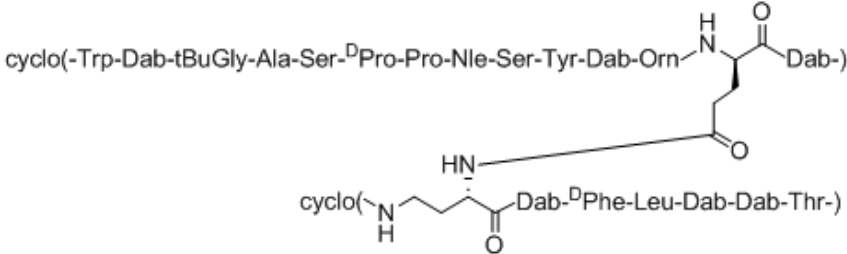
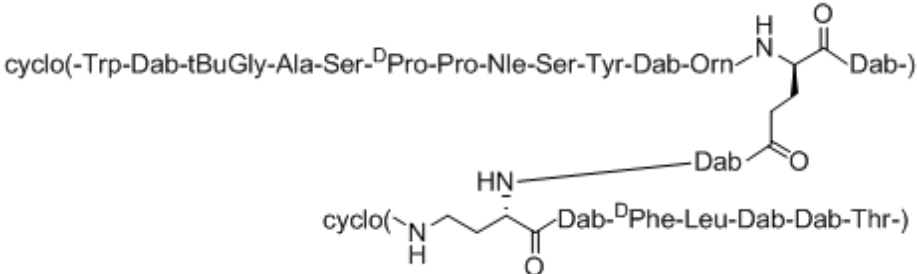
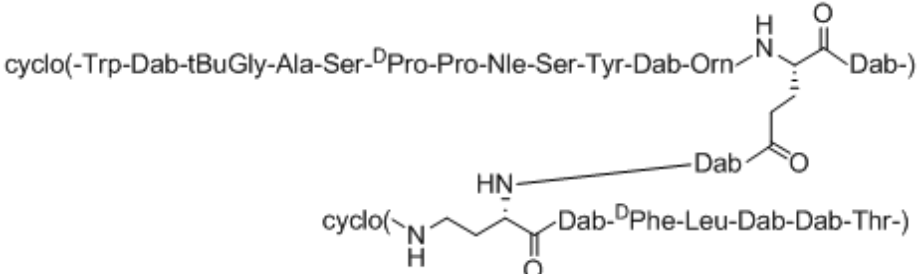
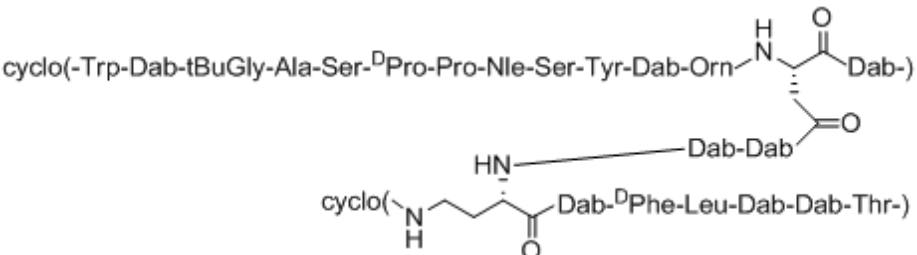


CLAIMS

1. A compound of formula (I) or a pharmaceutically acceptable salt thereof, wherein the compound is selected from the group consisting of **Ex. 1** to **216**:

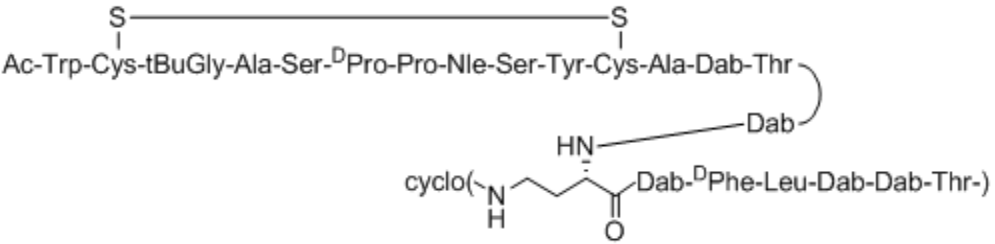
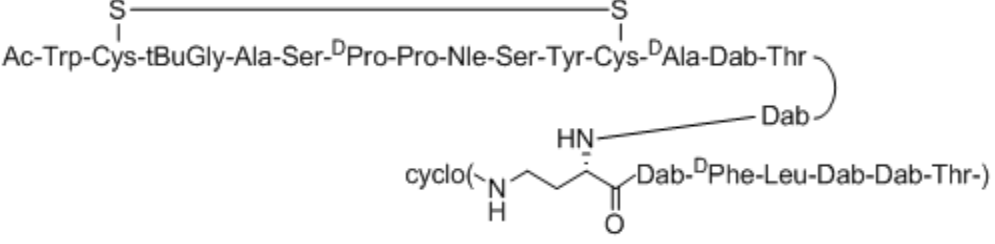
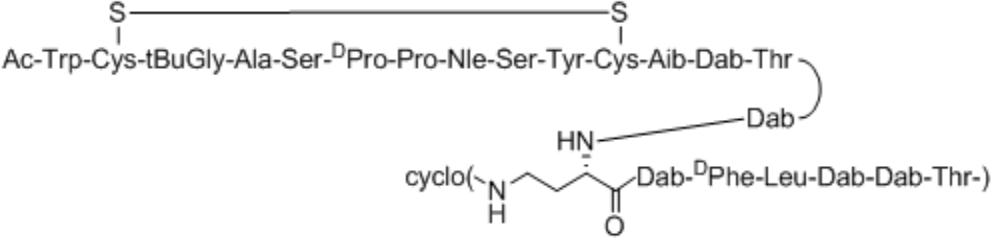
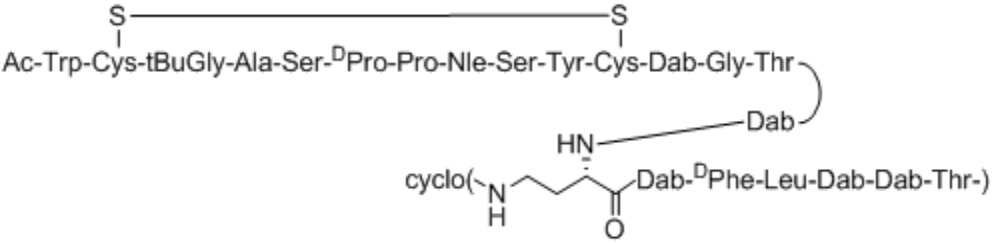
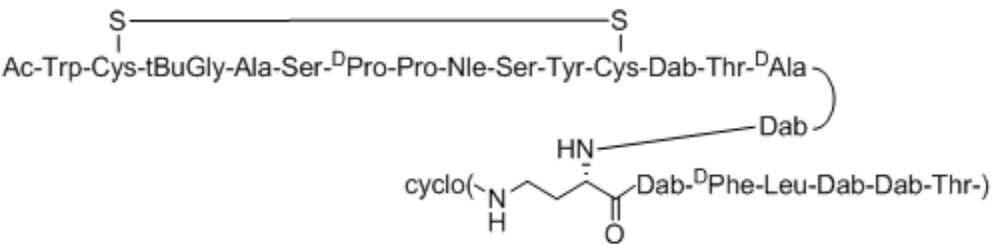
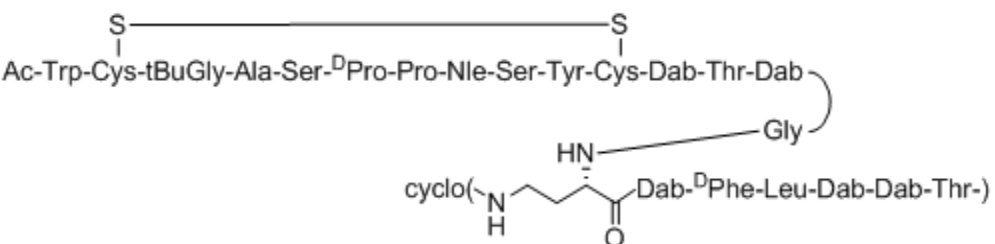
Ex. No.	
Ex. 1	
Ex. 2	
Ex. 3	
Ex. 4	

Ex. 5	<p>cyclo(-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-Orn-NH-CH(CH₂CH₂CO-NH-Dab-Thr)-CO-Dab-)</p> <p>cyclo(-NH-CH(CH₂CH₂CO-NH-Dab-Thr)-CO-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>
Ex. 6	<p>cyclo(-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-Orn-NH-CH(CH₂CH₂CO-NH-Dab-Thr)-CO-Dab-)</p> <p>cyclo(-NH-CH(CH₂CH₂CO-NH-Dab-Thr)-CO-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>
Ex. 7	<p>cyclo(-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-NH-CH(CH₂CH₂CO-NH-Dab-Thr-Dab)-CO-^DDab-Dab-)</p> <p>cyclo(-NH-CH(CH₂CH₂CO-NH-Dab-Thr-Dab)-CO-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>
Ex. 8	<p>cyclo(-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Val-Ser-Tyr-Dab-Orn-NH-CH(CH₂CH₂CO-NH-Dab-Thr-Dab)-CO-Dab-)</p> <p>cyclo(-NH-CH(CH₂CH₂CO-NH-Dab-Thr-Dab)-CO-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>
Ex. 9	<p>cyclo(-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-Orn-NH-CH(CH₂CH₂CO-NH-Dab-Thr-Dab)-CO-Dab-)</p> <p>cyclo(-NH-CH(CH₂CH₂CO-NH-Dab-Thr-Dab)-CO-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>

Ex. 10	<p>cyclo(-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-Orn-^DDab-)</p> <p>HN-CH(CH₂-CH₂-C(=O)-Dab-Thr-Dab)-</p> <p>cyclo(-NH-CH(CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>
Ex. 11	<p>cyclo(-Trp-NH-CH(CH₂-CH₂-C(=O)-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-Orn-^DDab-Dab-)</p> <p>O=C-CH(CH₂-CH₂-NH-)-Dab-Thr-Dab-</p> <p>cyclo(-NH-CH(CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>
Ex. 12 a)	<p>S-S-</p> <p>Ac-Trp-Cys-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Cys-Dab-Thr-Dab-</p> <p>HN-CH(CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p> <p>cyclo(-NH-CH(CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>
Ex. 13 a)	<p>S-S-</p> <p>Ac-Trp-Cys-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Cys-Orn-Thr-Dab-</p> <p>HN-CH(CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p> <p>cyclo(-NH-CH(CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>
Ex. 14 b)	<p>O=C-NH-</p> <p>Ac-Trp-Glu-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-Dab-Thr-Dab-</p> <p>HN-CH(CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p> <p>cyclo(-NH-CH(CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>
Ex. 15 a)	<p>S-S-</p> <p>Trp-Cys-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Cys-Dab-Thr-Dab-</p> <p>HN-CH(CH₂-CH₂-C(=O)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p> <p>cyclo(-NH-CH(CH₂-CH₂-C(=O)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p>

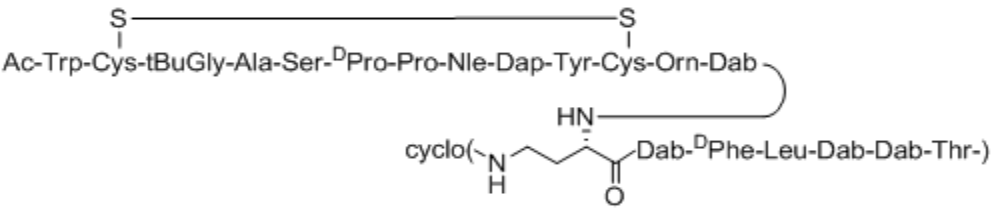
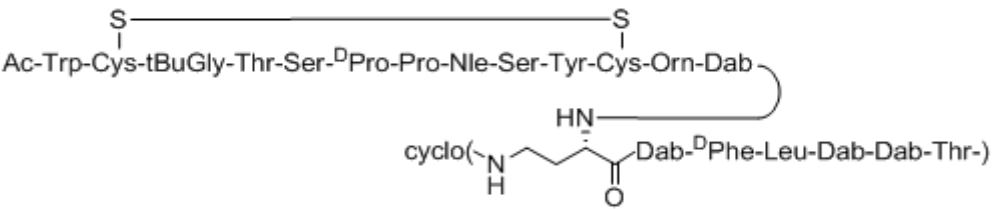
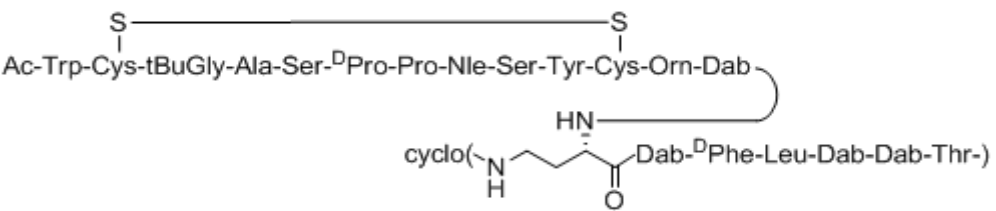
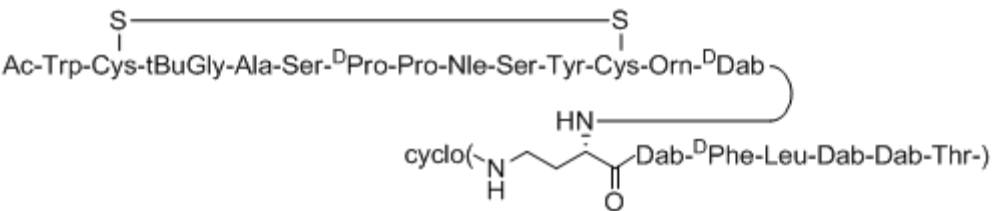
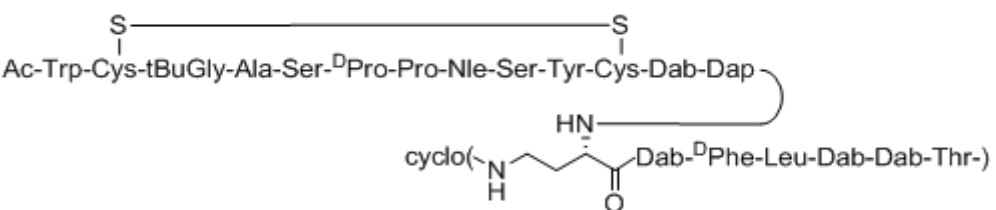
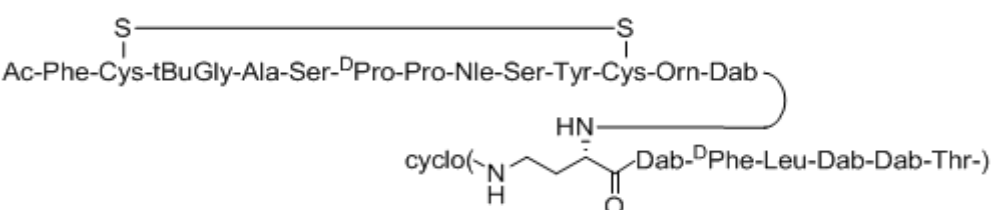
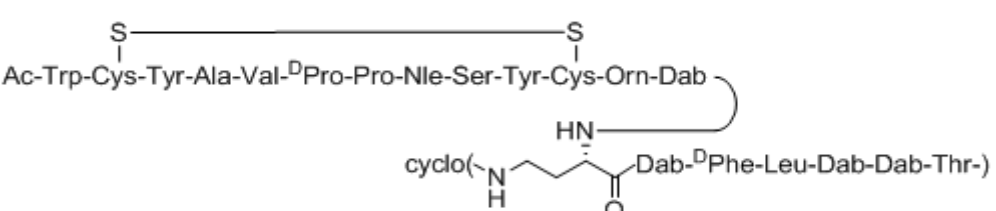
Ex. 16 a)	<p>Trp-Cys-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Cys-Dab-Thr-Dap</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>
Ex. 17 a)	<p>Ac-Trp-Cys-tBuGly-Ala-Ser-^DAla-Pro-Nle-Ser-Tyr-Cys-Dab-Thr-Dab</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>
Ex. 18 a)	<p>Ac-Trp-Cys-tBuGly-Ser-Thr-^DPro-Pro-Nle-Ala-Tyr-Cys-Dab-Thr-Dab</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>
Ex. 19 a)	<p>Ac-Trp-Cys-tBuGly-Dap-Thr-^DPro-Pro-Nle-Gly-Tyr-Cys-Dab-Thr-Dab</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>
Ex. 20	<p>Ac-Trp-Cys-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Cys-Dab-Thr-Dap</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>
Ex. 21 a)	<p>Ac-Trp-Cys-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Cys-Dab-<i>allo</i>Thr-Dab</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>
Ex. 22 a)	<p>Ac-Trp-Cys-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Cys-Dab-Gly-Dab</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>

