

## Supplementary Information

### Urotensin-II Peptidomimetic Incorporating a Non-Reducible 1,5-Triazole Disulfide Bond Reveals a Pseudo-Irreversible Covalent Binding Mechanism to the Urotensin G-Protein Coupled Receptor

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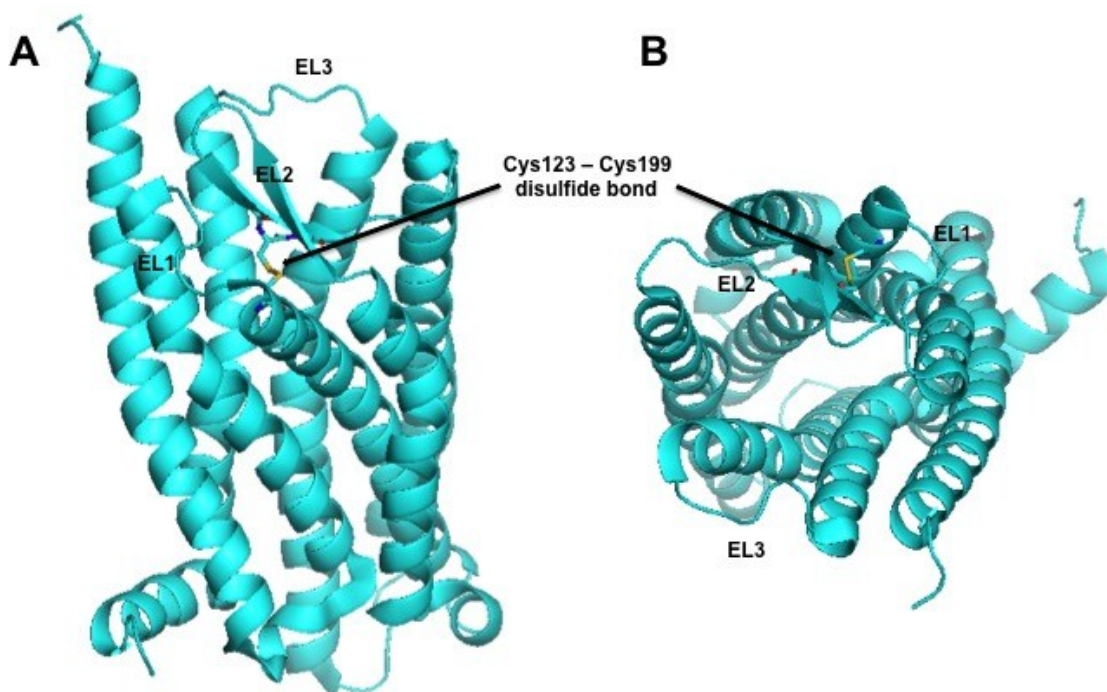
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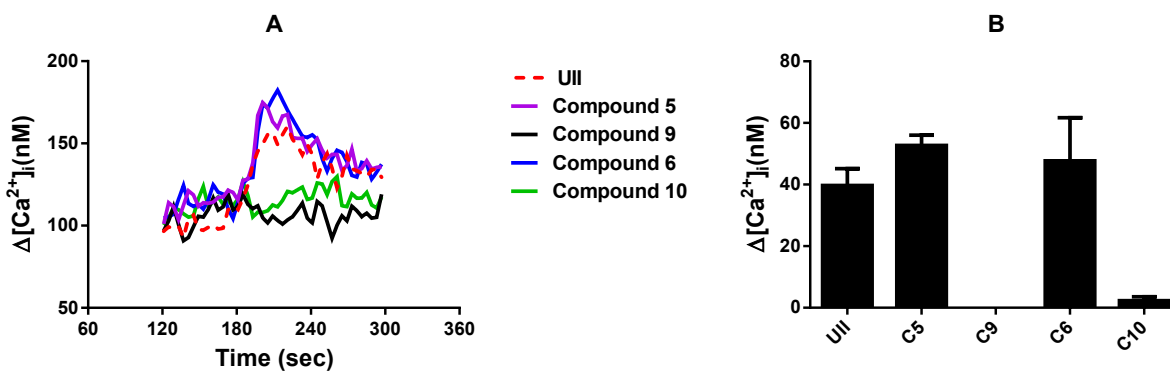
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## 1. Supplementary Figures



**Figure S1.** Crystal structure of h-DOR (PDB ID: 4n6h) indicating the disulfide bond between EL1 and EL2. (A) h-DOR side-view; (B) top, extracellular view



**Figure S2.** Intracellular  $\text{Ca}^{2+}$  data. U-II, compounds 5 and 6 increased intracellular  $\text{Ca}^{2+}$  in SJCRH30 cells. Data are Mean  $\pm$  SEM (n = 3).

