

Supplementary Information

Synthesis, structural and behavioral studies of D2AAK5, D2AAK6 and D2AAK7 as serotonin 5-HT_{1A} and 5-HT_{2A} receptor ligands

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Supplementary Table S1 and S2.

Supplementary Figure S1- S4.

¹H and ¹³C NMR, HRMS and IR spectra of the investigated compounds.

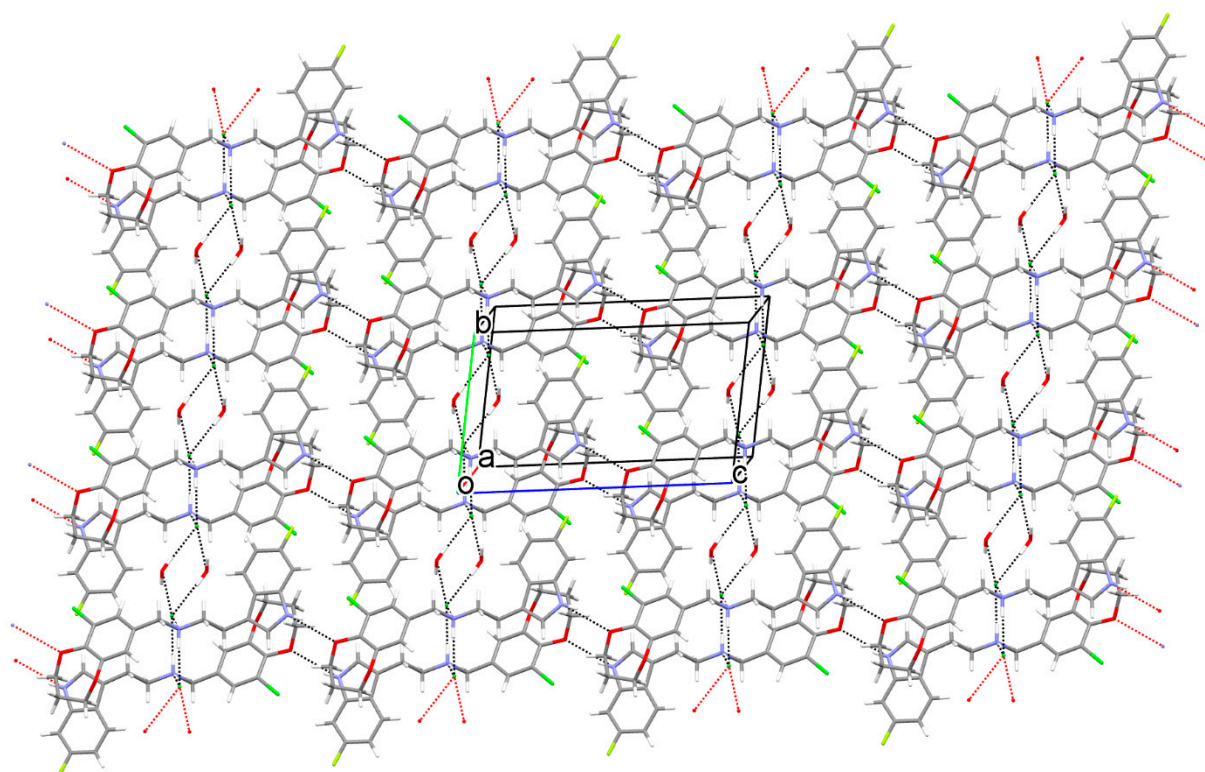
Table S1. Interatomic distances and selected bond angles for D2AAK5.

<i>Bond lengths (Å)</i>			
C1-C2	1.364(4)	C11-C12	1.500(3)
C1-C8	1.438(4)	C12-C13	1.384(4)
C1-C9	1.501(3)	C12-C20	1.394(4)
C2-N1	1.379(3)	C13-C14	1.387(3)
C3-N1	1.376(4)	C14-C15	1.389(4)
C3-C4	1.398(4)	C14-Cl1	1.732(3)
C3-C8	1.419(3)	C15-O1	1.373(3)
C4-C5	1.360(4)	C15-C19	1.404(4)
C5-F1	1.367(3)	C16-O1	1.450(3)
C5-C6	1.386(4)	C16-C17	1.503(4)
C6-C7	1.382(4)	C17-C18	1.515(4)
C7-C8	1.395(4)	C18-O2	1.449(3)
C9-C10	1.522(4)	C19-O2	1.372(3)
C10-N2	1.495(3)	C19-C20	1.388(3)
C11-N2	1.492(3)		
<i>Bond angles (°)</i>			
C2-C1-C8	107.3(2)	N2-C10-C9	112.0(2)
C1-C2-N1	109.6(3)	N2-C11-C12	113.4(2)
C3-N1-C2	109.1(2)	C11-N2-C10	111.9(2)
N1-C3-C8	107.6(2)	C15-C14-Cl1	119.1(2)
C4-C5-F1	118.1(2)	C13-C14-Cl1	118.9(2)
F1-C5-C6	117.0(3)	C15-O1-C16	116.0(2)
C3-C8-C1	106.4(2)	C19-O2-C18	117.7(2)
C1-C9-C10	110.6(2)		
<i>Torsion angles (°)</i>			
O1-C16-C17-C18	68.1(3)	C17-C16-O1-C15	-78.9(3)
C16-C17-C18-O2	-66.6(3)	C15-C19-O2-C18	-59.2(4)
O1-C15-C19-O2	-1.2(4)	C17-C18-O2-C19	76.8(3)
C19-C15-O1-C16	60.9(4)		

Table S2. Hydrogen bonding and C-H...Cg interactions geometry.

<i>Hydrogen bond [\AA, $^\circ$]</i>				
D-H...A	d(D-H)	d(H...A)	d(D...A)	\angle DHA
N1-H1N...O1 ⁱ	0.82(3)	2.26(3)	3.034(3)	157(3)
O3-H1W...Cl2	0.89	2.35	3.209(2)	164
O3-H2W...Cl2 ⁱⁱ	0.92	2.33	3.255(2)	176
N2-H2N...Cl2	0.98(3)	2.26(3)	3.217(3)	169(3)
N2-H3N...Cl2 ⁱⁱⁱ	0.97(3)	2.17(3)	3.130(3)	175(2)
C2-H2...F1 ^{iv}	0.95	2.40	3.338(3)	168
C10-H10B...O2 ^v	0.99	2.55	3.424(3)	147
C11-H11B...O3	0.99	2.57	3.337(4)	134
<i>C-H...Cg interactions [\AA, $^\circ$]</i>				
C-H...Cg	d(D-H)	d(H...Cg)	d(C...Cg)	\angle CHCg
C16-H16B...Cg1 ⁱⁱ	0.99	2.43	3.399(3)	165
C16-H16B...Cg5 ⁱⁱ	0.99	2.78	3.651(3)	147

Symmetry codes: (i) $x, y, z-1$; (ii) $-x+1, -y+1, -z$; (iii) $-x+1, -y+2, -z$; (iv) $x, y+1, z$; (v) $-x+2, -y+2, -z$; Cg1 and Cg5 are the centroids of the N1/C2/C1/C8/C3 and N1/C2/C1/C8/C7/C6/C5/C4/C3 rings, respectively.

