

## SUPPORTING INFORMATION

### Protein tyrosine phosphatase inactivation by electrophilic tyrosine modification

Madeleine L. Ware<sup>1</sup>, David M. Leace<sup>1,2</sup>, Zihan Qu<sup>3</sup>, Quentin Schaefer<sup>1</sup>, Sagar D. Vaidya<sup>1</sup>, Mikayla L. Horvath<sup>1</sup>, Zhihong Li<sup>1</sup>, Yunpeng Bai<sup>4</sup>, Zhong-Yin Zhang<sup>\*3,4</sup> and Ku-Lung Hsu<sup>\*1</sup>

<sup>1</sup>Department of Chemistry, University of Texas at Austin, Austin, TX, 78712, USA

<sup>2</sup>Department of Pharmacology, University of Virginia School of Medicine, Charlottesville, VA, 22908, USA.

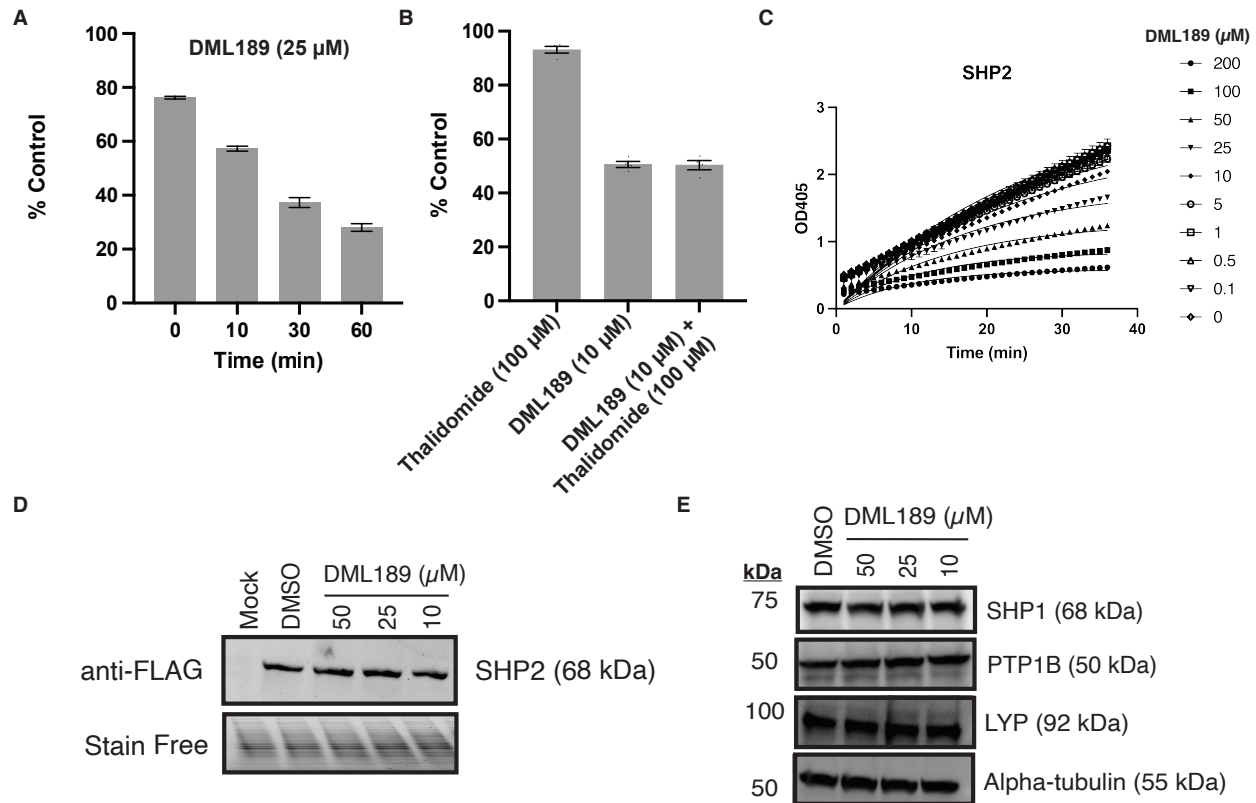
<sup>3</sup>James Tarpo Jr. and Margaret Tarpo Department of Chemistry, <sup>4</sup>Borch Department of Medicinal Chemistry and Molecular Pharmacology, Purdue University, West Lafayette, IN, 47907, USA

\*Corresponding author. Email: zhang-zy@purdue.edu (Z.-Y.Z.); ken.hsu@austin.utexas.edu (K.-L.H.)

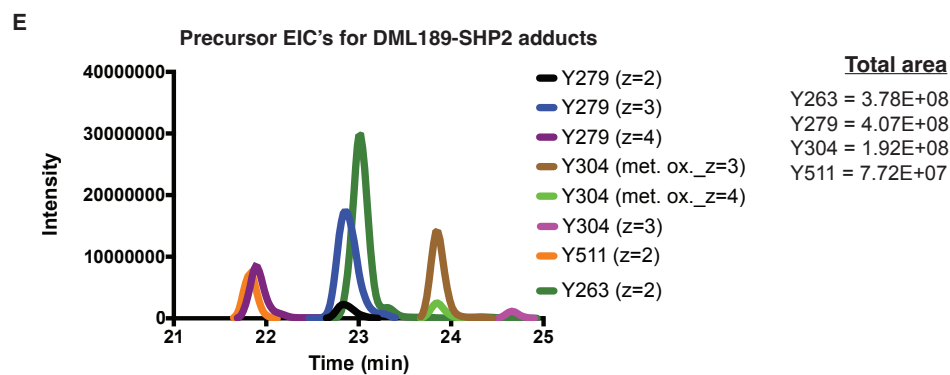
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## 1. SUPPORTING FIGURES

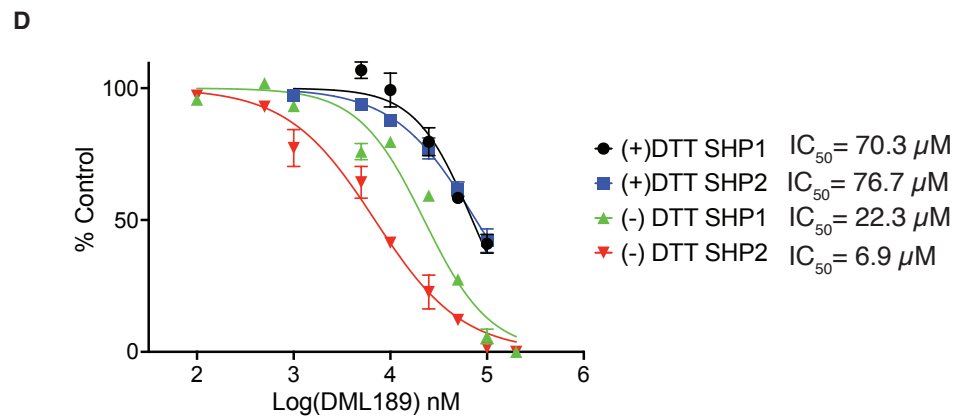
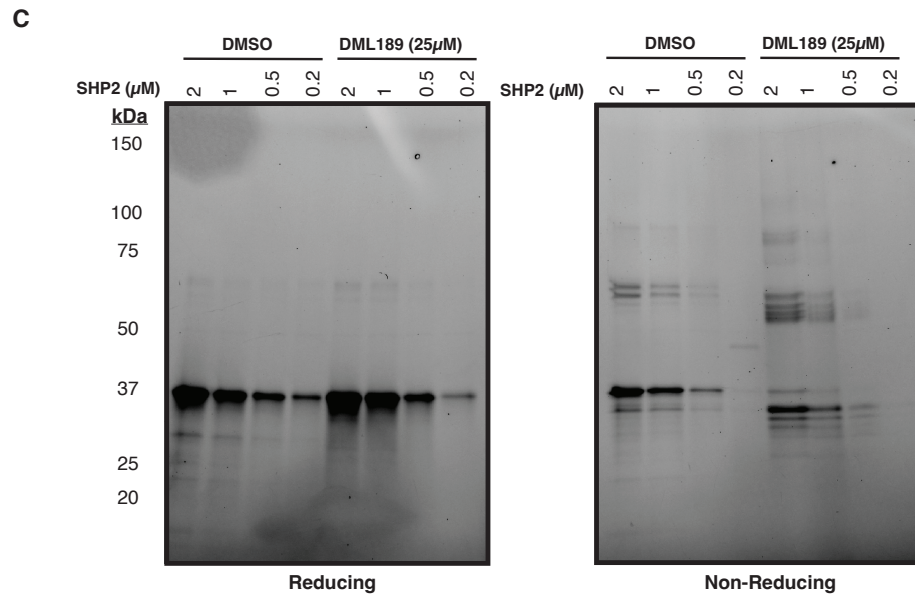
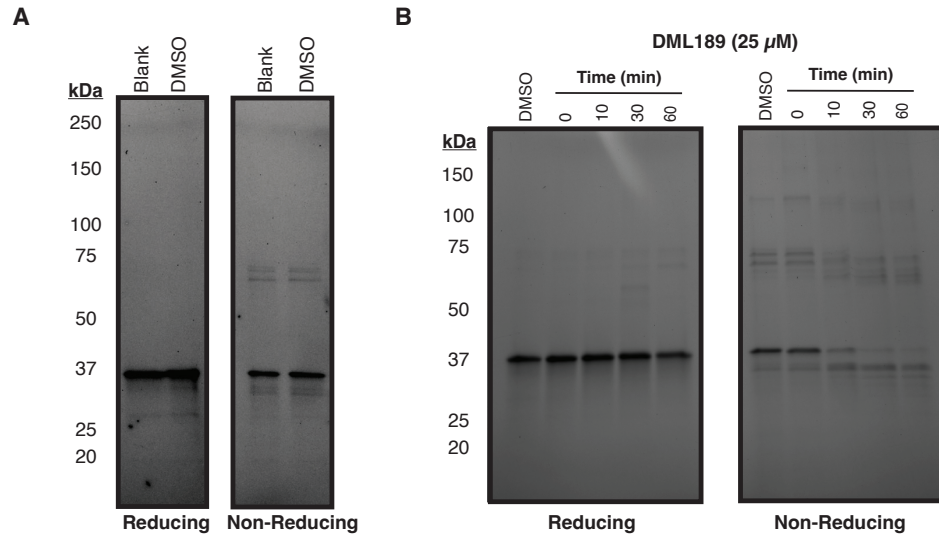


**Supplementary Figure 1.** (A) Time dependent inhibition (% vehicle control) of SHP2 by DML189 (25  $\mu$ M). Data shown as mean  $\pm$  SEM of % control from n =3 replicates. (B) SHP2 activity after treatment with thalidomide alone (100  $\mu$ M, 1 h) and after co-incubation with DML189 (10  $\mu$ M) and thalidomide (100  $\mu$ M, 1 h). Data shown as mean  $\pm$  SEM of % control from n =4 replicates. (C) SHP2 and pNPP (20 mM) reaction progress curves with DML189 (200-0.1  $\mu$ M) treatment. Data are from n=3 replicates. (D) Western blot analysis of full length SHP2 protein levels after DML189 dose response (0-50  $\mu$ M, 24 h) in live HEK 293T cells overexpressing recombinant SHP2. (E) Western blot analysis of SHP1, PTP1B, and LYP protein levels after DML189 treatment (0-50  $\mu$ M, 24 h) in live Jurkat cells.



**Supplementary Figure 2.** (A) Bar plot of % protein sequence coverage of PTP catalytic domains after trypsin digest and bottom-up LC-MS/MS analysis. (B-D) HCD spectra for SHP2-DML189 adducts at Y263, Y304 and Y511. Peptide spectra met the following quality control; Byonic score  $\geq 300$ ,  $-5 \leq \text{ppm err} \leq 5$ , delta mod score  $\geq 20$ . (E) Extracted ion chromatograms (EICs) of detected DML189-SHP2 precursor adducts from a single sample injection. Precursor areas were integrated using Proteome Discoverer 3.0.





**Supplementary Figure 3.** (A) SHP2 was incubated with and without DMSO for 1 h at room temperature and analyzed by reducing and non-reducing SDS-PAGE. (B) Reducing and non-reducing SDS-PAGE analysis of DML189 (25  $\mu$ M) incubated with SHP2 for 0, 10, 30, and 60 min. (C) Reducing and non-reducing SDS-PAGE analysis of DML189 (25  $\mu$ M) incubated with the indicated concentrations of SHP2. (D) Reversal of DML189 inhibitory activity against SHP1 and SHP2 after treatment with DTT (5 mM, 30 min) as measured by pNPP substrate assay.

## 2. CHEMICAL SYNTHESIS

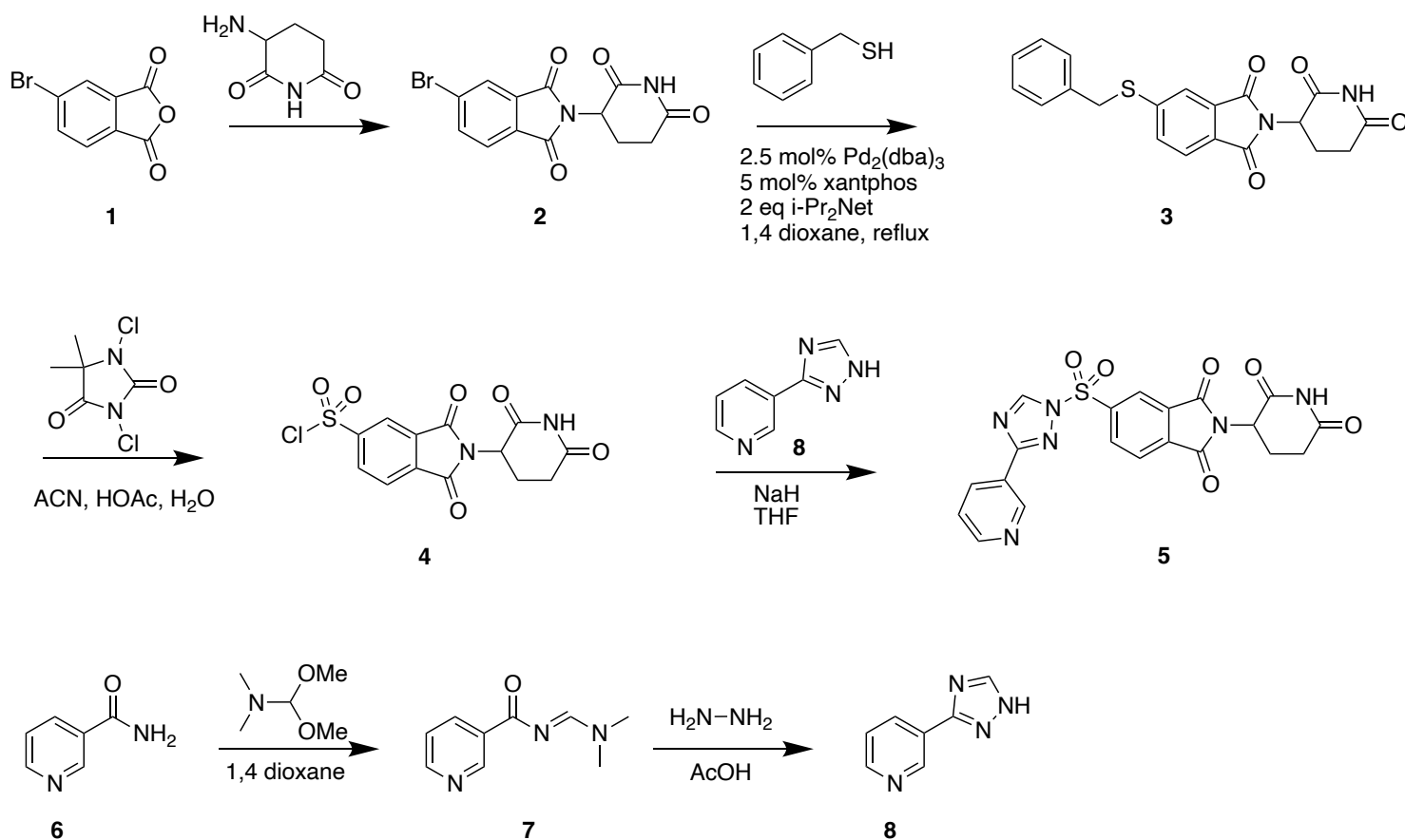
### General synthetic procedure for 1, 2, 3 sulfonyl triazoles

Synthesis was performed as previously described.<sup>1</sup>

### General synthetic procedure for 1, 2, 4 sulfonyl triazoles

Synthesis was performed as previously described.<sup>2</sup>

### Scheme 1. Synthesis of DML189



### **5-Bromo-2-(2,6-dioxo-3-piperidinyl)-1*H*-isoindole-1,3(2*H*)-dione (2)**

10 g (60.8 mmol, 1.2 equiv) of aminoglutarimide HCl was dissolved in 550 mL of glacial acetic acid followed by addition of 11.5 g (50.7 mmol, 1 equiv) of bromo phthalic anhydride **1**. Next, 17.8 mL (127.7 mmol, 2.1 equiv) of triethylamine was added slowly, and the mixture was heated to reflux for 2h and monitored for starting material depletion. At completion, the reaction was cooled and poured onto ice. The precipitate was filtered through a glass frit, washed with cold water, and dried to yield 15.3 g of a purple powder (75%).

### **1*H*-Isoindole-1,3(2*H*)-dione, 2-(2,6-dioxo-3-piperidinyl)-5-[(phenylmethyl)thio]- (3)<sup>3</sup>**

**2** (4.37g, 13 mmol, 1 equiv), benzyl mercaptan (1.53 mL, 91 mmol, 7 equiv), Pd<sub>2</sub>(dba)<sub>3</sub> (299 mg, 0.325 mmol, 0.025 equiv), xantphos (377 mg, 0.65 mmol, 0.05 equiv), and *i*-Pr<sub>2</sub>Net (4.6 mL, 26 mmol, 2 equiv) were dissolved in 27 mL of 1,4 dioxane and heated to reflux overnight. The next morning, the reaction mixture was cooled and dried on celite. The material was loaded on a biotage column and eluted with 0-50% EtOAc/DCM. The product was collected, precipitated, and washed with DCM to yield an off white solid (3.37g, 66% yield).

### **1*H*-Isoindole-5-sulfonyl chloride, 2-(2,6-dioxo-3-piperidinyl)-2,3-dihydro-1,3-dioxo- (4)<sup>4</sup>**

**3** (1g, 2.5 mmol, 1 equiv), acetic acid (0.94 mL), and water (0.625 mL) were added to 25 mL of ACN and cooled to -10 °C. DCDMH (0.99g, 5 mmol, 2 equiv) was then added slowly. After 1 hour the reaction mixture was diluted with 25 mL of DCM and cooled to 0°C. 25 mL NaHCO<sub>3</sub> was then added, and the solution was stirred for 10 minutes. The organic layer was then extracted, washed with brine, dried over Na<sub>2</sub>SO<sub>4</sub>, concentrated, precipitated, and filtered. After washing with cold DCM, the product was obtained as a bright white powder. H-NMR showed the product with presence of the hydrolyzed sulfonic acid.

### **3-(1*H*-1,2,4-Triazol-5-yl)pyridine (8)**

Synthesis was performed as previously described.<sup>5</sup>

### **2-(2,6-Dioxopiperidin-3-yl)-5-((3-(pyridine-3-yl)-1*H*-1,2,4-triazol-1-yl)sulfonyl)isoindoline-1,3-dione (5, DML189)**

To a flame dried flask was dissolved 1, 2, 4 triazole **8** (82 mg, 0.56 mmol, 1 equiv) in 2.5 mL of dry THF. The solution was cooled to 0° C and NaH (22.4 mg, 0.56 mmol, 1 equiv) was added. The reaction was stirred for 30 minutes after which sulfonyl chloride **4** (200 mg, 0.56 mmol, 1 equiv) was added and the reaction was brought to room temperature and stirred overnight. The next morning, the precipitate was filtered, loaded onto celite, and purified with 0-20% MeOH/DCM on a biotage to yield a white solid (170 mg, 65% yield).

### 3. BIOLOGICAL METHODS

#### Purification of PTP catalytic domains

The cDNA encoding the catalytic domain of SHP1 (residues 245-543) was amplified by PCR and subcloned into pET21a(+) bacterial expression vector to allow expression of N-terminal His-tagged proteins. *E. coli* BL21(DE3) (Novagen) was used as an expression host and the induction of protein expression was carried out in LB Broth with 1 mM IPTG at 18 °C overnight. Cell pellets were stored at –80 °C for subsequent protein purification. Frozen cell pellets were lysed by sonication in 40 mL cold lysis buffer (50 mM Tris-HCl, pH 8.0, 150 mM NaCl, 5 mM imidazole, and 1 mM PMSF) per liter cell pellet. Cell lysates were clarified by centrifugation using a Beckman JA-18 rotor for 15 min at 5308 x g. The supernatant was incubated with HisPur Ni-NTA resin (Thermo Scientific) for 2 h and then packed onto a column and washed with 50 resin volume of buffer A (50 mM Tris-HCl, pH 8.0, 500 mM NaCl, 5 mM imidazole). The His-tagged protein was eluted with Buffer B (50 mM Tris-HCl, pH=8.0, 500 mM NaCl, 300 mM imidazole). Pooled His-protein-containing fractions were concentrated, loaded onto a HiLoad 26/600 Superdex 75 column (GE Healthcare Biosciences) and eluted with storage buffer (50 mM Tris-HCl, pH 8.0, 150 mM NaCl, 1 mM DTT, 10% glycerol). The purity of the protein was determined to be >95% by SDS-PAGE and Coomassie staining. The protein was aliquoted and stored at –80 °C. The catalytic domain of SHP2, PTP1B, and LYP were expressed and purified as described previously<sup>6-9</sup>.

#### PTP activity assay for SuTEx ligand screen

SuTEx compounds were dissolved in DMSO to make 50 mM stock and transferred to assay plate. Ligands (500 μM) were incubated with SHP1/SHP2/PTP1B/LYP (400 nM) in assay buffer (50 mM 3,3-dimethylglutaric acid, pH 7.0, 18 mM NaCl, 1 mM EDTA, 0.01% Triton X-100) for 30 min to allow covalent labeling. Excessive substrate para-nitrophenol phosphate (pNPP, 20 mM) was added to the mixture and incubated for 2/2/0.5/10 min, respectively to monitor remaining enzyme activity.

#### DML189 dose responses and time dependence assays

0.025 nmols of PTP catalytic domain in assay buffer (50 mM Tris, 150 mM NaCl, pH 7.5) was treated with ligands (0-200 μM, 1h). pNPP (20 mM) was then added and the enzyme activity monitored by reading absorbance at 405nm. The above assay was performed with SHP2 treated with DML189 (25μM) where enzyme activity was monitored at 0, 10, 30, and 60 min.

#### DML189-SHP2 $k_{inact}$ /K<sub>I</sub> calculation

DML189 (0-500 μM), PTP catalytic domain (0.025 nmols), and PNPP (20 mM) were incubated together in assay buffer (50 mM Tris, 150 mM NaCl, pH 7.5) and monitored for product formation using a plate reader for at least 30 min. Reaction progress curves were then fitted to the equation  $Y=(V_0/k_{obs})(1-\exp(-k_{obs}X))$  in order to obtain  $k_{obs}$  values for each concentration of DML189. These  $k_{obs}$  values are then plotted against compound concentration.  $k_{inact}/K_I$  was calculated using the equation  $k_{obs} = k_{inact}*[I]/(K_I+[I])$ .

#### Assessment of thalidomide activity against SHP2

0.025 nmols of SHP2 catalytic domain in assay buffer (50 mM Tris, 150 mM NaCl, pH 7.5) was treated with thalidomide (100  $\mu$ M, 1h) and enzyme activity was evaluated using pNPP (20 mM) as a substrate. Additionally, the above assay was performed after SHP2 was pre-incubated with thalidomide (100  $\mu$ M) followed by DML189 (10  $\mu$ M) for 1h at RT.

### **Testing DML189 degradative activity against PTPs**

HEK293T cells were transiently transfected with full length SHP2 using polyethylenimine (PEI). After 48h, cells were treated with DML189 (50, 25, 10  $\mu$ M) in serum free media for 24h. Cells were harvested and washed 3x in 1x DPBS. Jurkat cells were treated (DML189 (50, 25, 10  $\mu$ M), 10<sup>6</sup>/mL) and harvested to test for LYP, SHP1, and PTP1B degradation. Following whole cell lysis (50 mM Tris, 150 mM NaCl, pH 7.5, 1% Triton), lysates were normalized to 2 mg/mL and analyzed for PTP degradation via western blot: SHP2 (anti-FLAG, Sigma, Rabbit, 1:2500), SHP1 (proteintech, Cat No. 24546-1-AP, 1:2000), PTP1B (proteintech, Cat No. 11334-1-AP), LYP (proteintech, Cat No. 11783-1-AP, 1:500).

### **Docking**

A structure of SHP2 (PDB ID: 3ZM0) was obtained from PDB website. Residues within 10 Å of the binding site were defined as the docking pocket. Subsequently, compound was docked into this defined pocket using GOLD 5.1 with default parameters and 100 genetic algorithm (GA) runs. For each GA run, a maximum of 125,000 operations were performed. Docking was terminated when the top ten solutions had RMSD values within 1.5 Å. The resulting protein–ligand complexes were visualized using the PyMOL Molecular Graphics System (version 2.3.0).

### **Non-reducing / reducing SDS-PAGE of PTP catalytic domains**

Catalytic domains of PTPs (SHP1, SHP2, PTP1B or LYP (0.5 mg/mL) in assay buffer (50 mM Tris, 150 mM NaCl, pH 7.5) were treated for 1h at RT with DML189 (150  $\mu$ M) or DMSO. The samples were split in half and quenched with BME-containing or BME-free loading buffer and analyzed by stain free gel.

### **DML189 LIPT dose response on SHP2**

SHP2 in assay buffer (50 mM Tris, 150 mM NaCl, pH 7.5) at 0.132 mg/mL was treated for 1h at RT with DML189 (100-0.1  $\mu$ M) or DMSO. The samples were split in half and quenched with BME-containing or BME-free loading buffer and analyzed by stain free gel.

### **SHP2 dilution LIPT analysis with DML189**

Various concentrations of SHP2 (2-0.2  $\mu$ M) were treated with DMSO or DML189 (25  $\mu$ M) and analyzed by reducing/non-reducing SDS-PAGE followed by stain free imaging.

### **SHP2 thalidomide and H<sub>2</sub>O<sub>2</sub> LIPT response**

SHP2 in assay buffer (50 mM Tris, 150 mM NaCl, pH 7.5) at 0.132 mg/mL was treated at RT with thalidomide (100  $\mu$ M, 1h) or H<sub>2</sub>O<sub>2</sub> (500  $\mu$ M, 1h). The samples were split in half and quenched with BME-containing or BME-free loading buffer and analyzed by stain free gel.

### **DTT reversibility assay**

0.025 nmols SHP1 and SHP2 catalytic domain were treated with DML189 (0-100  $\mu$ M, 1h) in assay buffer (50 mM Tris, 150 mM NaCl, pH 7.5). After DML189 incubation, samples were treated with DTT (5 mM, 30 min) followed by PTP activity readout using pNPP (20 mM).

### **Sample preparation for DML189 adduct detection by LC-MS/MS**

1 nmol of SHP1, SHP2, LYP, and PTP1B catalytic domains were treated with DML189 (100  $\mu$ M, 1h, RT) in assay buffer (50 mM Tris, 150 mM NaCl, pH 7.4). Following treatment, samples were loaded onto a 10,000 Da MWCO filter and washed 3x with 25mM ammonium bicarbonate. After transfer to Eppendorf tubes, samples were reduced (DTT, 20 mM, 65°C, 15 min), alkylated (iodoacetamide, 40 mM, RT, 30 min), and trypsinized (3h, 37°C). Trypsin was quenched with 1 $\mu$ L acetic acid prior to injection (150 ng) on the LC-MS.

### **Sample preparation for crosslinking LC-MS/MS**

SHP1 and SHP2 (1 nmol) were treated with DML189 (75  $\mu$ M, 1h, RT) or DMSO and then immediately quenched with iodoacetamide (40mM, 30 min, RT) to prevent disulfide reshuffling. Samples were loaded onto a 10,000 Da MWCO filter and washed 3x with 25 mM ammonium bicarbonate. After washing, samples were trypsinized (1.5h, 37°C) and quenched with 1 $\mu$ L formic acid before injection onto the LC-MS.

### **LC-MS/MS acquisition and analysis for DML189 adduct detection on PTP catalytic domains**

Peptides from DML189 treated PTP's were evaluated by LC-MS/MS using a Vanquish Neo UHPLC coupled to an Orbitrap Eclipse Tribrid mass spectrometer. Peptides were separated on a 30 min gradient by reverse phase LC using 3  $\mu$ m C18 (20 cm) as follows: (A: 0.1% formic acid/H<sub>2</sub>O; B: 80% ACN, 0.1% formic acid in H<sub>2</sub>O): 0–1 min 15% B, 450 nL/min; 1–8.5 min 40% B, 300 nL/min; 8.5–16 min 60% B, 300 nL/min; 16–20 min 99% B, 300 nL/min; 20–30 min 1% B. Data were acquired using a top 20 ddMS2 method. MS1 spectra were acquired at 120K resolution with a max IT of 20 ms and MS2 spectra were taken with a normalized AGC of 50%, quadrupole isolation width of 1.6 m/z, and max IT of 100 ms. DML189 (+320.0181 Da) adducts on PTP domains were searched as modifications on tyrosine, lysine, histidine, and cysteine using Byonic. Peptides met the following quality criteria: PMI-Byonic Score  $\geq$  300, delta ppm err. 5, delta mod score  $\geq$  20.

### **Data acquisition and analysis for crosslinking LC-MS/MS.**

Non-reduced peptides from DML189 treated PTP's were evaluated by LC-MS/MS using a Vanquish Neo UHPLC coupled to an Orbitrap Eclipse Tribrid mass spectrometer. Peptides were separated on a 30 min gradient by reverse phase LC using 3  $\mu$ m C18 (20 cm) as follows: (A: 0.1% formic acid/H<sub>2</sub>O; B: 80% ACN, 0.1% formic acid in H<sub>2</sub>O): 0–1 min 15% B, 450 nL/min; 1–8.5 min 40% B, 300 nL/min; 8.5–16 min 60% B, 300 nL/min; 16–20 min 99% B, 300 nL/min; 20–30 min 1% B. Calibrated charge-state dependent ETD activation followed by a supplemental HCD at 30% CE was used to fragment disulfide-bound peptides. MS2 spectra were taken at 15k resolution, an absolute AGC of 1E5 and maximum injection time of 200 ms. Disulfide linked peptides were identified using Byonic S-S Xlink node (S-S cleavable linkage, –2.015650 Da) with the following quality control criteria: PMI-Byonic Score  $\geq$  300, delta ppm err. 5, and Xlink score of >100.

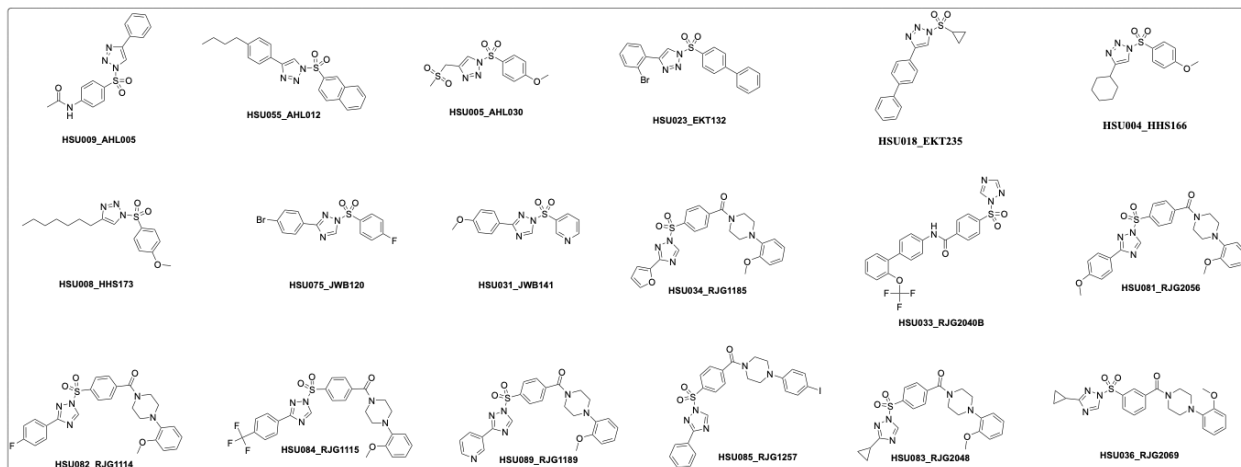
#### 4. REFERENCES

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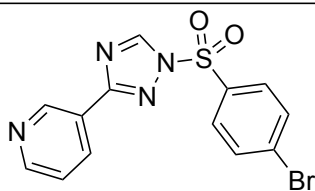


## 5. APPENDIX

**A. Published sulfonyl triazole compounds used for PTP screening. Synthesis and characterization are reported in References [1, 2, 5]:**



**B. Compound characterization of remaining sulfonyl-triazoles used in PTP screen:**



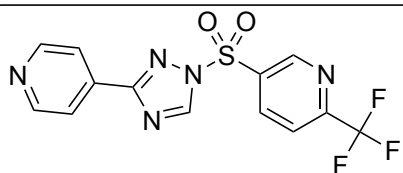
**HSU001-JWB179**

Chemical Formula:  $C_{13}H_{10}BrN_4O_2S$   
Exact Mass: 364.9708

**$^1H$  NMR** (800 MHz, DMSO- $D_6$ )  $\delta$  9.56 (s, 1H), 9.15 (d,  $J$  = 2.4 Hz, 1H), 8.70 (dd,  $J$  = 4.8, 1.7 Hz, 1H), 8.33 (dt,  $J$  = 7.9, 1.9 Hz, 1H), 8.10 – 8.07 (m, 2H), 7.98 – 7.94 (m, 2H), 7.55 (dd,  $J$  = 7.9, 4.8 Hz, 1H).

**$^{13}C$  NMR** (201 MHz, DMSO- $D_6$ )  $\delta$  162.06, 151.36, 148.15, 147.26, 134.34, 134.26, 133.47, 130.74, 130.24, 124.75, 124.23.

**HRMS** (ESI-TOF)  $m/z$ :  $[M + H]^+$  calculated for  $C_{13}H_{11}BrN_4O_2S$  364.9708, Found, 364.9695.



**HSU002\_JWB222**

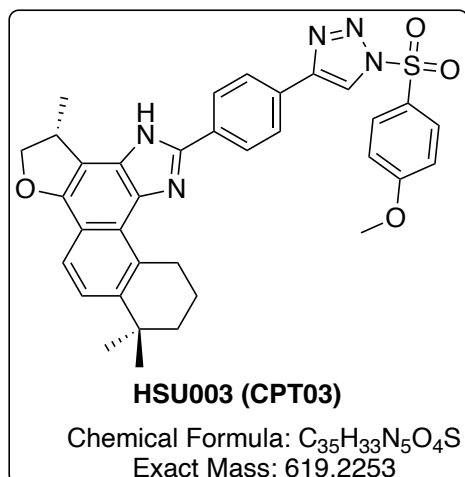
Chemical Formula:  $C_{13}H_8F_3N_5O_2S$   
Exact Mass: 355.0351

**$^1H$  NMR** (800 MHz, DMSO- $D_6$ )  $\delta$  9.63 (s, 1H), 9.51 (d,  $J$  = 2.3 Hz, 1H), 8.88 (dd,  $J$  = 8.4, 2.4 Hz, 2H), 8.77 (d,  $J$  = 6.2 Hz, 2H), 8.28 (dd,  $J$  = 8.4, 0.8 Hz, 2H), 8.01 (d,  $J$  = 6.2 Hz, 2H).

**$^{13}C$  NMR** (201 MHz, DMSO- $D_6$ )  $\delta$  162.25, 151.35 (q,  $J$  = 34.8 Hz), 149.84, 149.37, 148.83, 147.17, 139.90, 136.66, 135.19, 122.35, 121.01.

