CH_2CH_2-Se-G_1)_2$, or (I');

- optionally the deprotection of the polypeptide of formula (Ia) in order to produce the compound of formula (I),

or comprising the following steps a2 and b2:

a2) the use

- of the compound of formula (IV) or (IV'')



Or

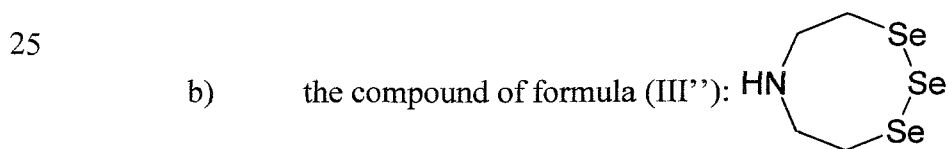
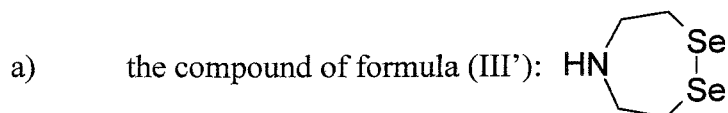
- of a peptide functionalized by a group bearing a thioester function (IV') = X_1-SR and R represents an optionally substituted aryl or alkyl group,

5 b2) the reaction between the compound of formula (IV) or (IV'') or the compound of formula (IV') with the amine compound of formula (III') or (III''):

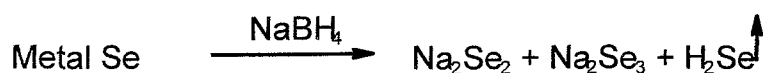


in liquid phase, in order to form the peptide of formula (I) or (I').

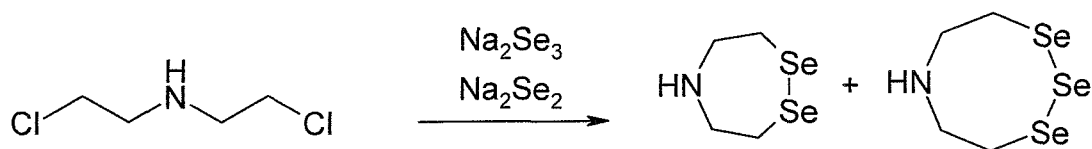
20 Another subject of the present invention relates to a compound capable of being used in the process for obtaining a functionalized peptide according to the invention, this compound being selected from:



30 The invention also relates to a process for producing the compounds of formulae (III') and (III'') according to the invention, comprising the following treatment reaction with a metal hydride illustrated in the diagram in the case of $NaBH_4$, but it is also possible to use $LiAlH_4$ or $LiBH_4$:

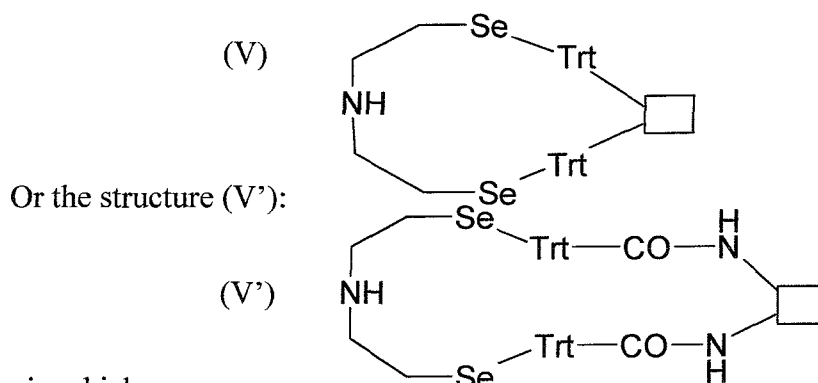


35 followed by the reaction:



Another subject of the present invention relates to a process for obtaining a functionalized peptide (I) or (I') according to the invention, comprising the following steps:

- a) Provision of a functionalized polymer resin with the structure (V):

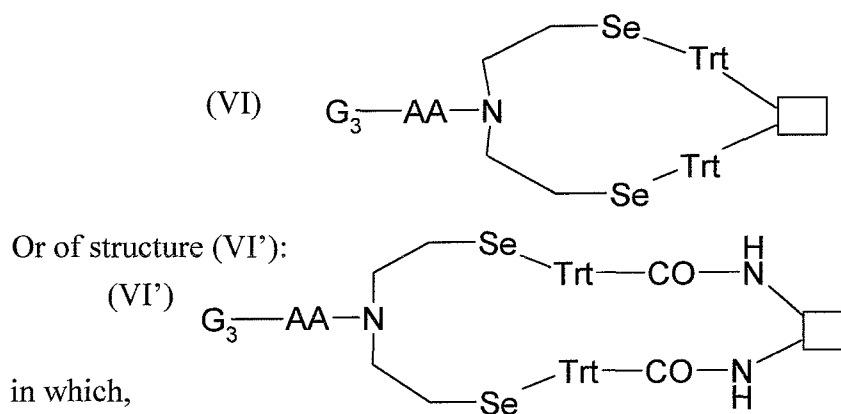


in which,

$\square$ represents a solid support,

Trt represents a triphenylmethyl group optionally substituted by one or more substituents, in particular selected from the substituents chlorine, methoxy, methyl, fluorine and cyano.

- b) Grafting of an amino acid onto the functionalized polymer resin obtained in step a) in order to produce the compound of structure (VI):



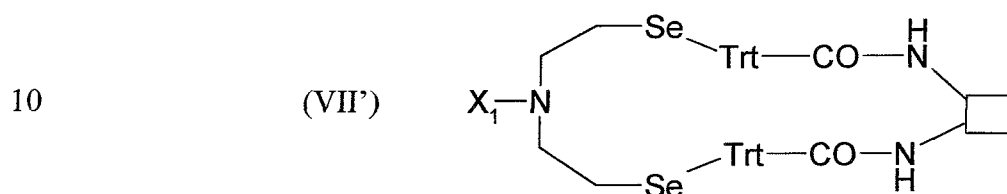
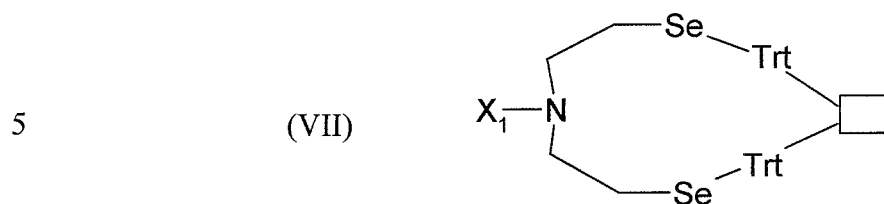
in which,

$\square$ and Trt have the same meaning as in compound (V) or (V'),

AA represents an amino acid residue optionally comprising one or more protective groups;

G_3 represents a hydrogen atom or a protective group of the N-terminal amine function of AA,

- c) Peptide synthesis starting from the amino acid AA in order to produce compound (VII) or (VII'):



in which,

□ represents a solid support,

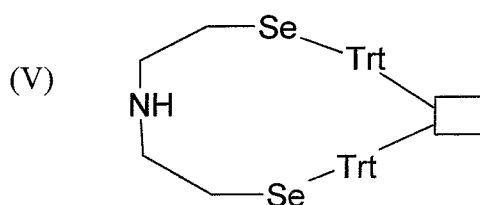
Trt represents a triphenylmethyl group optionally substituted by one or more substituents, in particular selected from the substituents chlorine, methoxy, methyl, fluorine and cyano,

X₁ represents a peptide fragment,

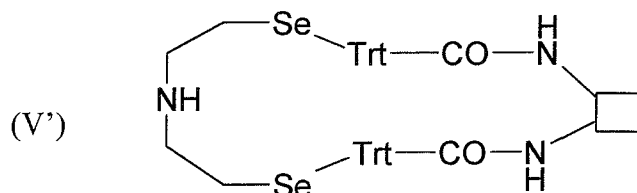
- d) Cleavage of the bond between Trt and Se in order to produce compound (I) or (I').

According to an embodiment of the process above, the grafting of an amino acid onto the polymer resin comprises bringing the functionalized resin (V) or (V') into contact with an amino acid halide or with an amino acid and an activation agent, preferably selected from HATU, PyBOP, BOP, PyBROP, more particularly preferably HATU.

Another subject of the present invention relates to a functionalized polymer resin capable of being used in the process according to the invention, with the structure (V):



Or the structure (V'):

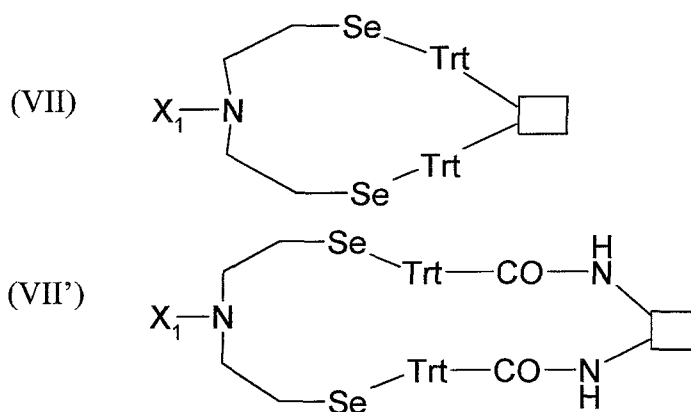


in which,

□ represents a solid support,

Trt represents a triphenylmethyl group optionally substituted by one or more substituents, in particular selected from the substituents chlorine, methoxy, methyl, fluorine and cyano.

Another subject of the present invention relates to a polymer resin comprising a peptide fragment corresponding to formula (VII) or (VII'):



in which,

□ represents a solid support,

Trt represents a triphenylmethyl group optionally substituted by one or more substituents, in particular selected from the substituents chlorine, methoxy, methyl, fluorine and cyano,

X_1 represents a peptide fragment.

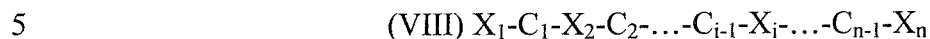
According to an embodiment, the solid support □ is selected from a polystyrene, a polyacrylamide, a polyethylene glycol, a cellulose, a polyethylene, a polyester, a latex, a polyamide, a polydimethylacrylamide, a polyethylene glycol-polystyrene copolymer, a polyethylene glycol-polyacrylamide copolymer and derivatives thereof.

Another subject of the present invention relates to a process for producing the polymer resin according to the invention, comprising:

- the provision of a solid support functionalized by -Trt or -NH-CO-Trt groups;
- reaction of the Trt function of the solid support with the amine compound of formula (III): $\text{NH}(\text{CH}_2\text{-CH}_2\text{-Se-H})_2$

Trt having the same definition as above.

Another subject of the present invention relates to a process for producing a polypeptide of formula (VIII):



in which,

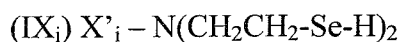
$X_1, X_2, \dots, X_i, \dots, X_n$ are peptide fragments,

$C_1, C_2, \dots, C_{i-1}, \dots, C_{n-1}$ are amino acid residues bearing a thiol or selenol function,

10 n is an integer greater than or equal to 2,

i is any integer in the range from 1 to n ,

this process comprising one or more steps of reaction between a compound of formula (IX_i):

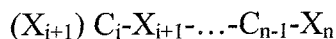


15 in which,

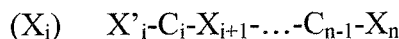
X'_1 represents X_1

X'_i represents $C_{i-1}-X_i$ for $i > 1$

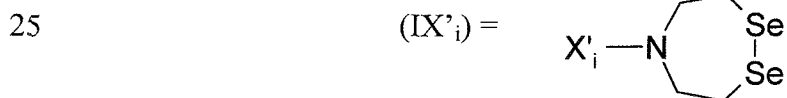
and a compound of formula (X_{i+1}):



20 in order to form a compound of formula (X_i):

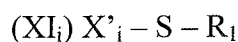


According to an embodiment, compound (IX_i) is formed from compound (IX'_i):

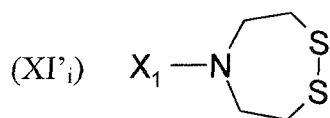


by bringing into contact with at least one compound that reduces diselenium bonds.

30 According to an embodiment, compound (IX'_i) is obtained by reaction between a compound of formula (XI_i), (XI'_i) or (XI''_i):

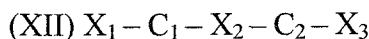


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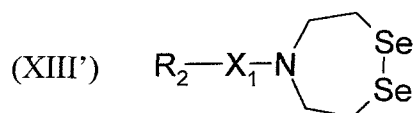
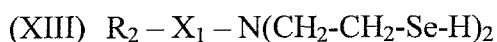
in which X'_i is a peptide fragment of the form $C_{i-1}-X_i$ for $i > 1$ and X'_1 represents X_1 , R_1 represents an optionally substituted alkyl or aryl group, and a compound of formula (III') or of formula (III'') according to the invention.

