

SUPPORTING INFORMATION FOR

**Identification of N-Acyl Hydrazones as New Non-Zinc-Binding MMP-13 Inhibitors
by Structure-Based Virtual Screening Studies and Chemical Optimization**

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Table S1. Combustion analysis data of final compounds 13, 13a-o.

Compound	Formula	Calculated			Found		
		% C	% H	% N	% C	% H	% N
13	C ₂₁ H ₁₇ BrN ₄ O ₅	51.97	3.53	11.55	51.99	3.54	11.58
13a	C ₂₁ H ₁₉ N ₅ O ₇ S	51.96	3.95	14.43	51.98	3.99	14.45
13b	C ₂₂ H ₂₀ N ₄ O ₆	60.55	4.62	12.84	60.58	4.64	12.87
13c	C ₂₀ H ₁₅ BrN ₄ O ₅	50.97	3.21	11.89	51.00	3.23	11.92
13d	C ₂₁ H ₁₈ N ₄ O ₆	59.71	4.30	13.26	59.72	4.34	13.28
13e	C ₂₄ H ₂₀ BrN ₅ O ₄	55.19	3.86	13.41	55.21	3.87	13.44
13f	C ₂₅ H ₂₃ N ₅ O ₅	63.42	4.90	14.79	63.45	4.92	14.80
13g	C ₂₄ H ₂₁ N ₅ O ₅	62.74	4.61	15.24	62.75	4.64	15.26
13h	C ₂₂ H ₁₈ BrN ₃ O ₅	54.56	3.75	8.68	51.57	3.77	8.70
13i	C ₂₃ H ₂₁ N ₃ O ₆	63.44	4.86	9.65	63.45	4.89	9.67
13l	C ₂₁ H ₁₆ BrN ₃ O ₅	53.63	3.43	8.94	53.65	3.47	8.95
13m	C ₂₂ H ₁₈ BrN ₃ O ₅	54.56	3.75	8.68	54.59	3.77	8.69
13n	C ₂₁ H ₁₆ BrN ₇ O ₃	51.03	3.26	19.84	51.05	3.29	19.86
13o	C ₂₂ H ₁₈ BrN ₇ O ₃	51.98	3.57	19.29	52.01	3.58	19.31

II. Supplementary figures

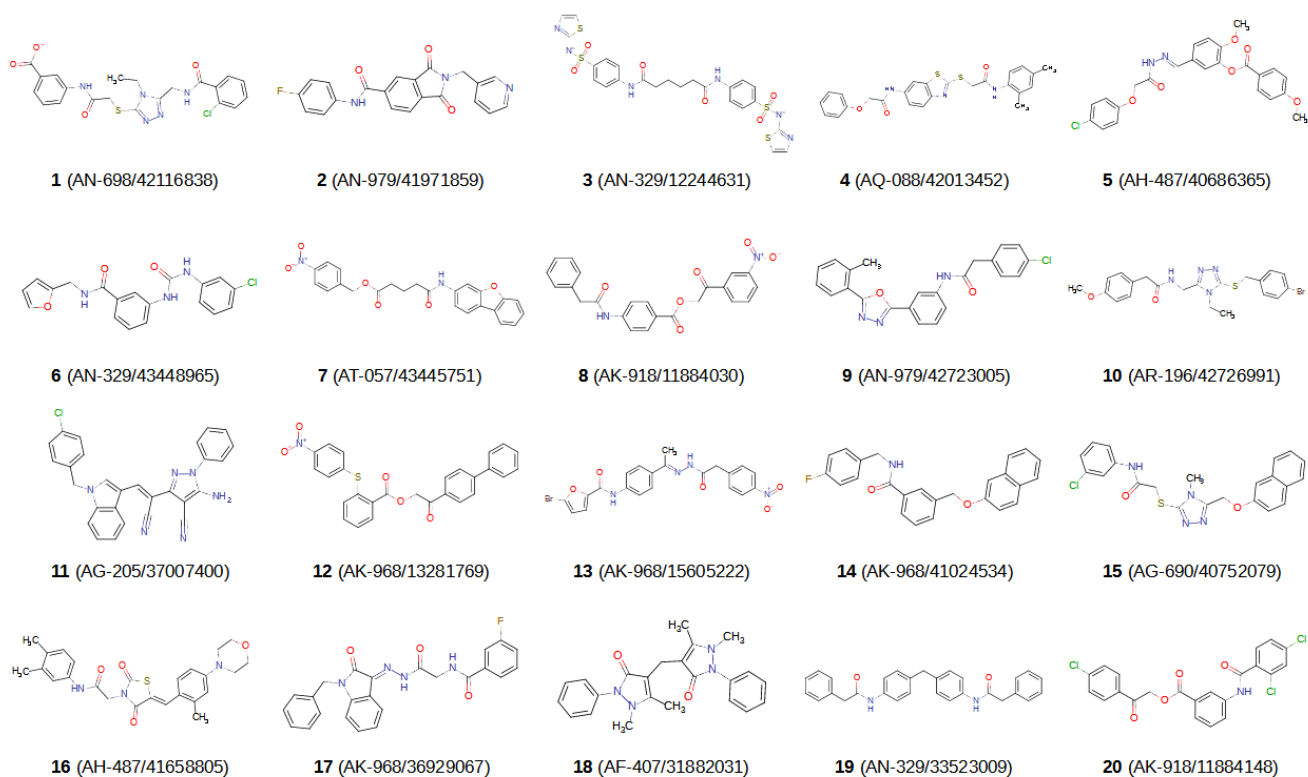
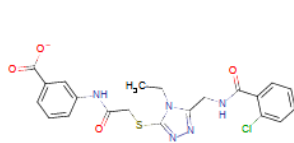
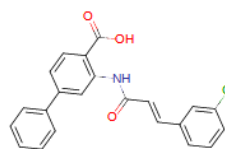


Figure S1. 2D structures of the 20 compounds selected for *in vitro* test. Each compound is identified with its Specs ID number. MarvinSketch¹ was used to draw the structures. The protonation state of each compound corresponds to the protonation state of the docked pose selected for that compound.

A)



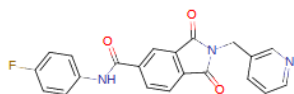
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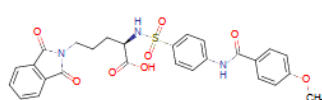
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Tanimoto: 0.39

B)



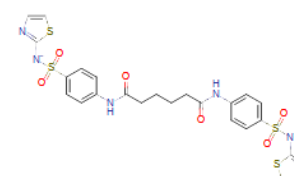
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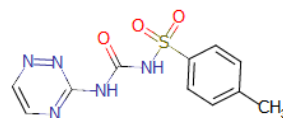
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C)



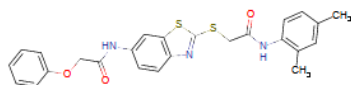
3 (AN-329/12244631)



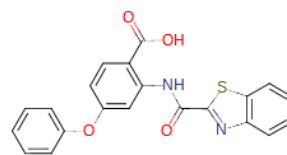
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D)



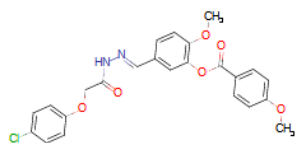
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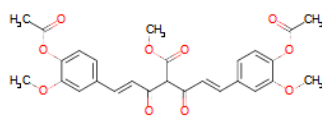
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E)



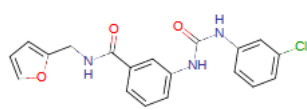
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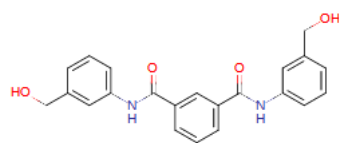
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F)



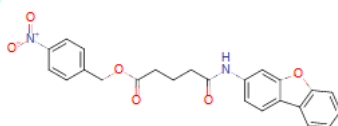
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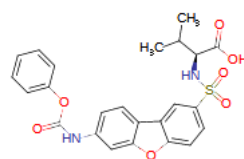
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G)



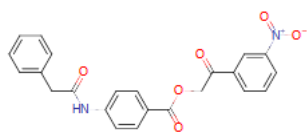
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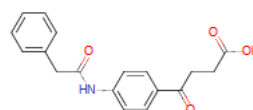
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H)



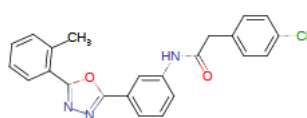
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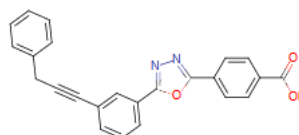
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I)



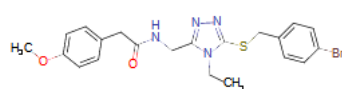
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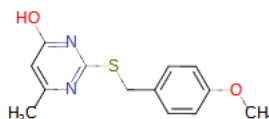
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J)



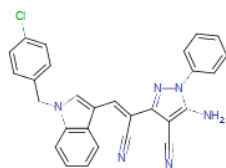
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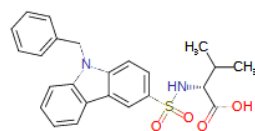
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K)



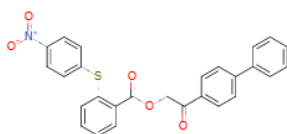
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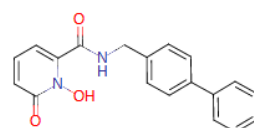
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L)



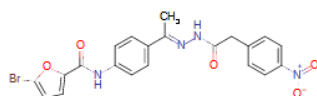
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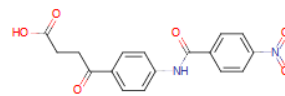
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M)



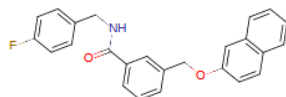
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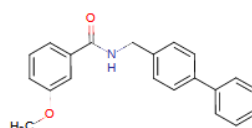
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N)



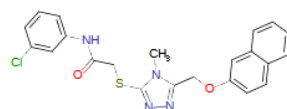
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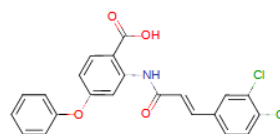
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O)



15 (AG-690/40752079)



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Tanimoto: 0.47

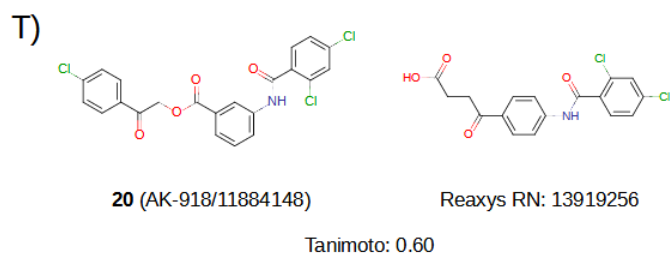
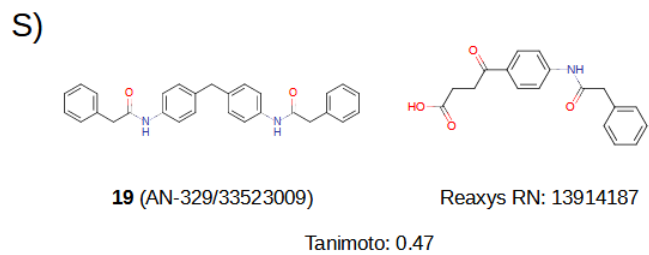
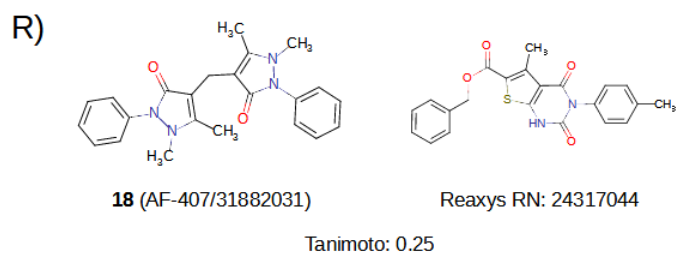
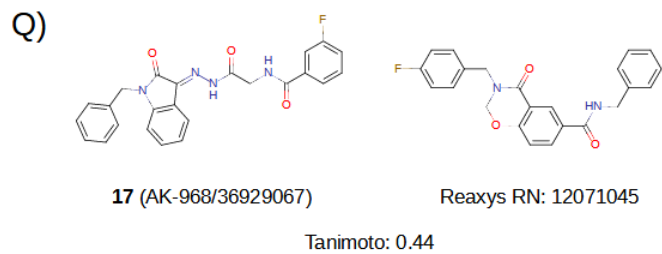
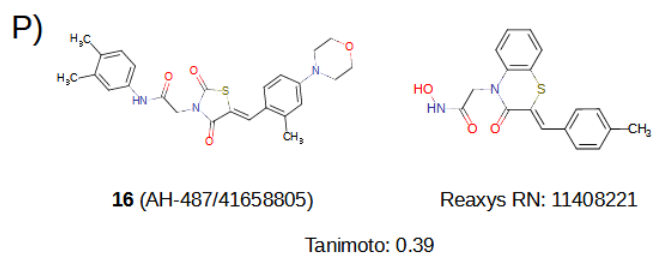


Figure S2. In each panel, a hit compound is represented in 2D, together with the most similar active compound found in the Reaxys² database, which is labeled with its Reaxys Registry Number. The Tanimoto similarity value resulting from the comparison of the OpenEyePath³ fingerprint of both compounds is shown below. MarvinSketch¹ was used to draw the structures.

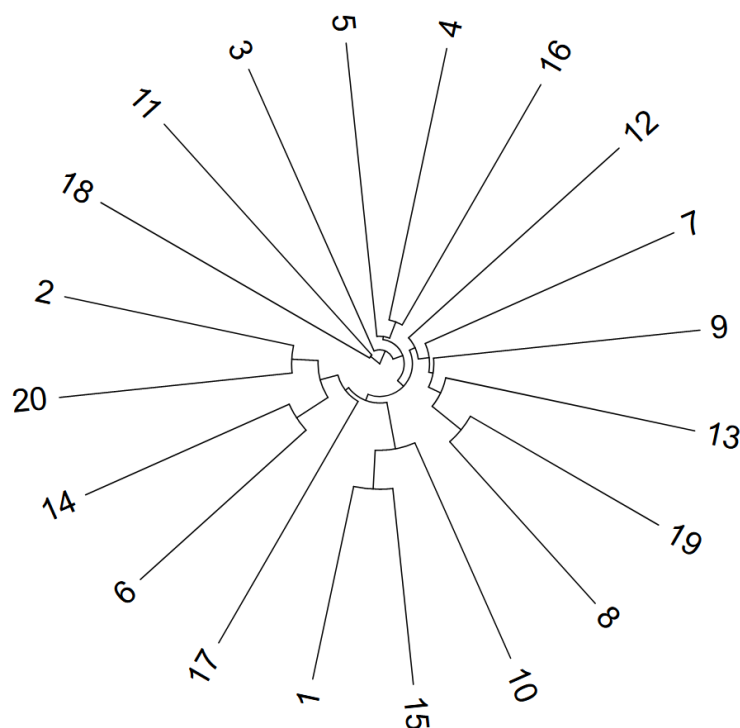


Figure S3. Dendrogram based on fingerprints showing the structural diversity of the 20 selected compounds. The fingerprint used to obtain the distance matrix was the OpenEyePath fingerprint.³ iTOL⁴ was used to draw the dendrogram.

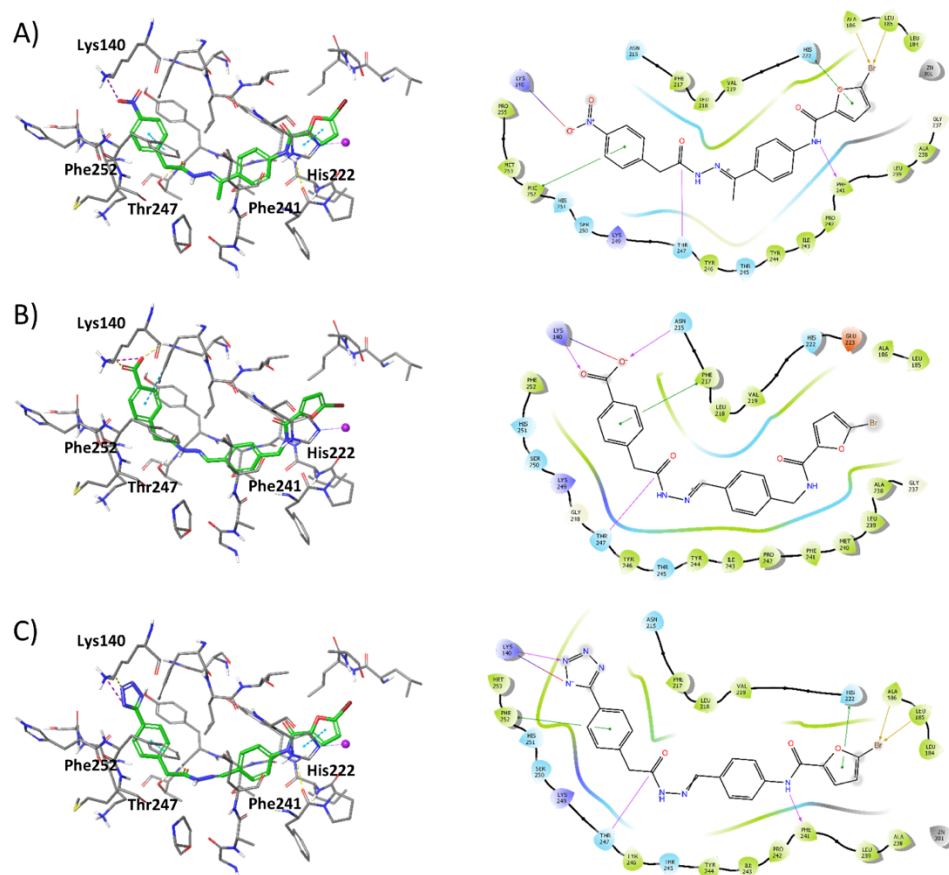


Figure S4. Protein-ligand interactions performed by the docked poses of compounds 13, 13m and 13n (panels A, B and C, respectively) with MMP-13. The first picture in each panel shows the intermolecular interactions predicted by docking in 3D (hydrogen bonds are colored in yellow, π stacking interactions are colored in cyan, and salt bridges are colored in magenta). The second picture shows the intermolecular interactions predicted by docking in 2D (hydrogen bonds, halogen bonds, π stacking interactions and salt bridges are represented by magenta arrows, yellow arrows, green lines and red and blue lines, respectively; positively charged, negatively charged, polar and hydrophobic residues are colored in blue, red, cyan and green, respectively). Figure obtained with Maestro (Schrödinger Release 2016-3: Maestro, Schrödinger, LLC, New York, NY, 2016).

