

Rational Design of Potent α -Conotoxin PeIA Analogues with Non-Natural Amino Acids for the Inhibition of Human $\alpha 9\alpha 10$ Nicotinic Acetylcholine Receptors

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Table S1. Pairwise interactions between PeIA (left) and PeIA[S4Dap, S9Dap] (right) at $\alpha 9(+)\alpha 9(-)$ binding sites of $h\alpha 9\alpha 10$ nAChR.

PeIA residue	$\alpha 9(+)\alpha 9(-)$		$\alpha 9(+)\alpha 9(-)$		PeIA[S4Dap,S9Dap] residue
	$\alpha 9(+)$	$\alpha 9(-)$	$\alpha 9(+)$	$\alpha 9(-)$	
S4		D169		D166, S168, D169	Dap4
P6	W149		W149		P6
S9				Q34	Dap9
N11			N154		N11
H12	Y197		Y197		H12
P13		I59, L115		I59, L115	P13
E14		R111		R111	E14

Contacts between $h\alpha 9\alpha 10$ nAChR and PeIA/PeIA[S4Dap, S9Dap] are defined as van der Waals interactions if the distance between heavy atoms of them is between 2 and 4 Å. The electronic interactions are in bold.

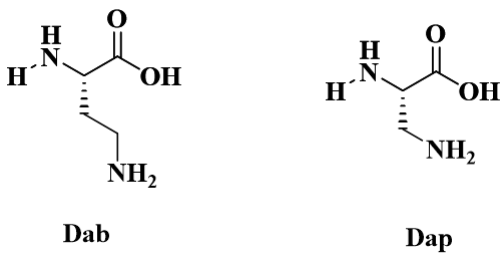


Figure S1. The structures of amino acids Dab-OH and Dap-OH. The non-proteinogenic amino acids Dab-OH and Dap-OH contain the same two and one methylene chain, respectively.

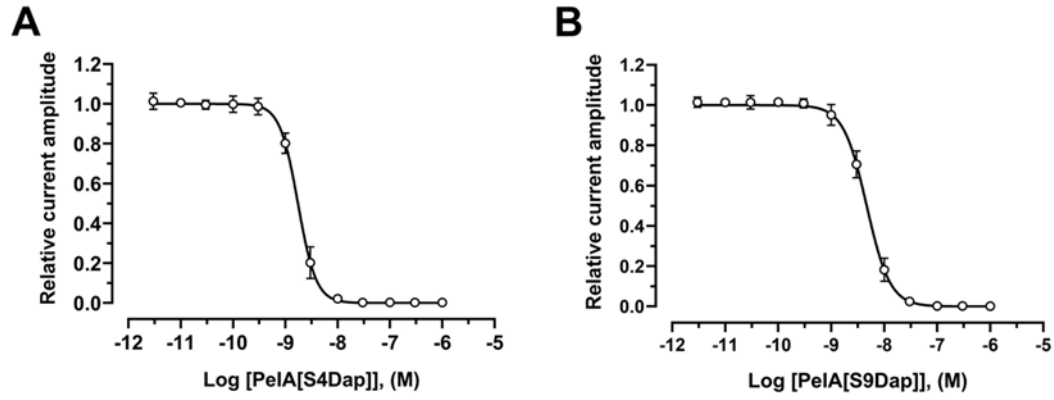


Figure S2. Concentration–response relationships of relative ACh-evoked current amplitude (mean \pm SD, $n = 7$) mediated by $\text{h}\alpha 9\alpha 10$ nAChRs in the presence of (A) PeIA[S4Dap] and (B) PeIA[S9Dap] giving IC_{50} 's of 1.74 nM (1.67 – 1.82; 95% CI) and 4.67 nM (4.45 – 4.90; 95% CI), respectively. Whole-cell currents at $\text{h}\alpha 9\alpha 10$ were activated by 6 μM ACh.