**$^{19}F$  NMR** (376 MHz, acetone- $D_6$ )  $\delta$  -69.04 (s, 3F).

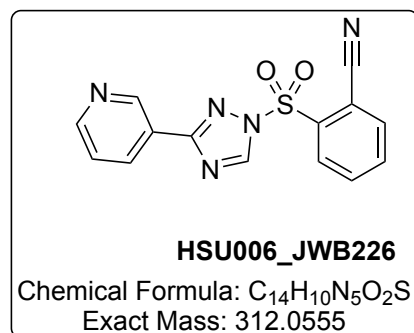
**HRMS** (ESI-TOF)  $m/z$ :  $[M + H]^+$  calculated for  $C_{13}H_9F_3N_5O_2S$  356.0424, Found, 356.0432.



**<sup>1</sup>H NMR** (800 MHz, Acetone-D<sub>6</sub>) δ 9.11 (s, 1H), 8.41 (s, 1H), 8.24 (d, *J* = 9.2 Hz, 2H), 8.17 (d, *J* = 8.0 Hz, 2H), 7.92 (s, 1H), 7.61 (d, *J* = 8.6 Hz, 1H), 7.33 (d, *J* = 9.2 Hz, 2H), 4.97 (s, 1H), 4.51 (s, 1H), 4.03 (s, 3H), 2.14 (p, *J* = 2.2 Hz, 2H), 2.10 (m, 3H), 1.92 – 1.85 (m, 2H), 1.58 (d, *J* = 6.9 Hz, 3H), 1.49 (s, 3H), 1.48 (s, 3H).

**<sup>13</sup>C NMR** (201 MHz, Acetone-D<sub>6</sub>) δ 166.62, 153.99, 147.62, 147.01, 144.11, 133.26, 132.04, 132.00, 130.31, 127.92, 127.27, 127.09, 123.93, 123.86, 121.25, 121.18, 120.61, 118.14, 116.34, 116.25, 111.13, 79.84, 79.76, 56.69, 39.75, 37.52, 35.19, 32.33, 31.75, 20.69, 19.99.

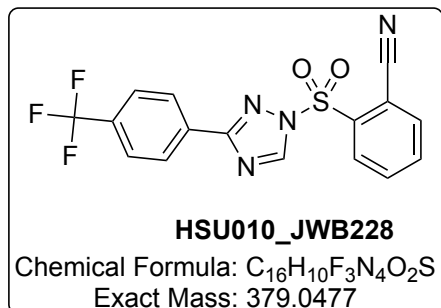
**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>35</sub>H<sub>34</sub>N<sub>5</sub>O<sub>4</sub>S 620.2332, Found, 620.2302.



**<sup>1</sup>H NMR** (400 MHz, Acetone-D<sub>6</sub>) δ 9.33 (s, 1H), 9.22 (s, 1H), 8.69 (d, *J* = 4.8 Hz, 1H), 8.55 (d, *J* = 7.6 Hz, 1H), 8.35 (d, *J* = 7.9 Hz, 1H), 8.21 (d, *J* = 7.3 Hz, 1H), 8.13 (m, 2H), 7.50 (dd, *J* = 8.0, 4.8 Hz, 1H).

**<sup>13</sup>C NMR** (100 MHz, Acetone-D<sub>6</sub>) δ 164.00, 152.53, 148.82, 148.81, 137.57, 137.20, 135.21, 134.81, 132.71, 125.95, 124.67, 115.33, 112.11.

**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>14</sub>H<sub>11</sub>N<sub>5</sub>O<sub>2</sub>S 312.0555, Found, 312.0559.

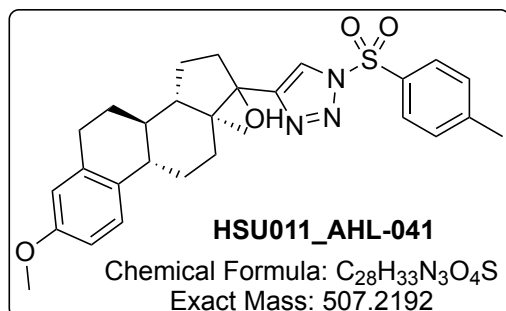


**$^1H$  NMR** (800 MHz, DMSO- $D_6$ )  $\delta$  8.65 (dd,  $J$  = 7.1, 3.9 Hz, 1H), 8.29 – 8.19 (m, 2H), 7.92 – 7.83 (m, 3H), 7.77 (dd,  $J$  = 7.6, 1.3 Hz, 1H), 7.64 (td,  $J$  = 7.7, 1.3 Hz, 1H), 7.49 (td,  $J$  = 7.6, 1.3 Hz, 1H).

**$^{13}C$  NMR** (201 MHz, DMSO- $D_6$ )  $\delta$  163.06, 150.38, 148.64, 146.04, 133.96, 132.25, 129.01, 127.36, 127.07, 126.48, 125.73(m, 1C), 125.70, 117.67, 108.95.

**$^{19}F$  NMR** (376 MHz, Acetone- $D_6$ )  $\delta$  -63.44 (s, 3F).

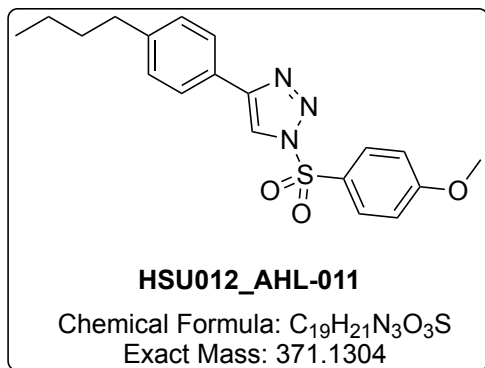
**HRMS** (ESI-TOF)  $m/z$ :  $[M + H]^+$  calculated for  $C_{16}H_{10}F_3N_4O_2S$  379.0471, Found, 379.0470.



**$^1H$  NMR** (400 MHz, Acetone- $D_6$ )  $\delta$  8.34 (s, 1H), 8.04 (d,  $J$  = 8.3 Hz, 2H), 7.55 (d,  $J$  = 8.1 Hz, 2H), 7.09 (d,  $J$  = 8.5 Hz, 1H), 6.64 (dd,  $J$  = 8.5, 2.7 Hz, 1H), 6.59 (s, 1H), 3.72 (s, 3H), 3.40 (q,  $J$  = 7.0 Hz, 1H), 3.11 (br s, 1H), 2.80 (m, 1H), 2.45 (s, 3H), 2.43 – 2.26 (m, 1H), 2.21 – 2.06 (m, 2H), 2.00 – 1.79 (m, 3H), 1.80 – 1.67 (m, 1H), 1.63 – 1.51 (m, 1H), 1.45 (q,  $J$  = 11.4 Hz, 1H), 1.38 – 1.23 (m, 2H), 1.11 (t,  $J$  = 7.0 Hz, 2H), 1.04 (s, 3H), 0.55 (td,  $J$  = 12.9, 4.2 Hz, 1H).

**$^{13}C$  NMR** (100 MHz, Acetone- $D_6$ )  $\delta$  158.49, 155.97, 148.47, 138.49, 134.43, 133.05, 131.50, 129.25, 126.99, 122.82, 114.39, 112.21, 82.57, 66.11, 55.27, 49.22, 48.19, 44.37, 40.47, 38.41, 33.80, 28.25, 27.06, 24.42, 21.67, 15.61, 14.72.

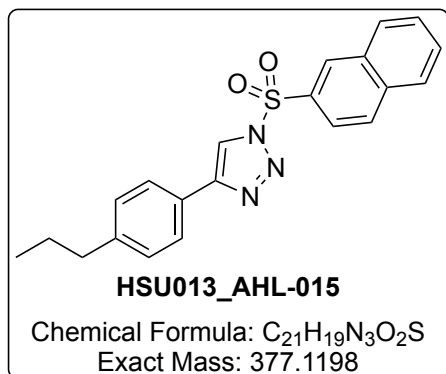
**HRMS** (ESI-TOF)  $m/z$ :  $[M + H]^+$  calculated for  $C_{28}H_{34}N_3O_4S$  508.2265, Found, 508.2257.



**<sup>1</sup>H NMR** (400 MHz, DMSO-D<sub>6</sub>) δ 9.31 (s, 1H), 8.10 (d, *J* = 9.1 Hz, 2H), 7.83 (d, *J* = 8.4 Hz, 2H), 7.29 (d, *J* = 8.5 Hz, 2H), 7.25 (d, *J* = 9.1 Hz, 2H), 3.87 (s, 3H), 2.60 (t, *J* = 7.6 Hz, 2H), 1.56 (p, *J* = 7.6 Hz, 2H), 1.29 (dq, *J* = 14.3, 7.2 Hz, 2H), 0.89 (t, *J* = 7.3 Hz, 3H).

**<sup>13</sup>C NMR** (151 MHz, DMSO-D<sub>6</sub>) δ 165.28, 147.03, 143.47, 130.99, 128.99, 126.21, 126.15, 125.79, 120.73, 115.77, 56.26, 34.61, 32.98, 21.74, 13.80.

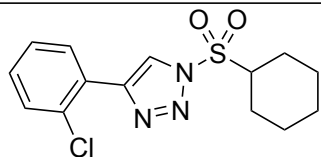
**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>19</sub>H<sub>22</sub>N<sub>3</sub>O<sub>3</sub>S 372.1376, Found, 372.1369.



**<sup>1</sup>H NMR** (400 MHz, DMSO-D<sub>6</sub>) δ 9.41 (s, 1H), 8.99 (d, *J* = 2.1 Hz, 1H), 8.33 (d, *J* = 8.1 Hz, 1H), 8.26 (d, *J* = 8.9 Hz, 1H), 8.16 – 8.08 (m, 1H), 8.05 (dd, *J* = 8.8, 2.1 Hz, 1H), 7.88 – 7.80 (m, 3H), 7.77 (ddd, *J* = 8.1, 7.0, 1.3 Hz, 1H), 7.28 (d, *J* = 8.1 Hz, 2H), 2.56 (t, *J* = 7.5 Hz, 2H), 1.58 (h, *J* = 7.4 Hz, 2H), 0.87 (t, *J* = 7.3 Hz, 3H).

**<sup>13</sup>C NMR** (100 MHz, Acetone-D<sub>6</sub>) δ 148.31, 144.50, 136.90, 134.02, 132.87, 131.88, 131.49, 131.42, 130.75, 129.85, 129.29, 129.03, 127.55, 126.76, 122.85, 120.91, 38.26, 25.14, 13.95.

**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>21</sub>H<sub>20</sub>N<sub>3</sub>O<sub>2</sub>S 378.1271, Found, 378.1272.



**HSU014\_EKT116**

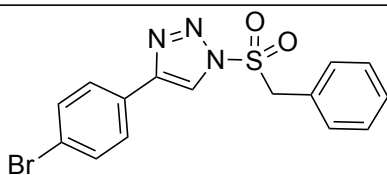
Chemical Formula:  $C_{14}H_{16}ClN_3O_2S$

Exact Mass: 325.0652

**$^1H$  NMR** (400 MHz, DMSO- $D_6$ )  $\delta$  9.03 (s, 1H), 8.04 (dd,  $J$  = 6.7, 2.8 Hz, 1H), 7.64 (dd,  $J$  = 7.2, 2.1 Hz, 1H), 7.57 – 7.46 (m, 2H), 4.10 (tt,  $J$  = 11.9, 3.6 Hz, 1H), 1.99 (d,  $J$  = 11.0 Hz, 2H), 1.80 (d,  $J$  = 13.2 Hz, 2H), 1.60 (d,  $J$  = 13.0 Hz, 1H), 1.45 (m, 2H), 1.31 (m, 2H), 1.13 (m, 1H).

**$^{13}C$  NMR** (151 MHz, DMSO- $D_6$ )  $\delta$  143.17, 131.21, 130.70, 130.46, 130.35, 127.71, 127.46, 124.94, 63.41, 25.46, 24.19, 23.91.

**HRMS** (ESI-TOF)  $m/z$ :  $[M + H]^+$  calculated for  $C_{14}H_{17}ClN_3O_2S$  326.0725, Found, 326.0721.



**HSU015\_EKT151**

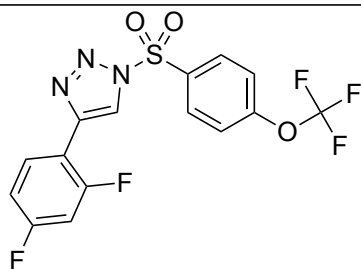
Chemical Formula:  $C_{15}H_{12}BrN_3O_2S$

Exact Mass: 376.9834

**$^1H$  NMR** (400 MHz,  $CDCl_3$ )  $\delta$  7.78 (s, 1H), 7.58 (q,  $J$  = 8.5 Hz, 4H), 7.36 (t,  $J$  = 7.4 Hz, 1H), 7.29 (t,  $J$  = 7.3 Hz, 2H), 7.12 (d,  $J$  = 7.4 Hz, 2H), 4.87 (s, 2H).

**$^{13}C$  NMR** (100 MHz, DMSO- $D_6$ )  $\delta$  145.32, 132.07, 130.88, 129.45, 128.82, 128.07, 127.77, 126.30, 122.79, 122.15, 60.07.

**HRMS** (ESI-TOF)  $m/z$ :  $[M + H]^+$  calculated for  $C_{15}H_{12}BrN_3O_2S$  377.9906, Found, 377.9888.



**HSU017\_EKT165**

Chemical Formula:  $C_{15}H_8F_5N_3O_3S$

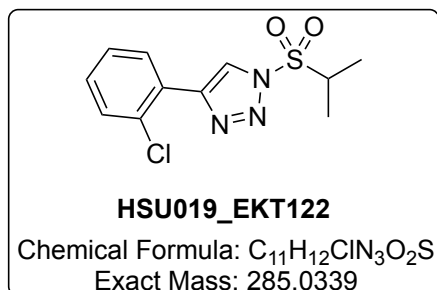
Exact Mass: 405.0207

**$^1H$  NMR** (400 MHz, Acetone- $D_6$ )  $\delta$  8.82 (d,  $J$  = 2.0 Hz, 1H), 8.42 (d,  $J$  = 7.7 Hz, 2H), 8.23 (q,  $J$  = 8.7 Hz, 1H), 7.72 (d,  $J$  = 9.0 Hz, 2H), 7.40 – 7.02 (m, 2H).

**<sup>13</sup>C NMR** (100 MHz, Acetone-D<sub>6</sub>) δ 164.08 (d, *J* = 237.5 Hz), 160.47 (dd, *J* = 251.3, 12.0 Hz), 155.07 (d, *J* = 2.0 Hz), 141.23, 135.35, 132.45, 130.31 (dd, *J* = 10.0, 4.6 Hz), 123.33 (d, *J* = 11.7 Hz), 122.82, 121.06 (d, *J* = 258.8 Hz), 114.63 (d, *J* = 13.4 Hz), 113.13 (dd, *J* = 21.9, 3.6 Hz), 105.30 (t, *J* = 26.0 Hz).

**<sup>19</sup>F NMR** (376 MHz, Acetone-D<sub>6</sub>) δ -58.51 (s, 3F), -109.77 – -110.07 (m, 1F), -110.25 – -110.58 (m, 1F).

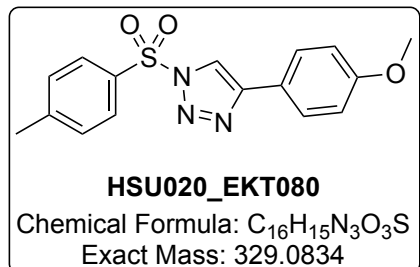
**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>15</sub>H<sub>9</sub>F<sub>5</sub>N<sub>3</sub>O<sub>3</sub>S 406.0279, Found, 406.0281.



**<sup>1</sup>H NMR** (400 MHz, CDCl<sub>3</sub>) δ 8.69 (s, 1H), 8.26 (dd, *J* = 7.8, 2.5 Hz, 1H), 7.50 (d, *J* = 8.0 Hz, 1H), 7.42 (t, *J* = 7.5 Hz, 1H), 7.35 (td, *J* = 7.6, 1.4 Hz, 1H), 3.90 (hept, *J* = 7.2 Hz, 1H), 1.49 (s, 3H), 1.47 (s, 3H).

**<sup>13</sup>C NMR** (100 MHz, DMSO-D<sub>6</sub>) δ 143.20, 131.28, 130.75, 130.53, 130.34, 127.73, 127.49, 125.15, 56.99, 15.58.

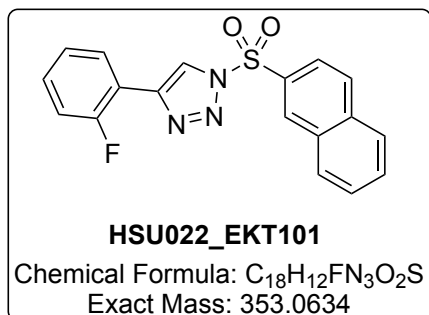
**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>11</sub>H<sub>13</sub>ClN<sub>3</sub>O<sub>2</sub>S 286.0412, Found, 286.0400.



**<sup>1</sup>H NMR** (400 MHz, DMSO-D<sub>6</sub>) δ 8.22 (s, 1H), 7.78 (d, *J* = 8.7 Hz, 2H), 7.48 (d, *J* = 8.1 Hz, 2H), 7.12 (d, *J* = 7.9 Hz, 2H), 7.02 (d, *J* = 8.8 Hz, 2H), 3.79 (s, 3H), 2.29 (s, 3H).

**<sup>13</sup>C NMR** (151 MHz, DMSO-D<sub>6</sub>) δ 160.81, 151.47, 147.06, 137.31, 132.10, 130.65, 128.21, 128.12, 120.05, 114.67, 55.35, 21.21.

**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>16</sub>H<sub>16</sub>N<sub>3</sub>O<sub>3</sub>S 364.0907, Found, 364.0907.

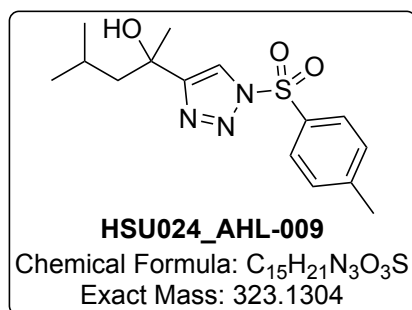


**<sup>1</sup>H NMR** (400 MHz, CDCl<sub>3</sub>) δ 8.79 (s, 1H), 8.55 (d, *J* = 3.4 Hz, 1H), 8.25 (t, *J* = 7.6 Hz, 1H), 8.02 (m, 3H), 7.92 (d, *J* = 8.3 Hz, 1H), 7.72 (t, *J* = 7.5 Hz, 1H), 7.67 (t, *J* = 7.5 Hz, 1H), 7.34 (q, *J* = 6.8 Hz, 1H), 7.23 (t, *J* = 7.6 Hz, 1H), 7.19 – 7.10 (m, 1H).

**<sup>13</sup>C NMR** (100 MHz, DMSO-D<sub>6</sub>) δ 159.98, 157.50, 140.72 (d, *J* = 2.3 Hz), 135.66, 131.96, 131.54, 131.33, 131.11 (d, *J* = 8.5 Hz), 130.92, 130.75, 130.03, 128.53, 128.14, 125.12 (d, *J* = 3.4 Hz), 123.32 (d, *J* = 10.4 Hz), 121.89, 116.24 (d, *J* = 21.0 Hz).

**<sup>19</sup>F NMR** (376 MHz, Acetone-D<sub>6</sub>) δ -114.71 (dtd, *J* = 11.4, 8.0, 3.8 Hz, 1F).

**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>18</sub>H<sub>13</sub>FN<sub>3</sub>O<sub>2</sub>S 354.0707, Found, 354.0709.

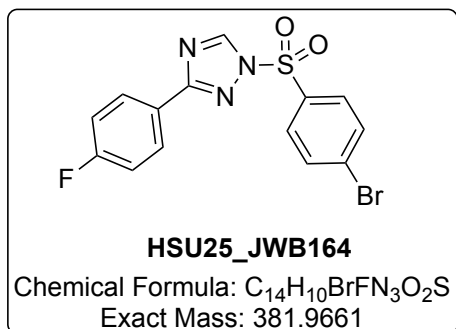


**<sup>1</sup>H NMR** (400 MHz, DMSO-D<sub>6</sub>) δ 8.59 (s, 1H), 7.67 (s, 1H), 7.48 (d, *J* = 8.1 Hz, 2H), 7.12 (d, *J* = 8.3 Hz, 2H), 2.29 (s, 3H), 1.67 – 1.62 (m, 2H), 1.58 (dt, *J* = 12.7, 6.0 Hz, 1H), 1.45 (s, 3H), 0.80 (d, *J* = 6.5 Hz, 3H), 0.70 (d, *J* = 6.5 Hz, 3H).

**<sup>13</sup>C NMR** (151 MHz, DMSO-D<sub>6</sub>) δ 145.32, 137.90, 128.16, 125.52, 69.66, 51.56, 29.65, 24.38, 24.15, 23.82, 20.81.

**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>15</sub>H<sub>22</sub>N<sub>3</sub>O<sub>3</sub>S 324.1376, Found, 324.1377.



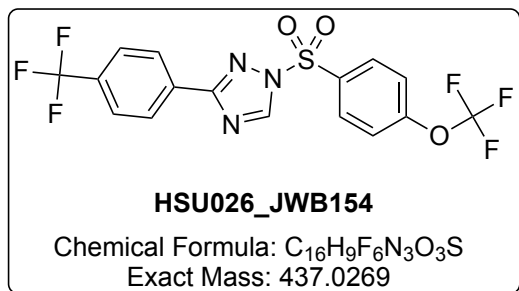


**$^1H$  NMR** (800 MHz, DMSO- $D_6$ )  $\delta$  8.59 (s, 1H), 8.15 – 7.98 (m, 2H), 7.60 – 7.47 (m, 4H), 7.34 (m, 2H).

**$^{13}C$  NMR** (201 MHz, DMSO- $D_6$ )  $\delta$  163.43, 162.22, 147.70, 146.27, 130.61, 128.23, 127.76, 121.57, 115.96, 115.85.

**$^{19}F$  NMR** (376 MHz, DMSO- $D_6$ )  $\delta$  -113.88 (td,  $J$  = 8.9, 4.5 Hz, 1F).

**HRMS** (ESI-TOF)  $m/z$ :  $[M + H]^+$  calculated for  $C_{14}H_{11}BrFN_3O_2S$  382.9656, Found, 382.9648.

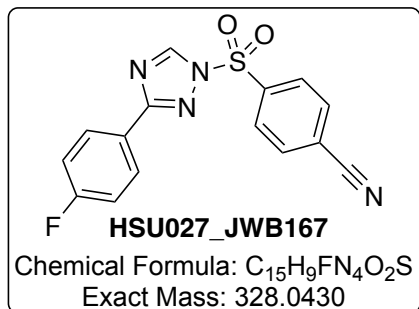


**$^1H$  NMR** (600 MHz, DMSO- $D_6$ )  $\delta$  9.57 (s, 1H), 8.43 – 8.28 (m, 2H), 8.19 (d,  $J$  = 8.1 Hz, 2H), 7.86 (d,  $J$  = 8.1 Hz, 2H), 7.77 – 7.66 (m, 2H).

**$^{13}C$  NMR** (201 MHz, DMSO- $D_6$ )  $\delta$  163.39, 158.41, 153.34, 148.66, 147.95, 146.48, 128.11, 127.80, 127.04, 126.62 (m, 1C), 126.32 (m, 1C), 120.74.

**$^{19}F$  NMR** (376 MHz, Acetone- $D_6$ )  $\delta$  -58.51 (s, 3F), -63.43 (s, 3F).

**HRMS** (ESI-TOF)  $m/z$ :  $[M + H]^+$  calculated for  $C_{16}H_{10}F_6N_3O_3S$  438.0342, Found, 438.0343.

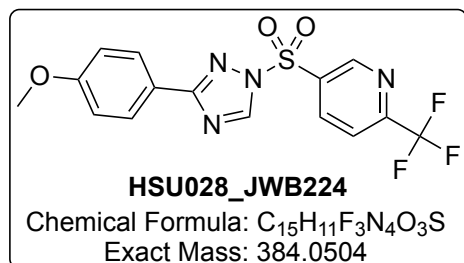


**$^1H$  NMR** (800 MHz, DMSO- $D_6$ )  $\delta$  8.72 (s, 1H), 8.12 – 7.99 (m, 2H), 7.86 – 7.78 (m, 2H), 7.78 – 7.68 (m, 2H), 7.47 – 7.24 (m, 2H).

**<sup>13</sup>C NMR** (201 MHz, DMSO-D6)  $\delta$  163.61, 162.39, 156.82, 152.55, 146.02, 132.10, 128.42 (d,  $J$  = 8.8 Hz), 126.43, 118.69, 116.02 (d,  $J$  = 22.0 Hz), 111.09.

**<sup>19</sup>F NMR** (376 MHz, Acetone-D6)  $\delta$  -111.70 (ddd,  $J$  = 13.7, 8.9, 5.3 Hz, 1F).

**HRMS** (ESI-TOF)  $m/z$ : [M + H]<sup>+</sup> calculated for C<sub>15</sub>H<sub>10</sub>FN<sub>4</sub>O<sub>2</sub>S 329.0503, Found, 329.0498.

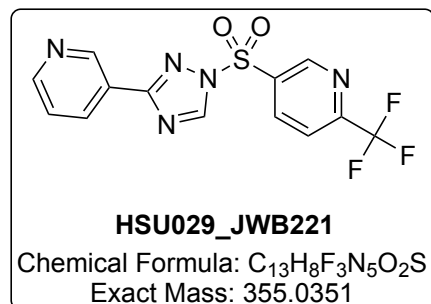


**<sup>1</sup>H NMR** (400 MHz, Acetone-D6)  $\delta$  9.48 (s, 1H), 9.17 (s, 1H), 8.89 (d,  $J$  = 8.3 Hz, 1H), 8.24 (d,  $J$  = 8.3 Hz, 1H), 8.02 (d,  $J$  = 8.9 Hz, 2H), 7.03 (d,  $J$  = 8.9 Hz, 2H), 3.85 (s, 3H).

**<sup>13</sup>C NMR** (100 MHz, Acetone-D6)  $\delta$  166.13, 162.81, 153.09 (d,  $J$  = 35.2 Hz), 150.26, 148.01, 140.23, 137.07, 129.38, 122.79 (q,  $J$  = 2.9 Hz), 122.34, 120.39, 115.07, 55.77.

**<sup>19</sup>F NMR** (376 MHz, Acetone-D6)  $\delta$  -69.04 (s, 1F).

**HRMS** (ESI-TOF)  $m/z$ : [M + H]<sup>+</sup> calculated for C<sub>15</sub>H<sub>12</sub>F<sub>3</sub>N<sub>4</sub>O<sub>3</sub>S 385.0577, Found, 385.0590.

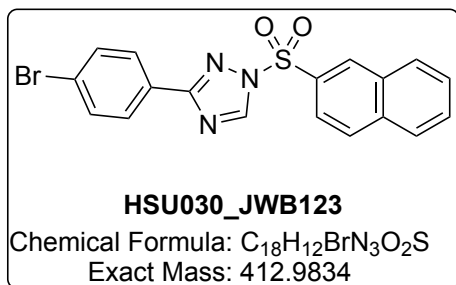


**<sup>1</sup>H NMR** (800 MHz, DMSO-D6)  $\delta$  9.59 (s, 1H), 9.50 (d,  $J$  = 2.4 Hz, 1H), 9.18 (d,  $J$  = 1.2 Hz, 1H), 8.88 (dd,  $J$  = 8.4, 2.4 Hz, 1H), 8.72 (dd,  $J$  = 4.8, 1.7 Hz, 1H), 8.39 – 8.32 (m, 1H), 8.28 (d,  $J$  = 8.4 Hz, 1H), 7.58 – 7.51 (m, 1H).

**<sup>13</sup>C NMR** (201 MHz, DMSO-D6)  $\delta$  162.38, 151.63, 151.29 (d,  $J$  = 35.0 Hz), 149.27, 148.58, 147.49, 139.80, 135.32, 134.31, 124.62, 124.20, 122.33 (q,  $J$  = 2.9 Hz), 121.31, 119.94.

**<sup>19</sup>F NMR** (376 MHz, acetone)  $\delta$  -69.04 (s, 3F).

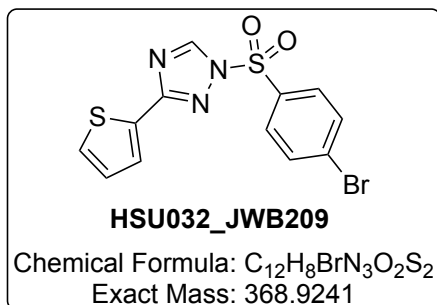
**HRMS** (ESI-TOF)  $m/z$ : [M + H]<sup>+</sup> calculated for C<sub>13</sub>H<sub>9</sub>F<sub>3</sub>N<sub>5</sub>O<sub>2</sub>S 356.0424, Found, 356.0419.



**<sup>1</sup>H NMR** (800 MHz, DMSO-D<sub>6</sub>) δ 8.61 (s, 1H), 8.14 (d, *J* = 0.8 Hz, 1H), 7.99 – 7.94 (m, 3H), 7.92 – 7.88 (m, 1H), 7.86 (d, *J* = 8.3 Hz, 1H), 7.73 – 7.69 (m, 3H), 7.54 – 7.50 (m, 2H).

**<sup>13</sup>C NMR** (201 MHz, DMSO-D<sub>6</sub>) δ 146.20, 145.58, 132.72, 132.15, 131.89, 128.46, 127.93, 127.45, 127.31, 126.43, 126.29, 124.04, 123.97, 122.73.

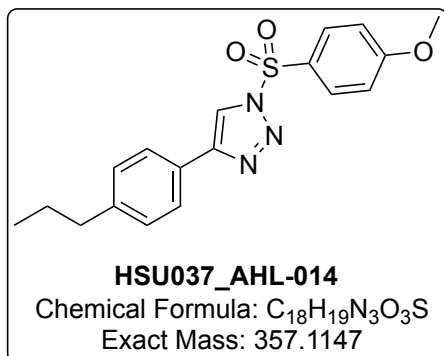
**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>18</sub>H<sub>13</sub>BrN<sub>3</sub>O<sub>2</sub>S 413.9906, Found, 413.9907.



**<sup>1</sup>H NMR** (400 MHz, DMSO-D<sub>6</sub>) δ 9.45 (s, 1H), 8.06 (d, *J* = 8.8 Hz, 2H), 7.97 (d, *J* = 9.0 Hz, 2H), 7.76 (dd, *J* = 5.0, 1.2 Hz, 1H), 7.71 (dd, *J* = 3.7, 1.2 Hz, 1H), 7.18 (dd, *J* = 5.0, 3.7 Hz, 1H).

**<sup>13</sup>C NMR** (100 MHz, DMSO-D<sub>6</sub>) δ 160.38, 148.03, 134.40, 133.55, 130.76, 130.23, 129.82, 128.77, 128.53, 127.83.

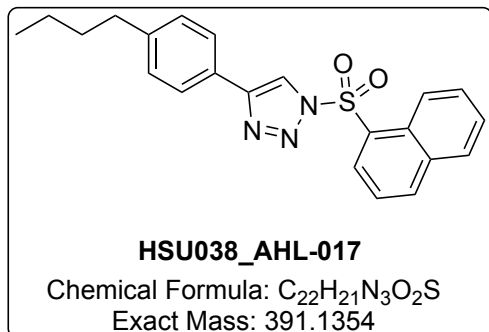
**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>12</sub>H<sub>9</sub>BrN<sub>3</sub>O<sub>2</sub>S<sub>2</sub> 369.9314, Found, 369.9309.



**<sup>1</sup>H NMR** (400 MHz, DMSO-D<sub>6</sub>) δ 9.32 (s, 1H), 8.10 (d, *J* = 9.1 Hz, 2H), 7.83 (d, *J* = 8.3 Hz, 2H), 7.29 (d, *J* = 8.6 Hz, 2H), 7.25 (d, *J* = 9.1 Hz, 2H), 3.87 (s, 3H), 2.58 (t, *J* = 7.5 Hz, 2H), 1.59 (h, *J* = 7.4 Hz, 2H), 0.88 (t, *J* = 7.3 Hz, 3H).

**<sup>13</sup>C NMR** (100 MHz, DMSO-D<sub>6</sub>) δ 165.22, 146.98, 143.18, 130.94, 128.99, 126.23, 126.12, 125.72, 120.72, 115.72, 56.21, 36.97, 23.91, 13.57.

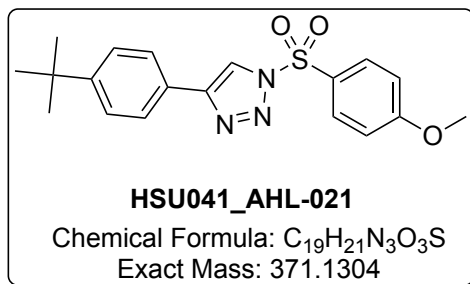
**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>18</sub>H<sub>20</sub>N<sub>3</sub>O<sub>3</sub>S 358.1220, Found, 358.1220.



**<sup>1</sup>H NMR** (400 MHz, DMSO-D<sub>6</sub>) δ 9.53 (s, 1H), 8.65 (t, *J* = 7.3 Hz, 2H), 8.51 (d, *J* = 8.3 Hz, 1H), 8.17 (d, *J* = 7.3 Hz, 1H), 7.85 (t, *J* = 7.9 Hz, 1H), 7.79 (d, *J* = 8.2 Hz, 3H), 7.71 (t, *J* = 7.5 Hz, 1H), 7.25 (d, *J* = 8.3 Hz, 2H), 2.56 (t, *J* = 7.7 Hz, 2H), 1.52 (p, *J* = 7.6 Hz, 2H), 1.26 (h, *J* = 7.3 Hz, 2H), 0.85 (t, *J* = 7.3 Hz, 3H).

**<sup>13</sup>C NMR** (100 MHz, DMSO-d<sub>6</sub>) δ 146.90, 143.51, 138.16, 133.82, 132.86, 129.94, 128.95, 127.90, 127.08, 126.00, 125.77, 125.10, 122.95, 121.26, 34.57, 32.94, 21.69, 13.76.

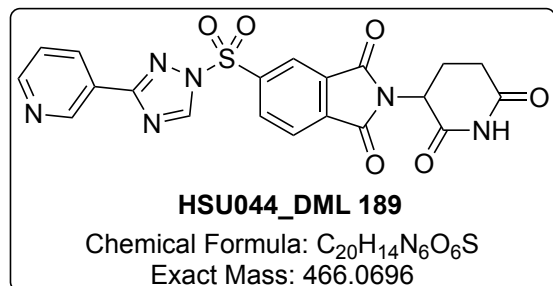
**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>22</sub>H<sub>22</sub>N<sub>3</sub>O<sub>2</sub>S 392.1427, Found, 392.1427.



**<sup>1</sup>H NMR** (400 MHz, DMSO-D<sub>6</sub>) δ 9.33 (s, 1H), 8.10 (d, *J* = 9.1 Hz, 2H), 7.85 (d, *J* = 8.5 Hz, 2H), 7.49 (d, *J* = 8.5 Hz, 2H), 7.25 (d, *J* = 9.1 Hz, 2H), 3.88 (s, 3H), 1.29 (s, 9H).

**<sup>13</sup>C NMR** (100 MHz, DMSO-D<sub>6</sub>) δ 165.25, 151.64, 146.88, 130.96, 126.10, 125.98, 125.83, 125.59, 120.79, 115.75, 56.23, 34.49, 31.00.

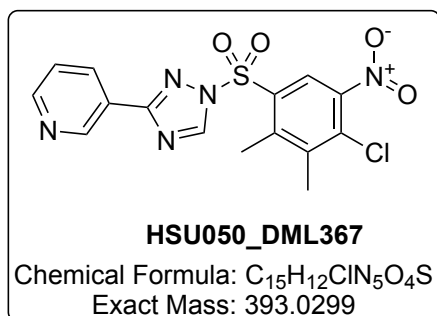
**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>19</sub>H<sub>22</sub>N<sub>3</sub>O<sub>3</sub>S 372.1376, Found, 372.1377.



**<sup>1</sup>H NMR** (400 MHz, Acetone-D<sub>6</sub>) δ 10.04 (s, 1H), 9.35 (s, 1H), 9.25 (s, 1H), 8.81 – 8.64 (m, 2H), 8.59 (s, 1H), 8.42 (d, *J* = 8.0 Hz, 1H), 8.25 (d, *J* = 8.0 Hz, 1H), 7.56 (dd, *J* = 8.0, 4.9 Hz, 1H), 5.23 (dd, *J* = 12.9, 5.4 Hz, 1H), 2.98 (ddd, *J* = 17.6, 14.4, 5.4 Hz, 1H), 2.84 – 2.68 (m, 2H), 2.25 (ddt, *J* = 13.0, 5.7, 2.7 Hz, 2H).