## 2. 3D Modeling and Calculations

UrotensinII\_NMR.pdb was imported into Schrödinger Maestro (*Schrödinger*, LLC). This is used as a reference structure for of U-II <sup>4-11</sup>. The disulfide connection between residues 5 and 10 was removed, retaining the respective C $\beta$  atoms. Then, the corresponding triazole linkages were modelled into the molecule. A conformational search was performed (MMFFs + mixed torsional/low-mode sampling, in water) for the triazole linkages and C $\beta$  atoms in **5-6** and **9-10**, whilst keeping the rest of Urotensin frozen. The lowest energy conformer for each respective case underwent an energy minimization (MMFFs + PRCG, in water) for the full structure. The obtained structures were aligned at the respective C $\alpha$  and C $\beta$  atoms of residue 5 and 10 using the “Atom pair superposition” task. Distances between C $\alpha$  atoms of residues 5 and 10 were calculated using the “Measurement” tool. The root mean square deviations (RMSD) were calculated for the respective C $\alpha$  and C $\beta$  atoms of residues 5 and 10 at the compared structures using the “RMSD” superposition tool.

Entry	d(C $\alpha_5$ -C $\alpha_{10}$ ) [Å]	RMSD
U-II <sup>4-11</sup>	3.95	
<b>9</b>	6.23	1.173
<b>10</b>	5.19	0.783
<b>5</b>	3.97	0.607
<b>6</b>	4.23	0.182

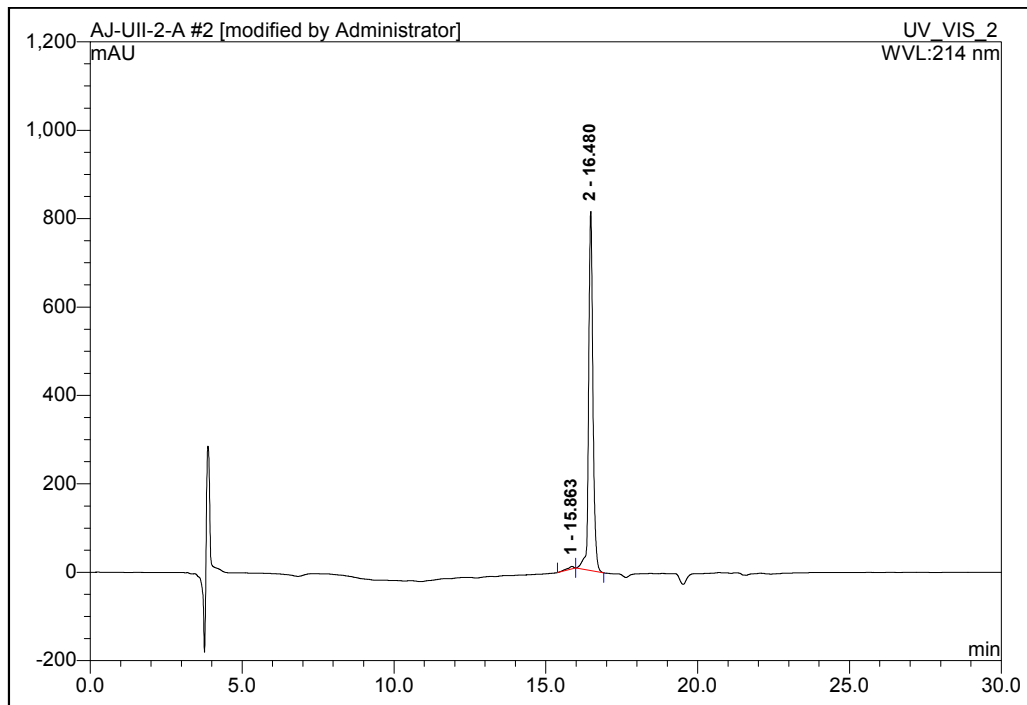
**3. Table S1: Yields of purified analogues**