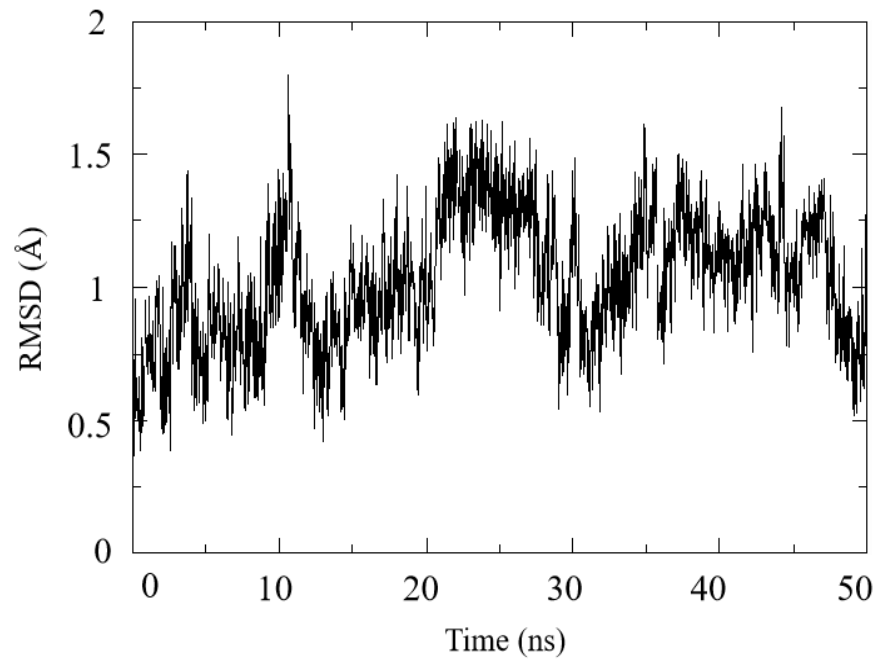


Figure S3. Evolution of root mean square deviation (RMSD) of PeIA.

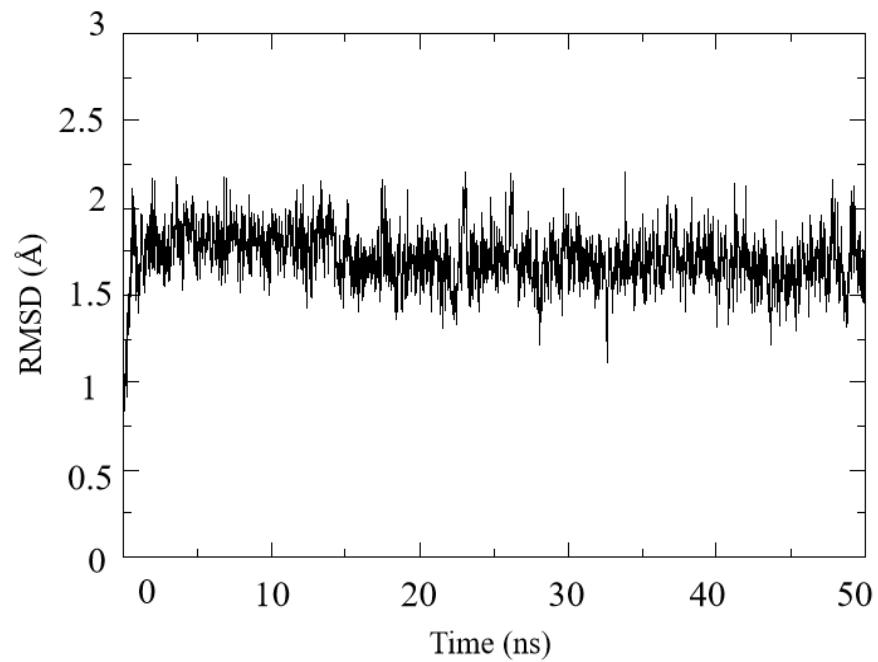


Figure S4. Evolution of root mean square deviation (RMSD) of PeIA[S4Dap, S9Dap].