**Figure S1.** Packing diagrams of D2AAK5 showing formation sheets viewed along the *a*-axis. Hydrogen bonds are indicated by dotted lines.

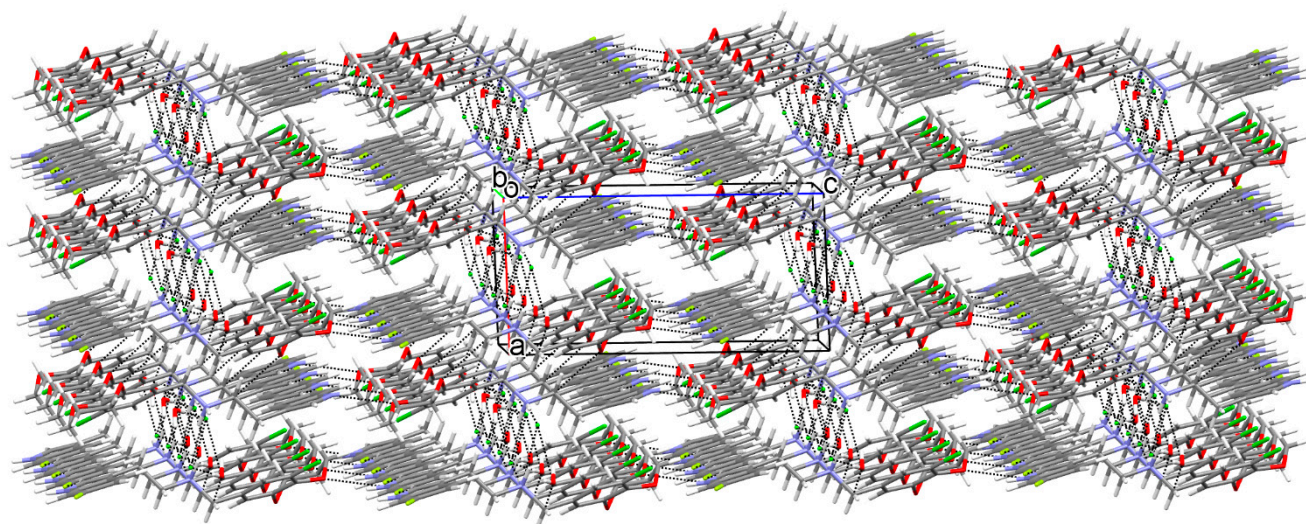


Figure S2. The 3D supramolecular structure of the D2AAK5 constructed by the connection of adjacent sheets through C10–H1B···O2 hydrogen bonds. The hydrogen bonds shown as dotted lines. Hanging contact are omitted for clarity.

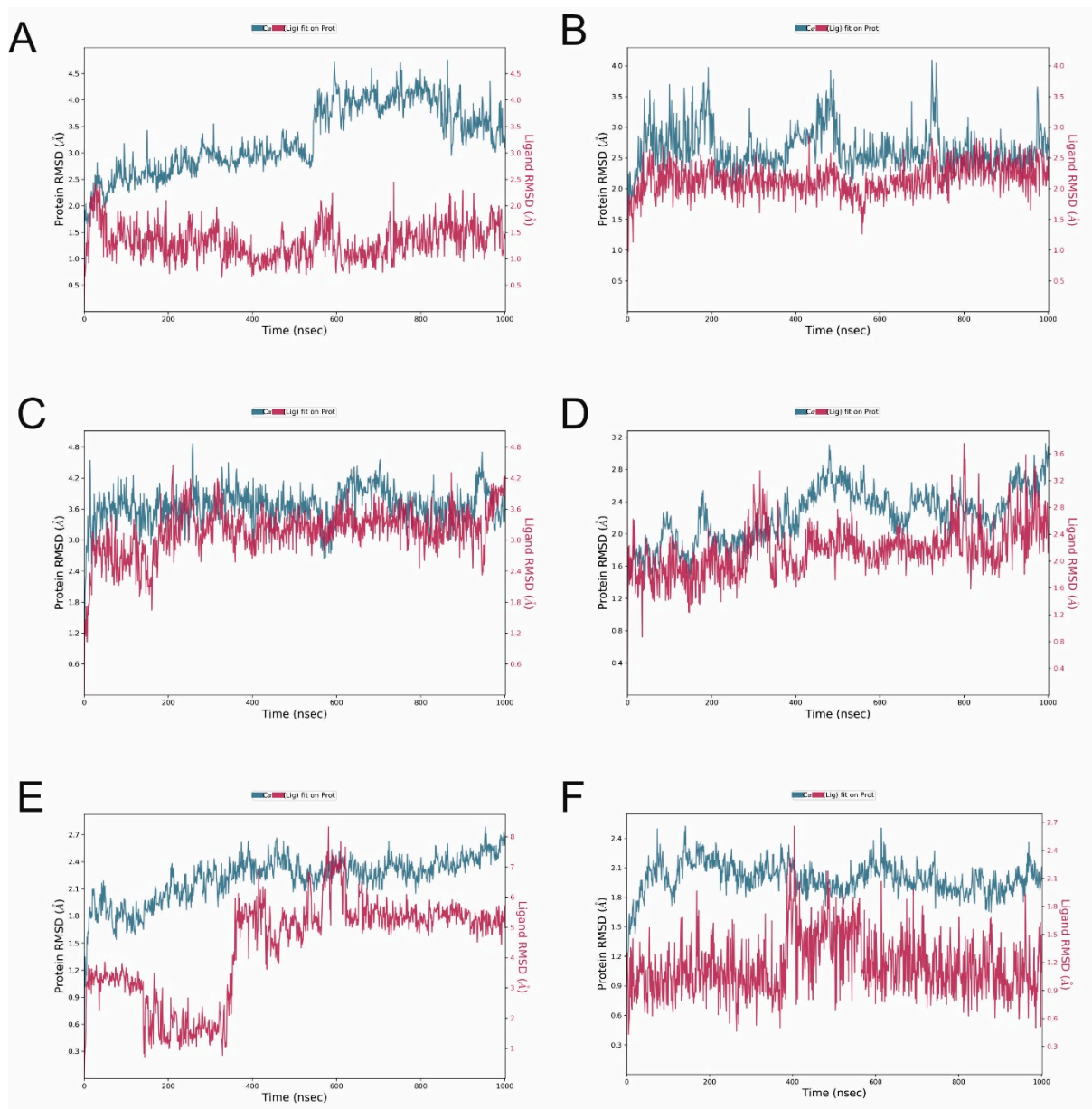


Figure S3. Ligand (purple) and protein (blue) C α RMSD during 1 μ s MD simulations for D2AAK5-5HT_{1A} (A), D2AAK6-5HT_{1A} (B), D2AAK7-5HT_{1A} (C), D2AAK5-5-HT_{2A} (D), D2AAK6-5HT_{2A} (E) and D2AAK7-5HT_{2A} (F) ligand-receptor systems.