III. References

¹ Marvin 16.10.10.0, 2016, ChemAxon <http://www.chemaxon.com>.

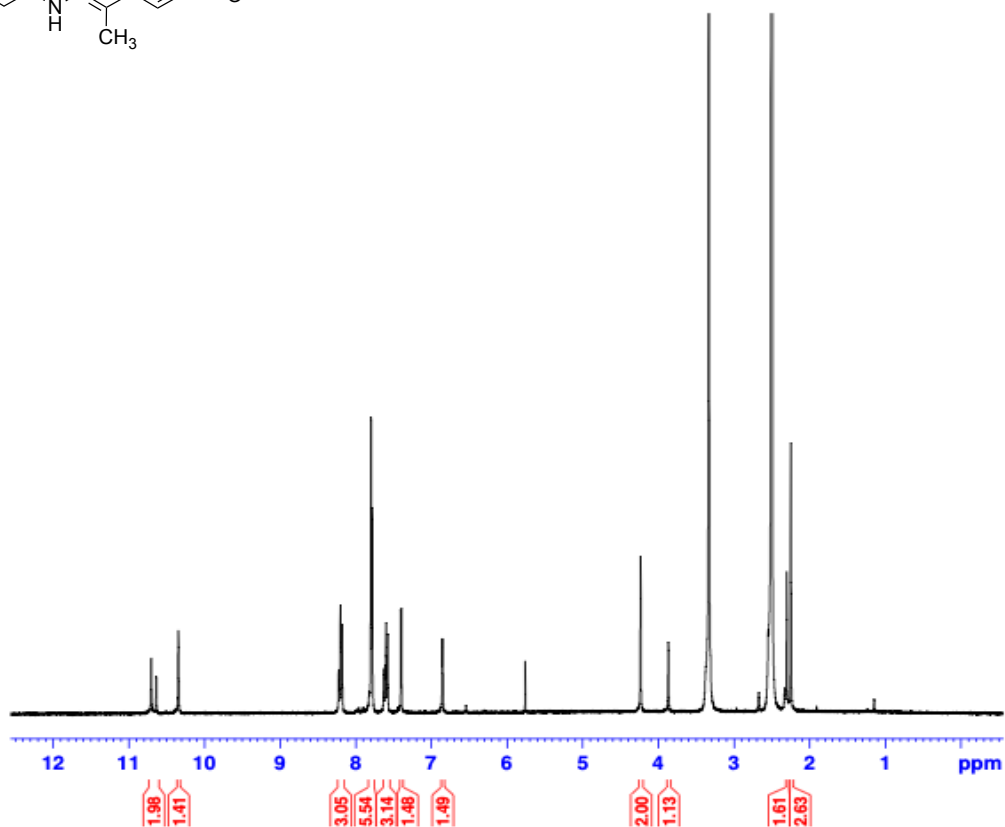
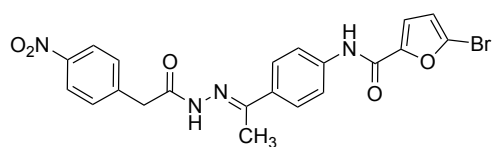
² Reaxys, <https://www.reaxys.com/>

³ OpenEye scientific software, OEChem, 2013.

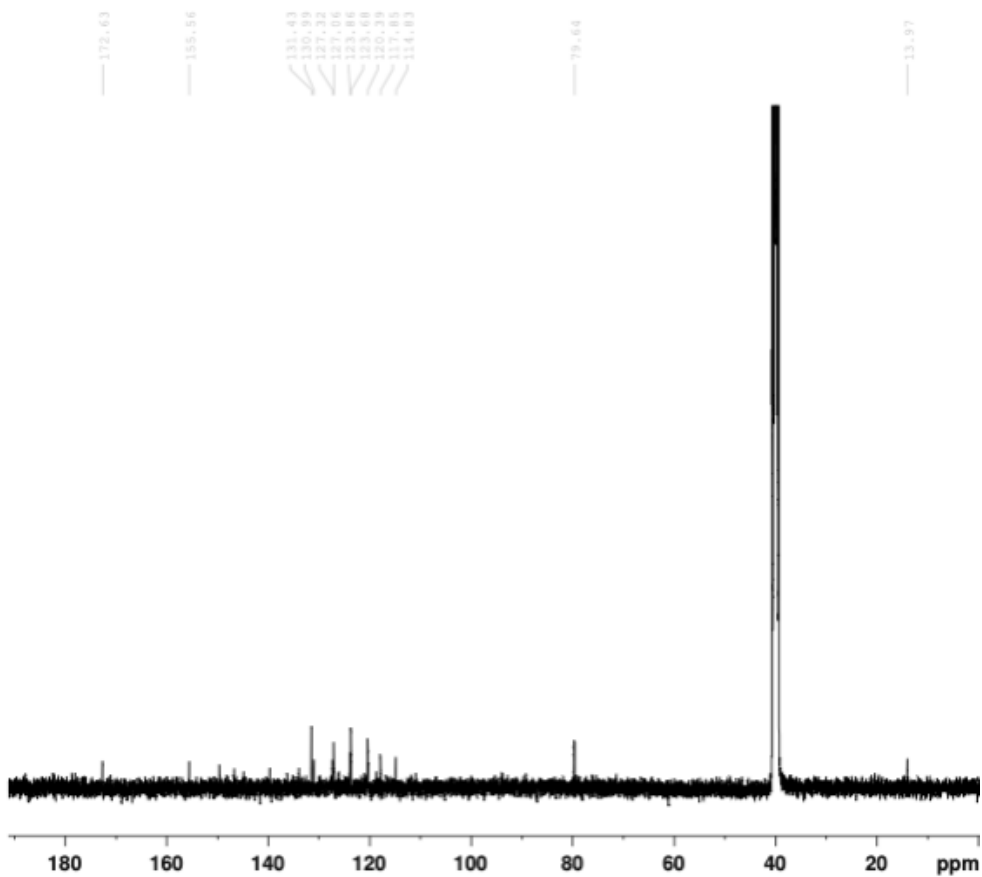
⁴ Letunic, I.; Bork, P. Interactive Tree of Life (ITOL) v3: An Online Tool for the Display and Annotation of Phylogenetic and Other Trees. *Nucleic Acids Res.* 2016, 44 (W1), W242-5.

IV. Representative NMR spectra

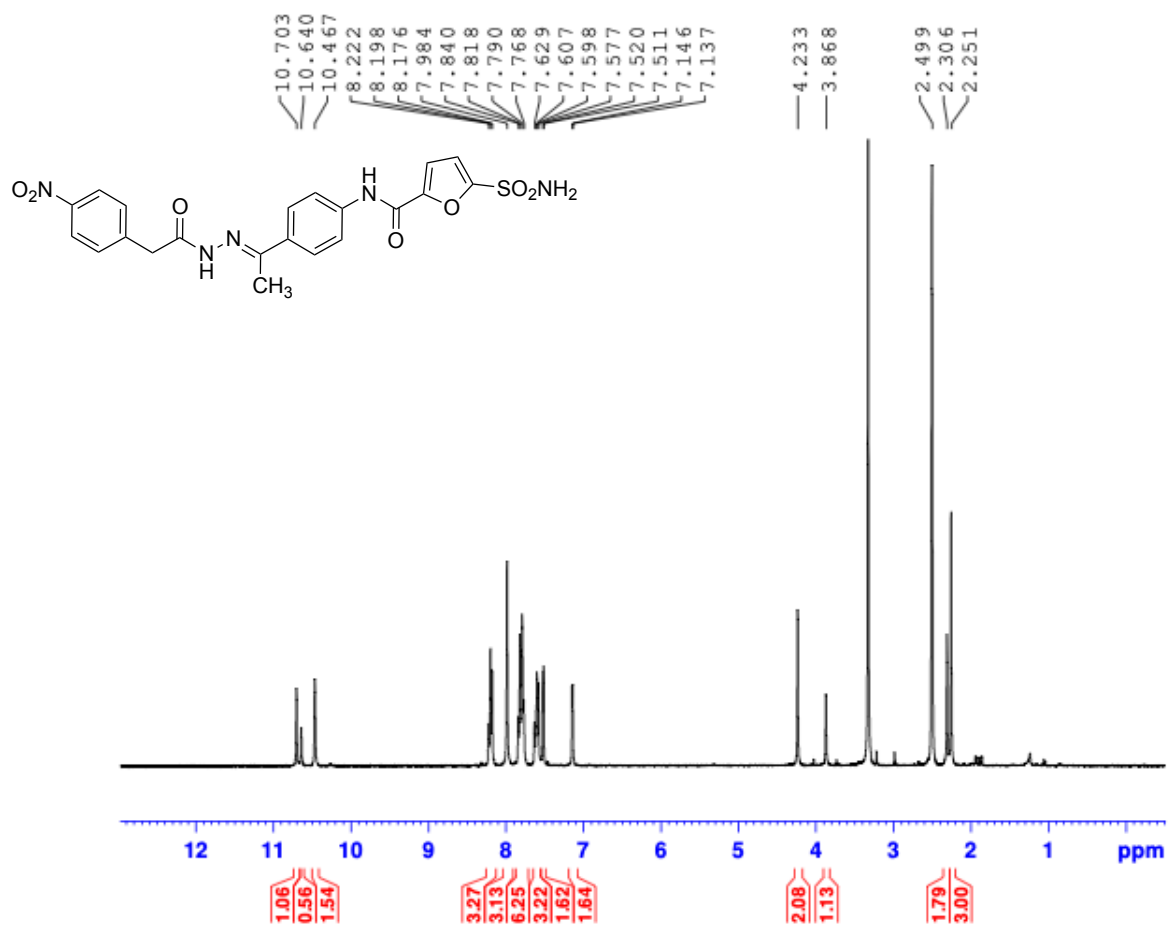
^1H NMR of compound **13** (400MHz, DMSO- d_6)



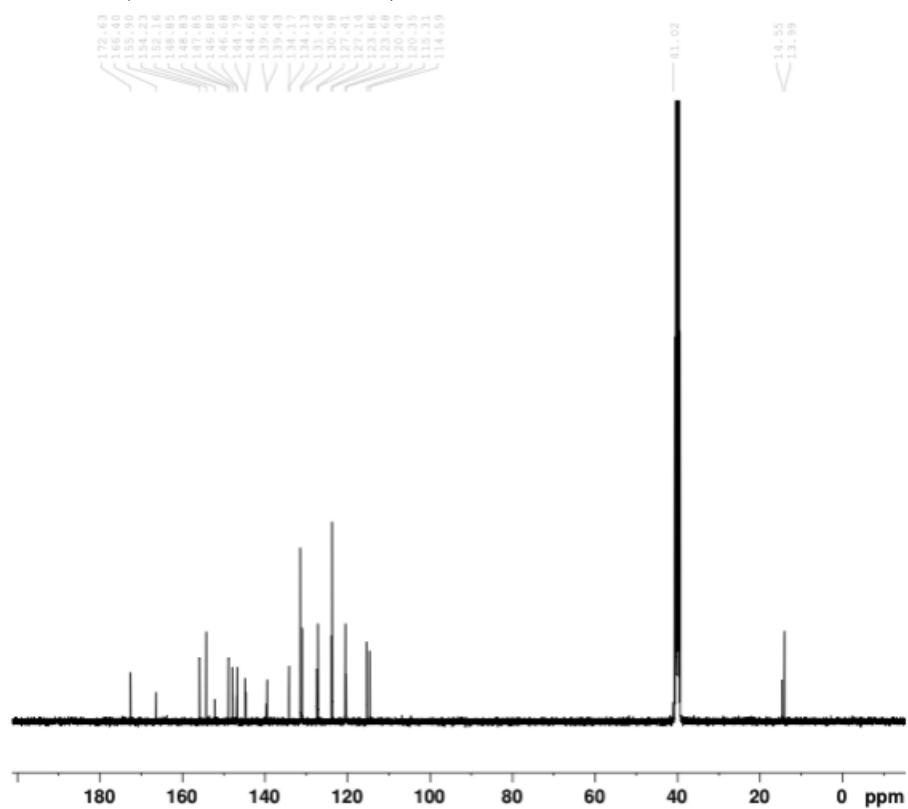
^{13}C NMR of compound **13** (100MHz, DMSO- d_6)



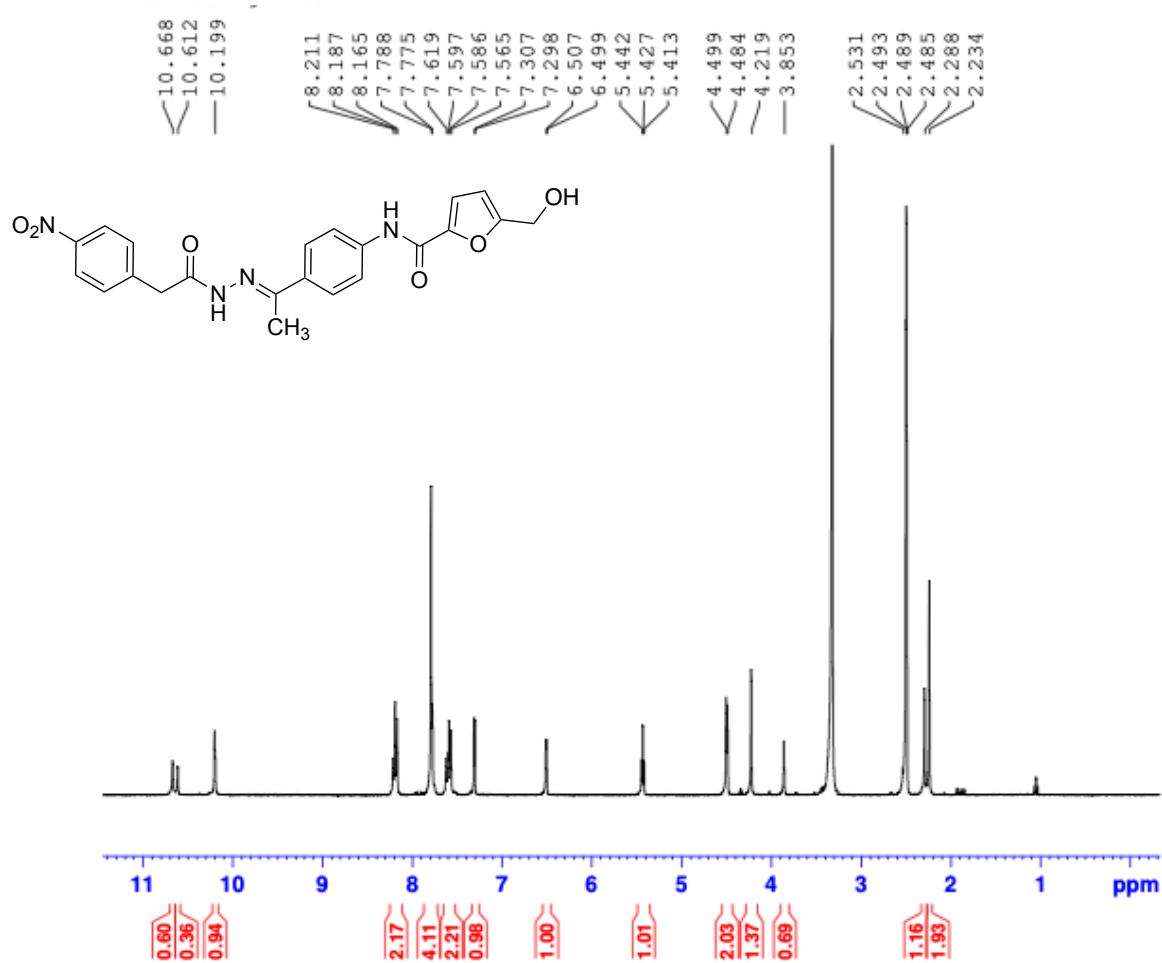
^1H NMR of compound **13a** (400MHz, DMSO- d_6)