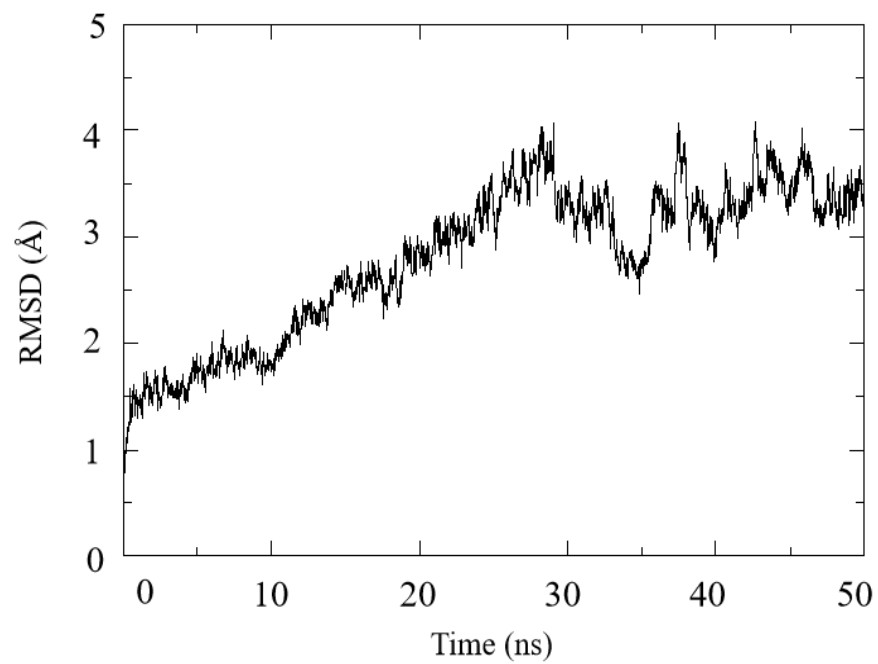


Figure S5. Evolution of root mean square deviation (RMSD) of PeIA bound to h α 9 α 10 nAChR.