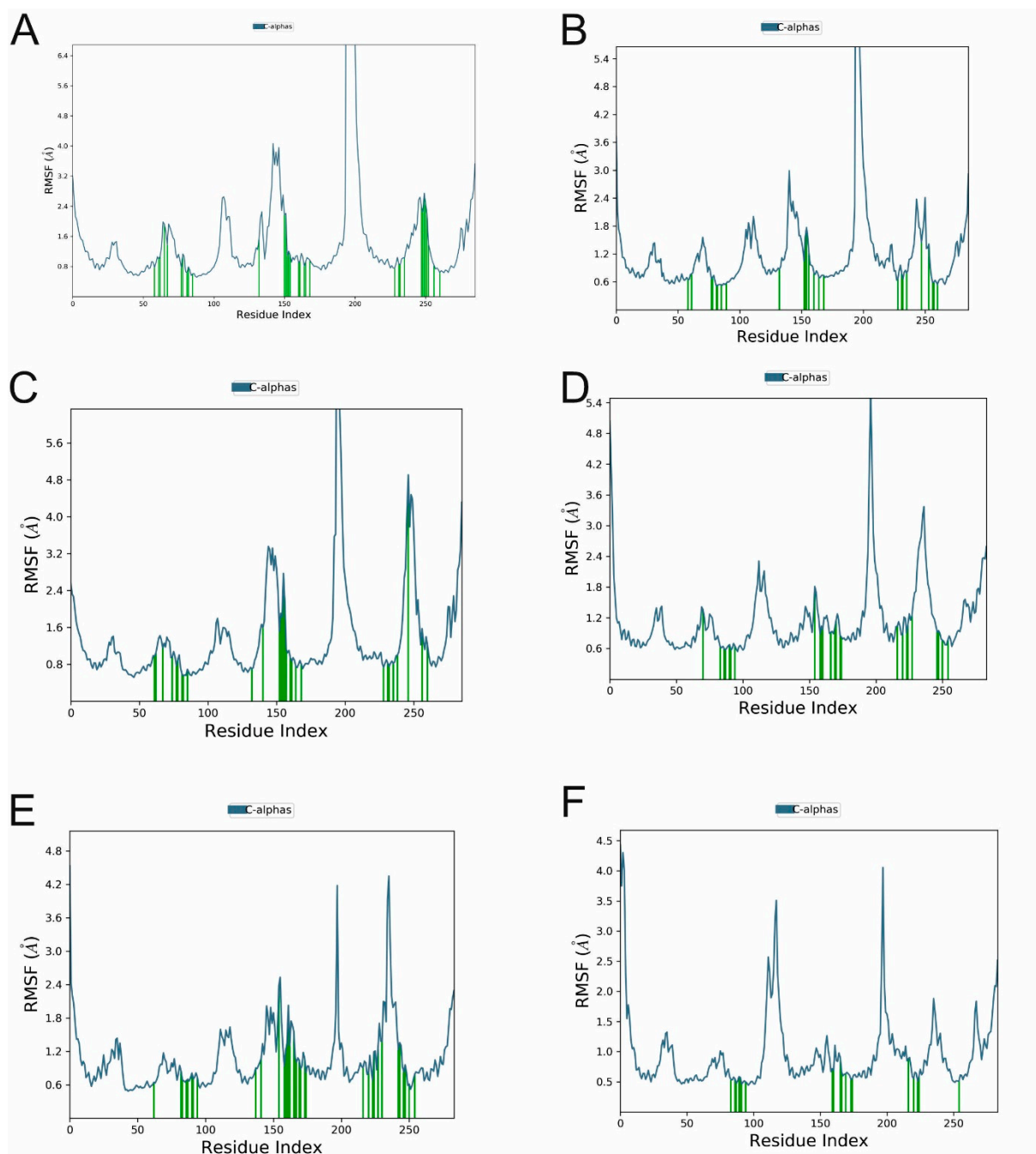
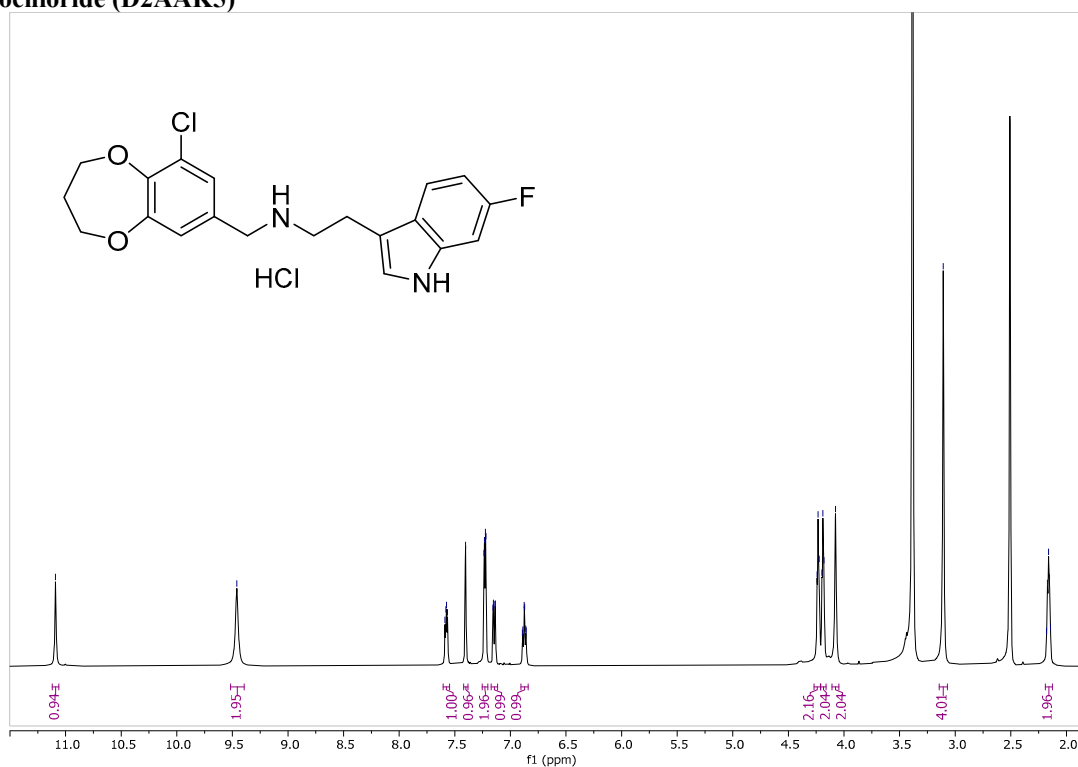
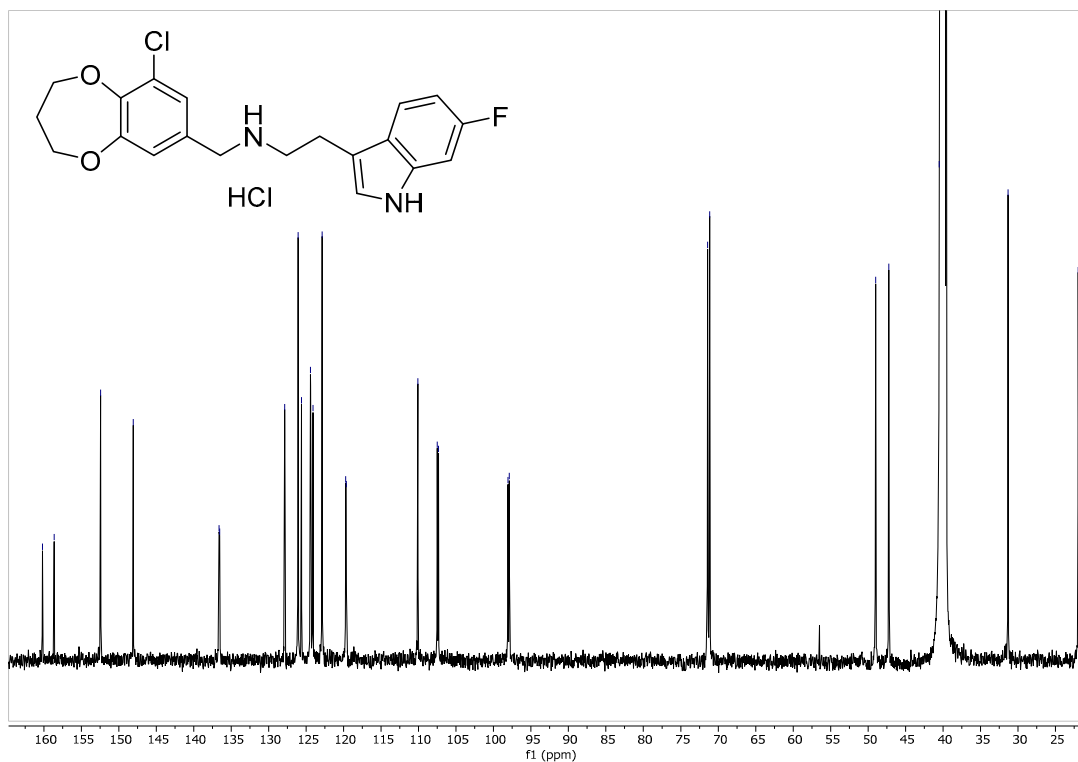


