

Supporting Information

Structure-based Design and Development of Chemical Probes Targeting Putative MOR-CCR5 Heterodimers to Inhibit Opioid Exacerbated HIV-1 Infectivity

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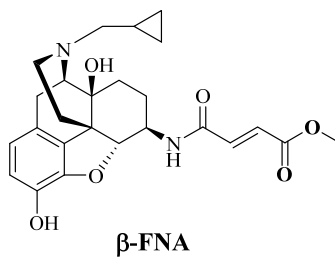
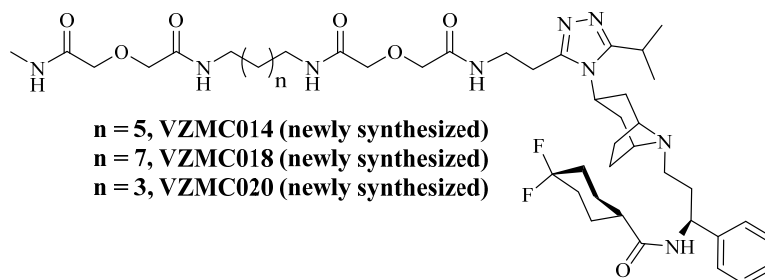
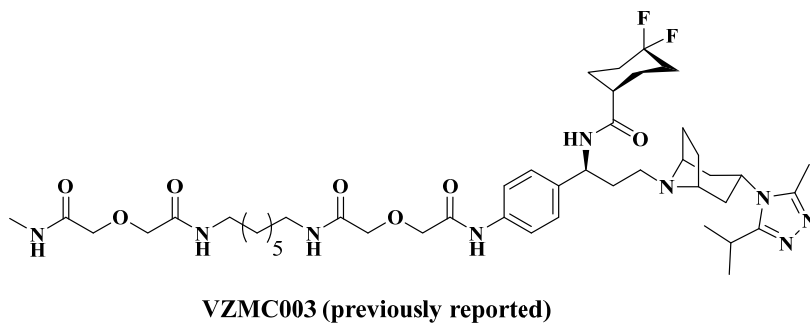
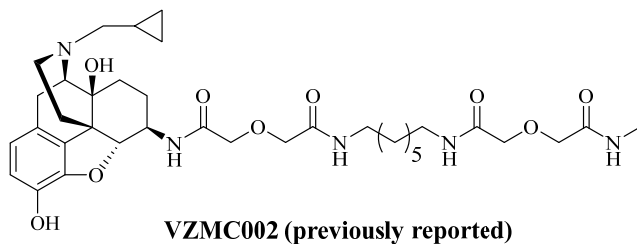


Figure S1. Chemical structures of corresponding bivalent ligands and monovalent control ligands and β -FNA.

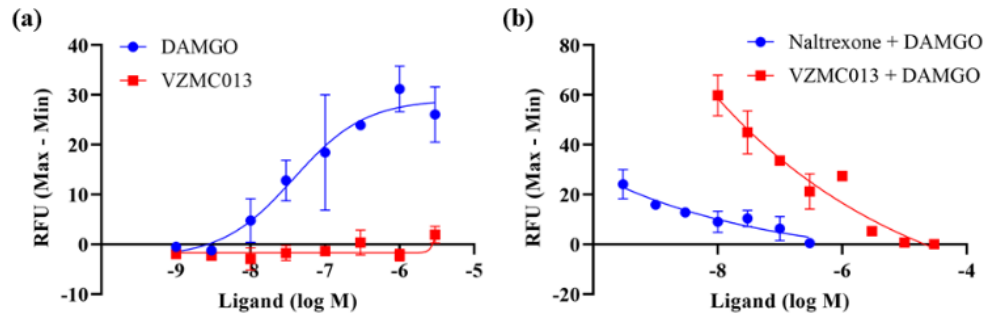


Figure S2. Effects of bivalent compound **VZMC013** on DAMGO-induced Ca^{2+} mobilization in Gqi5-transfected mMOR-CHO cells. (a) **VZMC013** exhibited no agonism to increase the intracellular calcium level. DAMGO was used as a control. (b) **VZMC013** dose-dependently antagonized the DAMGO-induced intracellular calcium increase. Naltrexone was used as a control.

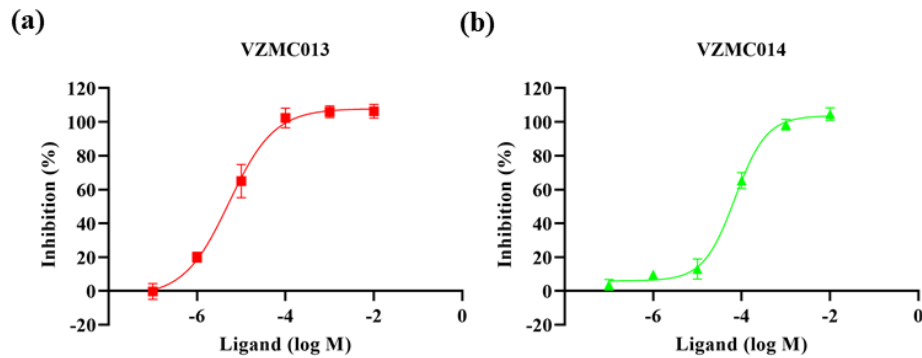


Figure S3. CCR5 Inhibition (%) corresponds to the ability of a tested compound to displace the radiolabeled ligand. (a) The competitive radioligand binding curve of **VZMC013**. (b) The competitive radioligand binding curve of **VZMC014**.

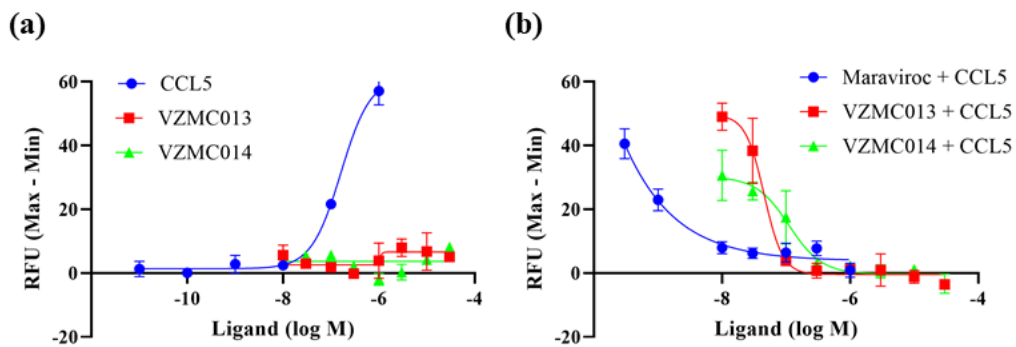


Figure S4. Effects of bivalent compound **VZMC013** and monovalent compound **VZMC014** on CCL5-induced Ca^{2+} mobilization in Gqi5-transfected HOS-CCR5 cells. (a) **VZMC013** and **VZMC014** failed to increase the intracellular calcium levels, suggesting a complete lack of agonist properties at CCR5. CCL5 was used as a control. (b) **VZMC013** and **VZMC014** dose-dependently antagonized CCL5-induced intracellular calcium increases. Maraviroc was used as a control.

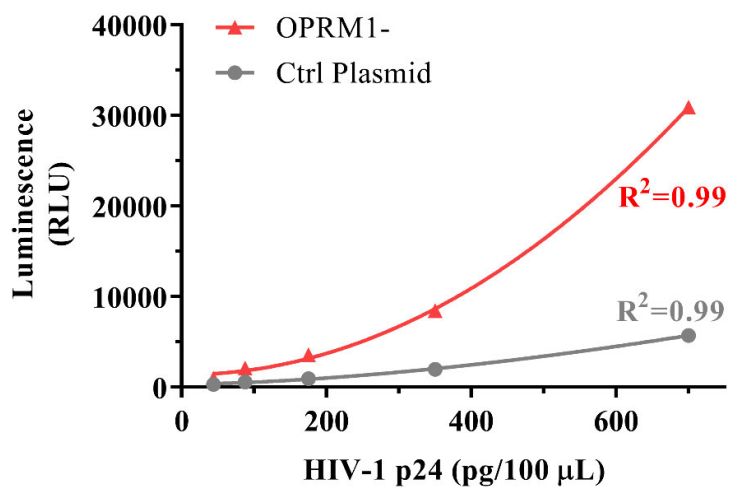


Figure S5. Comparison of intracellular concentration of HIV-1_{BaL} (RLU) in OPRM1- and control plasmid transfected TZM-bl cells.

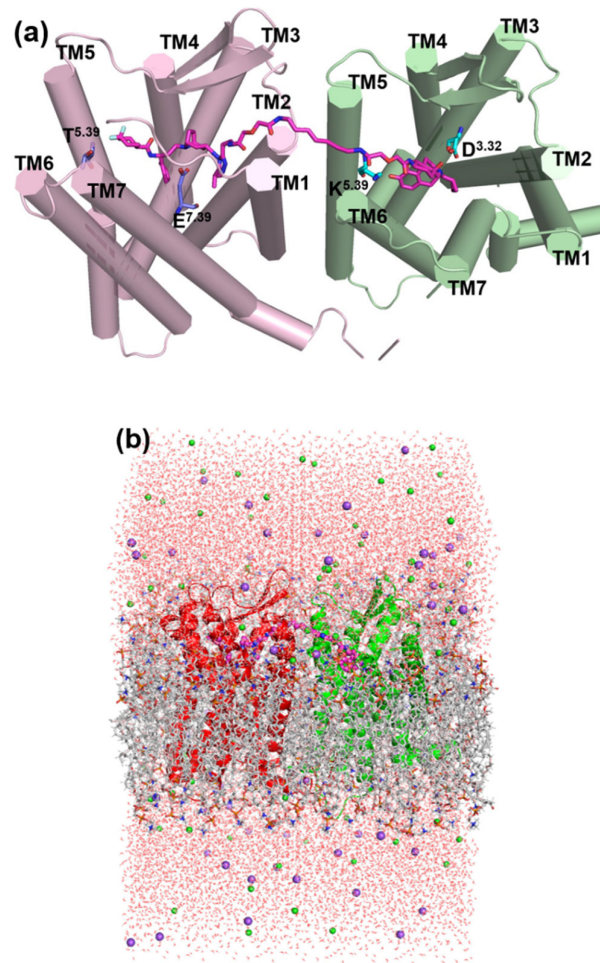
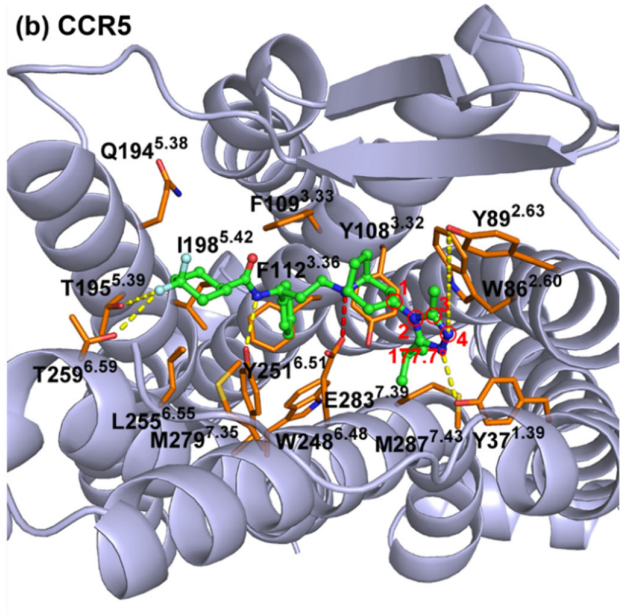
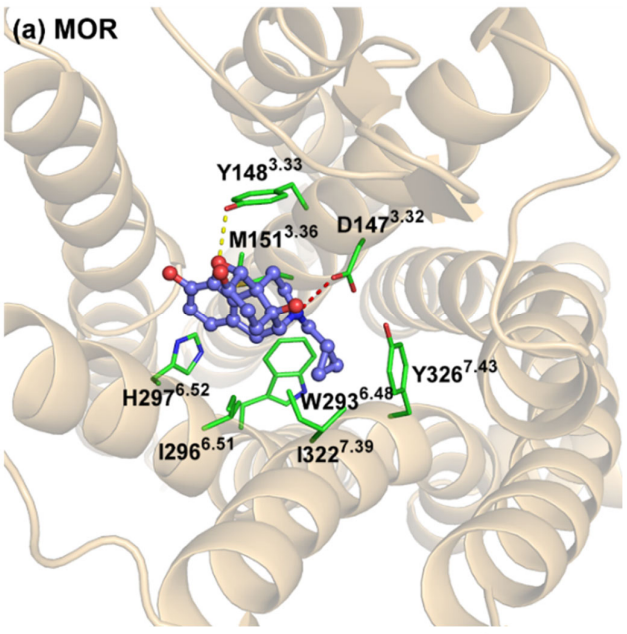


Figure S6. (a) The binding mode of VZMC013 in the MOR-CCR5 heterodimer complex. (b) The MOR-CCR5_VZMC013 complex in the membrane-aqueous sodium chloride solution system.