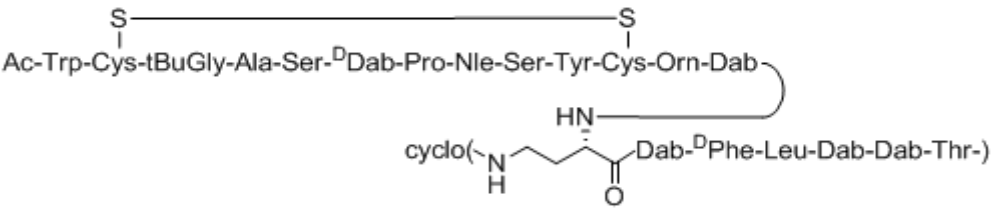
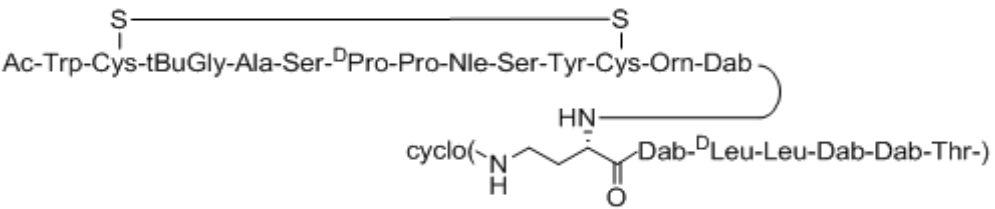
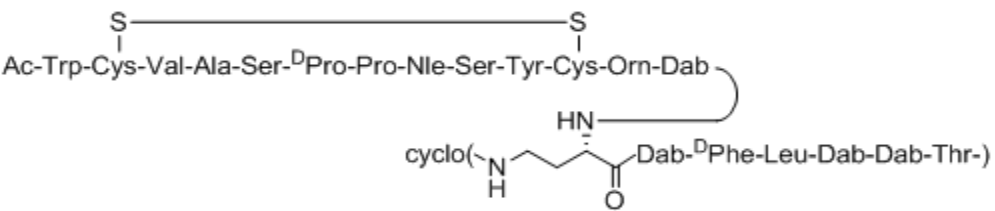
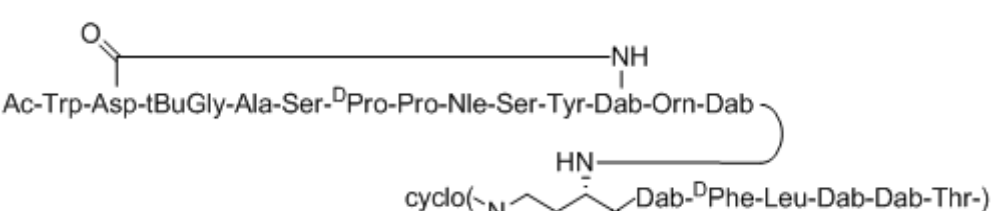
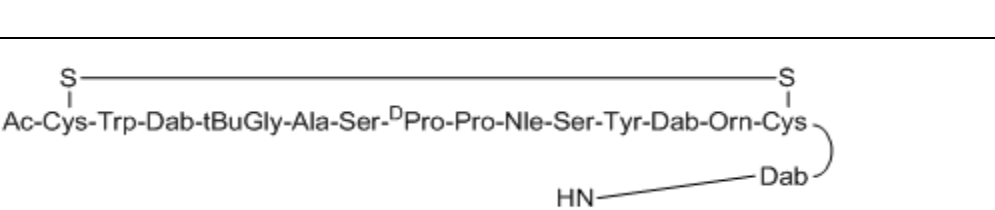
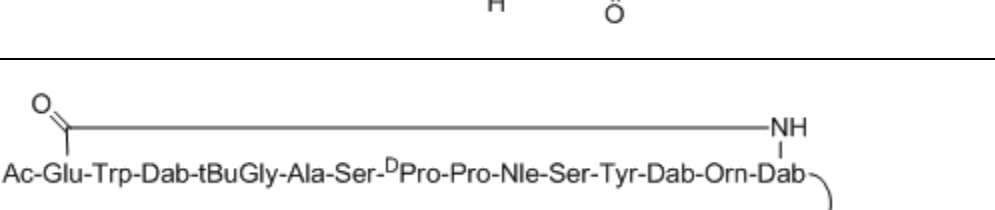
<p>Ex. 23 a)</p>	
<p>Ex. 24 a) c)</p>	
<p>Ex. 25 a) c)</p>	
<p>Ex. 26 a)</p>	
<p>Ex. 27 a)</p>	

<p>Ex. 28 a)</p>	 <p>Chemical structure of a peptide with a disulfide bridge (S-S) connecting the Cys residues. The main chain is Ac-Trp-Cys-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Cys-Ala-Dab-Thr. A side chain is attached to the second Cys residue, consisting of a cyclic peptide: cyclo(-NH-CH₂-CH₂-CH₂-NH-CO-Dab-^DPhe-Leu-Dab-Dab-Thr-). The side chain is linked to the main chain via an amide bond (HN) to the Dab residue in the cycle.</p>
<p>Ex. 29 a)</p>	 <p>Chemical structure of a peptide with a disulfide bridge (S-S) connecting the Cys residues. The main chain is Ac-Trp-Cys-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Cys-^DAla-Dab-Thr. A side chain is attached to the second Cys residue, consisting of a cyclic peptide: cyclo(-NH-CH₂-CH₂-CH₂-NH-CO-Dab-^DPhe-Leu-Dab-Dab-Thr-). The side chain is linked to the main chain via an amide bond (HN) to the Dab residue in the cycle.</p>
<p>Ex. 30 a)</p>	 <p>Chemical structure of a peptide with a disulfide bridge (S-S) connecting the Cys residues. The main chain is Ac-Trp-Cys-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Cys-Aib-Dab-Thr. A side chain is attached to the second Cys residue, consisting of a cyclic peptide: cyclo(-NH-CH₂-CH₂-CH₂-NH-CO-Dab-^DPhe-Leu-Dab-Dab-Thr-). The side chain is linked to the main chain via an amide bond (HN) to the Dab residue in the cycle.</p>
<p>Ex. 31 a)</p>	 <p>Chemical structure of a peptide with a disulfide bridge (S-S) connecting the Cys residues. The main chain is Ac-Trp-Cys-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Cys-Dab-Gly-Thr. A side chain is attached to the second Cys residue, consisting of a cyclic peptide: cyclo(-NH-CH₂-CH₂-CH₂-NH-CO-Dab-^DPhe-Leu-Dab-Dab-Thr-). The side chain is linked to the main chain via an amide bond (HN) to the Dab residue in the cycle.</p>
<p>Ex. 32 a)</p>	 <p>Chemical structure of a peptide with a disulfide bridge (S-S) connecting the Cys residues. The main chain is Ac-Trp-Cys-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Cys-Dab-Thr-^DAla. A side chain is attached to the second Cys residue, consisting of a cyclic peptide: cyclo(-NH-CH₂-CH₂-CH₂-NH-CO-Dab-^DPhe-Leu-Dab-Dab-Thr-). The side chain is linked to the main chain via an amide bond (HN) to the Dab residue in the cycle.</p>
<p>Ex. 33 a)</p>	 <p>Chemical structure of a peptide with a disulfide bridge (S-S) connecting the Cys residues. The main chain is Ac-Trp-Cys-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Cys-Dab-Thr-Dab. A side chain is attached to the second Cys residue, consisting of a cyclic peptide: cyclo(-NH-CH₂-CH₂-CH₂-NH-CO-Dab-^DPhe-Leu-Dab-Dab-Thr-). The side chain is linked to the main chain via an amide bond (HN) to the Gly residue in the cycle.</p>

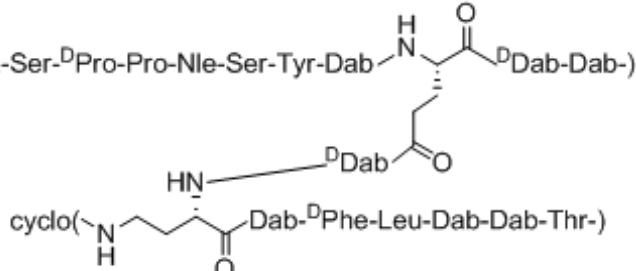
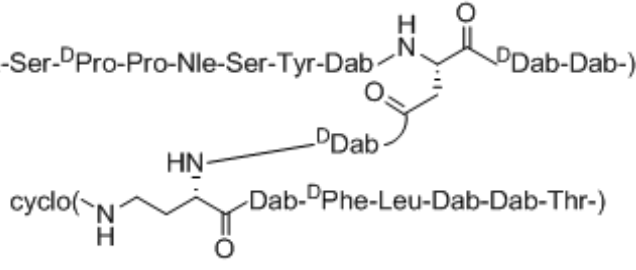
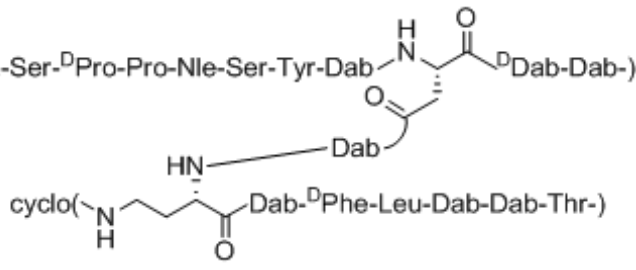
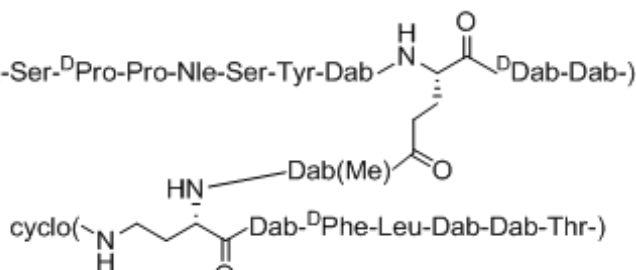
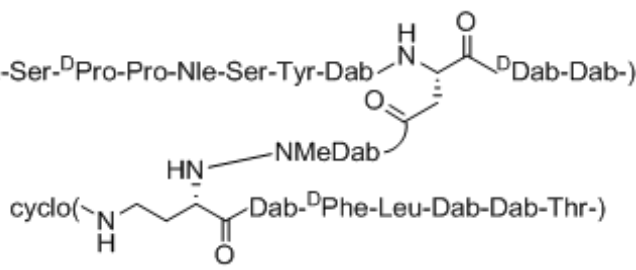
<p>Ex. 34 a)</p>	
<p>Ex. 35 a)</p>	
<p>Ex. 36 a)</p>	
<p>Ex. 37 a)</p>	
<p>Ex. 38 a)</p>	
<p>Ex. 39 a)</p>	