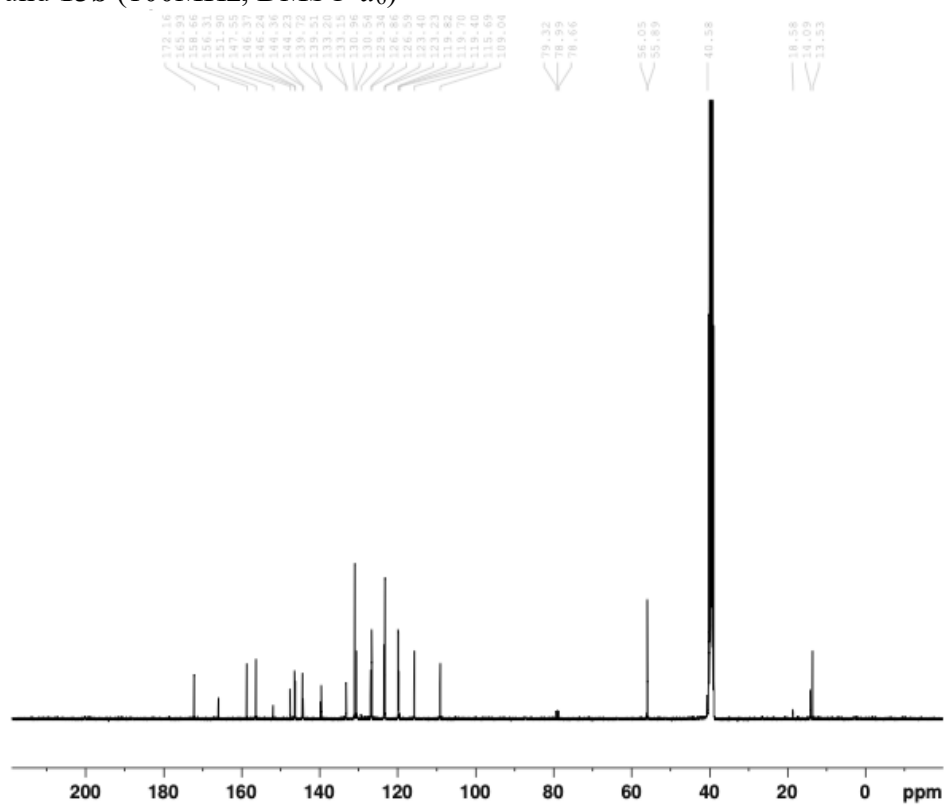
^{13}C NMR of compound **13a** (100MHz, DMSO- d_6)



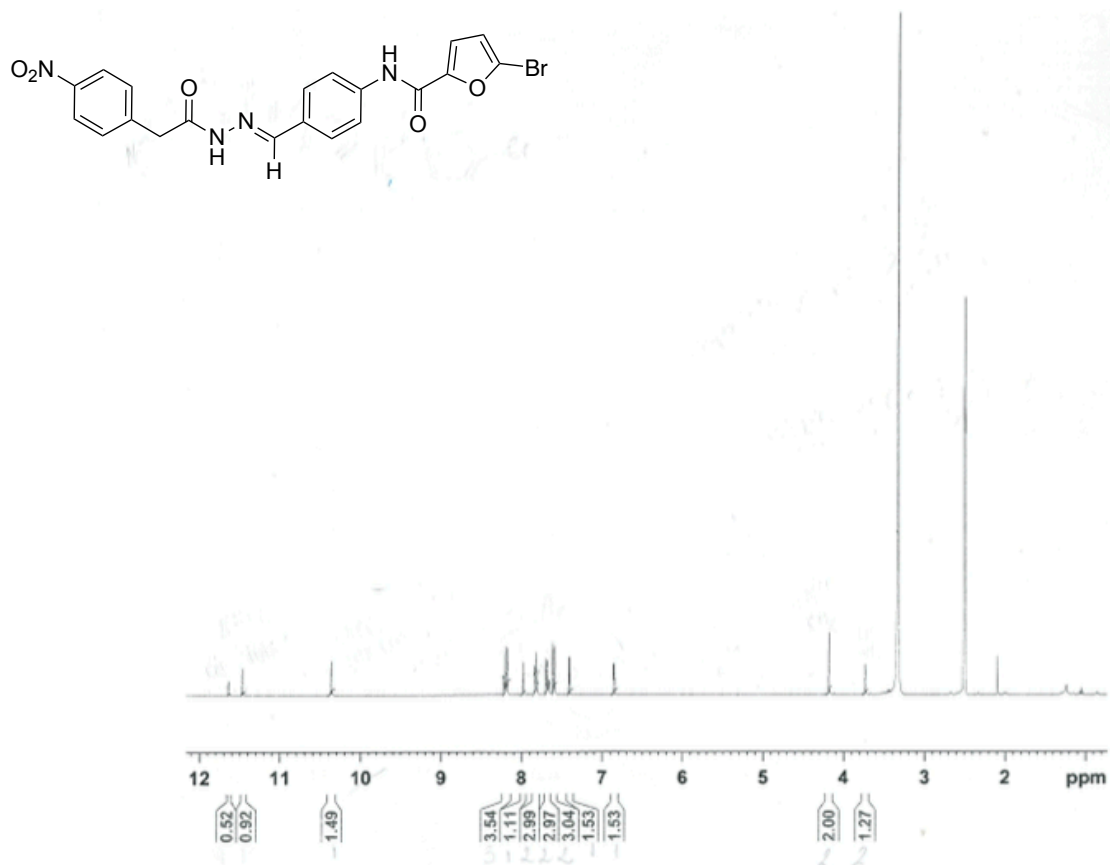
^1H NMR of compound **13b** (400MHz, DMSO- d_6)



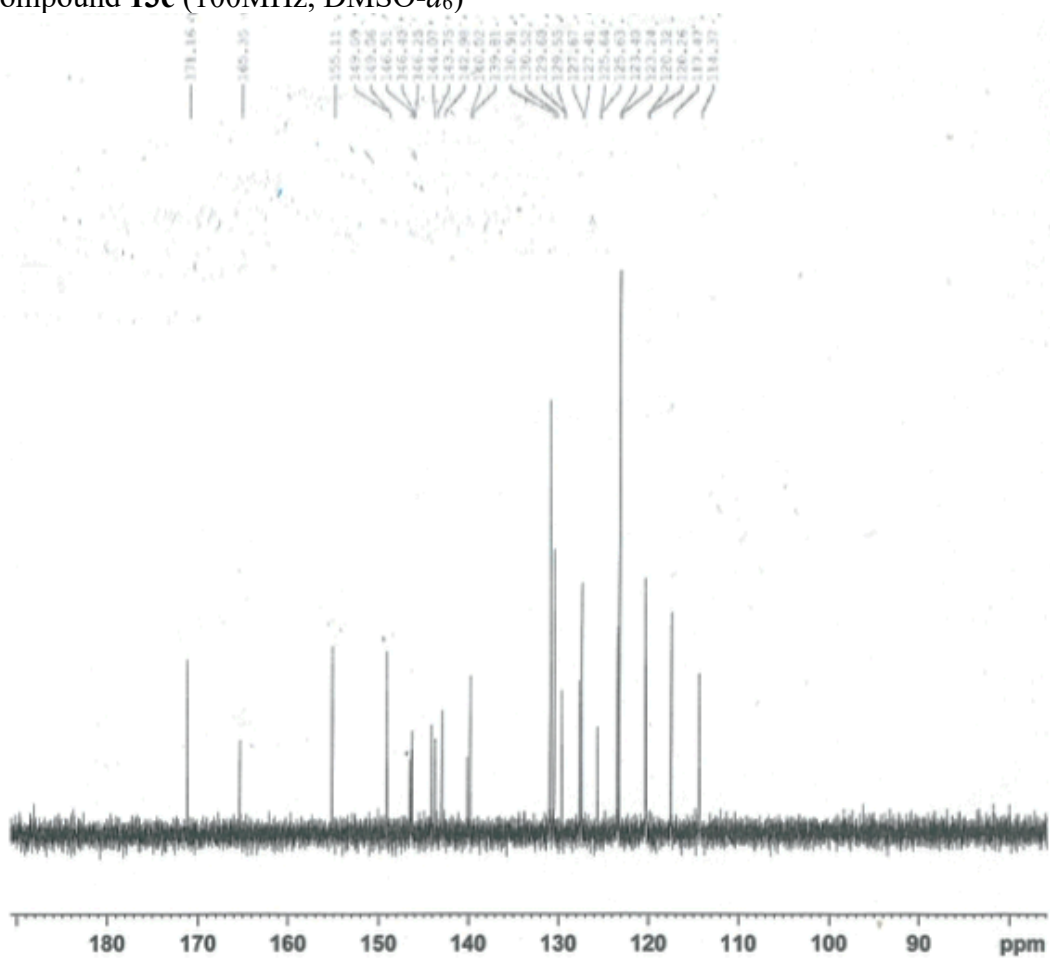
^{13}C NMR of compound **13b** (100MHz, DMSO- d_6)



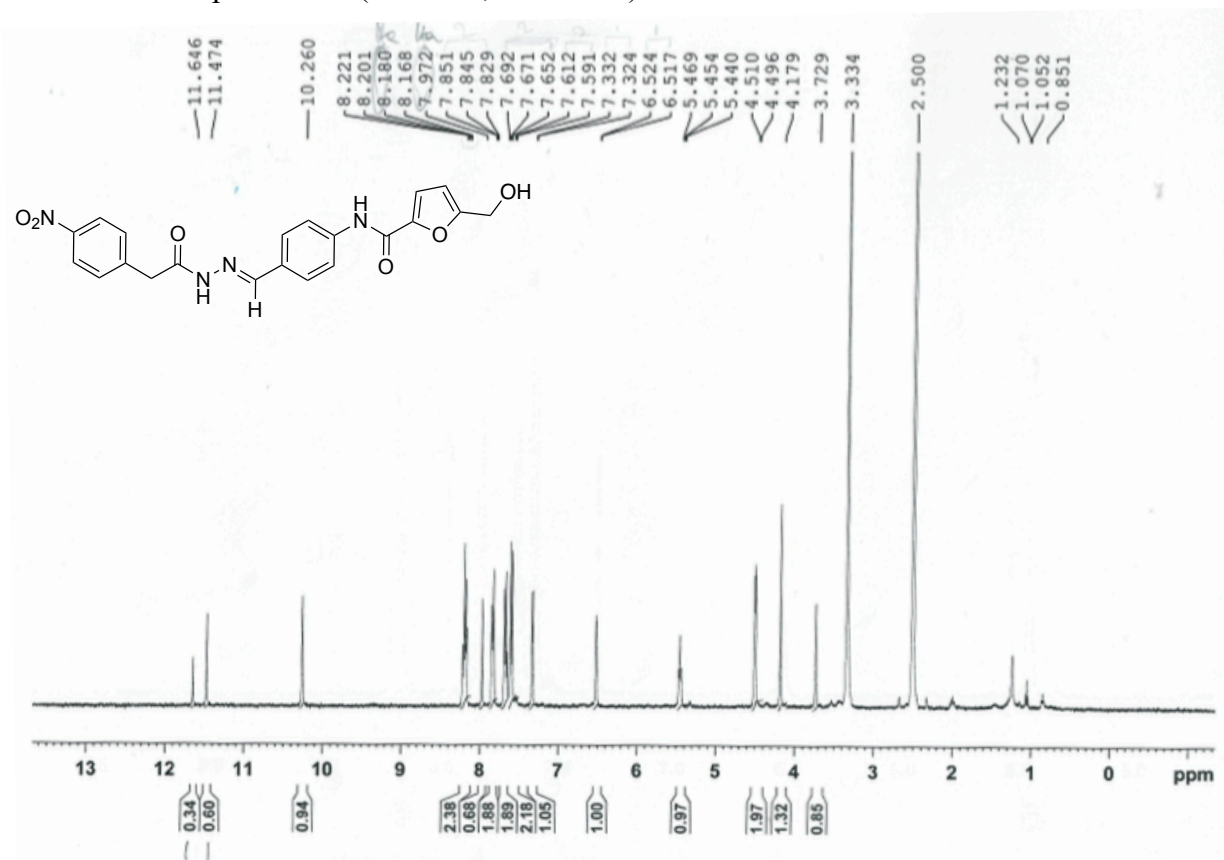
^1H NMR of compound **13c** (400MHz, $\text{DMSO-}d_6$)



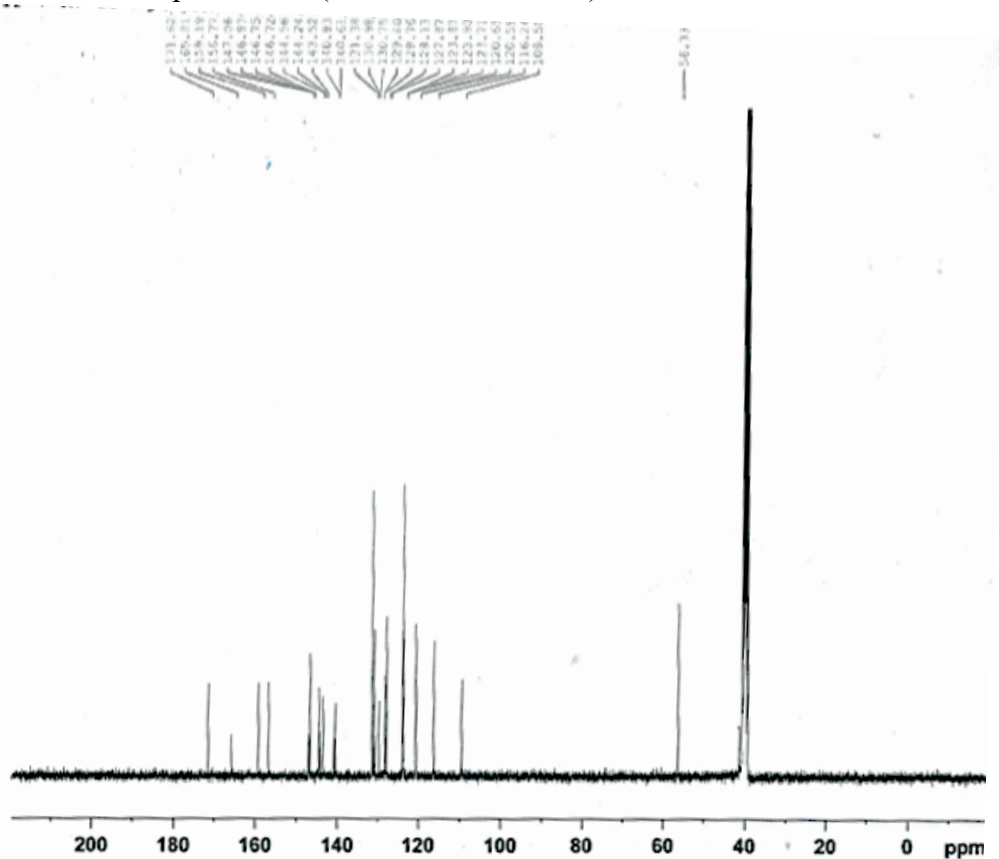
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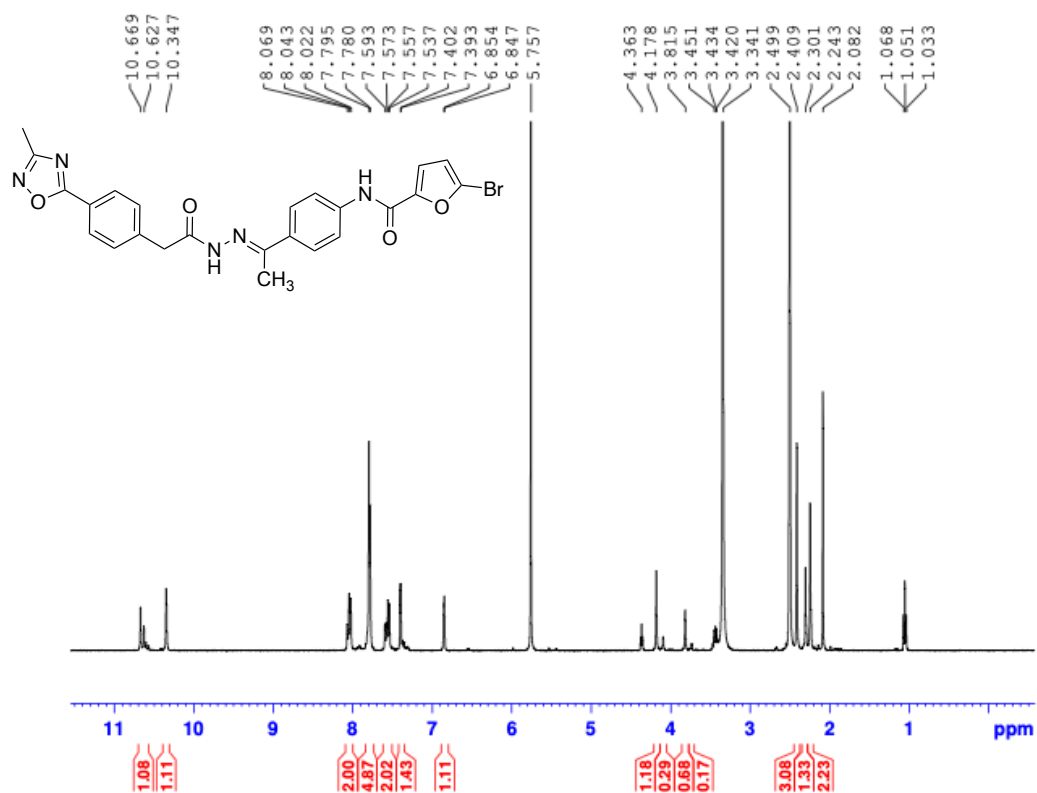
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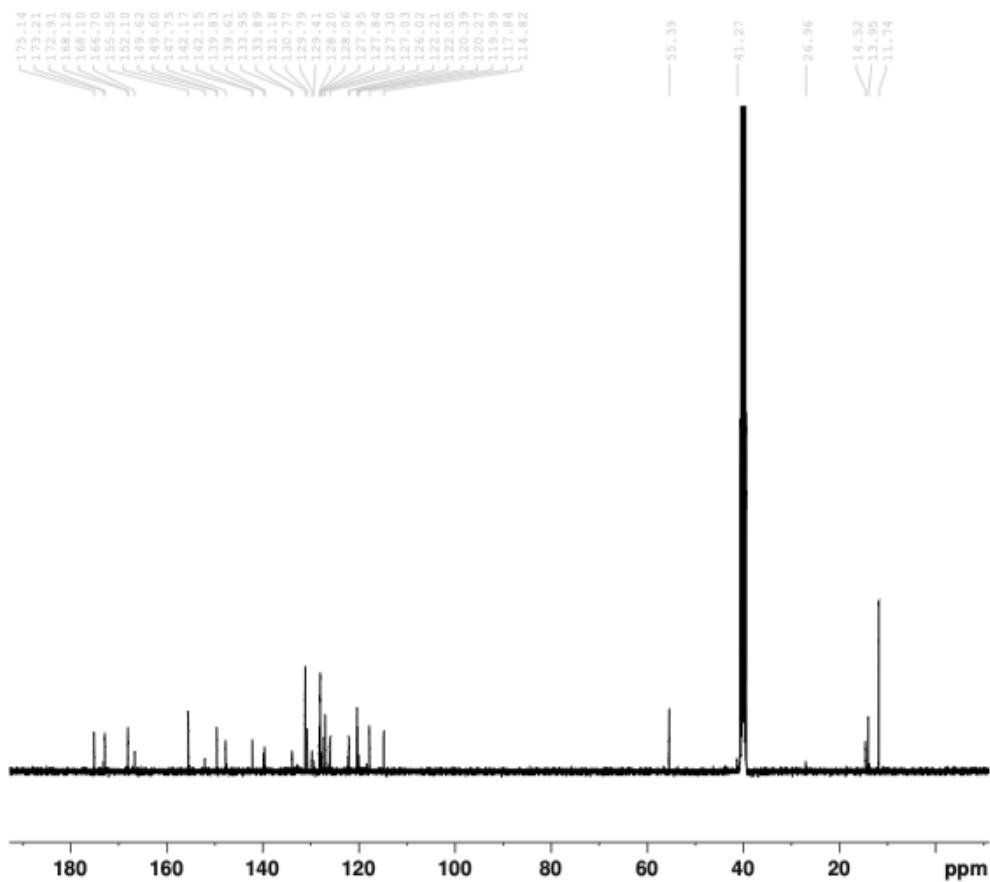
^{13}C NMR of compound **13d** (100MHz, DMSO- d_6)