According to an embodiment, the process above is implemented for the production of a polypeptide represented by formula (XII):

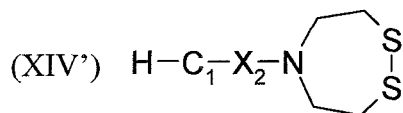
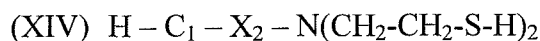


comprising the following steps:

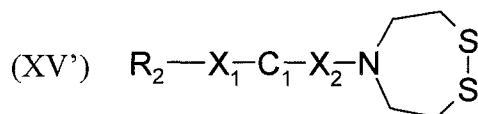
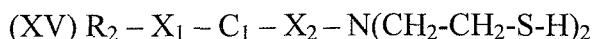
a) reaction between a compound of formula (XIII) or (XIII'):



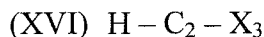
and a compound of formula (XIV) or (XIV'):



in order to form the compound of formula (XV) or (XV'):



b) reaction between a compound of formula (XV) or (XV') and a polypeptide of formula (XVI):



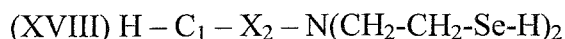
in order to form the compound of formula (XII),

in which, R_2 represents H or a protective group of the N-terminal function of X_1 ; X_1 , X_2 , X_3 represent, independently of each other, peptide fragments; C_1 and C_2 represent, independently of each other, amino acid residues bearing a thiol or selenol function.

Another subject of the present invention relates to a process for producing a cyclic polypeptide of formula (XVII):

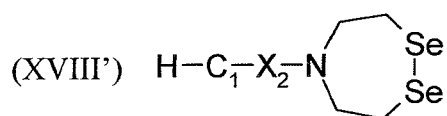


X₂ representing a peptide fragment, and C₁ representing an amino acid residue comprising a thiol or selenol function, said process comprising at least one reaction step of ligation of a polypeptide of formula (XVIII):



with itself.

According to an embodiment of the process for producing a cyclic polypeptide according to the invention, the ligation reaction is carried out by placing a polypeptide of formula (XVIII'):



in contact with at least one compound that reduces diselenium bonds.

Another subject of the present invention relates to a polypeptide synthesis kit comprising one or more peptides of formulae (IX_i) or (IX'_i) capable of being used in the ligation process according to the invention, or at least one compound (III') or (III'') according to the invention.

Advantages of the present invention are the following:

- the native ligation of the invention makes it possible to achieve the binding of difficult junctions, such as hindered junctions,
- the kinetics of native ligation of the invention are very rapid,
- the native ligation of the invention can be combined with other ligation methods having different kinetics,
- the native ligation of the invention is simple to implement,
- the SeEA peptides can be produced from a solid support,
- The SeEA peptides can be produced from SEA segments or thioester in solution
- the native ligation of the invention can be implemented starting from peptide fragments which can be prepared in solid phase or in liquid phase,
- the native ligation of the invention can be combined with other types of native ligations of the prior art in a one-pot process comprising several successive ligations,

- the native ligation of the invention can be carried out by implementing a ligation process of the prior art by the simple addition of a particular selenium-containing compound to the reaction mixture.

5 BRIEF DESCRIPTION OF THE FIGURES

Other features and advantages of the invention will become apparent on reading the following description of a preferred embodiment of the invention, given by way of example and with reference to the attached drawings.

10 Figures 1a and 1b represent respectively the HPLC chromatogram and the mass spectrum of a selenium-containing compound according to the invention.

Figures 2a and 2b represent respectively the HPLC chromatogram and the mass spectrum of a selenium-containing compound according to the invention.

Figures 3a and 3b represent HPLC chromatograms for monitoring the synthesis of a functionalized peptide according to the invention.

15 Figure 3c represents the mass spectrum of a functionalized peptide according to the invention.

Figures 4a and 4b represent respectively the HPLC chromatogram and the mass spectrum of a functionalized peptide according to the invention.

20 Figures 5a and 5b represent respectively the HPLC chromatogram and the mass spectrum of a functionalized peptide according to the invention after purification.

Figures 6a and 6b represent respectively the HPLC chromatogram and the mass spectrum of a polypeptide obtained by a ligation process according to the invention.

Figure 7 represents a diagram measuring the reaction kinetics of a ligation process according to the invention and according to the prior art.

25 Figure 8 represents a diagram measuring the reaction kinetics of a ligation process according to the invention and according to the prior art.

Figures 9a and 9b represent respectively the HPLC chromatogram and the mass spectrum of a functionalized polypeptide obtained by a process according to the invention.

30 Figures 10a and 10b represent respectively the chromatogram and the mass spectrum of a polypeptide obtained by a process according to the invention.

Figures 11a and 11b represent respectively the chromatogram and the mass spectrum of a polypeptide obtained by a process according to the invention after purification.

35 Figure 12 represents a diagram measuring the reaction kinetics of a ligation process according to the invention and according to the prior art.

Figure 13 represents an HPLC chromatogram of a solution originating from the synthesis of a functionalized peptide according to the invention.

Figures 14a and 14b represent respectively the HPLC chromatogram and the mass spectrum of a functionalized peptide according to the invention.

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DESCRIPTION OF EMBODIMENTS OF THE INVENTION

The invention is now described in more detail and non-limitatively in the following description.

By “peptide or polypeptide” is meant, in the context of the present application, a linear chain of amino acid residues (of a number greater than or equal to two) linked by peptide bonds. The “peptides or polypeptides” within the meaning of the present application can therefore for example be oligopeptides, peptides or proteins according to the conventional acceptance of these terms. The amino acid residues present in the polypeptides according to the invention can be selected from the proteinogenic or non-proteinogenic amino acid residues. Preferably, they are selected from the twenty proteinogenic amino acid residues and selenocysteine.

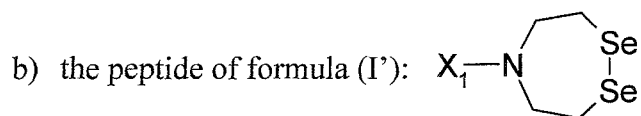
The polypeptide notation is carried out from the N-terminal end towards the C-terminal end. The usual one-letter or three-letter code is used for notation of the amino acid residues present along the polypeptide chain. An amino acid residue is a polypeptide fragment of formula -NH-(CH-R)-(C=O)- , in which R represents a side chain, which differs from one amino acid to another.

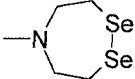
By “peptide fragment” is meant, in the context of the present application, a portion of polypeptide comprising at least one amino acid residue. A peptide fragment, within the meaning of the present application, can therefore be for example: a sequence of amino acid residues (such as -AHG- or -Ala-His-Gly-) if the peptide fragment comprises neither the N-terminal end nor the C-terminal end of the polypeptide; or a sequence of amino acid residues with a group at its N-terminal end (such as H-AHG- or H-Ala-His-Gly-) if the peptide fragment comprises the N-terminal end of the polypeptide; or a sequence of amino acid residues with a group at its C-terminal end (such as -AHG-OH or -Ala-His-Gly-OH) if the peptide fragment comprises the C-terminal end of the polypeptide.

Functionalized peptide

A first subject of the invention relates to a peptide functionalized by a selenium group selected from:

a) the peptide of formula (I): $\text{X}_1\text{-N(CH}_2\text{CH}_2\text{SeH)}_2$;



5 in which X_1 represents a peptide fragment and the $-N(CH_2CH_2SeH)_2$ group or  forms an amide bond with the C=O termination of the amino acid residue of the peptide fragment X_1 which is in the C-terminal position.

10 The peptide fragment X_1 is in the form $Y_1-AA_1-AA_2-...-AA_n$ in which Y_1 is an N-terminal group, preferably a hydrogen atom, but optionally also any group being substituted for the primary or secondary amines known to a person skilled in the art, for example an acyl group and in particular an acetyl group; n is an integer greater than or equal to 2; $AA_1, AA_2, ..., AA_n$ represent, independently of each other, amino acid residues.

15 The polypeptide of formula (I) or (I') preferably comprises from 2 to 300 amino acid residues, preferably from 5 to 100 amino acid residues, more particularly preferably from 8 to 50 amino acid residues.

20 The polypeptides of formula (I) or (I') preferably comprise only amino acid residues selected from the twenty proteinogenic amino acid residues and selenocysteine. However, according to a particular embodiment, the polypeptides of formula (I) or (I') comprise one or more non-proteinogenic amino acid residues.

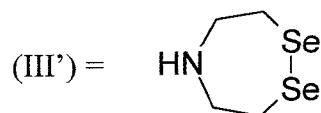
The amino acid residues of the polypeptides of formula (I) or (I') can optionally be protected by side chain protective groups.

25 Amine compound of formula (III') and (III'')

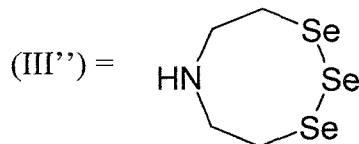
Another subject of the invention relates to the amine compounds of formula (III') and (III'') below and their production process.

Compounds (III') and (III'') correspond to the formulae:

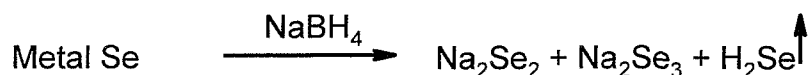
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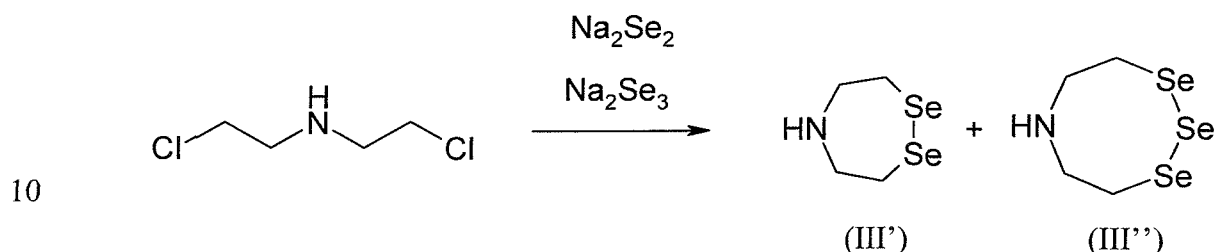
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Compounds (III') and (III'') can be obtained according to the process comprising the following step a):



5 followed by the following step b):



The first step above (step a) is preferably carried out under a nitrogen atmosphere. Preferably, the reagents are introduced cold, then the mixture is taken to reflux.

15 According to another embodiment, in step a), NaBH₄ is replaced with another metal hydride, such as LiAlH₄ or LiBH₄. In the case of the two metal hydrides mentioned, the compounds Li₂Se₂ and Li₂Se₃ are formed. These two compounds Li₂Se₂ and Li₂Se₃ are then used in the following step b).

Preferably, the second step (step b) is carried out at ambient temperature.

20 Preferably, the organic phase comprising the products (III') and (III'') is purified at the end of the process, for example, by high performance liquid chromatography.

Process for obtaining compounds (I) and (I') in liquid phase

25 Another subject of the invention relates to a process for obtaining the functionalized polypeptide (I) or (I') in liquid phase, this process comprises a first step of peptide synthesis and a second step of functionalization.

The first step (peptide synthesis step) makes it possible to obtain the polypeptide of formula:



This peptide synthesis step can be carried out according to any method known to a person skilled in the art. It can in particular be carried out in liquid phase or, preferably, in solid phase.

35 In principle, peptide synthesis comprises a succession of couplings of amino acids starting from a primer (initial amino acid or peptide fragment resulting from the preceding amino acid additions) and from deprotections. More precisely, the peptide synthesis can successively comprise:

- (a) the provision of a peptide fragment with an unprotected N-terminal end, and of an amino acid protected at its N-terminal end;
- (b) the establishment of a peptide bond between the amino acid and the peptide fragment, at the N-terminal end of said peptide fragment;
- 5 (c) the deprotection of the N-terminal end of the linked amino acid, in order to provide the peptide fragment of the following step (a).

During the different coupling reactions, it is advantageous to use an activator compound, in particular a carbodiimide (for example dicyclohexylcarbodiimide or diisopropylcarbodiimide) in the presence of a triazole (for example 1-hydroxy-10 benzotriazole or 1-hydroxy-7-azabenzotriazole), or a phosphonium or uronium salt of a non-nucleophilic anion (for example HBTU, HATU, TBTU or PyBOP); or also to use activated amino acids in the form of acid halides, for example acid fluorides.

During the different coupling reactions, the N-terminal ends of the amino acids are advantageously protected by a protective group, preferably a Fmoc group (9H-fluoren-9-ylmethoxycarbonyl) or also a t-Boc group (tert-butoxycarbonyl) or NSC group (2-(4-nitrophenylsulphonyl)ethoxycarbonyl).

Similarly, the side chains of the amino acid residues are preferably protected during the different coupling reactions by one or more appropriate protective groups, for example a tert-butyl group for the chains bearing a carboxylic acid function.

20 In this case, once the last amino acid coupling reaction has been carried out, a deprotection reaction of the side chains can be provided. It should however be noted that it may be preferable to retain all of the protections (with the exception of an optional protection of the COOH function at the C-terminal end) for the functionalization step.

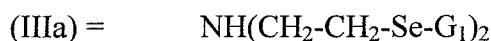
25

Functionalization

The functionalization step comprises successively:

- Optionally, the protection of the N-terminal end of the polypeptide of formula (II) with a protective group, preferably selected from the family of the carbamates, amides or alkyl groups, and in particular a tert-butoxycarbonyl group (t-Boc), Fmoc (9H-fluoren-9-ylmethoxycarbonyl), trifluoroacetyl or triphenylmethyl.
- 30 Optionally also, the protection of the functions of the side chains of the amino acid residues of the polypeptide of formula (II), and quite particularly of the amine functions (preferably by means of the protective groups above) and carboxylic acid functions (by means for example of tert-butyl groups).