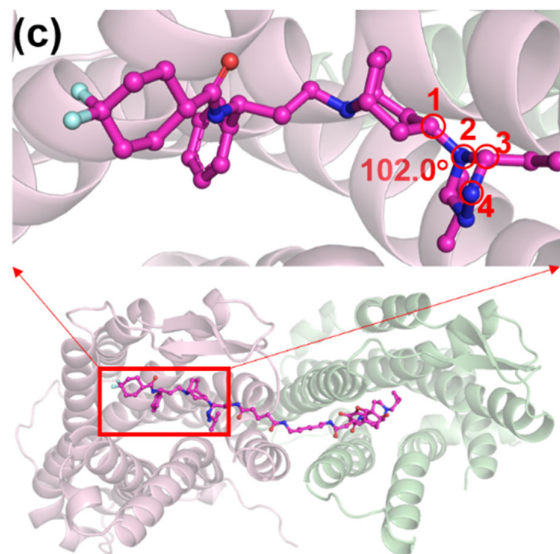


Figure S7. (a) The docking pose of naltrexone in the inactive MOR. (b) The binding mode of maraviroc in the crystal structure of the CCR5. The MOR and CCR5 shown as cartoon models in light-orange and light-blue, respectively. The key residues in the MOR and CCR5 shown as stick models in green and orange, respectively. Naltrexone and maraviroc shown as stick and sphere models in light-blue and green, respectively. The ionic interaction and hydrogen bond shown as dashed lines in red and yellow, respectively. (c) The rotation of the triazole moiety of **VZMC013** in the MOR-CCR5_ **VZMC013** complex after 100 ns MD simulation. The atoms related to the dihedral angle were labeled with red circle.

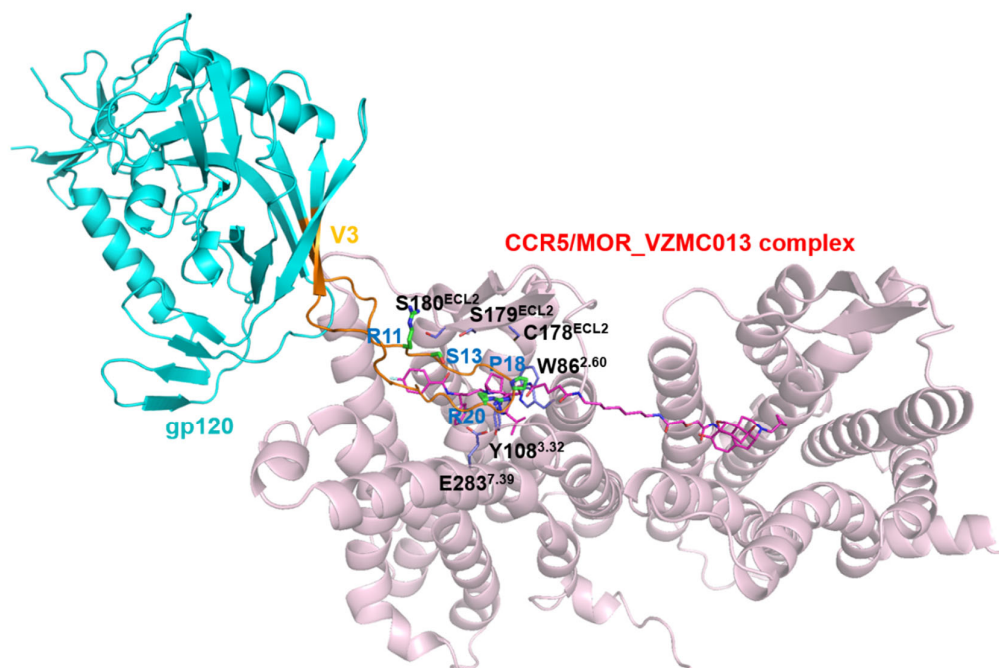


Figure S8. The putative binding of gp120 with the MOR-CCR5_VZMC013 complex. gp120 shown as a cartoon model in cyan. V3 loop of gp120 shown as a cartoon model in orange. MOR-CCR5 heterodimer shown as a cartoon model in pink. Compound VZMC013 shown as a stick model in magentas. Key residues of the CCR5 and V3 loop shown as stick models in light-blue and green, which labeled in black and blue text, respectively.

Table S1. MOR radioligand binding affinity.

Compounds	Ki (nM)
VZMC002	9.2 ± 3.4 ^a
Naltrexone	0.7 ± 0.1 ^a

^aData have been reported in Reference 1, and are presented here for comparison.

Table S2. CCR5 radioligand binding affinity.^a

Compounds	Ki (nM) ^b
VZMC014	41 ± 1
VZMC018	7.15 ± 0.26
VZMC020	ND ^c
MIP-1β	0.056 ± 0.006

^a[¹²⁵I]MIP-1α was used as the radioligand in the binding assay.

^bKi values were calculated using the Cheng-Prusoff equation. The values are the mean ± SEM of at least three independent experiments.

^cNot determined.

Table S3. Inhibition of CCL5-stimulated intracellular Ca²⁺ mobilization.^a

Compounds	IC ₅₀ (nM)
VZMC014	116 ± 2.53
VZMC018	39.6 ± 4.01
VZMC020	919 ± 45.4
VZMC003	622 ± 36 ^b
Maraviroc	0.77 ± 0.20

^aThe values are the mean ± SEM of at least three independent experiments.

^bData have been reported in Reference 1, and are presented here for comparison purposes.

Table S4. Inhibition of HIV-1_{BaL} and cytotoxicity in GHOST CCR5 cells.^a

Compounds	EC ₅₀ (μM)	TC ₅₀ (μM)	TI
VZMC014	2.62 ± 1.25	> 100	> 38
VZMC002	> 100	> 100	-
Maraviroc	0.018 ± 0.002	> 0.5	> 28

^aMeasured in triplicate.

Table S5. Residues within 5 Å proximity of the bivalent ligand in association with the MOR-CCR5_VZMC013 complex after 100 ns MD simulations.^a

Portions of the bivalent ligand VZMC013	Interactional residues
The CCR5 pharmacophore	Y37 ^{1.39} , W86 ^{2.60} , Y89 ^{2.63} , Y108 ^{3.32} , F109 ^{3.33} , F112 ^{3.36} , N163 ^{4.64} , C178 ^{ECL2} , S179 ^{ECL2} , S180 ^{ECL2} , F182 ^{ECL2} , T195 ^{5.39} , I198 ^{5.42} , W248 ^{6.48} , Y251 ^{6.51} , L255 ^{6.55} , T259 ^{6.59} , E283 ^{7.39} , M287 ^{7.43}
The MOR pharmacophore	D147 ^{3.32} , Y148 ^{3.33} , M151 ^{3.36} , L232 ^{5.38} , V236 ^{5.42} , I296 ^{6.51} , H297 ^{6.52} , V300 ^{6.55} , W318 ^{7.35} , I322 ^{7.39}
Spacer	A30 ^{1.32} , R31 ^{1.33} , A87 ^{2.61} , A90 ^{2.64} , A91 ^{2.65} , W226 ^{5.32} , N230 ^{5.36} , E229 ^{5.35} , K233 ^{5.39} , K303 ^{6.58}

^aThe residues in **Bold** represented the residues playing important roles in the binding of compound VZMC013 with the MOR-CCR5 heterodimer and these residues also presented in the binding of maraviroc and naltrexone in their respective receptors, the CCR5 and MOR.

Table S6. Residues with 5 Å around maraviroc and naltrexone in the respective receptors, CCR5 and MOR.^a

Complex	Residues
CCR5_Maraviroc	Y37 ^{1.39} , W86 ^{2.60} , Y89 ^{2.63} , Y108 ^{3.32} , F109 ^{3.33} , F112 ^{3.36} , Q194 ^{5.38} , T195 ^{5.39} , I198 ^{5.42} , W248 ^{6.48} , Y251 ^{6.51} , L255 ^{6.55} , T259 ^{6.59} , M279 ^{7.35} , E283 ^{7.39} , M287 ^{7.43}
MOR_Naltrexone	D147 ^{3.32} , Y148 ^{3.33} , M151 ^{3.36} , W293 ^{6.48} , I296 ^{6.51} , H297 ^{6.52} , I322 ^{7.39} , Y326 ^{7.43}

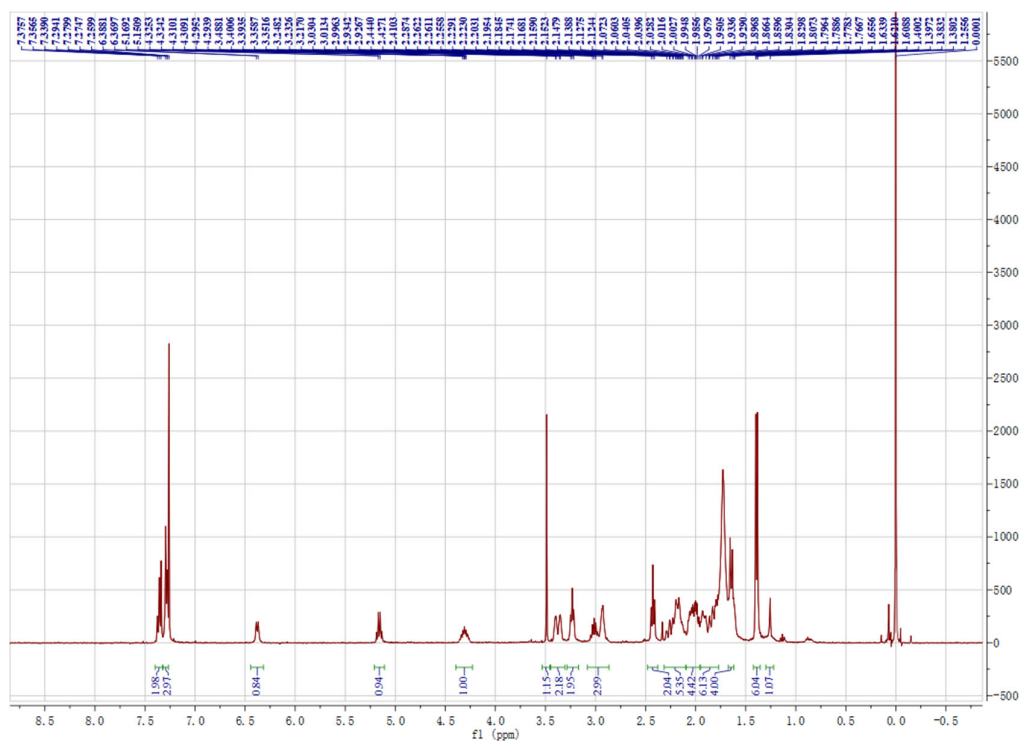
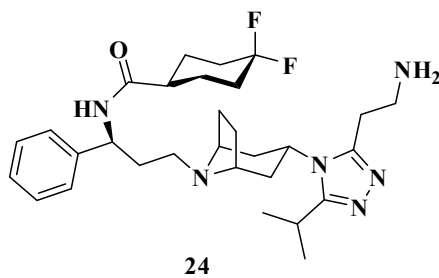
^aThe residues in **Bold** represented the residues played important roles in the binding of maraviroc and naltrexone with their respective receptors, CCR5 and MOR, also existed in the binding of compound VZMC013 with the MOR-CCR5 heterodimer.