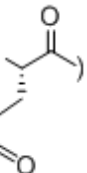
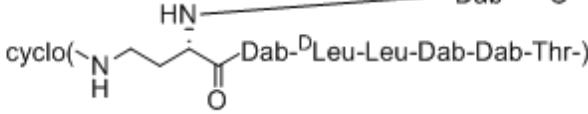
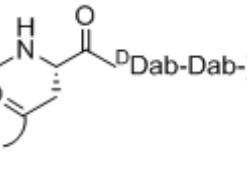
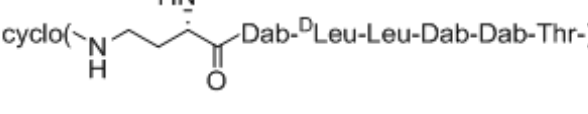
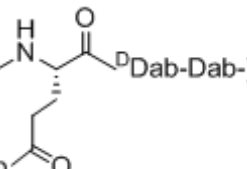
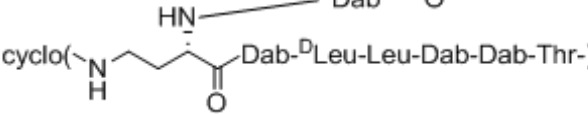
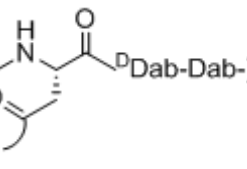
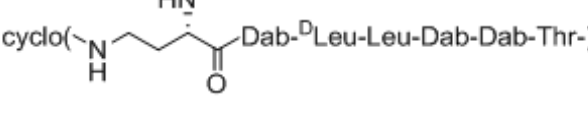
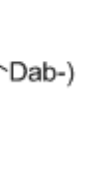
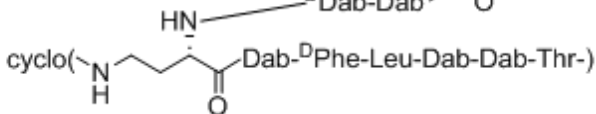
Ex. 40 a)	<p>Chemical structure of Ex. 40a: A linear peptide chain Glu-Trp-Cys-tBuGly-Ala-Ser-DPro-Pro-Nle-Ser-Tyr-Cys-Dab-Thr with a disulfide bond between the two Cys residues. A side chain Dab is attached to the Cys residue. A cyclic peptide is attached to the Dab residue: cyclo(-NH-CH₂-CH₂-CH(NH-Dab)-C(=O)-Dab-DPhe-Leu-Dab-Dab-Thr-).</p>
Ex. 41 a)	<p>Chemical structure of Ex. 41a: A linear peptide chain Ac-Ala-Trp-Cys-tBuGly-Ala-Ser-DPro-Pro-Nle-Ser-Tyr-Cys-Dab-Thr with a disulfide bond between the two Cys residues. A side chain Dab is attached to the Cys residue. A cyclic peptide is attached to the Dab residue: cyclo(-NH-CH₂-CH₂-CH(NH-Dab)-C(=O)-Dab-DPhe-Leu-Dab-Dab-Thr-).</p>
Ex. 42 a)	<p>Chemical structure of Ex. 42a: A linear peptide chain Dab-Trp-Cys-tBuGly-Ala-Ser-DPro-Pro-Nle-Ser-Tyr-Cys-Dab-Thr with a disulfide bond between the two Cys residues. A side chain Dab is attached to the Cys residue. A cyclic peptide is attached to the Dab residue: cyclo(-NH-CH₂-CH₂-CH(NH-Dab)-C(=O)-Dab-DPhe-Leu-Dab-Dab-Thr-).</p>
Ex. 43 a)	<p>Chemical structure of Ex. 43a: A linear peptide chain Thr-Trp-Cys-tBuGly-Ala-Ser-DPro-Pro-Nle-Ser-Tyr-Cys-Dab-Thr with a disulfide bond between the two Cys residues. A side chain Dab is attached to the Cys residue. A cyclic peptide is attached to the Dab residue: cyclo(-NH-CH₂-CH₂-CH(NH-Dab)-C(=O)-Dab-DPhe-Leu-Dab-Dab-Thr-).</p>
Ex. 44 a)	<p>Chemical structure of Ex. 44a: A linear peptide chain Aib-Trp-Cys-tBuGly-Ala-Ser-DPro-Pro-Nle-Ser-Tyr-Cys-Dab-Thr with a disulfide bond between the two Cys residues. A side chain Dab is attached to the Cys residue. A cyclic peptide is attached to the Dab residue: cyclo(-NH-CH₂-CH₂-CH(NH-Dab)-C(=O)-Dab-DPhe-Leu-Dab-Dab-Thr-).</p>
Ex. 45 a)	<p>Chemical structure of Ex. 45a: A linear peptide chain Trp-Cys-tBuGly-Ala-Ser-DPro-Pro-Nle-Ser-Tyr-Cys-Orn-Dab with a disulfide bond between the two Cys residues. A side chain Dab is attached to the Cys residue. A cyclic peptide is attached to the Dab residue: cyclo(-NH-CH₂-CH₂-CH(NH-Dab)-C(=O)-Dab-DPhe-Leu-Dab-Dab-Thr-).</p>

<p>Ex. 46 a)</p>	 <p>Ac-Trp-Cys-tBuGly-Ala-Ser-^DPro-Pro-Nle-Dap-Tyr-Cys-Orn-Dab</p> <p>cyclo(-NH-CH(CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)-)</p>
<p>Ex. 47 a)</p>	 <p>Ac-Trp-Cys-tBuGly-Thr-Ser-^DPro-Pro-Nle-Ser-Tyr-Cys-Orn-Dab</p> <p>cyclo(-NH-CH(CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)-)</p>
<p>Ex. 48 a)</p>	 <p>Ac-Trp-Cys-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Cys-Orn-Dab</p> <p>cyclo(-NH-CH(CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)-)</p>
<p>Ex. 49 a)</p>	 <p>Ac-Trp-Cys-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Cys-Orn-^DDab</p> <p>cyclo(-NH-CH(CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)-)</p>
<p>Ex. 50 a)</p>	 <p>Ac-Trp-Cys-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Cys-Dab-Dap</p> <p>cyclo(-NH-CH(CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)-)</p>
<p>Ex. 51 a)</p>	 <p>Ac-Phe-Cys-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Cys-Orn-Dab</p> <p>cyclo(-NH-CH(CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)-)</p>
<p>Ex. 52 a)</p>	 <p>Ac-Trp-Cys-Tyr-Ala-Val-^DPro-Pro-Nle-Ser-Tyr-Cys-Orn-Dab</p> <p>cyclo(-NH-CH(CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)-)</p>

Ex. 53 a)	 <p>Chemical structure of a linear peptide with a disulfide bridge (S-S) connecting the Cys residues. The sequence is Ac-Trp-Cys-tBuGly-Ala-Ser-Dab-Pro-Nle-Ser-Tyr-Cys-Orn-Dab. A side chain is attached to the Ser-Dab residue, consisting of a cyclic peptide: cyclo(-NH-CH2-CH2-CH(NH-C(=O)-Dab-DPhe-Leu-Dab-Dab-Thr-)-). The side chain is connected to the Ser-Dab residue via its NH group.</p>
Ex. 54 a)	 <p>Chemical structure of a linear peptide with a disulfide bridge (S-S) connecting the Cys residues. The sequence is Ac-Trp-Cys-tBuGly-Ala-Ser-DPro-Pro-Nle-Ser-Tyr-Cys-Orn-Dab. A side chain is attached to the Ser-Dab residue, consisting of a cyclic peptide: cyclo(-NH-CH2-CH2-CH(NH-C(=O)-Dab-DLeu-Leu-Dab-Dab-Thr-)-). The side chain is connected to the Ser-Dab residue via its NH group.</p>
Ex. 55 a)	 <p>Chemical structure of a linear peptide with a disulfide bridge (S-S) connecting the Cys residues. The sequence is Ac-Trp-Cys-Val-Ala-Ser-DPro-Pro-Nle-Ser-Tyr-Cys-Orn-Dab. A side chain is attached to the Ser-Dab residue, consisting of a cyclic peptide: cyclo(-NH-CH2-CH2-CH(NH-C(=O)-Dab-DPhe-Leu-Dab-Dab-Thr-)-). The side chain is connected to the Ser-Dab residue via its NH group.</p>
Ex. 56 b)	 <p>Chemical structure of a linear peptide with an amide bridge (NH-C(=O)-) connecting the Asp and Tyr residues. The sequence is Ac-Trp-Asp-tBuGly-Ala-Ser-DPro-Pro-Nle-Ser-Tyr-Dab-Orn-Dab. A side chain is attached to the Ser-Dab residue, consisting of a cyclic peptide: cyclo(-NH-CH2-CH2-CH(NH-C(=O)-Dab-DPhe-Leu-Dab-Dab-Thr-)-). The side chain is connected to the Ser-Dab residue via its NH group.</p>
Ex. 57 a)	 <p>Chemical structure of a linear peptide with a disulfide bridge (S-S) connecting the Cys residues. The sequence is Ac-Cys-Trp-Dab-tBuGly-Ala-Ser-DPro-Pro-Nle-Ser-Tyr-Dab-Orn-Cys. A side chain is attached to the Ser-Dab residue, consisting of a cyclic peptide: cyclo(-NH-CH2-CH2-CH(NH-C(=O)-Dab-DPhe-Leu-Dab-Dab-Thr-)-). The side chain is connected to the Ser-Dab residue via its NH group.</p>
Ex. 58 b)	 <p>Chemical structure of a linear peptide with an amide bridge (NH-C(=O)-) connecting the Glu and Tyr residues. The sequence is Ac-Glu-Trp-Dab-tBuGly-Ala-Ser-DPro-Pro-Nle-Ser-Tyr-Dab-Orn-Dab. A side chain is attached to the Ser-Dab residue, consisting of a cyclic peptide: cyclo(-NH-CH2-CH2-CH(NH-C(=O)-Dab-DPhe-Leu-Dab-Dab-Thr-)-). The side chain is connected to the Ser-Dab residue via its NH group.</p>

<p>Ex. 59 b)</p>	<p>HN— Ac-Dab-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-Orn-Glu— HN— cyclo(-NH-CH₂-CH₂-CH(NH-CO-Dab-^DPhe-Leu-Dab-Dab-Thr)-)</p>
<p>Ex. 60 a)</p>	<p>S— Ac-Cys-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-Orn-Cys— HN— cyclo(-NH-CH₂-CH₂-CH(NH-CO-Dab-^DPhe-Leu-Dab-Dab-Thr)-)</p>
<p>Ex. 61 a)</p>	<p>S— Ac-Cys-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-Orn-Cys— HN— cyclo(-NH-CH₂-CH₂-CH(NH-CO-Dab-^DPhe-Leu-Dab-Dab-Thr)-)</p>
<p>Ex. 62 a)</p>	<p>S— Ac-Cys-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-Orn-Cys— HN— cyclo(-NH-CH₂-CH₂-CH(NH-CO-Dab-^DPhe-Leu-Dab-Dab-Thr)-)</p>
<p>Ex. 63 a)</p>	<p>S— Ac-^DCys-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-Orn-Cys— HN— cyclo(-NH-CH₂-CH₂-CH(NH-CO-Dab-^DPhe-Leu-Dab-Dab-Thr)-)</p>
<p>Ex. 64</p>	<p>cyclo(-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-NH-CH₂-CH₂-CO-^DDab-Dab-) HN— cyclo(-NH-CH₂-CH₂-CH(NH-CO-Dab-^DPhe-Leu-Dab-Dab-Thr)-)</p>

Ex. 65	<p>cyclo(-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-NH-CH(^DDab)-CO-^DDab-Dab-)</p> 
Ex. 66	<p>cyclo(-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-NH-CH(^DDab)-CO-^DDab-Dab-)</p> 
Ex. 67	<p>cyclo(-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-NH-CH(^DDab)-CO-^DDab-Dab-)</p> 
Ex. 68	<p>cyclo(-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-NH-CH(^DDab)-CO-^DDab-Dab-)</p> 
Ex. 69	<p>cyclo(-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-NH-CH(^DDab)-CO-^DDab-Dab-)</p> 

Ex. 70	<p>cyclo(-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-Orn-^DDab-NH-)</p> <p></p>
Ex. 71	<p>cyclo(-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-NH-)</p> <p></p>
Ex. 72	<p>cyclo(-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-NH-)</p> <p></p>
Ex. 73	<p>cyclo(-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-NH-)</p> <p></p>
Ex. 74	<p>cyclo(-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-Orn-NH-)</p> <p></p>

Ex. 75	<p>cyclo(-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-Orn-^DDab-NH-CH(CH₂CO-)-)</p> <p>HN-CH(CH₂CO-)-Dab-Thr-CH(CH₂CO-)-</p> <p>cyclo(-NH-CH(CH₂CO-)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>
Ex. 76	<p>cyclo(-Trp-Dab-Val-Ala-Ser-^DPro-Pro-Leu-Ser-Tyr-Dab-NH-CH(CH₂CO-)-^DDab-Dab-)</p> <p>O=C-CH(CH₂CO-)-Dab-Sar-CH(CH₂CO-)-</p> <p>cyclo(-NH-CH(CH₂CO-)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p>
Ex. 77	<p>cyclo(-Trp-Dab-tBuGly-Ala-Tyr-^DPro-Pro-Leu-Ser-Tyr-Dab-Orn-NH-CH(CH₂CO-)-Dab-)</p> <p>O=C-CH(CH₂CO-)-Dab-Sar-CH(CH₂CO-)-</p> <p>cyclo(-NH-CH(CH₂CO-)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p>
Ex. 78	<p>cyclo(-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-Orn-NH-CH(CH₂CO-)-Dab-)</p> <p>CH(CH₂CO-)-^DDab-Dab-CH(CH₂CO-)-</p> <p>cyclo(-NH-CH(CH₂CO-)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p>
Ex. 79 a)	<p>S-CH₂-CH₂-S-CH₂-CH₂-</p> <p>cyclo(-Trp-Cys-Val-Ala-Ser-^DPro-Pro-Leu-Ser-Tyr-Cys-Gly-NH-CH(CH₂CO-)-Dab-)</p> <p>CH(CH₂CO-)-Dab-Dab-CH(CH₂CO-)-</p> <p>cyclo(-NH-CH(CH₂CO-)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p>