- Alternatively, and according to a simpler embodiment, it is possible to provide the polypeptide of formula (II) directly in a form in which all of the amine and carboxylic acid functions are protected, by providing selective deprotection of the COOH function of the C-terminal end.
- The liquid phase coupling reaction of the polypeptide of formula (II) with an amine compound selected from compounds (IIIa) and (III') where:



in which G₁ represents a selenium-protective group,

- Optionally the deprotection of the polypeptide.

According to an embodiment, the protective group G₁ of compound (IIIa) is selected from the groups of triphenylmethyl and benzyl type and in particular from para-methylbenzyl or para-methoxybenzyl.

In the case where compound (IIIa) is used as starting reagent for coupling with compound (II), a subsequent step of deprotection of the group G₁ is provided in order to obtain compound (I) or (I'). Preferably, the deprotection is carried out by treatment with an acid such as trifluoroacetic acid or hydrofluoric acid, or an oxidant such as iodine.

An activator is advantageously present during the coupling reaction, for example benzotriazol-1-yl-oxytrypyrrolidinophosphonium hexafluorophosphate (PyBOP) or bromo-trispyrrolidinophosphonium hexafluorophosphate (PyBROP) or the C-terminal amino acid itself is activated in the form of a halogenated derivative (in particular in the form of an amino acid fluoride or amino acid chloride), the latter being able to be pre-formed or formed *in situ* using appropriate reagents known to a person skilled in the art. Among the amino acid halides, the amino acid fluorides are preferred, pre-formed by reaction with 1,3,5-trifluorotriazine or formed *in situ* using TFFH (tetramethylfluoroformamidinium hexafluorophosphate).

Generally, it is also possible to envisage any reagent allowing the activation of the carboxylic acid function of the amino acid known to a person skilled in the art such as HBTU, TBTU, HATU, BOP etc. (reference may be made for example to *Chemical approaches to the synthesis of peptides and proteins* by Lloyd-Williams, P., Albericio, F., Giralt, E., 1997, CRC Press). PyBOP, PyBROP, BOP (benzotriazole-1-yl-oxy-tris-(dimethylamino)-phosphonium) hexafluorophosphate or more generally the phosphoniums are preferred.

The functionalized polypeptide (I) or (I') according to the invention can be obtained according to a second variant.

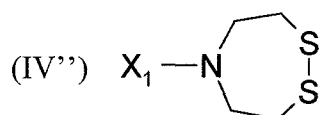
5 The SeEA segments can be formed separately by exchange starting from an SEA segment of $X_1-N(CH_2CH_2-S-G_4)_2$ or X_1-S-R_1 thioester type using the reagents of structure (III') or (III''), G_4 representing a hydrogen atom, a thiol function protective group or a disulphide bond and R_1 representing a hydrogen atom or an optionally substituted alkyl or aryl group.

10 The SeEA segments can also be formed *in situ*, during the exchange ligation starting from an SEA segment of X_1-SEA or X_1-SR_1 thioester type using reagents (III') or (III''). The segment of H-Cys- X_2 type present in solution consumes the X_1 -SeEA segment formed by the exchange. In general, the X_1 -SeEA segment formed *in situ* is not observed since it reacts very rapidly with the H-Cys- X_2 segment. Due to the rapidity of the exchange and of the SeEA ligation, this results in a very
15 significant acceleration of the ligation, once the reagent (III') or (III'') is introduced into the ligation reaction medium.

Thus, according to a second variant, compounds (I) and (I') are obtained by the following successive steps:

a2) the provision

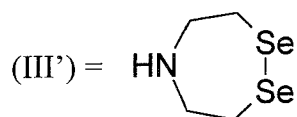
20 - of a peptide functionalized on its C-terminal end by a group bearing a thiol function corresponding to formula (IV) or (IV''):



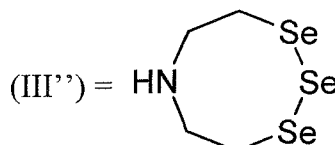
25

- of a peptide functionalized by a group bearing a thioester function
(IV') = X_1-SR and R represents an optionally substituted aryl or alkyl group,

30 b2) the liquid-phase reaction between the compound of formula (IV) or the compound of formula (IV'') or the compound of formula (IV') and an amine compound of formula (III') or (III''):



35



optionally in the presence of a compound that reduces the disulphide bond in the case where compound (IV'') is used.

The compound of formula (IV) or of formula (IV'') can in particular be obtained according to the protocol described in the document WO 2011/051906.

In formula (IV'), the R group preferably represents an optionally substituted C1-C12 alkyl or an optionally substituted C6-C12 aryl. Such compounds are described in particular by WO96/34878 and WO98/28434.

According to an embodiment, the compound that reduces the disulphide bonds can be selected from the thiol compounds, such as 4-mercaptophenylacetic acid (MPAA), dithiothreitol (DTT), thiophenol and its derivatives, an alkylthiol (in particular benzylmercaptan), from the selenol compounds such as selenophenol or diphenyl selenide or from the phosphines such as tris(2-carboxyethyl)phosphine (TCEP), or mixtures thereof.

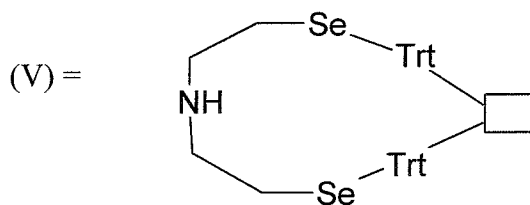
At the end of the previous coupling reaction, compound (I') is obtained. Compound (I) is obtained by placing compound (I') in contact with a compound that reduces diselenium or triselenium bonds.

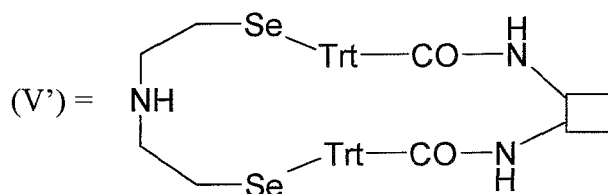
Among the known compounds that reduce diselenium or triselenium bonds, there may be mentioned: tris(2-carboxyethyl)phosphine (TCEP), the reducing thiols such as 4-mercaptophenylacetic acid (MPAA), thiophenol, benzylmercaptan, dithiothreitol (DTT), selenophenol or diphenyl selenide plus a diselenium bond reducing agent, metallic zinc, and hydrides such as NaBH₄, LiBH₄, NaBH₃CN, NaBH(OAc)₃.

At the end of the functionalization step, the polypeptide of formula (I) or (I') is obtained. It is advantageous to provide at this step a compound purification step, for example by liquid chromatography.

Functionalized polymer resin

Another subject of the invention relates to the functionalized polymer resins corresponding to structure (V) or (V'):





in which,

□ represents a solid support,

Trt represents a triphenylmethyl group optionally substituted by one or more substituents, in particular selected from the substituents chlorine, methoxy, methyl, fluorine and cyano.

These polymer resins are then used to prepare the functionalized SeEA peptide.

By way of solid support □, a soluble or insoluble polymer is used, the insoluble polymer preferably being in the form of particles (beads). It is possible for example to use a resin based on polystyrene (preferably) or also based on polyacrylamide, polyethylene glycol, cellulose, polyethylene, polyester, latex, polyamide, polydimethylacrylamide, polyethylene glycol-polystyrene copolymer, polyethylene glycol-polyacrylamide copolymer or also a resin derived therefrom.

In order to prepare the functionalized polymer resin (V) or (V'), a solid support □ bearing a suitable functionalization, □-Z is used, in which Z is a linker group. Preferably, the solid support has chloro-triphenylmethyl (or chlorotrityl) groups, where the triphenylmethyl is optionally substituted by one or more substituents, in particular selected from the substituents chlorine, methoxy, methyl, fluorine and cyano.

By way of examples of a functionalized solid support □-Z, there may be mentioned the polystyrene resins with trityl chloride, 2-chlorotrityl chloride, 4-methyltrityl chloride or 4-methoxytrityl chloride linker groups. Such solid supports are commercially available, for example from Glycopep.

According to another embodiment, the functionalized solid support □-Z has linker groups which are groups of the trityl-alcohol type, i.e. OH-Trt-CO-NH- groups, where the triphenylmethyl (Trt) is optionally substituted by one or more substituents, in particular selected from the substituents chlorine, methoxy, methyl, fluorine and cyano. Solid supports with such linker groups have been described for example in Quibell, JACS 1995, 117, 11656-11668, and are commercially available, for example in the ChemMatrix® range of solid polyethylene glycol supports.

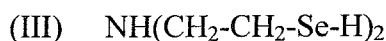
The use of solid supports of this type requires a preliminary step of activation of the solid support:

- either with a brominated agent (in particular acetyl bromide) in order to modify the linker groups in the Br-Trt-CO-NH- form;

- or with a chlorinated agent (in particular oxalyl chloride) in order to modify the linker groups in the Cl-Trt-CO-NH- form;
- or with an acid such as trifluoroacetic acid, in order to modify the linker groups in the H_2O^+ -Trt-CO-NH- form;
- 5 – or with $\text{BF}_3 \cdot \text{Et}_2\text{O}$, in order to modify the linker groups in the $\text{BF}_3 \cdot \text{HO}$ -Trt-CO-NH- form. according to Singh, S. et al., *J. Org. Chem.* **2004**, 69, 4551-4554.

An activation step of this type is described for example in Harre et al., *Reactive & functional polymers* 1999, 41, 111-114.

10 The preparation of the functionalized polymer resin is carried out by coupling the amine compound of formula:



15 with the above functionalized solid support, $\square\text{-Z}$. The amine compound (III) can itself be obtained by deprotection of the above amine compound of formula (IIIa), in which G_1 is a selenium protective group.

The coupling is carried out in an acid medium in order to limit the risks of unmasking the secondary amine, which would lead to secondary reactions.

20 The excess trityl chloride groups are preferably neutralized, for example with methanol. It is then important to add a base in order to neutralize the HCl formed.

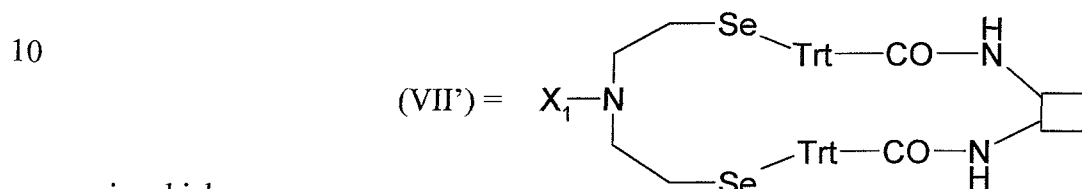
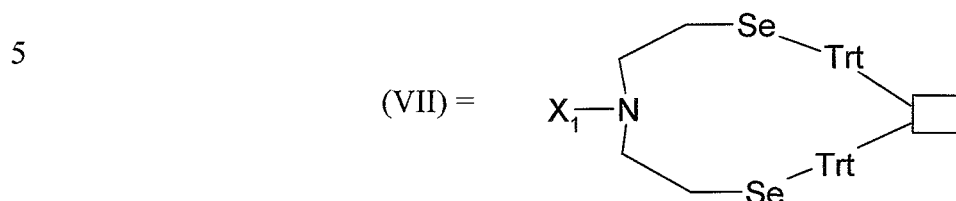
It can be provided that the preparation of the functionalized polymer resin forms an integral part of the solid-phase production process of the polypeptides of formula (I) and (I').

25 Alternatively, the functionalized polymer resin can be prepared in advance and separately, in order to be ready for use within the context of the solid phase production process of the peptides of formula (I) and (I').

30 The use of solid supports of this type can make it possible to use resin backbones of the polyethylene glycol or polyethylene glycol-polystyrene type, more suitable for the preparation of long polypeptides than resins of the polystyrene type.

35 Preferably, the resin particles have a $\text{Dv}50$ comprised between 1 and 1000 μm . The $\text{Dv}50$ is defined as being the 50th centile of the particle size distribution, i.e. 50% of the particles have a size smaller than the $\text{Dv}50$ and 50% have a size greater than the $\text{Dv}50$. Generally, the $\text{Dv}50$ is characteristic of the granulometric profile (volumetric distribution) of the particles, and it can be determined by laser granulometry (in the case of a size smaller than 200 μm), or by sieving (in the case of a size greater than 200 μm).

Another subject of the invention relates to a peptide grafted onto a functionalized polymer resin with the structure (VII) or (VII'):



in which,

represents a solid support,

15 Trt represents a triphenylmethyl group optionally substituted by one or more substituents, in particular selected from the substituents chlorine, methoxy, methyl, fluorine and cyano,

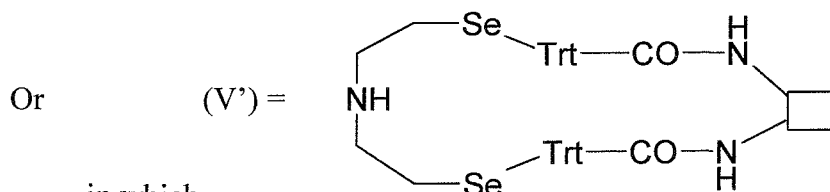
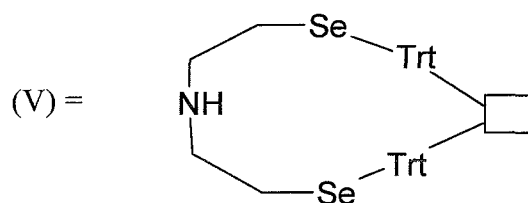
X₁ represents a peptide fragment functionalized on its C-terminal end.