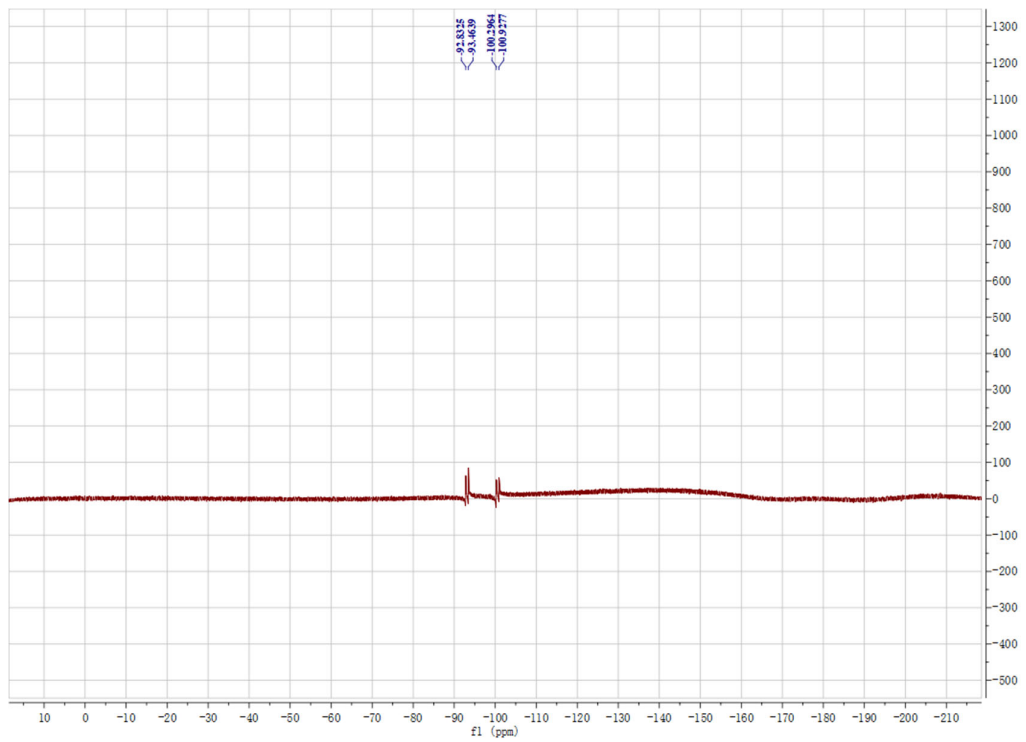
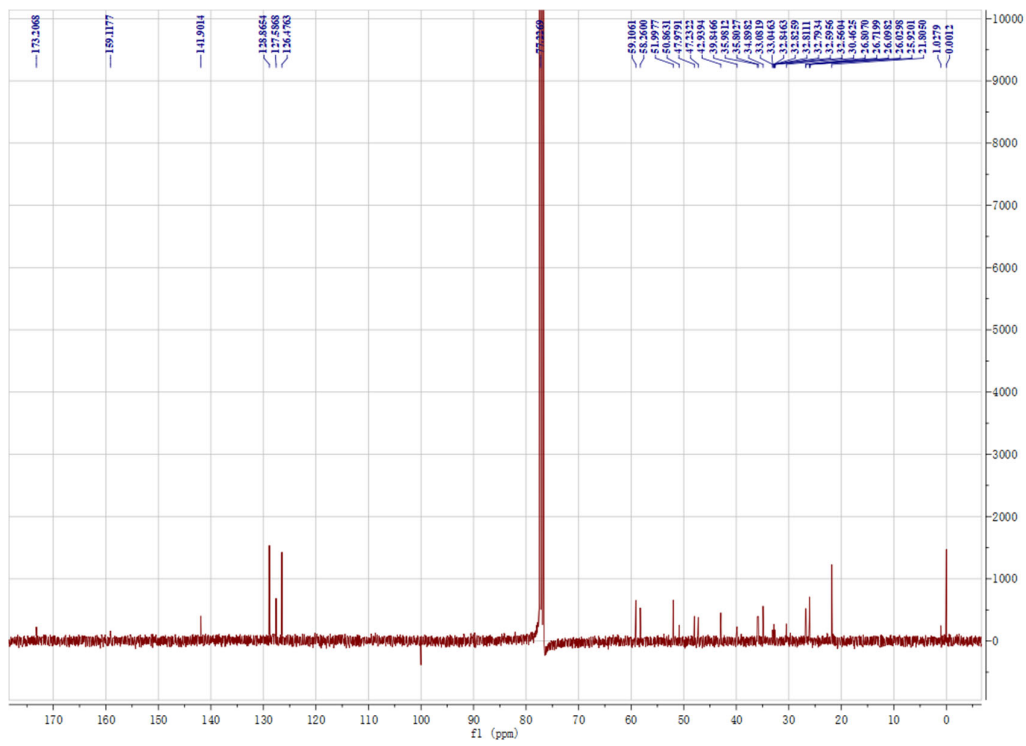
Table S7. The distances between atom of ligand and atom of receptor in the MOR_naltrexone, CCR5_maraviroc, and MOR-CCR5_VZMC013 complexes.

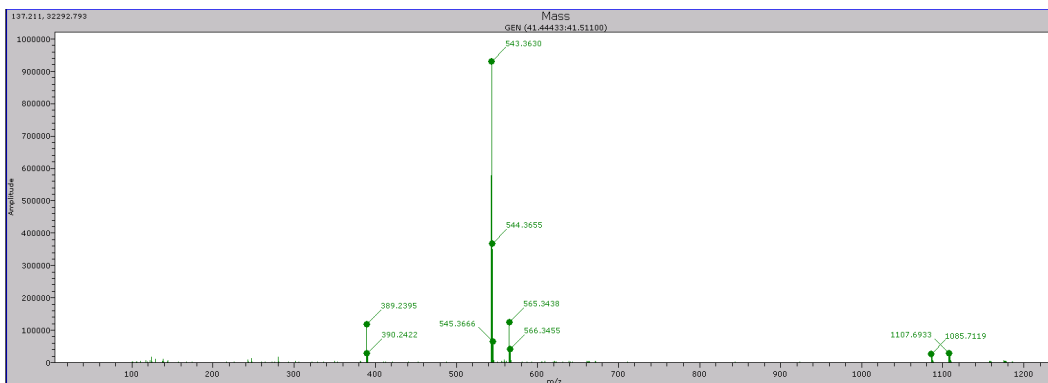
Complex	Atom@Ligand	Atom@Receptor	Distance (Å)
MOR_naltrexone	N17	OD2@D147 ^{3.32}	3.0
	O15	OH@Y148 ^{3.33}	2.8
	C5	CZ3@W293 ^{6.48}	4.6
	C36	CZ@Y326 ^{7.43}	3.6
CCR5_maraviroc	C9	CB@C178 ^{ECL2}	8.1
	C22	CZ@F182 ^{ECL2}	5.4
	O1	OG@S179 ^{ECL2}	6.0
	O1	OG@S180 ^{ECL2}	5.8
MOR-CCR5_VZMC013	N17	OD2@D147 ^{3.32}	7.1
	O15	OH@Y148 ^{3.33}	6.4
	C5	CZ3@W293 ^{6.48}	8.2
	C36	CZ@Y326 ^{7.43}	7.3
	C9	CB@C178 ^{ECL2}	5.2
	C22	CZ@F182 ^{ECL2}	3.5
	O1	OG@S179 ^{ECL2}	3.4
	O1	OG@S180 ^{ECL2}	3.6

Spectra of key intermediate 24 and target compounds

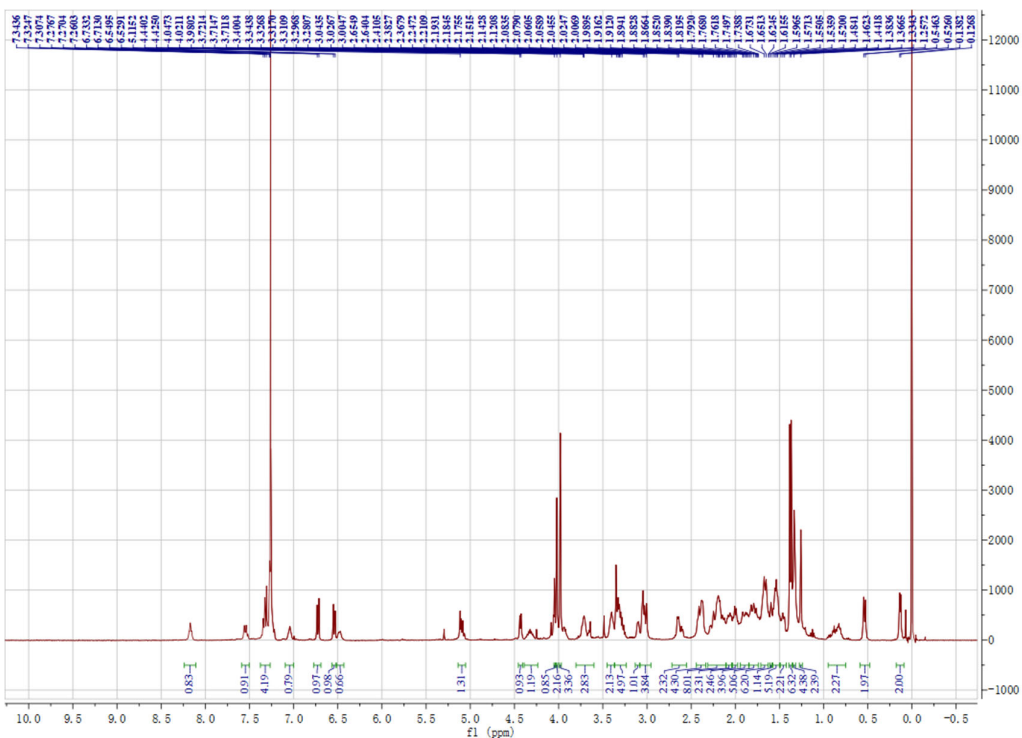
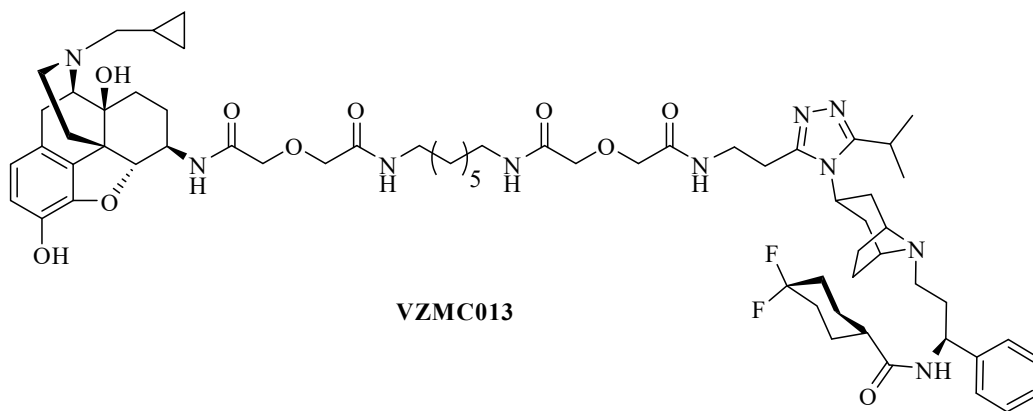
^1H NMR, ^{13}C NMR, ^{19}F NMR, and HRMS spectra for 24