Ex. 80	<p>cyclo(-Trp-Dab-tBuGly-Ala-Ser-^DAzt-Pro-Val-Thr-Tyr-Dab-NH-CH(^D)-C(=O)-^DDab-Dab-)</p> <p>cyclo(-NH-CH(^D)-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p> <p>HN-Dab-Thr-Dab</p>
Ex. 81	<p>cyclo(-Trp-Dab-tBuGly-Ala-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-NH-CH(^D)-C(=O)-^DDab-Dab-)</p> <p>cyclo(-NH-CH(^D)-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p> <p>HN-Dab-^DThr-Dab</p>
Ex. 82	<p>cyclo(-Trp-Dab-tBuGly-Ala-Ser-^DAzt-Pro-Val-Thr-Tyr-Dab-NH-CH(^D)-C(=O)-^DDab-Dab-)</p> <p>cyclo(-NH-CH(^D)-C(=O)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p> <p>HN-Dab-Thr-Dab</p>
Ex. 83	<p>cyclo(-Trp-Dab-tBuGly-Ser-Ser-^DPro-Pro-Nle-Dap-Tyr-Dab-NH-CH(^D)-C(=O)-^DDab-Dab-)</p> <p>cyclo(-NH-CH(^D)-C(=O)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p> <p>HN-Dab-Thr-Dab</p>
Ex. 84	<p>cyclo(-Trp-Dab-tBuGly-Dap-Ser-^DPro-Pro-Nle-Ser-Tyr-Dab-NH-CH(^D)-C(=O)-^DDab-Dab-)</p> <p>cyclo(-NH-CH(^D)-C(=O)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p> <p>HN-Dab-Thr-Dab</p>

Ex. 85	<p>cyclo(-Trp-Dab-Val-Ala-Ser-^DPro-Pro-Val-Dap-Tyr-Dab-NH-CH(CH₂CH₂CO^DDab-Dab)-CO-CH₂CH₂CO-NH-Dab-Thr-Dab-)-</p> <p>cyclo(-NH-CH₂CH₂CH₂CO-Dab-^DLeu-Leu-Dab-Dab-Thr-)-</p>
Ex. 86 b)	<p>cyclo(-Trp-Dab-tBuGly-Asp-Thr-^DAzt-Pro-Val-Dab-Tyr-Dab-NH-CH(CH₂CH₂CO^DDab-Dab)-CO-CH₂CH₂CO-NH-Dab-Thr-Dab-)-</p> <p>cyclo(-NH-CH₂CH₂CH₂CO-Dab-^DPhe-Leu-Dab-Dab-Thr-)-</p>
Ex. 87 b)	<p>cyclo(-Trp-Dab-tBuGly-Asp-Thr-^DAzt-Pro-Val-Dab-Tyr-Dab-NH-CH(CH₂CH₂CO^DDab-Dab)-CO-CH₂CH₂CO-NH-Dab-Thr-Dab-)-</p> <p>cyclo(-NH-CH₂CH₂CH₂CO-Dab-^DLeu-Leu-Dab-Dab-Thr-)-</p>
Ex. 88	<p>cyclo(-Trp-Dab-Val-Thr-Ser-^DAla-Pro((4R)OH)-Leu-Ser-Tyr-Dab-NH-CH(CH₂CH₂CO^DDab-Dab)-CO-CH₂CH₂CO-NH-Dab-Thr-Dab-)-</p> <p>cyclo(-NH-CH₂CH₂CH₂CO-Dab-^DLeu-Leu-Dab-Dab-Thr-)-</p>
Ex. 89	<p>cyclo(-Trp-Dab-Val-Thr-Ser-^DPro-Pro-Leu-Ser-Tyr-Dab-NH-CH(CH₂CH₂CO^DDab-Dab)-CO-CH₂CH₂CO-NH-Dab-Thr-NMeDab-)-</p> <p>cyclo(-NH-CH₂CH₂CH₂CO-Dab-^DLeu-Leu-Dab-Dab-Thr-)-</p>

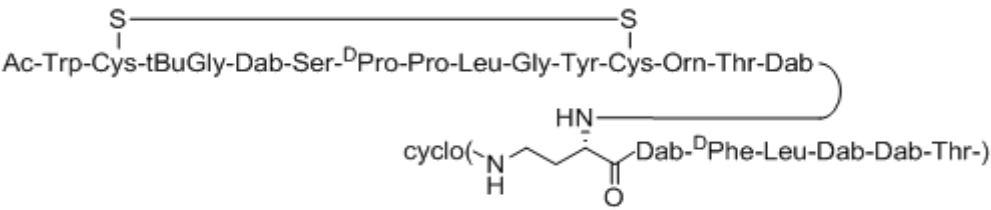
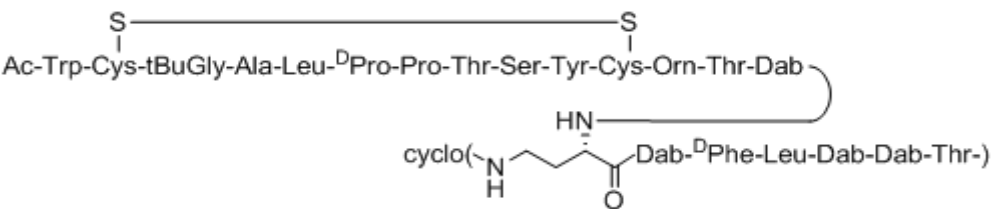
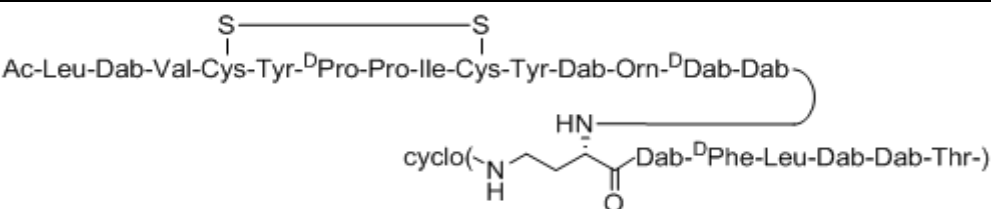
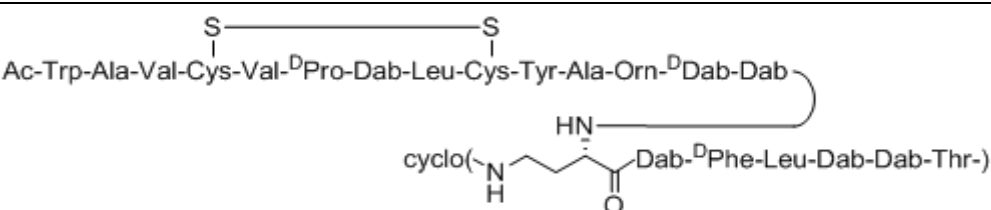
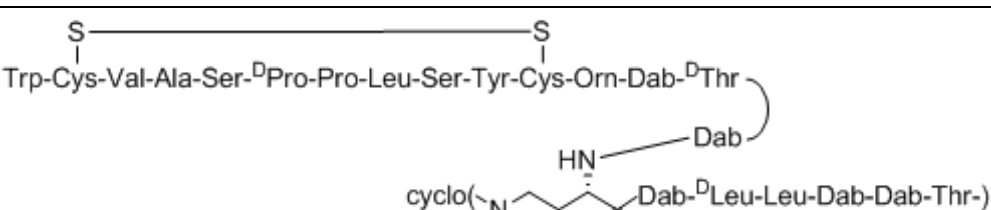
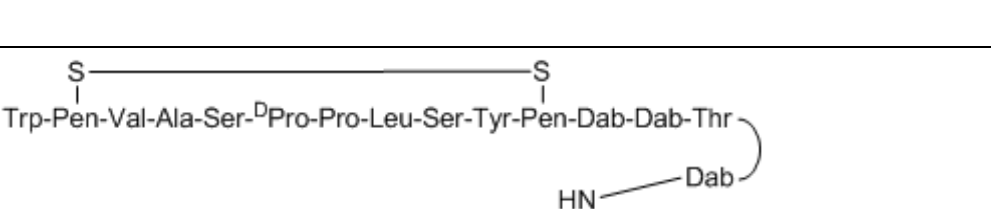
Ex. 90	<p>cyclo(-Trp-Dab-Val-Thr-Ser-^DPro-Pro-Leu-Ser-Tyr-Gly-NH-CH(CH₂CH₂CO^DDab-Dab-)-CO-CH₂CH₂CO-NH-Dab-Thr-Dab-)-CO-CH₂CH₂CO-NH-cyclo(-NH-CH₂CH₂CH₂CO-Dab-^DLeu-Leu-Dab-Dab-Thr-)-CO-CH₂CH₂CO-NH-)</p>
Ex. 91	<p>cyclo(-Trp-Dab-Val-Thr-Ser-^DPro-Pro-Leu-Thr-Tyr-Dab-NH-CH(CH₂CH₂CO^DDab-Dab-)-CO-CH₂CH₂CO-NH-Dab-Thr-Dab-)-CO-CH₂CH₂CO-NH-cyclo(-NH-CH₂CH₂CH₂CO-Dab-^DLeu-Leu-Dab-Dab-Thr-)-CO-CH₂CH₂CO-NH-)</p>
Ex. 92	<p>cyclo(-Trp-Dab-Val-Thr-Ser-^DPro-Pro-Leu-Ser-Tyr-Dab-NH-CH(CH₂CH₂CO^DDab-Dab-)-CO-CH₂CH₂CO-NH-Dab-Thr-Dab-)-CO-CH₂CH₂CO-NH-cyclo(-NH-CH₂CH₂CH₂CO-Dab-^DLeu-Leu-Dab-Dab-Thr-)-CO-CH₂CH₂CO-NH-)</p>
Ex. 93	<p>cyclo(-Trp-Dab-Val-Ser-Ser-^DAzt-Pro((4R)OH)-Val-Thr-Tyr-Dab-NH-CH(CH₂CH₂CO^DDab-Dab-)-CO-CH₂CH₂CO-NH-Dab-Thr-Dab-)-CO-CH₂CH₂CO-NH-cyclo(-NH-CH₂CH₂CH₂CO-Dab-^DLeu-Leu-Dab-Dab-Thr-)-CO-CH₂CH₂CO-NH-)</p>
Ex. 94	<p>cyclo(-Trp-Dab-Val-Thr-Ser-^DPro-Pro-Leu-Ser-Tyr-Dab-Orn-Gly-NH-CH(CH₂CH₂CO^DDab-Dab-)-CO-CH₂CH₂CO-NH-Dab-Thr-Dab-)-CO-CH₂CH₂CO-NH-cyclo(-NH-CH₂CH₂CH₂CO-Dab-^DLeu-Leu-Dab-Dab-Thr-)-CO-CH₂CH₂CO-NH-)</p>

Ex. 95	<p>cyclo(-Trp-Dab-Val-Tyr-Ser-^DPro-Pro-Leu-Ser-Tyr-Dab-NH-CH(^DCH₂CO^DDab-Dab)-CO-CH₂-CH₂-CO-NH-Dab-Thr-Dab-)-</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-CO-NH-Dab-Thr-Dab-)-</p>
Ex. 96	<p>cyclo(-Trp-Dab-Val-Thr-Tyr-^DPro-Pro-Leu-Ser-Tyr-Dab-NH-CH(^DCH₂CO^DDab-Dab)-CO-CH₂-CH₂-CO-NH-Dab-Thr-Dab-)-</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-CO-NH-Dab-Thr-Dab-)-</p>
Ex. 97	<p>cyclo(-Trp-Dab-Tyr-Ala-Val-^DDab-Arg-Leu-Ser-Tyr-Dab-NH-CH(^DCH₂CO^DDab-Dab)-CO-CH₂-CH₂-CO-NH-Dab-Thr-Dab-)-</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-CO-NH-Dab-Thr-Dab-)-</p>
Ex. 98 a)	<p>cyclo(-Trp-Cys-tBuGly-Thr-Ser-^DPro-Pro-Leu-Ser-Tyr-Cys-NH-CH(^DCH₂CO^DDab-Dab)-CO-CH₂-CH₂-CO-NH-Dab-Thr-Dab-)-</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-CO-NH-Dab-Thr-Dab-)-</p>
Ex. 99	<p>cyclo(-Leu-Dab-Val-Thr-Tyr-^DPro-Pro-Leu-Ser-Tyr-Dab-NH-CH(^DCH₂CO^DDab-Dab)-CO-CH₂-CH₂-CO-NH-Dab-Thr-Dab-)-</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-CO-NH-Dab-Thr-Dab-)-</p>

Ex. 100	<p>cyclo(-Trp-Thr-tBuGly-Ala-Thr-^DAzt-Pro-Val-Thr-Tyr-Thr-NH-CH(CH₂CH₂CO-D-Thr-Thr)-CO-)</p> <p>cyclo(-NH-CH(CH₂CH₂CO-Dab-D^DLeu-Leu-Dab-Dab-Thr)-CO-NH-Dab-Thr-Dab-CH(CH₂CH₂CO-D-Thr-Thr)-CO-)</p>
Ex. 101	<p>cyclo(-Trp-Dab-tBuGly-Ala-Ser-^DAzt-Pro-Val-Thr-Tyr-Dab-NH-CH(CH₂CH₂CO-Dab-Dab)-CO-)</p> <p>cyclo(-NH-CH(CH₂CH₂CO-Dab-D^DLeu-Leu-Dab-Dab-Thr)-CO-NH-Dab-Thr-Dab-CH(CH₂CH₂CO-Dab-Dab)-CO-)</p>
Ex. 102	<p>cyclo(-Trp-Hse-tBuGly-Ala-Ser-^DAla-Pro-Val-Thr-Tyr-Hse-NH-CH(CH₂CH₂CO-Dab-Hse)-CO-)</p> <p>cyclo(-NH-CH(CH₂CH₂CO-Dab-D^DLeu-Leu-Dab-Dab-Thr)-CO-NH-Dab-Thr-Dab-CH(CH₂CH₂CO-Dab-Hse)-CO-)</p>
Ex. 103	<p>cyclo(-Trp-Hse-tBuGly-Ala-Ser-^DAla-Pro-Val-Thr-Tyr-Dab-NH-CH(CH₂CH₂CO-Dab-Hse)-CO-)</p> <p>cyclo(-NH-CH(CH₂CH₂CO-Dab-D^DLeu-Leu-Dab-Dab-Thr)-CO-NH-Dab-Thr-Dab-CH(CH₂CH₂CO-Dab-Hse)-CO-)</p>
Ex. 104 a)	<p>Trp-Cys-tBuGly-Ser-Ser-^DPro-Pro-Nle-Ser-Tyr-Cys-Orn-^DAla-^DDab</p> <p>cyclo(-NH-CH(CH₂CH₂CO-Dab-D^DPhe-Leu-Dab-Dab-Thr)-CO-NH-)</p>