Analogue	Yield (mg)	Yield	Purity	MS [M+H] <sup>+</sup>	
				Calcd.	Found
1,5-Triazole Bridged Urotensin II(4-11) <b>5</b>	15 mg	14 %	98%	1106.5060	1106.5077
1,5-Triazole Bridged Urotensin II(4-11) <b>6</b>	10 mg	8 %	97%	1120.5216	1120.5220
1,4-Triazole Bridged Urotensin II(4-11) <b>9</b>	20 mg	18 %	99%	1106.5060	1106.5054
1,4-Triazole Bridged Urotensin II(4-11) <b>10</b>	17 mg	15 %	97%	1120.5216	1120.5195

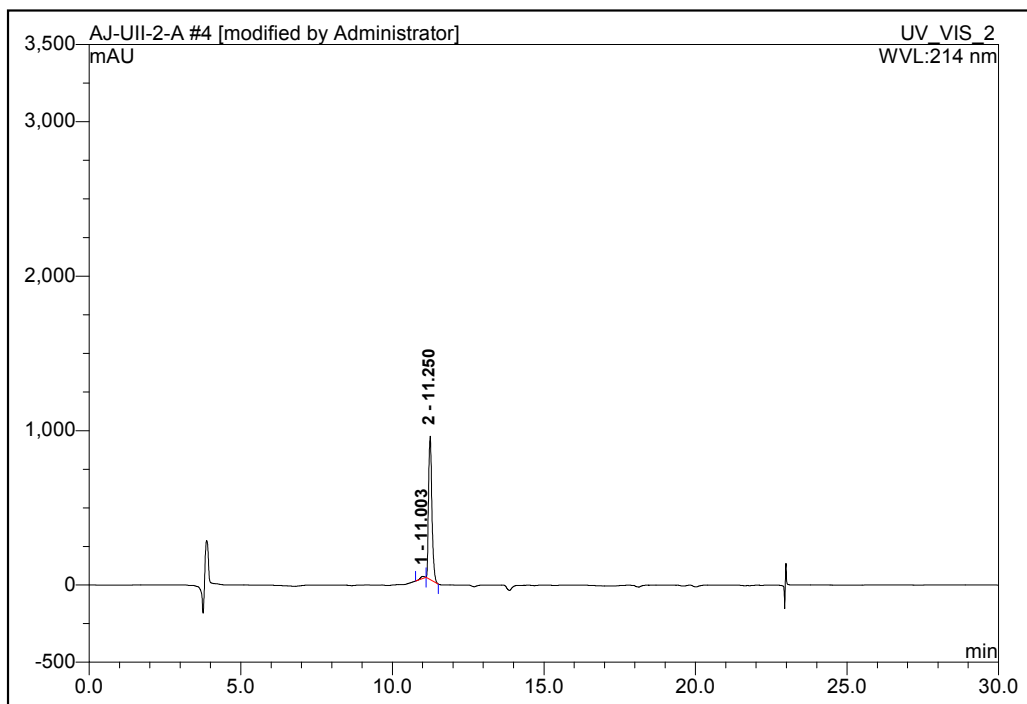
#### 4. RP-HPLC and MS data

1,5-Triazole Bridged Urotensin II<sup>(4-11)</sup> **5**

Analytical RP-HPLC: Gradient: 5-100% B, over 15 min.