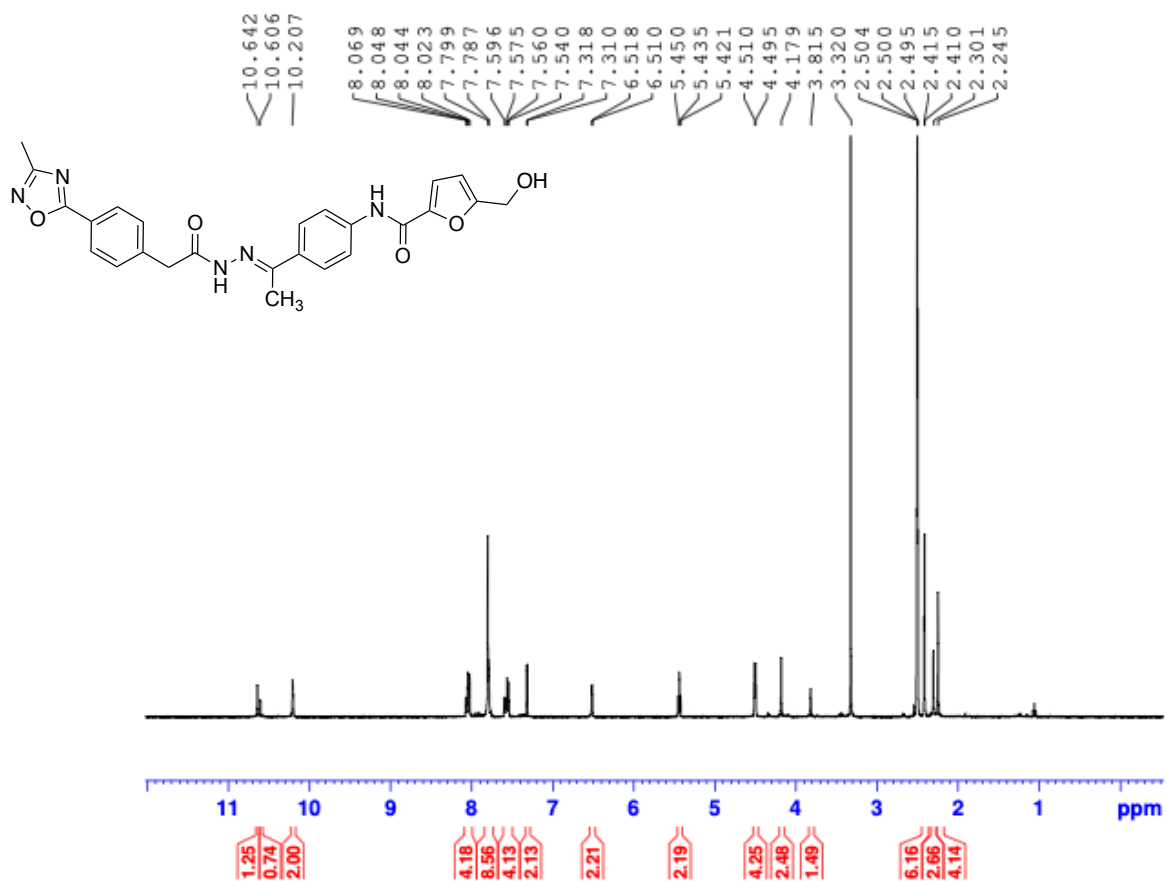
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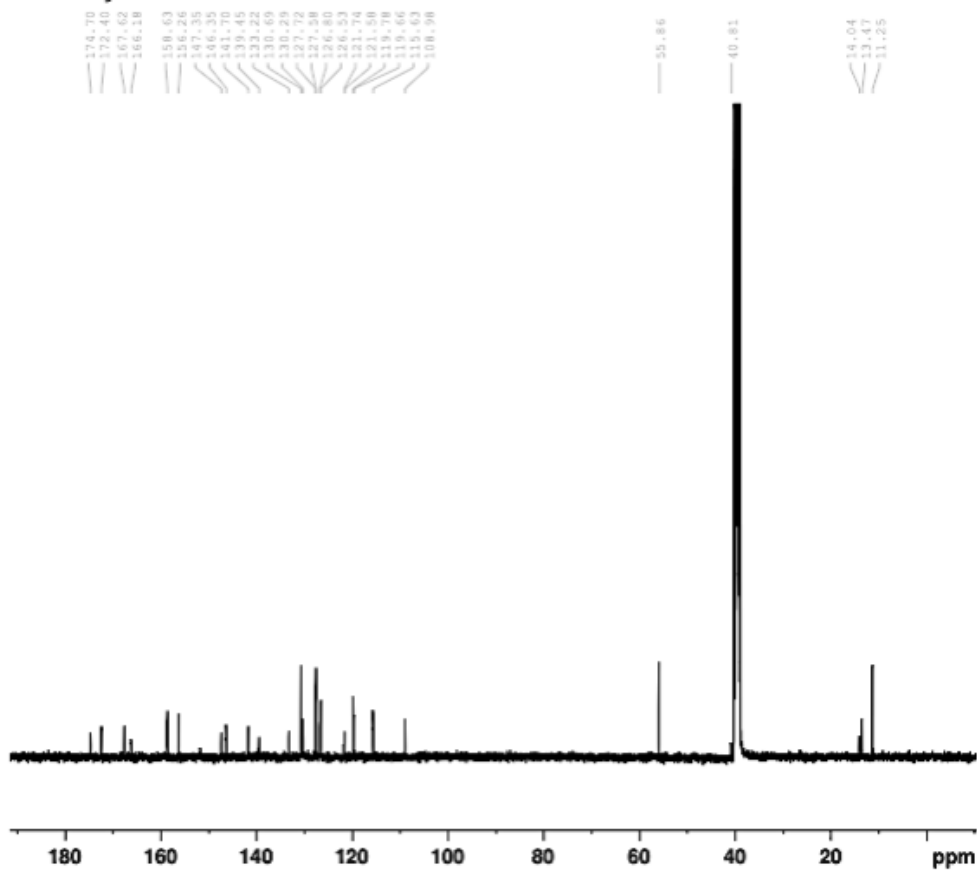
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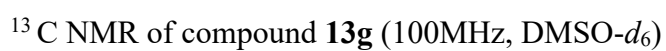


^1H NMR of compound **13f** (400MHz, $\text{DMSO}-d_6$)

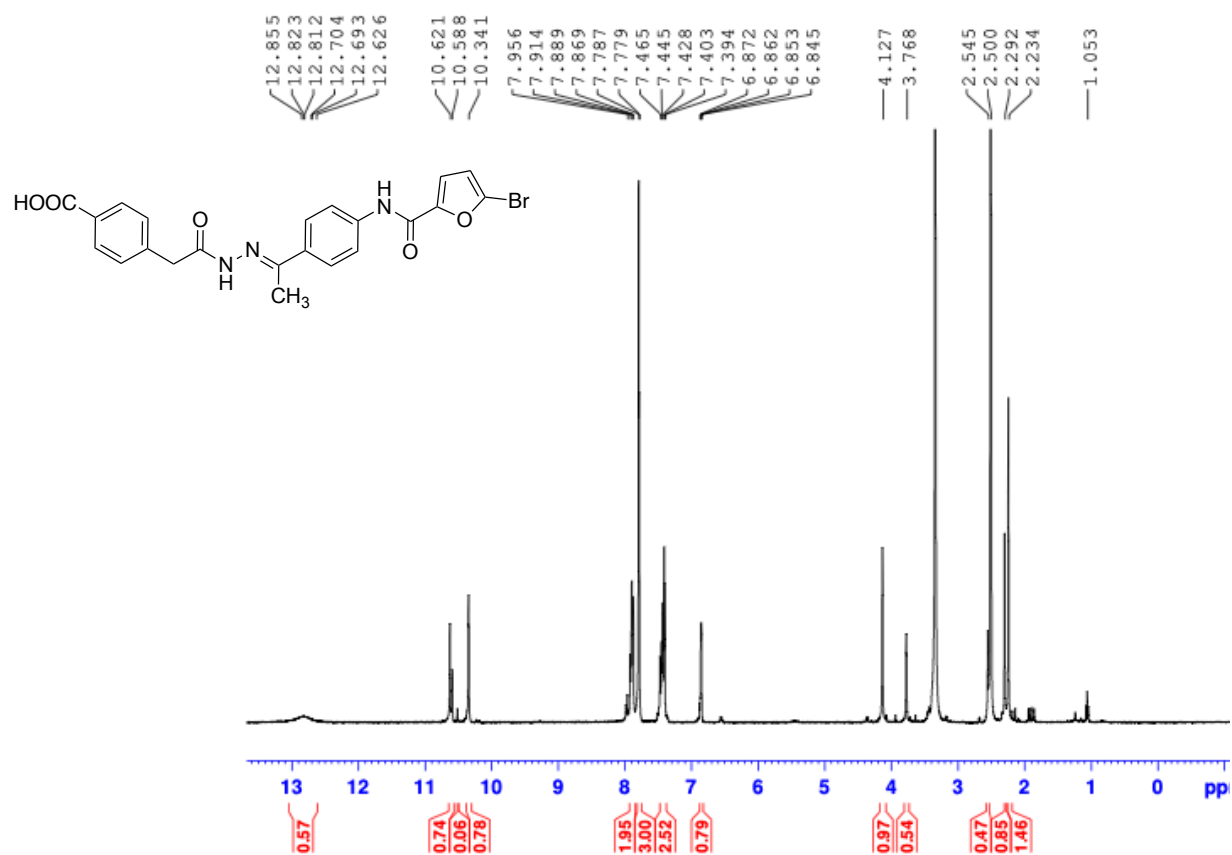


^{13}C NMR of compound **13f** (100MHz, $\text{DMSO}-d_6$)

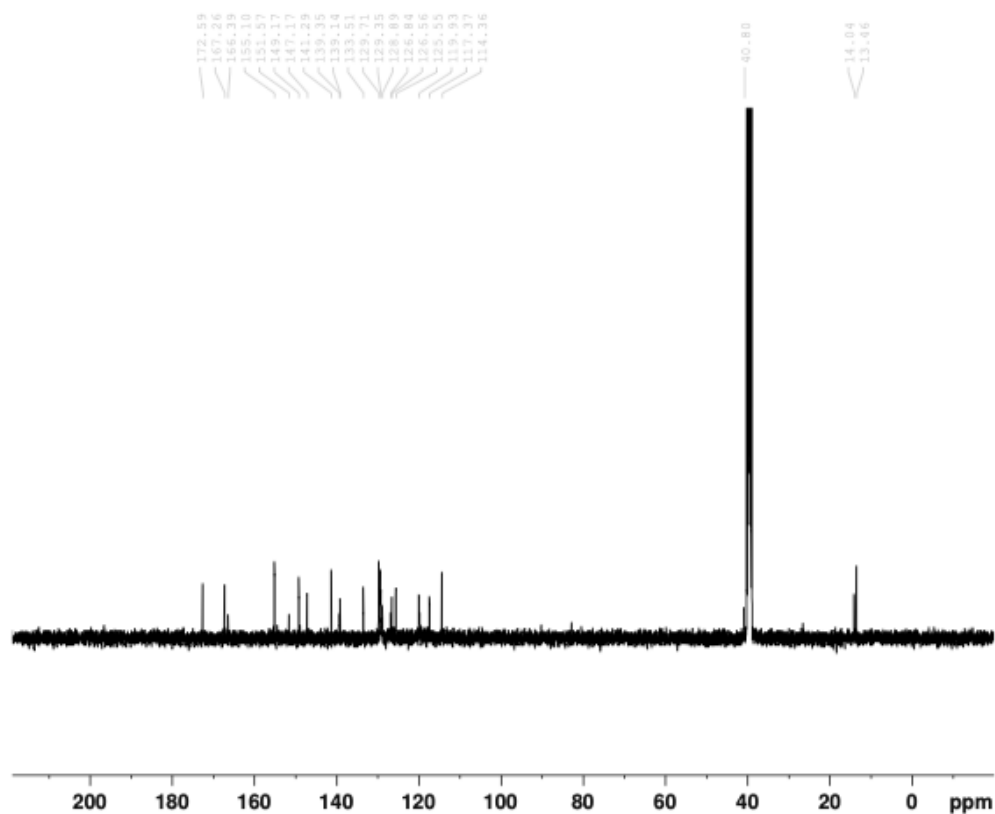


Cc1nc2cc(ccc2o1)CC(=O)NN=Cc3ccc(NC(=O)c4cc(O)oc4)cc3

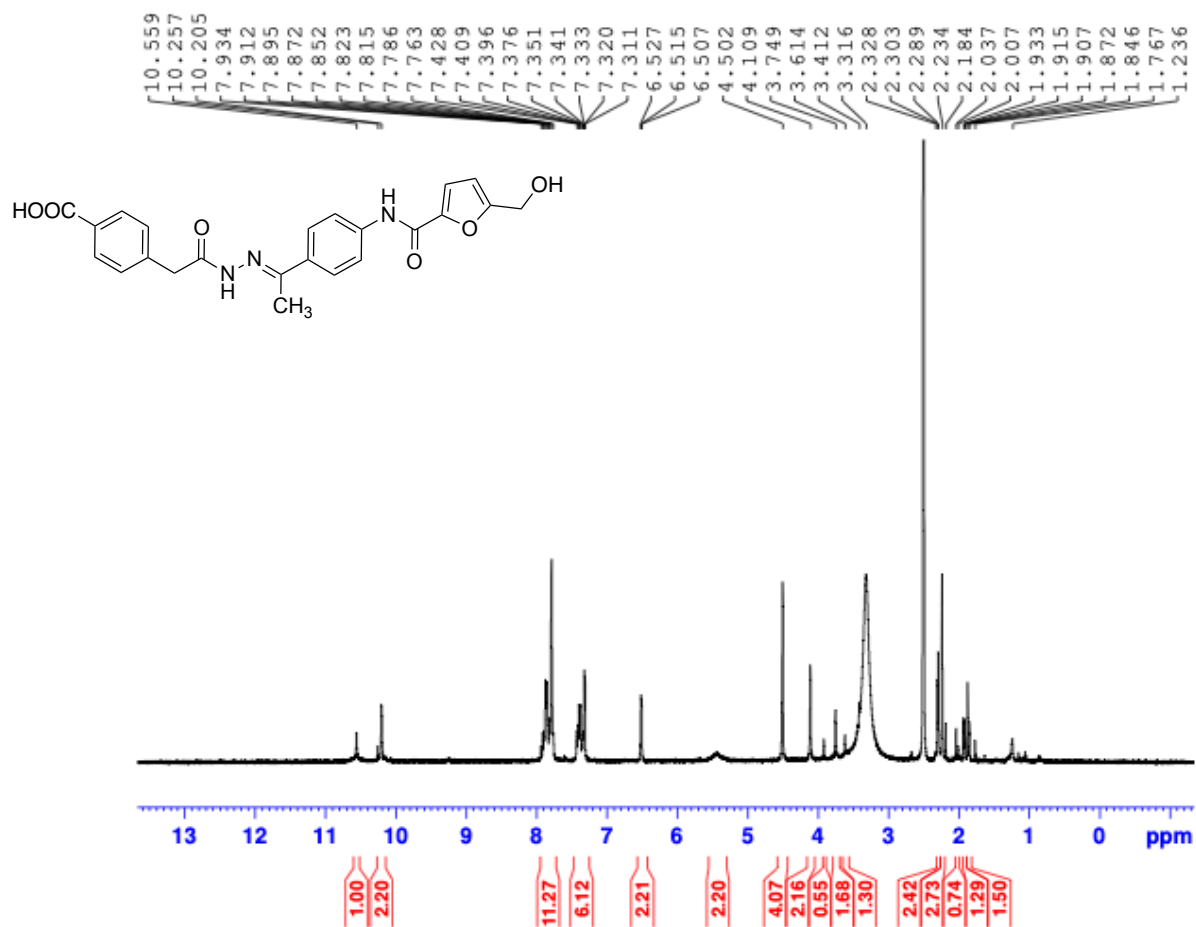
^1H NMR of compound **13h** (400MHz, DMSO- d_6)