20 Solid-phase process for obtaining the peptides of formulae (I) and (I')

Another subject of the invention relates to a solid phase production process of the previously described peptides (I) and (I').

This solid-phase process comprises a functionalization phase followed by a peptide synthesis phase.

25 The first step comprises the provision of a functionalized polymer resin with the structure:

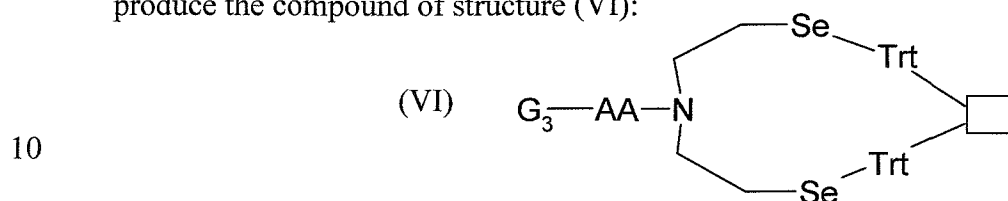


35 in which,

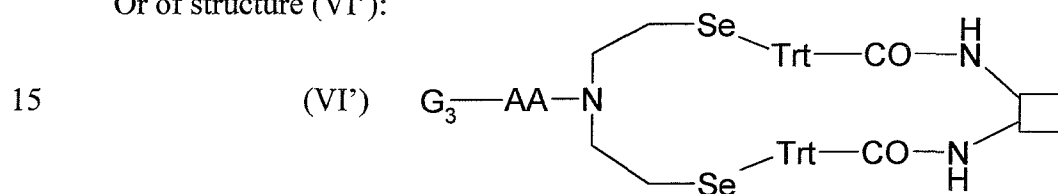
represents a solid support,

Trt represents a triphenylmethyl group optionally substituted by one or more substituents, in particular selected from the substituents chlorine, methoxy, methyl, fluorine and cyano.

5 The second step of the process comprises the grafting of an amino acid $G_3 - AA$ onto the functionalized polymer resin (V) or (V') obtained in the first step in order to produce the compound of structure (VI):



Or of structure (VI'):



in which,

20  and Trt have the same meaning as in compound (V) or (V'),
AA represents an amino acid residue optionally comprising one or more protective groups;
 G_3 represents a hydrogen atom or a protective group of the N-terminal amine function of AA.

25 Preferably the amino acid AA is protected at its N-terminal end by a protective group G_3 which is labile in the presence of a base, preferably selected from the 2-(4-nitrophenylsulphonyl)ethoxycarbonyl (NSC) or Fmoc groups.

30 Preferably, the amino acid AA comprises protective groups on all or some (preferably all) of the functions present on its side chain, and in particular the carboxylic acid (aspartic and glutamic acids), amine (for lysine), alcohol (for serine or threonine), phenol (for tyrosine), thiol or selenol (for cysteine or selenocysteine), guanidine (for arginine) and imidazole (for histidine) functions. Such protective groups are known to a person skilled in the art. Reference may be made for example to the reference work *Protective groups in organic synthesis*, 2nd edition, T. Greene and P. Wuts, John Wiley & Sons, Inc.

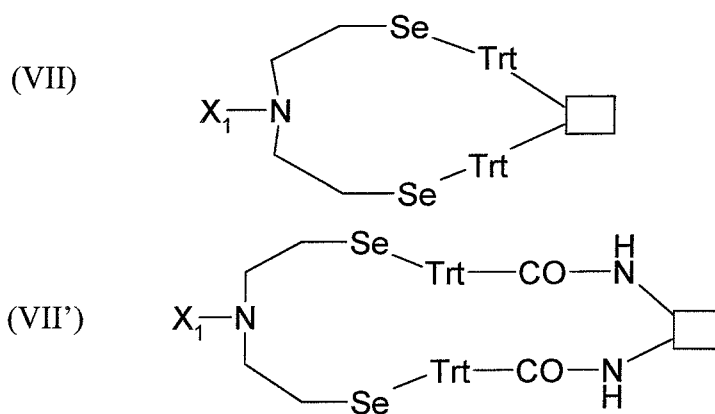
35 Preferably, the amino acid is activated in the presence of HATU (2-(1H-7-Azabenzotriazol-1-yl)--1.1,3,3-tetramethyl uronium hexafluorophosphate Methanaminium), of PyBOP or of BOP or more particularly preferably in the

presence of PyBROP, or also in the form of a halide, in particular a fluoride (i.e. a fluorine atom is linked to the acyl group of the amino acid residue).

The amino acid reacts with the secondary amine function present on the functionalized solid support in order to form an amide bond. After coupling of the amino acid, the latter may optionally be deprotected.

Thus, in this embodiment, the functionalization step consists of creating a primer solid support from a previously functionalized solid support.

The third step comprises peptide synthesis starting from the amino acid AA grafted onto the polymer resin in order to produce compound (VII) or (VII'):



The peptide synthesis step is carried out analogously to what has already been described previously, relating to liquid-phase peptide synthesis, the initial primer being provided by the amino acid AA grafted onto the solid support.

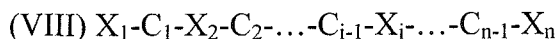
Finally, once the peptide synthesis is completed, a reaction of separation (or cleavage) of the polypeptide from its support is provided, by cleavage of the bond between Trt and Se in order to produce compound (I) or (I'). Preferably, the cleavage is carried out with a trifluoroacetic acid solution comprising the carbocation traps known to a person skilled in the art such as water, triisopropylsilane, thiophenol, thioanisole, anisole, dimethyl sulphide, etc.

Suitable deprotections of the side chains of the amino acids may also optionally be provided in order to obtain a functionalized peptide of formula (I) or (I').

As for the liquid-phase process, it is advantageous to provide at this stage a compound purification step, for example by liquid chromatography.

Native ligation process

Another subject of the invention relates to a process for producing a polypeptide of formula (VIII):



in which,

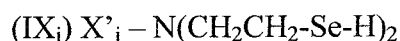
$X_1, X_2, \dots, X_i, \dots, X_n$ are peptide fragments,

$C_1, C_2, \dots, C_{i-1}, \dots, C_{n-1}$ are amino acid residues bearing a thiol, or selenol function

5 n is an integer greater than or equal to 2,

i is any integer in the range from 1 to n ,

this process comprising one or more steps of reaction between a compound of formula (IX_i) :

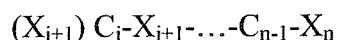


10 in which:

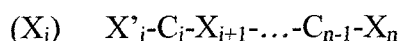
X'_1 represents X_1

and X'_i represents $C_{i-1}-X_i$ for $i > 1$

and a compound of formula (X_{i+1}) :



15 in order to form a compound of formula (X_i) :



The residue C_i is an amino acid residue comprising a thiol or selenol function. This thiol or selenol function can be in particular a beta-amino thiol function or beta-amino selenol function (in which case the C_i residue preferably represents the cysteine residue or the selenocysteine residue), a gamma-amino thiol or gamma-amino selenol function (in which case the C_i residue preferably represents the homocysteine or homoselenocysteine residue).

The invention makes it possible to produce polypeptides by successively carrying out several ligation reactions. This can prove appropriate in order to obtain large-sized polypeptides, for example polypeptides comprising more than approximately 100 amino acid residues. In fact, in such cases the production of polypeptides of formulae (IX_i) and (X_i) by direct synthesis can have a low yield, and it is therefore advantageous to use two or more than two successive ligations, in order to have to directly synthesize only polypeptides comprising for example less than approximately 50 amino acid residues.

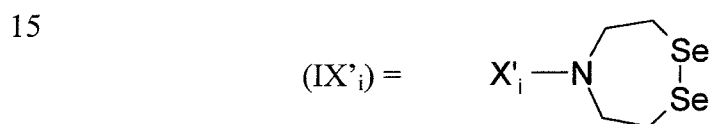
By way of example, the use of two successive ligations makes it possible to obtain a polypeptide comprising approximately 150 amino acid residues without having to directly synthesize polypeptide comprising more than approximately 50 amino acid residues; the use of three successive ligations makes it possible to obtain a polypeptide comprising approximately 200 amino acid residues without having to directly synthesize polypeptide comprising more than approximately 50 amino acid residues.

The polypeptide of formula (VIII) can be obtained by using several ligation methods. Other ligation methods are for example methods involving thioesters or involving particular sulphur-containing compounds as described in WO 2011/051906.

5 The process for producing the polypeptide of formula (VIII) according to the invention comprises at least one reaction step between a compound of formula (IX_i) and a compound of formula (X_i).

Thus, the process according to the invention makes it possible to obtain a polypeptide of formula (VIII); each X_i, for i an integer between 1 and n, being a peptide fragment; and each C_i, for i an integer between 1 and n, being an amino acid residue comprising
10 a thiol or selenol function, and in particular a cysteine or selenocysteine residue according to a particular embodiment, using n-1 successive ligations.

According to an embodiment of the invention, compound (IX_i) can be obtained from compound (IX'_i):



in the presence of at least one compound that reduces diselenium and/or triselenium bonds. In fact, the polypeptide of formula (IX'_i) is then reduced *in situ* and provides
20 the polypeptide of formula (IX_i) for the ligation reaction.

According to an embodiment, the reducing compound is selected from tris(2-carboxyethyl)phosphine (TCEP), 4-mercaptophenylacetic acid (MPAA), benzylmercaptan, thiophenol, DTT, sodium 2-mercaptoethylsulphonate (MESNa), selenophenol.

25 Analogously to what was described previously for compounds (I) and (I'), according to an embodiment, the compounds (IX_i) and (IX'_i) can be formed by exchange, starting from an SEA segment of structure X'_i – N(CH₂CH₂–S–G₅)₂ or from a thioester of structure X'_i – S – R₁ using the selenium-containing compounds of structure (III') or (III''), optionally in the presence of compounds that reduce
30 disulphide and/or diselenium bonds, G₅ representing H, a protective group of the thiol function or a disulphide bond and R₁ representing H or a protective group of sulphur.

In the case where i is equal to 1, compounds (IX_i) and (IX'_i) correspond to compounds (I) and (I') respectively.

35 Generally, the ligation reaction starting from the polypeptide of formula (IX'_i) as reagent can be more practical to implement than the ligation reaction directly with the polypeptide of formula (IX_i). In fact, the polypeptide of formula (IX_i) has a natural tendency to oxidize to the polypeptide of formula (IX'_i), in particular under the action

of the oxygen in the air. For example, it is possible that the open-form polypeptide (IX_i) oxidizes over time when it is stored in the lyophilized form. In other words, a preparation of the polypeptide of formula (IX_i) generally necessarily partly contains the polypeptide of formula (IX'_i). The presence of these two forms can complicate characterization and purification. This is why it may be simpler to implement the ligation reaction by placing the polypeptide of formula (IX'_i) in contact with the polypeptide of formula (X_i), the polypeptide with the cyclic termination of formula (IX'_i) being reduced *in situ* to the open polypeptide of formula (IX_i).

For the same reasons, even when the ligation reaction is carried out directly by placing the polypeptide of formula (IX_i) in contact with the polypeptide of formula (X_i), it is preferable to use one or more of the above-mentioned compounds that reduce diselenium bonds during the reaction.

The ligation reaction preferably takes place in liquid phase, and in particular in an aqueous medium, for example in a phosphate buffer. Preferably, this reaction is carried out at a pH comprised between 4 and 8.5, more particularly preferably at a pH comprised between 5 and 7.5 and ideally at a pH close to 5.5.

The ligation reaction is preferably carried out at a temperature comprised between 0 and 50°C, and ideally at a temperature of approximately 37°C. The duration of the reaction is adjusted as a function of the selection of the reagents and other conditions of the reaction. The appropriate duration can also be adjusted according to the results of a liquid chromatography– mass spectrometry analysis during the reaction. The appropriate duration is typically from a few hours to a few days.

Each of the polypeptides of formula (IX_i) and (X_i) is preferably present in a concentration comprised between 0.01 and 50 mM, during the reaction. The molar concentration ratio between the polypeptides of formula (IX_i) and (X_i) during the reaction is preferably comprised between 2:3 and 3:2.

The ligation reaction described above can be followed by a step of purification of the polypeptide of formula (X_i) obtained, for example by liquid chromatography or by any other usual technique.

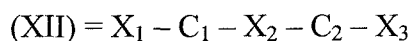
The ligation kinetics according to the present invention are significantly superior to the ligation kinetics obtained with sulphur-containing compounds described in the document WO 2011/051906 or with thioesters such as those taught by Kent and Dawson.

Due to the rapidity of the SeEA ligation, it can be carried out in the presence of other groups capable of participating in a native ligation reaction, such as the SEA or thioester segments.

The combination of the SeEA and SEA ligations constitute a very efficient tool for the total synthesis of proteins. For the production of a protein which must be synthesized by ligation of several fragments it is possible for example to implement SEA ligation for the easy junctions, and SeEA ligation for the difficult junctions. In fact, SeEA ligation makes it possible to easily form hindered junctions, for example in 4 hours for the Val-Cys junction, whereas this type of junction requires ligation times of 96 hours in the case of SEA ligation or 8 hours in the case of NCL ligation.

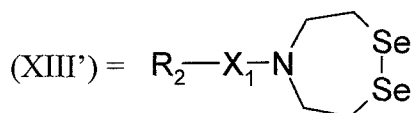
Another useful way of using SeEA and SEA ligations is one-pot sequential SeEA-SEA ligation. The latter can be implemented in order to link three fragments without isolation and intermediate purification (known as "one-pot reaction") when the junctions have an analogous steric hindrance. An embodiment example of this variant is explained below.