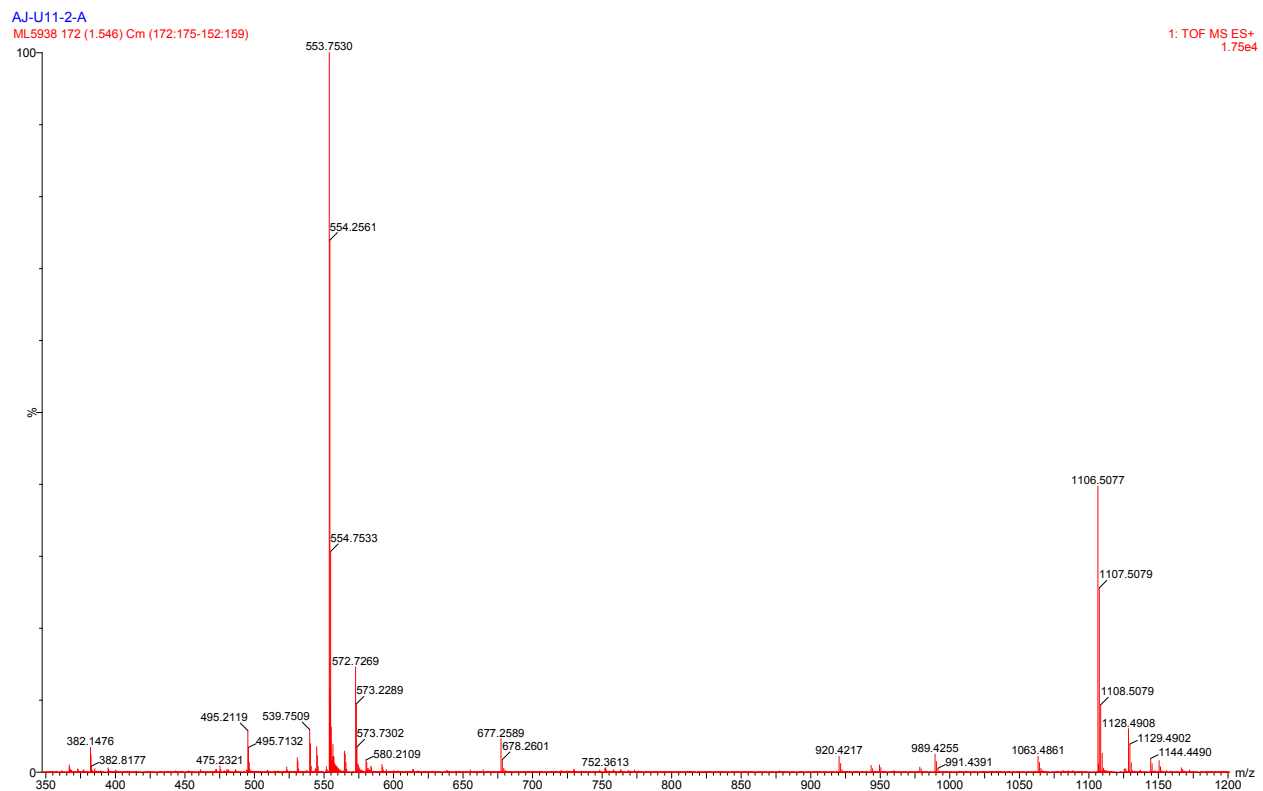


Analytical RP-HPLC: Gradient: 5-50% B, over 15 min.



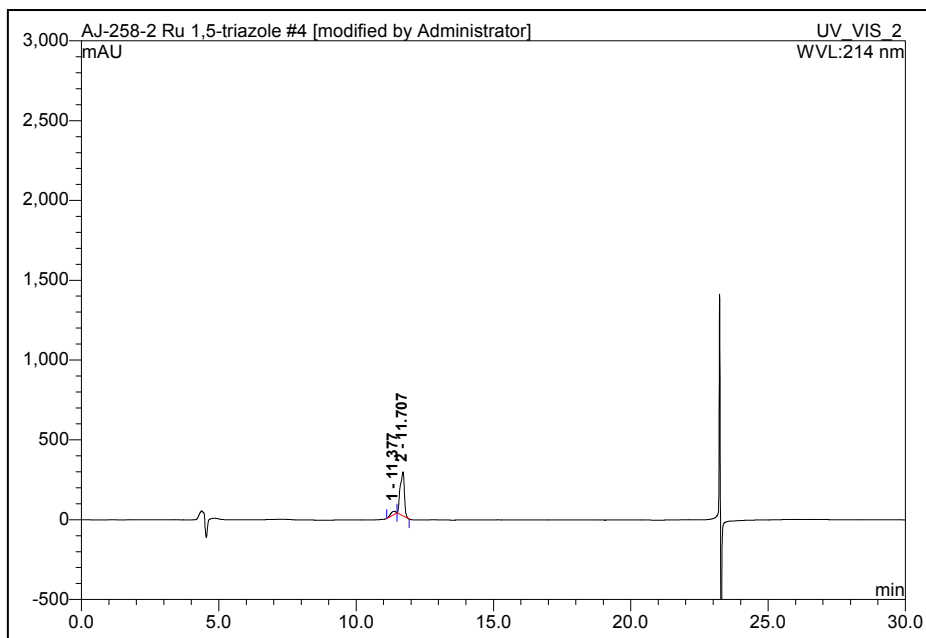
# ESI Mass Spectrometry

1,5-triazole bridged Urotensin II(4-11) **5** - Chemical Formula:  $C_{54}H_{67}N_{13}O_{13}$ ; MW 1106