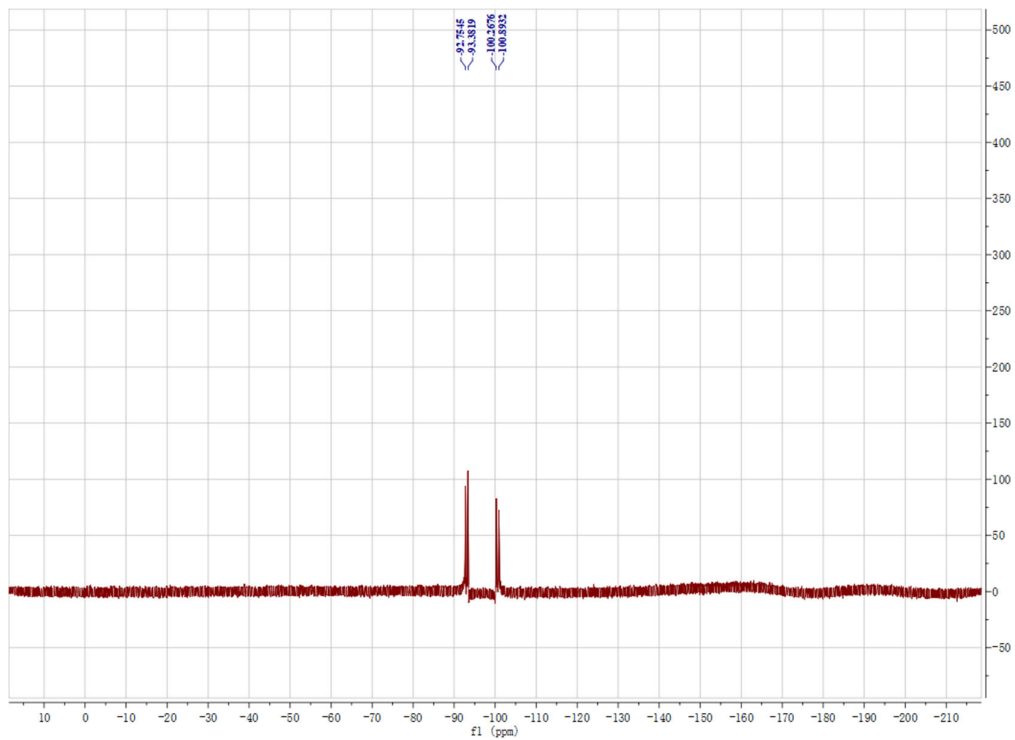
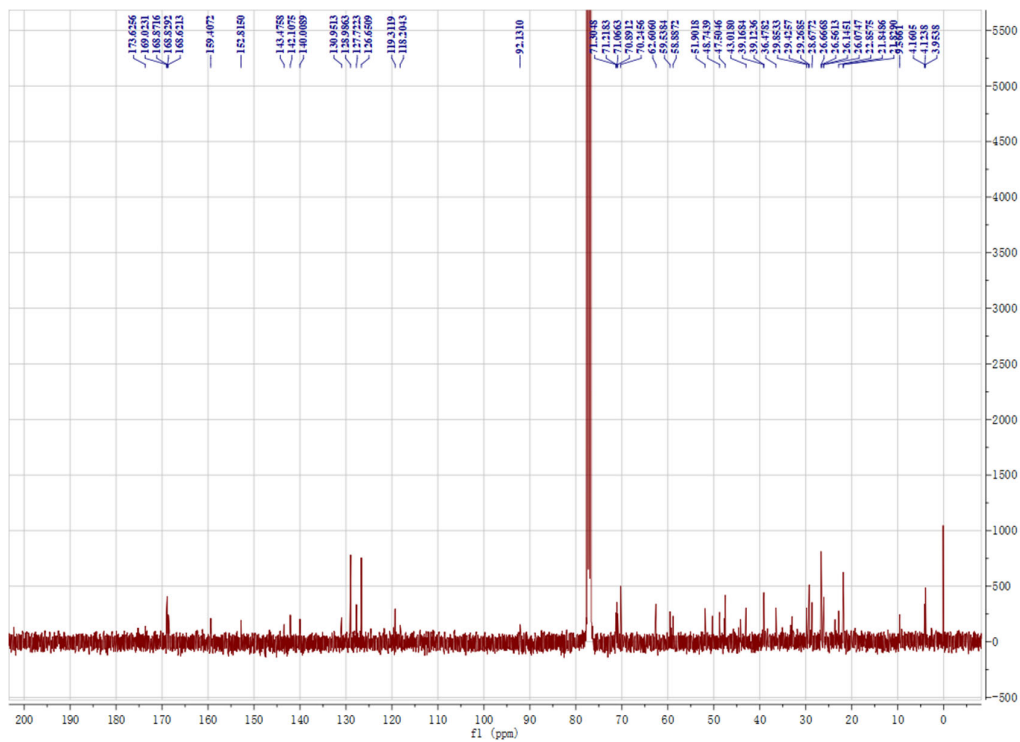


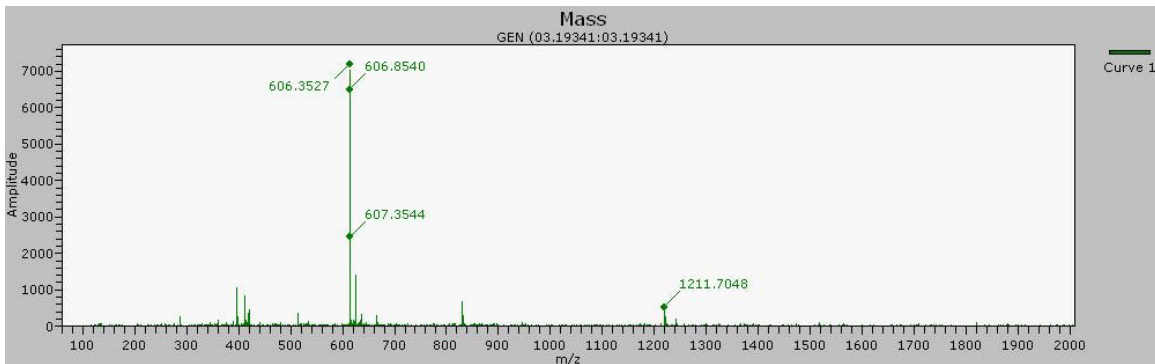




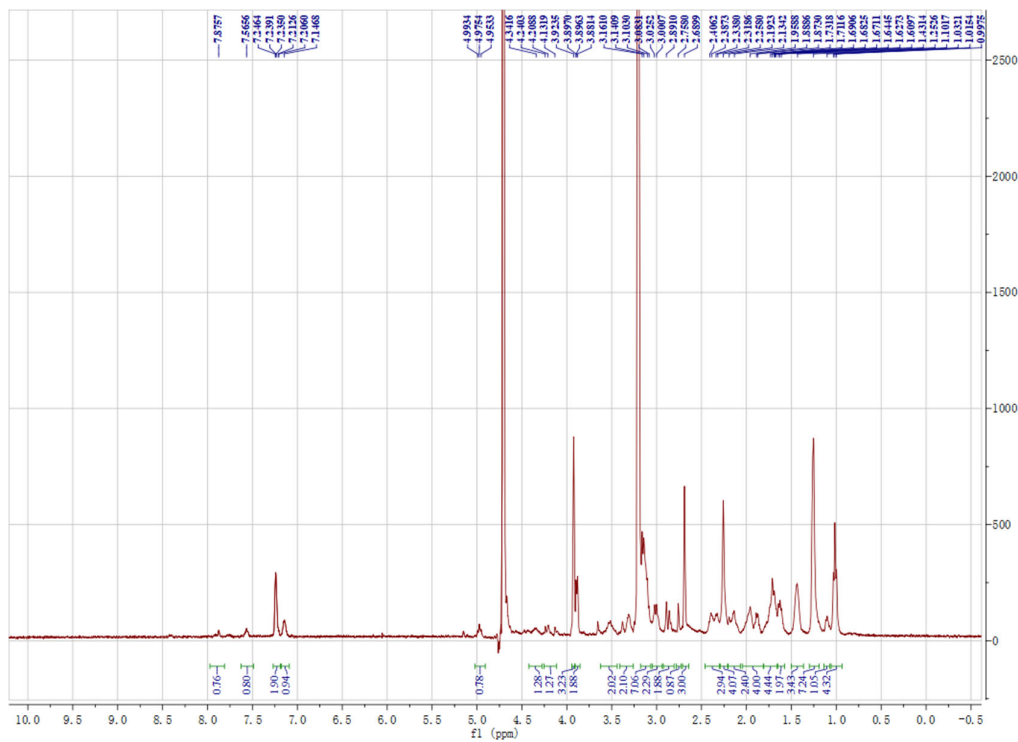
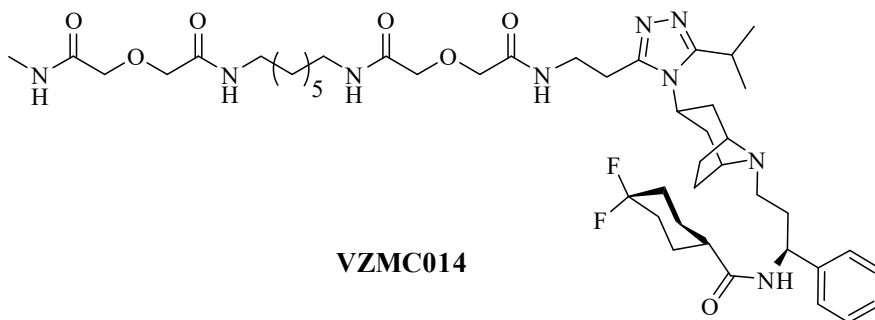
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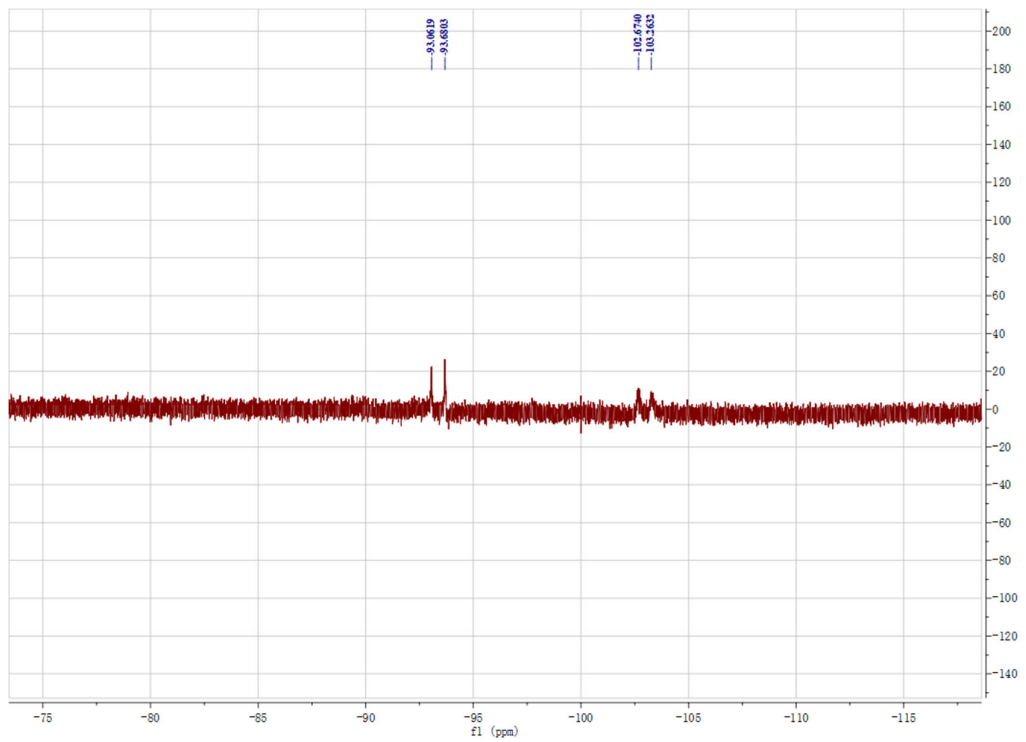
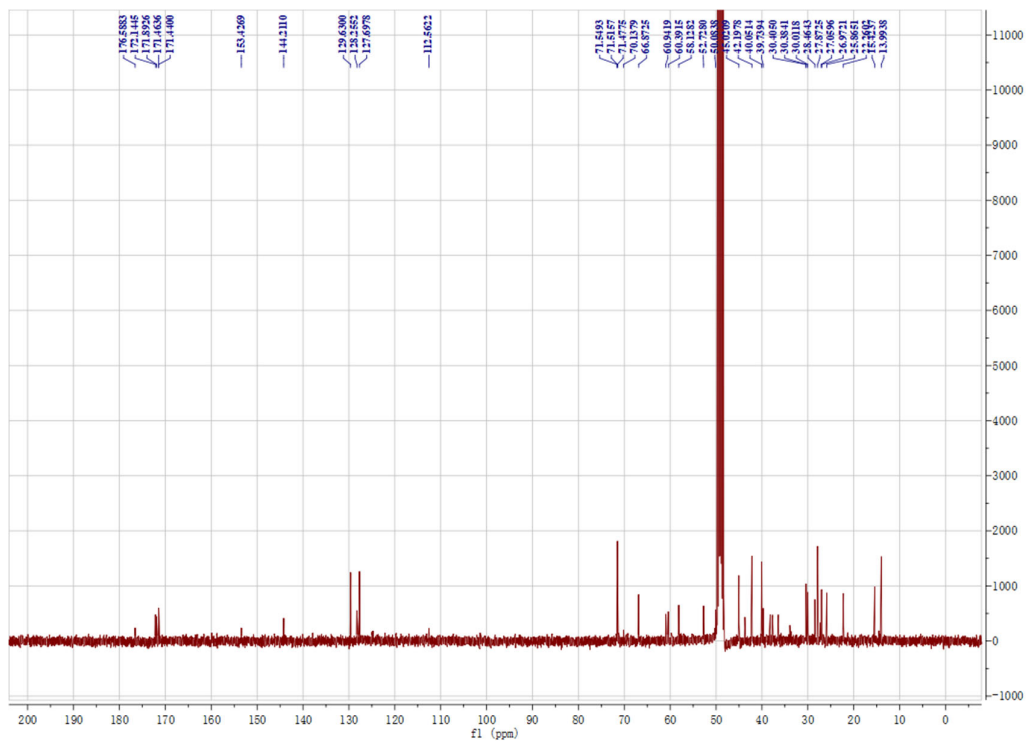