According to an embodiment of the invention, the ligation process of the invention allows the production from three peptide fragments, of a polypeptide represented by formula (XII):

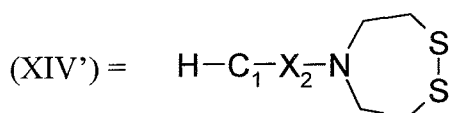
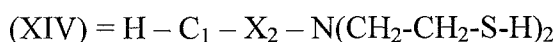


said polypeptide being obtained by a process comprising the following steps:

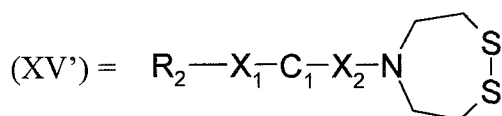
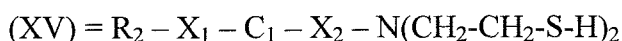
a) reaction between a compound of formula (XIII) or (XIII'):



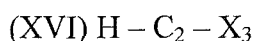
and a compound of formula (XIV) or (XIV'):



in order to form the compound of formula (XV) or (XV'):



b) reaction between a compound of formula (XV) or (XV') and a polypeptide of formula (XVI):



in order to form the compound of formula (XII),

in which,

R₂ represents a protective group of the N-terminal function of X₁,

X₁, X₂, X₃ represent, independently of each other, peptide fragments,

C₁ and C₂ represent, independently of each other, amino acid residues bearing a thiol or selenol function.

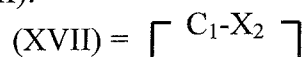
5 In a manner analogous to what is described previously for obtaining compounds (I) and (I'), the above compounds (XIII) and (XIII') can be formed *in situ*, by exchange starting from reagents of the (IIIa), (III') or (III'') type in the presence of SEA compounds of structure R₂ – X₁ – SEA or thioester of structure R₂ – X₁ – S – R₁, in which R₁ represents H or an optionally substituted alkyl or aryl, thus making it
10 possible to accelerate the second ligation step.

This one-pot three-segment ligation reaction can be implemented in order to link three fragments without isolation and intermediate purification (known as “one-pot reaction”) when the junctions have an analogous steric hindrance. As the hindrance of the junctions is analogous, the ligation between compound (XIII) or
15 (XIII') and compound (XIV) or (XIV') is more rapid than the cyclization of compound (XIV) or (XIV') by intramolecular SEA ligation, or the polymerization of the segment (XIV) or (XIV') by intermolecular SEA ligation. The intramolecular or intermolecular SEA ligations are described in WO 2011/051906.

20 Production of a cyclic polypeptide by native self-ligation

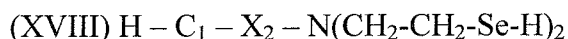
The principles used for native ligation described above can also be used to produce cyclic polypeptides, by native self-ligation of a polypeptide (ligation of one end of the polypeptide with the other end of the same polypeptide).

Another subject of the invention relates to a process for producing a cyclic
25 polypeptide of formula (XVII):



X₂ representing a peptide fragment, and C₁ representing an amino acid residue comprising a thiol or selenol function,

said process comprising at least one reaction step of ligation of a polypeptide of
30 formula (XVIII):



with itself.

The polypeptide of formula (XVIII) preferably comprises between 2 and 300 amino acid residues, preferably between 5 and 100 amino acid residues, more
35 particularly preferably between 8 and 50 amino acid residues.

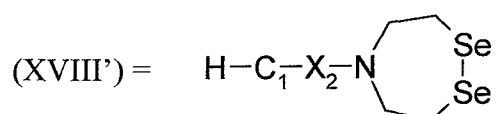
The polypeptide of formula (XVIII) comprises a hydrogen atom and a C₁ residue at the N-terminal end, C₁ having the meaning indicated above. X₂ here represents a

peptide fragment of the form $AA_1AA_2...AA_n$ where n is an integer greater than or equal to 1 and each AA_i represents an amino acid residue.

The polypeptide of formula (XVIII) preferably comprises only amino acid residues selected from the twenty proteinogenic amino acid residues and selenocysteine. However, according to a particular embodiment, the polypeptide of formula (XVIII) comprises one or more non-proteinogenic amino acid residues, other than selenocysteine.

The amino acid residues of the polypeptide of formula (XVIII) can optionally be protected by side chain protective groups.

It is possible to carry out the above ligation reaction by placing the polypeptide of formula (XVIII'):



in contact with at least one compound that reduces diselenium bonds, which is preferably as described above. In fact, the polypeptide of formula (XVIII') is then reduced *in situ* and provides the polypeptide of formula (XVIII) for the ligation reaction.

Generally, even when the ligation reaction is carried out directly from the polypeptide of formula (XVIII), it is preferable to use one or more of the above-mentioned compounds that reduce diselenium bonds during the reaction.

In general, the cyclization of the polypeptide of formula (XVIII) is not impeded by concurrent multimerizations, if the reaction is carried out under sufficiently dilute conditions. It is possible for example to use a concentration of polypeptide of formula (XVIII) comprised between 0.01 and 50 mM, typically of approximately 1 mM (or optionally between 0.01 and 0.1 mM in the case of serious risks of multimerizations).

Moreover, the preferred conditions for implementation of the ligation reaction are the same as those described above for the reaction starting from the polypeptides of formula (IX) and (X).

The ligation reaction described above can be followed by a step of purification of the cyclic polypeptide of formula (XVII) obtained, for example by liquid chromatography or by any other usual technique.

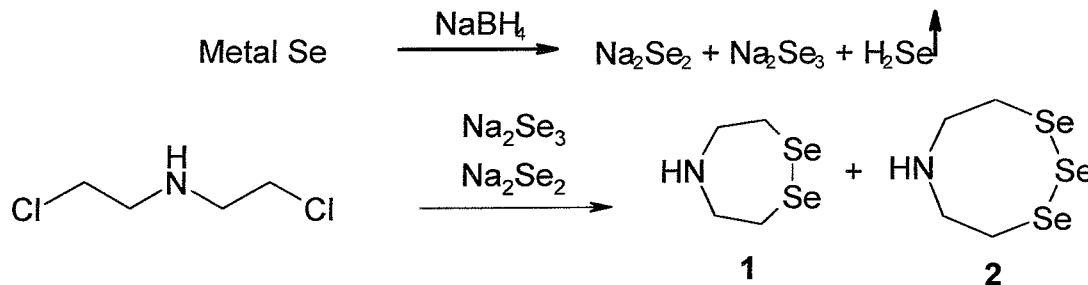
Synthesis kit

Another subject of the invention relates to a polypeptide synthesis kit comprising one or more peptides of formulae (IX_i) or (IX'_i) , or at least one compound

of formula (III') or (III''), and optionally one or more coupling, protection, deprotection reagents, or solvents.

EXAMPLES

5 Example 1: Synthesis of compounds **1** (Se₂EA) and **2** (Se₃EA)



Under nitrogen flushing, absolute ethanol V = 19 mL is added dropwise, under stirring, to a mixture of selenium Se (940 mg, 11.9 mmol, 1.5 eq) and sodium borohydride (NaBH₄) (300 mg, 7.9 mmol) cooled down in an ice bath. After a vigorous reaction, the reaction medium is taken to reflux for 1 hour 30 minutes. The sodium diselenide Na₂Se₂ formed in solution is reddish brown in colour.

At ambient temperature, bis(2-chloroethyl)amine hydrochloride (847 mg, 4.7 mmol, 0.5eq) in solution in an NaOHaq (0.5 M)/EtOH (1/1) mixture V = 3 mL is added dropwise to the above solution of Na₂Se₂ and Na₂Se₃ over 10 minutes. The reaction medium is stirred for 45 minutes at ambient temperature.

The reaction medium is then diluted V = 10 mL with a solution of NaOHaq (0.5M) then extracted with CH₂Cl₂ (3 × 30 mL). The organic phases are then combined, dried with Na₂SO₄, then concentrated to 1/3 of the total volume.

The organic concentrate obtained is then extracted with an aqueous solution of trifluoroacetic acid (TFA) (10%) then purified directly by RP-HPLC: Xbridge C18 Column (d = 1.9 cm, L = 20 cm, 130 Å, 5 µm), UV detection (λ = 215 nm), buffer A: H₂O/TFA (1:0.05% v/v), buffer B: CH₃CN/H₂O/TFA (4:1:0.05% v/v/v), isocratic: buffer A (3 min), then gradient: buffer B (0 to 50% over 9 min, 25 mL/min).

After lyophilization, compounds **1** (Se₂EA) 230 mg (Yield Y = 21 %) and compounds **2** (Se₃EA) 240 mg (Yield Y = 16 %) are obtained in the form of a yellow powder.

LC-MS analysis, RP-HPLC chromatogram of compounds **1** (Se₂EA) and **2** (Se₃EA) respectively

LC-MS: Buffer A: H₂O/TFA (1/0.1% v/v), Buffer B: CH₃CN/H₂O/TFA (4/1:0.1% v/v/v).

RP-HPLC on an XBridge BEH C18 column (300 Å, 3.5 µm, 4.6 x 150 mm) using a linear gradient 0-100% B over 30 minutes, flow rate 1 mL/min, UV detection ($\lambda = 215$ nm).

MS: Electrospray ionization positive mode, voltage cone 30V, quadripolar analyser.

The chromatogram and the mass spectrum of compound **1** are shown in Figures 1a and 1b respectively.

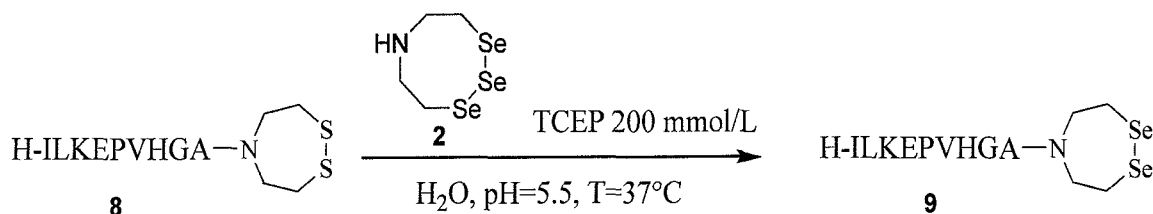
The chromatogram and the mass spectrum of compound **2** are shown in Figures 2a and 2b respectively.

Example 2: Synthesis and isolation of the SeEA segments according to the invention by exchange, starting from SEA segments.

Synthesis of peptide 8: H-ILKEPVHGA-SEA

The method for the synthesis of this peptide is described in the document Ollivier, N.; Dheur, J.; Mhidia, R.; Blanpain, A.; Melnyk, O. *Organic letters* **2010**, *12*, 5238-41.

Synthesis of peptide 9: H-ILKEPVHGA-SeEA



The reaction is carried out in a glove box (the dioxygen concentration is less than 50 ppm) under a nitrogen atmosphere. A 200 mmol/L solution of TCEP is prepared in a 0.1M phosphate buffer at pH = 7.2. The pH of the TCEP/MPAA solution obtained is adjusted to pH = 5.5.

Peptide **8** (3.8 mg, 3 µmol) and Se₃EA linker **2** (8.4 mg, 28 µmol, 10 eq) are dissolved in the above solution of TCEP V = 297 µL. The reaction medium is stirred for 24 hours at 37°C.

At the end of the reaction, the reaction medium is diluted V = 2 mL with a solution of TFA (0.1%) then purified directly by reversed-phase high-performance liquid chromatography (RP-HPLC) (C18 Nucleosil column (d = 1 cm, L = 20 cm, 120 Å, 5 µm), UV detection ($\lambda = 215$ nm), buffer A: H₂O/TFA (1:0.05% v/v), buffer B: CH₃CN/H₂O/TFA (4:1:0.05% v/v/v), gradient: buffer B (0 to 20% over 5 min then 20 to 42% over 60 min, 6 mL/min)). 1.96 mg of peptide **9** is obtained (yield 62%).

LC-MS analysis, RP-HPLC chromatogram of peptide 9

LC-MS: Buffer A: H₂O/TFA (1/0.1% v/v), Buffer B: CH₃CN/H₂O/TFA (4/1:0.1% v/v/v).

RP-HPLC on an XBridge BEH C18 column (300 Å, 3.5 µm, 4.6 x 150 mm) using a linear gradient 0-100% B over 30 minutes, flow rate 1 mL/min, UV detection ($\lambda = 215$ nm).

MS: Electrospray ionization positive mode, voltage cone 30V, quadrupolar analyser.

Figure 3a corresponds to the chromatogram of the solution after a reaction time t of 0.8 hour. For $t = 0.8$ hour, the exchange reaction is not completed, in fact, the peak corresponding to peptide **8** is still present in a significant quantity.

Figure 3b corresponds to the chromatogram of the solution after a reaction time t of 6 hours and Figure 3c is the corresponding mass spectrum. This chromatogram shows that the exchange reaction is completed.