**<sup>13</sup>C NMR** (100 MHz, Acetone-D<sub>6</sub>) δ 172.50, 169.55, 166.24, 166.02, 163.78, 151.90, 148.55, 148.26, 142.24, 138.13, 136.12, 135.47, 134.00, 132.98, 125.89, 124.95, 124.13, 50.79, 31.84, 22.98.

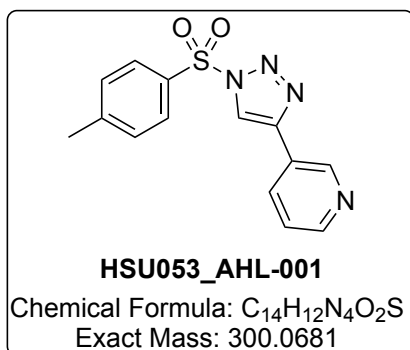
**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>20</sub>H<sub>15</sub>N<sub>6</sub>O<sub>6</sub>S 467.0768, Found, 467.0768.



**<sup>1</sup>H NMR** (400 MHz, Acetone-D<sub>6</sub>) δ 9.38 (s, 1H), 9.23 (s, 1H), 8.69 (s, 2H), 8.37 (dd, *J* = 8.1, 1.9 Hz, 1H), 7.52 (dd, *J* = 8.0, 4.6 Hz, 1H), 2.86 (s, 3H), 2.56 (s, 3H).

**<sup>13</sup>C NMR** (100 MHz, Acetone-D<sub>6</sub>) δ 163.54, 152.49, 148.82, 148.35, 145.02, 143.31, 135.09, 134.86, 133.33, 126.03, 125.44, 124.71, 18.39, 17.85.

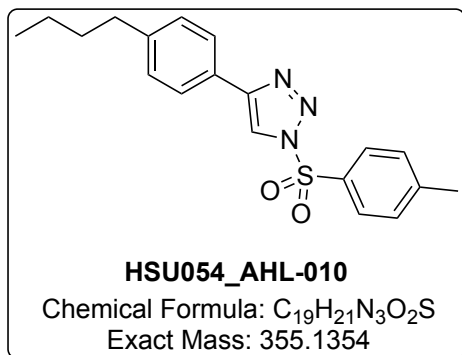
**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>15</sub>H<sub>12</sub>ClN<sub>5</sub>O<sub>4</sub>S 394.0371, Found, 394.0358.



**<sup>1</sup>H NMR** (400 MHz, Acetone-D<sub>6</sub>) δ 9.16 (s, 2H), 8.61 (d, *J* = 4.9 Hz, 1H), 8.31 (dd, *J* = 8.0, 1.8 Hz, 1H), 8.10 (d, *J* = 8.8 Hz, 2H), 7.58 (d, *J* = 8.1 Hz, 2H), 7.48 (dd, *J* = 8.0, 4.8 Hz, 1H), 2.48 (s, 3H).

**<sup>13</sup>C NMR** (100 MHz, Acetone-D<sub>6</sub>) δ 150.85, 148.86, 148.14, 145.42, 133.99, 133.87, 131.63, 129.45, 126.24, 124.67, 122.07, 21.69.

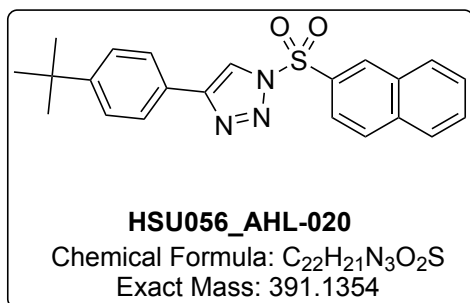
**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>14</sub>H<sub>13</sub>BrN<sub>4</sub>O<sub>2</sub>S 301.0754, Found, 301.0755.



**<sup>1</sup>H NMR** (400 MHz, DMSO-D<sub>6</sub>) δ 9.34 (s, 1H), 8.05 (d, *J* = 8.5 Hz, 2H), 7.83 (d, *J* = 8.3 Hz, 2H), 7.56 (d, *J* = 8.8 Hz, 2H), 7.29 (d, *J* = 8.2 Hz, 2H), 2.60 (t, *J* = 7.6 Hz, 2H), 2.42 (s, 3H), 1.56 (p, *J* = 7.3 Hz, 2H), 1.30 (h, *J* = 7.1 Hz, 2H), 0.89 (t, *J* = 7.3 Hz, 3H).

**<sup>13</sup>C NMR** (151 MHz, DMSO-D<sub>6</sub>) δ 147.68, 147.07, 143.44, 132.40, 130.87, 128.92, 128.22, 126.09, 125.76, 120.87, 34.56, 32.92, 21.68, 21.26, 13.74.

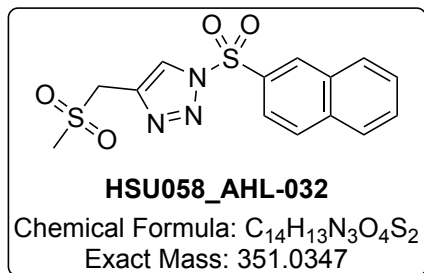
**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>19</sub>H<sub>22</sub>N<sub>3</sub>O<sub>2</sub>S 356.1427, Found, 356.1426.



**<sup>1</sup>H NMR** (400 MHz, DMSO-D<sub>6</sub>) δ 9.42 (s, 1H), 9.00 (d, *J* = 2.2 Hz, 1H), 8.33 (d, *J* = 8.3 Hz, 1H), 8.27 (d, *J* = 9.1 Hz, 1H), 8.12 (d, *J* = 7.1 Hz, 1H), 8.05 (dd, *J* = 8.8, 2.1 Hz, 1H), 7.91 – 7.80 (m, 3H), 7.77 (ddd, *J* = 8.2, 7.0, 1.4 Hz, 1H), 7.48 (d, *J* = 8.8 Hz, 2H), 1.28 (s, 9H).

**<sup>13</sup>C NMR** (100 MHz, DMSO-D<sub>6</sub>) δ 151.74, 147.05, 135.62, 132.19, 131.55, 131.02, 130.92, 130.81, 130.04, 128.58, 128.17, 125.87, 125.84, 125.66, 121.75, 121.16, 34.50, 30.99.

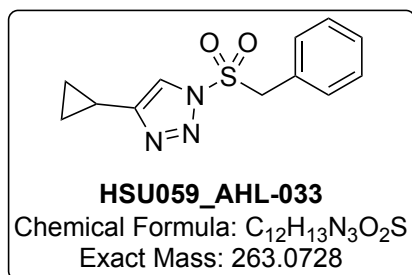
**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>22</sub>H<sub>22</sub>N<sub>3</sub>O<sub>2</sub>S 392.1427, Found, 392.1428.



**<sup>1</sup>H NMR** (400 MHz, DMSO-D<sub>6</sub>) δ 9.03 (d, *J* = 2.1 Hz, 1H), 8.96 (s, 1H), 8.33 (d, *J* = 8.0 Hz, 1H), 8.27 (d, *J* = 8.8 Hz, 1H), 8.13 (d, *J* = 7.9 Hz, 1H), 8.06 (dd, *J* = 8.8, 2.1 Hz, 1H), 7.84 (ddd, *J* = 8.3, 6.9, 1.4 Hz, 1H), 7.78 (ddd, *J* = 8.2, 6.9, 1.4 Hz, 1H), 4.68 (s, 2H), 3.03 (s, 3H).

**<sup>13</sup>C NMR** (100 MHz, DMSO-D<sub>6</sub>) δ 137.10, 135.69, 131.76, 131.55, 131.36, 131.00, 130.82, 130.05, 128.59, 128.16, 125.81, 121.82, 50.38.

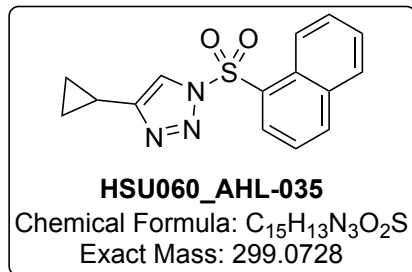
**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>14</sub>H<sub>14</sub>N<sub>3</sub>O<sub>4</sub>S<sub>2</sub> 352.0420, Found, 352.0419.



**<sup>1</sup>H NMR** (400 MHz, DMSO-D<sub>6</sub>) δ 8.10 (s, 1H), 7.60 – 7.25 (m, 3H), 7.10 (d, *J* = 6.5 Hz, 2H), 5.29 (s, 2H), 1.97 (ddd, *J* = 13.4, 8.4, 5.0 Hz, 1H), 1.13 – 0.85 (m, 3H), 0.84 – 0.55 (m, 3H).

**<sup>13</sup>C NMR** (151 MHz, DMSO-D<sub>6</sub>) δ 149.70, 130.81, 129.39, 128.77, 126.48, 121.68, 59.98, 7.89, 6.15.

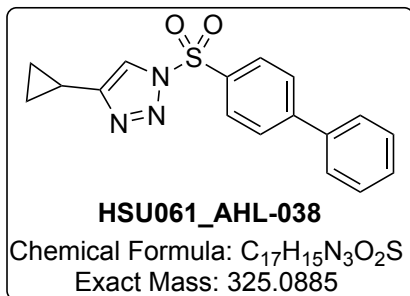
**HRMS** (ESI-TOF) *m/z*: [M + Na]<sup>+</sup> calculated for C<sub>12</sub>H<sub>13</sub>N<sub>3</sub>O<sub>2</sub>SNa 286.0621, Found, 286.0616.



**<sup>1</sup>H NMR** (400 MHz, DMSO-D<sub>6</sub>) δ 8.99 – 8.73 (m, 1H), 7.94 (d, *J* = 7.0 Hz, 1H), 7.91 – 7.83 (m, 2H), 7.60 (s, 1H), 7.53 – 7.47 (m, 2H), 7.43 (t, *J* = 7.7 Hz, 1H), 1.96 (ddd, *J* = 13.3, 8.1, 4.8 Hz, 1H), 0.99 – 0.89 (m, 2H), 0.75 – 0.67 (m, 2H).

**<sup>13</sup>C NMR** (151 MHz, DMSO-D<sub>6</sub>) δ 143.93, 133.54, 129.42, 129.03, 127.75, 127.62, 125.58, 125.56, 124.46, 124.35, 7.95, 5.84.

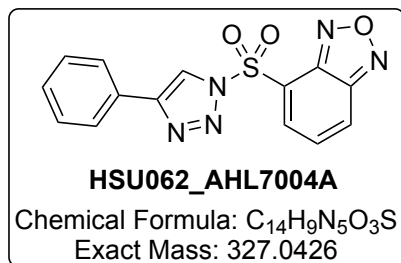
**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>15</sub>H<sub>14</sub>N<sub>3</sub>O<sub>2</sub>S 300.0801, Found, 300.0788.



**<sup>1</sup>H NMR** (400 MHz, DMSO-D<sub>6</sub>) δ 9.17 (s, 1H), 7.69 – 7.65 (m, 3H), 7.63 – 7.58 (m, 3H), 7.47 (t, *J* = 7.7 Hz, 2H), 7.37 (t, *J* = 7.6 Hz, 1H), 1.96 (tt, *J* = 8.5, 5.1 Hz, 1H), 0.97 – 0.89 (m, 2H), 0.75 – 0.65 (m, 2H).

**<sup>13</sup>C NMR** (100 MHz, DMSO-D<sub>6</sub>) δ 147.17, 147.00, 140.29, 139.73, 128.99, 127.63, 127.41, 126.78, 126.19, 126.11, 8.00, 5.84.

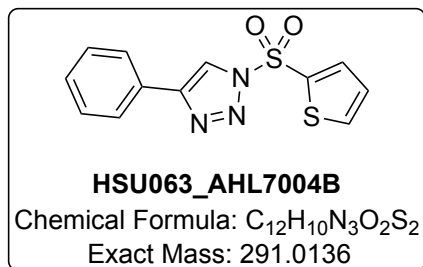
**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>17</sub>H<sub>16</sub>N<sub>3</sub>O<sub>2</sub>S 326.0958, Found, 326.0946.



**<sup>1</sup>H NMR** (400 MHz, Acetone-D<sub>6</sub>) δ 9.37 (s, 1H), 8.69 (dd, *J* = 6.9, 0.8 Hz, 1H), 8.51 (dd, *J* = 9.1, 0.8 Hz, 1H), 8.09 – 7.85 (m, 3H), 7.63 – 7.35 (m, 3H).

**<sup>13</sup>C NMR** (100 MHz, Acetone-D<sub>6</sub>) δ 165.86, 150.68, 149.09, 144.74, 139.51, 132.13, 131.58, 129.99, 129.66, 127.67, 125.77, 125.25.

**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>14</sub>H<sub>10</sub>N<sub>5</sub>O<sub>3</sub>S 328.0499, Found, 328.0488.

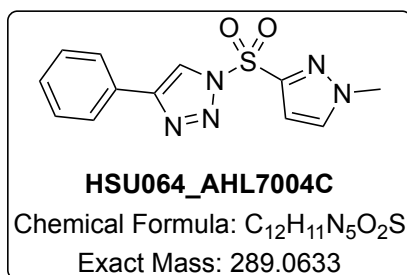


**<sup>1</sup>H NMR** (400 MHz, Acetone-D<sub>6</sub>) δ 9.16 (s, 1H), 8.25 (d, *J* = 5.0 Hz, 1H), 8.17 – 7.94 (m, 3H), 7.49 (s, 3H), 7.35 (t, *J* = 4.6 Hz, 1H).

**<sup>13</sup>C NMR** (100 MHz, Acetone-D<sub>6</sub>) δ 165.51, 147.26, 139.25, 138.25, 136.02, 131.43, 130.22, 129.66, 129.65, 127.59.

**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>12</sub>H<sub>11</sub>N<sub>3</sub>O<sub>2</sub>S<sub>2</sub> 292.0209, Found, 292.0196.

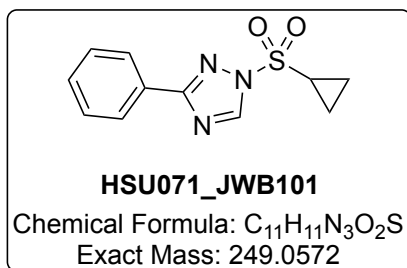




**<sup>1</sup>H NMR** (400 MHz, Acetone-D<sub>6</sub>) δ 9.14 (s, 1H), 8.10 (ddt, *J* = 5.5, 2.2, 0.7 Hz, 2H), 7.93 (d, *J* = 2.4 Hz, 1H), 7.49 (dd, *J* = 3.8, 2.8 Hz, 3H), 7.07 (d, *J* = 2.5 Hz, 1H), 4.01 (s, 3H).

**<sup>13</sup>C NMR** (100 MHz, Acetone-D<sub>6</sub>) δ 165.41, 147.80, 146.46, 134.77, 131.38, 130.34, 129.67, 127.59, 110.22, 40.55.

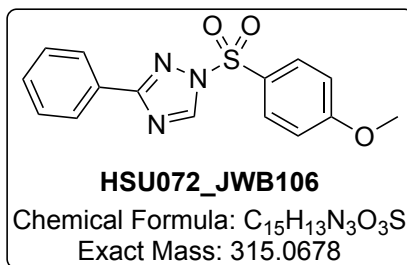
**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>12</sub>H<sub>12</sub>N<sub>5</sub>O<sub>2</sub>S 290.0706, Found, 290.0695



**<sup>1</sup>H NMR** (800 MHz, DMSO-D<sub>6</sub>) δ 9.27 (s, 1H), 8.32 – 7.96 (m, 2H), 7.72 – 7.31 (m, 3H), 3.37 (tt, *J* = 7.8, 4.5 Hz, 1H), 1.44 – 1.41 (m, 2H), 1.39 – 1.22 (m, 2H).

**<sup>13</sup>C NMR** (201 MHz, DMSO-D<sub>6</sub>) δ 163.49, 147.48, 130.64, 129.07, 128.99, 126.53, 30.99, 7.25.

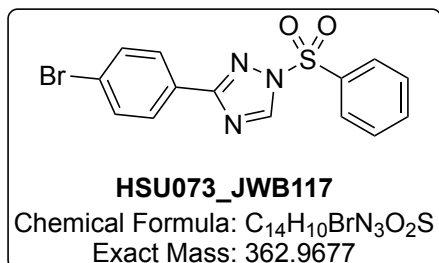
**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>11</sub>H<sub>12</sub>N<sub>3</sub>O<sub>2</sub>S 250.0645, Found, 250.0641.



**<sup>1</sup>H NMR** (800 MHz, DMSO-D<sub>6</sub>) δ 9.42 (s, 1H), 8.10 – 8.07 (m, 2H), 8.00 – 7.96 (m, 2H), 7.49 (dd, *J* = 5.1, 2.0 Hz, 3H), 7.26 – 7.20 (m, 2H), 3.86 (s, 3H).

**$^{13}\text{C}$  NMR** (201 MHz, DMSO- $\text{D}_6$ )  $\delta$  164.98, 163.79, 147.34, 130.95, 130.68, 129.02, 128.77, 126.47, 126.05, 115.54, 56.13.

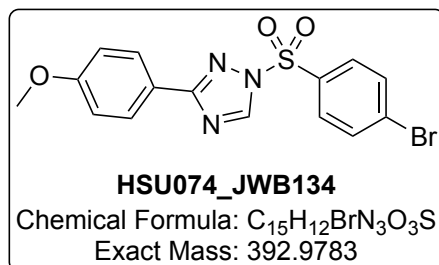
**HRMS** (ESI-TOF)  $m/z$ :  $[\text{M} + \text{H}]^+$  calculated for  $\text{C}_{15}\text{H}_{14}\text{N}_3\text{O}_3\text{S}$  316.0750, Found, 316.0748.



**$^1\text{H}$  NMR** (600 MHz, DMSO- $\text{D}_6$ )  $\delta$  9.51 (s, 1H), 8.15 (d,  $J = 7.5$  Hz, 2H), 7.92 (d,  $J = 8.4$  Hz, 2H), 7.88 (t,  $J = 7.5$  Hz, 1H), 7.75 (t,  $J = 7.8$  Hz, 2H), 7.70 (d,  $J = 8.4$  Hz, 2H).

**$^{13}\text{C}$  NMR** (201 MHz, DMSO- $\text{D}_6$ )  $\delta$  163.38, 152.71, 148.19, 132.23, 131.92, 128.52, 128.48, 128.00, 127.98, 127.67, 125.48.

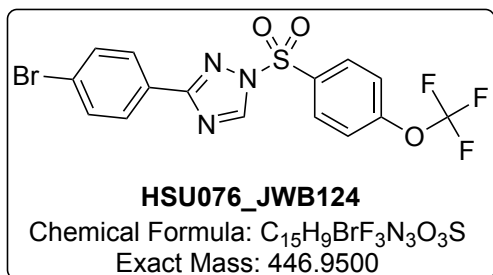
**HRMS** (ESI-TOF)  $m/z$ :  $[\text{M} + \text{H}]^+$  calculated for  $\text{C}_{14}\text{H}_{11}\text{BrN}_3\text{O}_2\text{S}$  363.9750, Found, 363.9744.



**$^1\text{H}$  NMR** (600 MHz, DMSO- $\text{D}_6$ )  $\delta$  9.42 (s, 1H), 8.06 (d,  $J = 8.6$  Hz, 2H), 7.94 (dd,  $J = 17.3, 8.7$  Hz, 4H), 7.04 (d,  $J = 8.7$  Hz, 2H), 3.80 (s, 3H).

**$^{13}\text{C}$  NMR** (201 MHz, DMSO- $\text{D}_6$ )  $\delta$  164.22, 161.33, 152.35, 147.60, 130.67, 128.24, 127.76, 121.66, 121.23, 114.62, 55.45.

**HRMS** (ESI-TOF)  $m/z$ :  $[\text{M} + \text{H}]^+$  calculated for  $\text{C}_{15}\text{H}_{13}\text{BrN}_3\text{O}_3\text{S}$  393.9856, Found, 393.9849.

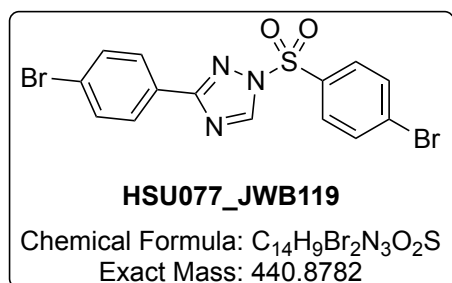


**$^1H$  NMR** (800 MHz, DMSO- $D_6$ )  $\delta$  8.63 (s, 1H), 8.04 – 7.89 (m, 2H), 7.80 – 7.60 (m, 4H), 7.43 – 7.16 (m, 2H).

**$^{13}C$  NMR** (201 MHz, DMSO- $D_6$ )  $\delta$  148.61, 148.09, 146.67, 132.37, 129.43, 128.41, 128.11, 123.24, 121.14, 120.73, 119.87.

**$^{19}F$  NMR** (376 MHz, Acetone- $D_6$ )  $\delta$  -58.50 (s, 3F).

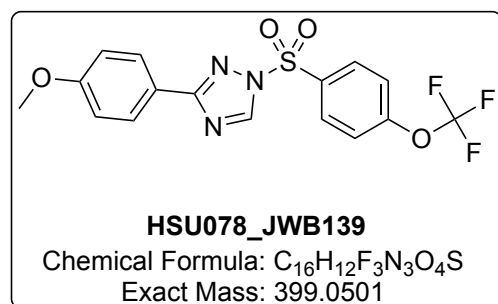
**HRMS** (ESI-TOF)  $m/z$ :  $[M + H]^+$  calculated for  $C_{15}H_{10}BrF_3N_3O_3S$  447.9573, Found, 447.9562.



**$^1H$  NMR** (800 MHz, DMSO- $D_6$ )  $\delta$  8.62 (s, 1H), 8.06 – 7.84 (m, 2H), 7.76 – 7.65 (m, 2H), 7.61 – 7.44 (m, 4H).

**$^{13}C$  NMR** (201 MHz, DMSO- $D_6$ )  $\delta$  157.69, 147.68, 146.17, 131.90, 130.62, 128.95, 127.95, 127.78, 122.77, 121.60.

**HRMS** (ESI-TOF)  $m/z$ :  $[M + H]^+$  calculated for  $C_{14}H_{10}Br_2N_3O_2S$  441.8855, Found, 441.8853.

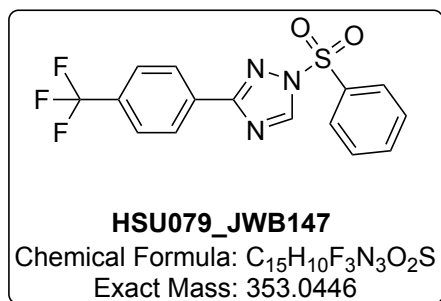


**$^1H$  NMR** (800 MHz, DMSO- $D_6$ )  $\delta$  9.41 (s, 1H), 8.57 – 8.09 (m, 2H), 8.04 – 7.84 (m, 2H), 7.71 (dd,  $J$  = 9.1, 1.1 Hz, 2H), 7.25 – 6.76 (m, 2H), 3.81 (s, 3H).

**$^{13}C$  NMR** (201 MHz, DMSO- $D_6$ )  $\delta$  164.08, 161.21, 153.06, 147.66, 133.93, 131.14, 128.17, 121.92, 120.92, 114.37, 55.26.

**$^{19}F$  NMR** (376 MHz, Acetone- $D_6$ )  $\delta$  -58.50 (s, 3F).

**HRMS** (ESI-TOF)  $m/z$ :  $[M + H]^+$  calculated for  $C_{16}H_{11}F_3N_3O_4S$  400.0573, Found, 400.0568.

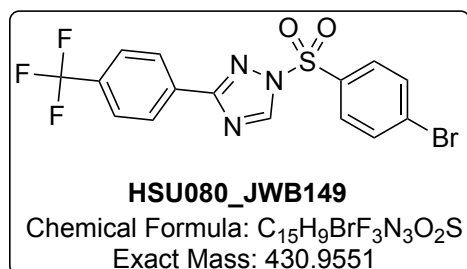


**$^1H$  NMR** (600 MHz, DMSO- $D_6$ )  $\delta$  9.57 (s, 1H), 8.18 (t,  $J = 9.3$  Hz, 4H), 7.87 (m, 3H), 7.75 (t,  $J = 7.9$  Hz, 2H).

**$^{13}C$  NMR** (201 MHz, DMSO- $D_6$ )  $\delta$  162.92, 158.08, 152.90, 148.18, 146.10, 128.49, 127.68, 127.35, 126.56, 126.20, 125.86 (q,  $J = 3.7$  Hz), 125.49.

**$^{19}F$  NMR** (376 MHz, acetone)  $\delta$  -63.41 (s, 3F).

**HRMS** (ESI-TOF)  $m/z$ :  $[M + H]^+$  calculated for  $C_{15}H_{11}F_3N_3O_2S$  354.0519, Found, 354.0519.

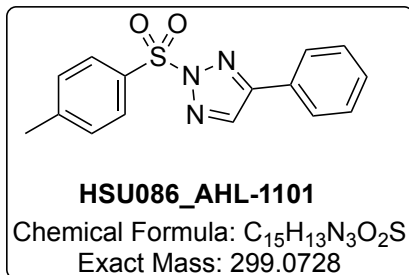


**$^1H$  NMR** (800 MHz, DMSO- $D_6$ )  $\delta$  8.67 (s, 1H), 8.23 (d,  $J = 8.0$  Hz, 2H), 7.86 (d,  $J = 7.9$  Hz, 2H), 7.61 – 7.36 (m, 4H).

**$^{13}C$  NMR** (201 MHz, DMSO- $D_6$ )  $\delta$  158.12, 147.65, 146.10, 134.08, 130.64, 129.30 (q,  $J = 31.6$  Hz), 127.76, 126.55, 125.86 (q,  $J = 3.9$  Hz), 124.18 (d,  $J = 272.4$  Hz), 121.63.

**$^{19}F$  NMR** (376 MHz, Acetone- $D_6$ )  $\delta$  -63.41 (s, 3F).

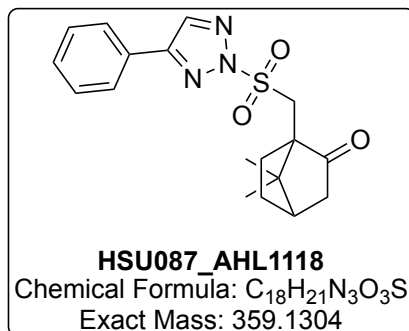
**HRMS** (ESI-TOF)  $m/z$ :  $[M + H]^+$  calculated for  $C_{15}H_{10}BrF_3N_3O_2S$  431.9624, Found, 431.9614.



**$^1H$  NMR** (400 MHz, Aceton-D6)  $\delta$  8.53 (d,  $J$  = 1.0 Hz, 1H), 7.99 (d,  $J$  = 8.5 Hz, 2H), 7.94 (dd,  $J$  = 6.9, 1.4 Hz, 2H), 7.55 – 7.39 (m, 5H), 2.42 (s, 3H).

**$^{13}C$  NMR** (100 MHz, Acetone-D6)  $\delta$  152.51, 148.04, 137.40, 133.97, 131.30, 130.89, 129.98, 129.32, 127.48, 21.62.

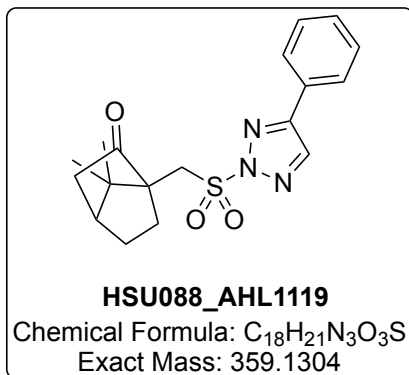
**HRMS** (ESI-TOF)  $m/z$ :  $[M + H]^+$  calculated for  $C_{15}H_{14}N_3O_2S$  300.0801, Found, 300.0793.



**$^1H$  NMR** (400 MHz, Acetone-D6)  $\delta$  8.59 (s, 1H), 8.00 (d,  $J$  = 6.5 Hz, 2H), 7.60 – 7.45 (m, 3H), 4.03 (d,  $J$  = 15.2 Hz, 1H), 3.72 (d,  $J$  = 15.2 Hz, 1H), 2.43 – 2.25 (m, 2H), 2.16 (t,  $J$  = 4.5 Hz, 1H), 2.13 – 2.07 (m, 1H), 1.94 (d,  $J$  = 18.4 Hz, 1H), 1.76 (ddd,  $J$  = 13.8, 9.4, 4.6 Hz, 1H), 1.52 (ddd,  $J$  = 12.9, 9.3, 3.8 Hz, 1H), 1.12 (s, 3H), 0.91 (s, 3H).

**$^{13}C$  NMR** (100 MHz, Aceton-D6)  $\delta$  213.17, 151.90, 136.63, 130.80, 130.01, 129.60, 127.51, 59.19, 52.69, 49.06, 43.48, 42.69, 27.37, 25.76, 19.72, 19.70.

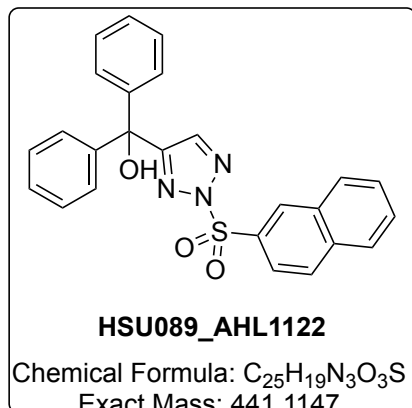
**HRMS** (ESI-TOF)  $m/z$ :  $[M + H]^+$  calculated for  $C_{18}H_{22}N_3O_3S$  360.1376, Found, 360.1363.



**<sup>1</sup>H NMR** (400 MHz, CDCl<sub>3</sub>) δ 8.18 (s, 1H), 7.90 (dd, *J* = 8.0, 1.7 Hz, 2H), 7.53 – 7.42 (m, 3H), 3.99 (d, *J* = 15.0 Hz, 1H), 3.43 (d, *J* = 14.9 Hz, 1H), 2.49 (ddd, *J* = 14.9, 11.7, 4.0 Hz, 1H), 2.39 (dt, *J* = 18.6, 4.0 Hz, 1H), 2.19 – 2.03 (m, 2H), 1.94 (d, *J* = 18.6 Hz, 1H), 1.80 (ddd, *J* = 13.9, 9.3, 4.5 Hz, 1H), 1.47 (ddd, *J* = 13.1, 9.2, 3.8 Hz, 1H), 1.18 (s, 3H), 0.91 (s, 3H).

**<sup>13</sup>C NMR** (151 MHz, DMSO-D<sub>6</sub>) δ 213.02, 150.63, 136.47, 130.10, 129.25, 128.11, 126.60, 58.14, 51.78, 48.35, 42.06, 41.79, 26.24, 24.72, 19.09, 18.95.

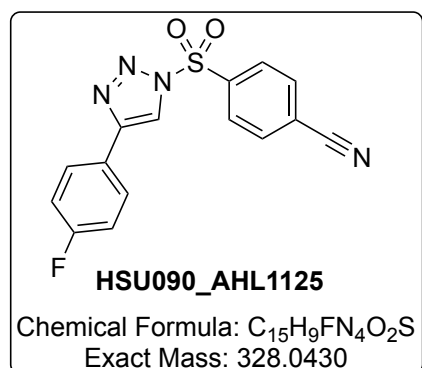
**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>18</sub>H<sub>22</sub>N<sub>3</sub>O<sub>3</sub>S 360.1376, Found, 360.1363.



**<sup>1</sup>H NMR** (400 MHz, DMSO-D<sub>6</sub>) δ 8.80 (d, *J* = 2.5 Hz, 1H), 8.31 (d, *J* = 7.4 Hz, 1H), 8.21 (d, *J* = 9.0 Hz, 1H), 8.17 (s, 1H), 8.11 (d, *J* = 8.1 Hz, 1H), 7.88 – 7.79 (m, 2H), 7.76 (ddd, *J* = 8.2, 6.9, 1.4 Hz, 1H), 7.23 – 7.11 (m, 10H), 6.97 (s, 1H).

**<sup>13</sup>C NMR** (100 MHz, DMSO-D<sub>6</sub>) δ 160.07, 145.11, 139.77, 135.39, 131.81, 131.53, 130.67, 130.57, 130.29, 129.95, 128.39, 128.08, 127.80, 127.28, 126.74, 121.78, 75.93.

**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>25</sub>H<sub>20</sub>N<sub>3</sub>O<sub>3</sub>S 442.1220, Found, 442.1204.

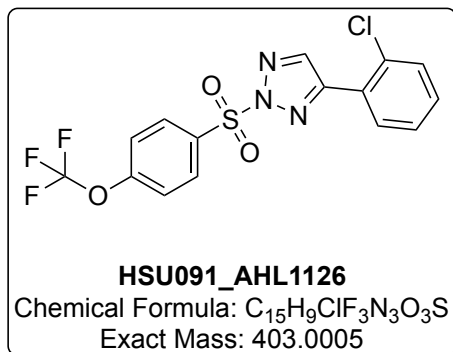


**<sup>1</sup>H NMR** (400 MHz, Acetone-D<sub>6</sub>) δ 8.63 (s, 1H), 8.30 (d, *J* = 8.9 Hz, 2H), 8.14 (d, *J* = 8.8 Hz, 2H), 8.02 (dd, *J* = 8.8, 5.3 Hz, 2H), 7.29 (t, *J* = 8.8 Hz, 2H).

**<sup>13</sup>C NMR** (100 MHz, Acetone-D<sub>6</sub>) δ 164.74 (d, *J* = 248.7 Hz), 163.50, 152.52, 140.57, 138.43, 134.77, 130.01, 130.00 (d, *J* = 8.7 Hz), 125.50 (d, *J* = 3.2 Hz), 125.48, 119.60, 117.60, 117.03 (d, *J* = 22.2 Hz).

**<sup>19</sup>F NMR** (376 MHz, Acetone-D<sub>6</sub>) δ -111.64 – -111.76 (m, 1F).

**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>15</sub>H<sub>10</sub>FN<sub>4</sub>O<sub>2</sub>S 329.0503, Found, 329.0499.

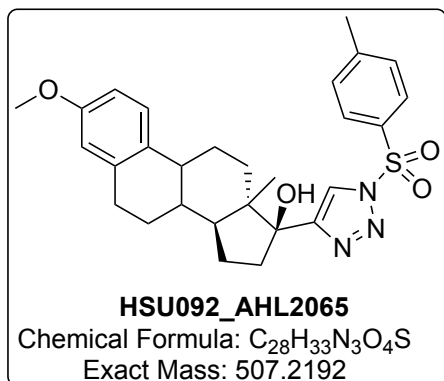


**$^1H$  NMR** (400 MHz,  $CDCl_3$ )  $\delta$  8.38 (s, 1H), 8.20 (d,  $J = 9.1$  Hz, 2H), 7.98 – 7.81 (m, 1H), 7.57 – 7.43 (m, 1H), 7.43 – 7.32 (m, 4H).

**$^{13}C$  NMR** (151 MHz, DMSO- $D_6$ )  $\delta$  153.24, 149.41, 139.95, 133.25, 131.79, 131.73, 131.34, 131.24, 131.21, 130.45, 127.84, 126.65, 122.02.

**$^{19}F$  NMR** (376 MHz, Acetone- $D_6$ )  $\delta$  -58.51 (s, 3F).

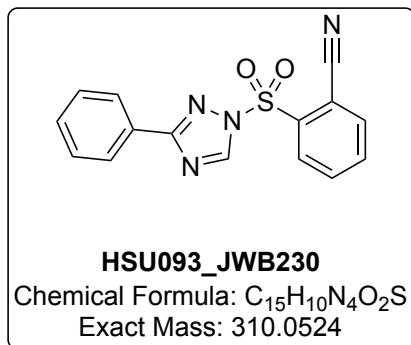
**HRMS** (ESI-TOF)  $m/z$ :  $[M + H]^+$  calculated for  $C_{15}H_{10}ClF_3N_3O_3S$  404.0078, Found, 404.0076.