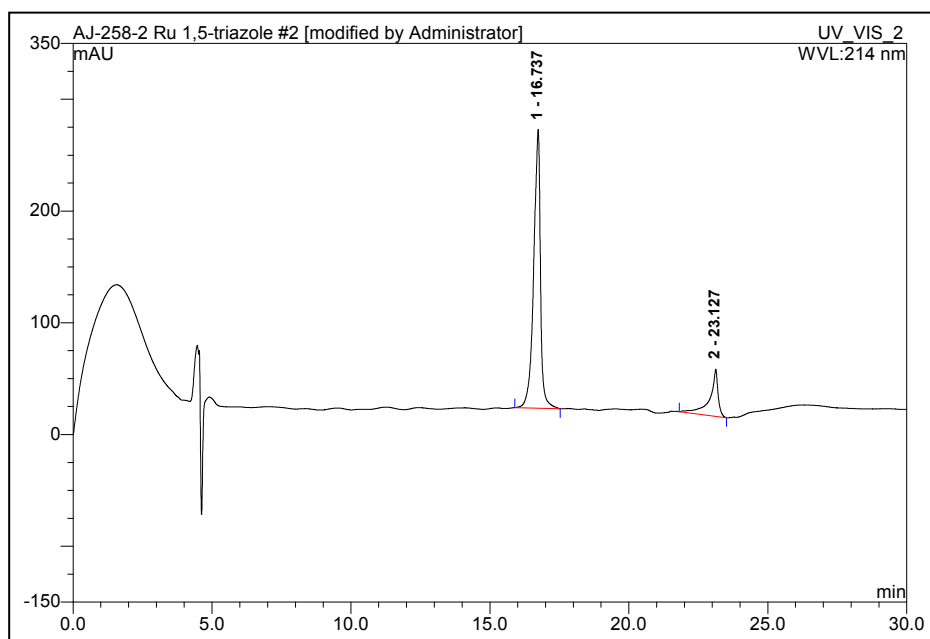


# 1,5-Triazole Bridged Urotensin II(4-11) 6

Analytical RP-HPLC: Gradient: 5-100% B, over 15 min.