Figure S4. Protein C α RMSF during 1 μ s MD simulations for D2AAK5-5HT_{1A} (A), D2AAK6-5HT_{1A} (B), D2AAK7-5HT_{1A} (C), D2AAK5-5-HT_{2A} (D), D2AAK6-5HT_{2A} (E) and D2AAK7-5HT_{2A} (F) ligand-receptor systems. Protein residues that interact with the ligand are marked with green-colored vertical bars.

N-((9-chloro-3,4-dihydro-2H-benzo[b][1,4]dioxepin-7-yl)methyl)-2-(6-fluoro-1H-indol-3-yl)ethan-1-amine hydrochloride (D2AAK5)

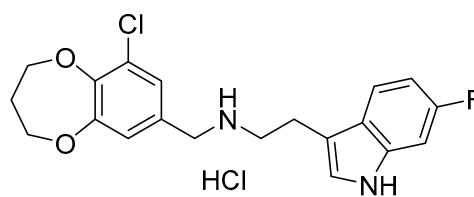
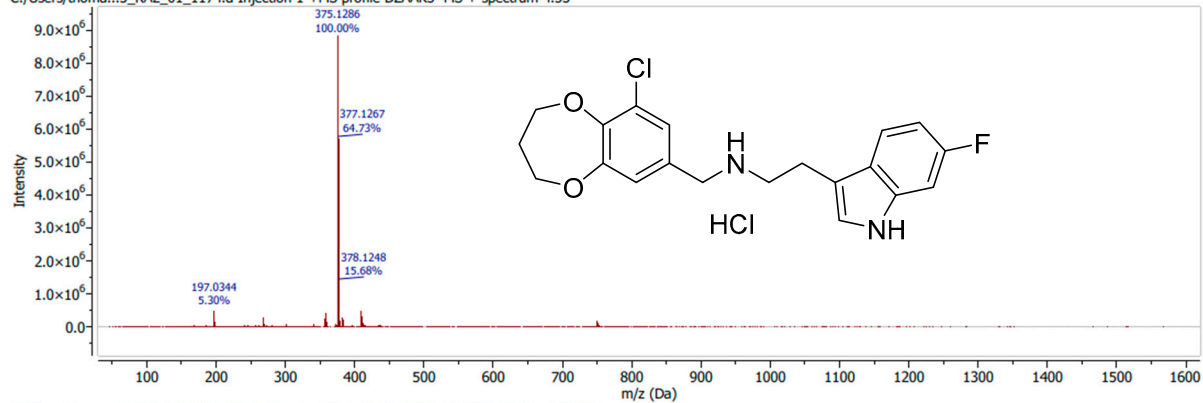


^1H NMR (600 MHz, DMSO) δ 11.09 (s, 1H), 9.46 (s, 2H), 7.58 (dd, $J = 8.7, 5.3$ Hz, 1H), 7.40 (s, 1H), 7.23 (dd, $J = 7.3, 2.3$ Hz, 2H), 7.15 (dd, $J = 10.1, 2.4$ Hz, 1H), 6.87 (td, $J = 9.2, 2.3$ Hz, 1H), 4.23 (t, $J = 5.5$ Hz, 2H), 4.19 (t, $J = 5.4$ Hz, 2H), 4.08 (s, 2H), 3.11 (s, 4H), 2.16 (p, $J = 5.5$ Hz, 2H).

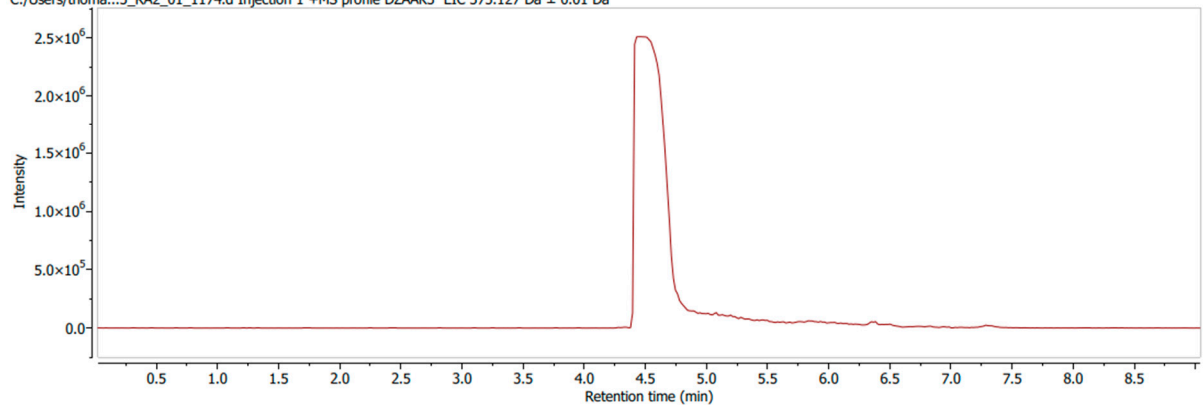


^{13}C NMR (151 MHz, DMSO) δ 159.41 (d, $J = 234.2$ Hz), 152.43, 148.07, 136.59 (d, $J = 12.7$ Hz), 127.86, 126.08, 125.62, 124.42, 124.09, 122.86, 119.68 (d, $J = 10.2$ Hz), 110.09, 107.42 (d, $J = 24.4$ Hz), 97.97 (d, $J = 25.5$ Hz), 71.42, 71.15, 49.00, 47.23, 31.33, 21.96.