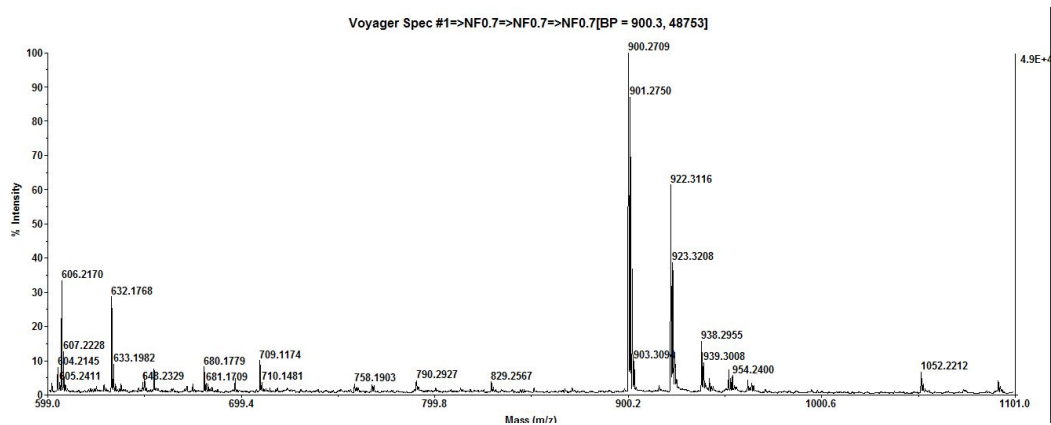




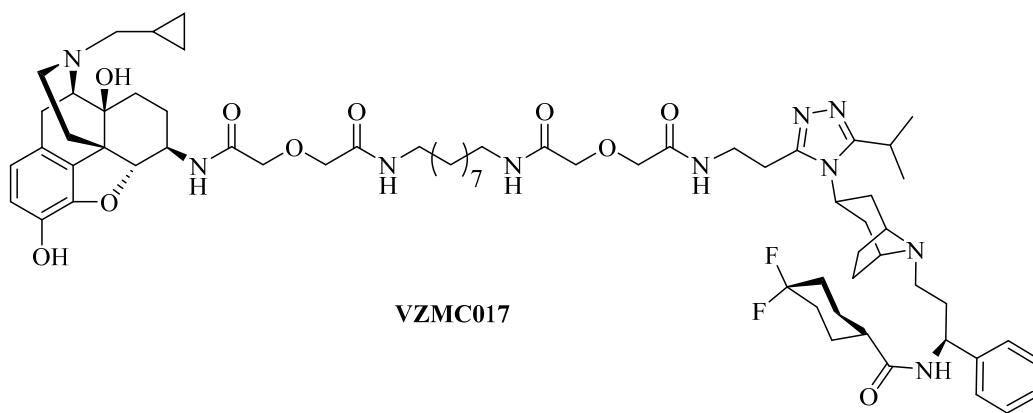
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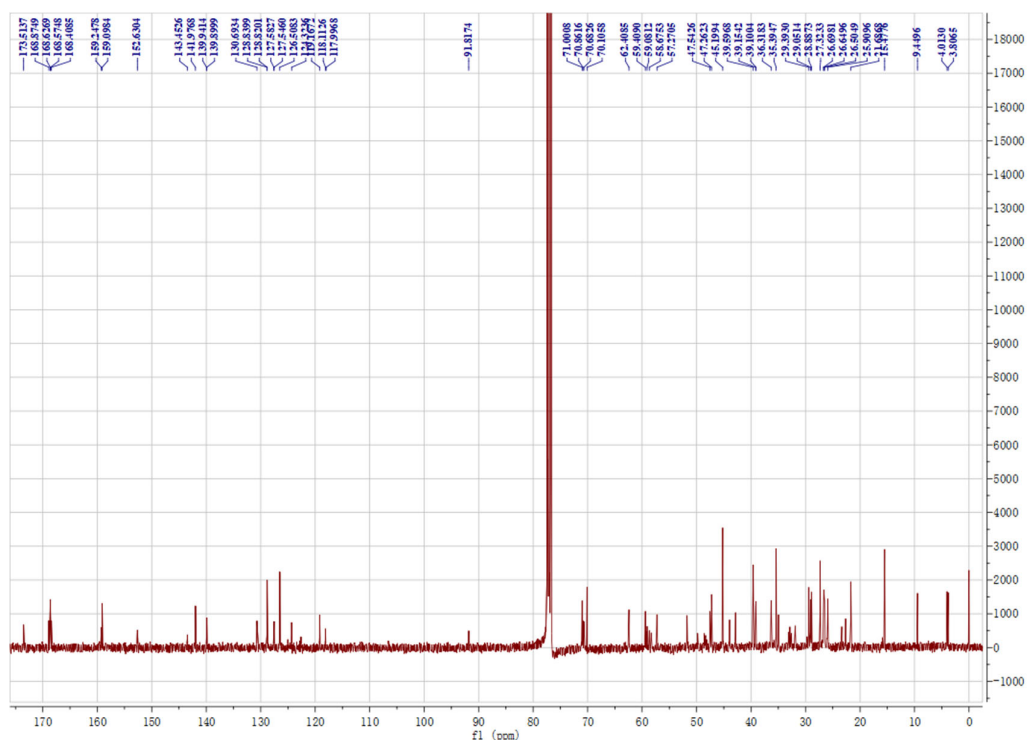
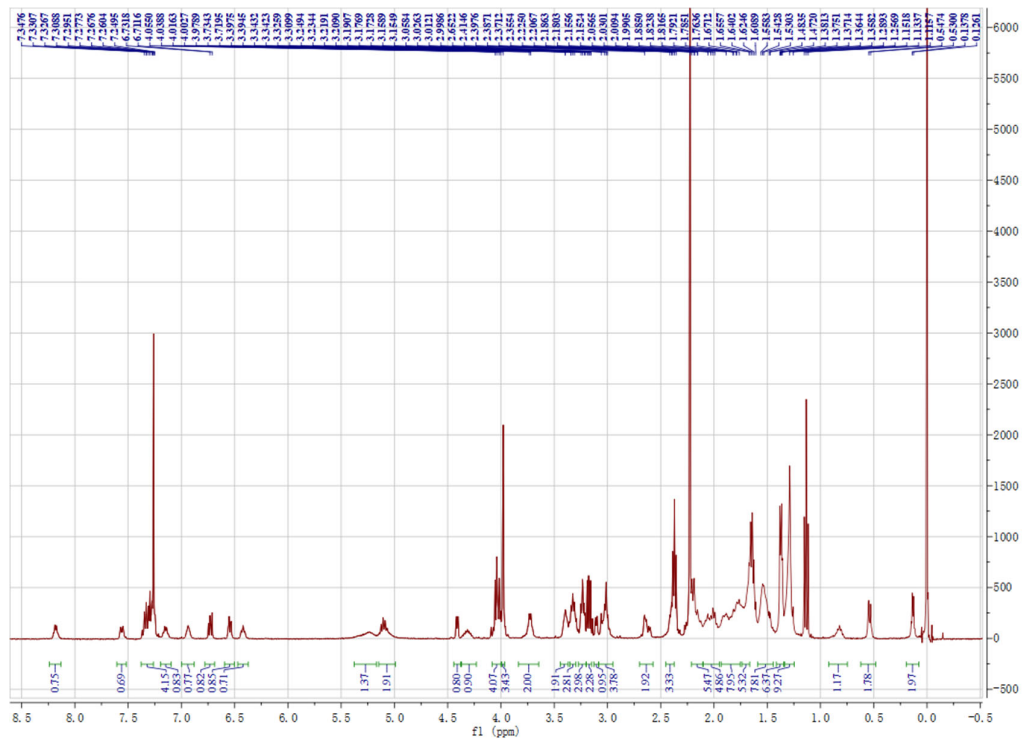


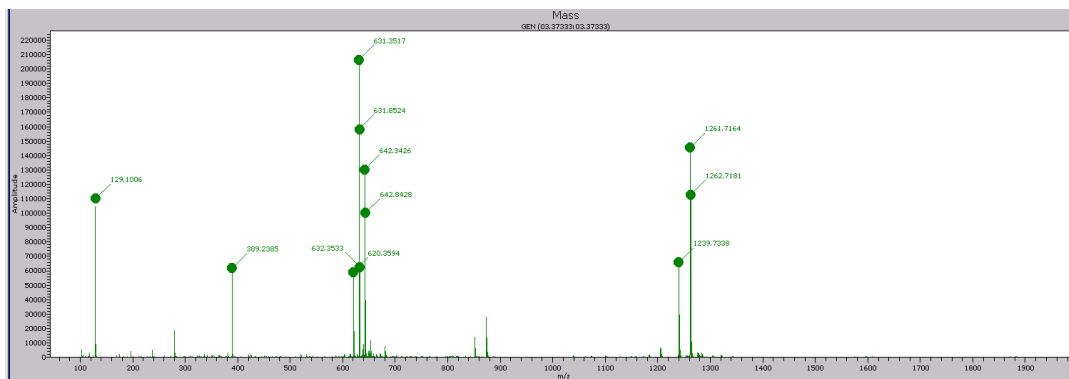
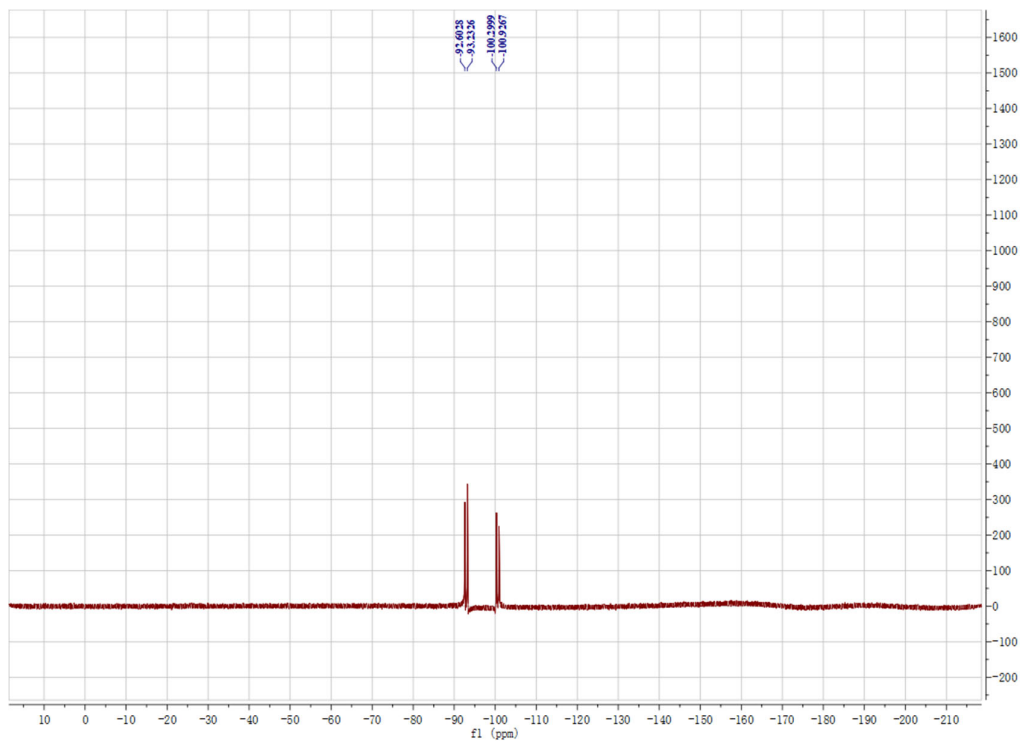