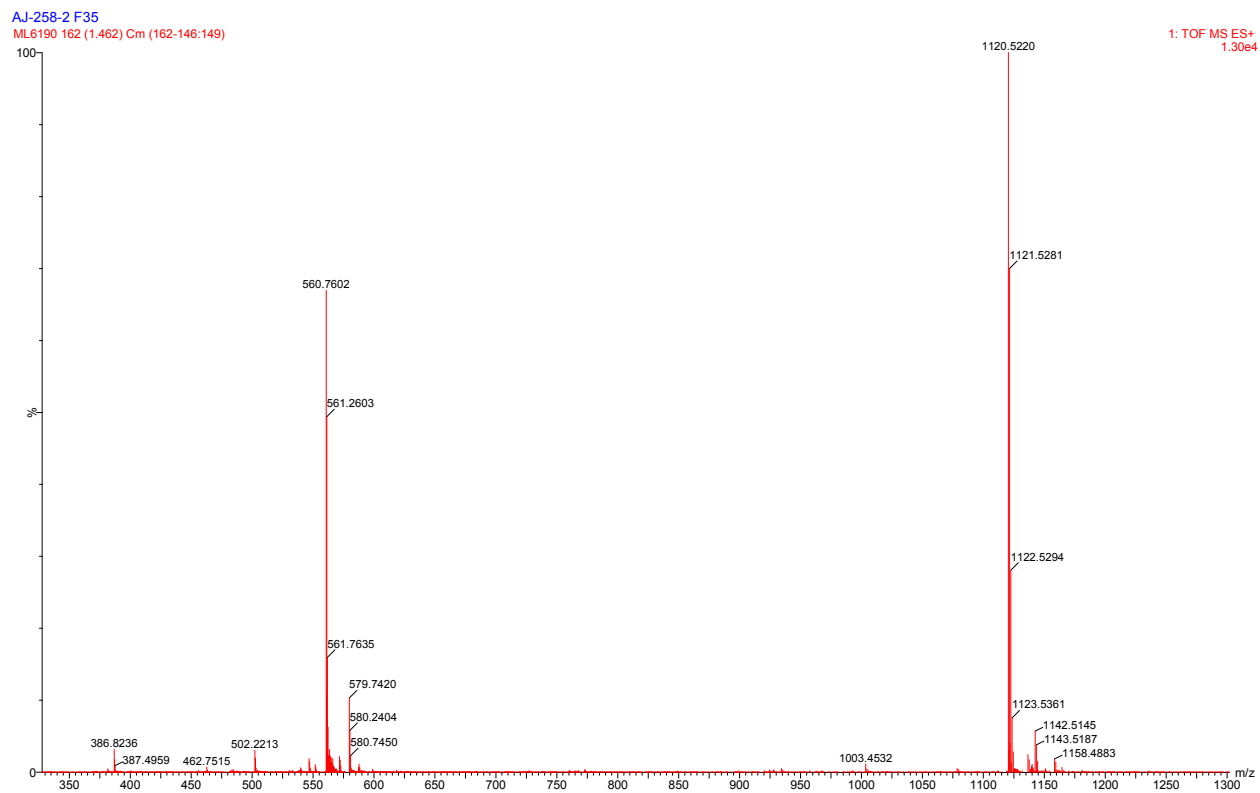


Analytical RP-HPLC: Gradient: 5-50% B, over 15 min.



# ESI Mass Spectrometry

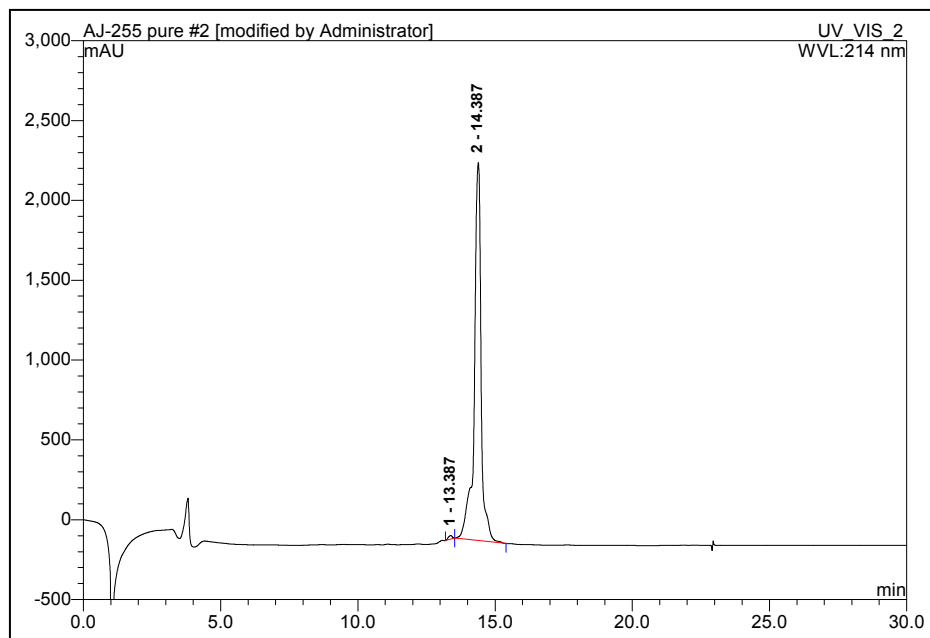
1,5-triazole bridged Urotensin II(4-11) **6** - Chemical Formula:  $C_{55}H_{69}N_{13}O_{13}$ ; MW 1120