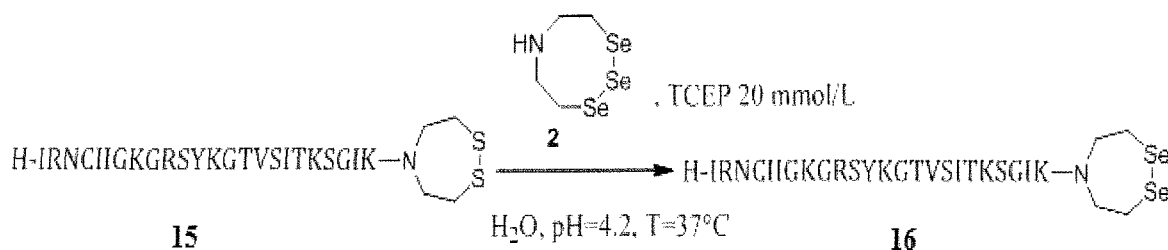
Figures 4a and 4b correspond to the chromatogram and the mass spectrum of peptide **9** after purification.

Example 2a: Synthesis and isolation of large SeEA segments according to the invention by exchange, starting from SEA segments.

Synthesis of peptide 15: H-IRNCIIGKGRSYKGTVSITKSGIK-SEA

The method for the synthesis of this peptide is described in the document Ollivier, N.; Dheur, J.; Mhidia, R.; Blanpain, A.; Melnyk, O. *Organic letters* **2010**, 12, 5238-41.

Synthesis of peptide 16: H-IRNCIIGKGRSYKGTVSITKSGIK-SeEA



The reaction is carried out under a nitrogen atmosphere. A 20 mmol/L solution of TCEP is prepared in a 0.1M acetate buffer at pH = 5.5. The pH of the solution TCEP obtained is adjusted to pH = 4.2.

Peptide **15** (4 mg, 1.4 µmol) and Se₃EA linker **2** (4.4 mg, 14.3 µmol, 10 eq) are dissolved in the above solution of TCEP V = 719 µl. The reaction medium is stirred for 24 hours at 37°C.

At the end of the reaction, the reaction medium is diluted V = 3 mL with a solution of TFA (0.1%) then purified directly by reversed-phase high-performance liquid chromatography (RP-HPLC) (C18 Nucleosil column (d = 1 cm, L = 20 cm, 120

Å, 5 µm), UV detection ($\lambda = 215$ nm), buffer A: H₂O/TFA (1:0.05% v/v), buffer B: CH₃CN/H₂O/TFA (4:1:0.05% v/v/v), gradient: buffer B (0 to 20% over 5 min then 20 to 42% over 60 min, 6 mL/min)). 1.8 mg of peptide 16 is obtained (Yield 38%).

LC-MS analysis, RP-HPLC chromatogram of peptide 16

5 LC-MS: Buffer A: H₂O/TFA (1/0.1% v/v), Buffer B: CH₃CN/H₂O/TFA (4/1:0.1% v/v/v).

RP-HPLC on an XBridge BEH C18 column (300 Å, 3.5 µm, 4.6 x 150 mm) using a linear gradient 0-100% B over 30 min, flow rate 1 mL/min, UV detection ($\lambda = 215$ nm).

10 MS: Electrospray ionization positive mode, voltage cone 30V, quadripolar analyser.

Figure 13 corresponds to the chromatogram of the solution after a reaction time *t* of 24 hours. The peak corresponding to peptide 16 is clearly present and indicates that the exchange reaction is completed.

15 Figures 14a and 14b correspond respectively to the chromatogram and to the mass spectrum of peptide 16 after purification.

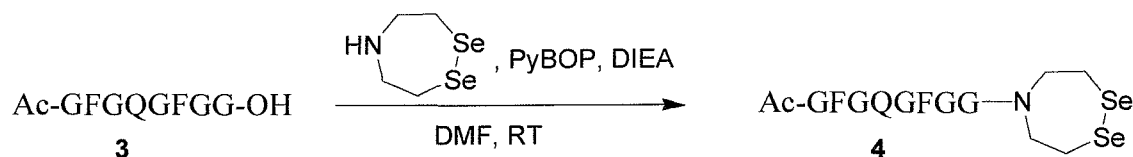
Example 3: Native ligation with selenium-based compounds according to the invention or sulphur-based compounds according to the prior art

20

Synthesis of peptide 3, SEQ ID NO: 1 Ac-GFGQGFGG-OH

The method for the synthesis of this peptide is described in the document Ollivier, N.; Dheur, J.; Mhidia, R.; Blanpain, A.; Melnyk, O. *Organic letters* **2010**, *12*, 5238-41.

25 *Synthesis of peptide 4: Ac-GFGQGFGG-SeEA*



30

Peptide 3 (50 mg, 0.07 mmol) and molecule 1 (30 mg, 0.13 mmol, 2 eq) are dissolved in *V* = 3.5 mL of DMF (*N,N*-dimethylformamide) followed by the addition of benzotriazol-1-yloxy-tripyrrolidinophosphoniumhexafluorophosphate (PyBOP) (68 mg, 0.14 mmol, 2 eq). *N,N*-diisopropylethylamine (DIEA) is then added (45 µL, 0.28 mmol, 4 eq). The reaction medium is stirred for 1.5 hours at ambient temperature then precipitated from a cold mixture of diethyl ether/heptane (75 mL, 1:1 v/v), centrifuged, then dissolved in a minimum amount of water and lyophilized.

35

Raw peptide **4** is purified directly by reversed-phase high-performance liquid chromatography (RP-HPLC) (C18 Nucleosil column (d = 1 cm, L = 20 cm, 120 Å, 5 μm), UV detection (λ = 215 nm), buffer A: H₂O/TFA (1:0.05% v/v), buffer B: CH₃CN/H₂O/TFA (4:1:0.05% v/v/v), gradient: buffer B (0 to 20% over 5 min then 20 to 42% over 60 min, 6 mL/min)). 15 mg of peptide **4** is obtained (Yield 24 %).

LC-MS analysis, RP-HPLC chromatogram of peptide 4

LC-MS: Buffer A: H₂O/TFA (1/0.1% v/v), Buffer B: CH₃CN/H₂O/TFA (4/1:0.1% v/v/v).

RP-HPLC on an XBridge BEH C18 column (300 Å, 3.5 μm, 4.6 x 150 mm) using a linear gradient 0-100% B over 30 minutes, flow rate 1 mL/min, UV detection (λ = 215 nm).

MS: Electrospray ionization positive mode, voltage cone 30V, quadrupolar analyser.

Figures 5a and 5b represent respectively the chromatogram and the mass spectrum of peptide **4** after purification.

Synthesis of peptide 5: Ac-GFGQGFGG-SEA

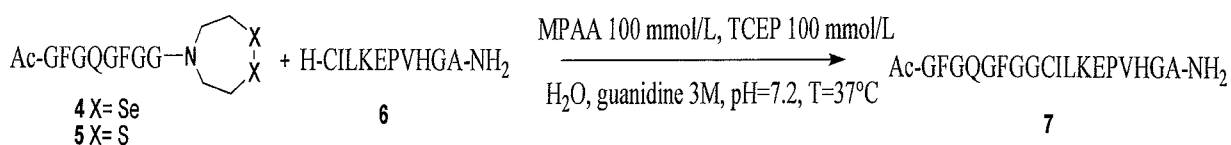
The method for the synthesis of this peptide is described in the document Ollivier, N.; Dheur, J.; Mhidia, R.; Blanpain, A.; Melnyk, O. *Organic letters* **2010**, *12*, 5238-41.

Synthesis of peptide 6, SEQ ID NO: 2 H-CILKEPVHGA-NH₂

The method for the synthesis of this peptide is described in the document Ollivier, N.; Dheur, J.; Mhidia, R.; Blanpain, A.; Melnyk, O. *Organic letters* **2010**, *12*, 5238-41.

Synthesis of peptide 7, SEQ ID NO: 3 Ac-GFGQGFGGCILKEPVHGA-NH₂ by SeEA ligation. Comparison of the reactivity of peptide 4 Ac-GFGQGFGG-SeEA and peptide

5 Ac-GFGQGFGG-SEA



a- Ligation kinetics and yield of peptide **4** Ac-GFGQGFGG-SeEA with peptide **6** H-CILKEPVHGA-NH₂

The SeEA ligation of peptide **4** with peptide **6** is carried out in a glove box (dioxygen concentration less than 50 ppm) under a nitrogen atmosphere. A 3M solution of guanidine is prepared in a 0.2M phosphate buffer at pH = 7.3. Tris(2-carboxyethyl)phosphine (TCEP, 28.6 mg, 0.1 mmol) and 3-mercaptophenylacetic acid (MPAA, 16.8 mg, 0.1 mmol) are then added. The pH of the TCEP/MPAA/Guanidine solution obtained is adjusted to pH = 7.2.

Peptides **4** (4.9 mg, 5 μ mol) and **6** (10.6 mg, 7.6 μ mol, 1.5 eq) are dissolved in the above TCEP/MPAA/Guanidine solution (1.5 mL). The reaction medium is stirred for 24 hours at 37°C.

Once the ligation reaction is completed, the reaction medium is diluted with water V = 3 mL then acidified with a solution of TFA (10%) V = 1 mL then the MPAA is extracted with ether using 3 times 6.5 mL.

The ligation product **7** in aqueous phase is purified directly by reversed-phase high-performance liquid chromatography (RP-HPLC) (C18 Nucleosil column (d = 1 cm, L = 20 cm, 120 Å, 5 μ m), UV detection (λ = 215 nm), buffer A: H₂O/TFA (1:0.05% v/v), buffer B: CH₃CN/H₂O/TFA (4:1:0.05% v/v/v), gradient: buffer B (0 to 20% over 5 min then 20 to 42% over 60 min, 6 mL/min). 5.6 mg of peptide **7** is obtained (Yield 54 %).

LC-MS analysis, RP-HPLC chromatogram of peptide **7**

LC-MS: Buffer A: H₂O/TFA (1/0.1% v/v), Buffer B: CH₃CN/H₂O/TFA (4/1:0.1% v/v/v).

RP-HPLC on an XBridge BEH C18 column (300 Å, 3.5 μ m, 4.6 x 150 mm) using a linear gradient 0-100% B over 30 minutes, flow rate 1 mL/min, UV detection (λ = 215 nm).

MS: Electrospray ionization positive mode, voltage cone 30V, quadripolar analyser.

Figures 6a and 6b represent respectively the chromatogram and the mass spectrum of peptide **7** after purification.

b- Kinetics of the ligation of peptide **5** Ac-GFGQGFGG-SEA with peptide **6** H-CILKEPVHGA-NH₂

The reaction of SEA ligation of peptide **5** with peptide **6** is carried out under strictly the same conditions.

Comparison of the kinetics

Figure 7 represents a comparison of the kinetics of the ligation reaction for the formation of peptide **7**, with a selenium-based compound (peptide **4**) according to the

invention, curve represented by the symbol ● or with a sulphur-based compound (peptide **5**) according to the state of the art, curve represented by the symbol ▲.

Figure 7 shows a higher reaction rate for the ligation reaction according to the invention, curve represented by the symbol ●.

5 Example 4: Comparison of the ligation kinetics for the SeEA, SEA and NCL (thioester chemistry) methods

*Synthesis of peptide **15a**: H-ILKEPVHGV-SEA*

10 The method for the synthesis of this peptide is described in the document Ollivier, N.; Dheur, J.; Mhidia, R.; Blanpain, A.; Melnyk, O. *Organic letters* **2010**, 12, 5238-41.

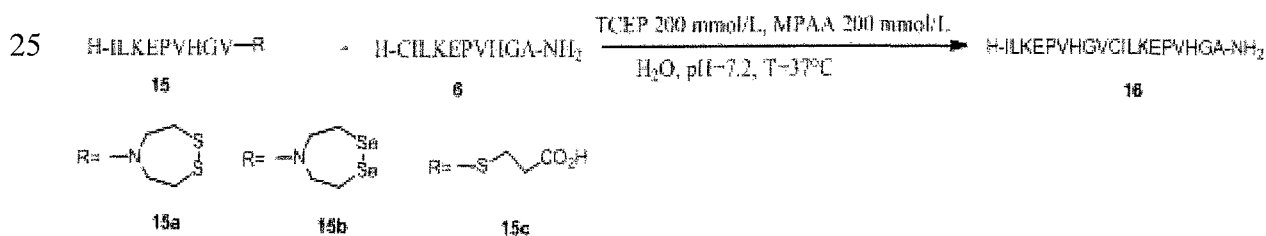
*Synthesis of peptide **15b**: H-ILKEPVHGV-SeEA*

15 Starting from peptide **15a**, the method for the synthesis of peptide **15b** is strictly identical to that used for the synthesis of peptide **9**.

*Synthesis of peptide **15c**: H-ILKEPVHGV-MPA (MPA = 3-mercapto propionic acid)*

20 The method for the synthesis of this peptide is described in the document Dheur, J.; Ollivier, N.; Vallin, A.; Melnyk, O. *Journal of Organic Chemistry* **2010**, 76, 3194-3202.

*Comparison of the ligation kinetics of peptide **15a/15b/15c** H-ILKEPVHGV-R with peptide **6** H-CILKEPVHGA-NH₂ for the SeEA, SEA ligation and NCL methods*



30 The different ligations of peptide **15a** or **15b** or **15c** with peptide **6** are carried out separately in a glove box ([O₂] <50 ppm) under a nitrogen atmosphere. A 200 mmol/L solution of TCEP is prepared in a 0.1M phosphate buffer at pH = 7.2. 3-Mercaptophenylacetic acid (MPAA 33 mg, 0.2 mmol) is then added. The pH of the TCEP/MPAA solution obtained is adjusted to pH = 7.2.

35 Peptides **15a** or **15b** or **15c** and **6** (1.5 eq) are dissolved separately in the above TCEP/MPAA solution at a concentration of 3 mmol/L. The reaction medium is stirred for 24 hours at 37°C. The formation kinetics of ligation product **16** are monitored by HPLC.