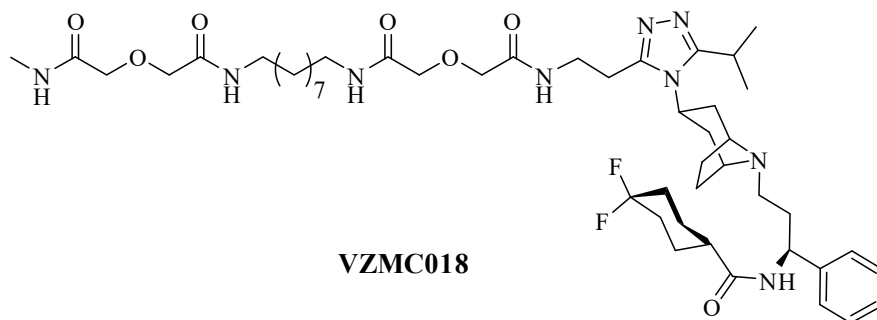
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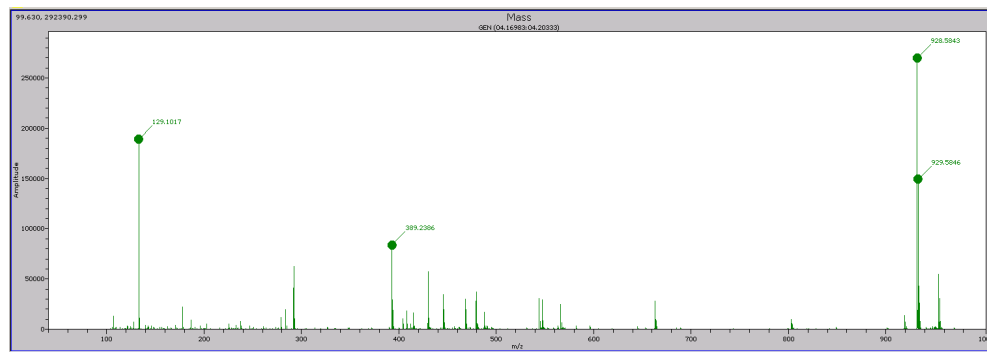
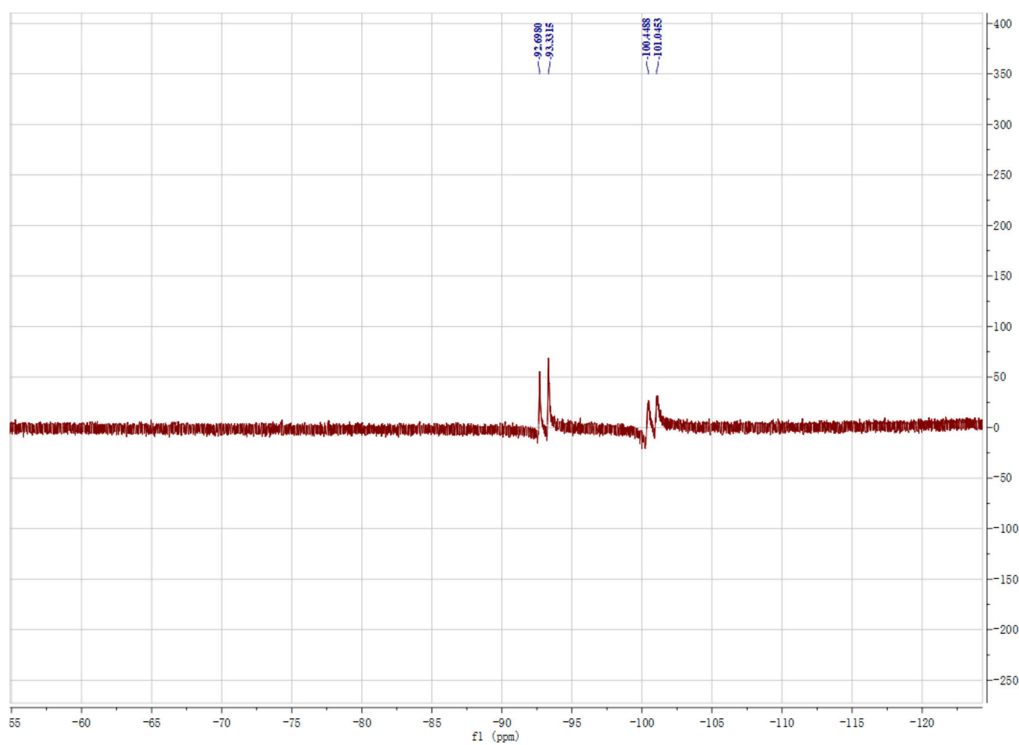




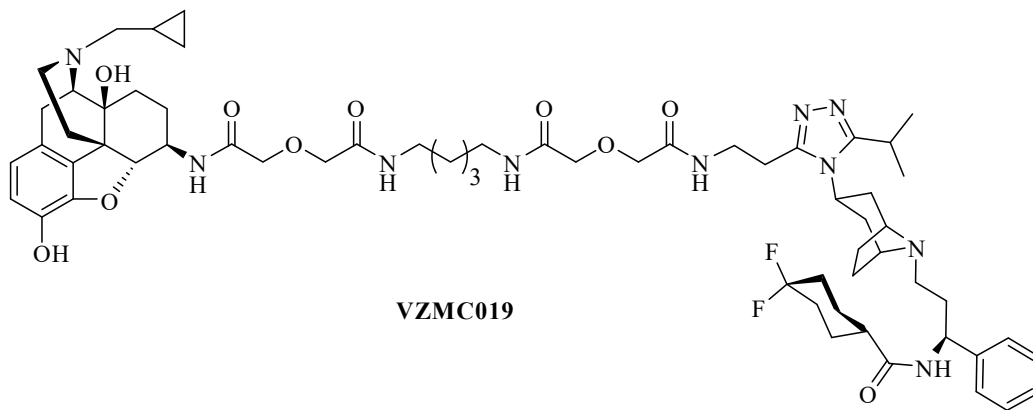


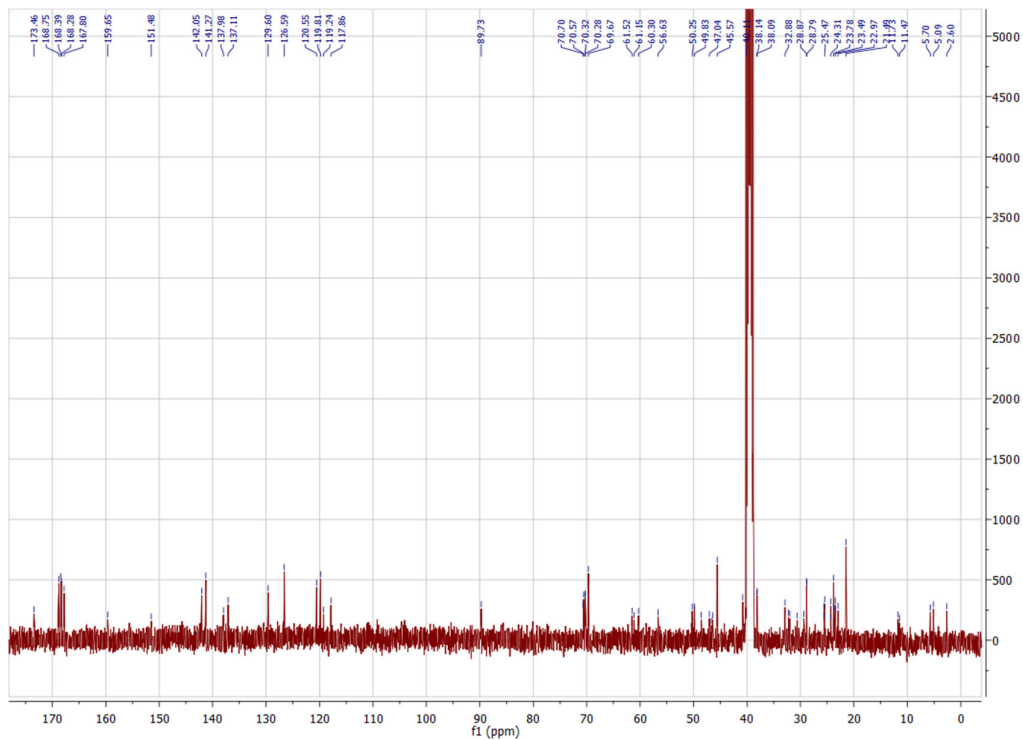
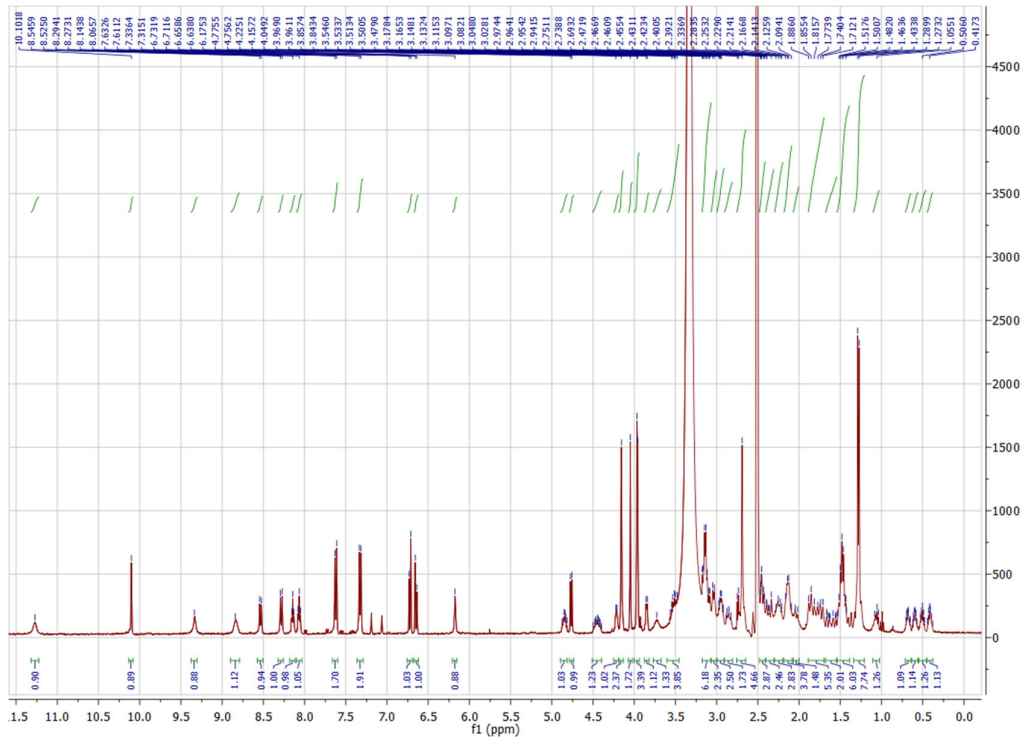
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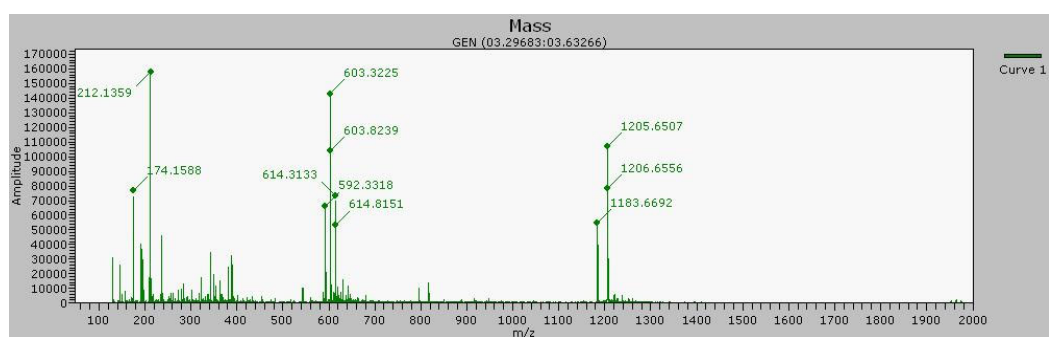
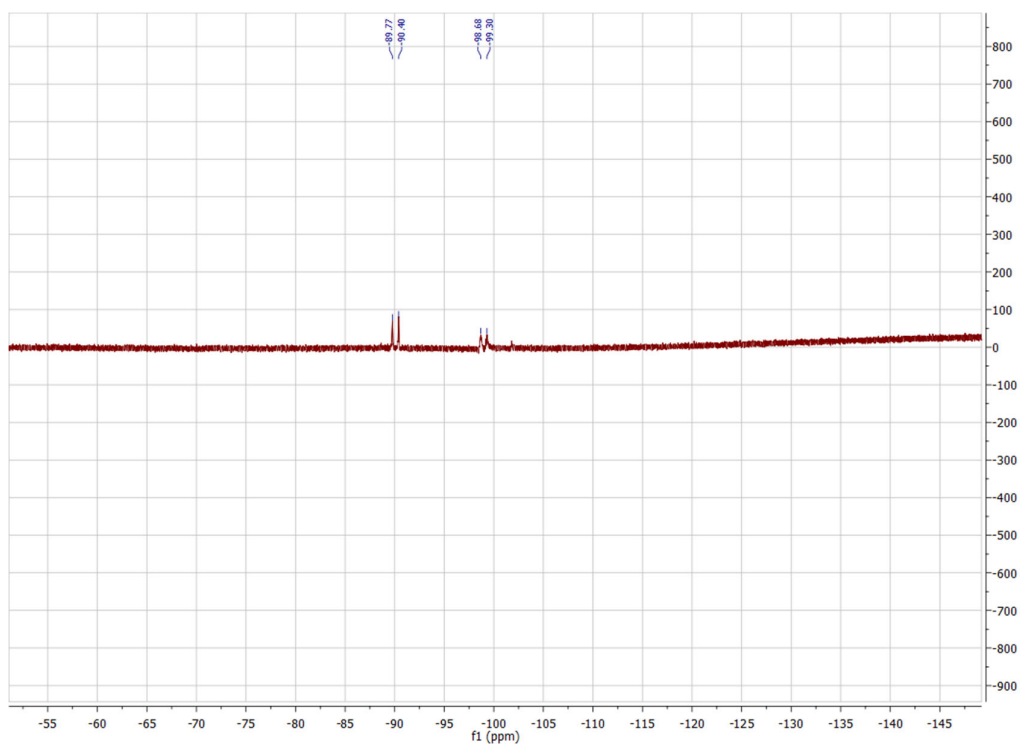