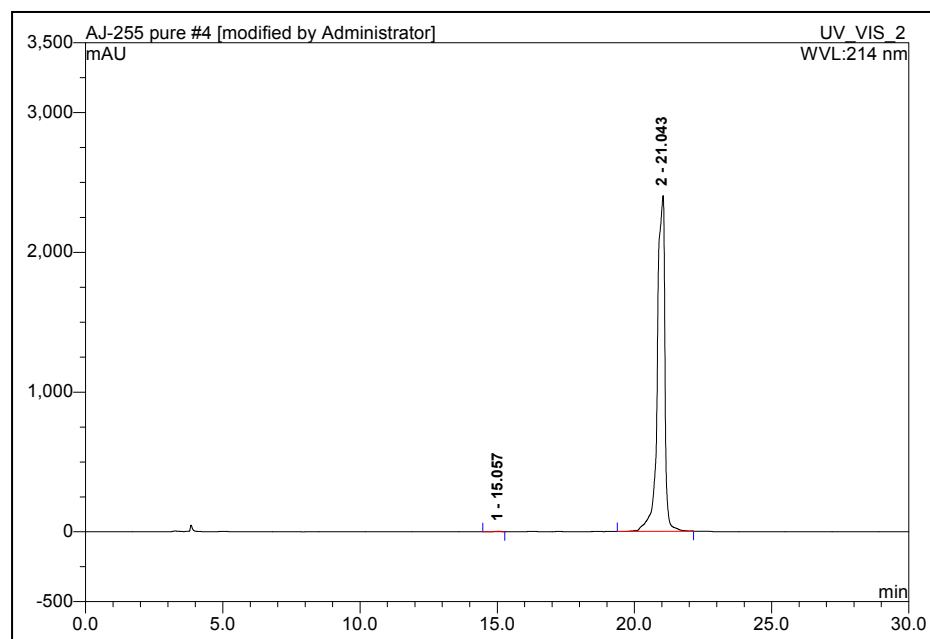


# 1,4-Triazole Bridged Urotensin II(4-11) **9**

Analytical RP-HPLC: Gradient: 5-100% B, over 15 min.

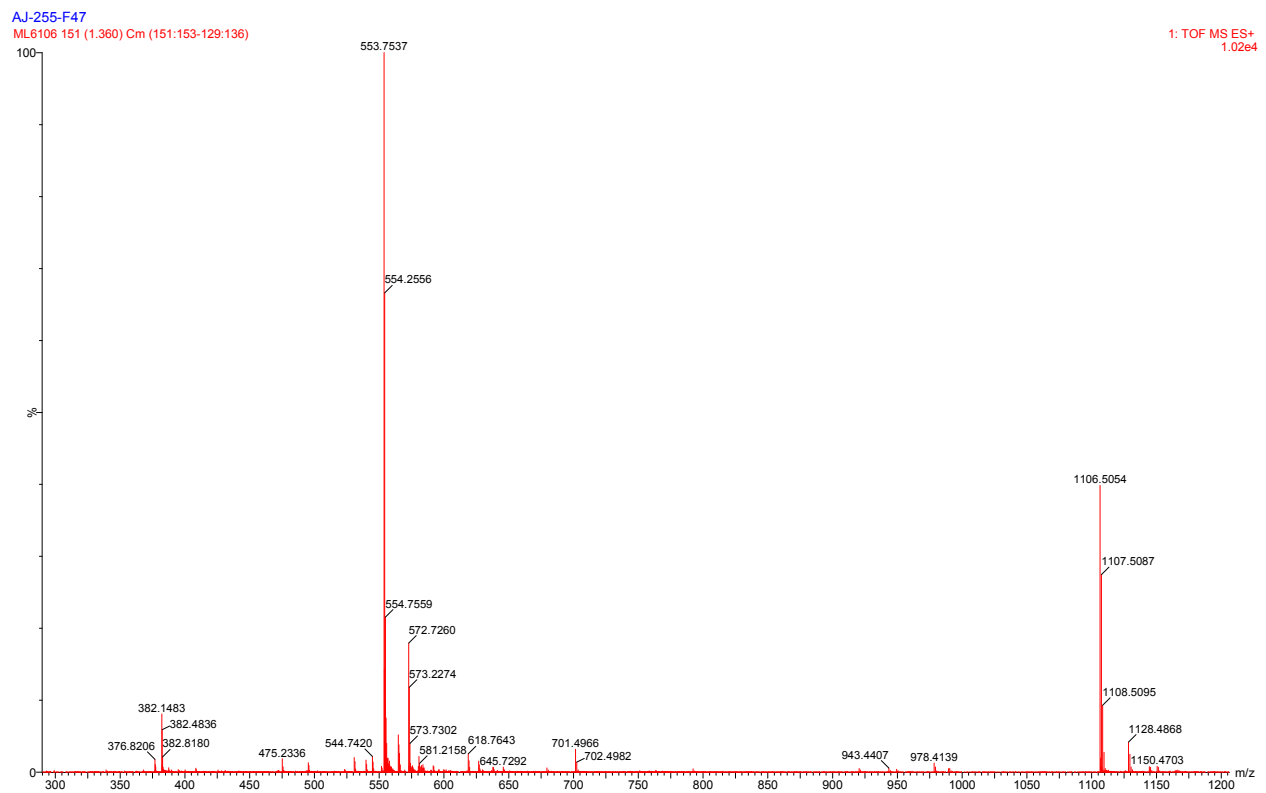


Analytical RP-HPLC: Gradient: 5-50% B, over 15 min.