Figure 12 allows comparison of the formation reaction kinetics of peptide **16** according to the SeEA, SEA ligation or NCL (thioester chemistry) method.

In Figure 12, the curve represented by ● corresponds to the formation of peptide **16** from peptide **15b**, the curve represented by ■ corresponds to the formation of peptide **16** from peptide **15a** and the curve represented by ♦ corresponds to the formation of peptide **16** from peptide **15c**.

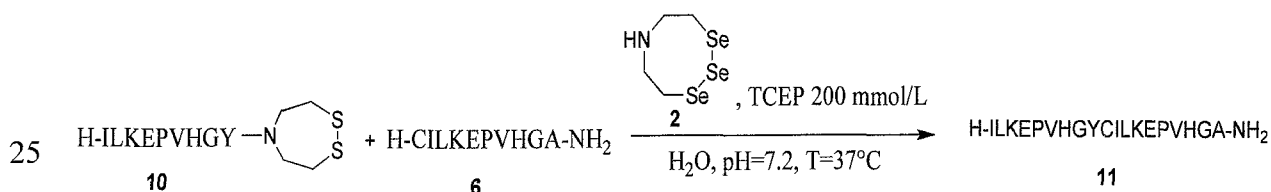
Figure 12 shows an SeEA ligation reaction rate (curve represented by ●) higher than that of the NCL method (curve represented by ■), itself significantly higher than that of the SEA method (curve represented by ♦).

Example 5: Formation of SeEA segments *in situ* by simultaneous exchange and ligation - Catalysis of the ligation reaction

11-Synthesis of peptide **10**: H-ILKEPVHGY-SEA

The method for the synthesis of this peptide is described in the document Ollivier, N.; Dheur, J.; Mhidia, R.; Blanpain, A.; Melnyk, O. *Organic letters* **2010**, *12*, 5238-41.

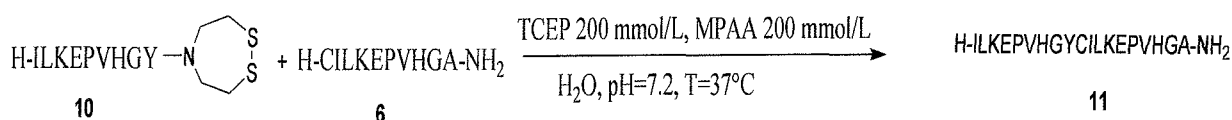
a- Ligation kinetics of peptide **10** H-ILKEPVHGY-SEA with peptide **6** H-CILKEPVHGA-NH₂ in the presence of Se₃EA linker **2** in order to produce peptide **11**, SEQ ID NO: 4 H-ILKEPVHGYCILKEPVHGA-NH₂



The SEA ligation of peptide **10** with peptide **6** is carried out in a glove box (dioxygen concentration less than 50 ppm) under a nitrogen atmosphere. A 200 mmol/L solution of TCEP is prepared in a 0.1 M phosphate buffer at pH 7.2.

Peptides **10** (1 mg, 0.6 μmol) and **6** (1.4 mg, 1 μmol, 1.5 eq) as well as the Se₃EA linker **2** (5.8 mg, 0.02 mmol, 28 eq) are dissolved in the above solution of TCEP (94 μL). The reaction medium is stirred for 24 hours at 37°C. The ligation formation kinetics of product **11** are monitored by HPLC.

b- Comparison of the ligation kinetics of peptide **10** H-ILKEPVHGY-SEA with peptide **6** H-CILKEPVHGA-NH₂ in the presence of MPAA



The SEA ligation of peptide **10** with peptide **6** is carried out in a glove box (dioxxygen concentration less than 50 ppm) under a nitrogen atmosphere. A 200 mmol/L solution of TCEP is prepared in a 0.1M phosphate buffer at pH 7.2. 3-Mercaptophenylacetic acid (MPAA 33 mg, 0.2 mmol) is then added. The pH of the TCEP/MPAA solution obtained is adjusted to pH = 7.2.

Peptides **10** (1 mg, 5 μ mol) and **6** (10.6 mg, 7.6 μ mol, 1.5 eq) are dissolved in the above TCEP/MPAA solution (94 μ L). The reaction medium is stirred for 24 hours at 37°C. The formation kinetics of ligation product **11** are monitored by HPLC.

Figure 8 allows comparison of the reaction kinetics of the formation of peptide **11** in the presence or not of compound **2** according to the invention.

In Figure 8, the curve represented by ● corresponds to the ligation according to the invention of peptide **10** in the presence of selenium-containing compound **2** and the curve represented by ▲ corresponds to the ligation according to the prior art of peptide **10** without the selenium-containing compound but in the presence of MPAA.

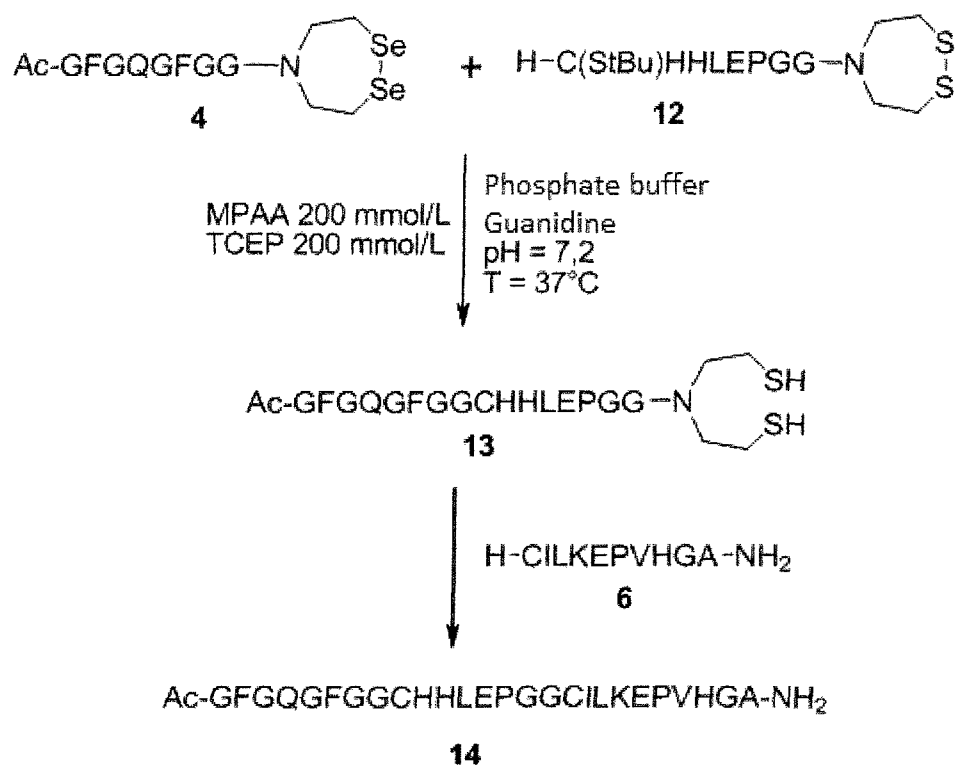
Figure 8 shows a higher reaction rate when compound **2** is present.

Example 6: One-pot sequential ligation of three segments using SeEA and SEA ligations

Synthesis of peptide 12: H-C(StBu)HHLEPGG-SEA

The method for the synthesis of this peptide is described in the document Ollivier, N.; Dheur, J.; Mhidia, R.; Blanpain, A.; Melnyk, O. *Organic letters* **2010**, 12, 5238-41.

Synthesis of peptide 14, SEQ ID NO: 5 Ac-GFGQGFGG-CHHLEPGG-CILKEPVHGA-NH₂ by ligation of segments 4 Ac-GFGQGFGG-SeEA, 12 H-C(StBu)HHLEPGG-SEA and 6 H-CILKEPVHGA-NH₂



- Step 1

Guanidine hydrochloride (GdnHCl, 573.18 mg, 6 mmol) is dissolved in 0.1 M phosphate buffer pH = 7.3 (1 mL final, 6M).

5 4-Mercaptophenylacetic acid (MPAA, 33.83 mg, 0.2 mmol) and *tris*(2-carboxyethyl)phosphine hydrochloride (TCEP-HCl, 57.37 mg, 0.2 mmol) are dissolved in this solution (1 mL). 6M NaOH is added to adjust the pH of the solution to 7.37.

10 Peptides **4** Ac-GFGQGFGG-SeEA (5.33 mg, 5.4 μ mol) and **12** H-C(StBu)HHLEPGG-SEA (7.59 mg, 5.4 μ mol) are dissolved together in the above solution (780 μ L). The reaction medium is placed in a thermostatically controlled bath at 37°C under an inert atmosphere.

15 The ligation is monitored by RP-HPLC on an XBridge BEH C18 column (4.6 x 250 mm, 300 Å, 5 μ m) (215 nm, 1 mL/min, 30°C, buffer A water containing 0.1% TFA, buffer B CH₃CN/water 4/1 containing 0.1% TFA, 0-100% B over 30 min). For this, an aliquot (2 μ L) of reaction medium is acidified with 10% aqueous TFA, extracted with Et₂O in order to remove MPAA before analysis.

Peptide **13** Ac-GFGQGFGG-CHHLEPGG-SEA in solution is obtained in this way.

20 Figures 9a and 9b represent respectively the chromatogram and the mass spectrum of peptide **13**.

- *Step 2*

After reaction for 4 hours 35 minutes, peptide **6** H-CILKEPVHGA-NH₂ (11.06 mg, 8.1 μmol) is added in solid form to the reaction medium. The reaction is placed under an inert atmosphere at 37°C.

5 The ligation is monitored by RP-HPLC on an XBridge BEH C18 column (4.6 × 250 mm, 300 Å, 5 μm) (215 nm, 1 mL/min, 30°C, buffer A water containing 0.1% TFA, buffer B CH₃CN/water 4/1 containing 0.1% TFA, 0-100% B over 30 min). For this, an aliquot (2 μL) of reaction medium is acidified with 10% aqueous TFA, extracted with Et₂O in order to remove MPAA before analysis.

10 When the reaction is complete, the reaction medium is diluted with water (4 mL) and 10% aqueous TFA (1 mL) is added. After 4 extractions with Et₂O (4 × 4 mL), the aqueous phase is degassed for 2 minutes by bubbling argon through it. After purification by RP-HPLC on a nucleosil C18 column, 120 Å, 5 μm (215 nm, 6 mL/min, buffer A water containing 0.05% TFA, buffer B CH₃CN/water 4/1
15 containing 0.05% TFA, 0-10% B over 5 min, then 10-100% B over 150 min), 6 mg (36%) of peptide **14** Ac-GFGQGFGG-CHHLEPGG-CILKEPVHGA-NH₂ is obtained.

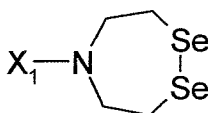
Figures 10a and 10b represent the chromatogram and the mass spectrum of compound **14** before purification respectively.

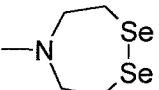
Figures 11a and 11b represent the chromatogram and the mass spectrum of compound
20 **14** after purification respectively.

CLAIMS

5 1. Functionalized peptide selected from:

a) the peptide of formula (I): $X_1-N(CH_2CH_2SeH)_2$; or

b) the peptide of formula (I'): X_1-N  ;

10 in which X_1 represents a peptide fragment and the $-N(CH_2CH_2SeH)_2$ group or  group forms an amide bond with the C=O termination

of the amino acid residue of the peptide fragment X_1 which is in the C-terminal position.

15

2. Functionalized peptide according to claim 1, in which X_1 comprises from 2 to 300 amino acid residues, preferably from 5 to 100 amino acid residues, more particularly preferably from 8 to 50 amino acid residues.

20 3. Process for obtaining a functionalized peptide according to claim 1 or according to claim 2, comprising the following steps a1 and b1:

a1) peptide synthesis for providing a polypeptide of formula (II):

(II) X_1-OH

preferably comprising protective groups on its amine and carboxylic acid functions, with the exception of its C-terminal carboxylic acid function,

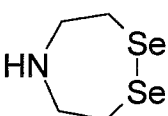
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b1) functionalization of the polypeptide obtained in step a) comprising:

— the reaction of the polypeptide of formula (II) with an amine compound selected from:

(IIIa) $NH(CH_2-CH_2-Se-G_1)_2$

30

(III') HN 

in which G_1 represents a selenium protective group or a hydrogen atom,

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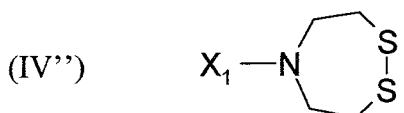
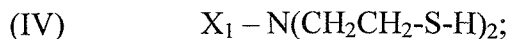
in liquid phase, in order to form the polypeptide of formula (Ia) $X_1-N(CH_2CH_2-Se-G_1)_2$, or (I');

– optionally the deprotection of the polypeptide of formula (Ia) in order to produce the compound of formula (I),
or the following steps a2 and b2

a2) the use

5

- of the compound of formula (IV) or (IV''):



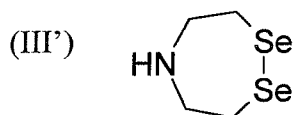
or

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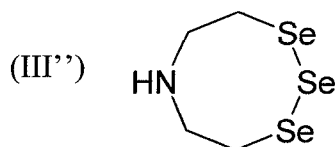
- of a peptide functionalized by a group bearing a thioester function (IV') = $X_1 - SR$ and R represents an optionally substituted aryl or alkyl group,

b2) the reaction between the compound of formula (IV) or (IV'') or the compound of formula (IV') with the amine compound of formula (III') or (III''):

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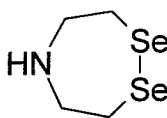
in liquid phase, in order to form the peptide of formula (I) or (I').