**$^1H$  NMR** (400 MHz, DMSO- $D_6$ )  $\delta$  7.92 (d,  $J = 8.6$  Hz, 2H), 7.72 (s, 1H), 7.50 (d,  $J = 9.4$  Hz, 2H), 7.10 (d,  $J = 8.2$  Hz, 1H), 6.65 (dd,  $J = 8.6, 2.9$  Hz, 1H), 6.58 (d,  $J = 2.9$  Hz, 1H), 5.51 (s, 1H), 3.67 (s, 3H), 3.38 (d,  $J = 7.0$  Hz, 1H), 2.76 (d,  $J = 5.4$  Hz, 2H), 2.43 (s, 3H), 2.29 – 2.05 (m, 4H), 1.96 – 1.77 (m, 3H), 1.52 – 1.18 (m, 5H), 0.98 (s, 3H), 0.59 – 0.49 (m, 1H).

**$^{13}C$  NMR** (100 MHz, DMSO- $D_6$ )  $\delta$  157.05, 146.39, 146.33, 137.33, 134.38, 133.80, 131.93, 130.07, 129.07, 126.16, 113.37, 111.48, 82.06, 54.86, 48.77, 47.89, 42.81, 40.73, 33.62, 29.25, 26.90, 26.14, 22.47, 21.25, 13.86.

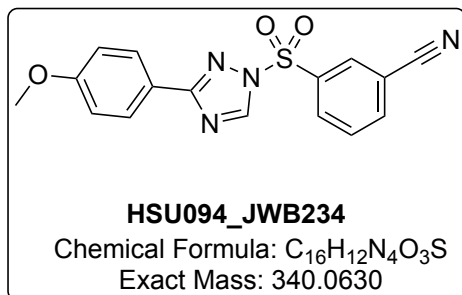
**HRMS** (ESI-TOF)  $m/z$ :  $[M + H]^+$  calculated for  $C_{28}H_{34}N_3O_4S$  508.2265, Found, 508.2258.



**<sup>1</sup>H NMR** (800 MHz, DMSO-D<sub>6</sub>) δ 8.79 (s, 1H), 8.07 – 7.98 (m, 2H), 7.87 (dd, *J* = 7.8, 1.3 Hz, 2H), 7.76 (dd, *J* = 7.6, 1.3 Hz, 2H), 7.64 (td, *J* = 7.7, 1.3 Hz, 2H), 7.56 – 7.52 (m, 2H), 7.52 – 7.47 (m, 2H).

**<sup>13</sup>C NMR** (201 MHz, DMSO-D<sub>6</sub>) δ 156.64, 150.35, 145.86, 133.95, 132.27, 130.10, 129.04, 128.93, 127.05, 126.55, 126.17, 117.67, 108.95.

**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>15</sub>H<sub>11</sub>N<sub>4</sub>O<sub>2</sub>S 311.0597, Found, 311.0596.

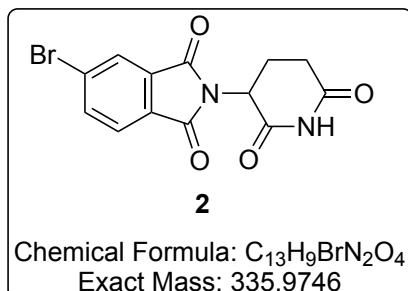


**<sup>1</sup>H NMR** (800 MHz, CDCl<sub>3</sub>) δ 8.72 (s, 1H), 8.42 (t, *J* = 1.7 Hz, 1H), 8.34 (ddd, *J* = 8.1, 1.9, 1.1 Hz, 1H), 8.03 (d, *J* = 8.9 Hz, 2H), 7.98 (dt, *J* = 7.7, 1.3 Hz, 1H), 7.76 (t, *J* = 7.9 Hz, 2H), 6.95 (d, *J* = 8.8 Hz, 2H), 3.85 (s, 3H).

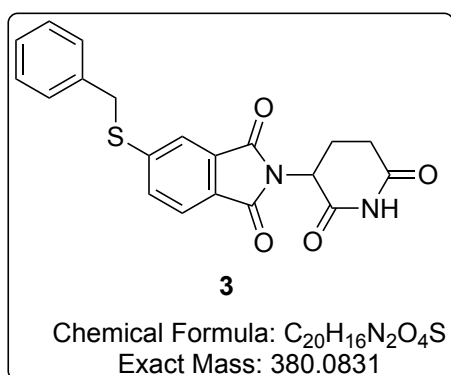
**<sup>13</sup>C NMR** (201 MHz, CDCl<sub>3</sub>) δ 165.85, 161.89, 145.65, 138.30, 138.08, 132.48, 132.26, 130.93, 128.92, 121.38, 116.49, 114.69, 114.27, 55.55.

**HRMS** (ESI-TOF) *m/z*: [M + H]<sup>+</sup> calculated for C<sub>16</sub>H<sub>12</sub>N<sub>4</sub>O<sub>3</sub>S 341.0703, Found, 341.0694.

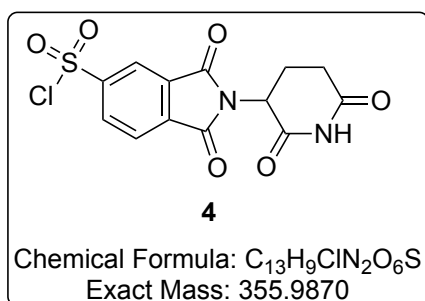




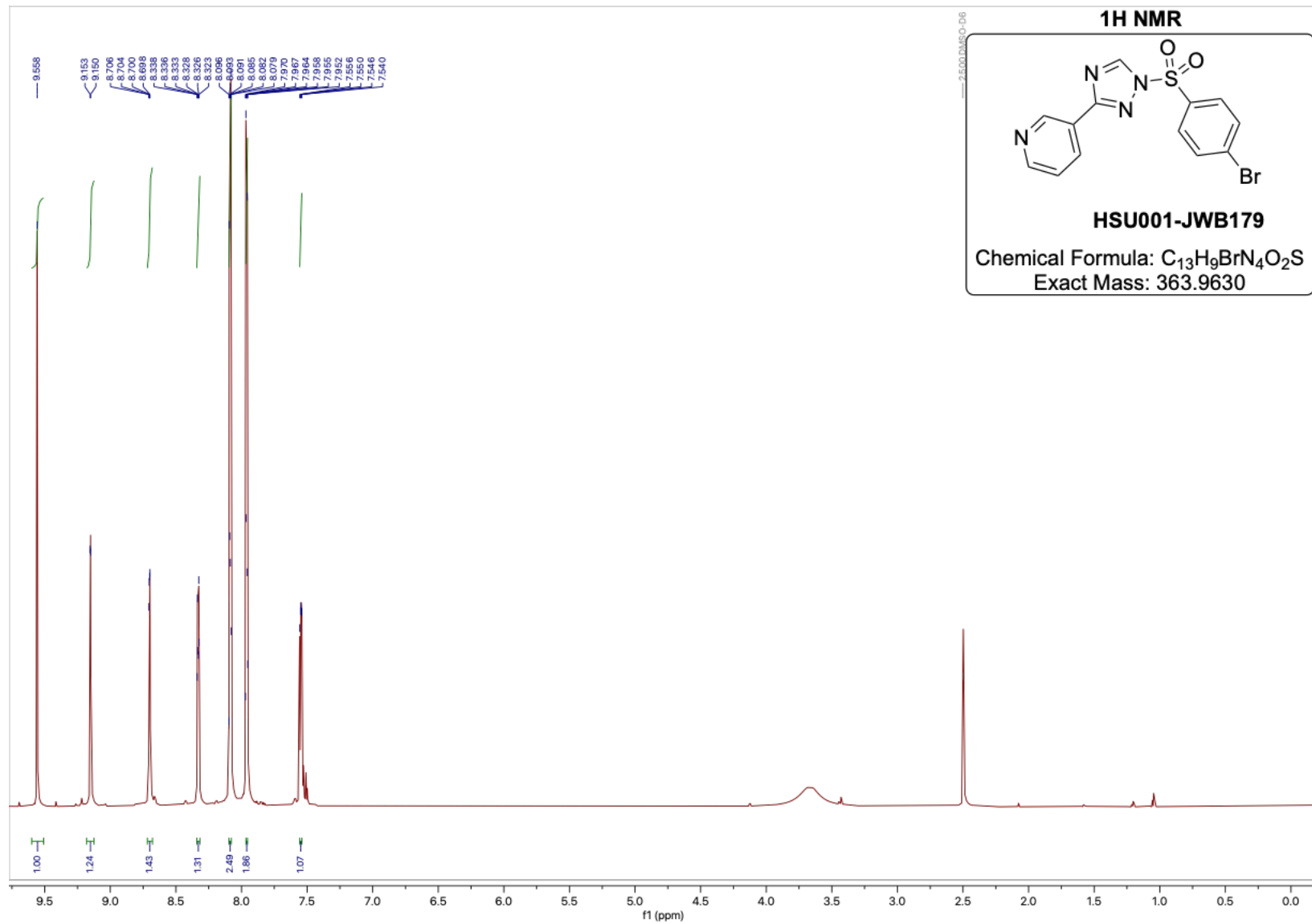
**$^1H$  NMR** (600 MHz, DMSO- $D_6$ )  $\delta$  11.14 (s, 1H), 8.15 (d,  $J$  = 1.8 Hz, 1H), 8.10 (dd,  $J$  = 7.9, 1.7 Hz, 1H), 7.87 (d,  $J$  = 7.9 Hz, 1H), 5.16 (dd,  $J$  = 13.0, 5.4 Hz, 1H), 2.89 (ddd,  $J$  = 17.1, 13.9, 5.4 Hz, 1H), 2.61 (dd,  $J$  = 17.4, 4.0 Hz, 1H), 2.56 – 2.48 (m, 1H), 2.06 (dtd,  $J$  = 13.0, 5.4, 2.4 Hz, 1H).

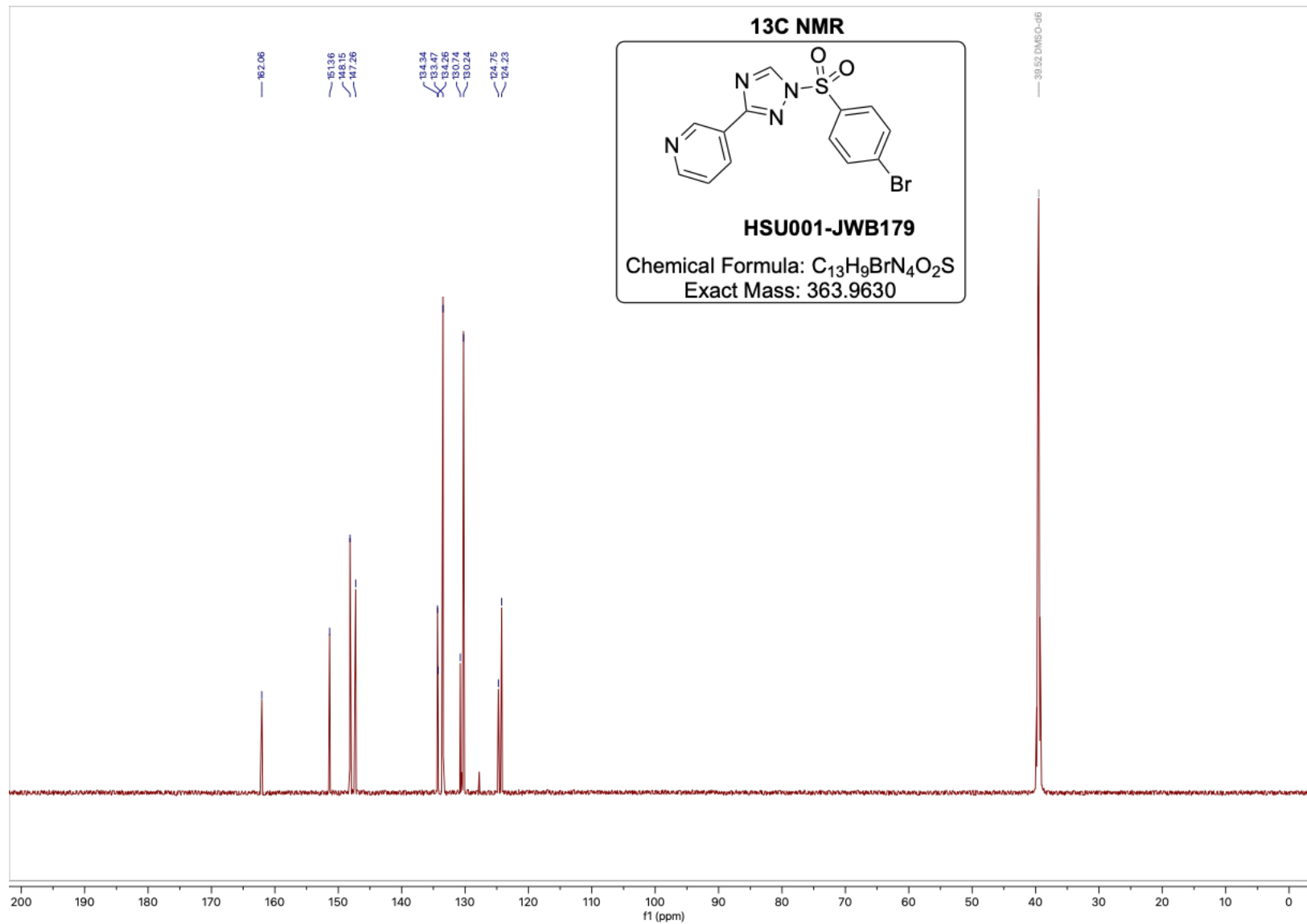


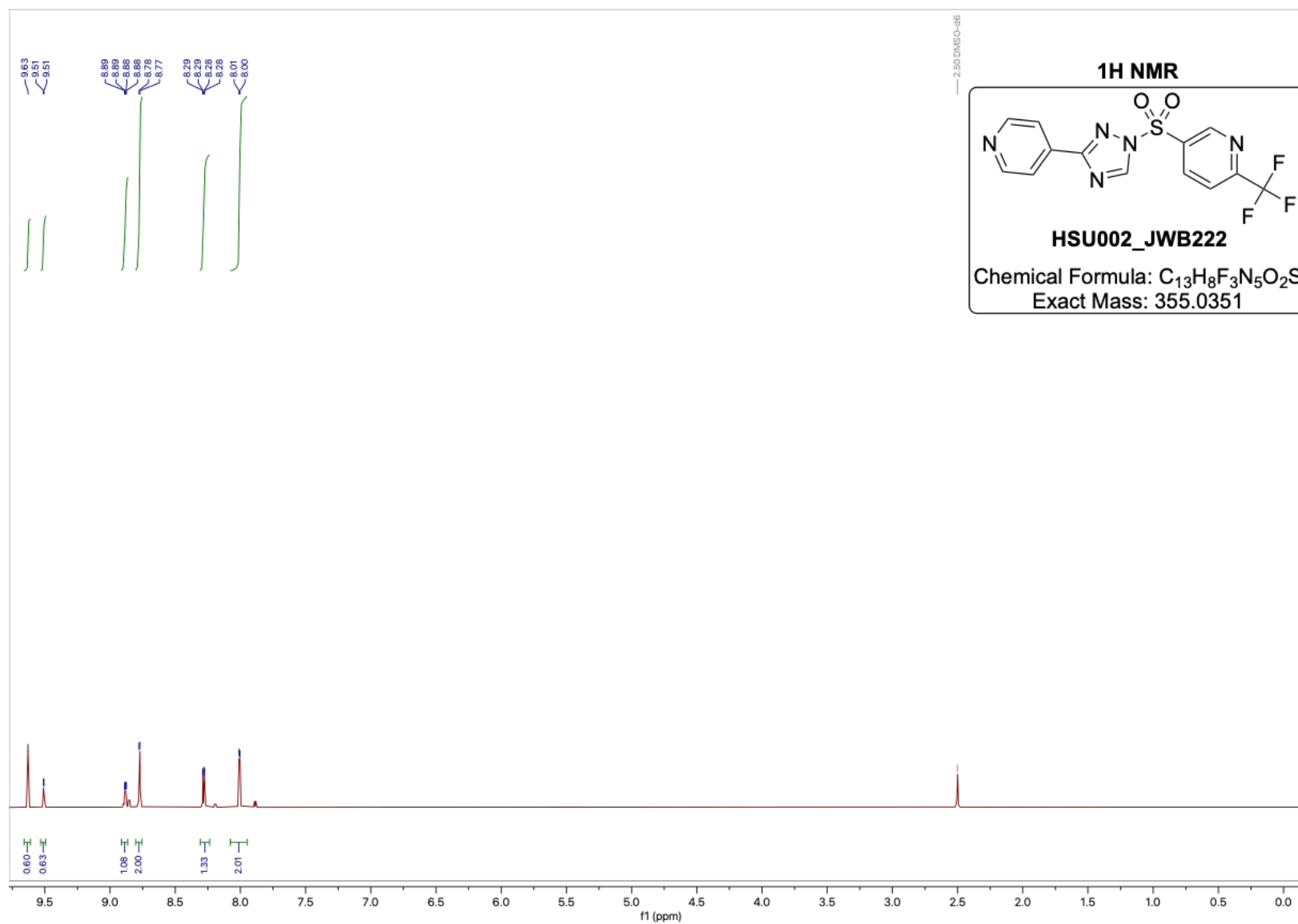
**$^1H$  NMR** (600 MHz, DMSO- $D_6$ )  $\delta$  11.09 (s, 1H), 7.93 – 7.57 (m, 3H), 7.53 – 7.36 (m, 2H), 7.31 (t,  $J$  = 7.6 Hz, 2H), 7.27 – 7.17 (m, 1H), 5.10 (dd,  $J$  = 12.9, 5.4 Hz, 1H), 4.47 (s, 2H), 2.85 (ddd,  $J$  = 17.1, 13.9, 5.4 Hz, 1H), 2.57 (d,  $J$  = 17.3 Hz, 1H), 2.49 (m, 2H), 2.12 – 1.97 (m, 1H).

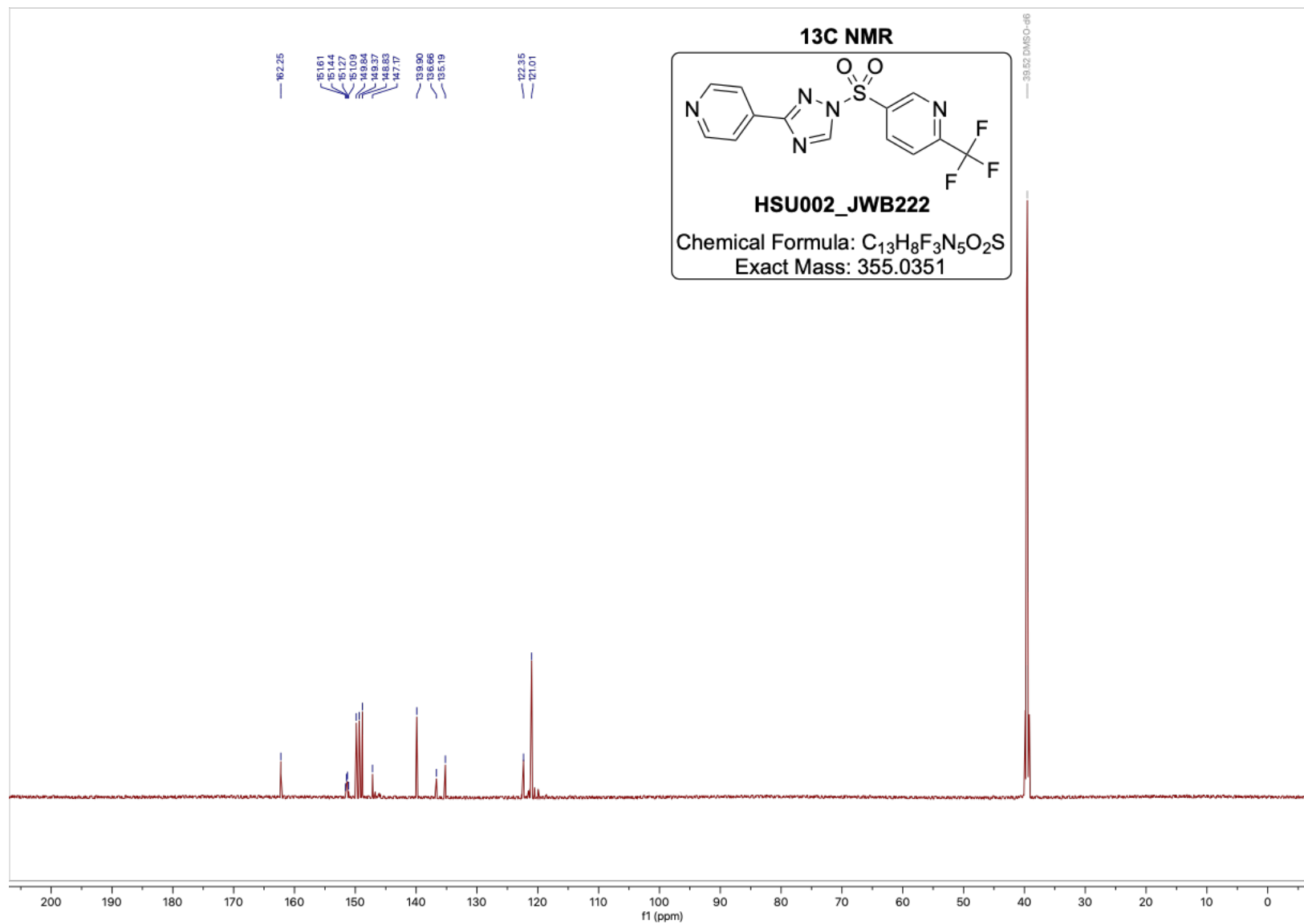


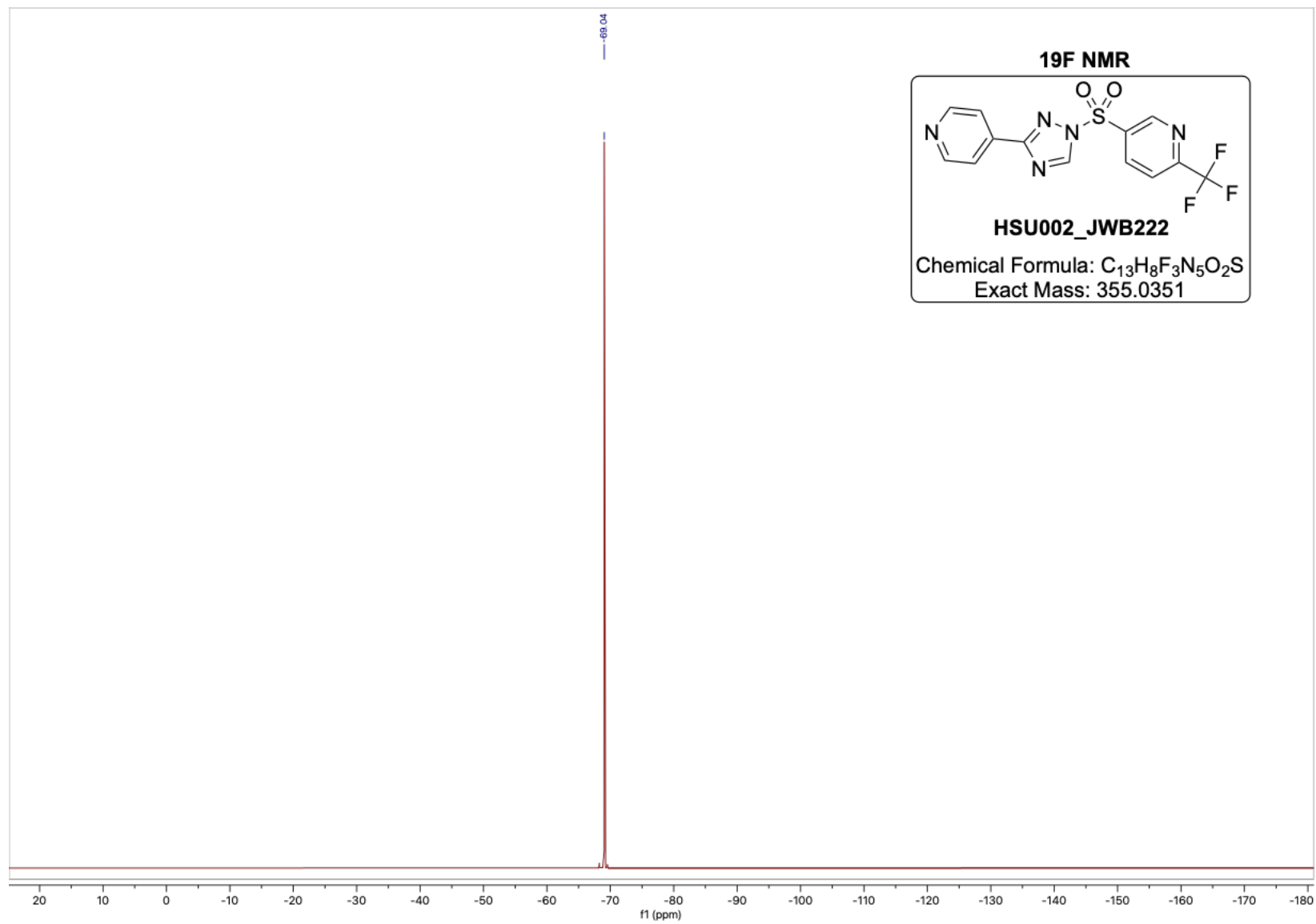
**$^1H$  NMR** (600 MHz, DMSO- $D_6$ )  $\delta$  11.12 (s, 1H), 8.07 (dd,  $J$  = 7.6, 1.3 Hz, 1H), 7.96 (s, 1H), 7.90 (d,  $J$  = 7.7 Hz, 1H), 5.15 (dd,  $J$  = 12.9, 5.5 Hz, 1H), 2.89 (ddd,  $J$  = 17.9, 13.8, 5.5 Hz, 1H), 2.70 – 2.52 (m, 2H), 2.16 – 2.03 (m, 1H).