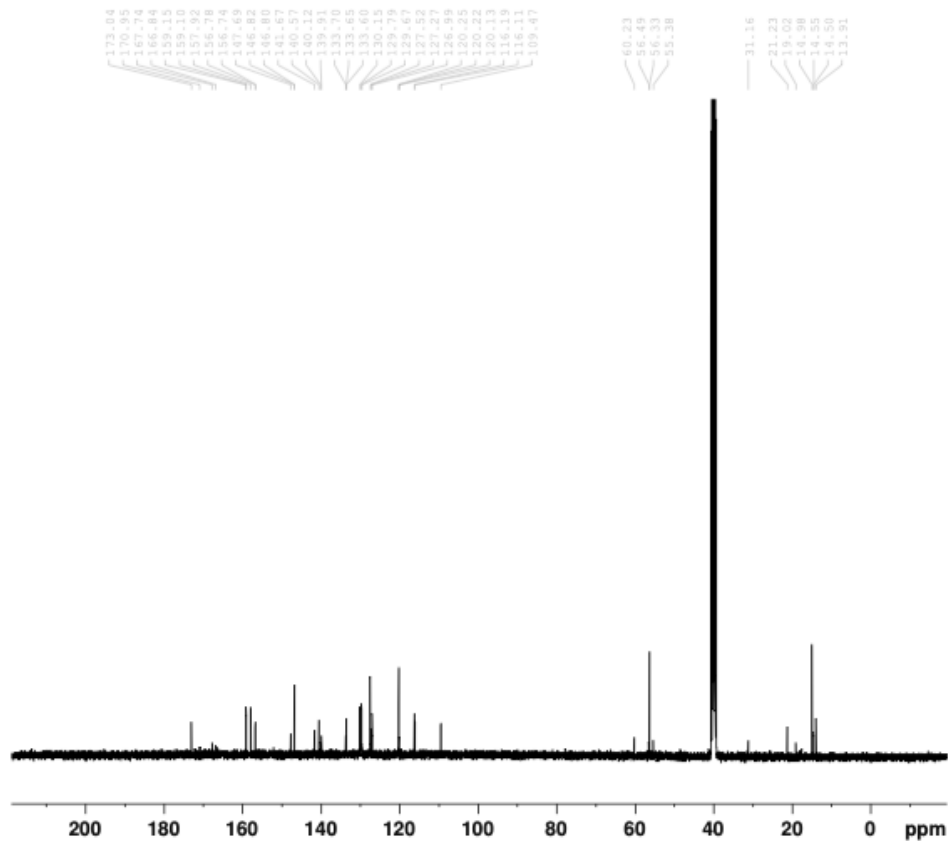
^{13}C NMR of compound **13h** (100MHz, DMSO- d_6)



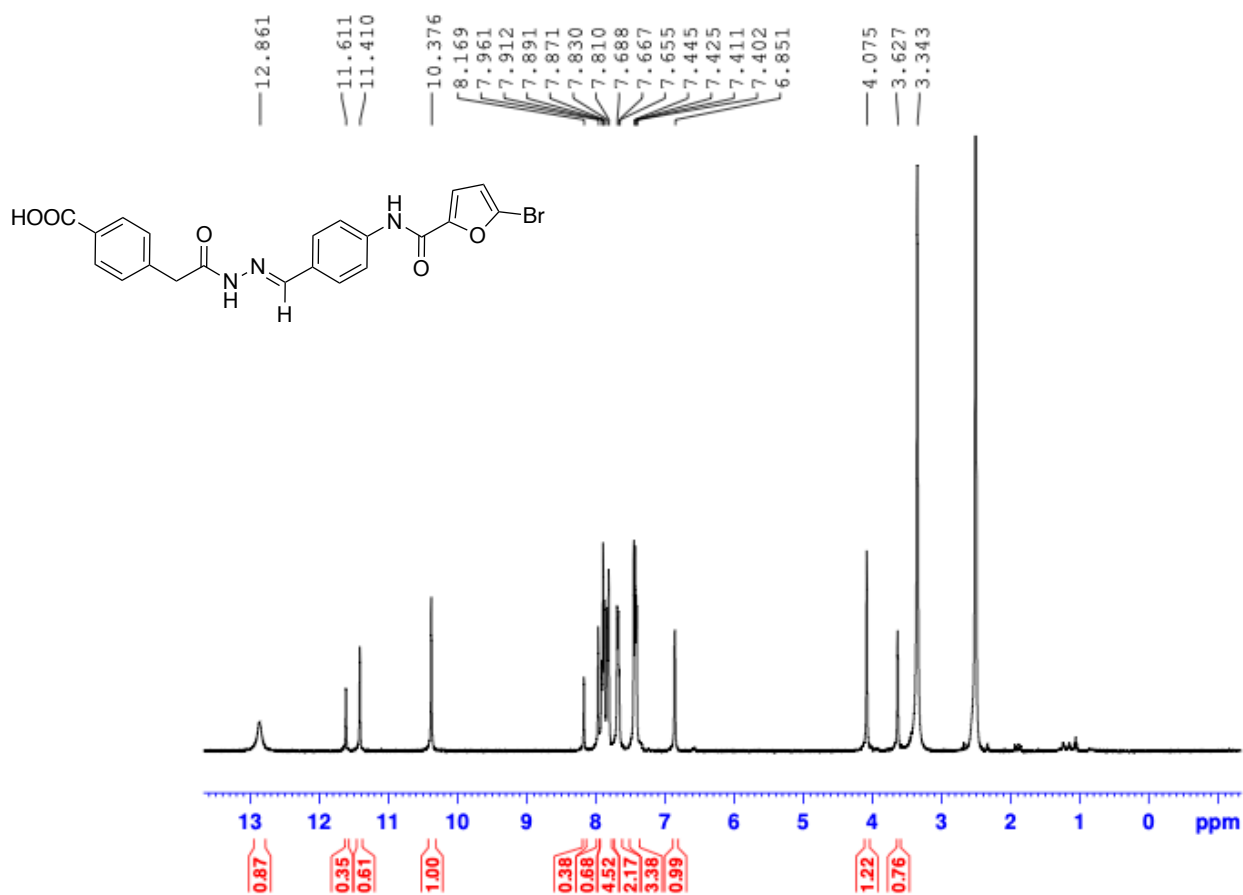
^1H NMR of compound **13i** (400MHz, $\text{DMSO}-d_6$)



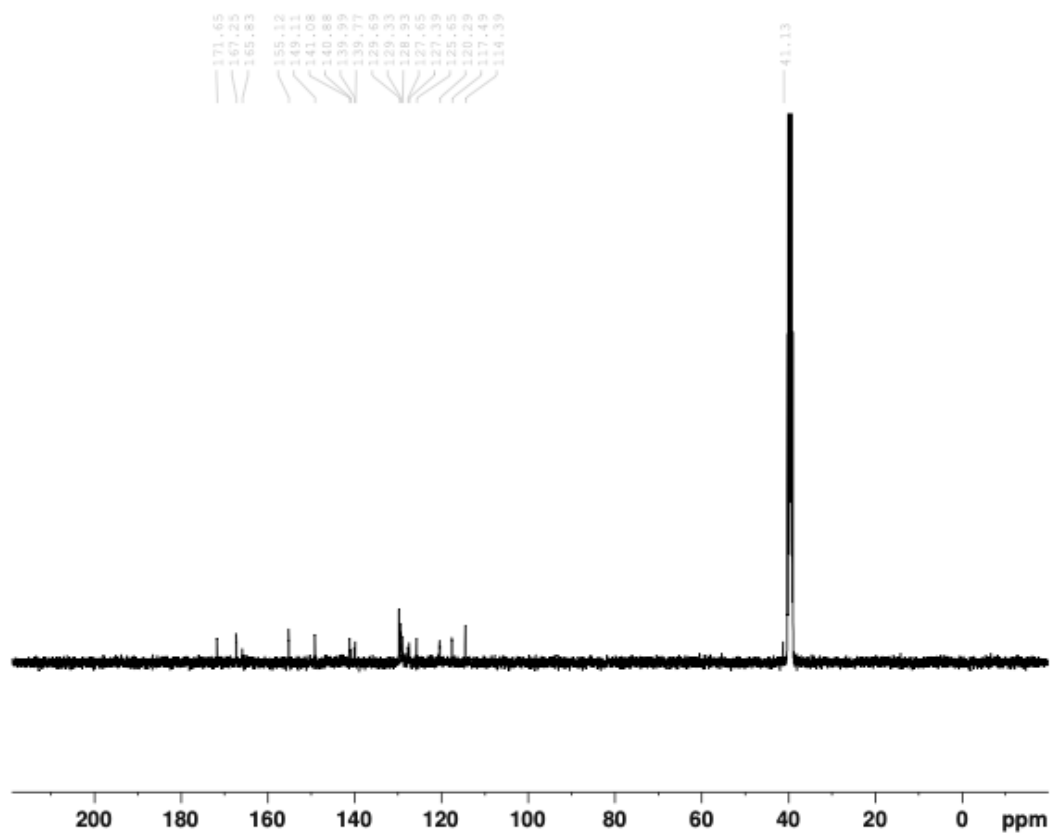
^{13}C NMR of compound **13i** (100MHz, $\text{DMSO}-d_6$)



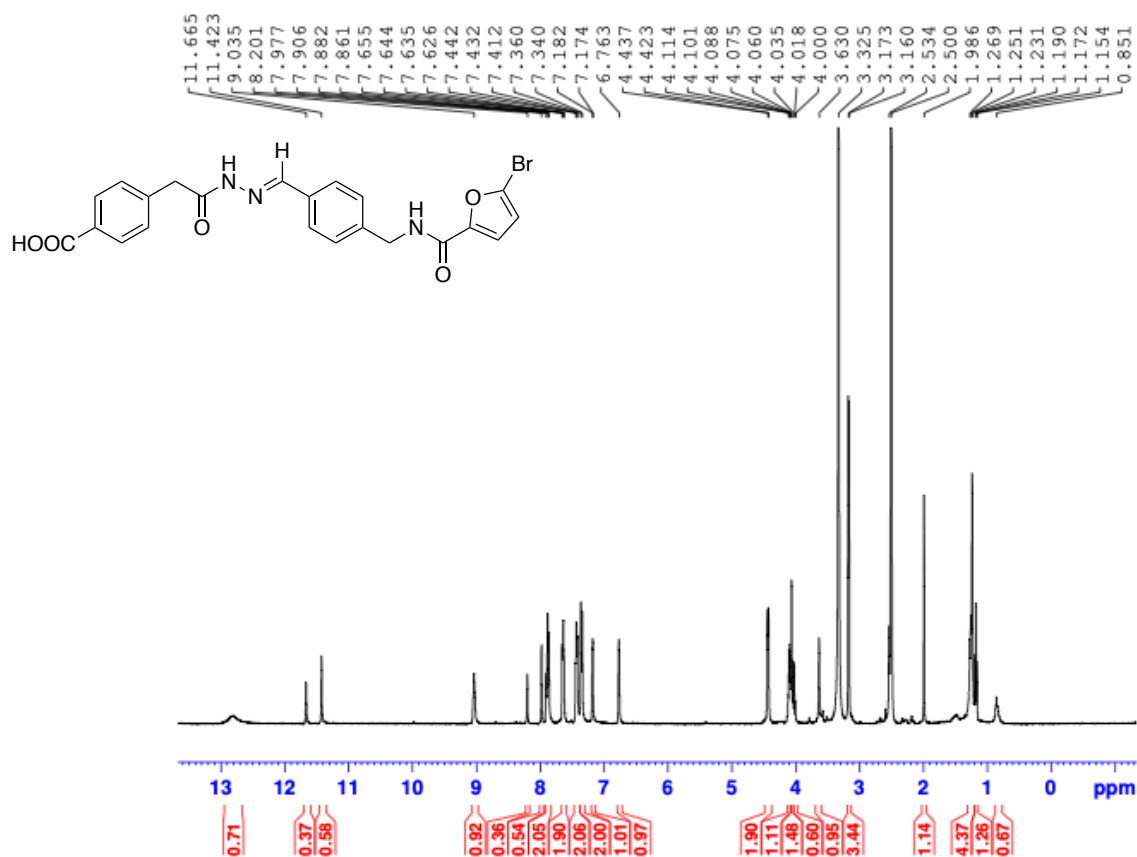
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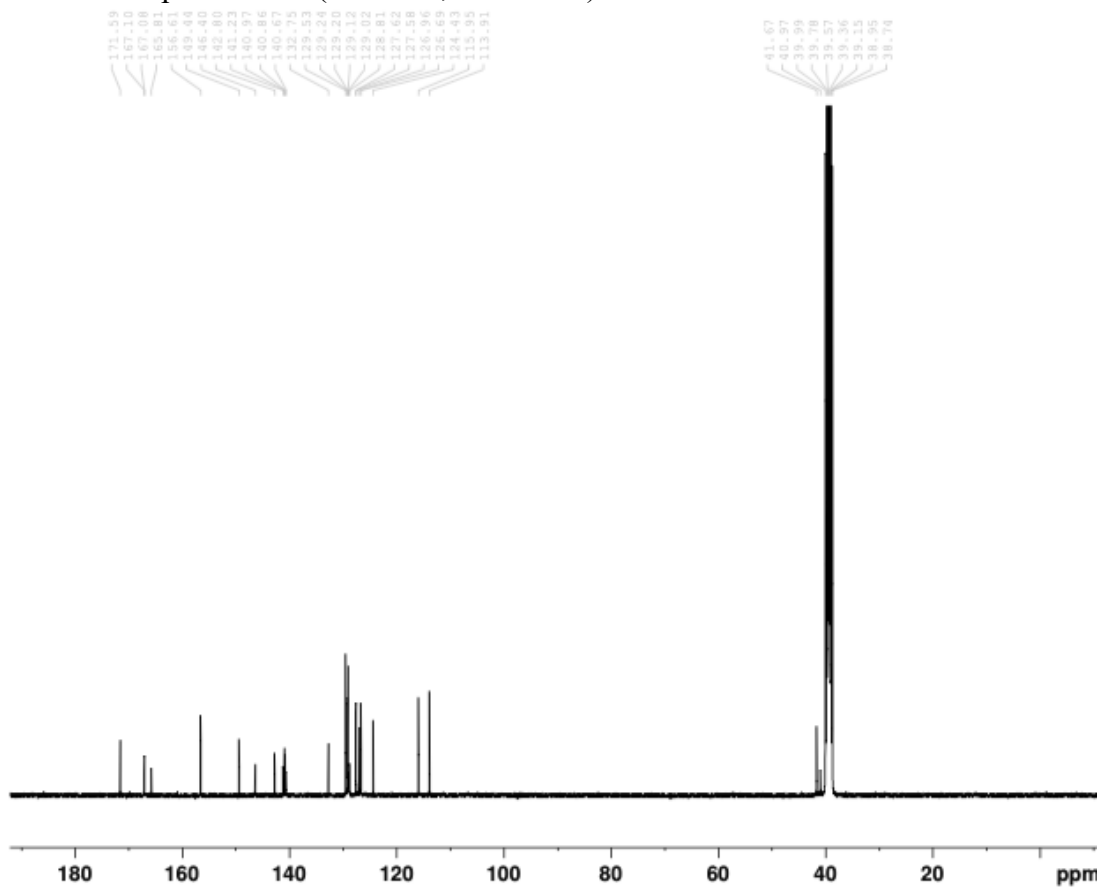
^{13}C NMR of compound **13I** (100MHz, $\text{DMSO}-d_6$)