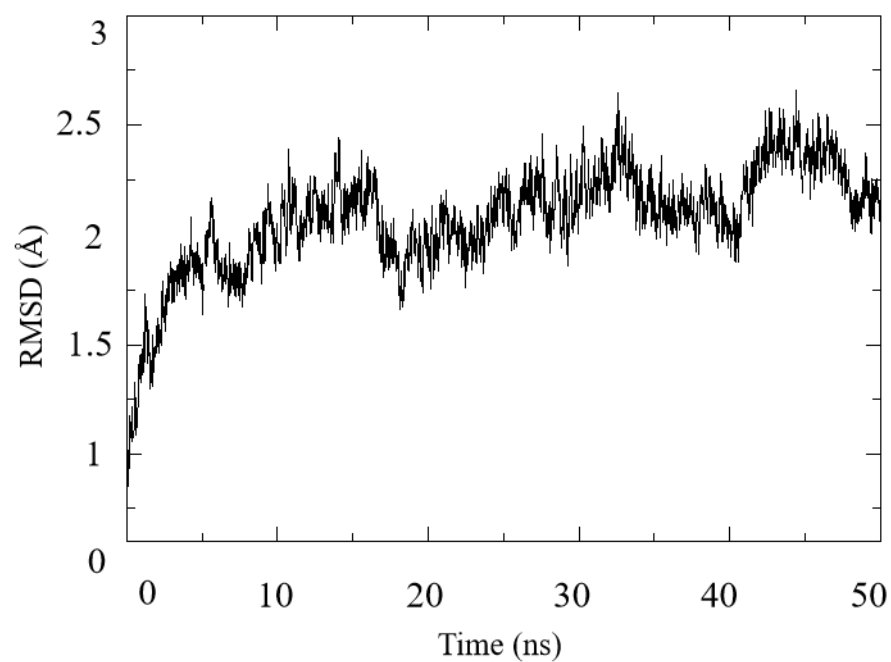
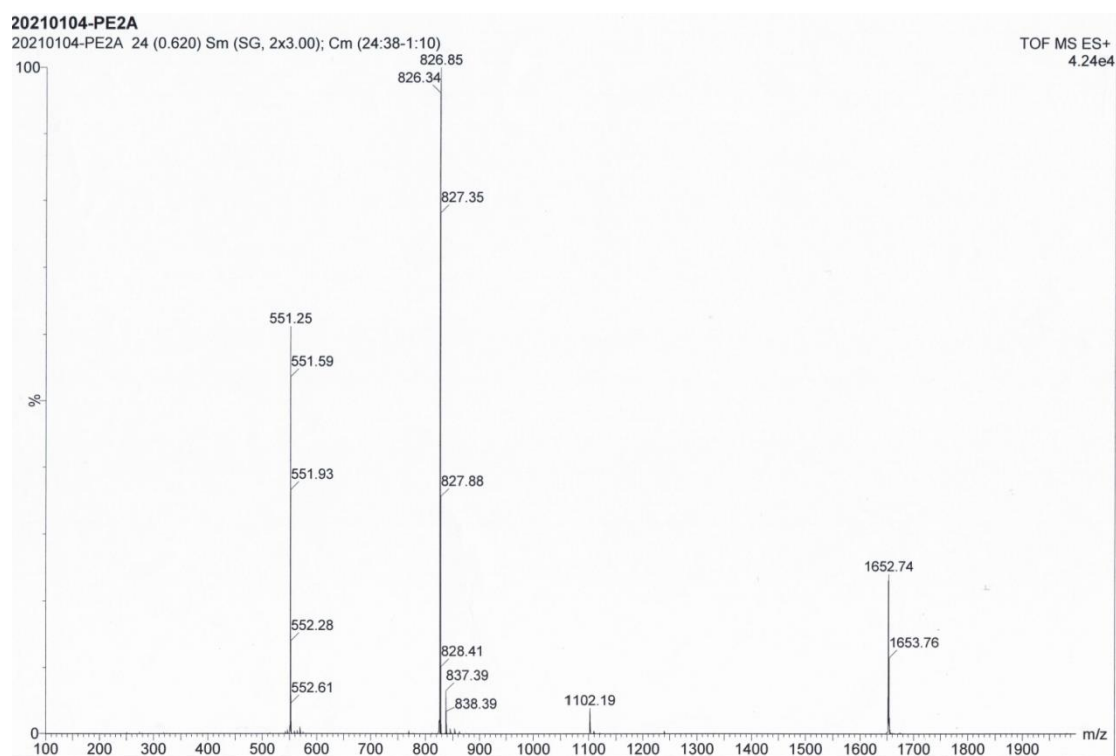


Figure S6. Evolution of root mean square deviation (RMSD) of PeIA[S4Dap, S9Dap] bound to hα9α10 nAChR.