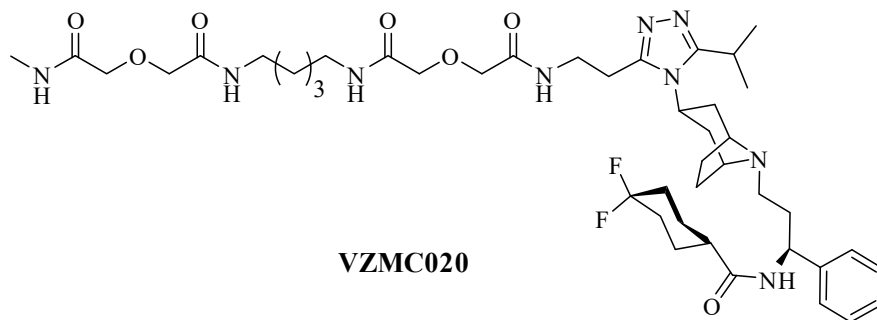
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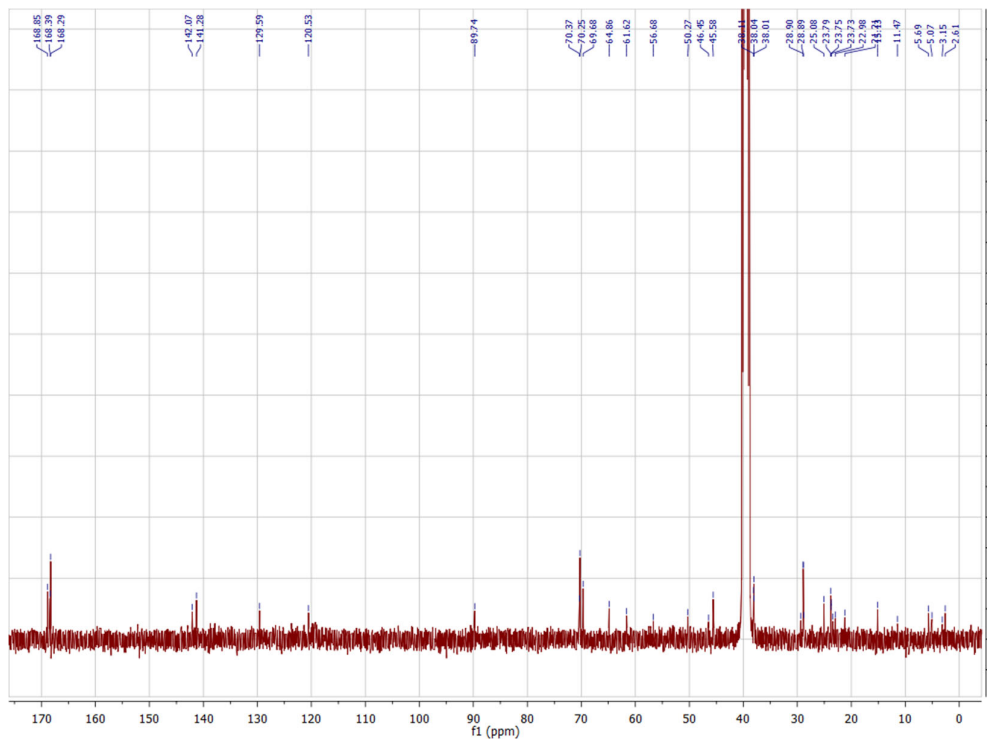
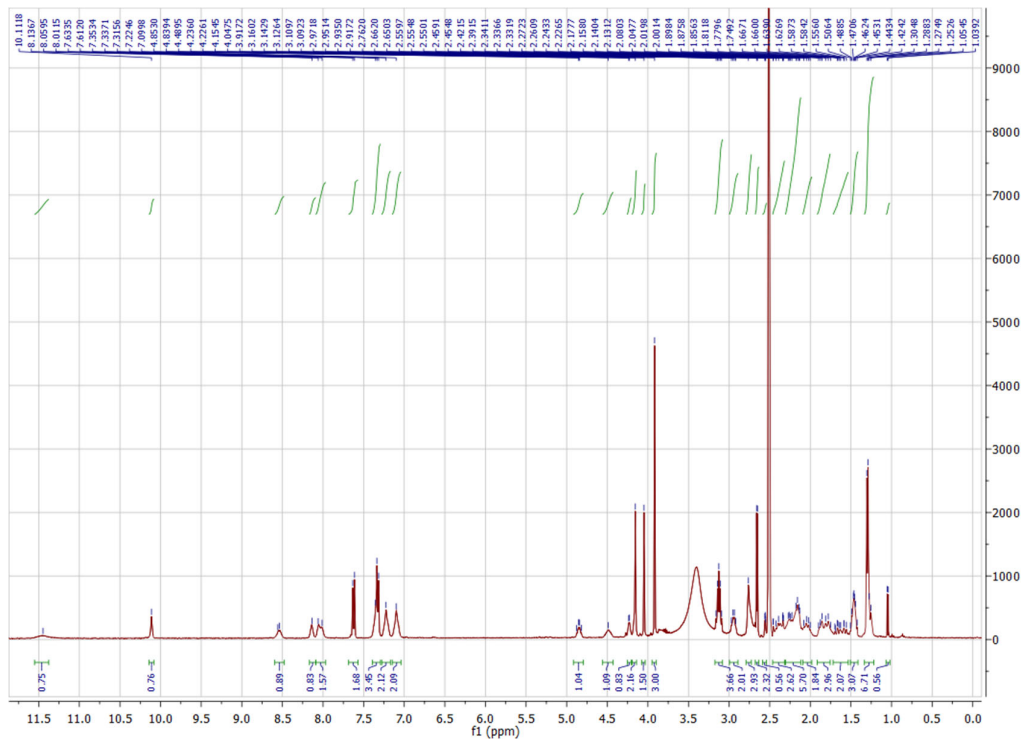