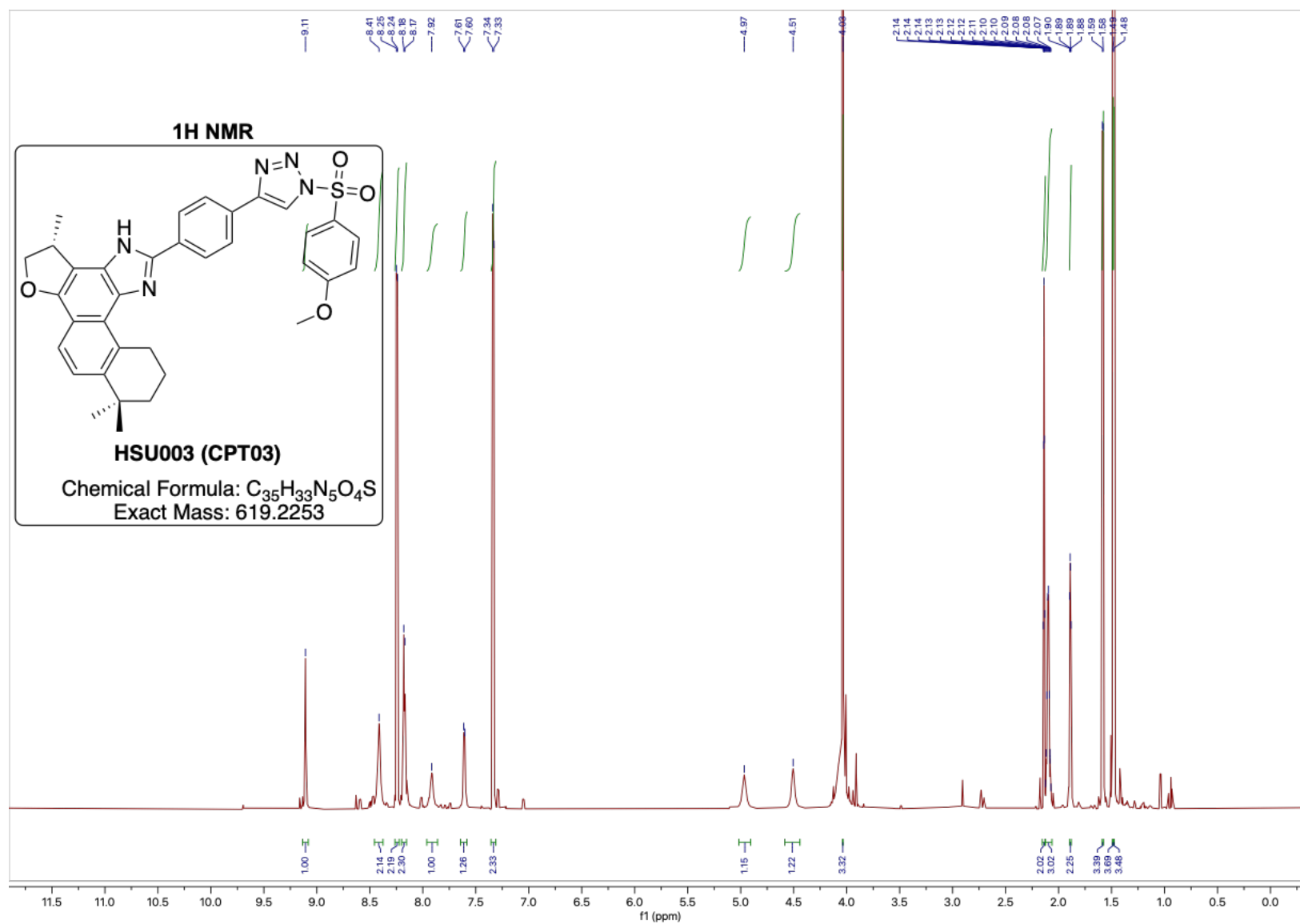


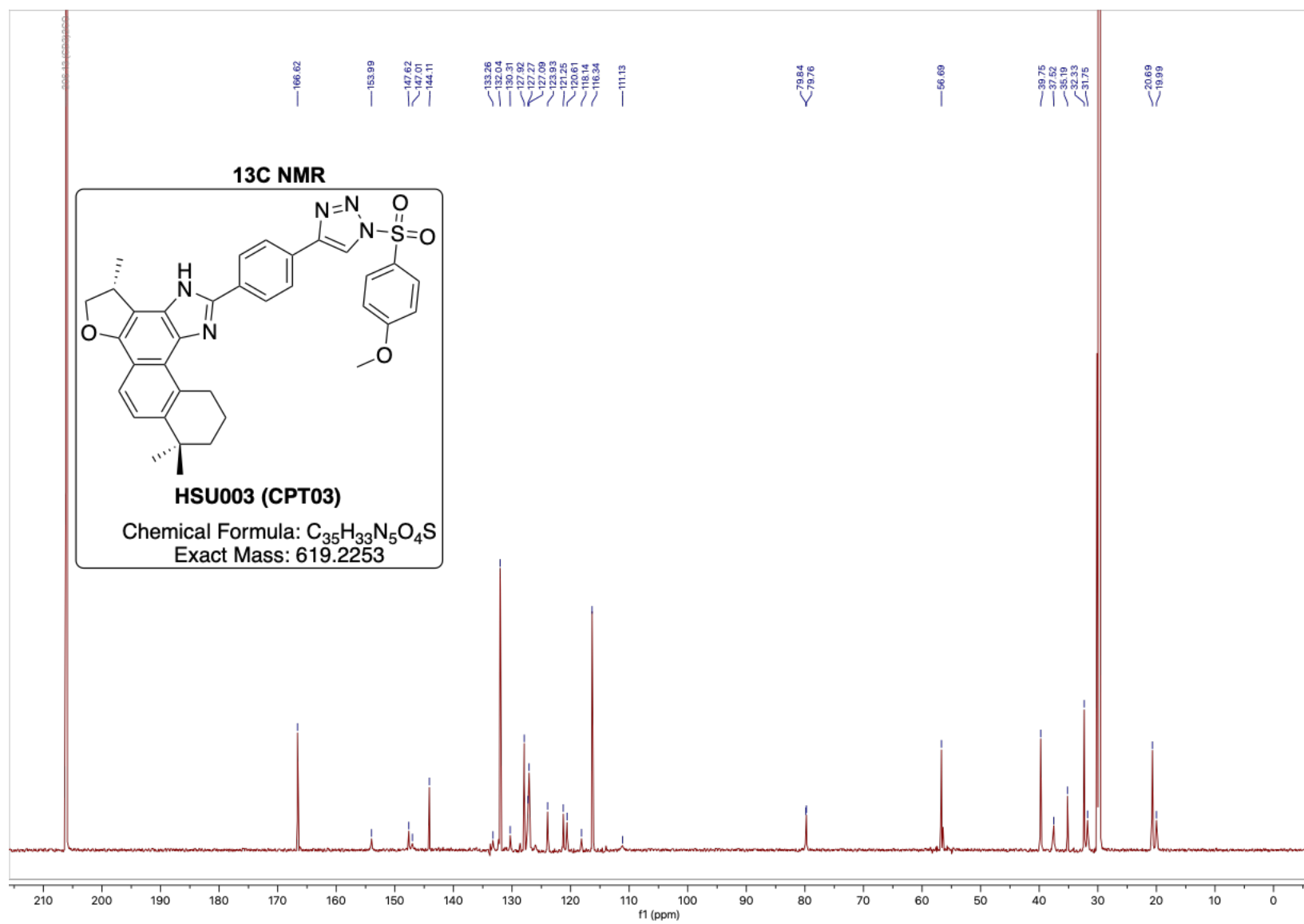




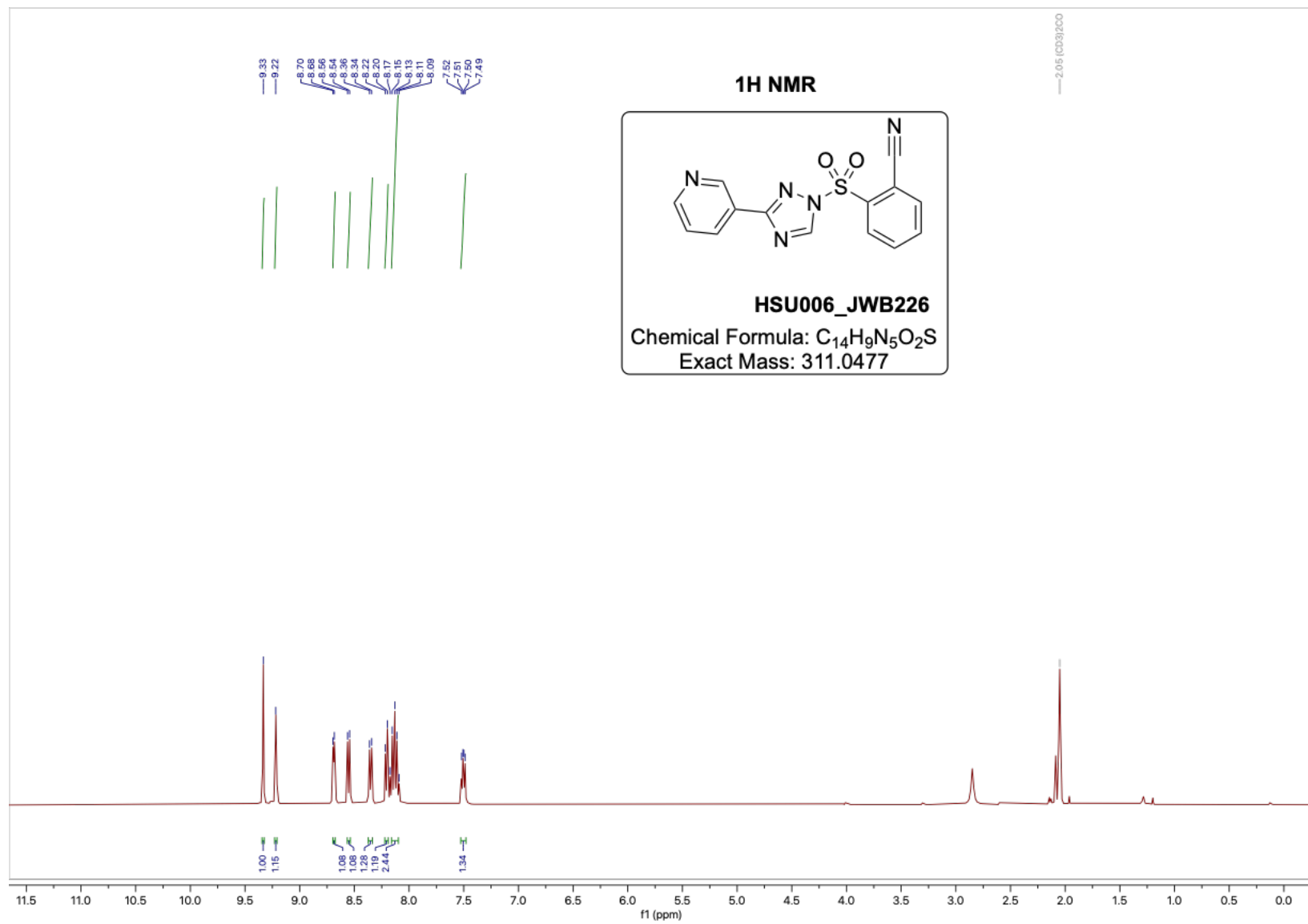


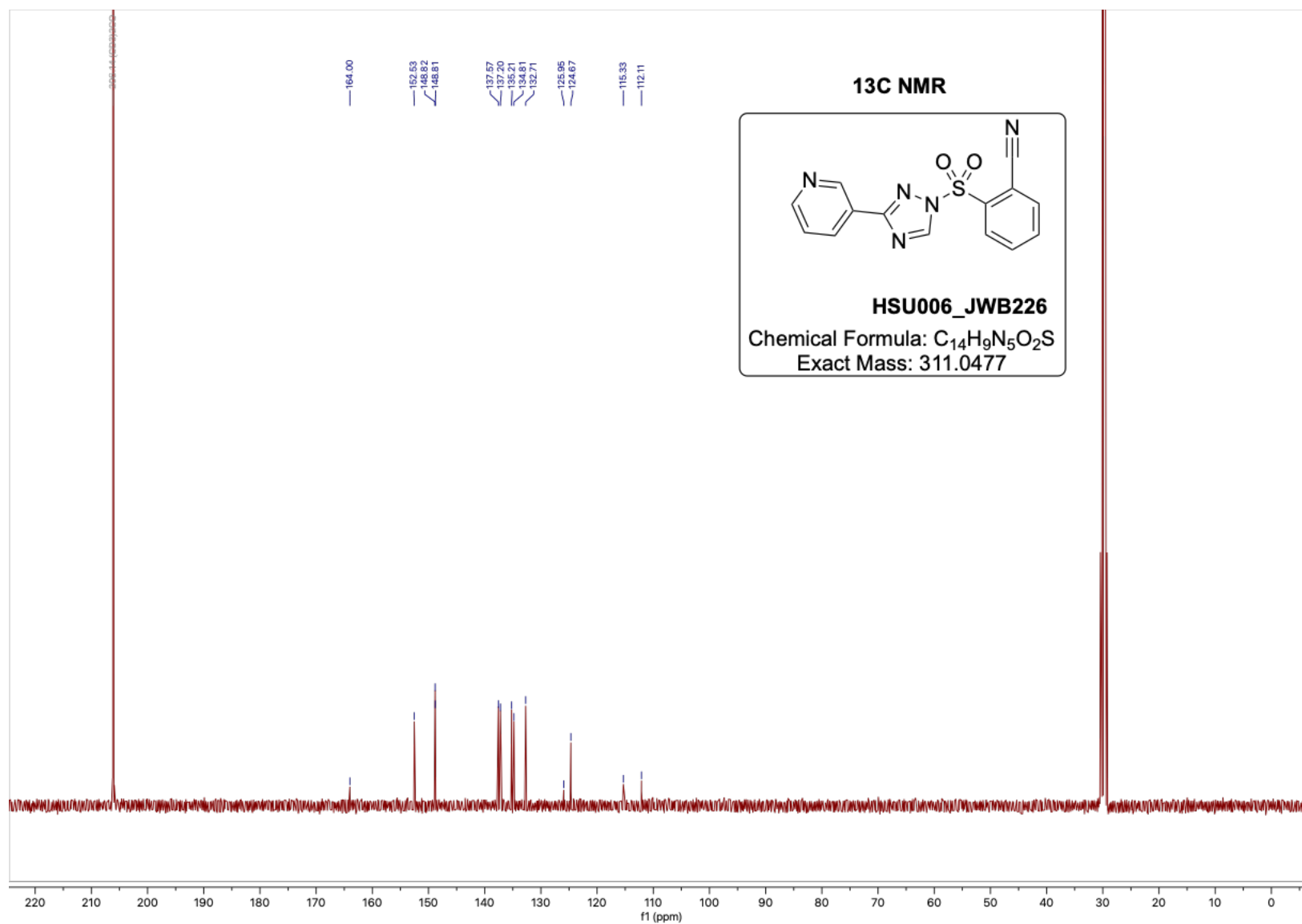


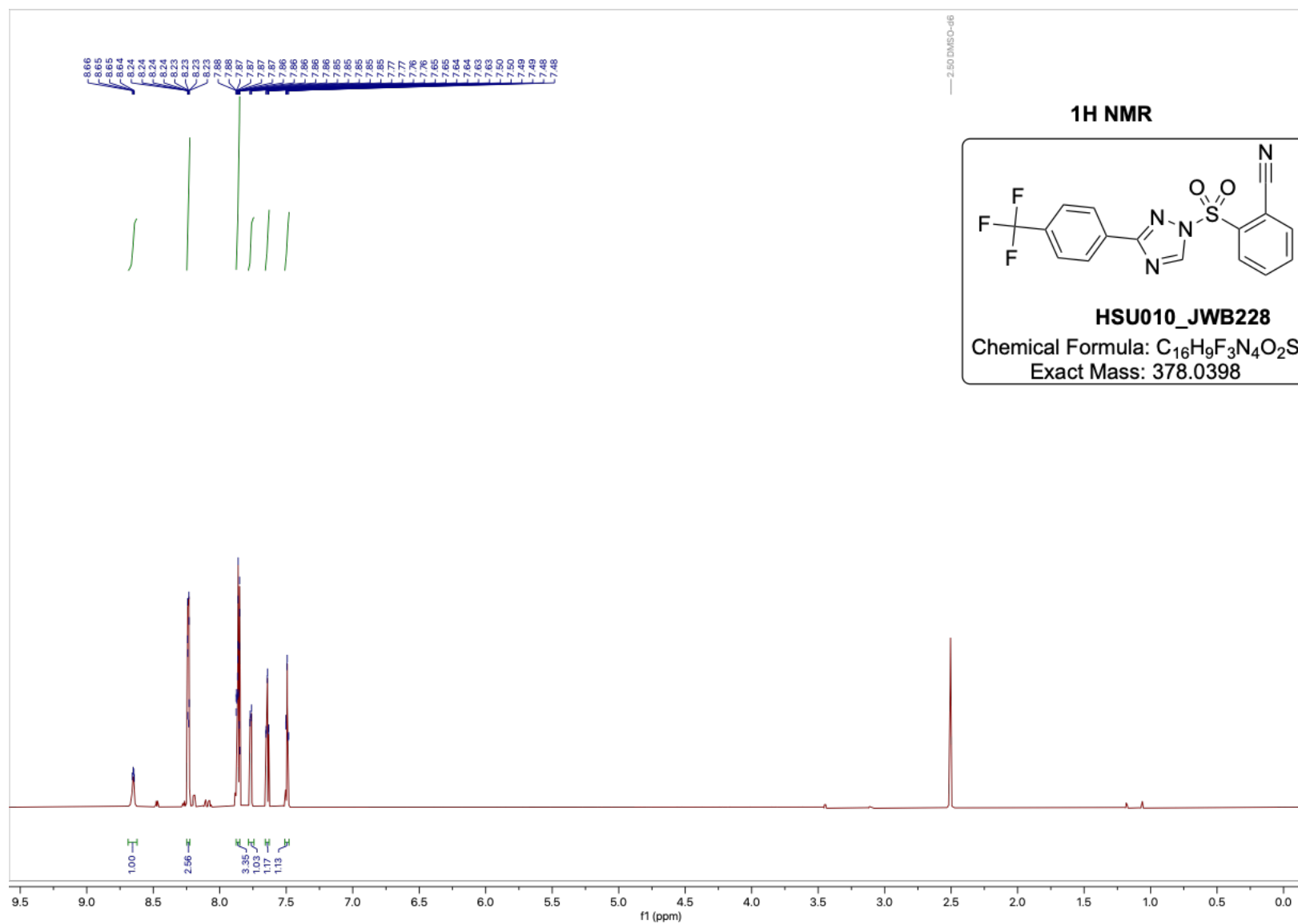




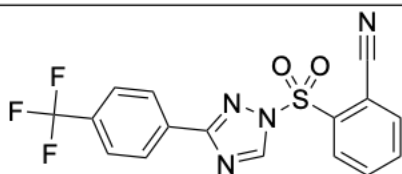






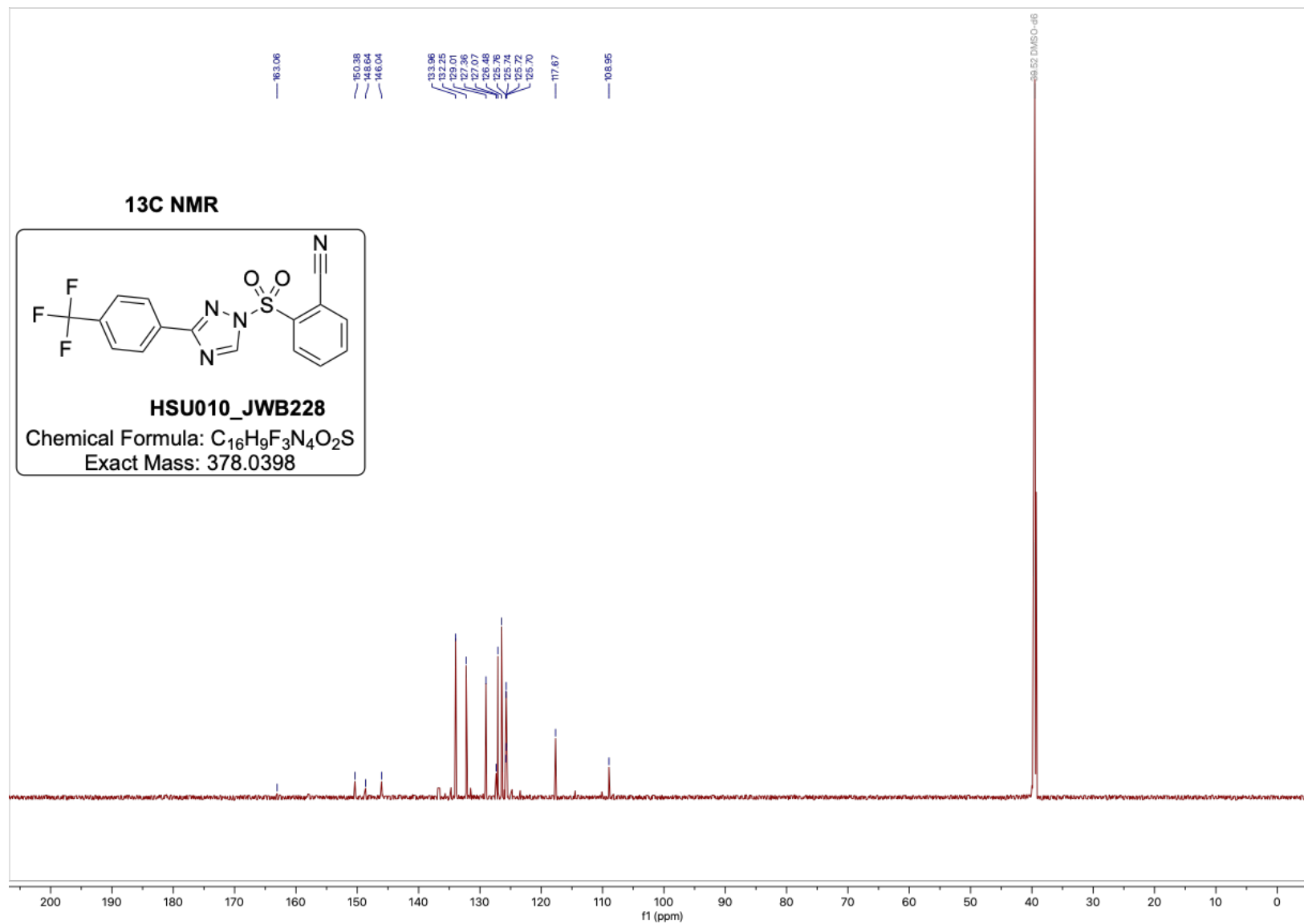


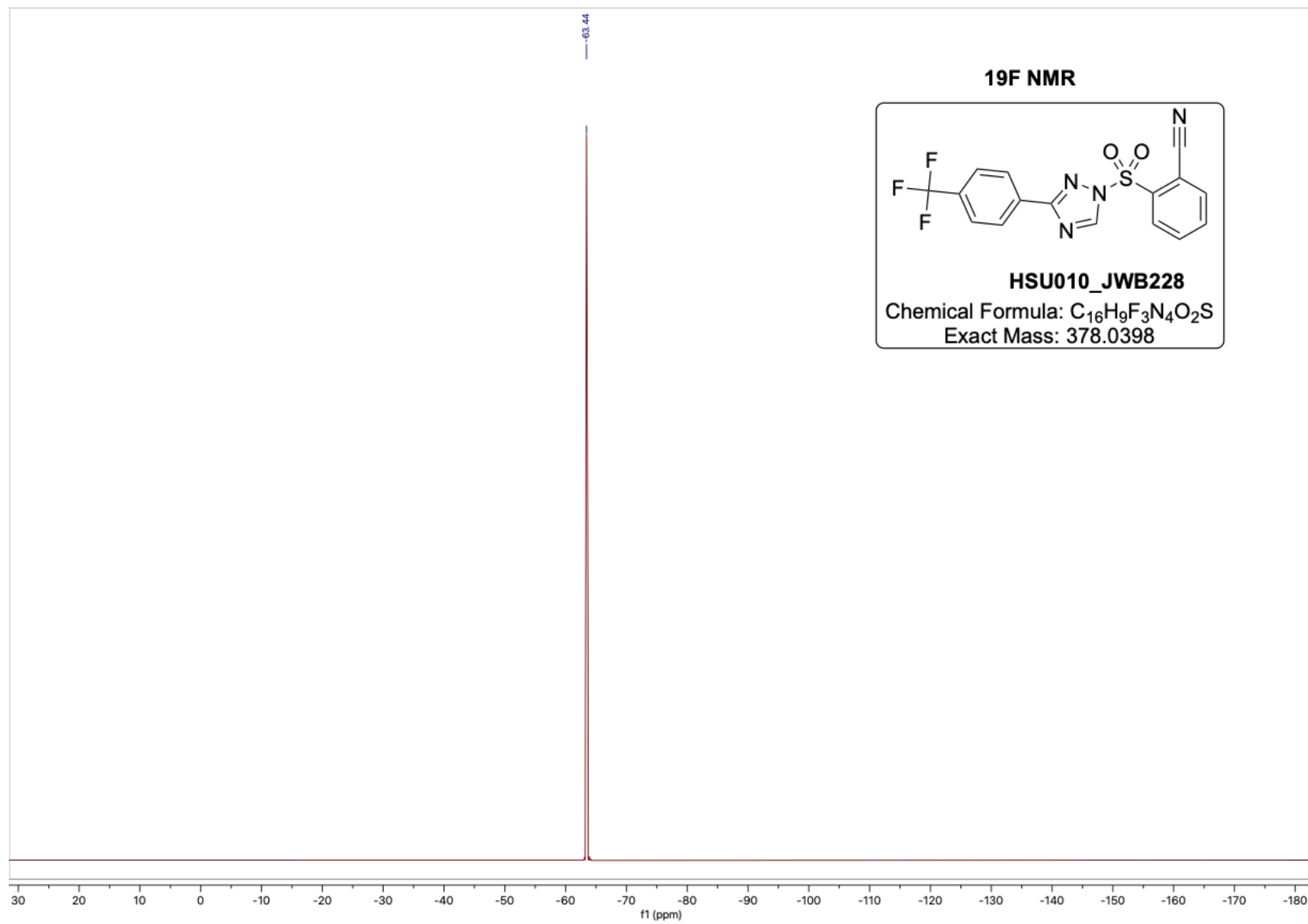
# **<sup>13</sup>C NMR**

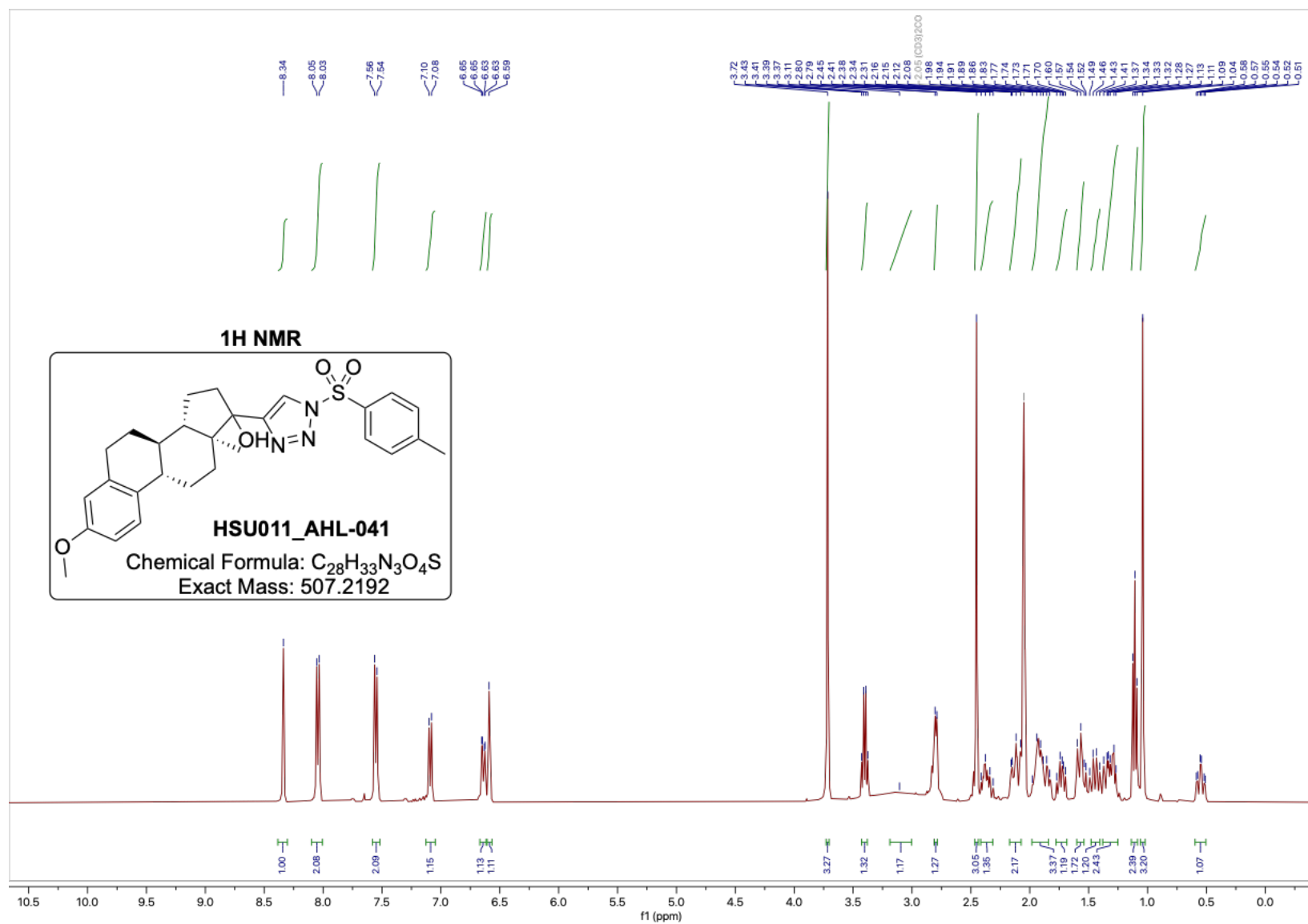


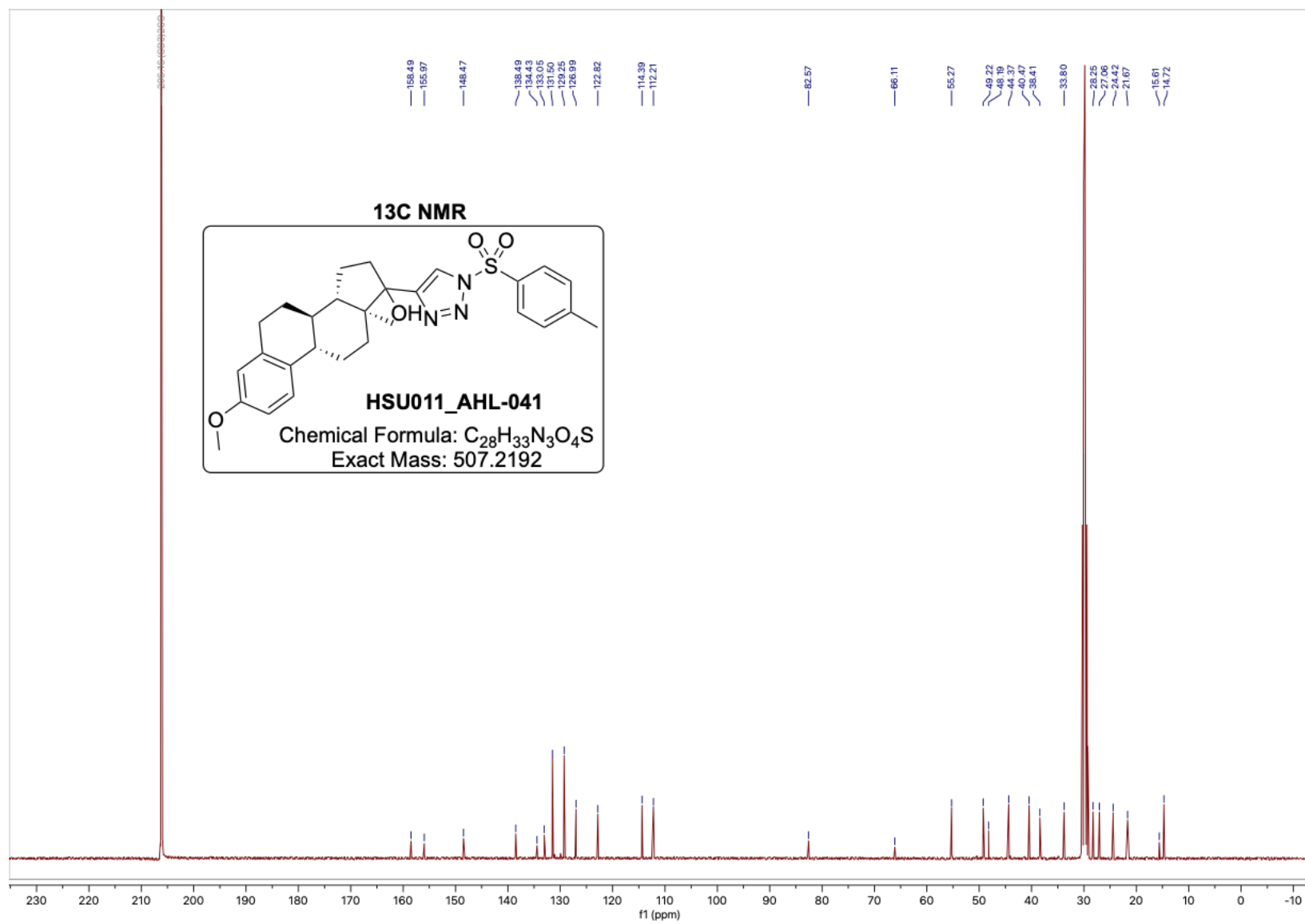
**HSU010\_JWB228**

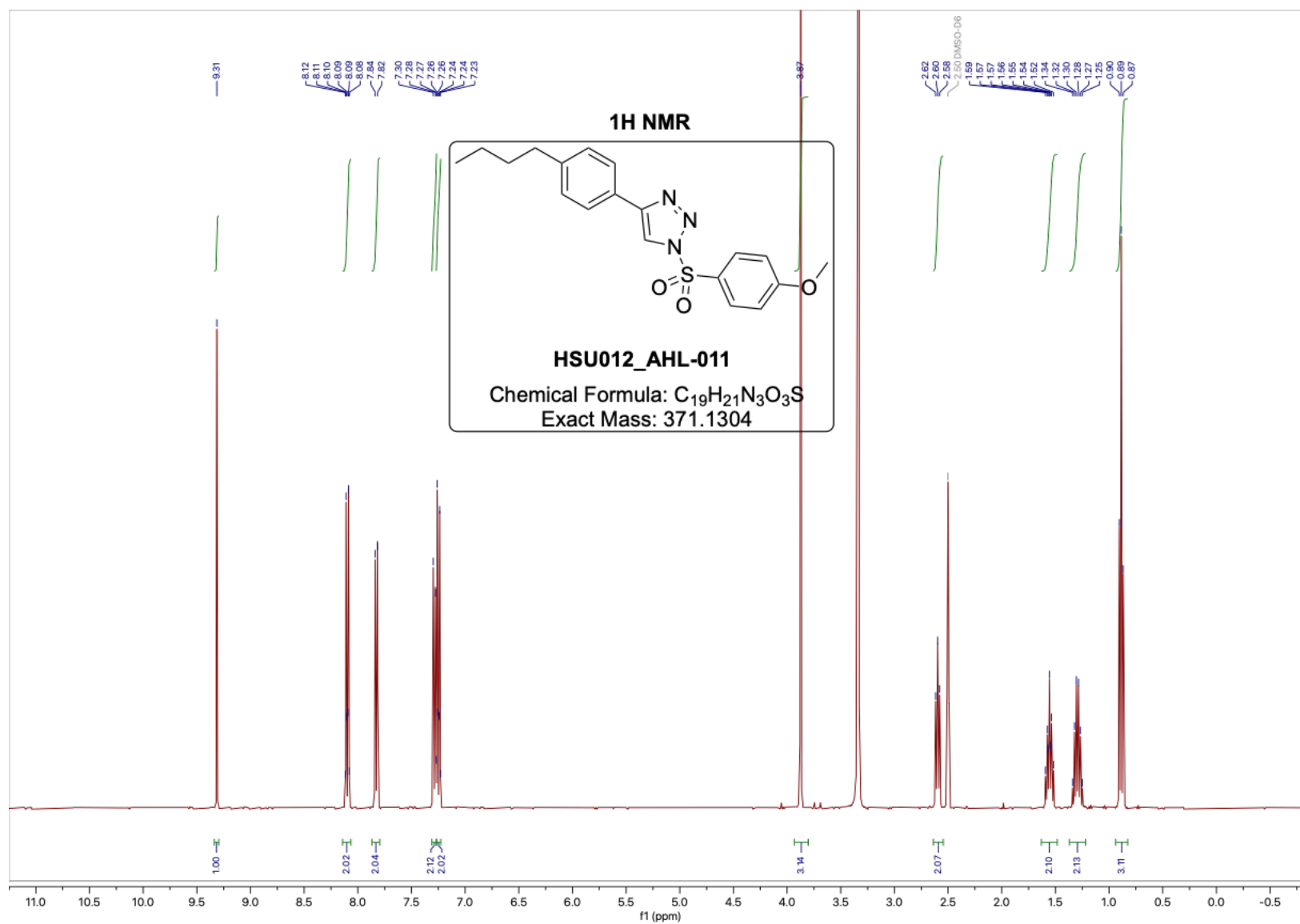
Chemical Formula: C<sub>16</sub>H<sub>9</sub>F<sub>3</sub>N<sub>4</sub>O<sub>2</sub>S  
Exact Mass: 378.0398