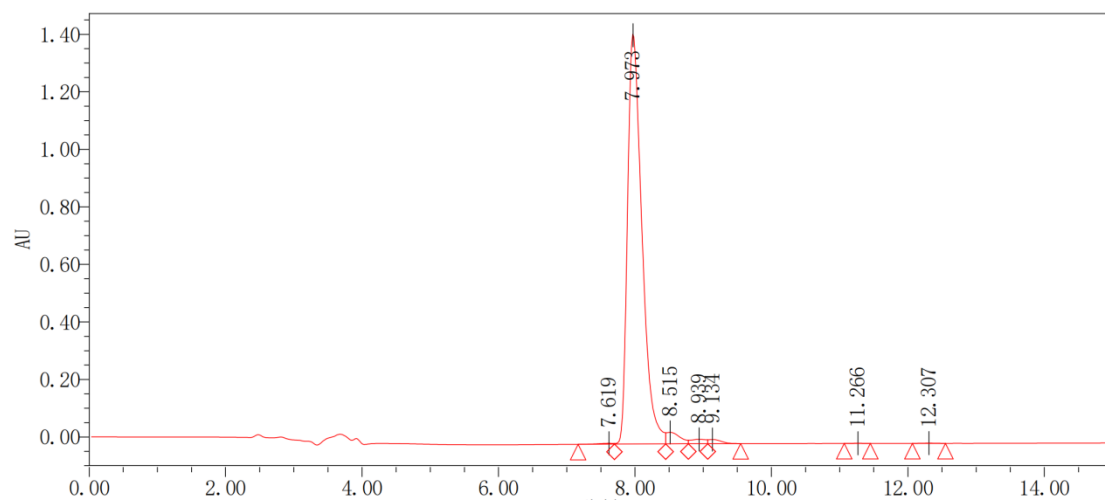
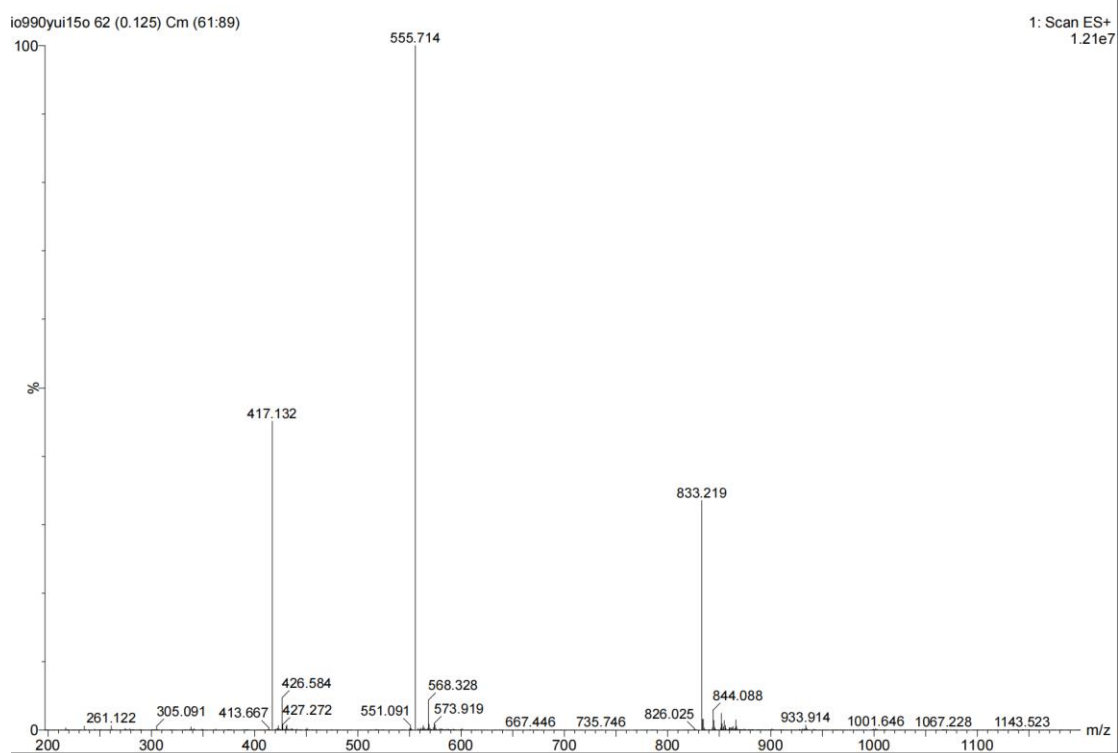


Figure S7. The ESI-MS and RP-HPLC of PeIA with a purity of 95.36%.