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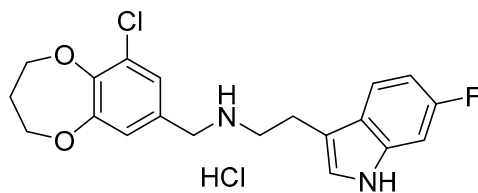
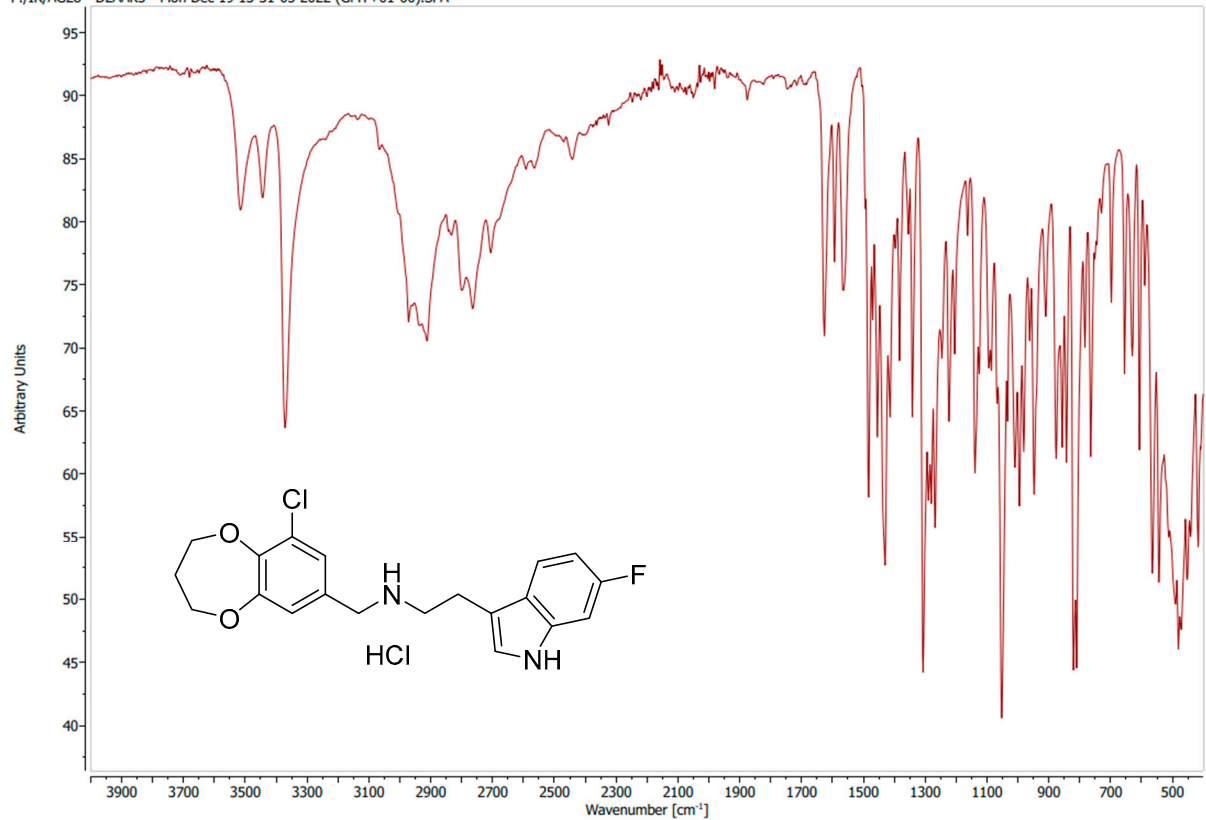


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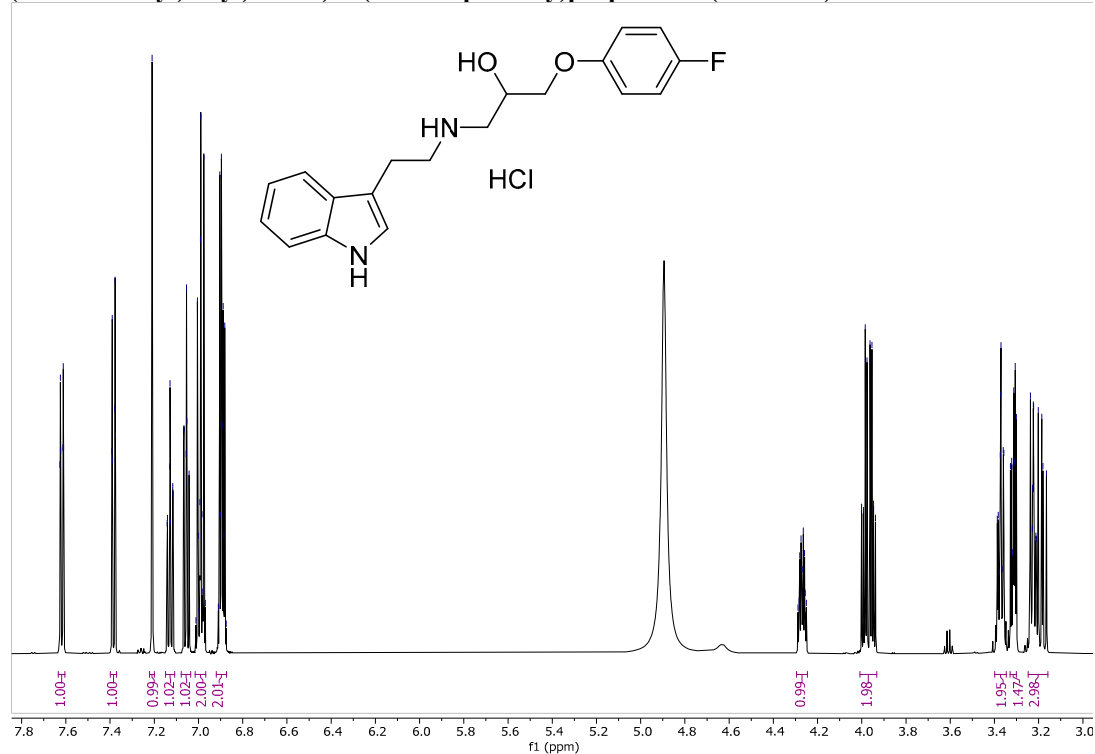