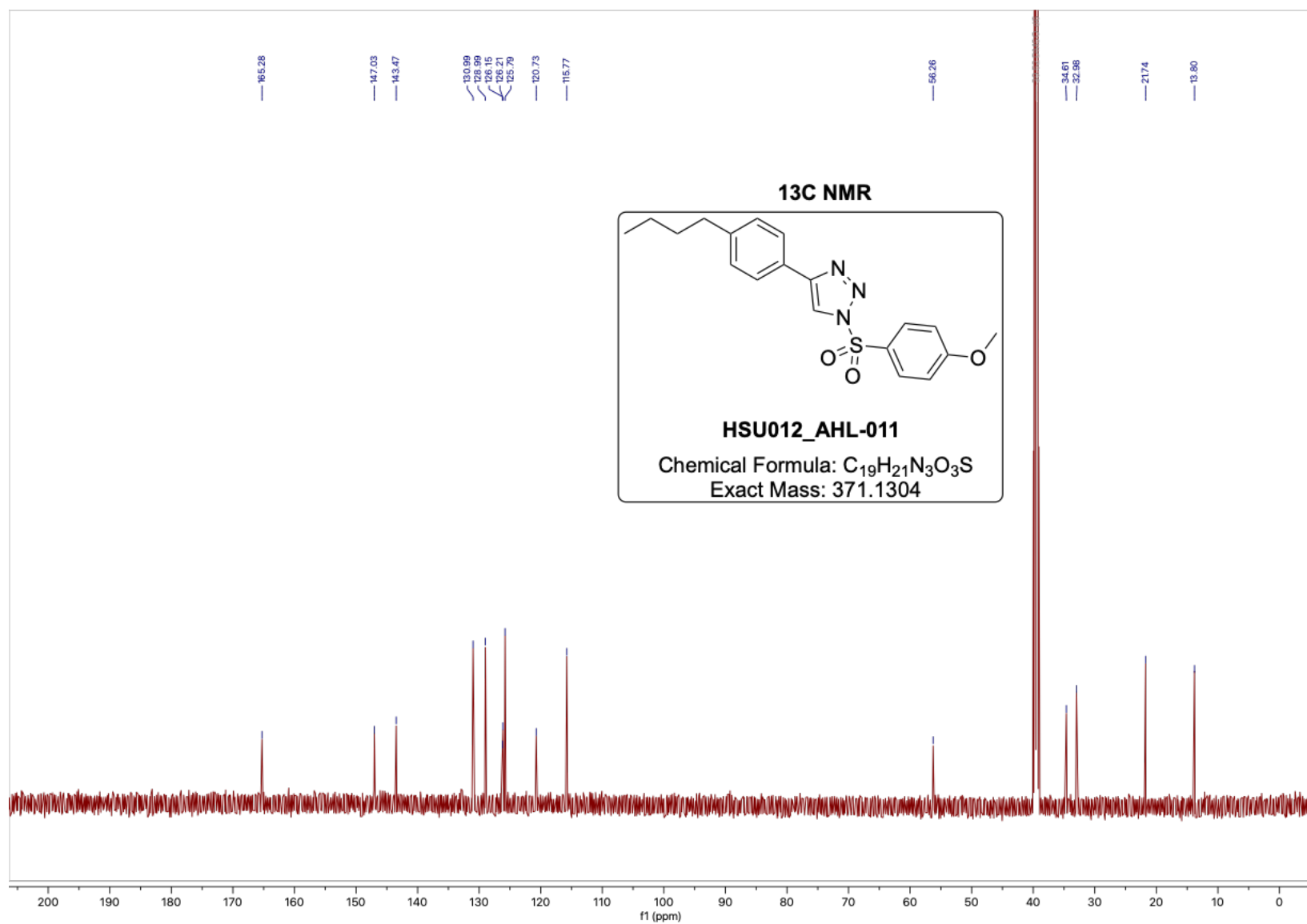


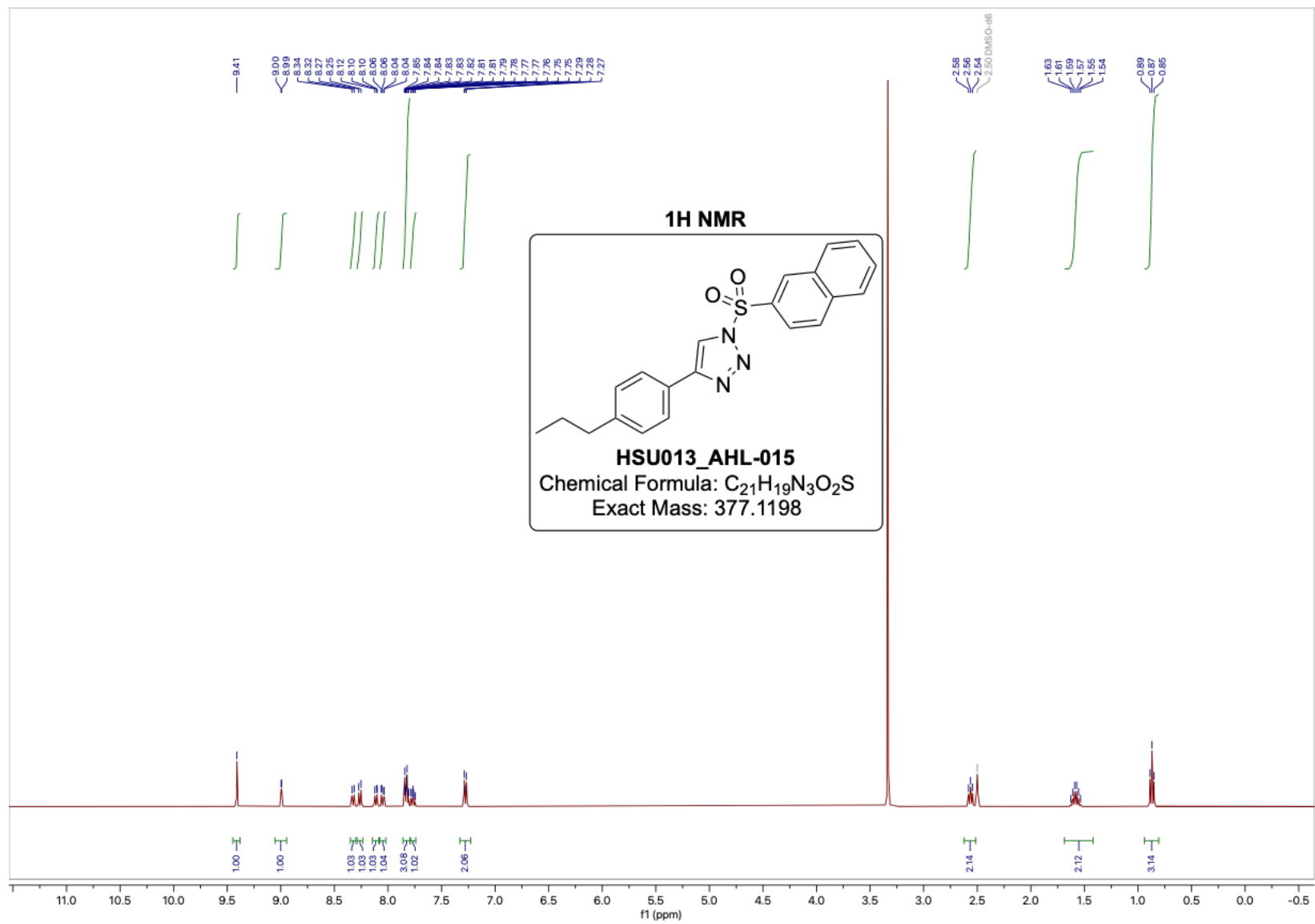


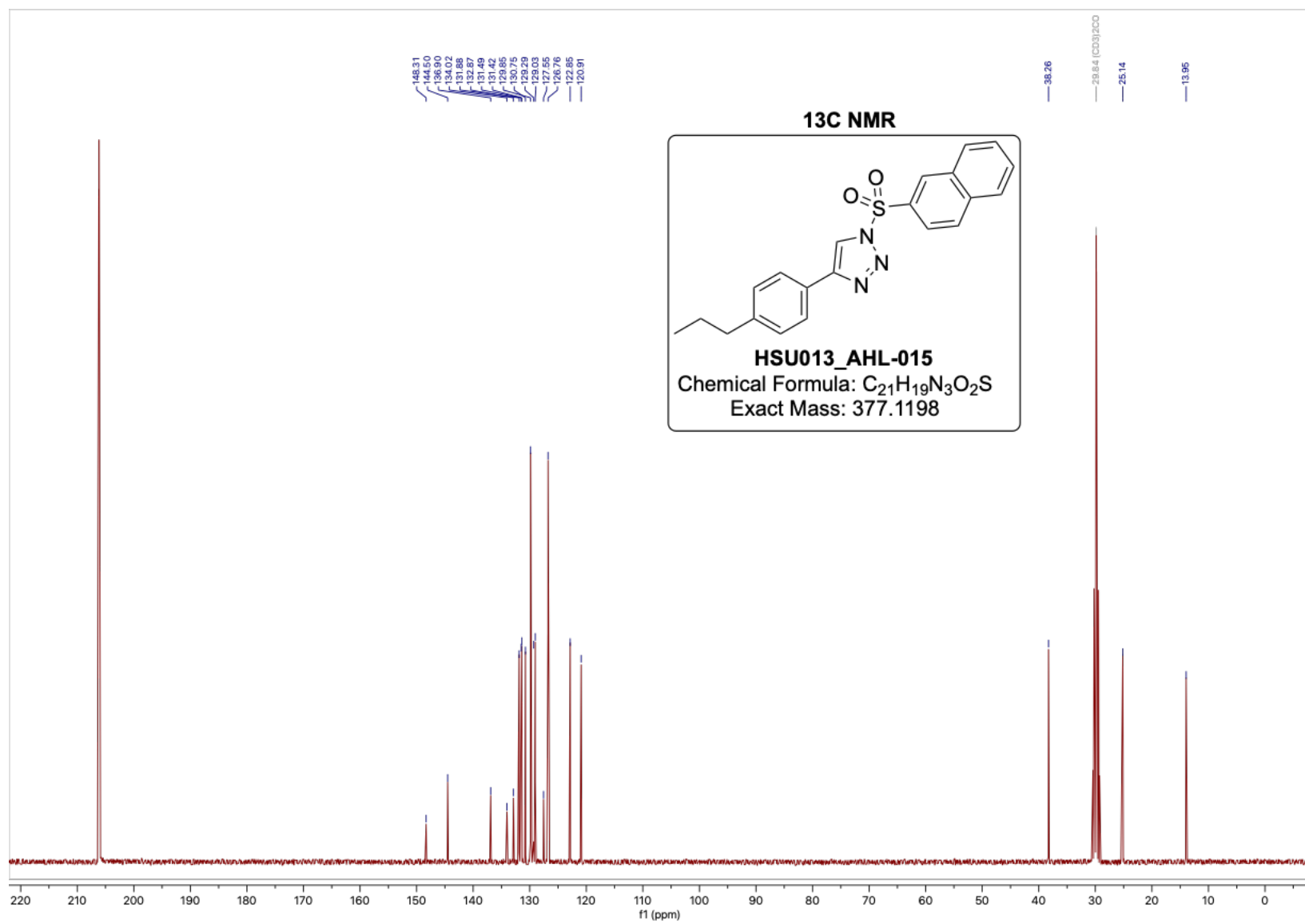


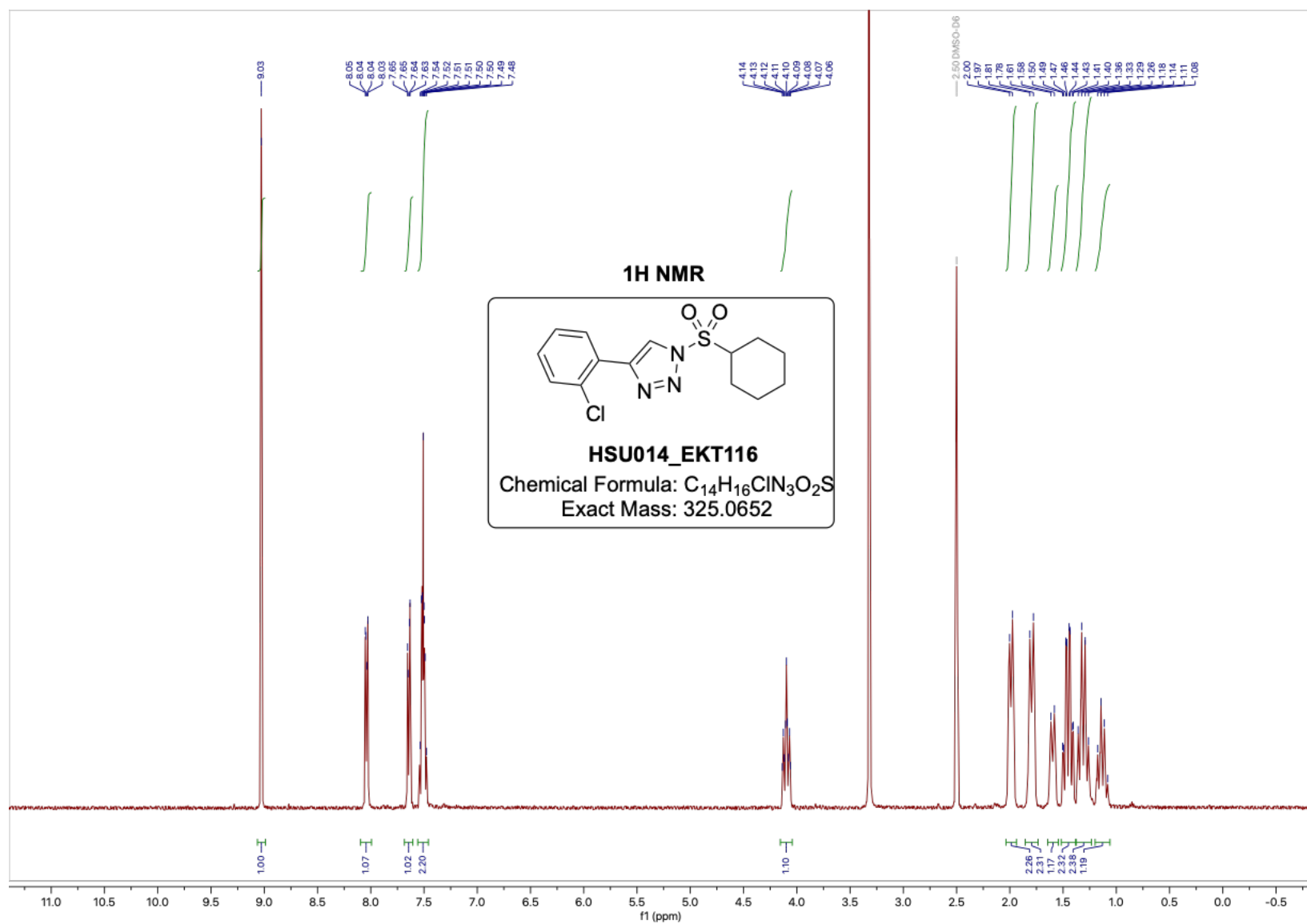


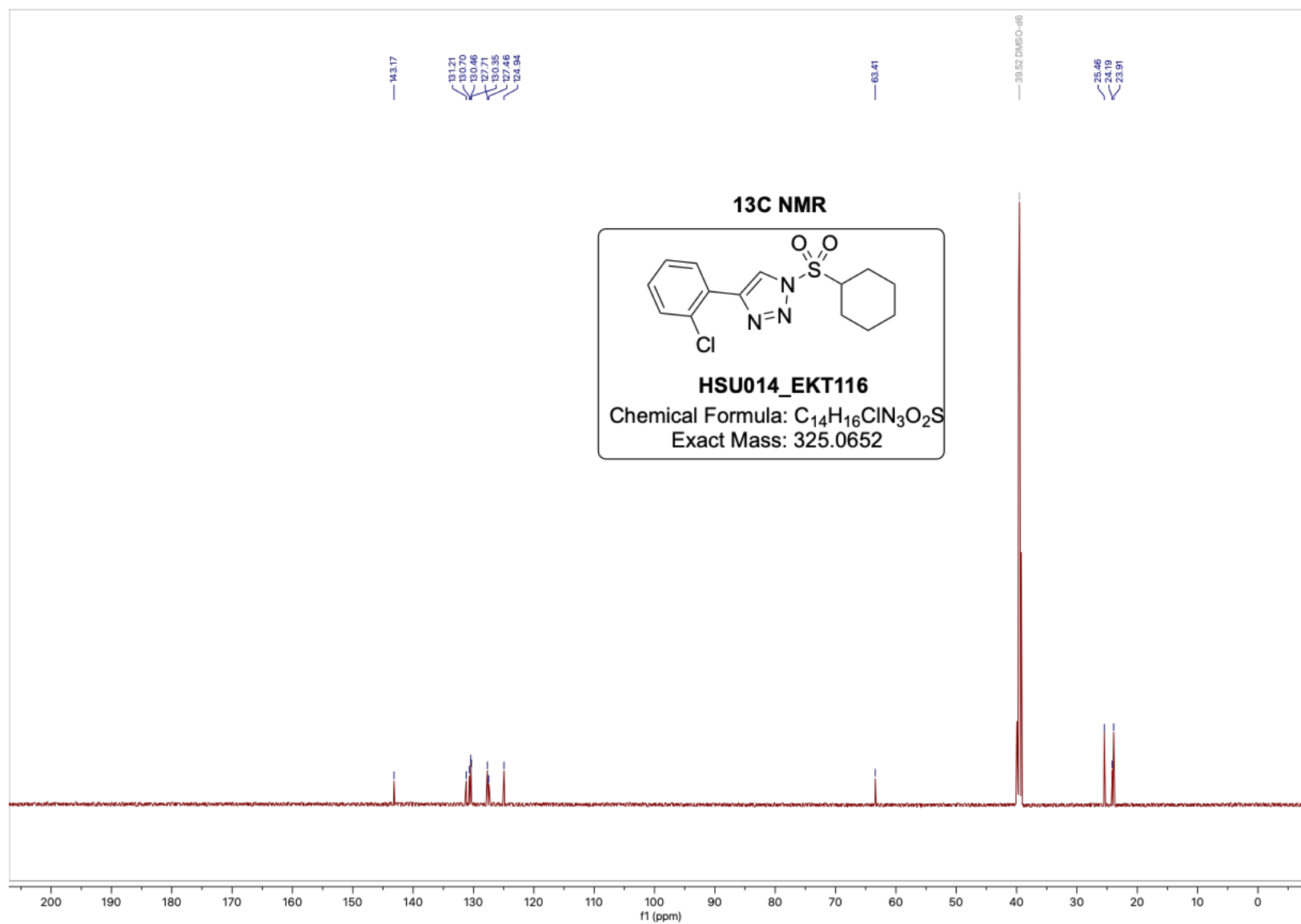


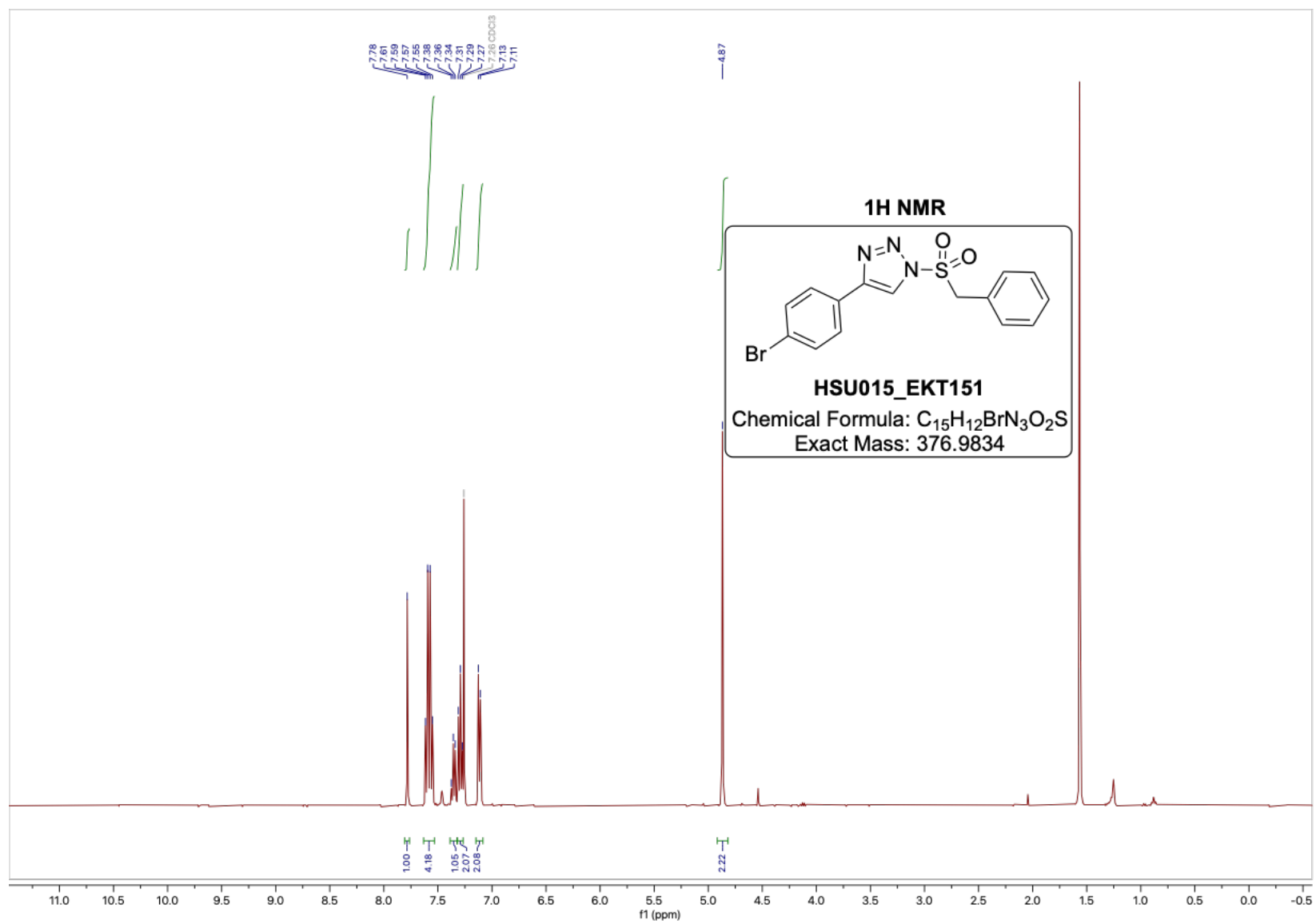


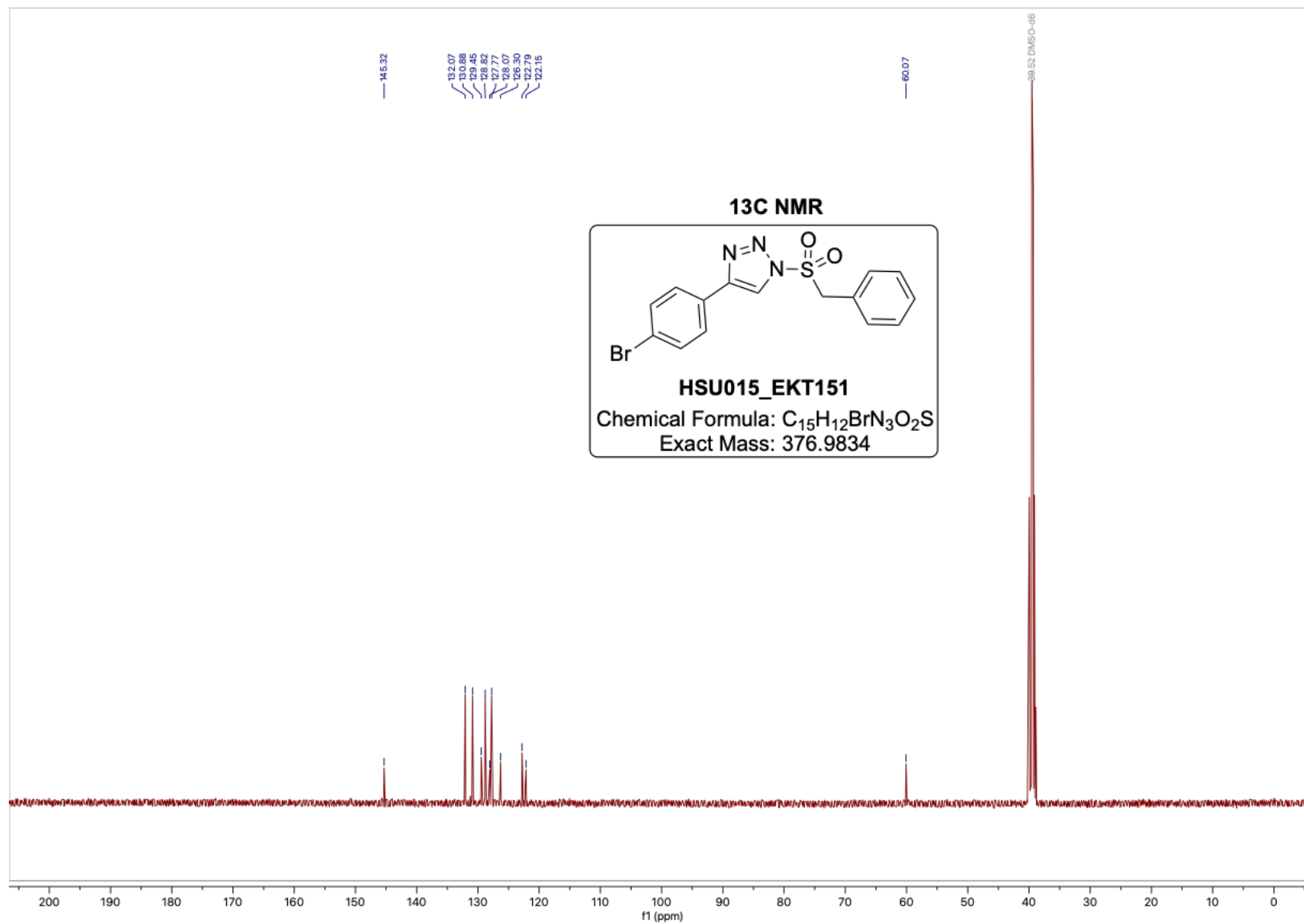


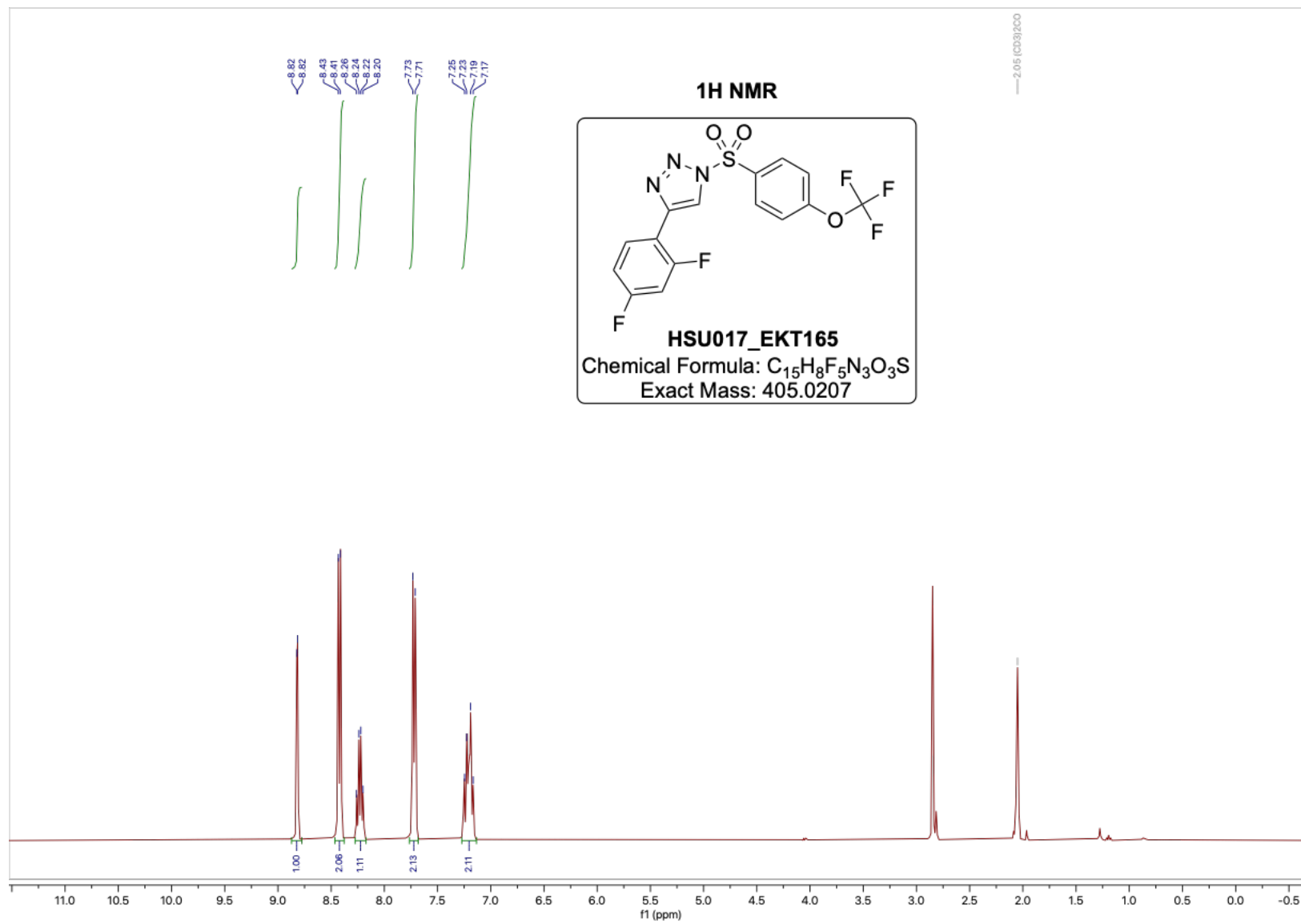




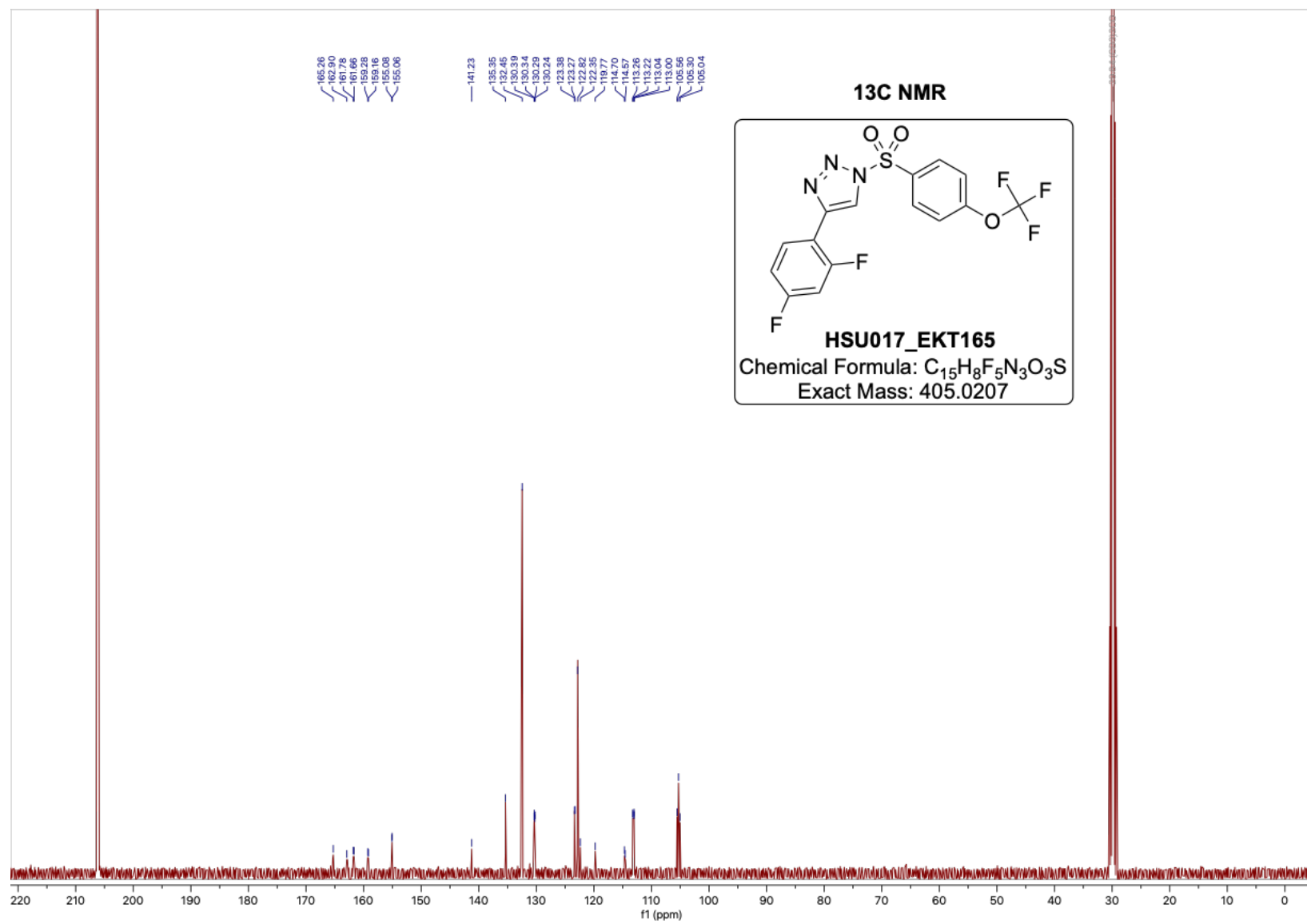


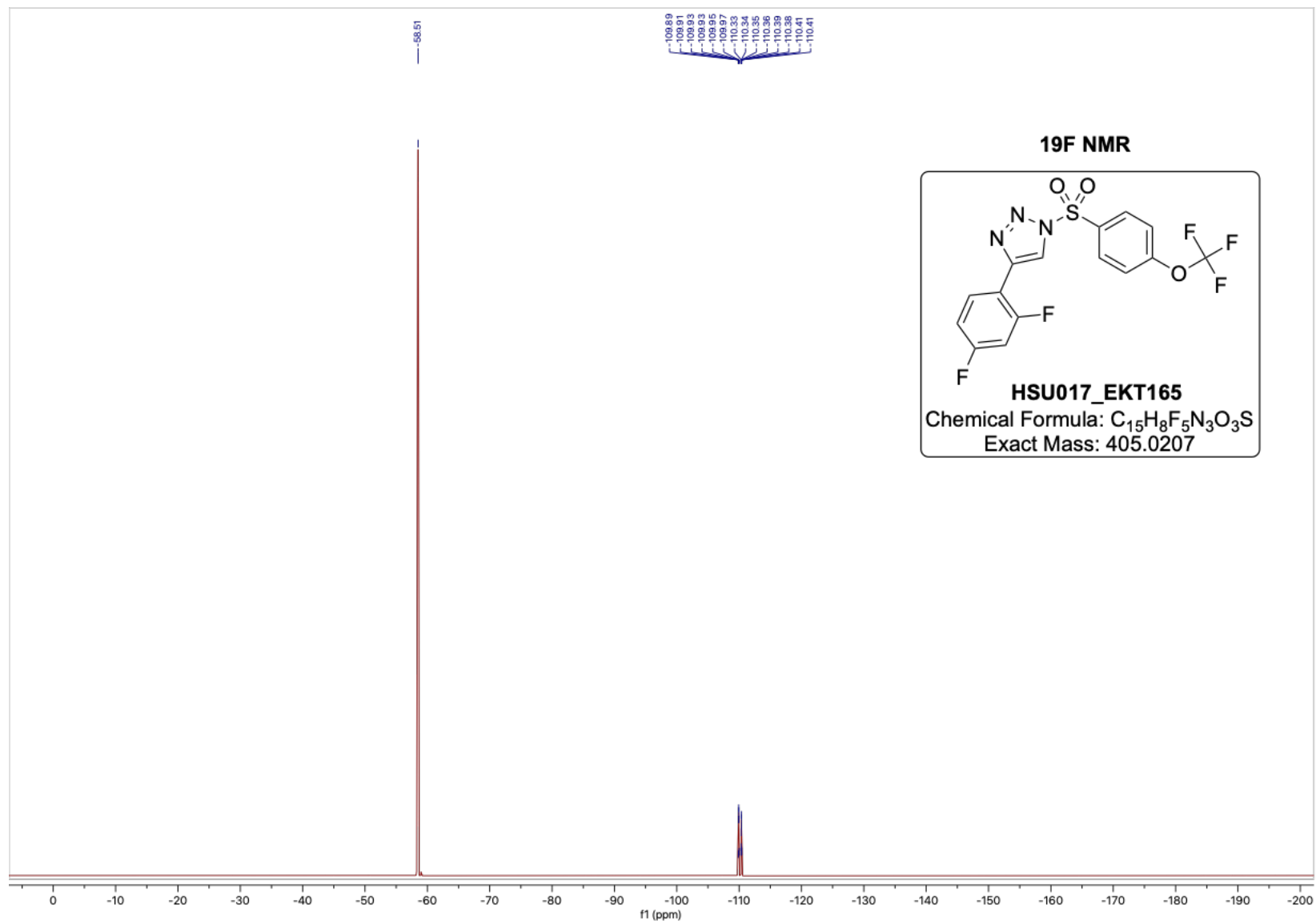




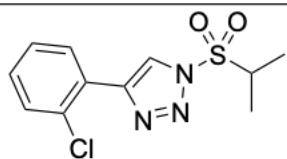






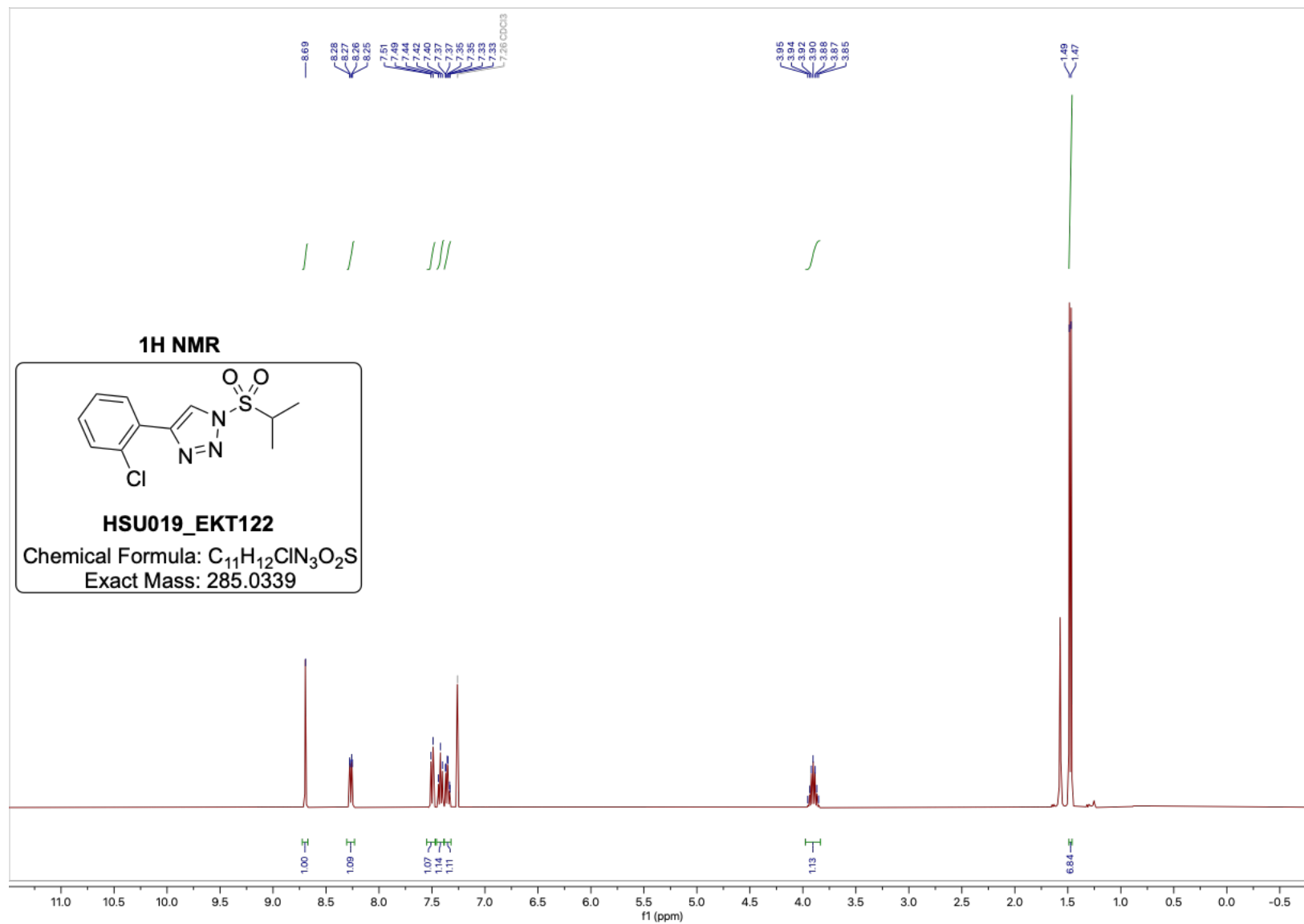


**<sup>1</sup>H NMR**

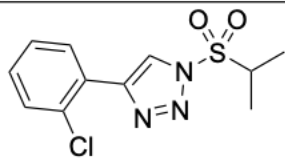


**HSU019\_EKT122**

Chemical Formula: C<sub>11</sub>H<sub>12</sub>ClN<sub>3</sub>O<sub>2</sub>S  
Exact Mass: 285.0339

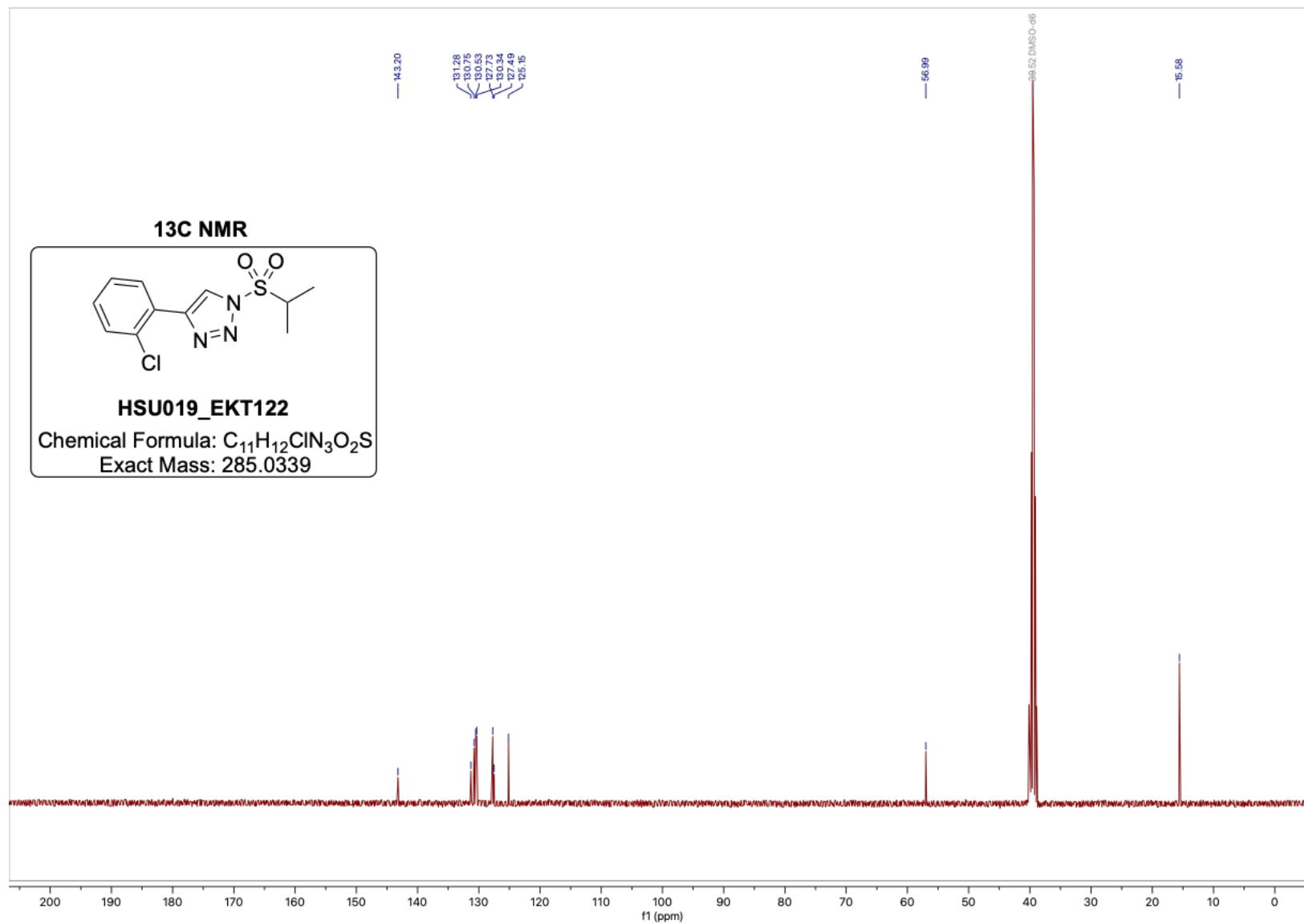


### **<sup>13</sup>C NMR**

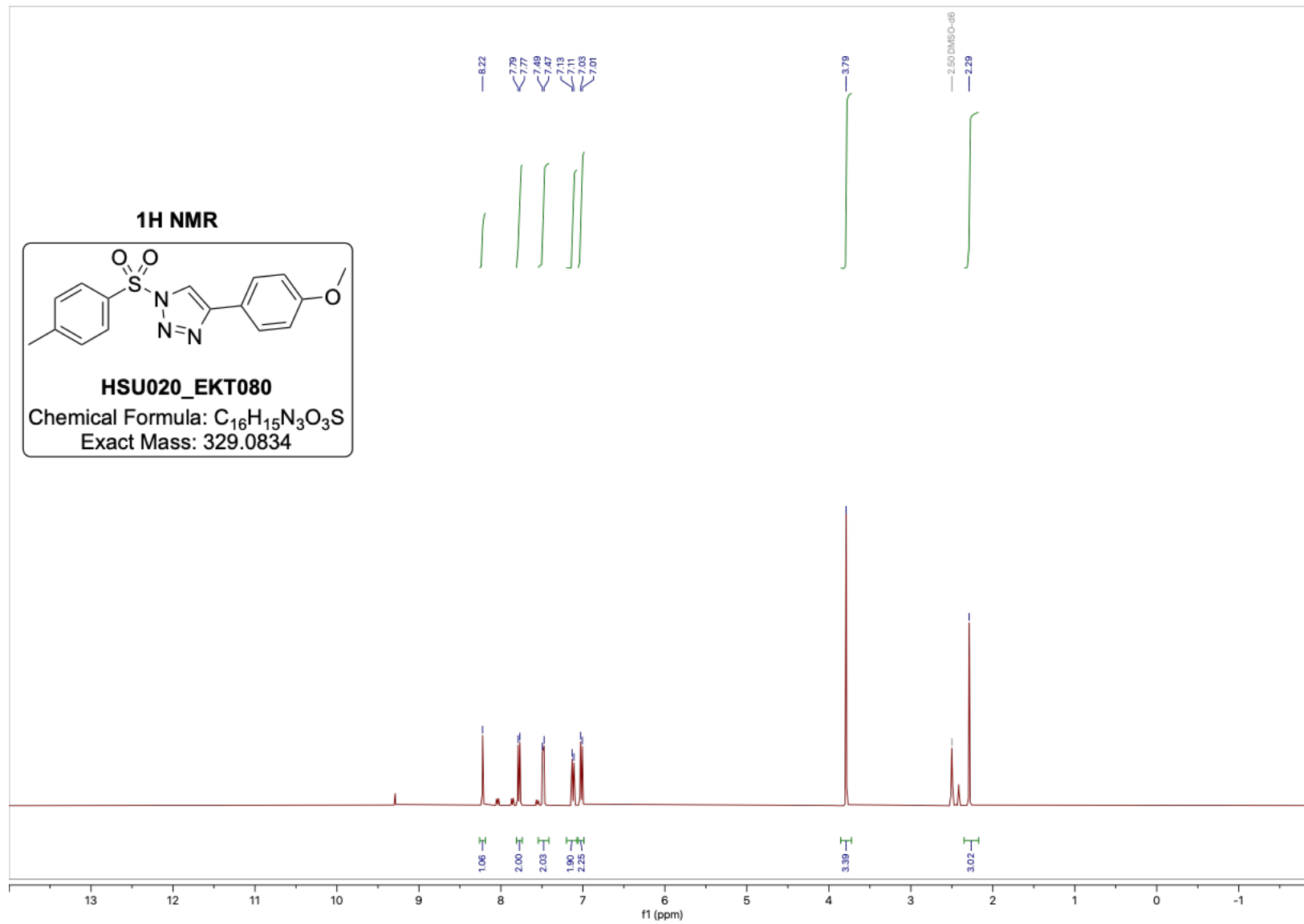
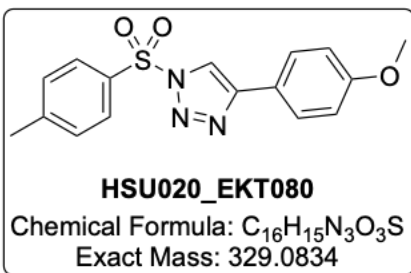