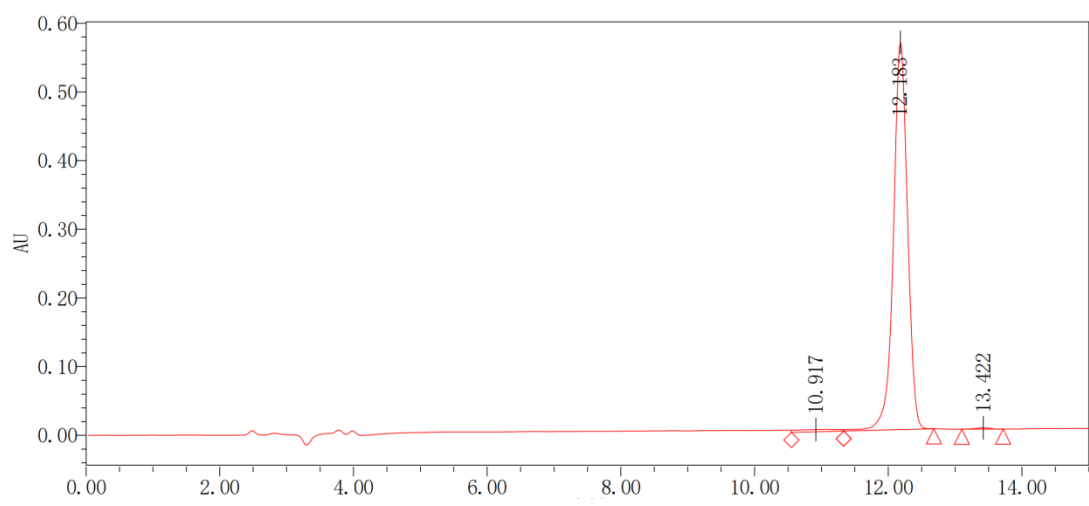


Figure S8. The ESI-MS and RP-HPLC of PeIA[S4Dab] with a purity of 99.67%.



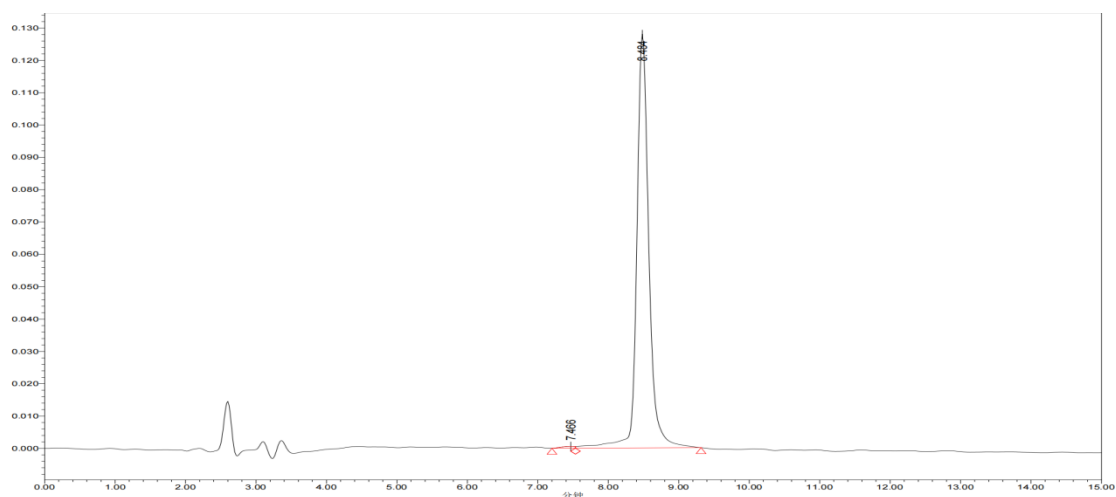


Figure S9. The ESI-MS and RP-HPLC of PeIA[S4Dap] with a purity of 99.53%.