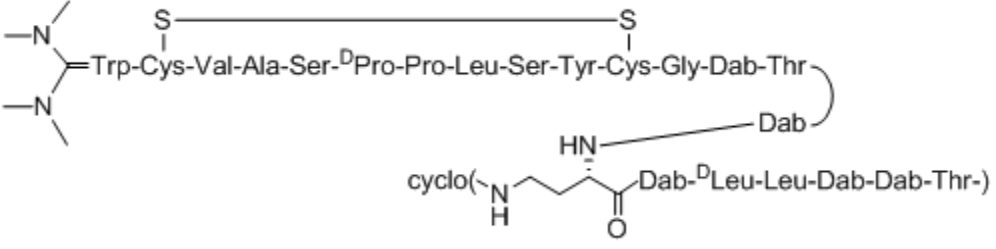
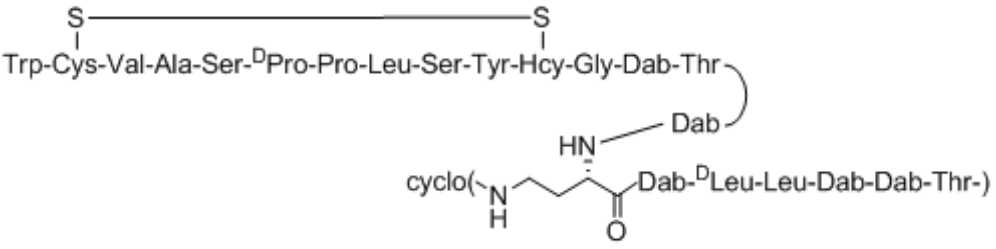
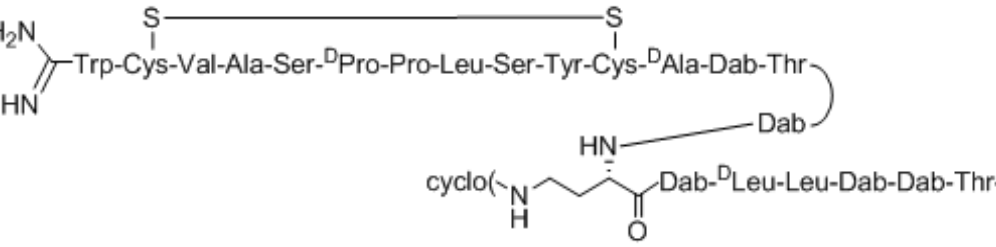
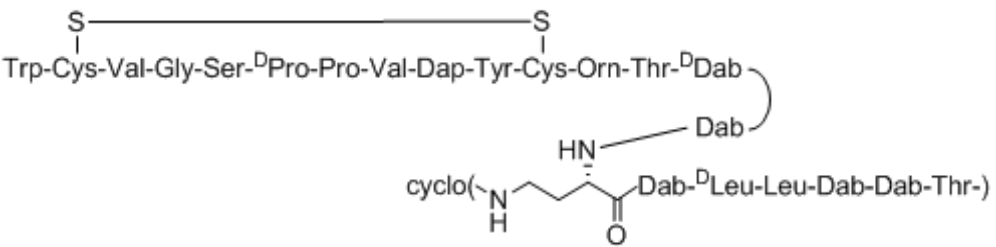
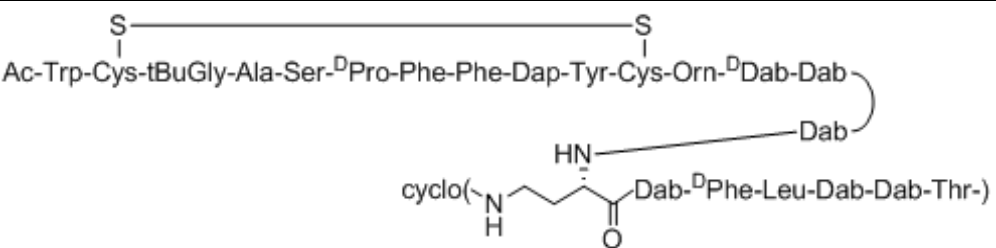
<p>Ex. 105 a)</p>	<p>Trp-Cys-tBuGly-Ser-Ser-^DPro-Pro-Nle-Ser-Tyr-Cys-Orn-^DPhe-^DDab</p> <p>cyclo($\text{HN}-\text{CH}_2-\text{CH}_2-\text{CH}(\text{Dab-}^{\text{D}}\text{Phe-Leu-Dab-Thr-})-\text{C}(=\text{O})$)</p>
<p>Ex. 106 a)</p>	<p>Trp-Cys-tBuGly-Ser-Ser-^DPro-Pro-Nle-Ser-Tyr-Cys-^DDab-Dab-^DDab</p> <p>cyclo($\text{HN}-\text{CH}_2-\text{CH}_2-\text{CH}(\text{Dab-}^{\text{D}}\text{Phe-Leu-Dab-Thr-})-\text{C}(=\text{O})$)</p>
<p>Ex. 107 a)</p>	<p>Trp-Cys-tBuGly-Ser-Ser-^DPro-Pro-Nle-Ser-Tyr-Cys-^DThr-Dab-^DDab</p> <p>cyclo($\text{HN}-\text{CH}_2-\text{CH}_2-\text{CH}(\text{Dab-}^{\text{D}}\text{Phe-Leu-Dab-Thr-})-\text{C}(=\text{O})$)</p>
<p>Ex. 108 a)</p>	<p>Trp-Cys-tBuGly-Ser-Ser-^DPro-Pro-Nle-Ser-Tyr-Cys-Glu-Dab-Dab</p> <p>cyclo($\text{HN}-\text{CH}_2-\text{CH}_2-\text{CH}(\text{Dab-}^{\text{D}}\text{Phe-Leu-Dab-Thr-})-\text{C}(=\text{O})$)</p>
<p>Ex. 109 a)</p>	<p>Trp-Cys-tBuGly-Ser-Ser-^DPro-Pro-Nle-Ser-Tyr-Cys-^DTyr-Dab-Dab</p> <p>cyclo($\text{HN}-\text{CH}_2-\text{CH}_2-\text{CH}(\text{Dab-}^{\text{D}}\text{Phe-Leu-Dab-Thr-})-\text{C}(=\text{O})$)</p>
<p>Ex. 110 a)</p>	<p>Ac-Trp-Cys-tBuGly-Ala-Ser-^DPro-Leu-Phe-Dap-Tyr-Cys-Orn-^DDab-Dab</p> <p>cyclo($\text{HN}-\text{CH}_2-\text{CH}_2-\text{CH}(\text{Dab-}^{\text{D}}\text{Phe-Leu-Dab-Thr-})-\text{C}(=\text{O})$)</p>

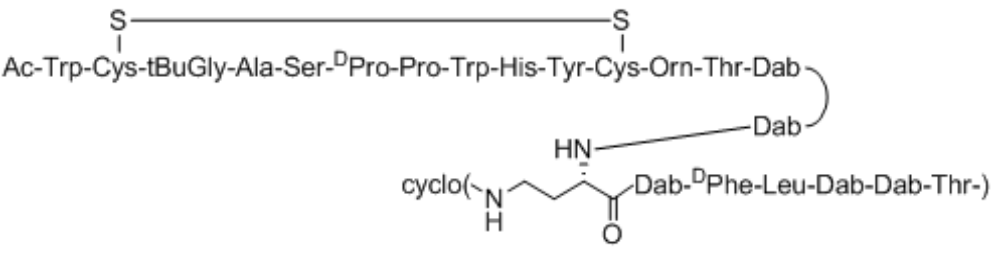
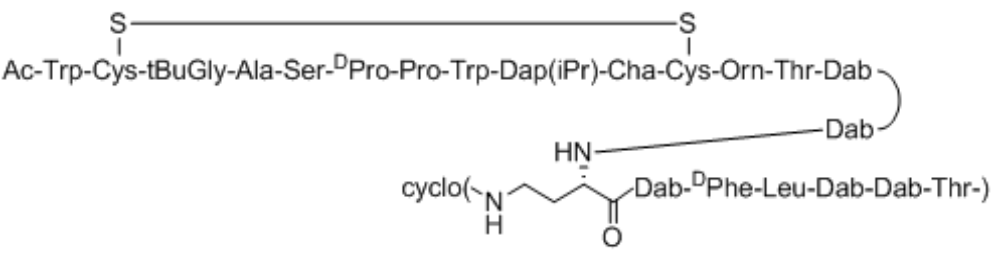
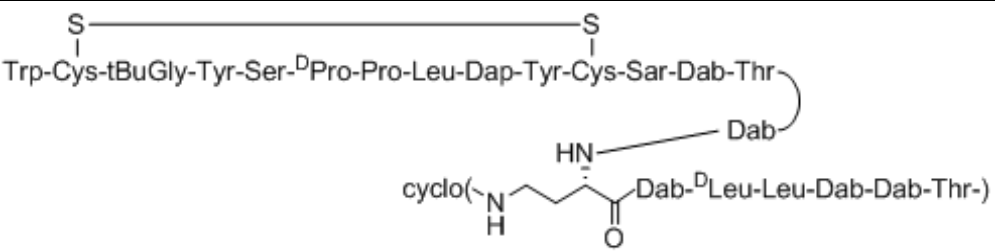
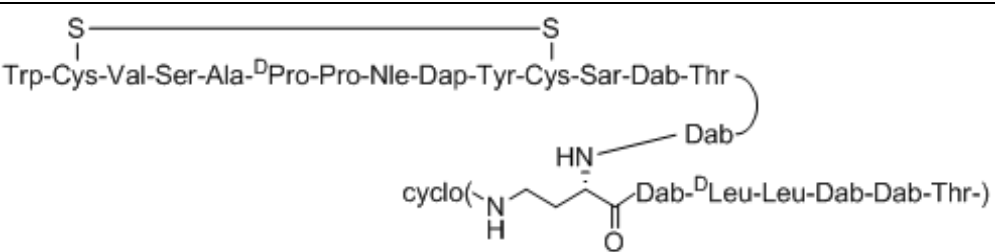
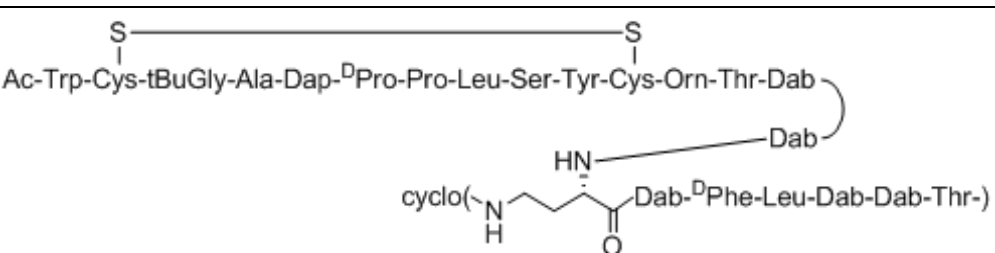
<p>Ex. 111 a)</p>	<p>Ac-Trp-Cys-Trp-Ala-Ser-^DPro-Pro-Leu-Ser-Tyr-Cys-Orn-Thr-Dab</p> <p>cyclo(NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>
<p>Ex. 112 a)</p>	<p>Ac-Trp-Cys-tBuGly-^DAla-Ser-^DPro-Pro-Leu-Ser-Tyr-Cys-Orn-Thr-Dab</p> <p>cyclo(NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>
<p>Ex. 113 a)</p>	<p>Ac-Trp-Cys-tBuGly-Ala-<i>allo</i>Thr-^DPro-Pro-Leu-His-Cha-Cys-Orn-Thr-Dab</p> <p>cyclo(NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>
<p>Ex. 114 a)</p>	<p>Trp-Cys-tBuGly-Ala-Ser-^DPic-Pro-Leu-Ser-Tyr-Cys-Dab-Thr-Dab</p> <p>cyclo(NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>
<p>Ex. 115 a)</p>	<p>Ac-Trp-Cys-Val-Thr-Val-^DPro-Arg-Leu-Thr-Tyr-Cys-Dab-Dab-Dab</p> <p>cyclo(NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>
<p>Ex. 116 a)</p>	<p>Ac-Trp-Cys-tBuGly-Dap-Thr-^DPro-Pro-Nle-Gly-Tyr-Cys-Dab-Thr-Dab</p> <p>cyclo(NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>