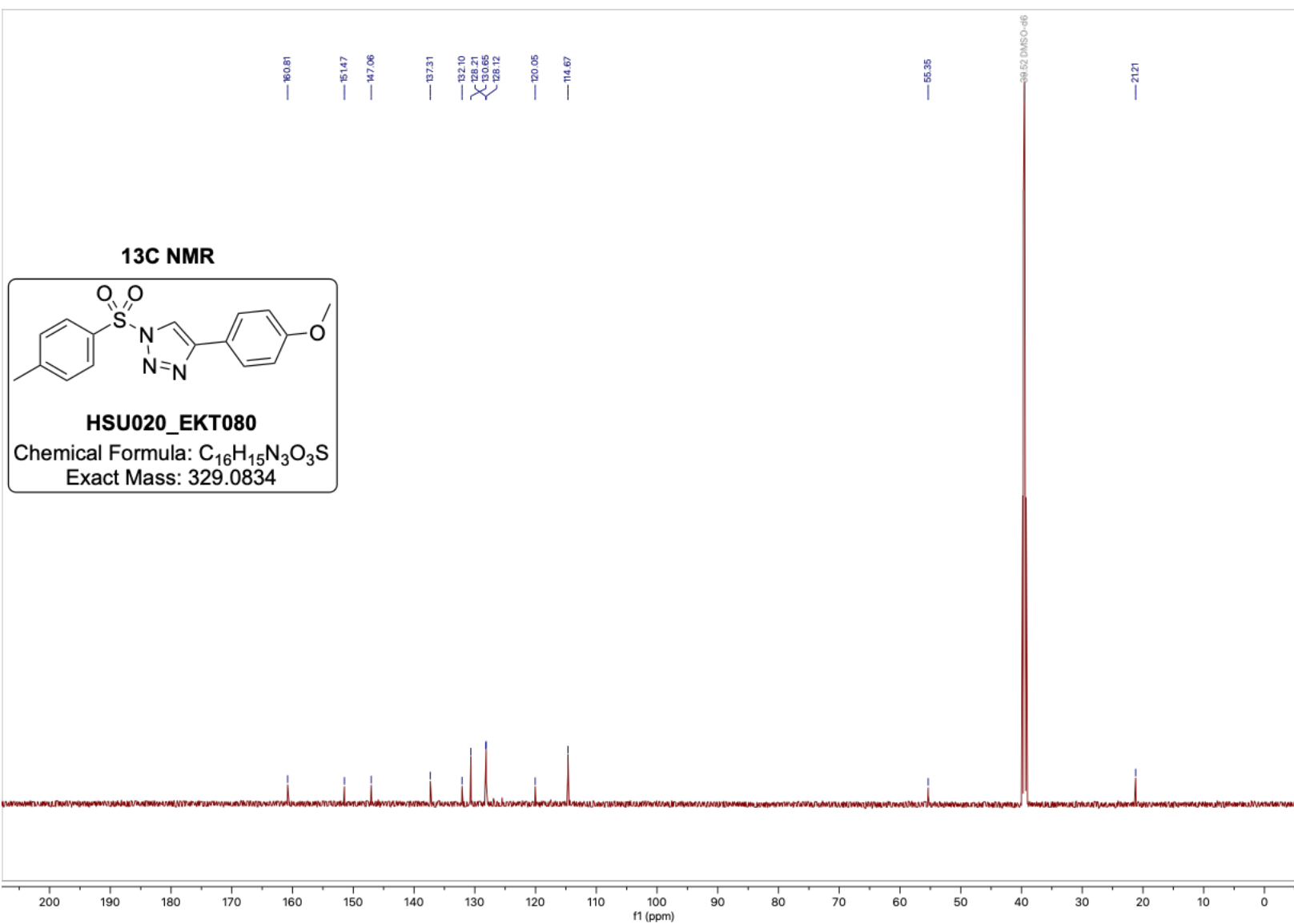
**HSU019\_EKT122**

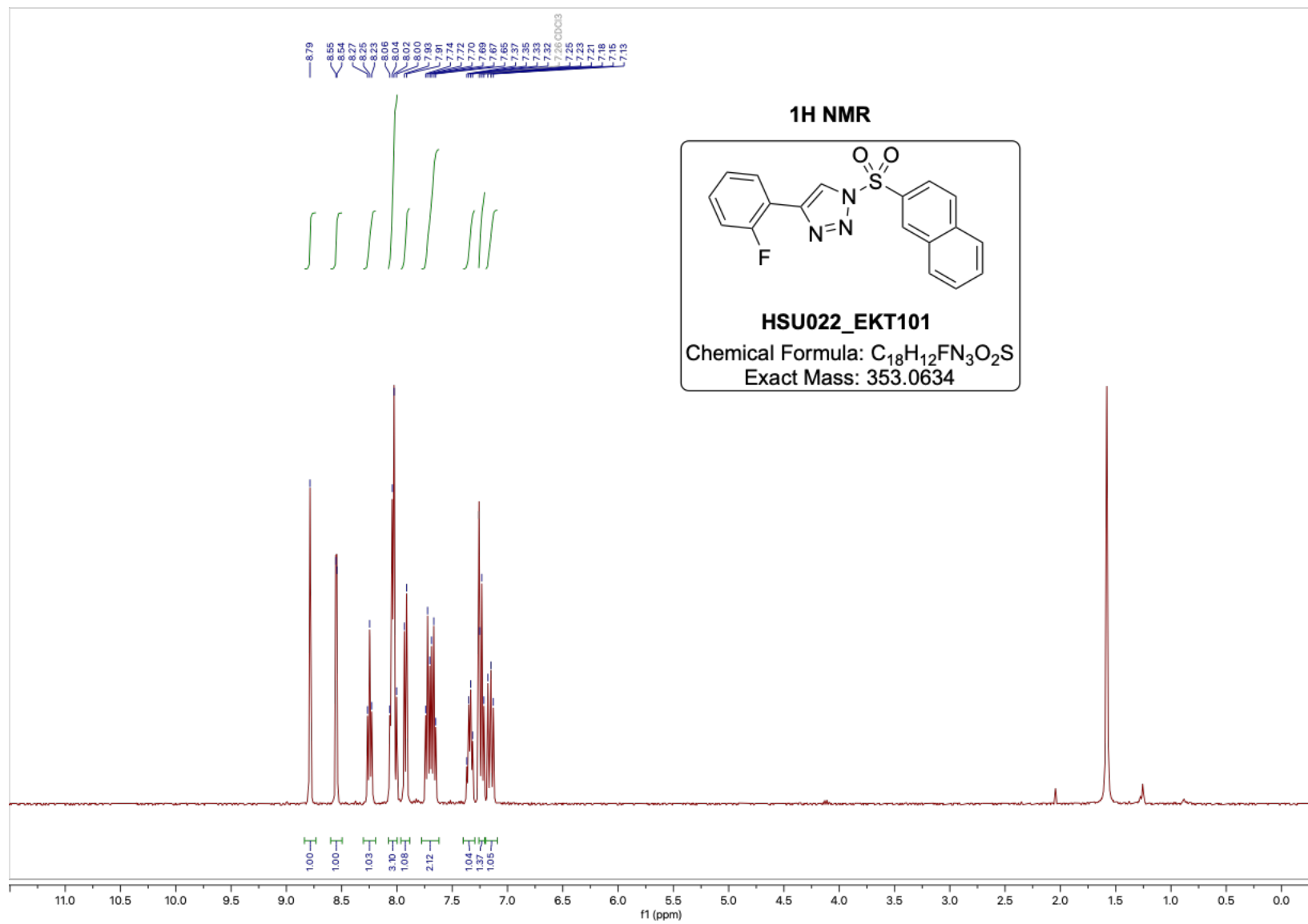
Chemical Formula: C<sub>11</sub>H<sub>12</sub>ClN<sub>3</sub>O<sub>2</sub>S  
Exact Mass: 285.0339