HRMS (ESI) calc. (M+H)⁺ 375.1270 exp. (M+H)⁺ 375.1286

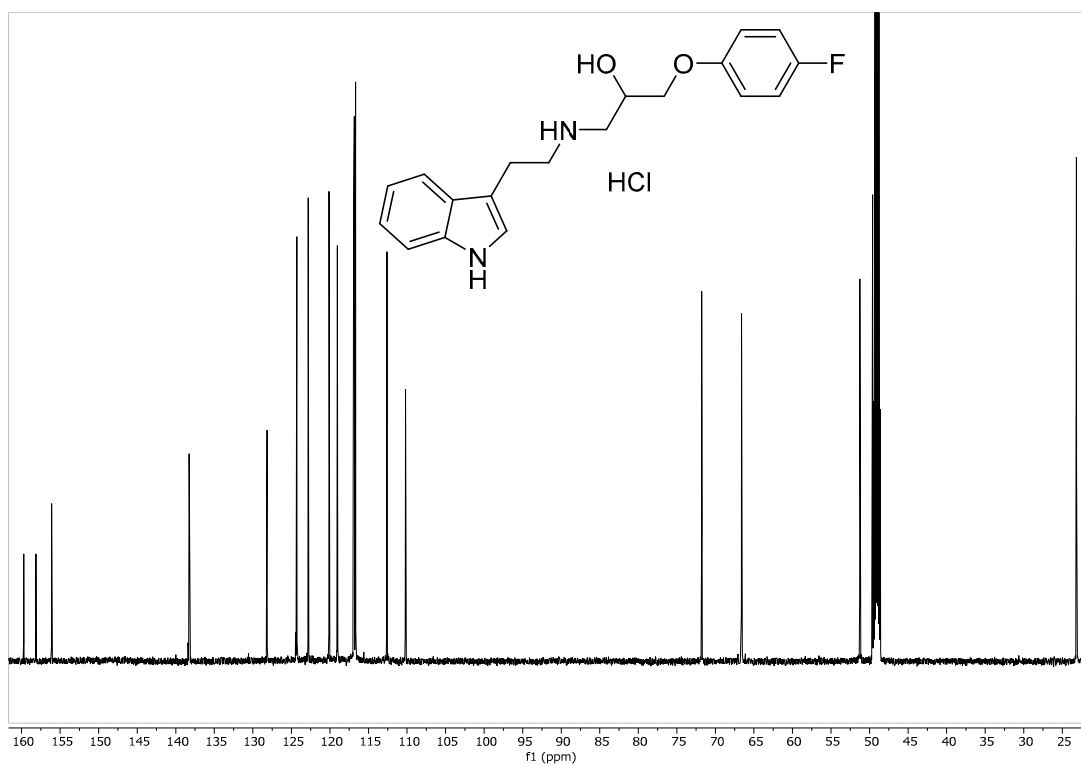
F:/IR/AG28 - D2AAK5 - Mon Dec 19 13-31-05 2022 (GMT+01-00).SPA -



1-((2-(1H-inden-3-yl)ethyl)amino)-3-(4-fluorophenoxy)propan-2-ol (D2AAK6)

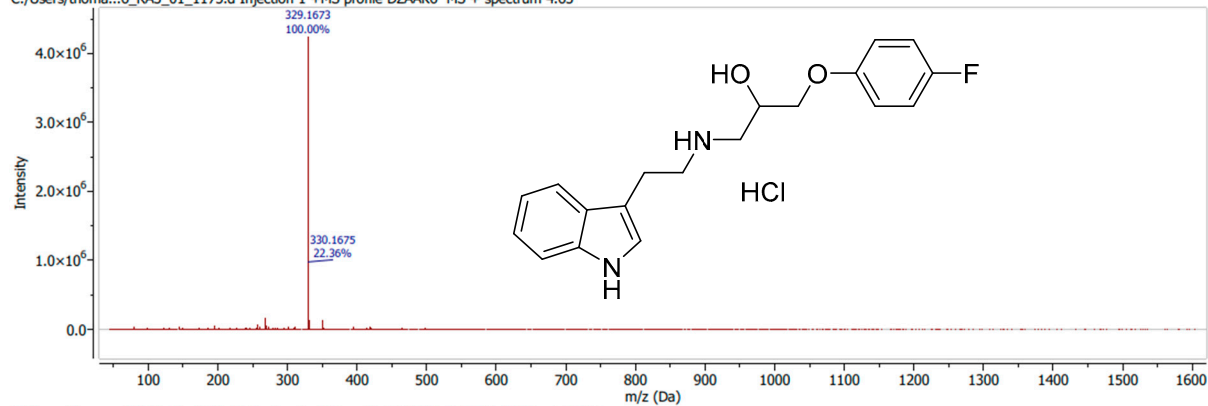


^1H NMR (600 MHz, MeOD) δ 7.62 (dt, J = 7.9, 1.0 Hz, 1H), 7.38 (dt, J = 8.2, 0.9 Hz, 1H), 7.21 (s, 1H), 7.13 (ddd, J = 8.1, 7.1, 1.1 Hz, 1H), 7.05 (ddd, J = 8.0, 7.0, 1.0 Hz, 1H), 7.02 – 6.97 (m, 2H), 6.92 – 6.87 (m, 2H), 4.27 (dtd, J = 8.3, 5.1, 3.2 Hz, 1H), 4.01 – 3.93 (m, 2H), 3.40 – 3.35 (m, 2H), 3.33 – 3.30 (m, 1H), 3.25 – 3.16 (m, 3H).

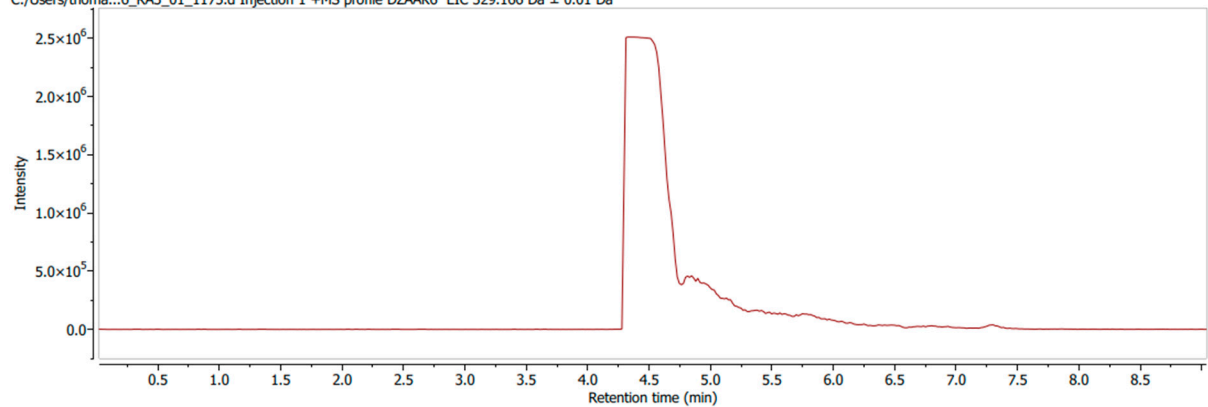


^{13}C NMR (151 MHz, MeOD) δ 159.7, 158.1, 156.1, 156.1, 138.3, 128.2, 124.3, 122.8, 120.1, 119.0, 116.9, 116.9, 116.8, 116.7, 112.6, 110.2, 71.8, 66.6, 51.3, 49.6, 23.2.