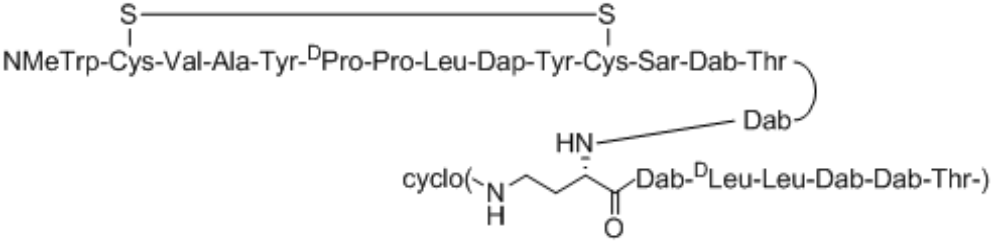
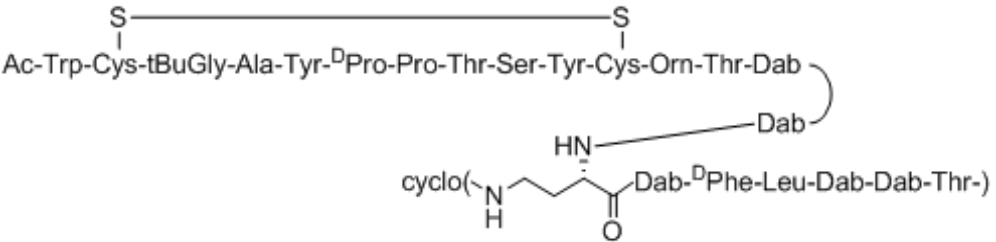
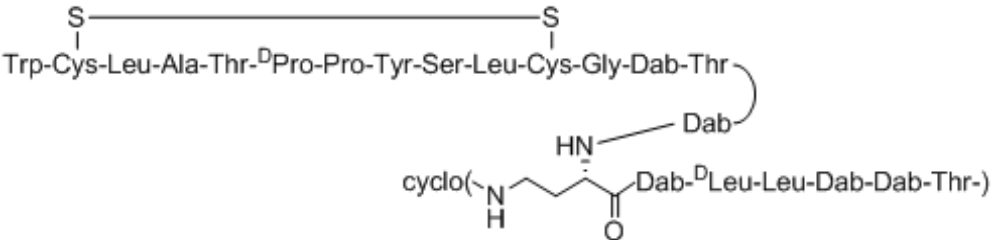
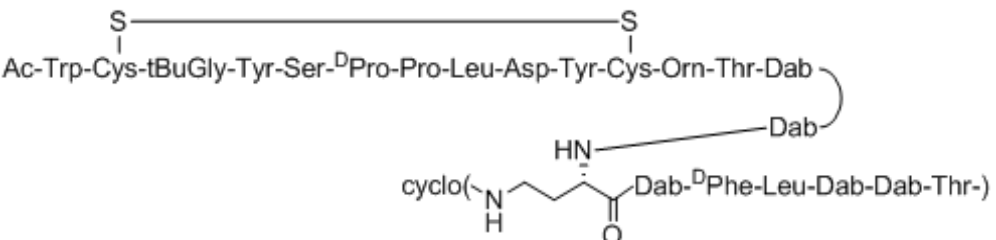
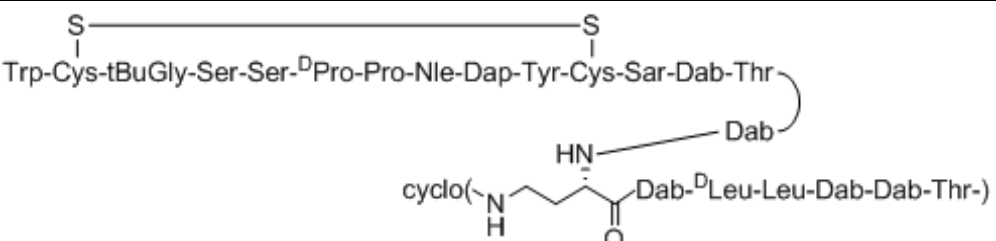
<p>Ex. 117 a)</p>	
<p>Ex. 118 a)</p>	
<p>Ex. 119 a)</p>	
<p>Ex. 120 a)</p>	
<p>Ex. 121 a)</p>	
<p>Ex. 122 a)</p>	

<p>Ex. 123 a)</p>	<p>Trp-Cys-Val-Ala-Ser-^DPro-Pro((4<i>R</i>)OH)-Leu-Ser-Tyr-Cys-Dab-Dab-Thr</p> <p>cyclo(-NH-CH₂-CH₂-CH(NH-Dab)-C(=O)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p>
<p>Ex. 124 a)</p>	<p>Trp-Cys-Val-Ala-Ser-^DPro-Pro((4<i>R</i>)F)-Leu-Ser-Tyr-Cys-Dab-Dab-Thr</p> <p>cyclo(-NH-CH₂-CH₂-CH(NH-Dab)-C(=O)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p>
<p>Ex. 125 a)</p>	<p>Trp-Cys-Val-Ala-Ser-^DPro-Ala-Leu-Ser-Tyr-Cys-Dab-Dab-Thr</p> <p>cyclo(-NH-CH₂-CH₂-CH(NH-Dab)-C(=O)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p>
<p>Ex. 126 a)</p>	<p>Trp-Cys-Val-Ala-Ser-^DPro-Thr-Leu-Ser-Tyr-Cys-Gly-Dab-Thr</p> <p>cyclo(-NH-CH₂-CH₂-CH(NH-Dab)-C(=O)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p>
<p>Ex. 127 a)</p>	<p>Trp-Cys-Val-Ala-Ser-^DPro-Dab-Leu-Ser-Tyr-Cys-Dab-Dab-Thr</p> <p>cyclo(-NH-CH₂-CH₂-CH(NH-Dab)-C(=O)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p>

<p>Ex. 128 a)</p>	
<p>Ex. 129 a)</p>	
<p>Ex. 130 a)</p>	
<p>Ex. 131 a)</p>	
<p>Ex. 132 a)</p>	

<p>Ex. 133 a) e)</p>	
<p>Ex. 134 a)</p>	
<p>Ex. 135 a) d)</p>	
<p>Ex. 136 a)</p>	
<p>Ex. 137 a)</p>	

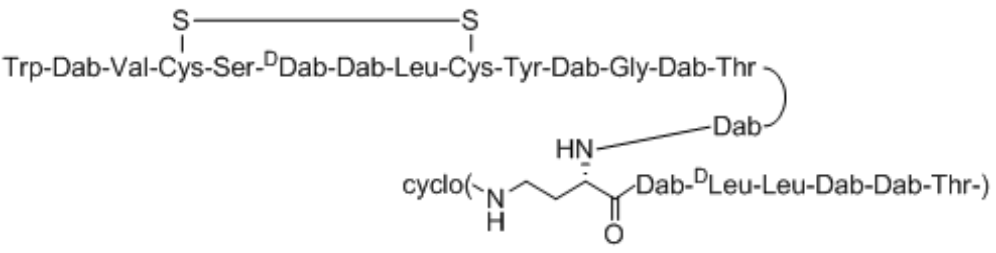
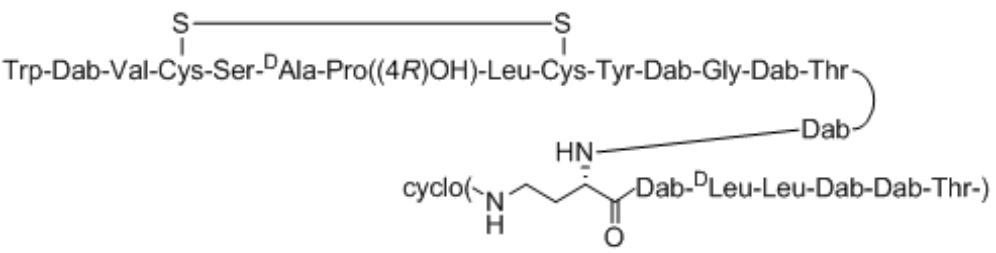
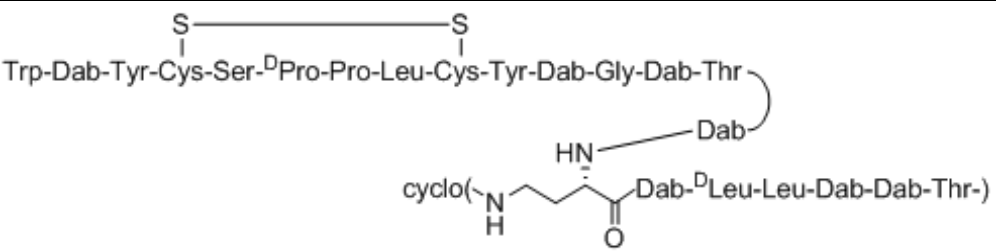
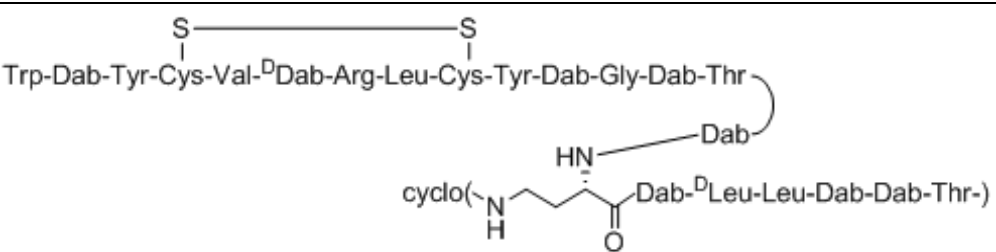
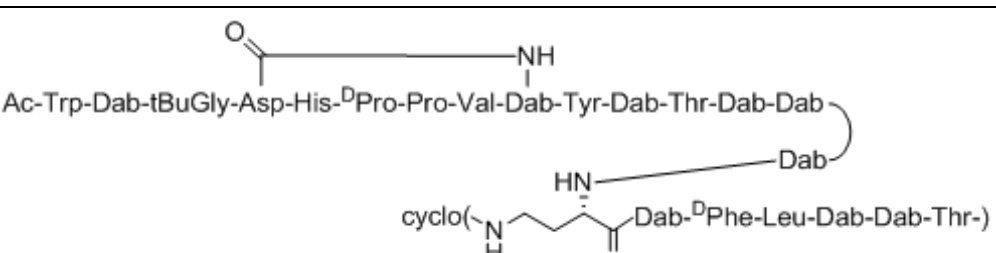
<p>Ex. 138 a)</p>	 <p>Ac-Trp-Cys-tBuGly-Ala-Ser-^DPro-Pro-Trp-His-Tyr-Cys-Orn-Thr-Dab</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>
<p>Ex. 139 a)</p>	 <p>Ac-Trp-Cys-tBuGly-Ala-Ser-^DPro-Pro-Trp-Dap(iPr)-Cha-Cys-Orn-Thr-Dab</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>
<p>Ex. 140 a)</p>	 <p>Trp-Cys-tBuGly-Tyr-Ser-^DPro-Pro-Leu-Dap-Tyr-Cys-Sar-Dab-Thr</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p>
<p>Ex. 141 a)</p>	 <p>Trp-Cys-Val-Ser-Ala-^DPro-Pro-Nle-Dap-Tyr-Cys-Sar-Dab-Thr</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p>
<p>Ex. 142 a)</p>	 <p>Ac-Trp-Cys-tBuGly-Ala-Dap-^DPro-Pro-Leu-Ser-Tyr-Cys-Orn-Thr-Dab</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>

<p>Ex. 143 a)</p>	
<p>Ex. 144 a)</p>	
<p>Ex. 145 a)</p>	
<p>Ex. 146 a)</p>	
<p>Ex. 147 a)</p>	

<p>Ex. 148 a)</p>	<p>Trp-Cys-Val-Ser-Ser-^DPro-Pro-Val-Dap-Tyr-Cys-Orn-^DDab-Dab</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p>
<p>Ex. 149 a)</p>	<p>Trp-Cys-Val-Ser-Ser-^DPro-Tyr-Val-Dap-Tyr-Cys-Orn-Thr-^DDab</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p>
<p>Ex. 150 a)</p>	<p>Trp-Cys-Val-Ser-Ser-^DPro-Pro-Nle-Dap-Ala-Cys-Sar-Dab-Thr</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p>
<p>Ex. 151 a)</p>	<p>Trp-Cys-Val-Tyr-Ser-^DPro-Pro-Leu-Asp-Tyr-Cys-Orn-Thr-Dab</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p>
<p>Ex. 152 a)</p>	<p>Trp-Cys-Val-Arg-Arg-^DPro-Pro-Leu-Ser-Tyr-Cys-Sar-Dab-Thr</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p>

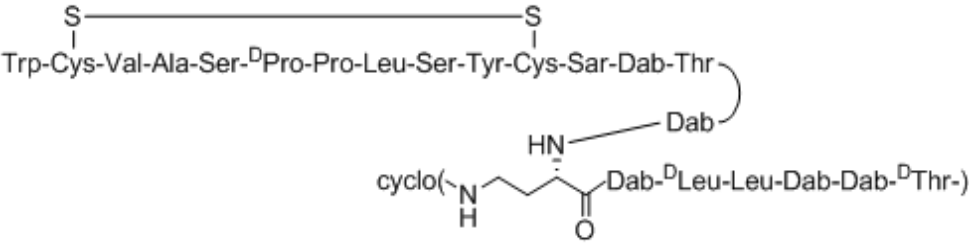
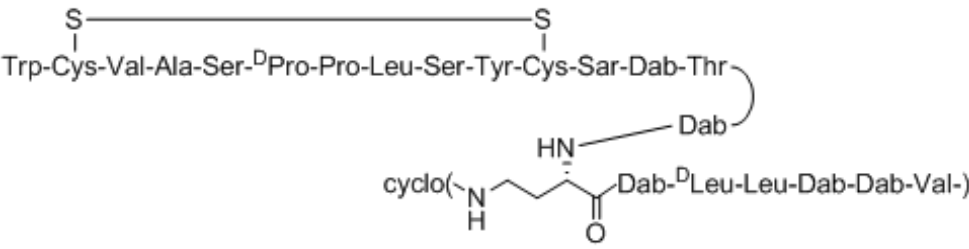
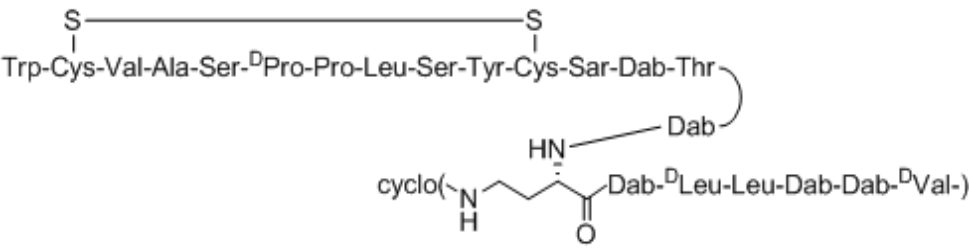
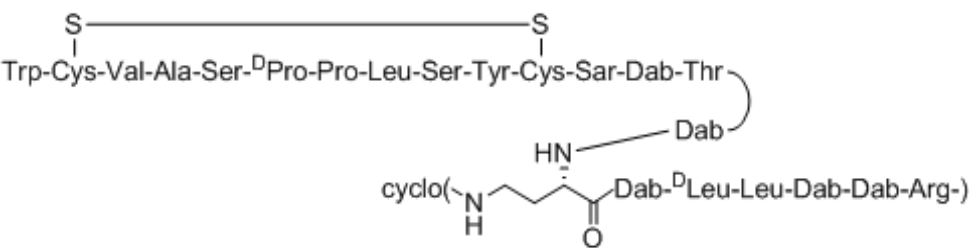
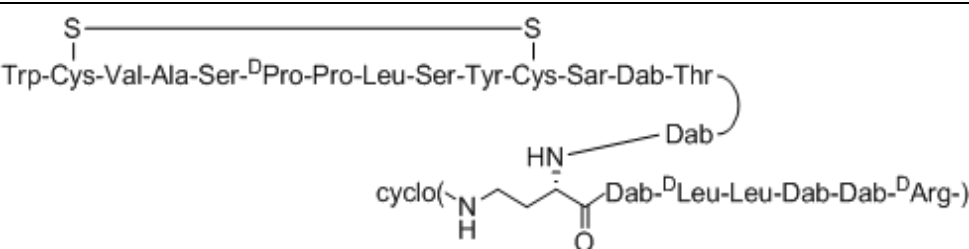
<p>Ex. 153 a)</p>	
<p>Ex. 154 a)</p>	
<p>Ex. 155 a)</p>	
<p>Ex. 156 a)</p>	
<p>Ex. 157 a)</p>	