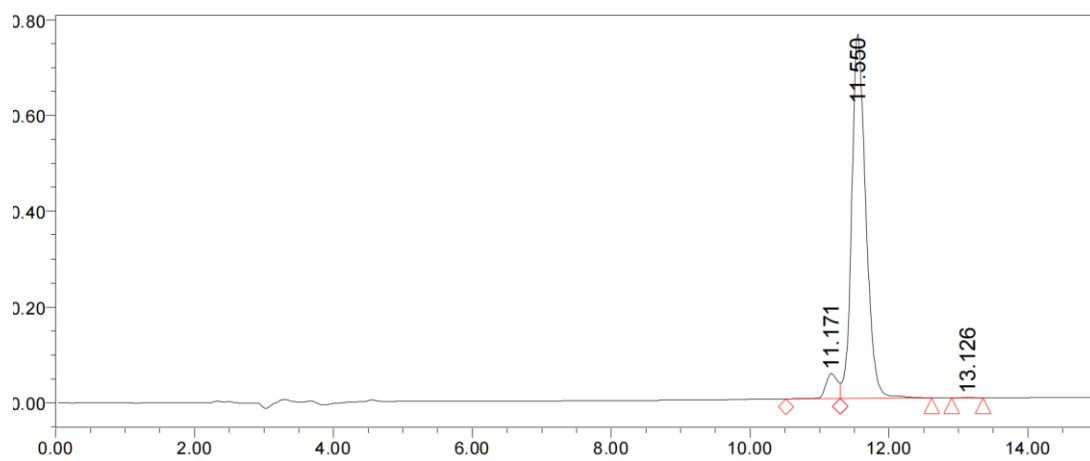
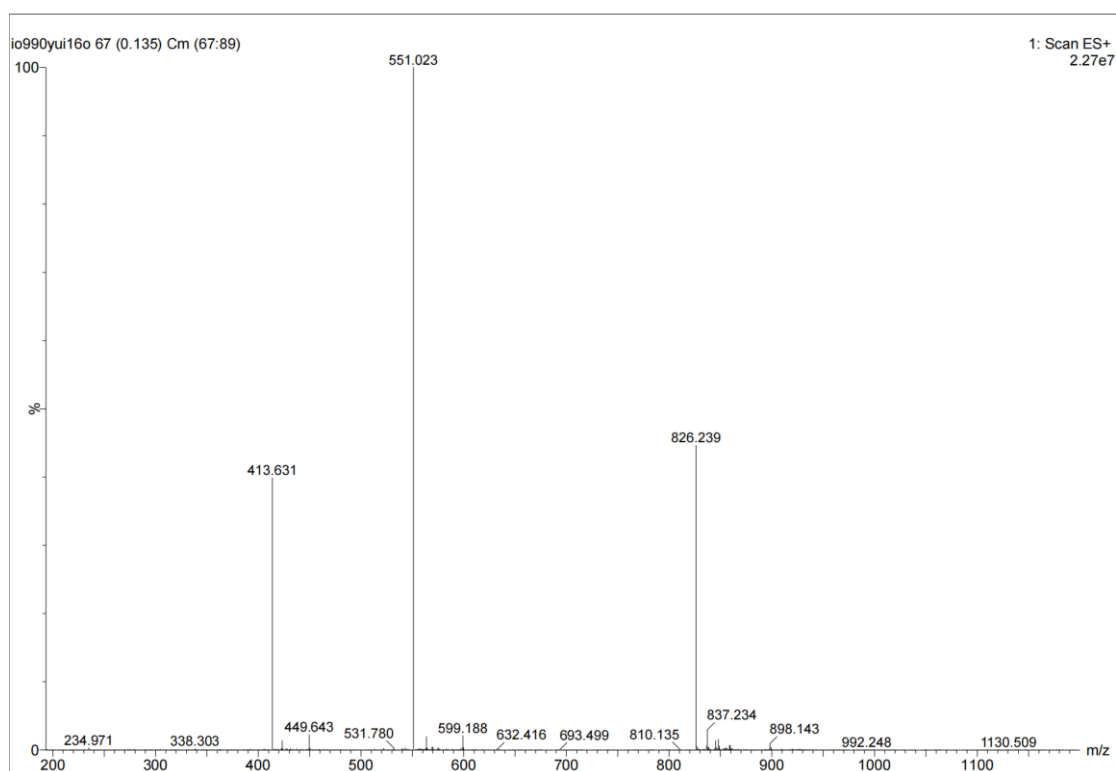


Figure S10. The ESI-MS and RP-HPLC of PeIA[S9Dap] with a purity of 94.68%.