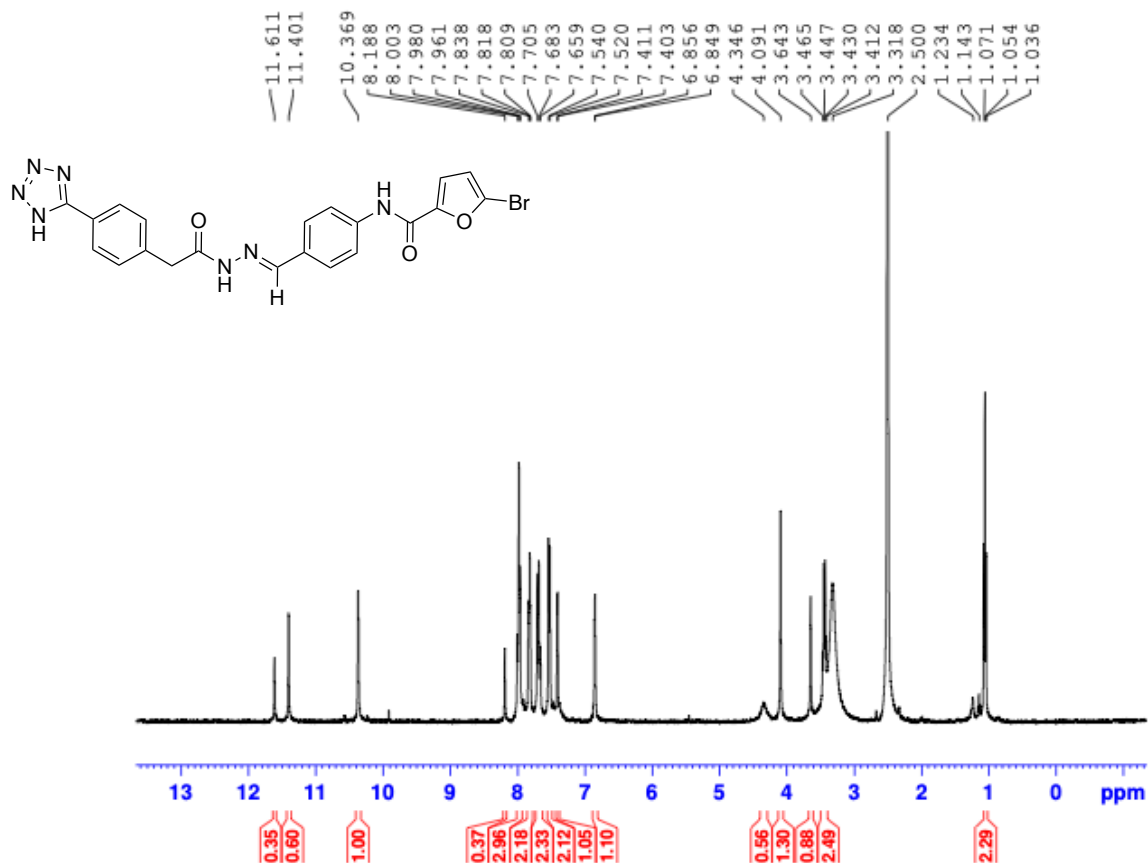
^1H NMR of compound **13m** (400MHz, $\text{DMSO}-d_6$)



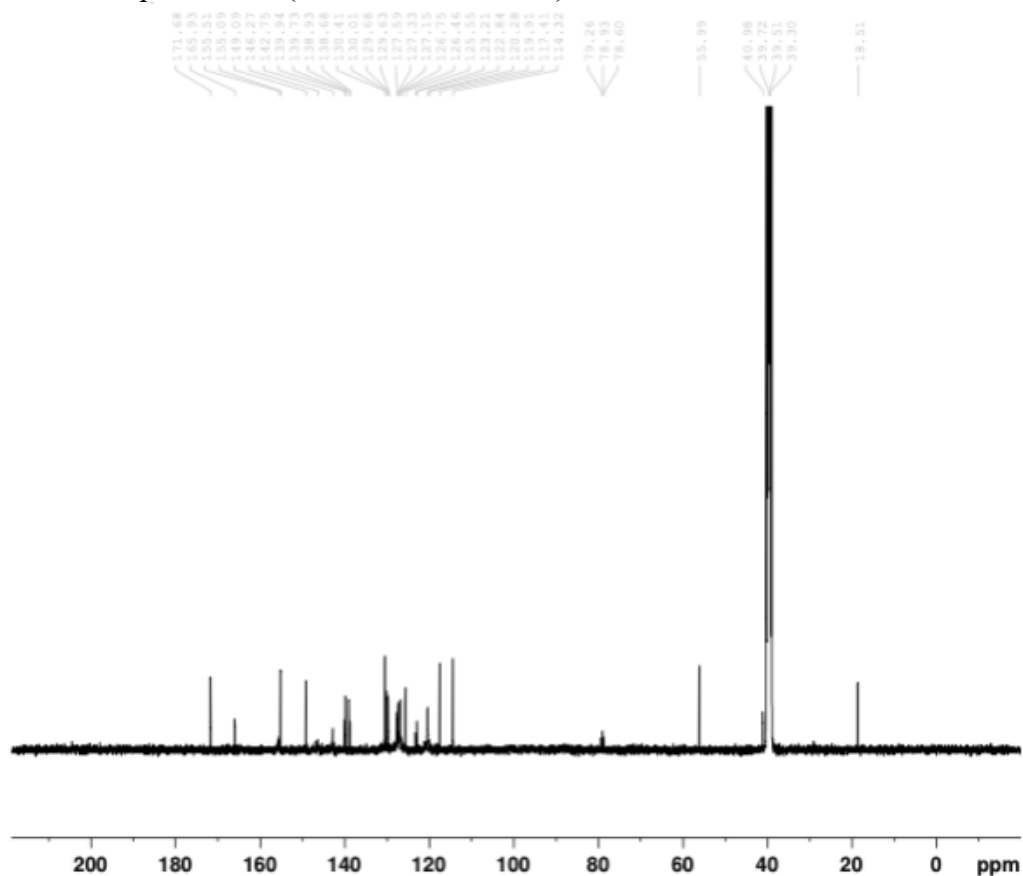
^{13}C NMR of compound **13m** (100MHz, $\text{DMSO}-d_6$)



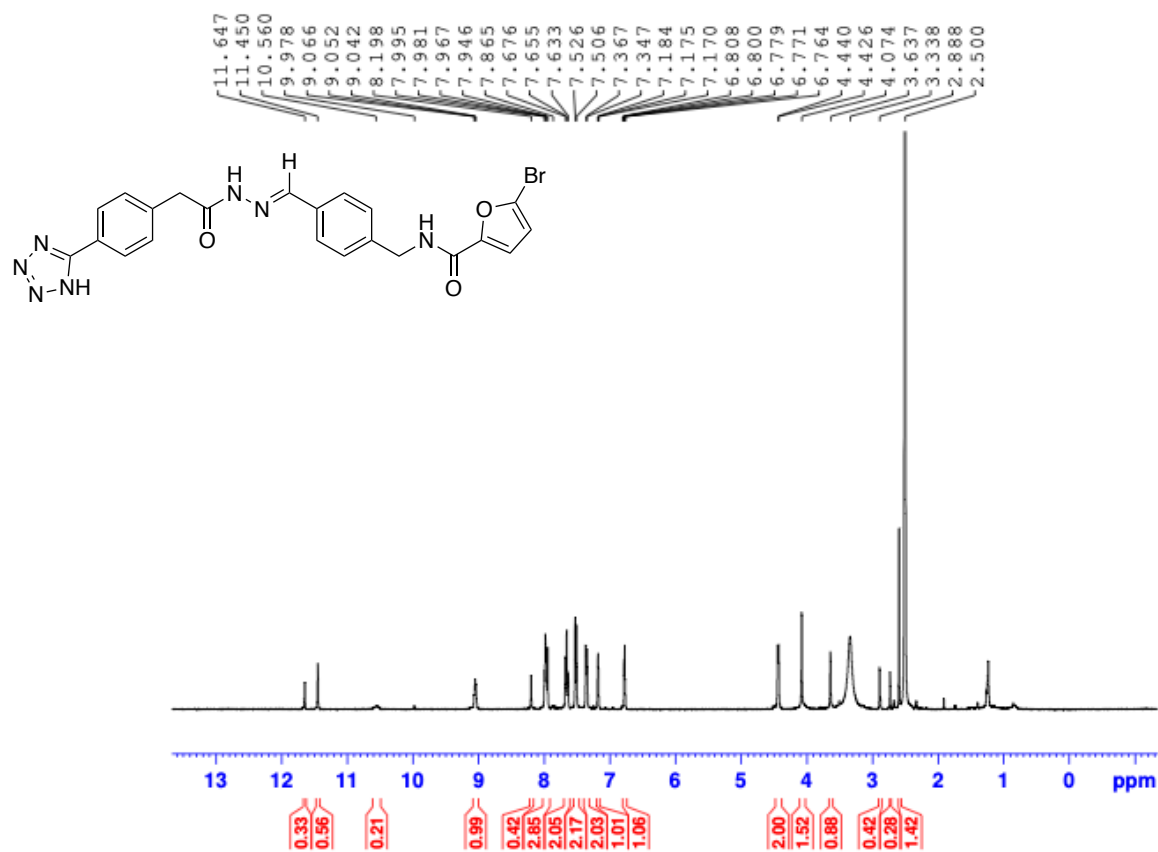
^1H NMR of compound **13n** (400MHz, DMSO- d_6)



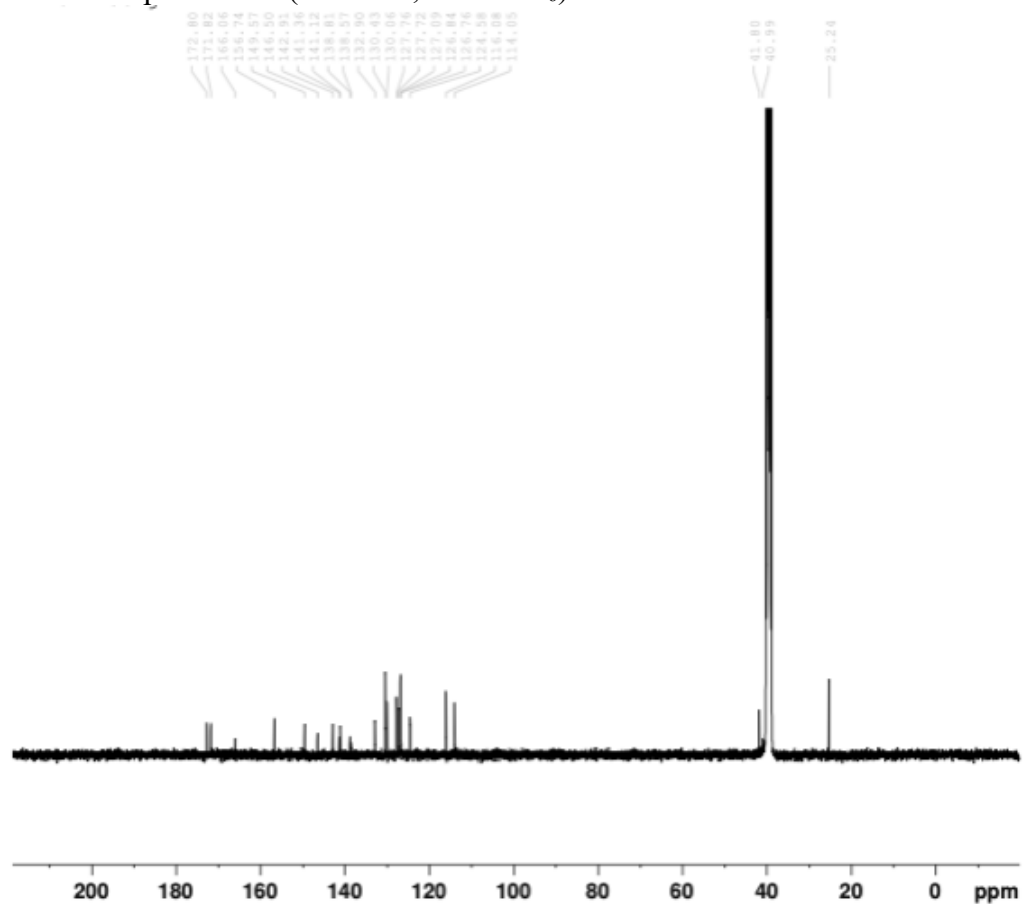
^{13}C NMR of compound **13n** (100MHz, DMSO- d_6)



^1H NMR of compound **13o** (400MHz, DMSO- d_6)



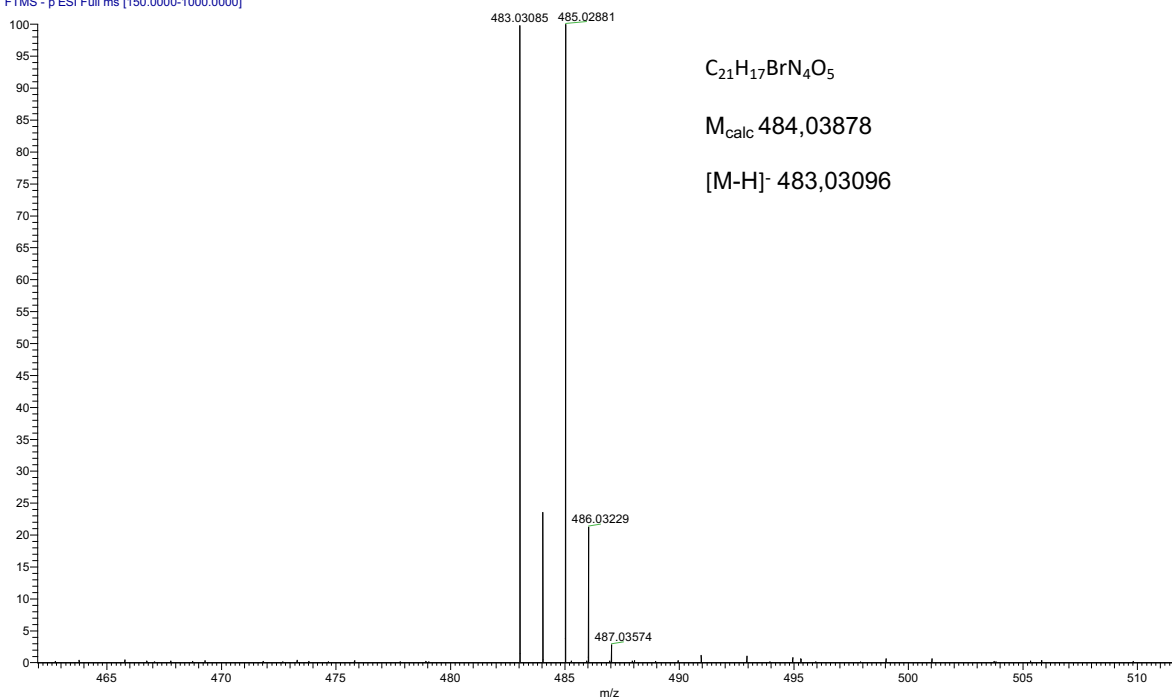
^{13}C NMR of compound **13o** (100MHz, DMSO- d_6)



V. Representative HRMS spectra

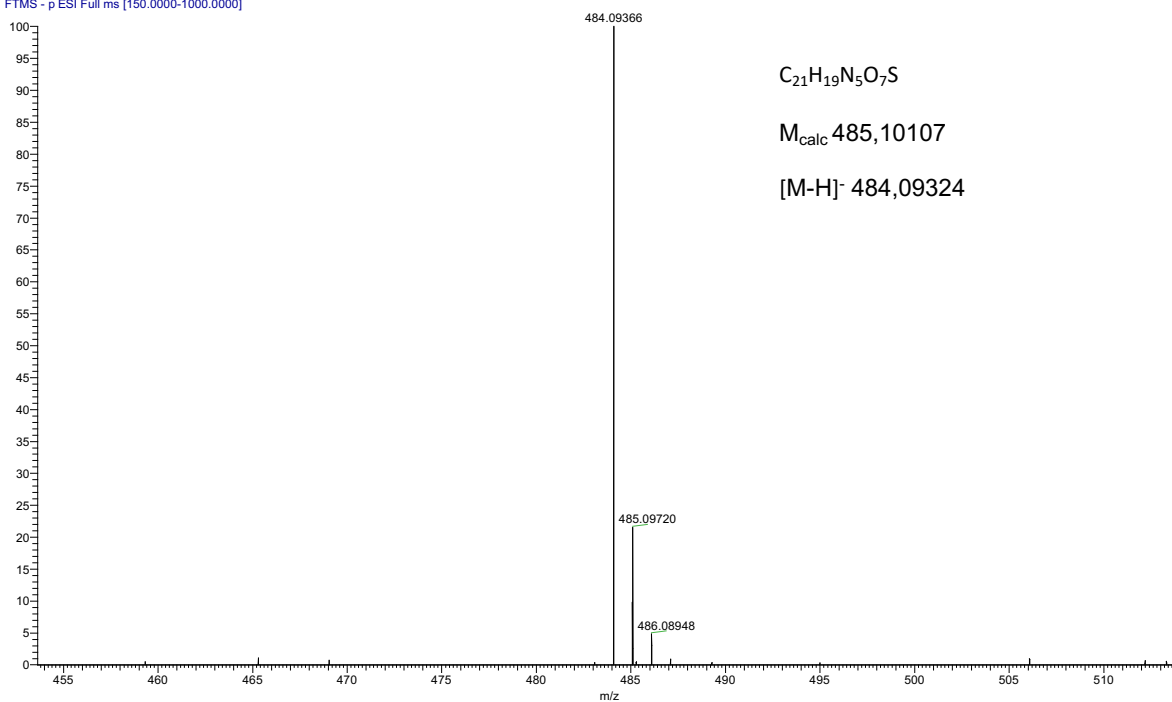
HRSM spectrum of compound **13**

AK #189 RT: 1.85 AV: 1 NL: 1.66E8
T: FTMS - p ESI Full ms [150.0000-1000.0000]