<p>Ex. 158 a)</p>	
<p>Ex. 159 a)</p>	
<p>Ex. 160 a)</p>	
<p>Ex. 161 a)</p>	
<p>Ex. 162 a)</p>	

<p>Ex. 163 a)</p>	 <p>Trp-Dab-Val-Cys-Ser-^DDab-Dab-Leu-Cys-Tyr-Dab-Gly-Dab-Thr</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p>
<p>Ex. 164 a)</p>	 <p>Trp-Dab-Val-Cys-Ser-^DAla-Pro((4<i>R</i>)-OH)-Leu-Cys-Tyr-Dab-Gly-Dab-Thr</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p>
<p>Ex. 165 a)</p>	 <p>Trp-Dab-Tyr-Cys-Ser-^DPro-Pro-Leu-Cys-Tyr-Dab-Gly-Dab-Thr</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p>
<p>Ex. 166 a)</p>	 <p>Trp-Dab-Tyr-Cys-Val-^DDab-Arg-Leu-Cys-Tyr-Dab-Gly-Dab-Thr</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DLeu-Leu-Dab-Dab-Thr-)</p>
<p>Ex. 167 b)</p>	 <p>Ac-Trp-Dab-tBuGly-Asp-His-^DPro-Pro-Val-Dab-Tyr-Dab-Thr-Dab-Dab</p> <p>cyclo(-NH-CH₂-CH₂-CH₂-C(=O)-Dab-^DPhe-Leu-Dab-Dab-Thr-)</p>

<p>Ex. 168 b)</p>	
<p>Ex. 169 a)</p>	
<p>Ex. 170 a)</p>	
<p>Ex. 171 a)</p>	
<p>Ex. 172 b) d)</p>	

<p>Ex. 173 a) d)</p>	
<p>Ex. 174 a) d)</p>	
<p>Ex. 175 a)</p>	
<p>Ex. 176 b)</p>	
<p>Ex. 177 a) c)</p>	

<p>Ex. 178 a)</p>	 <p>The structure shows a protein chain: Trp-Cys-Val-Ala-Ser-^DPro-Pro-Leu-Ser-Tyr-Cys-Sar-Dab-Thr. A disulfide bridge (S-S) connects the two Cys residues. The Cys at position 11 is linked to a cyclic peptide side chain: cyclo(-NH-CH₂-CH₂-CH(NH-Dab)-C(=O)-Dab-^DLeu-Leu-Dab-Dab-^DThr-). The side chain is enclosed in a circle with a dash at the N-terminus.</p>
<p>Ex. 179 a)</p>	 <p>The structure shows a protein chain: Trp-Cys-Val-Ala-Ser-^DPro-Pro-Leu-Ser-Tyr-Cys-Sar-Dab-Thr. A disulfide bridge (S-S) connects the two Cys residues. The Cys at position 11 is linked to a cyclic peptide side chain: cyclo(-NH-CH₂-CH₂-CH(NH-Dab)-C(=O)-Dab-^DLeu-Leu-Dab-Dab-Val-). The side chain is enclosed in a circle with a dash at the N-terminus.</p>
<p>Ex. 180 a)</p>	 <p>The structure shows a protein chain: Trp-Cys-Val-Ala-Ser-^DPro-Pro-Leu-Ser-Tyr-Cys-Sar-Dab-Thr. A disulfide bridge (S-S) connects the two Cys residues. The Cys at position 11 is linked to a cyclic peptide side chain: cyclo(-NH-CH₂-CH₂-CH(NH-Dab)-C(=O)-Dab-^DLeu-Leu-Dab-Dab-^DVal-). The side chain is enclosed in a circle with a dash at the N-terminus.</p>
<p>Ex. 181 a)</p>	 <p>The structure shows a protein chain: Trp-Cys-Val-Ala-Ser-^DPro-Pro-Leu-Ser-Tyr-Cys-Sar-Dab-Thr. A disulfide bridge (S-S) connects the two Cys residues. The Cys at position 11 is linked to a cyclic peptide side chain: cyclo(-NH-CH₂-CH₂-CH(NH-Dab)-C(=O)-Dab-^DLeu-Leu-Dab-Dab-Arg-). The side chain is enclosed in a circle with a dash at the N-terminus.</p>
<p>Ex. 182 a)</p>	 <p>The structure shows a protein chain: Trp-Cys-Val-Ala-Ser-^DPro-Pro-Leu-Ser-Tyr-Cys-Sar-Dab-Thr. A disulfide bridge (S-S) connects the two Cys residues. The Cys at position 11 is linked to a cyclic peptide side chain: cyclo(-NH-CH₂-CH₂-CH(NH-Dab)-C(=O)-Dab-^DLeu-Leu-Dab-Dab-^DArg-). The side chain is enclosed in a circle with a dash at the N-terminus.</p>

<p>Ex. 183 a)</p>	
<p>Ex. 184 a)</p>	
<p>Ex. 185 a)</p>	
<p>Ex. 186 a)</p>	
<p>Ex. 187 a) c)</p>	
<p>Ex. 188 a)</p>	

<p>Ex. 189 a)</p>	
<p>Ex. 190 b)</p>	
<p>Ex. 191 a) b)</p>	
<p>Ex. 192 a)</p>	
<p>Ex. 193 b)</p>	

<p>Ex. 194 a)</p>	
<p>Ex. 195 a)</p>	
<p>Ex. 196 b)</p>	
<p>Ex. 197 b)</p>	
<p>Ex. 198 b)</p>	

<p>Ex. 199 a)</p>	
<p>Ex. 200 a)</p>	
<p>Ex. 201 a)</p>	
<p>Ex. 202 a)</p>	
<p>Ex. 203 a)</p>	

<p>Ex. 204 a)</p>	
<p>Ex. 205 a)</p>	
<p>Ex. 206 a)</p>	
<p>Ex. 207 a)</p>	
<p>Ex. 208 a)</p>	

<p>Ex. 209 a)</p>	
<p>Ex. 210 a) b)</p>	
<p>Ex. 211 a)</p>	
<p>Ex. 212 a)</p>	
<p>Ex. 213 a)</p>	

<p>Ex. 214 a)</p>	
<p>Ex. 215 a)</p>	
<p>Ex. 216 a)</p>	

wherein a) designates compounds comprising one or more disulfide interstrand linkage(s) between indicated amino acid residues, wherein the interstrand linkage(s) comprise at least one disulfide bond(s) between a pair(s) of side-chain thiol groups comprised on the indicated amino acid residues;

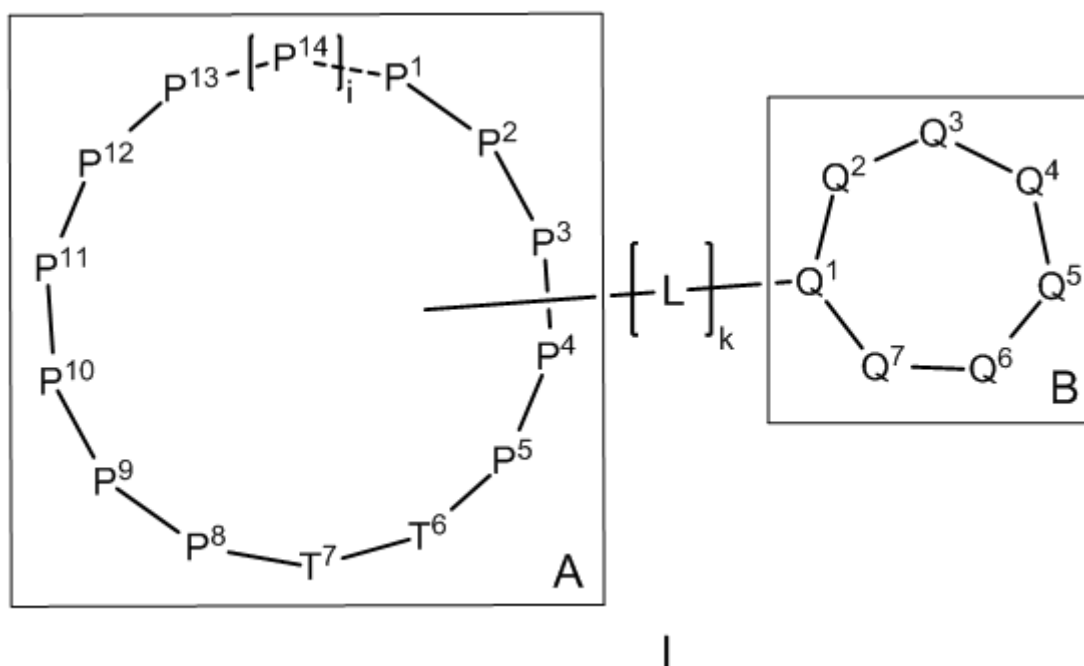
wherein b) designates compounds comprising a lactam interstrand linkage between two indicated amino acid residues, wherein the lactam interstrand linkage comprises an amide bond between a side-chain amino group and a side-chain carboxyl group;

wherein c) designates compounds comprising a dipeptidic amino acid residue at indicated position, wherein the dipeptidic amino acid residue comprises an amide bond between a α -carboxyl group of a terminal amino acid residue and a side-chain amino group of a second amino acid residue as specified;

wherein d) designates compounds comprising a guanidine group comprising an amino group of a N-terminal amino acid residue; and

wherein e) designates compounds comprising a tetramethylguanidine group comprising an amino group of the N-terminal amino acid residue;

wherein formula (I) is defined as



and comprises a **module A** comprising single elements P or T being connected in either direction from the carbonyl (C=O) point of attachment to the nitrogen (N) of the next element, wherein i is 0 or 1, with the proviso that, if i = 1, P¹³ and P¹⁴; or P¹⁴ and P¹ may not be connected as aforementioned; if i = 0, P¹³ and P¹ are not connected as aforementioned; and wherein,

if $i = 1$, and

P² and P¹¹ taken together and/or P⁴ and P⁹ taken together and/or P¹³ and P¹⁴ taken together may form naturally or non-naturally cross-linking α -amino acids containing each in total 1 to 12 carbon- and/or heteroatoms in a single side-chain which together are connecting P² and P¹¹ and/or P⁴ and P⁹ and/or P¹³ and P¹⁴ by covalent interaction (interstrand linkage); then

P¹ is a naturally or non-naturally occurring aromatic L α-amino acid containing 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring aliphatic L α-amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain;

P² is a naturally or non-naturally occurring basic L α-amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one amino function; or a naturally or non-naturally occurring L α-amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one amide function; or a naturally or non-naturally occurring alcoholic L α-amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring aliphatic L α-amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain;

P³ is a naturally or non-naturally occurring aliphatic L α-amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring aromatic L α-amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain;

P⁴ is Gly; Sar; or a naturally or non-naturally occurring aliphatic α-amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally

- occurring basic L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one amino function; or a naturally or non-naturally occurring alcoholic L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one amide function; or a naturally or non-naturally occurring aromatic L α -amino acid containing 1 to 25 carbon- and/or heteroatoms in a single side-chain;
- P⁵ is a naturally or non-naturally occurring aliphatic L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring alcoholic L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring basic L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one amino function; or a naturally or non-naturally occurring aromatic L α -amino acid containing 1 to 25 carbon- and/or heteroatoms in a single side-chain;
- T⁶ is a naturally or non-naturally occurring D α -amino acid containing an optionally substituted side-chain which forms a four-, five- or six-membered hetero-cycle or a bicyclic system comprising the α -carbon and the α -amino atom; or a naturally or non-naturally occurring aliphatic D α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring basic D α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one amino function; or a naturally or non-naturally occurring aromatic D α -amino acid containing 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring alcoholic D α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain;
- T⁷ is a naturally or non-naturally occurring L α -amino acid containing an optionally substituted side-chain which forms a five- or six-membered heterocycle or a bicyclic system comprising the α -carbon and the α -amino atom; or a naturally or non-naturally occurring aliphatic L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring aromatic L α -amino acid containing 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring basic L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one amino function; or a naturally or non-naturally occurring alcoholic L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain;
- P⁸ is a naturally or non-naturally occurring aliphatic L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring aromatic L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring alcoholic L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain;
- P⁹ is Gly; Sar; or a naturally or non-naturally occurring aliphatic L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally

- occurring basic L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one amino function; or a naturally or non-naturally occurring alcoholic L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one amide function; or a naturally or non-naturally occurring aromatic L α -amino acid containing 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one urea function, carboxylic acid function, amide function, ester function, sulfone function or ether function;
- P¹⁰ is a naturally or non-naturally occurring aliphatic L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring aromatic L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain;
- P¹¹ is Gly; Sar; or a naturally or non-naturally occurring basic L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one amino function; or a naturally or non-naturally occurring L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one amide function; or a naturally or non-naturally occurring aliphatic L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring alcoholic L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain;
- P¹² is Gly; Sar; Aib; or a naturally or non-naturally occurring aliphatic α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring basic α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one amino function; or a naturally or non-naturally occurring alcoholic α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring aromatic α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one urea function, carboxylic acid function, amide function, ester function, sulfone function or ether function;
- P¹³ is Gly; Sar; Aib; or a naturally or non-naturally occurring aliphatic α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring basic α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one amino function; or a naturally or non-naturally occurring alcoholic α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring aromatic α -amino acid containing 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally

occurring α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one amide function;