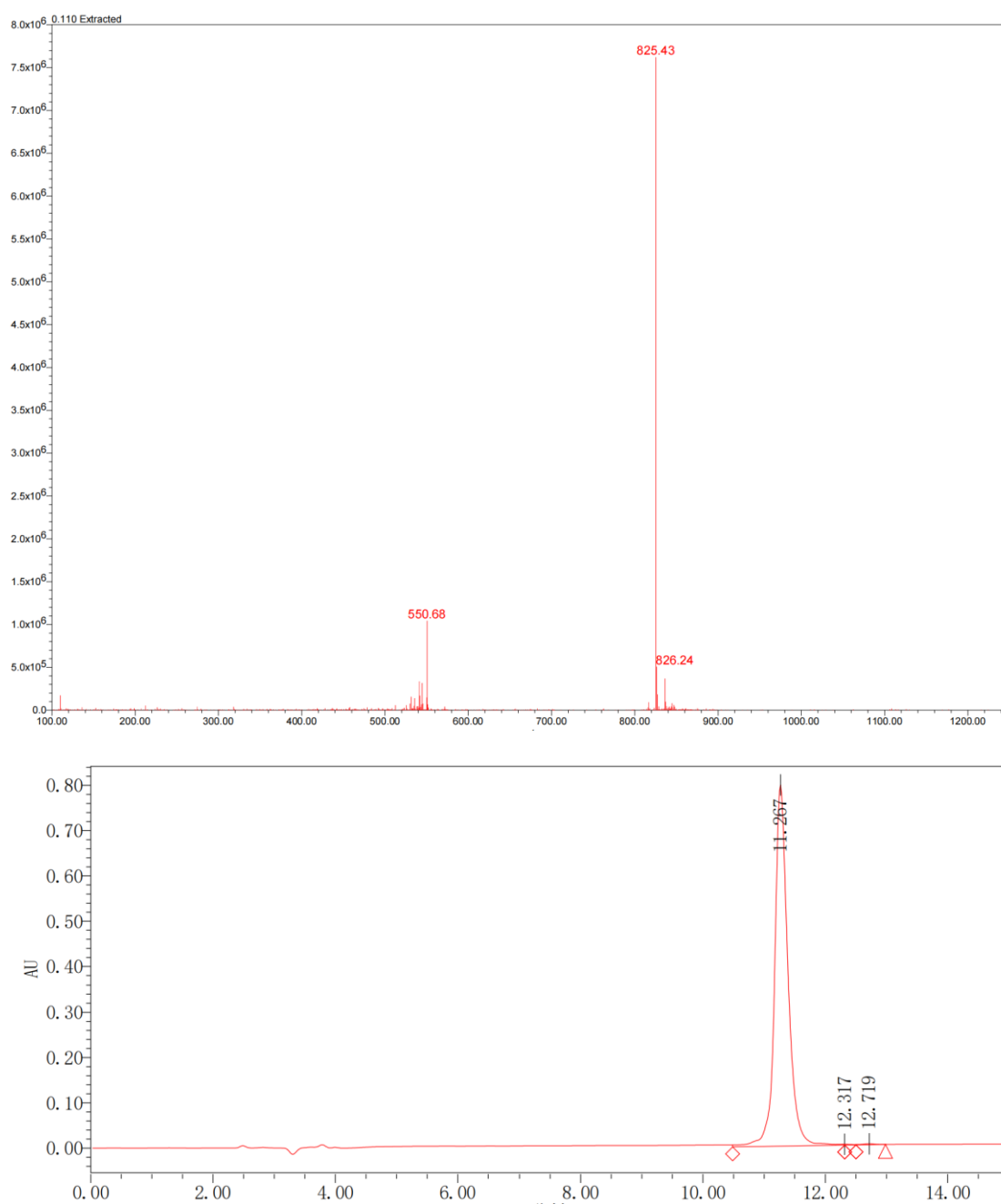


Figure S11. The ESI-MS and RP-HPLC of PeIA[S4Dap, S9Dap] with a purity of 99.28%.