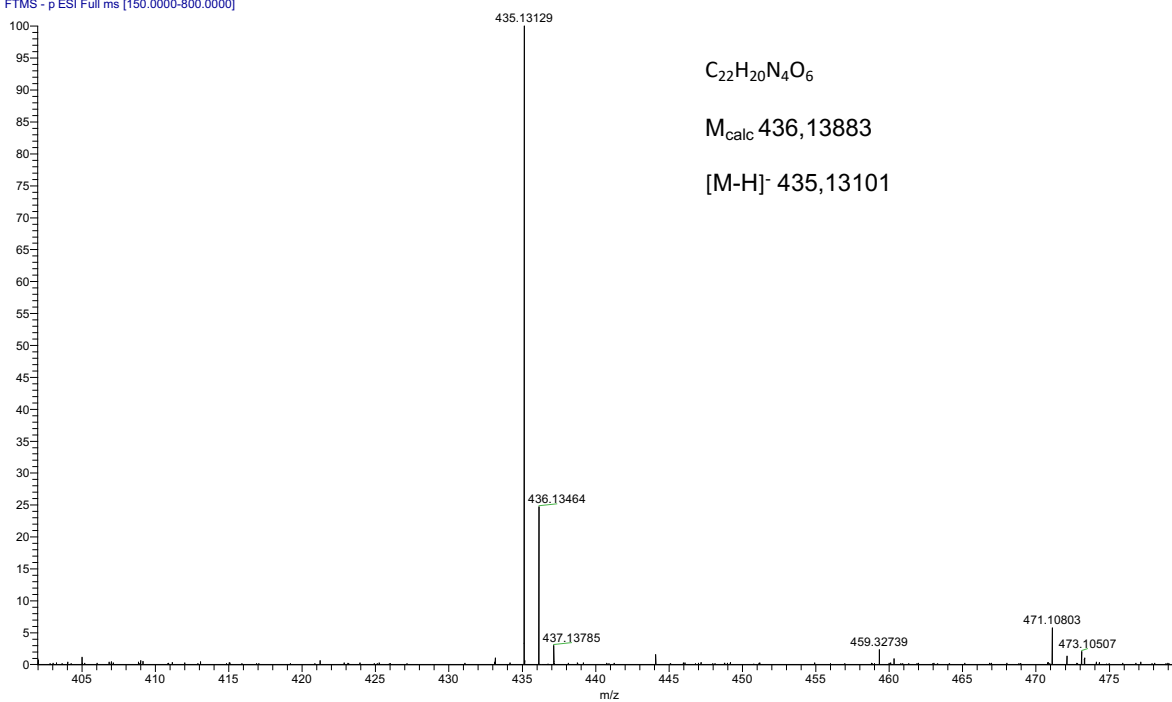
HRSM spectrum of compound **13a**

DC298 #1 RT: 0.01 AV: 1 NL: 6.35E6
T: FTMS - p ESI Full ms [150.0000-1000.0000]



HRSM spectrum of compound **13b**

BLB50 #389 RT: 2.14 AV: 1 NL: 1.54E8
T: FTMS - p ESI Full ms [150.0000-800.0000]



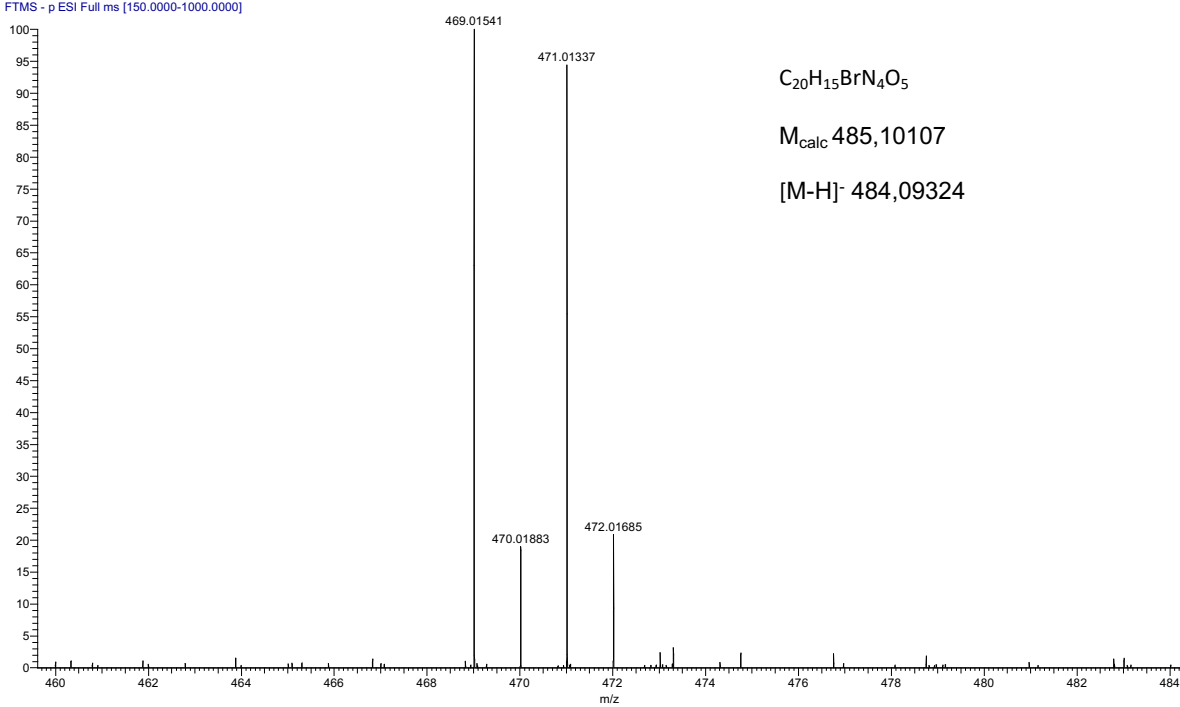
$C_{22}H_{20}N_4O_6$

M_{calc} 436,13883

$[M-H]^-$ 435,13101

HRSM spectrum of compound **13c**

TF3 #92 RT: 0.89 AV: 1 NL: 6.32E7
T: FTMS - p ESI Full ms [150.0000-1000.0000]



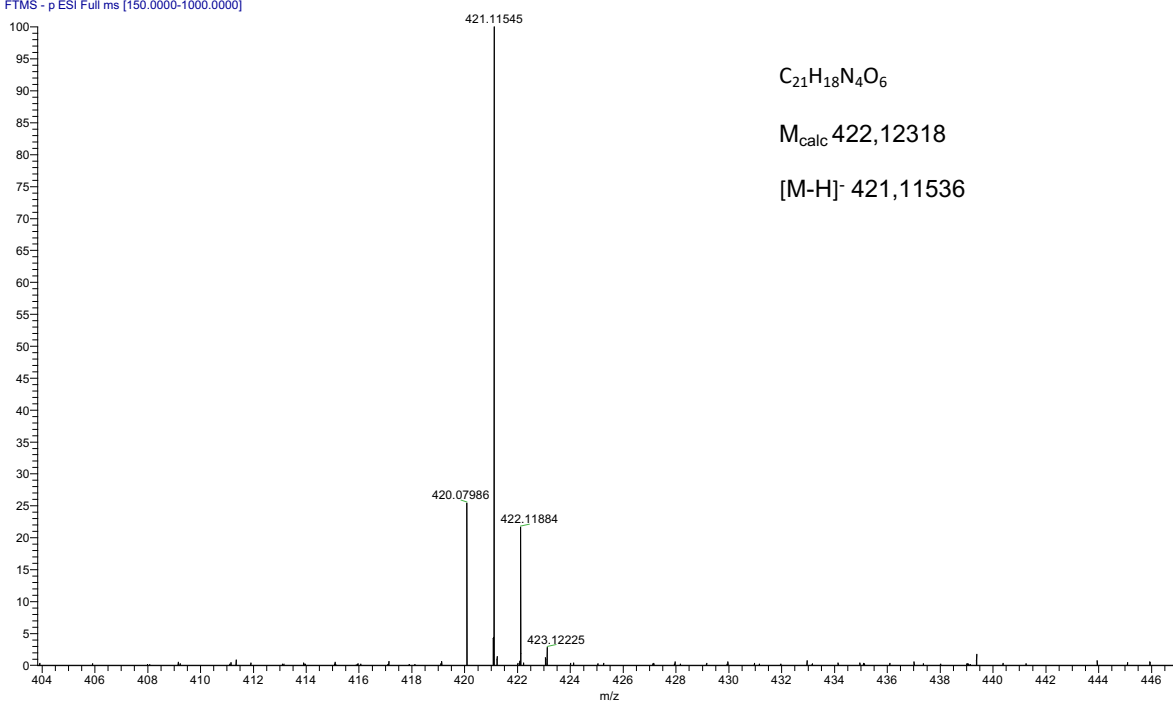
$C_{20}H_{15}BrN_4O_5$

M_{calc} 485,10107

$[M-H]^-$ 484,09324

HRSM spectrum of compound **13d**

TF7 #23 RT: 0.21 AV: 1 NL: 1.55E8
T: FTMS - p ESI Full ms [150.0000-1000.0000]



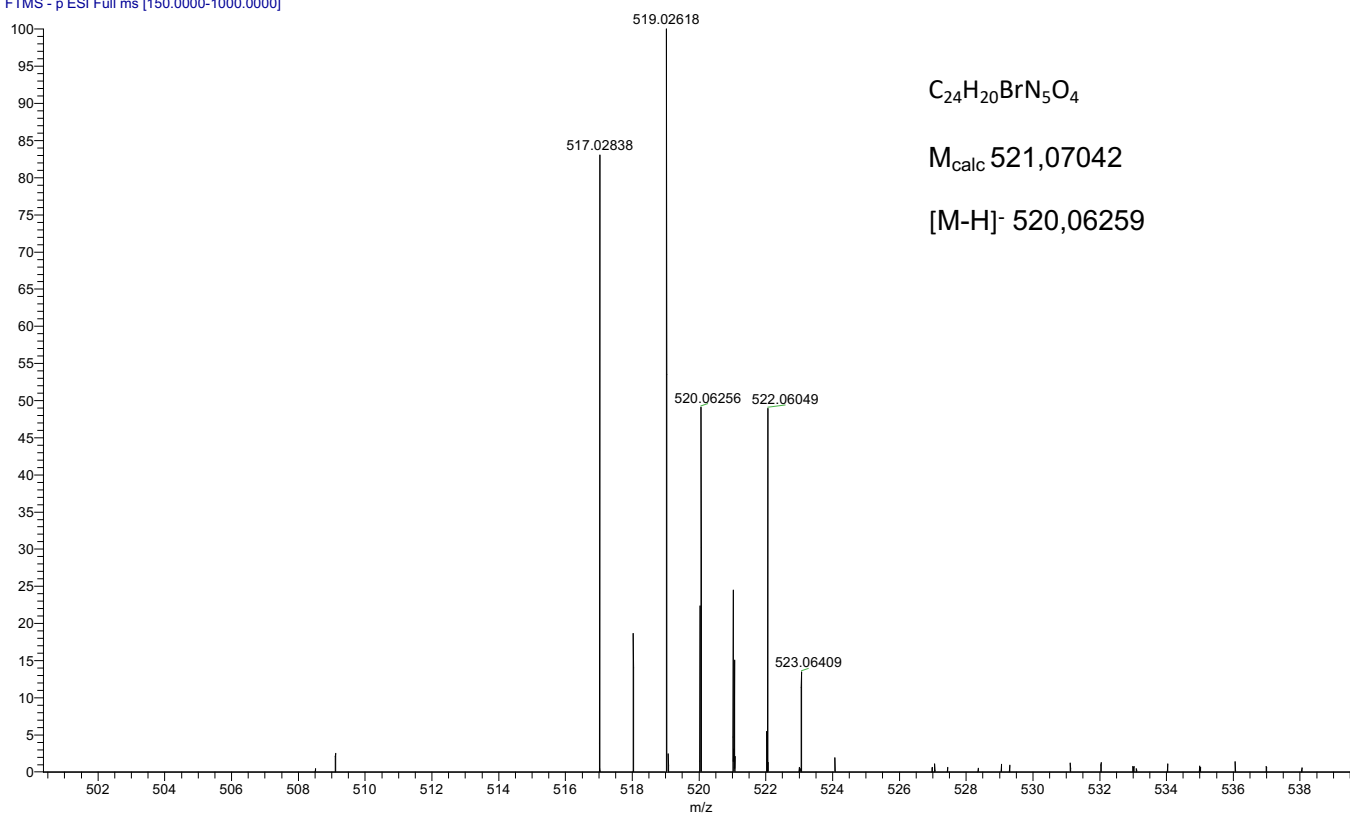
$C_{21}H_{18}N_4O_6$

M_{calc} 422,12318

$[M-H]^-$ 421,11536

HRSM spectrum of compound **13e**

BLB51 #26 RT: 0.25 AV: 1 NL: 6.21E7
T: FTMS - p ESI Full ms [150.0000-1000.0000]



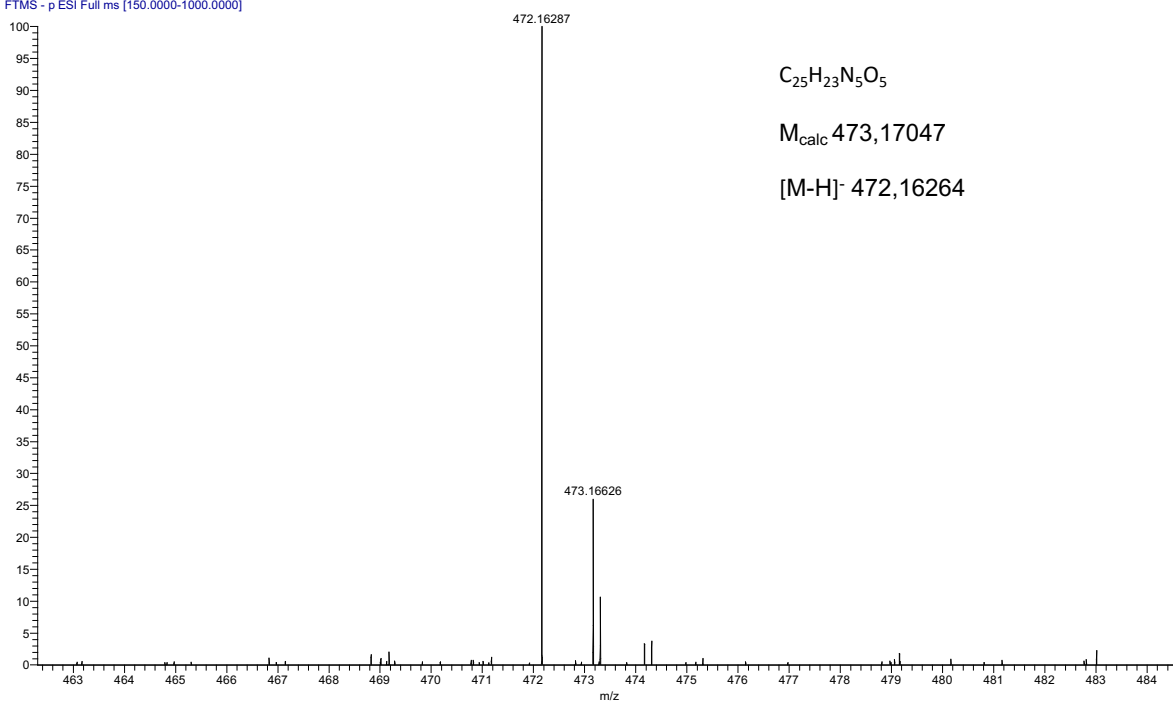
$C_{24}H_{20}BrN_5O_4$

M_{calc} 521,07042

$[M-H]^-$ 520,06259

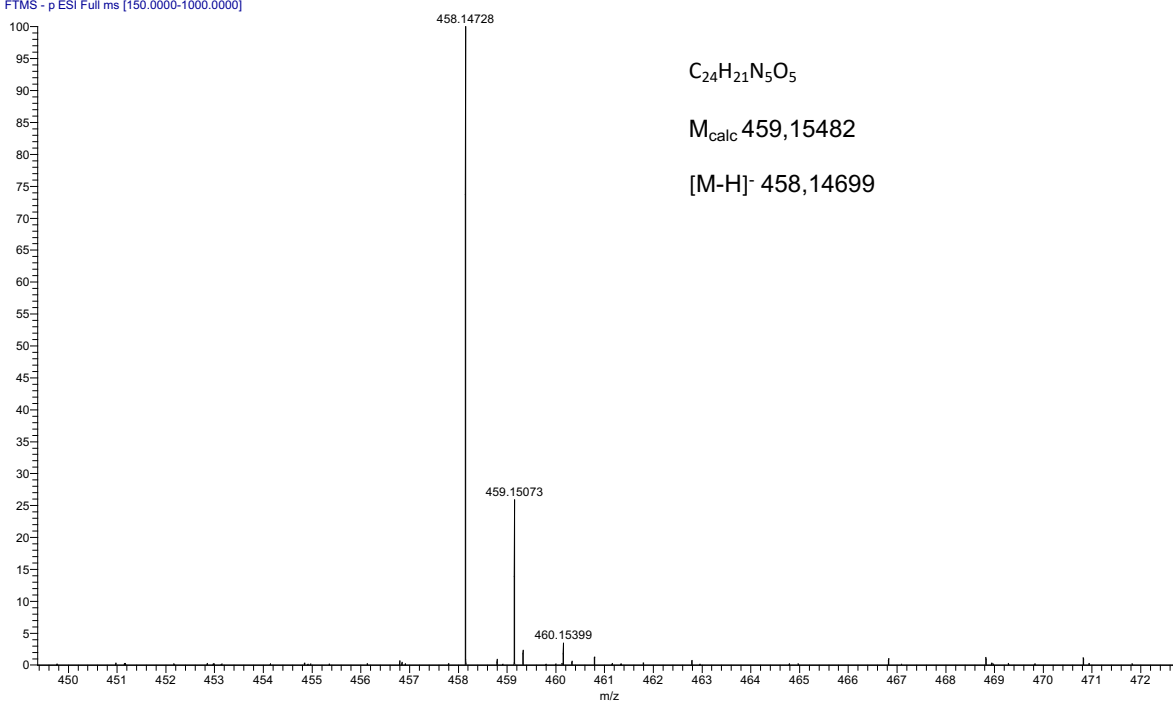
HRSM spectrum of compound **13f**

DC294 #5 RT: 0.05 AV: 1 NL: 6.17E7
T: FTMS - p ESI Full ms [150.0000-1000.0000]