P¹⁴ is Gly; Sar; Aib; or a naturally or non-naturally occurring aliphatic α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring basic α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one amino function; or a naturally or non-naturally occurring α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one urea function, carboxylic acid function, amide function, ester function, sulfone function or ether function; or a naturally or non-naturally occurring alcoholic α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain;

with the proviso that,

- if no interstrand linkage is formed, then P¹³ and P¹⁴, and P¹⁴ and P¹ are connected;
- if P¹³ and P¹⁴ taken together form an interstrand linkage, as defined above, then P¹³ and P¹⁴ are not additionally connected;

with the further proviso that,

- if linker L, as defined below, is connected with module A by a carbonyl (C=O) point of attachment of P²; P⁵; or P¹²; then
P²; P⁵; or P¹²; is a naturally or non-naturally occurring α -amino acid
containing in total 1 to 25 carbon- and/or heteroatoms in a
single side-chain comprising at least one carboxyl function;
- if linker L, as defined below, is connected with module A by a carbonyl (C=O) point of attachment of P¹³ and
P¹³ and P¹⁴, and P¹⁴ and P¹ are connected; or
P¹⁴ and P¹ are not connected; then
P¹³ is a naturally or non-naturally occurring α -amino acid containing in total 1 to 25
carbon- and/or heteroatoms in a single side-chain comprising at least one
carboxyl function;
- if linker L, as defined below, is connected with module A by a carbonyl (C=O) point of attachment of P¹⁴ and
P¹³ and P¹⁴, and P¹⁴ and P¹ are connected as aforementioned; or
P¹³ and P¹⁴ are not connected; then
P¹⁴ is a naturally or non-naturally occurring α -amino acid containing in total 1 to 25
carbon- and/or heteroatoms in a single side-chain comprising at least one
carboxyl function;

if i = 0, and

P² and P¹¹ taken together and/or P⁴ and P⁹ taken together form naturally or non-naturally cross-linking α -amino acids containing each in total 1 to 12 carbon- and/or heteroatoms in a single side-chain which together are connecting P² and P¹¹ and/or

P⁴ and P⁹ by covalent interaction (interstrand linkage); then

P¹ to P⁵; T⁶; T⁷; P⁸ to P¹³ are naturally or non-naturally occurring α -amino acids;

with the proviso that,

- if linker L, as defined below, is connected with module A by a carbonyl (C=O) point of attachment of P⁵; or P¹²; then

P⁵; or P¹²; is a naturally or non-naturally occurring α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one carboxyl function;

and a **module B** consisting of single elements Q being connected in either direction from the carbonyl (C=O) point of attachment to the nitrogen (N) of the next element with the proviso that Q⁷ is connected from the α -carbonyl (C=O) point of attachment to the ω -nitrogen (N) of Q¹, and wherein

Q¹ is a naturally or non-naturally occurring basic α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one amino function;

Q², Q⁵, and Q⁶ are independently

a naturally or non-naturally occurring basic L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one amino function;

Q³ is a naturally or non-naturally occurring aliphatic D α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or or a naturally or non-naturally occurring aromatic D α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain;

Q⁴ is a naturally or non-naturally occurring aliphatic L α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain;

Q⁷ is a naturally or non-naturally occurring alcoholic α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one amide function; or a naturally or non-naturally occurring aliphatic α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring basic α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one amino function;

and a **linker L** consisting of k = 0 - 3 single elements L being connected in either direction from the carbonyl (C=O) point of attachment to the nitrogen (N) of the next element, and wherein,

if k = 1,

L¹ is Gly; Sar; Aib; or a naturally or non-naturally occurring aliphatic α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring basic α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one amino function; or a naturally or non-naturally occurring alcoholic α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring aromatic α -amino acid containing 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally

occurring α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one amide function;

if $k = 2$, the additional element

L^2 is Gly; Sar; Aib; or a naturally or non-naturally occurring aliphatic α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring basic α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one amino function; or a naturally or non-naturally occurring alcoholic α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain;

if $k = 3$, the additional element

L^3 is Gly; Sar; Aib; or a naturally or non-naturally occurring aliphatic α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain; or a naturally or non-naturally occurring basic α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain comprising at least one amino function; or a naturally or non-naturally occurring alcoholic α -amino acid containing in total 1 to 25 carbon- and/or heteroatoms in a single side-chain;

said linker L being connected with module B from the carbonyl (C=O) point of attachment of L^k to the α -nitrogen (N) of Q^1 and,

if $k = 1 - 3$ and $i = 1$, being connected with module A from the carbonyl (C=O) point of attachment of P^2 ; P^5 ; P^{12} ; P^{13} ; or P^{14} ; to the nitrogen (N) of L^1 ; or,

if $k = 1 - 3$ and $i = 0$, being connected with module A from the carbonyl (C=O) point of attachment of P^5 ; P^{12} ; or P^{13} ; to the nitrogen (N) of L^1 ; or

if $k = 0$ and $i = 1$, then

Q^1 being directly connected with module A from the carbonyl (C=O) point of attachment of P^2 ; P^5 ; P^{12} ; P^{13} ; or P^{14} ; to the α -nitrogen (N) of Q^1 ; or

if $k = 0$ and $i = 0$, then

Q^1 being directly connected with module A from the carbonyl (C=O) point of attachment of P^5 ; P^{12} ; or P^{13} ; to the α -nitrogen (N) of Q^1 ;

the carbonyl (C=O) point of attachment of P^{13} ; or P^{14} ; and/or nitrogen (N) of P^1 ; or P^{14} ; not connected being saturated to form the corresponding naturally or non-naturally occurring terminal α -amino acids optionally having modified carbonyl (C=O) functional groups and/or nitrogen (N) functional groups; or a tautomer or rotamer thereof; or a salt; or a pharmaceutically acceptable salt; or a hydrate; or a solvate thereof.

2. A diastereomer or epimer of a compound of formula **(I)** according to claim 1 based on one or more chiral center(s) not explicitly specified in formula **(I)** or an enantiomer of a compound of formula **(I)**.

3. A pharmaceutical composition comprising a compound or a mixture of compounds according to claim 1 and at least one pharmaceutically inert carrier.
4. A pharmaceutical composition according to claim 3 in a form suitable for oral, topical, transdermal, injection, buccal, transmucosal, rectal, pulmonary or inhalation administration, especially in the form of tablets, dragees, capsules, solutions, liquids, gels, plaster, creams, ointments, syrup, slurries, suspensions, spray, nebulizer or suppositories.
5. Use of a pharmaceutically acceptable amount of a compound or compounds according to claim 1 in the manufacture of a medicament for the treatment of an infection caused by a Gram-negative bacteria selected from the group consisting of nosocomial infections, catheter-related infections, non-catheter-related infections, urinary tract infections, and bloodstream infections, or a disease or disorder associated with an infection caused by a Gram-negative bacteria, selected from the group consisting of ventilator-associated pneumonia (VAP), hospital-acquired pneumonia (HAP), healthcare-associated pneumonia (HCAP), cystic fibrosis, emphysema, asthma, pneumonia, epidemic diarrhea, necrotizing enterocolitis, typhilitis, gastroenteritis, pancreatitis, keratitis, endophthalmitis, otitis, brain abscess, meningitis, encephalitis, osteochondritis, pericarditis, epididymitis, prostatitis, urethritis, sepsis; surgical wounds, traumatic wounds, and burns.
6. A process for the preparation of compounds according to claim 1, comprising
 - (a) coupling a functionalized solid support with an N-protected derivative of that amino acid which in a desired end-product is at position Q⁷ of module B, any functional group which may be present in said N-protected amino acid derivative being protected;
 - (b) removing the N-protecting group from the product obtained in step (a);
 - (c) coupling the product of step (b) with an N-protected derivative of that amino acid which in the desired end-product corresponds to Q⁶, any functional group which may be present in said N-protected amino acid derivative being protected;
 - (d) further effecting steps substantially corresponding to steps (b) to (c) using N-protected derivatives of amino acids which in the desired end-product are at positions Q⁵ to Q¹, any functional group(s) which may be present in said N-protected amino acid derivatives being protected;
 - (e) if coupling to the solid support in step (a) is via a hydroxyl group of the amino acid residue at position Q⁷, performing the following chemical conversion as described below:
selectively removing an N-protecting group at position Q¹ and a carboxyl-protecting group at position Q⁷; and generating a macrolactam cycle-by formation of an amide bond between the thus liberated carboxyl group at position Q⁷ and the amino group at Q¹ of module B;
 - (f) if L is present (k = 1, 2, or 3), effecting steps substantially corresponding to steps (b) to (c) using N-protected derivatives of amino acids which in the desired end-product are at positions

L^k to L^1 , any functional group(s) which may be present in said N-protected amino acid derivatives being protected;

- (g) if P^{14} is present in module A ($i = 1$) effecting steps substantially corresponding to steps (b) to (c) using an N-protected derivative of an amino acid which in the desired end-product is at position P^n ($n = 2, 5, 12, 13$, or 14), any functional group(s) which may be present in said N-protected amino acid derivative being protected; and

if $n = 2$, further performing steps comprising:

- (h1) effecting steps substantially corresponding to steps (b) to (c) using N-protected derivatives of amino acids which in the desired end-product are at positions P^1 , P^{14} to P^8 , T^7 , T^6 , and P^5 to P^3 , any functional group(s) which may be present in said N-protected amino acid derivatives being protected and-optionally following each coupling, selectively deprotecting one or several protected functional group(s) present in the molecule and chemically transforming the reactive group(s) thus liberated to form an interstrand linkage(s); and
- (i1) selectively removing a carboxyl-protecting group at position P^2 and the N-protecting group at position P^3 ; and generating the macrolactam cycle-by formation of an amide bond between the thus liberated carboxyl and amino functions;

if $n = 5$, further performing steps which comprise:

- (h2) further effecting steps substantially corresponding to steps (b) to (c) using N-protected derivatives of amino acids which in the desired end-product are at positions P^4 to P^1 , P^{14} to P^8 , T^7 , and T^6 , any functional group(s) which may be present in said N-protected amino acid derivatives being protected and-optionally following each coupling, selectively deprotecting one or several protected functional group(s) present in the molecule and chemically transforming the reactive group(s) thus liberated to form an interstrand linkage(s); and
- (i2) selectively removing a carboxyl-protecting group at position P^5 and the N-protecting group at position T^6 ; and generating the macrolactam cycle by formation of an amide bond between the thus liberated carboxyl and amino functions;

if $n = 12$, further performing steps comprising:

- (h3) further effecting steps substantially corresponding to steps (b) to (c) using N-protected derivatives of amino acids which in the desired end-product are at positions P^{11} to P^8 , T^7 , T^6 , P^5 to P^1 , P^{14} and P^{13} , any functional group(s) which may be present in said N-protected amino acid derivatives being protected and-optionally following each coupling, selectively deprotecting one or several protected functional group(s) present in the molecule and chemically transforming the reactive group(s) thus liberated to form an interstrand linkage(s); and
- (i3) selectively removing a carboxyl-protecting group at position P^{12} and the N-protecting group at position P^{13} ; and generating the macrolactam cycle by formation of an amide bond between the thus liberated carboxyl and amino functions;

if $n = 13$, further performing steps comprising:

- (h4) further effecting steps substantially corresponding to steps (b) to (c) using N-protected derivatives of amino acids which in the desired end-product are at positions P¹² to P⁸, T⁷, T⁶, P⁵ to P¹, and P¹⁴, any functional group(s) which may be present in said N-protected amino acid derivatives being protected
 - (i4) selectively removing a carboxyl-protecting group at position P¹³ and an N-protecting group at position P¹⁴; and generating the macrolactam cycle-by formation of an amide bond between the thus liberated carboxyl and amino functions;
- if n = 14, performing steps comprising:
- (h5) further effecting steps substantially corresponding to steps (b) to (d) using N-protected derivatives of amino acids which in the desired end-product are at positions P¹³ to P⁸, T⁷, T⁶, P⁵ to P¹, any functional group(s) which may be present in said N-protected amino acid derivatives being selectively removing a carboxyl-protecting group at position P¹⁴ and an N-protecting group at position P¹; and generating the macrolactam cycle, as defined above, by formation of an amide bond between the thus liberated carboxyl and amino functions;
 - (j) if coupling to the solid support in step (a) is via a carboxyl group of the amino acid residue at position Q⁷ of module B, selectively removing an N-protecting group at position Q¹ of module B;
 - (k) detaching the product thus obtained from the solid support;
 - (l) if coupling to the solid support in step (a) is via a carboxyl group of the amino acid residue at position Q⁷ of module B, generating the macrolactam cycle by formation of an amide bond between the thus liberated carboxyl group at position Q⁷ and the amino group at position Q¹ of module B;
 - (m) removing any protecting groups present on functional groups of any members of the chain of amino acid residues and, optionally any protecting group(s) which may in addition be present in the molecule;
 - (n) optionally converting the product thus obtained into a pharmaceutically acceptable salt or converting a pharmaceutically acceptable, or unacceptable, salt thus obtained into the corresponding free compound of formula (I) or into a different, pharmaceutically acceptable salt.
7. The process according to claim 6, further comprising:
- (l1) selectively deprotecting one or more protected functional group(s) present in the molecule and chemically transforming the reactive group(s) thus liberated, wherein (l1) is between steps (l) and (m).
8. The process according to claim 6, further comprising:
- (n) implementing additional chemical transformations of one or more reactive group(s) present in the molecule, wherein (n) is after step (m).

9. The process according to claim 6, further comprising:
- (o) converting the product thus obtained into a pharmaceutically acceptable salt or converting a pharmaceutically acceptable, or unacceptable, salt thus obtained into the corresponding free compound of formula (I) or into a different, pharmaceutically acceptable salt, wherein (o) is after step (m).
10. The process according to claim 8, further comprising:
- (o) converting the product thus obtained into a pharmaceutically acceptable salt or converting a pharmaceutically acceptable, or unacceptable, salt thus obtained into the corresponding free compound of formula (I) or into a different, pharmaceutically acceptable salt, wherein (o) is after step (n).