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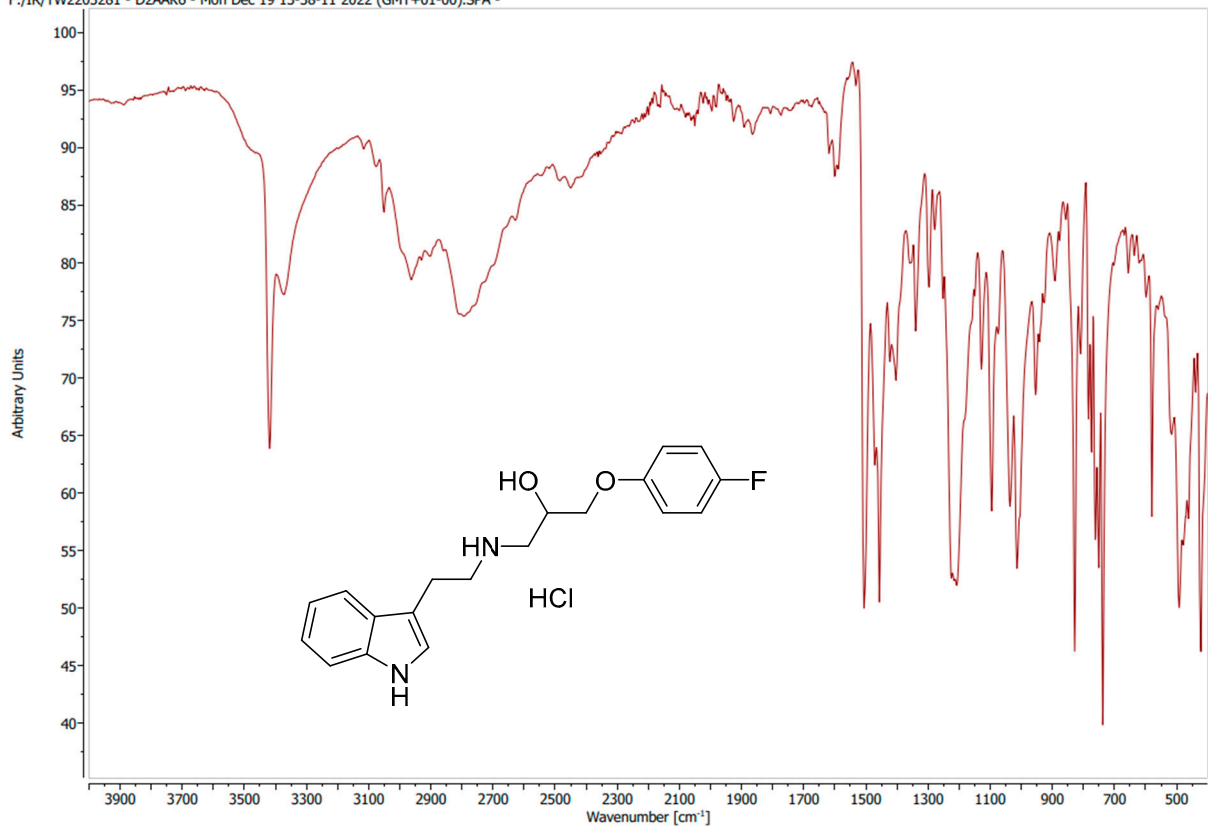


C:/Users/thoma...6_RA3_01_1175.d Injection 1 +MS profile DZAAK6 EIC 329.166 Da \pm 0.01 Da