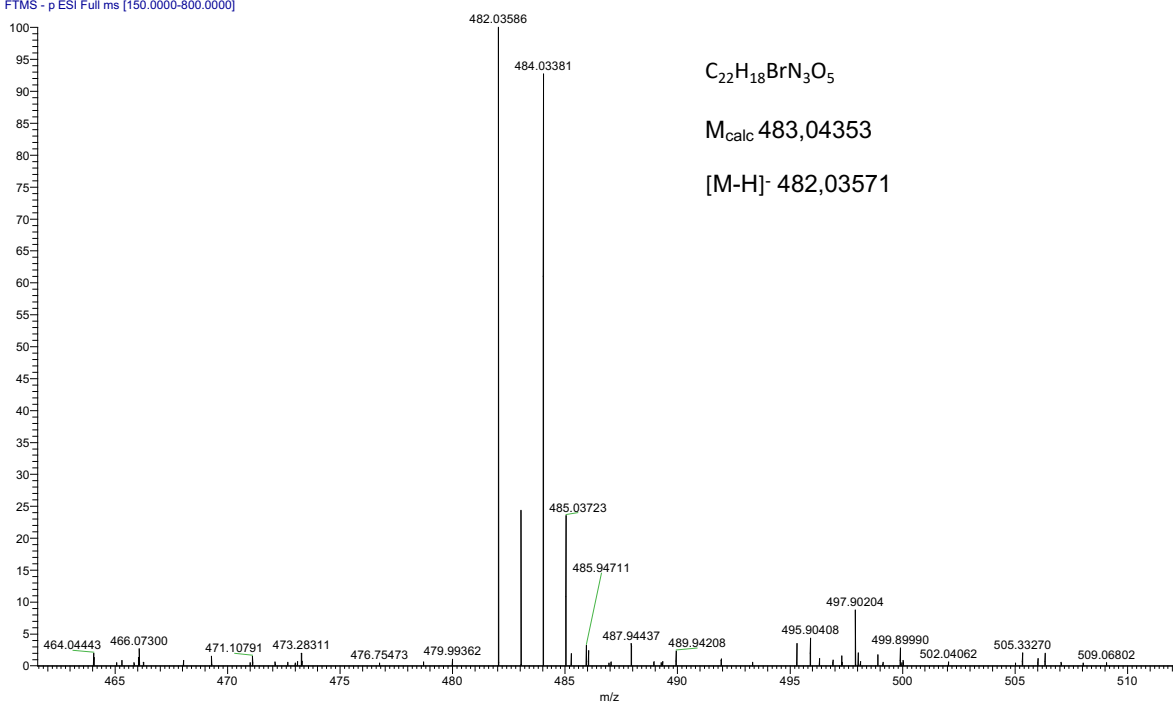
HRSM spectrum of compound **13g**

TF14 #1 RT: 0.01 AV: 1 NL: 1.58E8
T: FTMS - p ESI Full ms [150.0000-1000.0000]



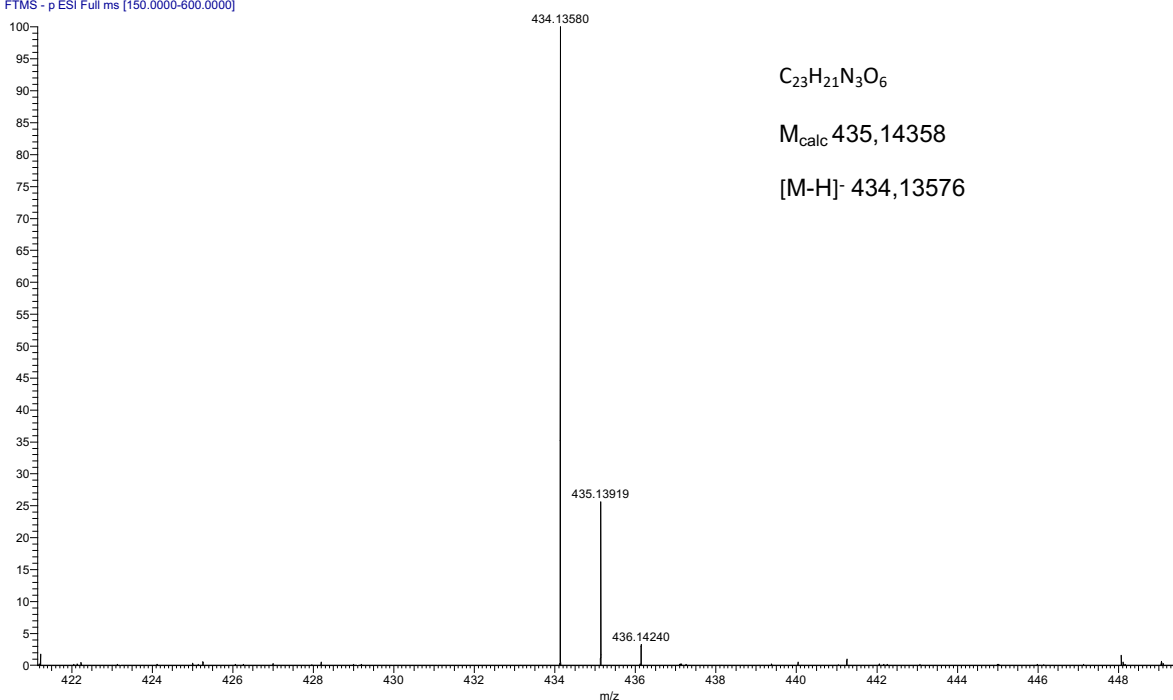
HRSM spectrum of compound 13h

BLB18 #20 RT: 0.20 AV: 1 NL: 3.27E7
T: FTMS - p ESI Full ms [150.0000-800.0000]



HRSM spectrum of compound **13i**

DC295 #1 RT: 0.01 AV: 1 NL: 1.34E8
T: FTMS - p ESI Full ms [150.0000-600.0000]



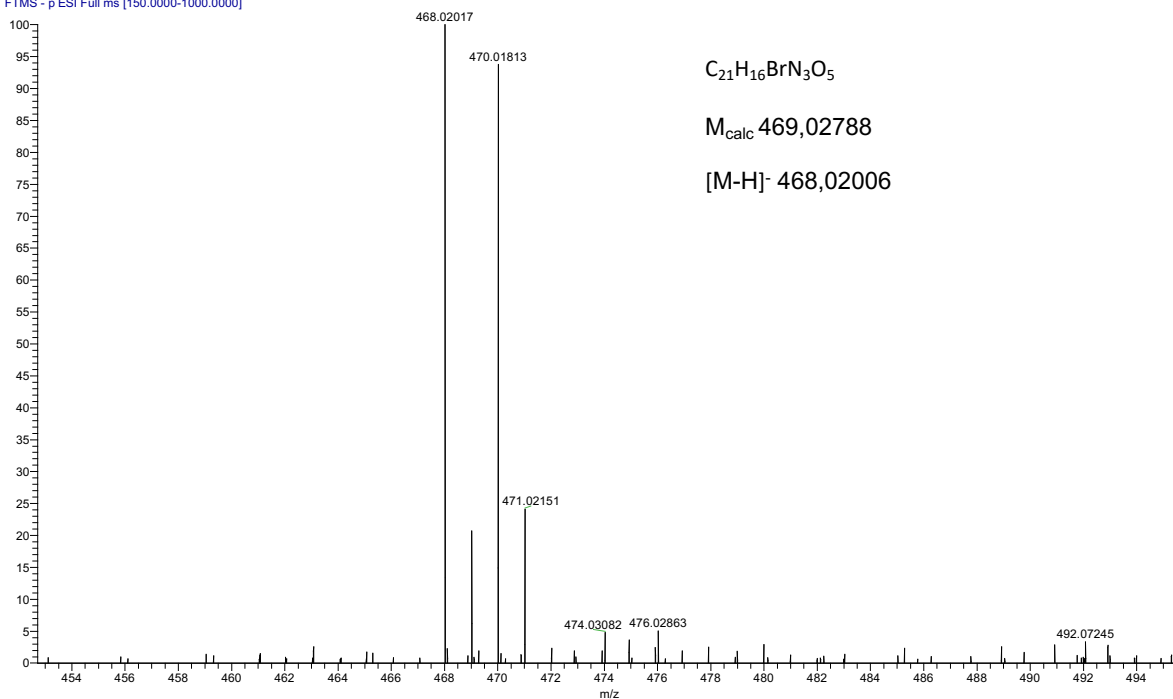
$C_{23}H_{21}N_3O_6$

M_{calc} 435,14358

$[M-H]^-$ 434,13576

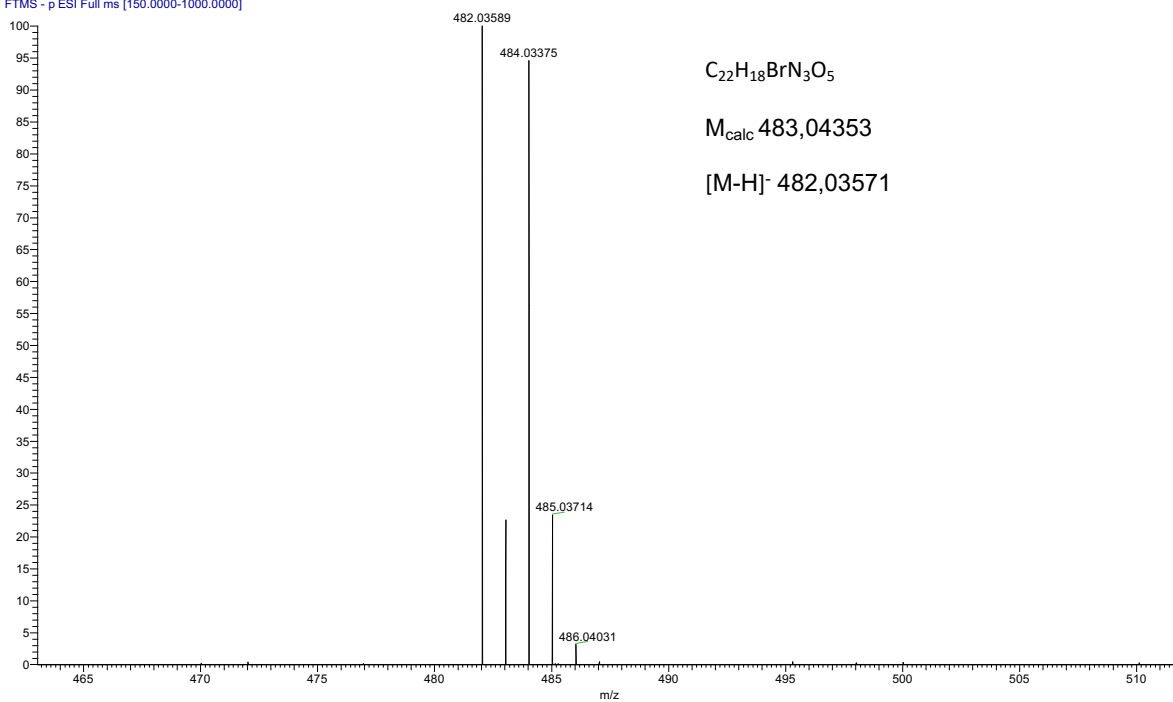
HRSM spectrum of compound **13l**

BLB19 #619 RT: 5.95 AV: 1 NL: 4.13E7
T: FTMS - p ESI Full ms [150.0000-1000.0000]



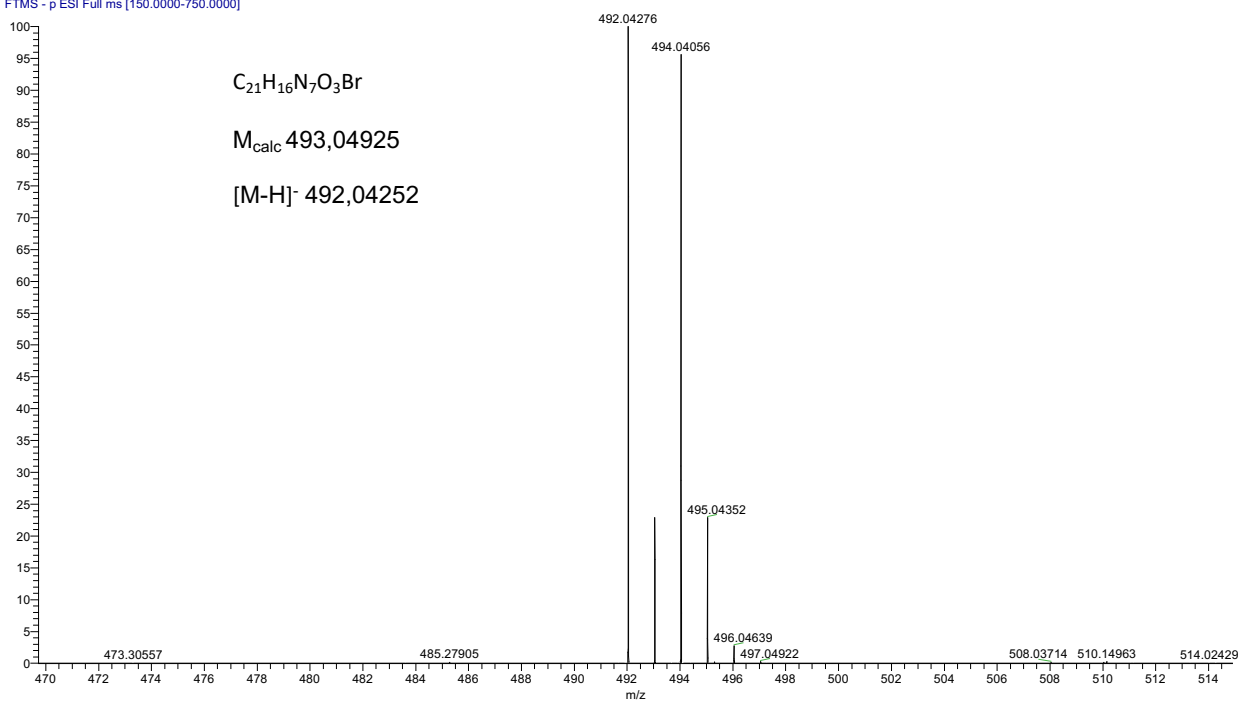
HRSM spectrum of compound 13m

RDL124 #42 RT: 0.40 AV: 1 NL: 4.36E8
T: FTMS - p ESI Full ms [150.0000-1000.0000]



HRSM spectrum of compound **13n**

BLB-29 #46 RT: 0.20 AV: 1 NL: 6.19E8
T: FTMS - p ESI Full ms [150.0000-750.0000]



HRSM spectrum of compound **13o**

RDL157 #215 RT: 1.54 AV: 1 NL: 2.50E8
T: FTMS - p ESI Full ms [150.0000-1500.0000]