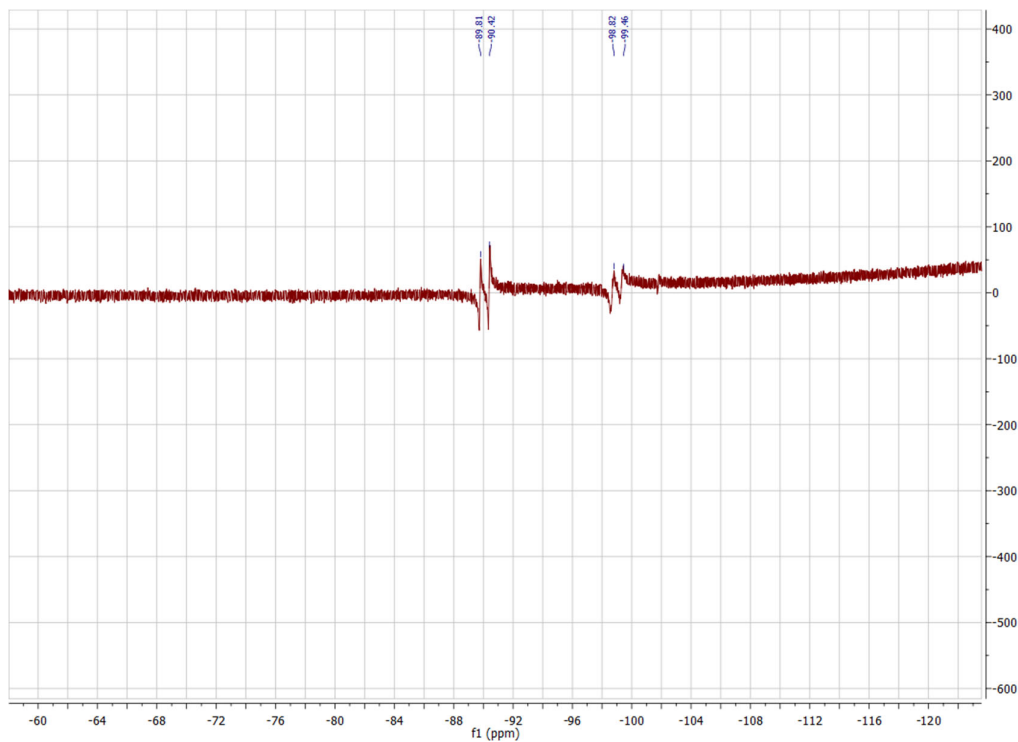




¹H NMR, ¹³C NMR, and ¹⁹F NMR spectra for **VZMC020**

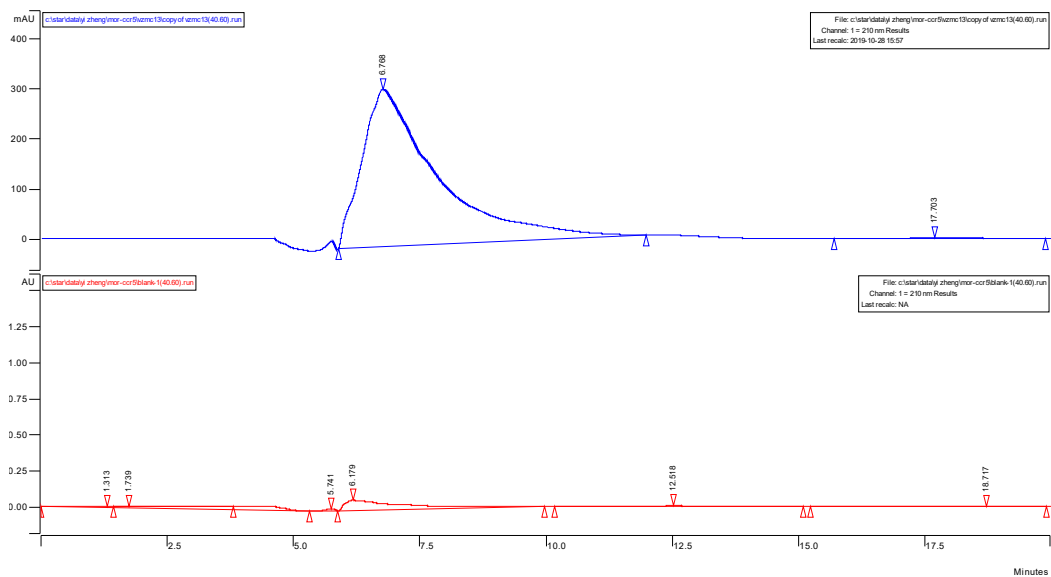




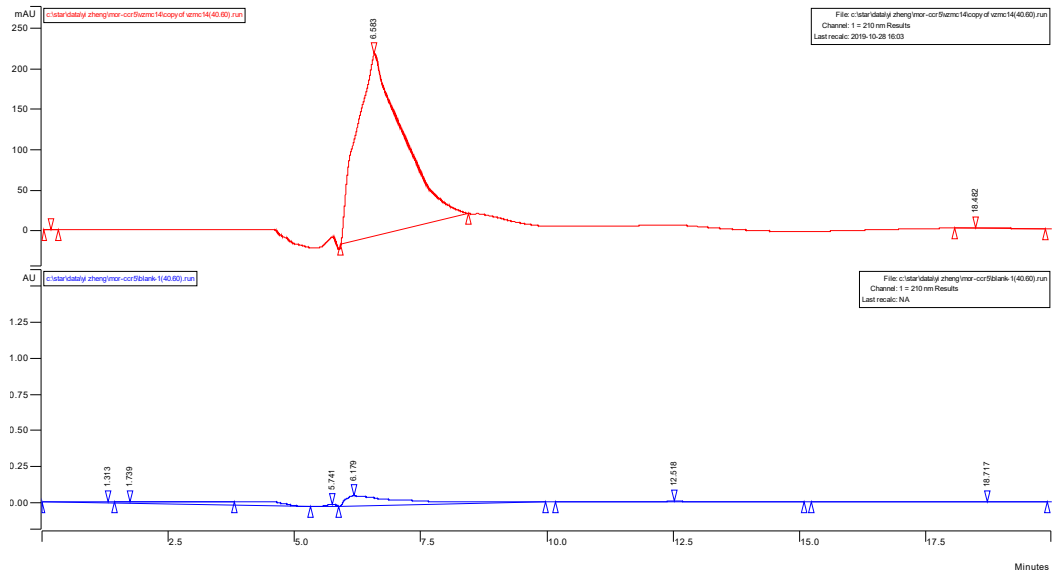


HPLC analysis

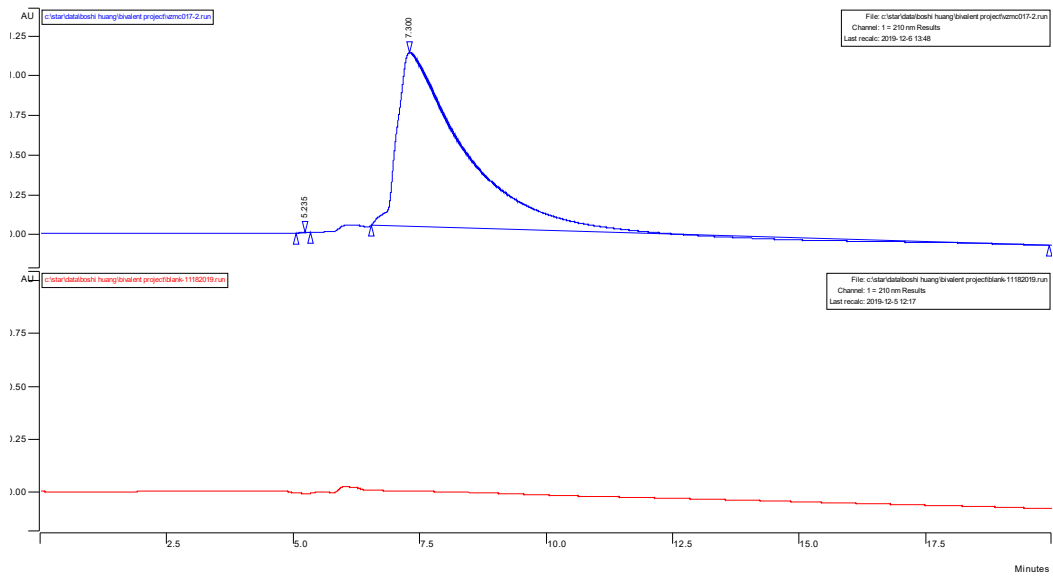
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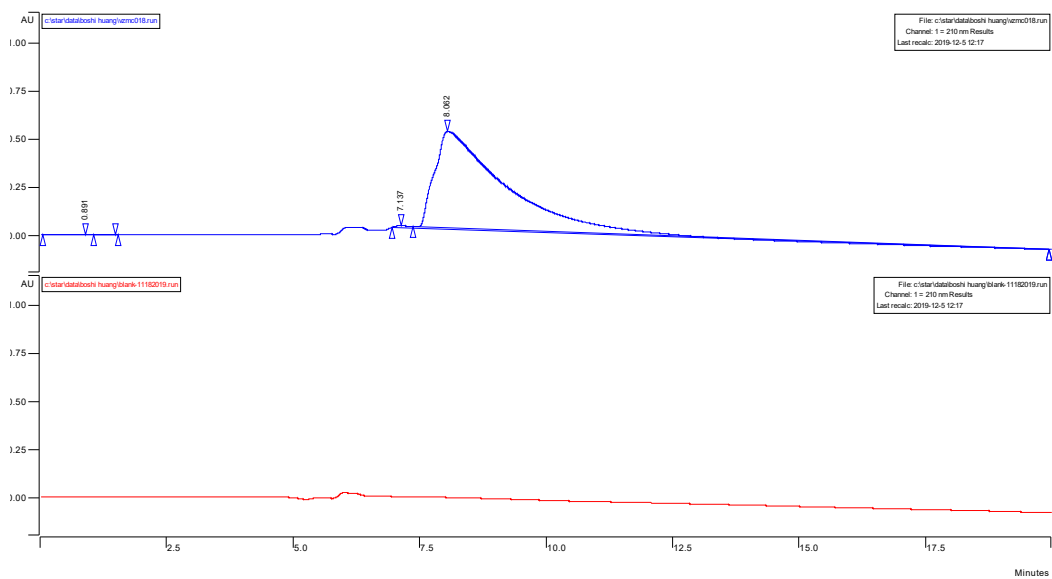
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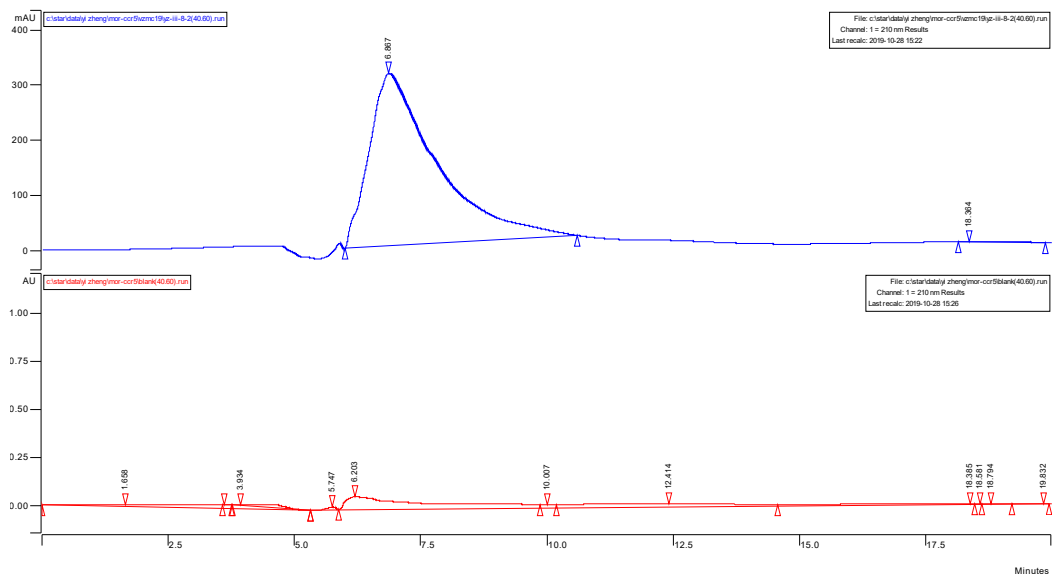
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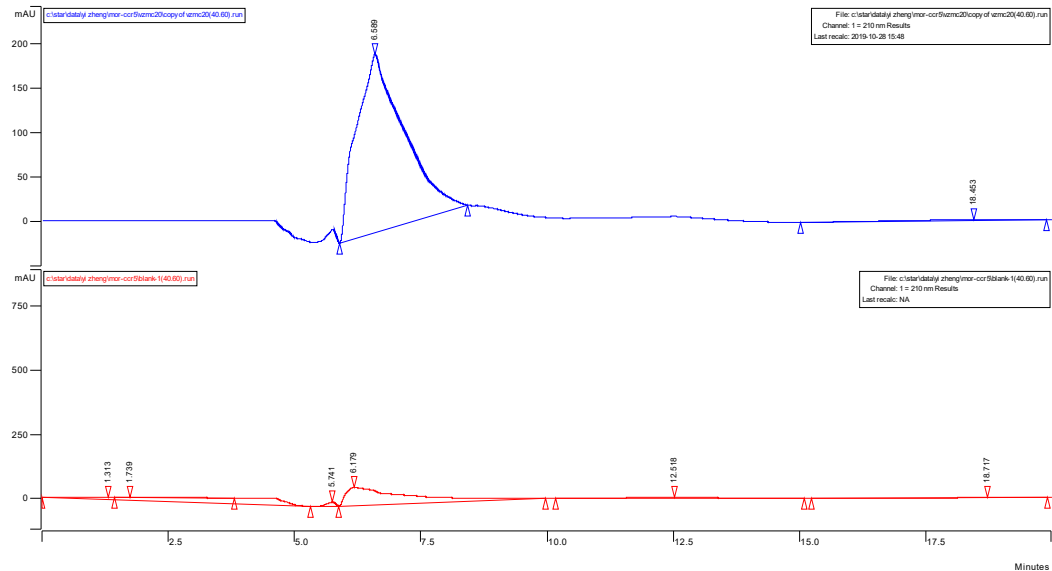
VZMC018



VZMC019



VZMC020



References

1. Yuan, Y.; Arnatt, C. K.; El-Hage, N.; Dever, S. M.; Jacob, J. C.; Selley, D. E.; Hauser, K. F.; Zhang, Y., A bivalent ligand targeting the putative mu opioid receptor and chemokine receptor CCR5 heterodimer: binding affinity versus functional activities. *MedChemComm* **2013**, *4* (5), 847-851.