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4. Compound capable of being used in the process according to claim 3 selected from:

a)

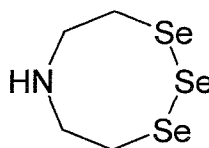
the compound of formula (III'):



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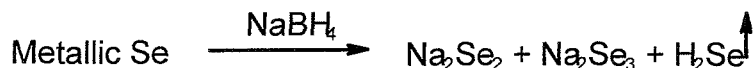
b)

the compound of formula (III''):

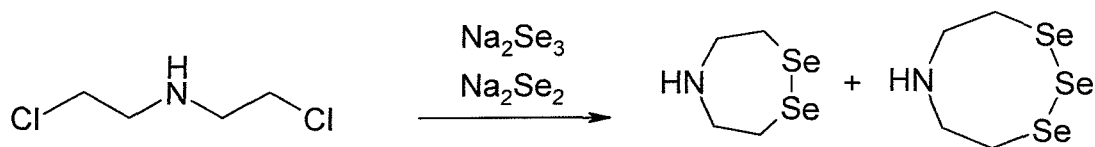


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5. Process for producing the compounds of formulae (III') and (III'') according to claim 4, comprising the following treatment reaction with a metal hydride $NaBH_4$, $LiBH_4$, or $LiAlH_4$:

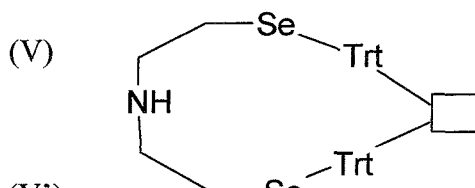


followed by the reaction:

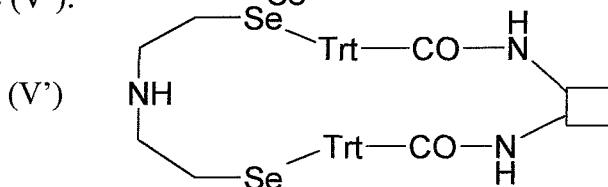


6. Process for obtaining a peptide according to claim 1 or 2, comprising the following steps:

a) Provision of a functionalized polymer resin with the structure (V):



Or the structure (V'):

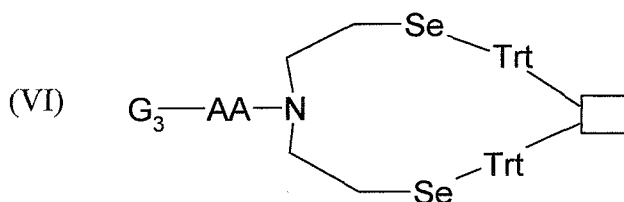


in which,

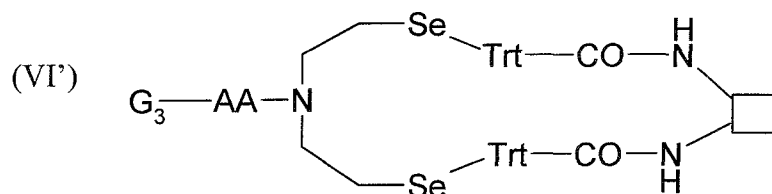
□ represents a solid support,

Trt represents a triphenylmethyl group optionally substituted by one or more substituents, in particular selected from the substituents chlorine, methoxy, methyl, fluorine and cyano.

b) Grafting of an amino acid onto the functionalized polymer resin obtained in step a) in order to produce the compound of structure (VI):



Or of structure (VI'):



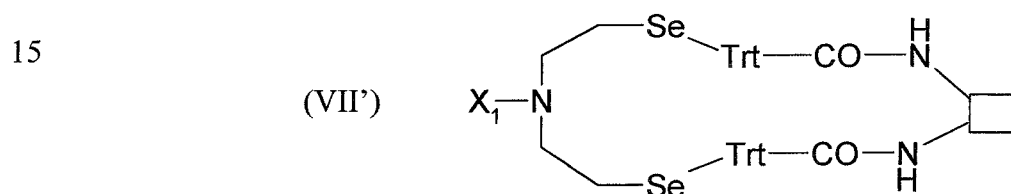
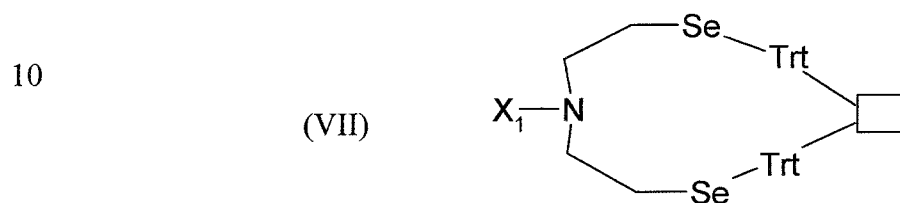
in which,

$\square$ and Trt have the same meaning as in compound (V) or (V'),

AA represents an amino acid residue optionally comprising one or more protective groups;

5 G_3 represents a hydrogen atom or a protective group of the N-terminal amine function of AA,

c) Peptide synthesis starting from the amino acid AA in order to produce compound (VII) or (VII'):



in which,

20 $\square$ represents a solid support,

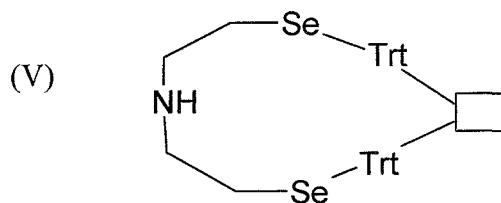
Trt represents a triphenylmethyl group optionally substituted by one or more substituents, in particular selected from the substituents chlorine, methoxy, methyl, fluorine and cyano,

X_1 represents a peptide fragment,

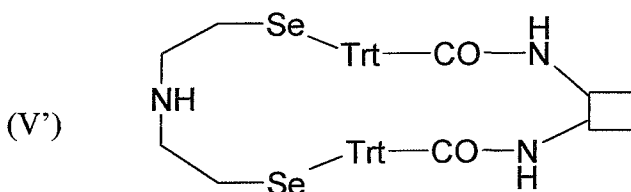
25 d) Cleavage of the bond between Trt and Se in order to produce compound (I) or (I').

7. Process according to claim 6, in which the grafting of an amino acid onto the polymer resin comprises placing the functionalized resin (V) or (V') in contact
30 with an amino acid halide or with an amino acid and an activation agent, preferably selected from HATU, PyBOP, BOP, PyBROP, more particularly preferably HATU.

8. Functionalized polymer resin capable of being used in the process according to
35 claim 6 or 7, with the structure (V):



Or the structure (V'):

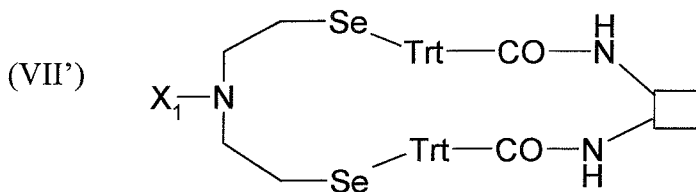
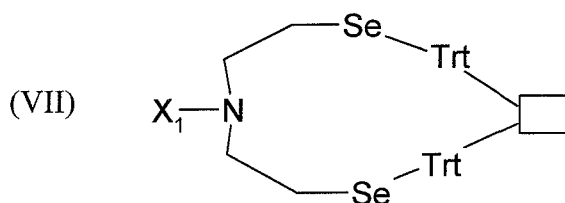


in which,

□ represents a solid support,

Trt represents a triphenylmethyl group optionally substituted by one or more substituents, in particular selected from the substituents chlorine, methoxy, methyl, fluorine and cyano.

9. Polymer resin comprising a peptide fragment corresponding to formula (VII) or (VII'):



in which,

□ represents a solid support,

Trt represents a triphenylmethyl group optionally substituted by one or more substituents, in particular selected from the substituents chlorine, methoxy, methyl, fluorine and cyano,

X₁ represents a peptide fragment.

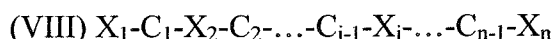
10. Polymer resin according to claim 8 or 9, in which the solid support □ is selected from a polystyrene, a polyacrylamide, a polyethylene glycol, a cellulose, a polyethylene, a polyester, a latex, a polyamide, a polydimethylacrylamide, a

polyethylene glycol-polystyrene copolymer, a polyethylene glycol-polyacrylamide copolymer and derivatives thereof.

11. Process for producing the polymer resin according to one of claims 8 to 10, comprising:

- a) the provision of a functionalized solid support by -Trt or -NH-CO-Trt groups;
 - b) reaction of the Trt function of the solid support with the amine compound of formula (III): $\text{NH}(\text{CH}_2\text{-CH}_2\text{-Se-H})_2$
- Trt having the same definition as in claims 8 to 10.

12. Process for producing a polypeptide of formula (VIII):



in which,

$X_1, X_2, \dots, X_i, \dots, X_n$ are peptide fragments,

$C_1, C_2, \dots, C_{i-1}, \dots, C_{n-1}$ are amino acid residues bearing a thiol or selenol function,

n is an integer greater than or equal to 2,

i is any integer in the range from 1 to n ,

this process comprising one or more steps of reaction between a compound of formula (IX_i) :

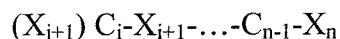


in which,

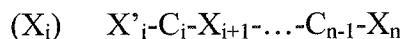
X'_1 represents X_1

X'_i represents $C_{i-1}\text{-X}_i$ for $i > 1$

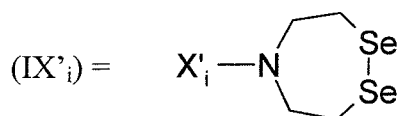
and a compound of formula (X_{i+1}) :



in order to form a compound of formula (X_i) :

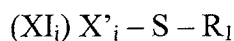


13. Process according to claim 12, in which compound (IX_i) is formed starting from compound (IX'_i) :

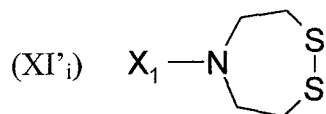


by placing in contact with at least one compound that reduces diselenium bonds.

14. Process according to claim 13, in which compound (IX'_i) is obtained by reaction between a compound of formula (XI_i), (XI'_i) or (XI''_i):



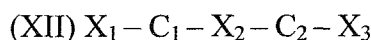
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in which X'_i is a peptide fragment of the form C_{i-1}-X_i for i > 1 and X'₁ represents X₁, R₁ represents an optionally substituted aryl or alkyl group, and a compound of formula (III') or of formula (III'') according to claim 4.

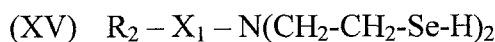
15. Process according to claim 13 or 14, for the production of a polypeptide represented by formula (XII):



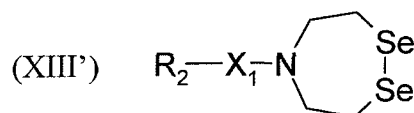
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comprising the following steps:

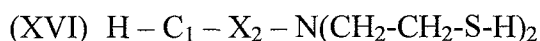
a) reaction between a compound of formula (XIII) or (XIII'):



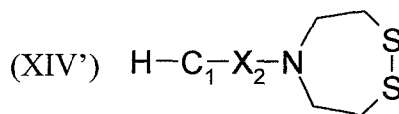
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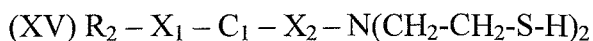
and a compound of formula (XIV) or (XIV'):



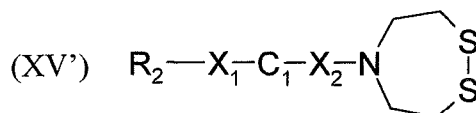
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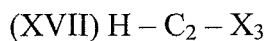
in order to form the compound of formula (XV) or (XV'):



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b) reaction between a compound of formula (XV) or (XV') and a polypeptide of formula (XVI):



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in order to form the compound of formula (XII),

in which, R_2 represents H or a protective group of the N-terminal function of X_1 ; X_1 , X_2 , X_3 represent, independently of each other, peptide fragments; C_1 and C_2 represent, independently of each other, amino acid residues bearing a thiol or selenol function.

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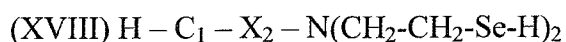
16. Process for producing a cyclic polypeptide of formula (XVII):



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X_2 representing a peptide fragment, and C_1 representing an amino acid residue comprising a thiol or selenol function,

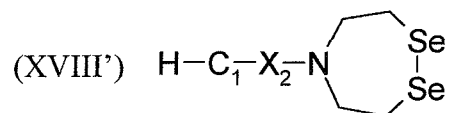
said process comprising at least one reaction step of ligation of a polypeptide of formula (XVIII):



with itself.

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17. Process according to claim 16, in which the ligation reaction is carried out by placing a polypeptide of formula (XVIII'):



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in contact with at least one compound that reduces diselenium bonds.

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18. Polypeptide synthesis kit comprising one or more peptides of formulae (IX_i) or (IX'_i) capable of being used in the process according to claim 12 or 13, or at least one compound according to claim 4.