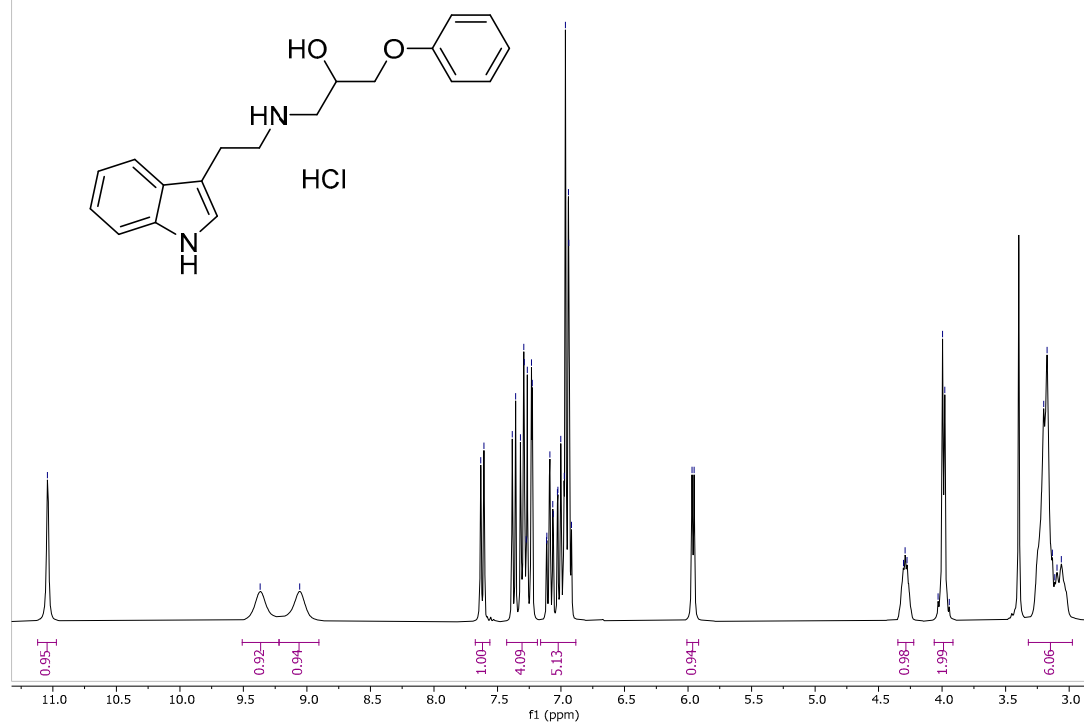


HRMS (ESI) calc. $(M+H)^+$ 329.1660 exp. $(M+H)^+$ 329.1673

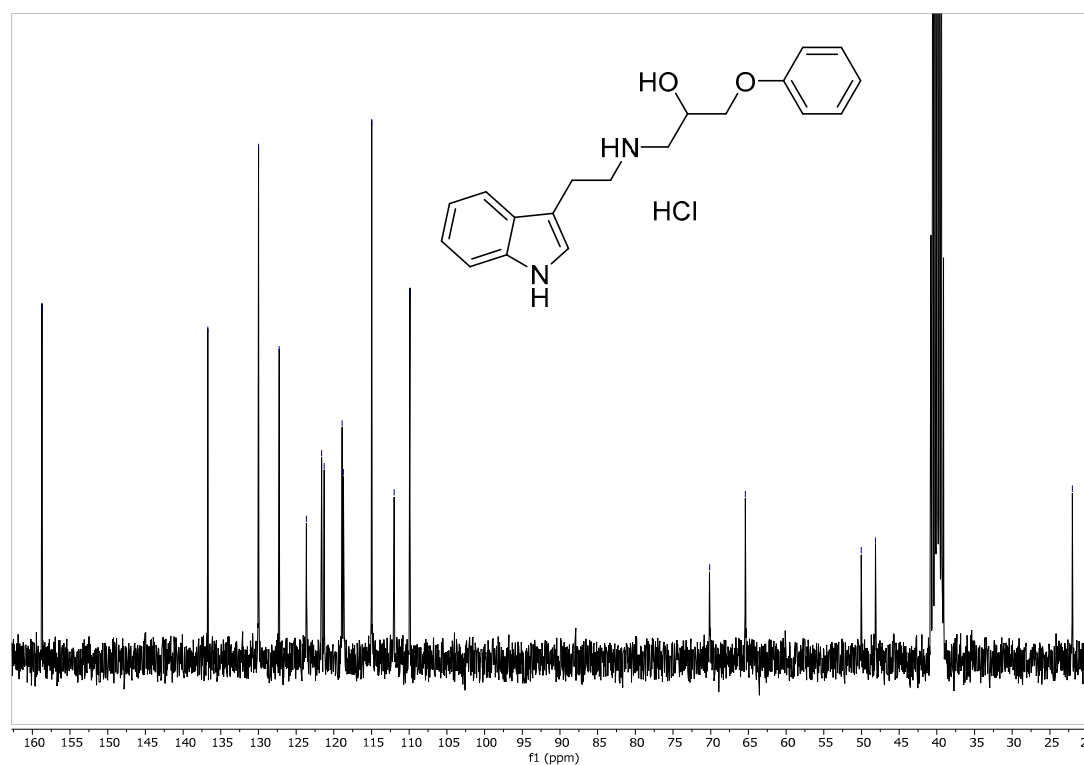
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1-((2-(1H-inden-3-yl)ethyl)amino)-3-phenoxypropan-2-ol (D2AAK7)

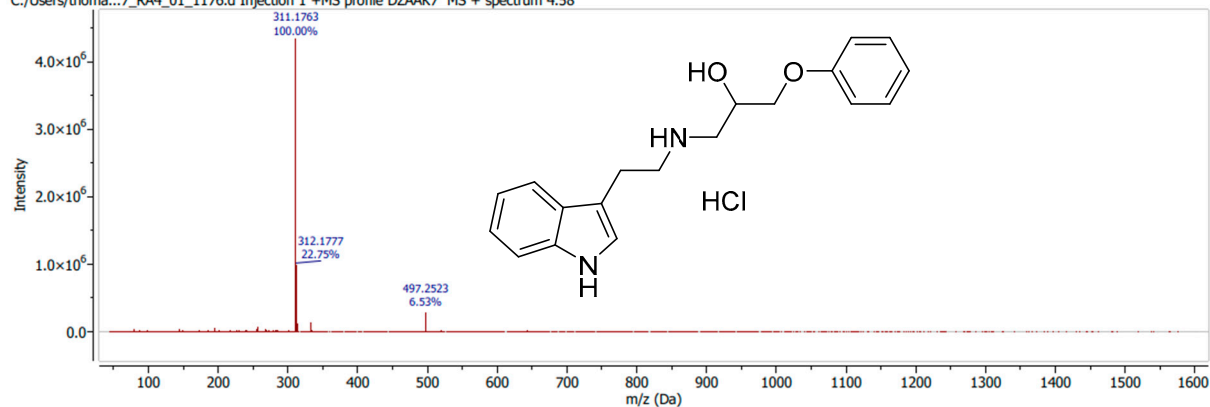


^1H NMR (300 MHz, DMSO) δ 11.04 (s, 1H), 9.37 (s, 1H), 9.06 (s, 1H), 7.62 (d, $J = 7.7$ Hz, 1H), 7.43 – 7.19 (m, 4H), 7.16 – 6.88 (m, 5H), 5.96 (d, $J = 4.9$ Hz, 1H), 4.35 – 4.22 (m, 1H), 4.06 – 3.91 (m, 2H), 3.32 – 2.97 (m, 6H).

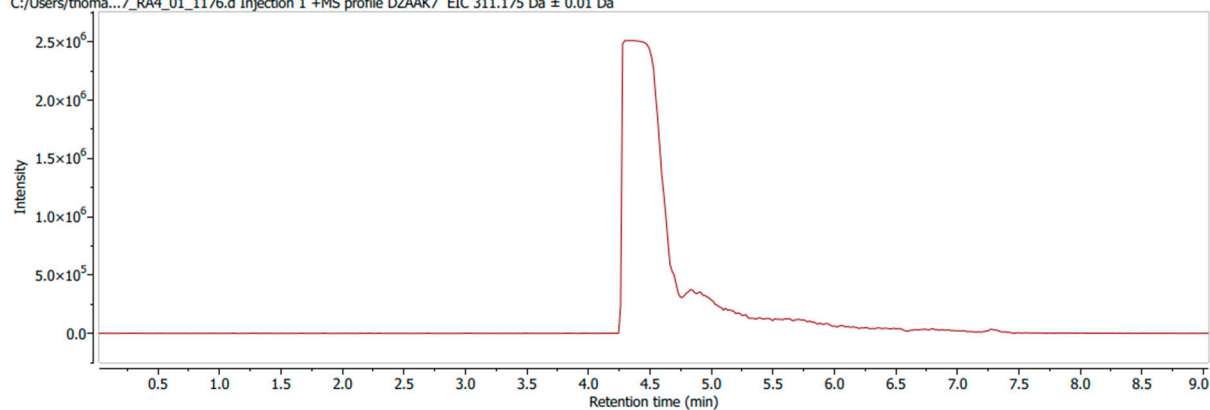


^{13}C NMR (75 MHz, DMSO) δ 158.7, 136.7, 130.0, 127.2, 123.6, 121.6, 121.3, 118.9, 118.7, 115.0, 112.0, 109.9, 70.2, 65.4, 50.0, 48.1, 22.0.

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HRMS (ESI) calc. (M+H)⁺ 311.1754 exp. (M+H)⁺ 311.1763

F:/IR/DZAAK7 - Mon Dec 19 13-42-50 2022 (GMT+01-00).SPA -