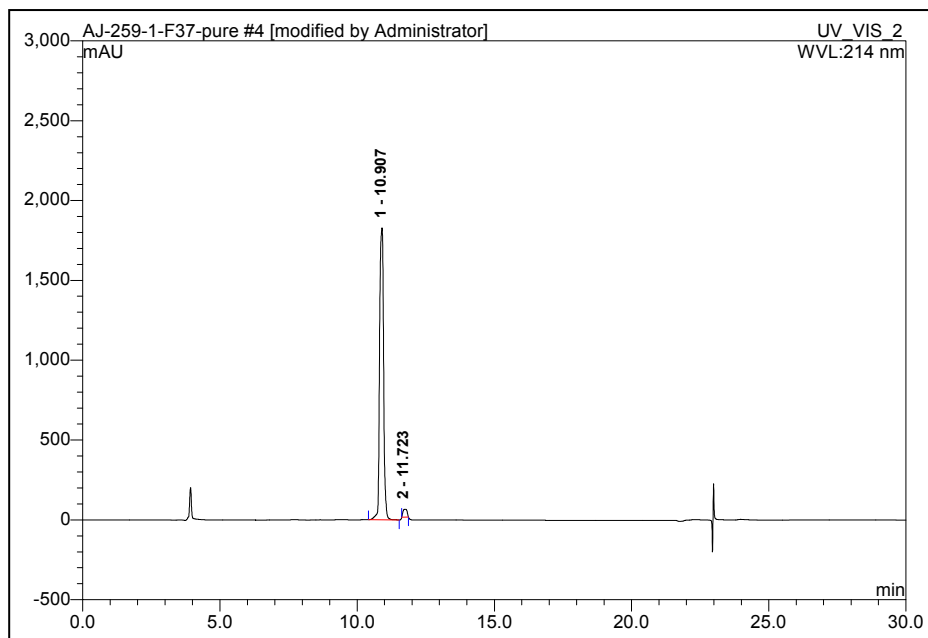
# ESI Mass Spectrometry

1,4-Triazole Bridged Urotensin II(4-11) **9** - Chemical Formula:  $C_{54}H_{67}N_{13}O_{13}$ ; MW 1106

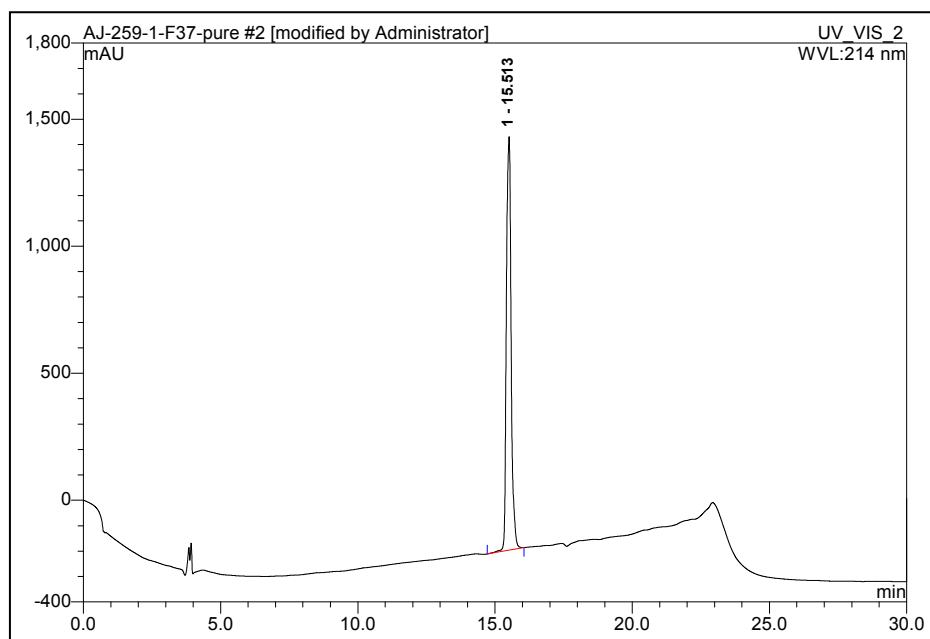


# 1,4-Triazole Bridged Urotensin II(4-11) **10**

Analytical RP-HPLC: Gradient: 5-100% B, over 15 min.



Analytical RP-HPLC: Gradient: 5-50% B, over 15 min.



# ESI Mass Spectrometry

1,4-Triazole Bridged Urotensin II(4-11) **10** - Chemical Formula:  $C_{55}H_{69}N_{13}O_{13}$ ; MW 1120