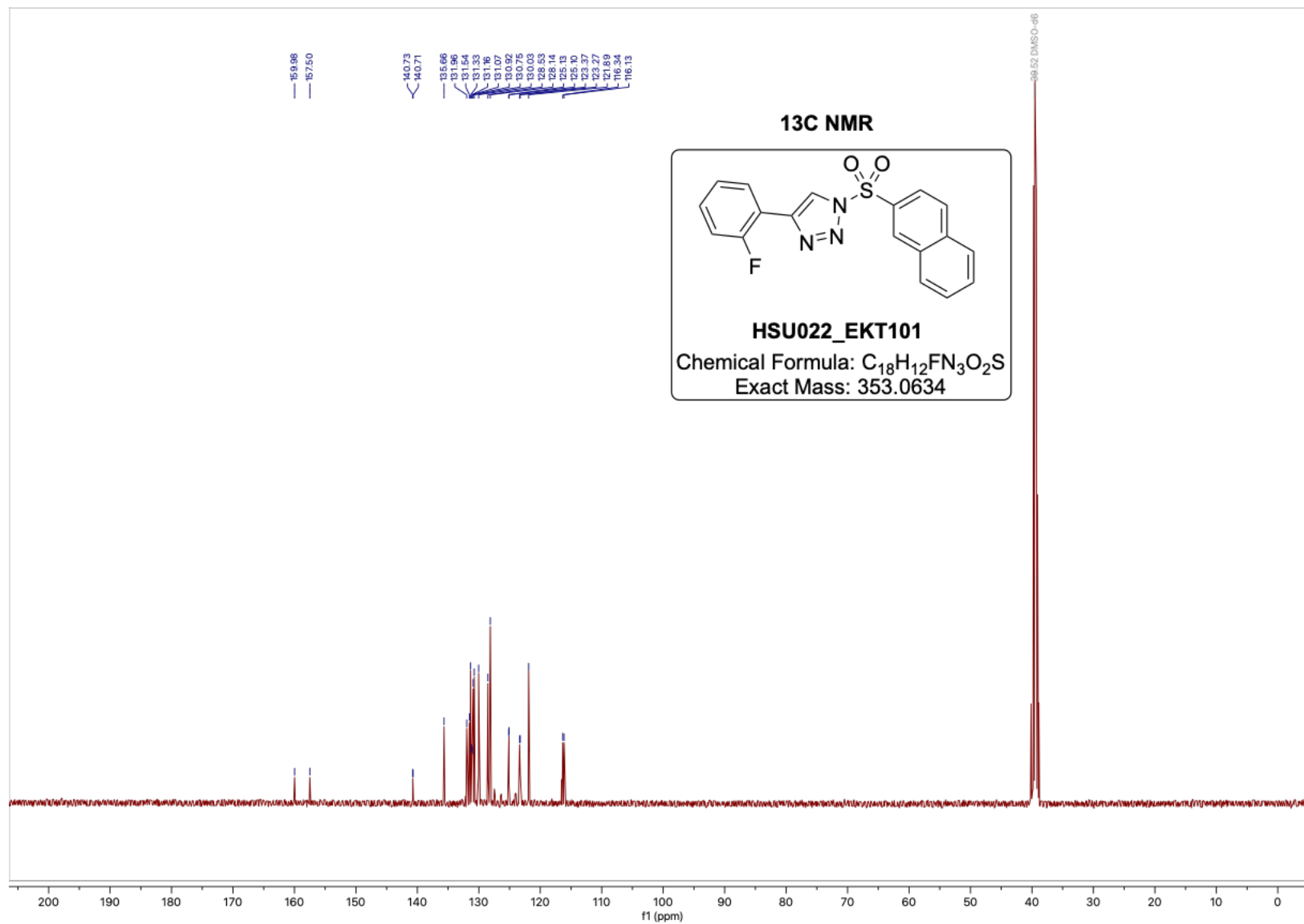


**<sup>1</sup>H NMR**

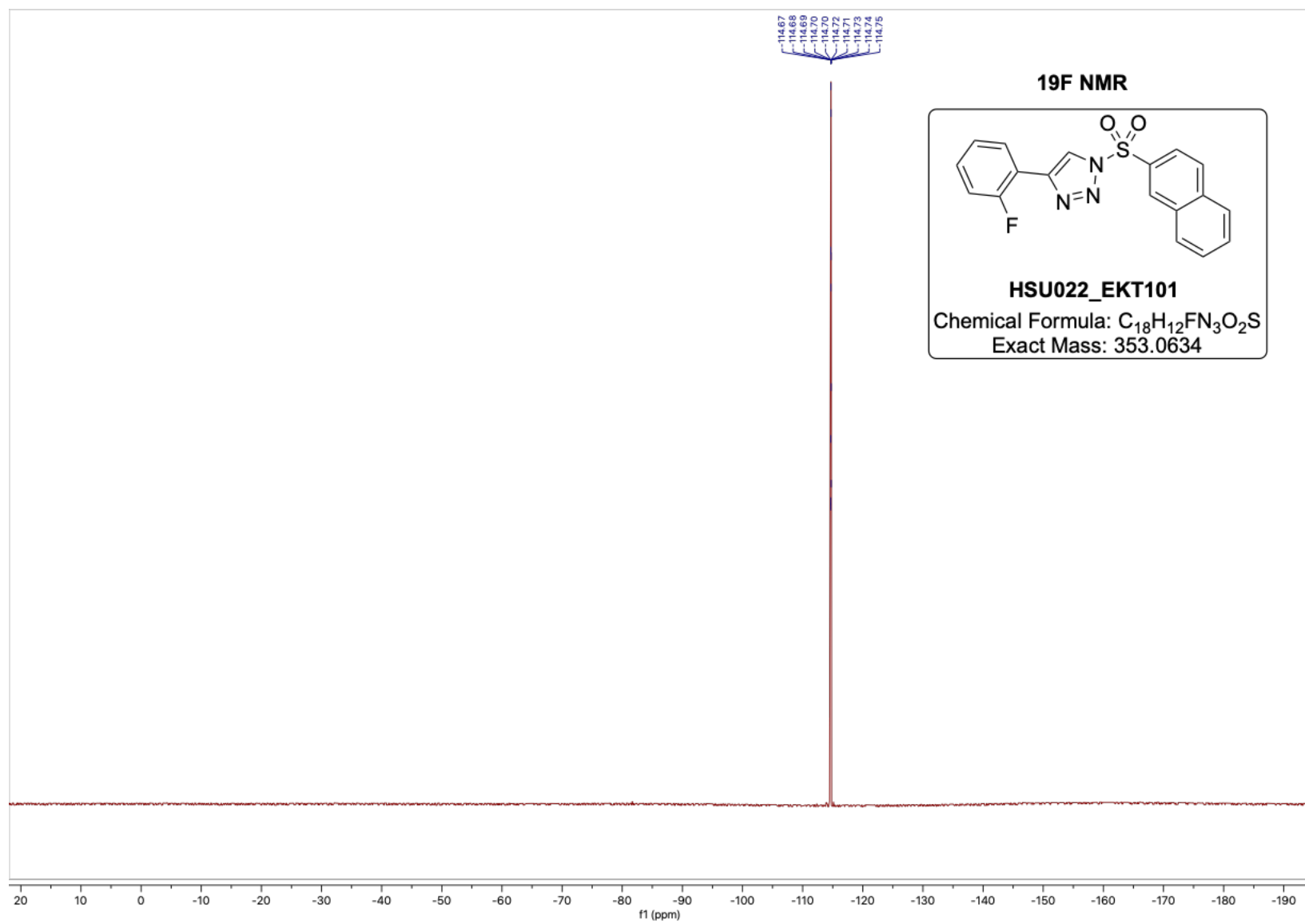


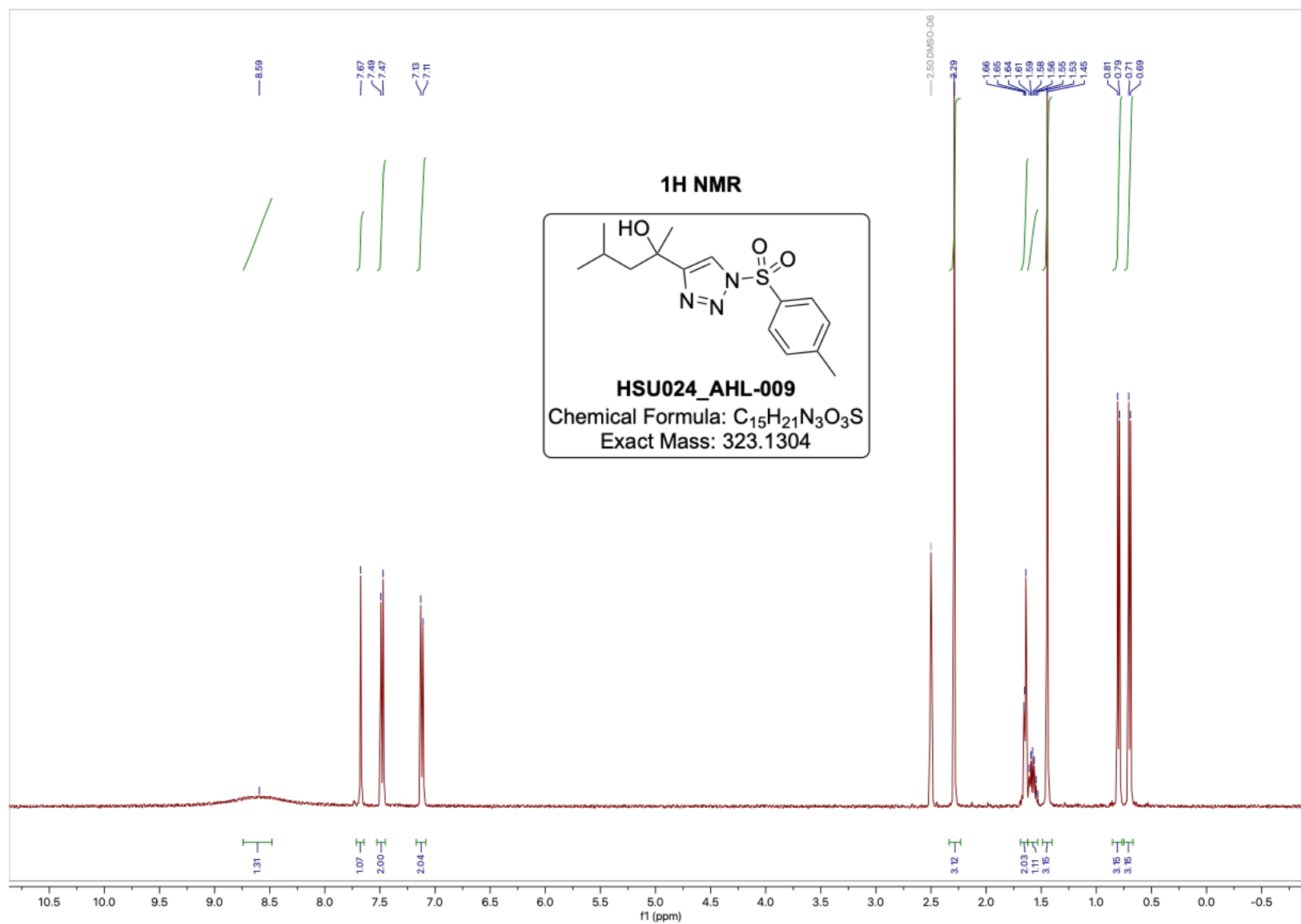


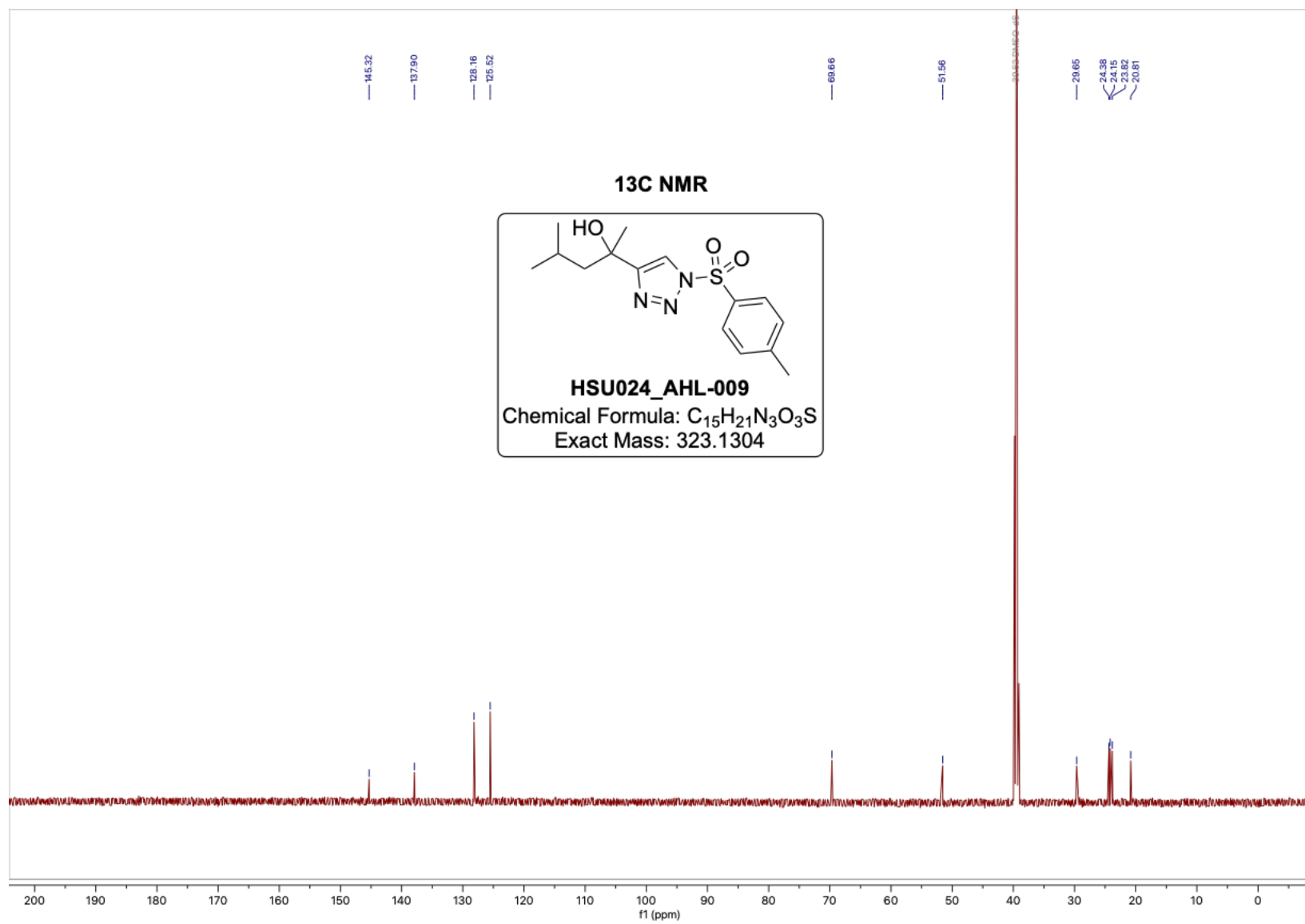


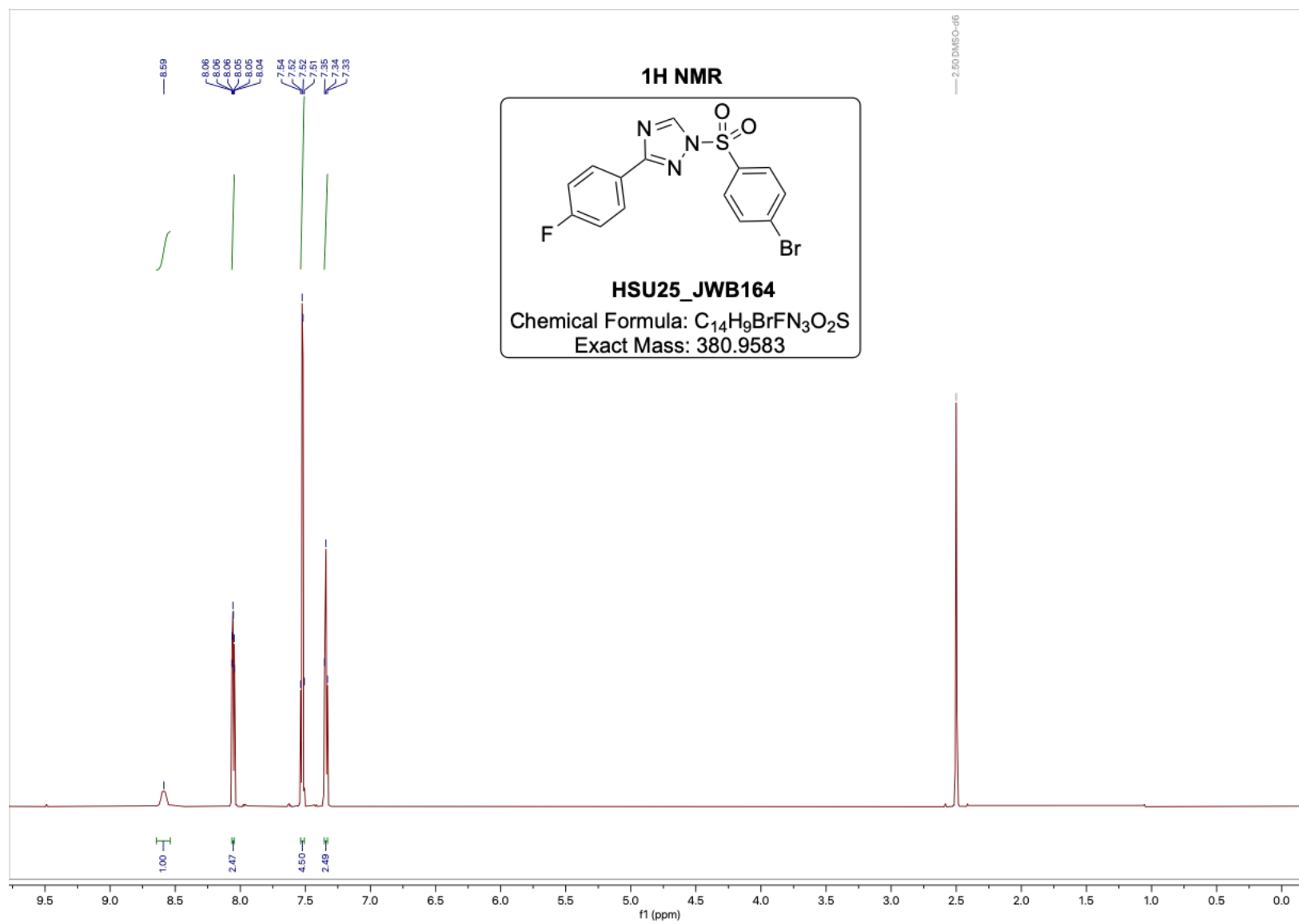


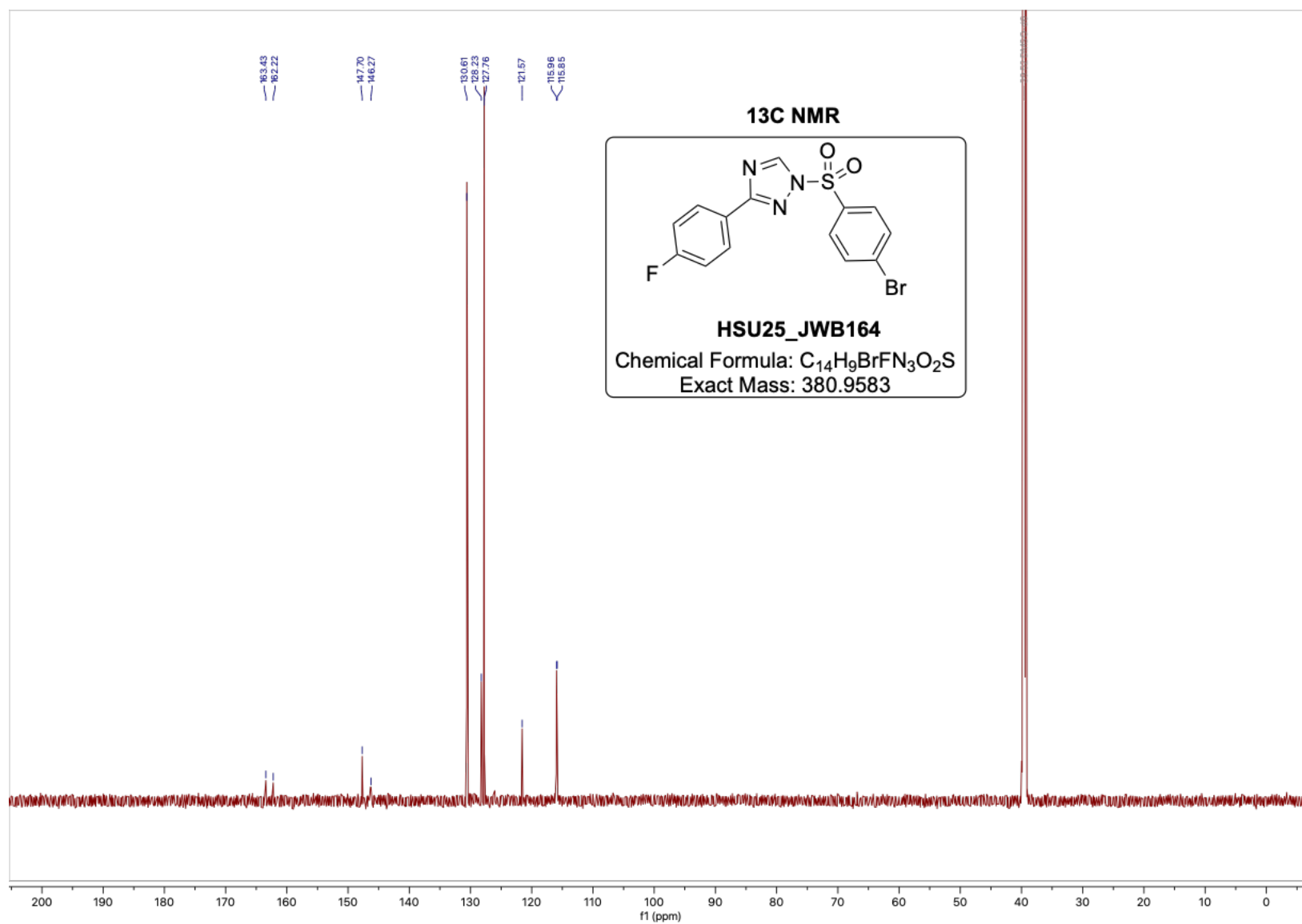




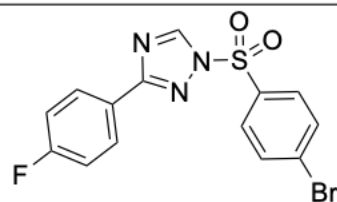








**<sup>19</sup>F NMR**

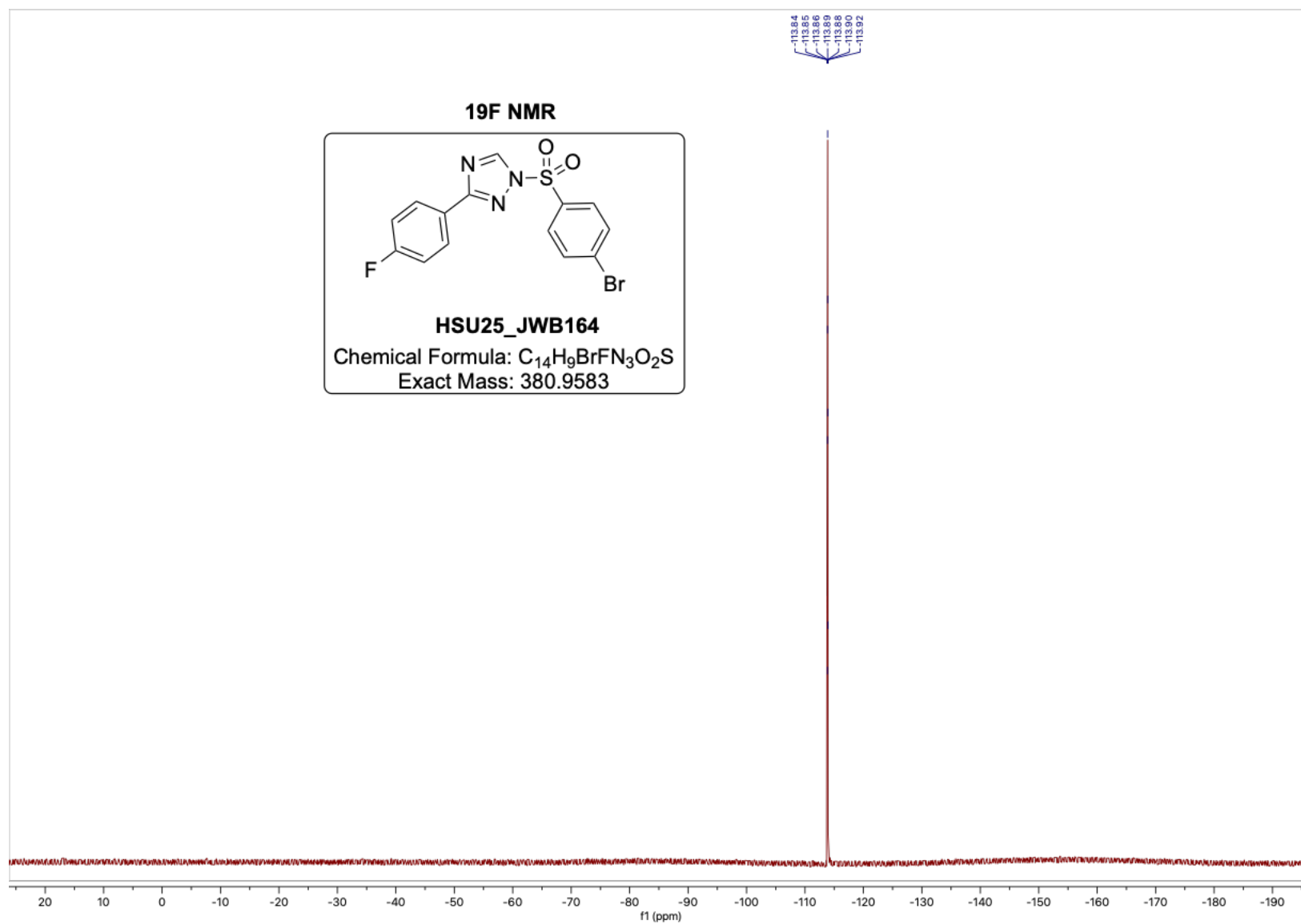


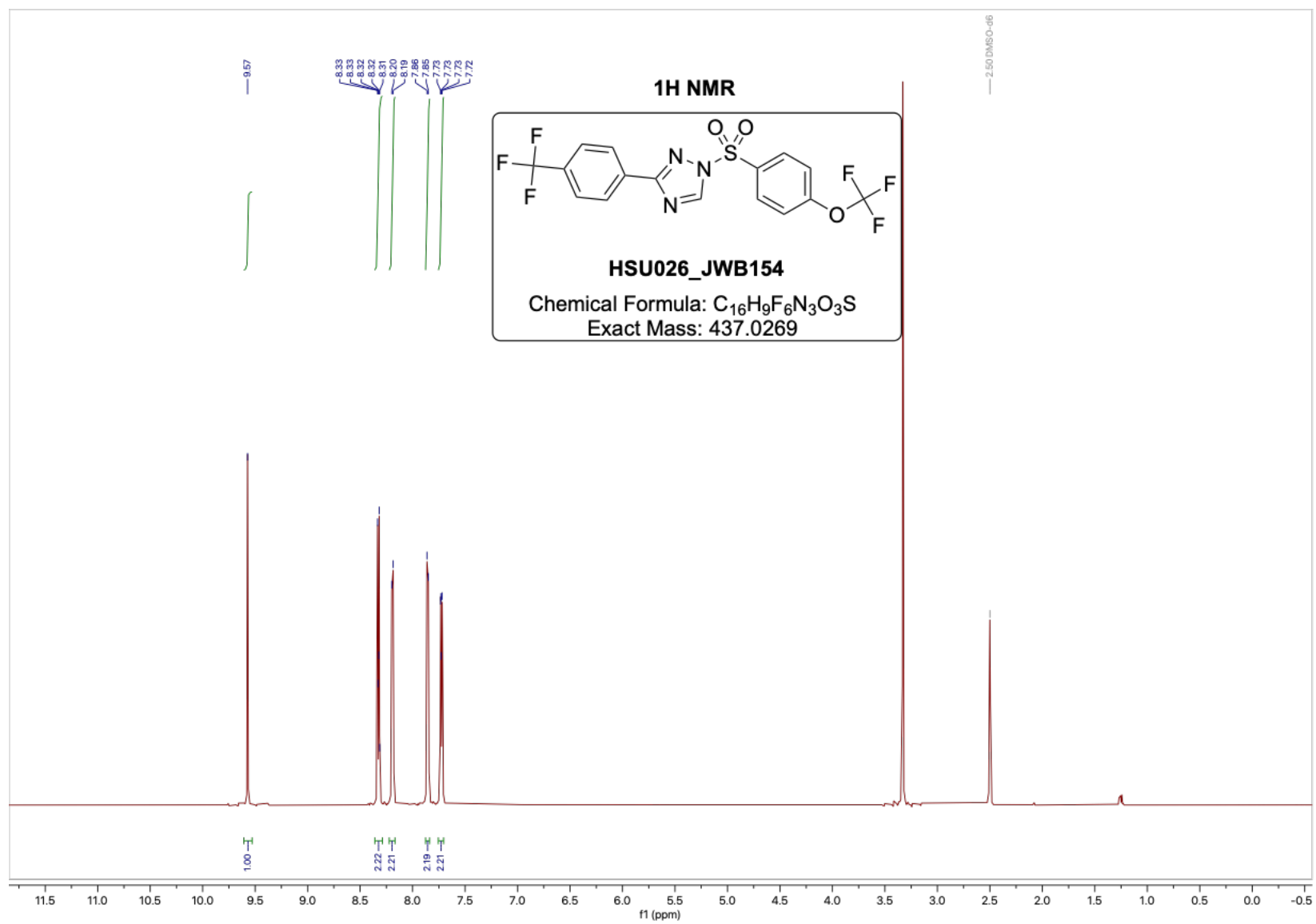
**HSU25\_JWB164**

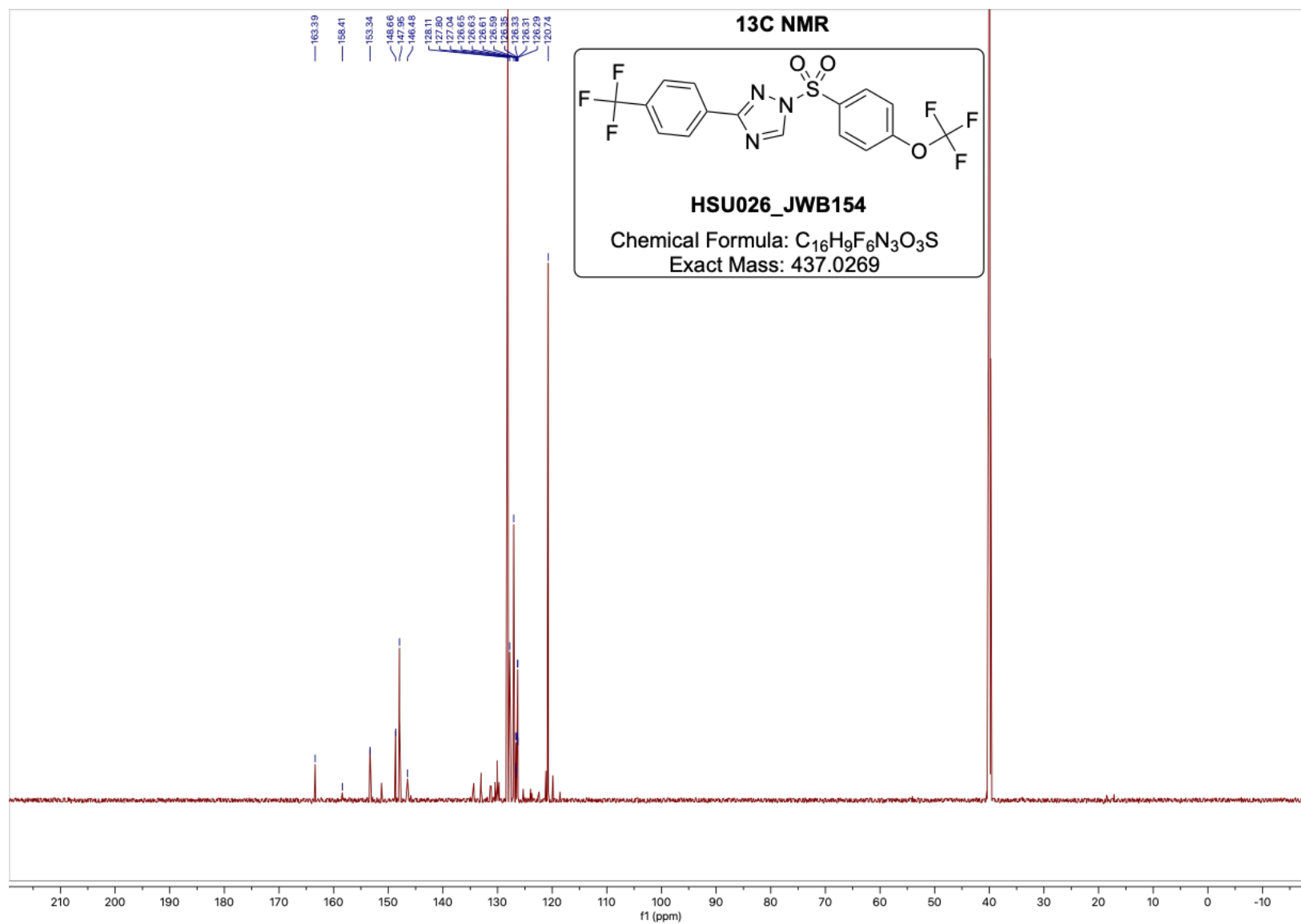
Chemical Formula: C<sub>14</sub>H<sub>9</sub>BrFN<sub>3</sub>O<sub>2</sub>S

Exact Mass: 380.9583

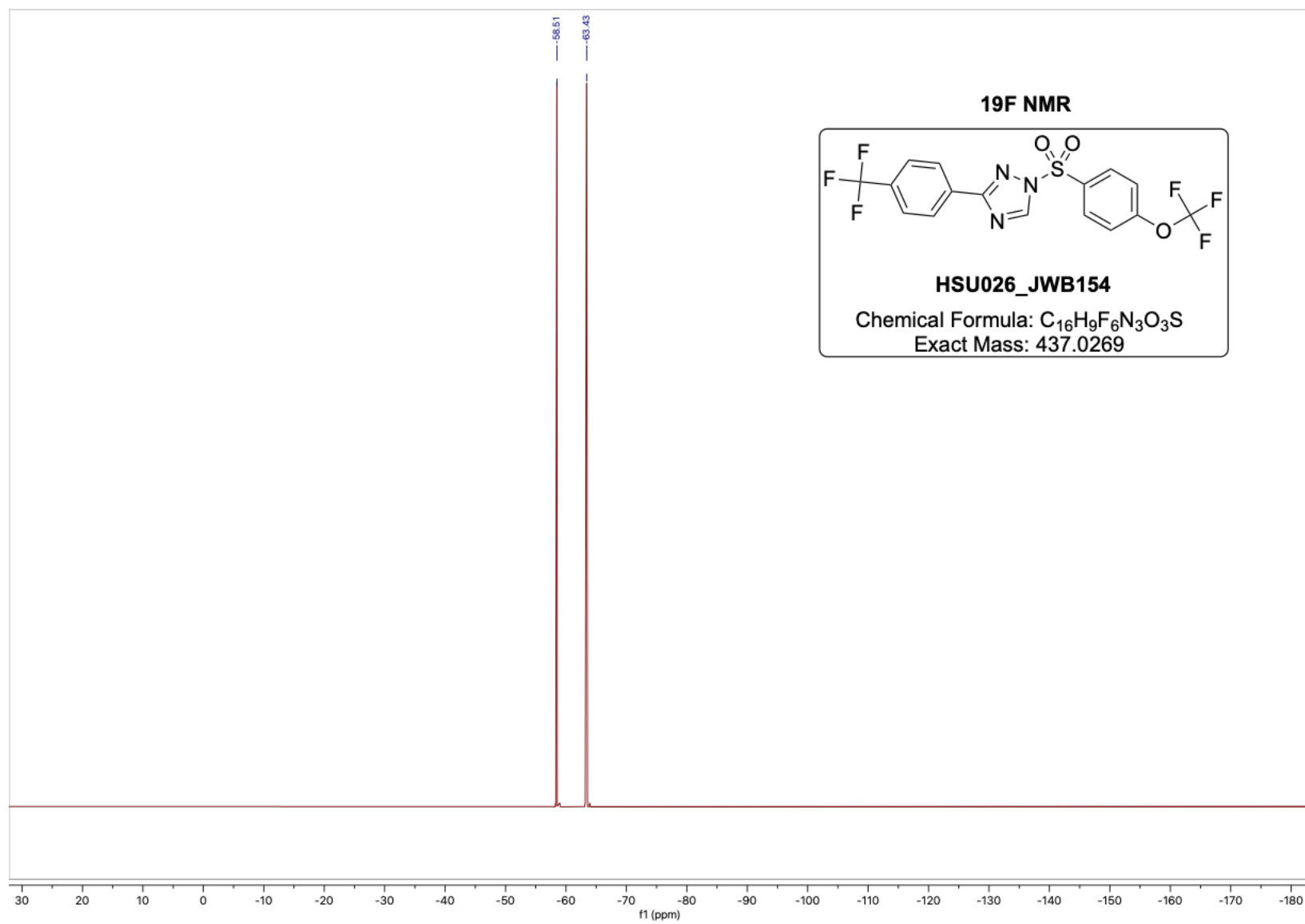
-113.84  
-113.85  
-113.86  
-113.87  
-113.88  
-113.89  
-113.90  
-113.92

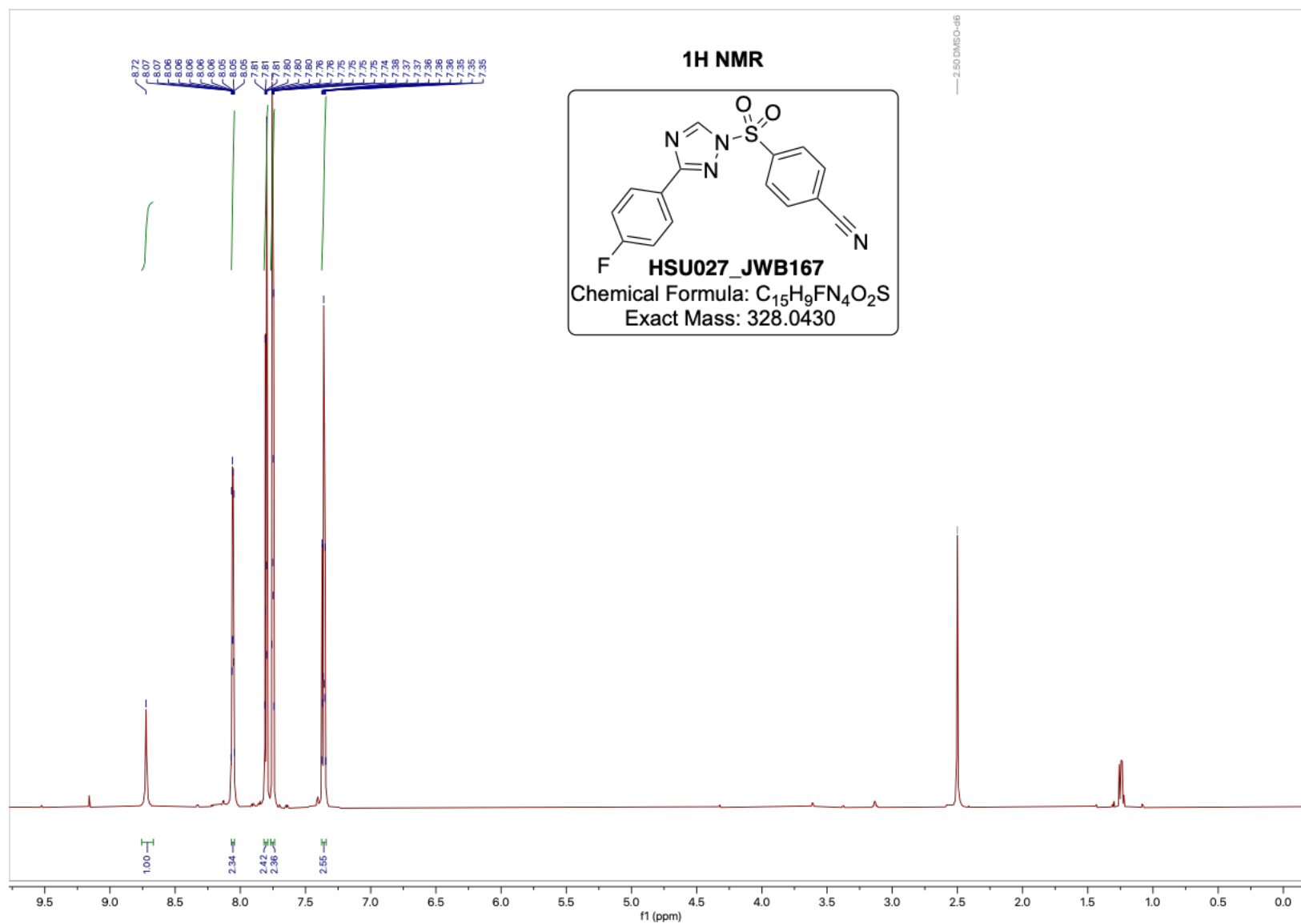


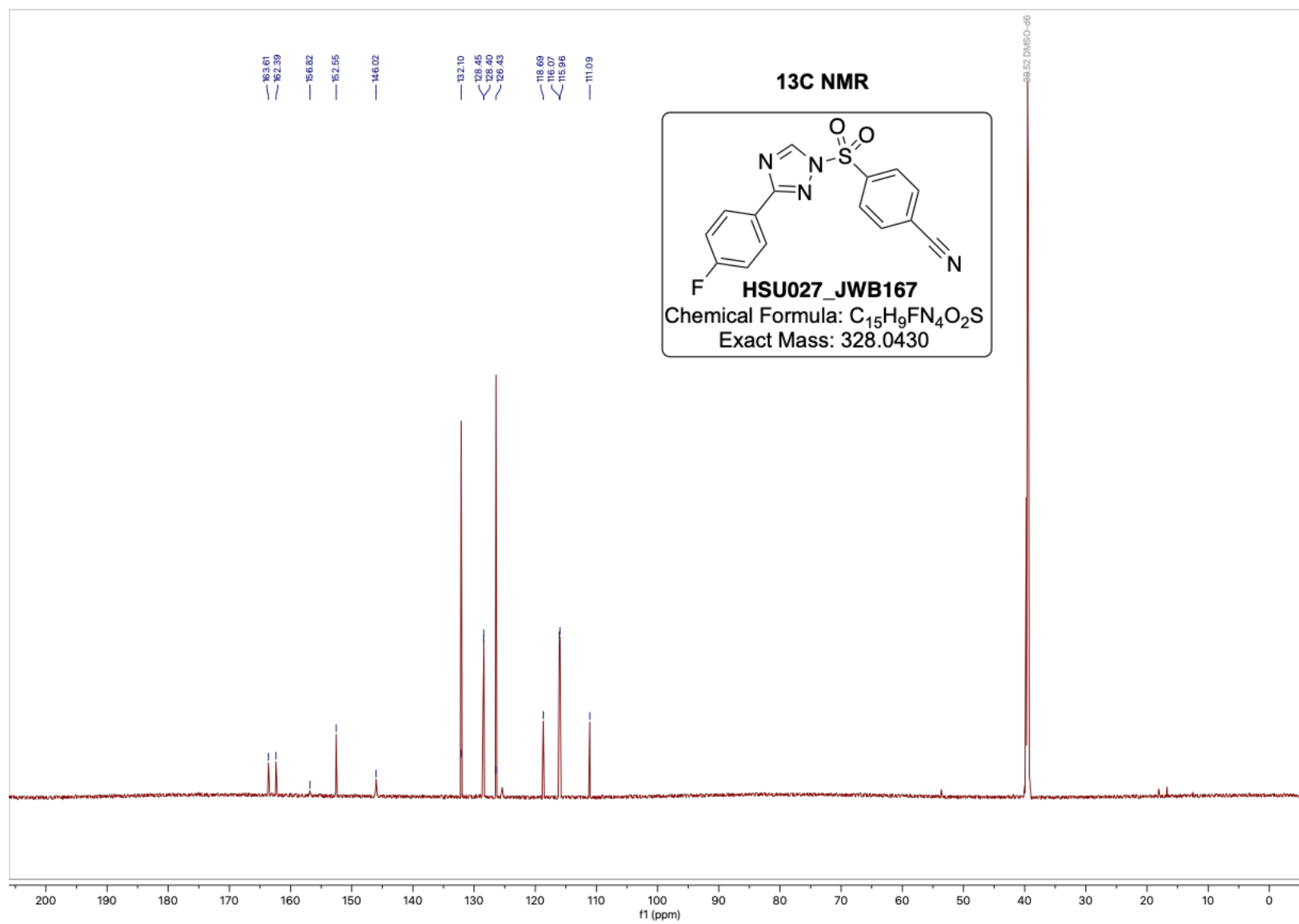






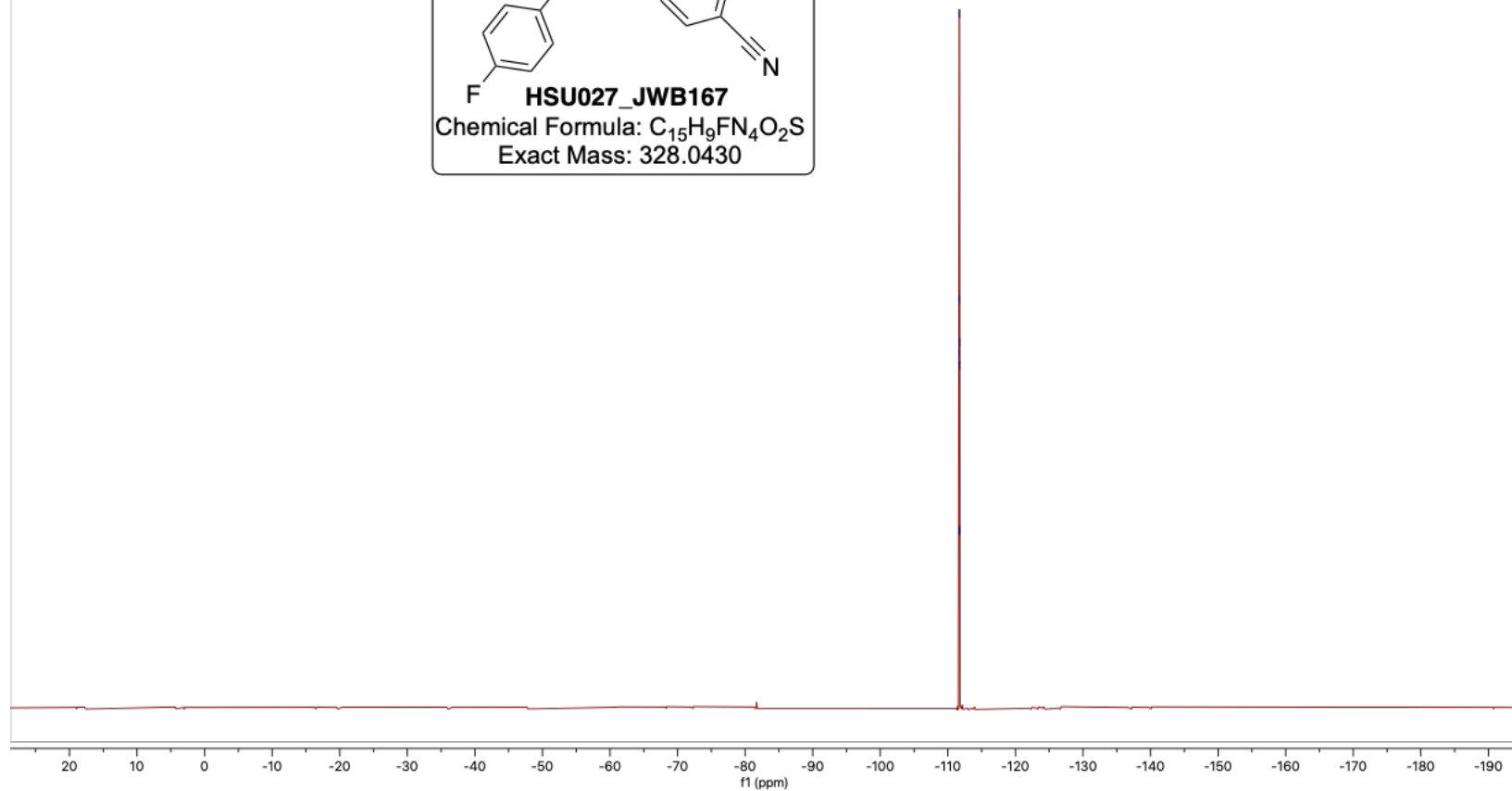
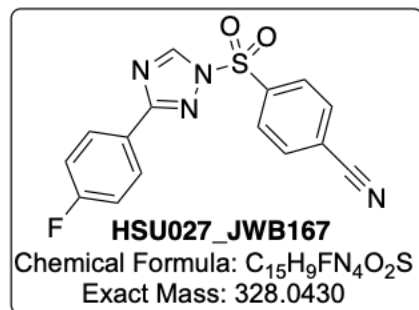


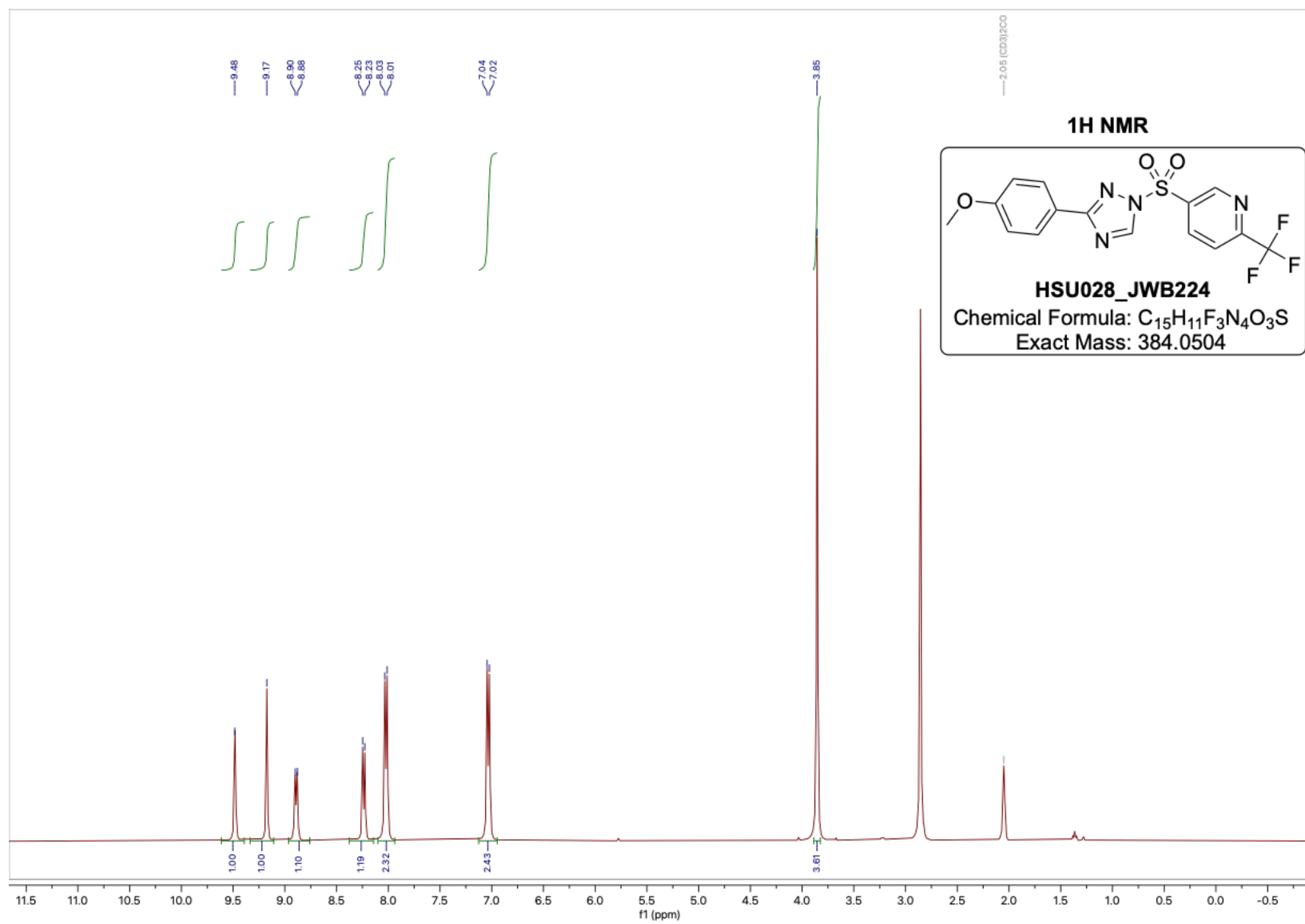


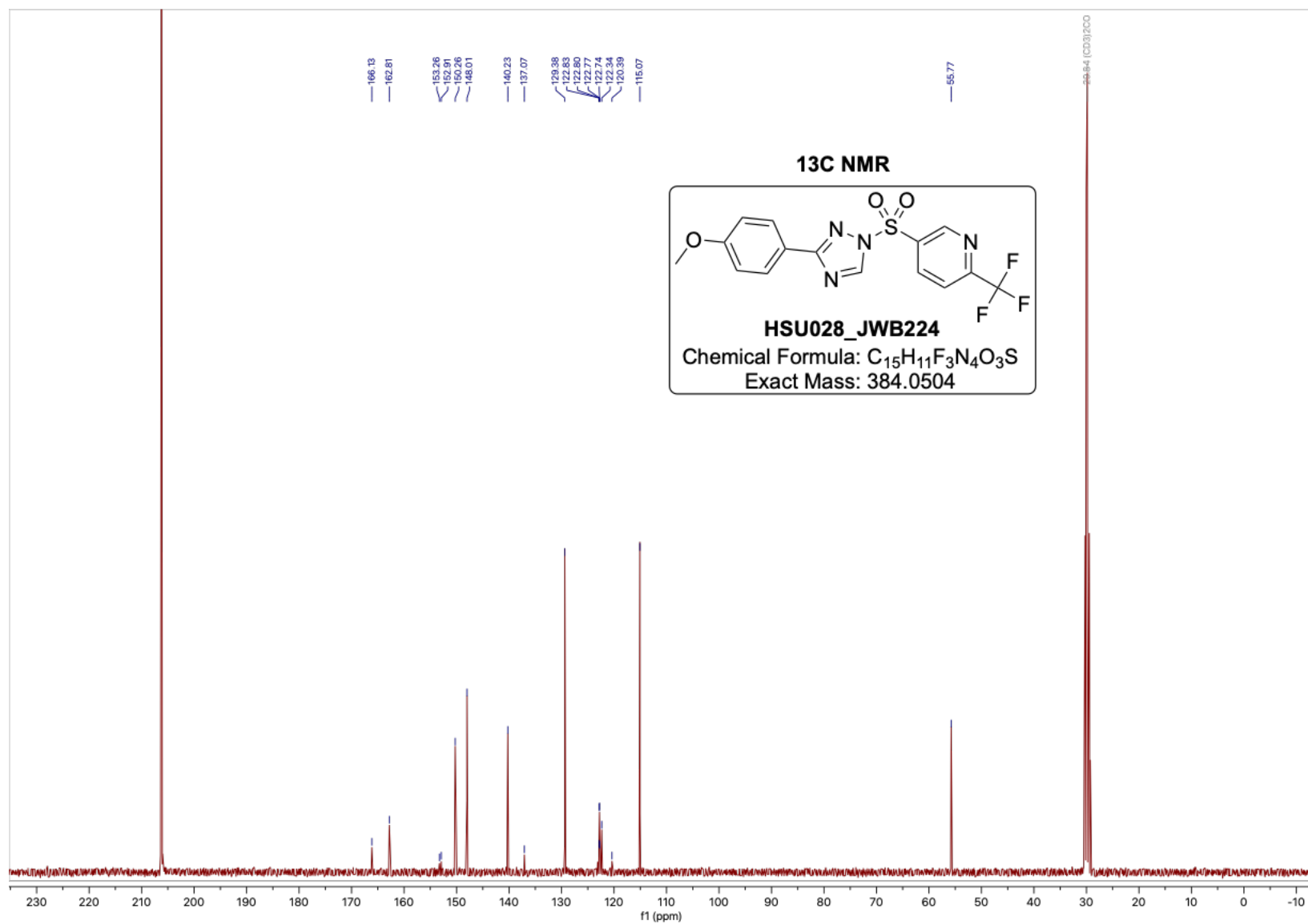


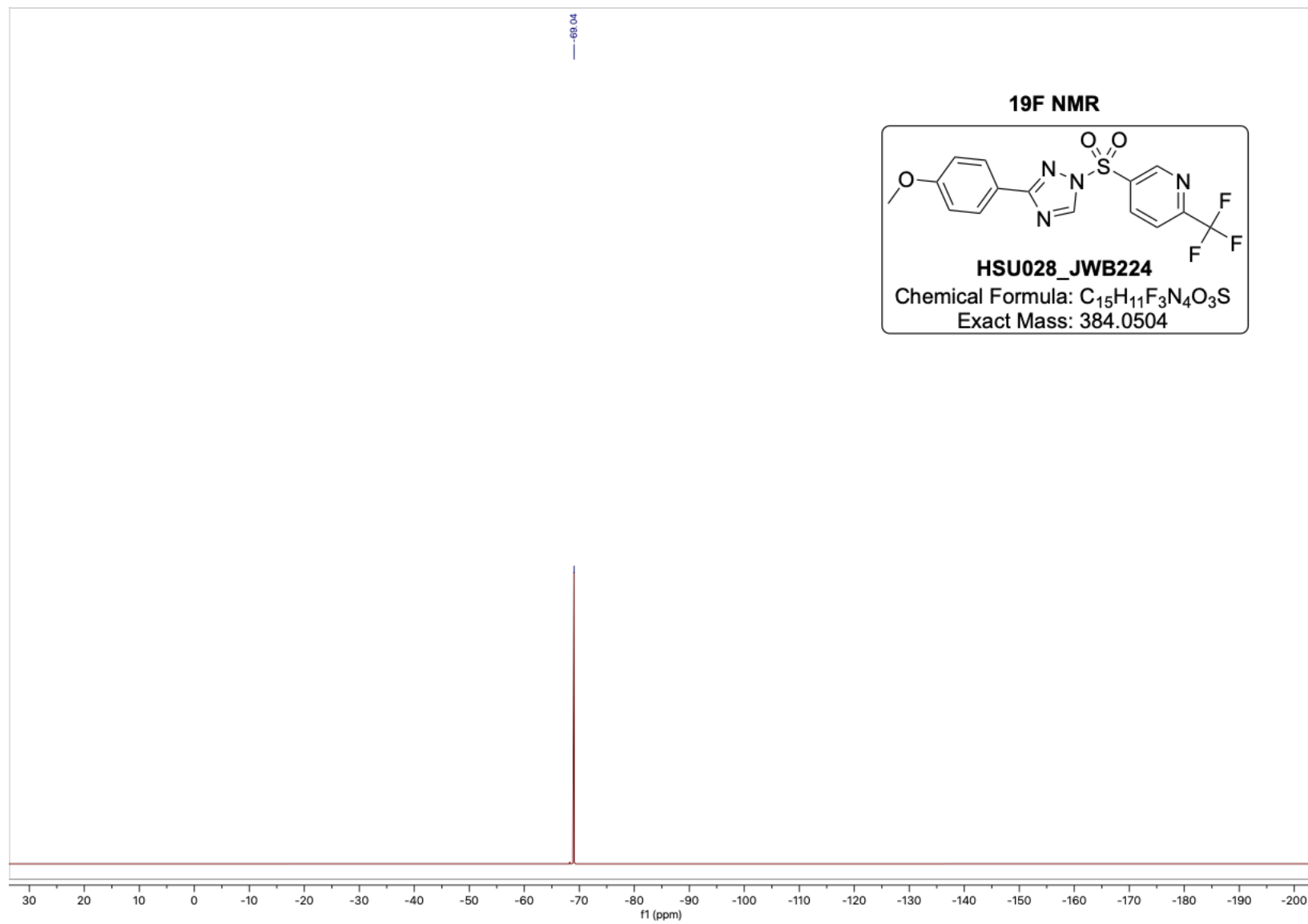
-111.66  
-111.68  
-111.69  
-111.70  
-111.72  
-111.74

# **<sup>19</sup>F NMR**



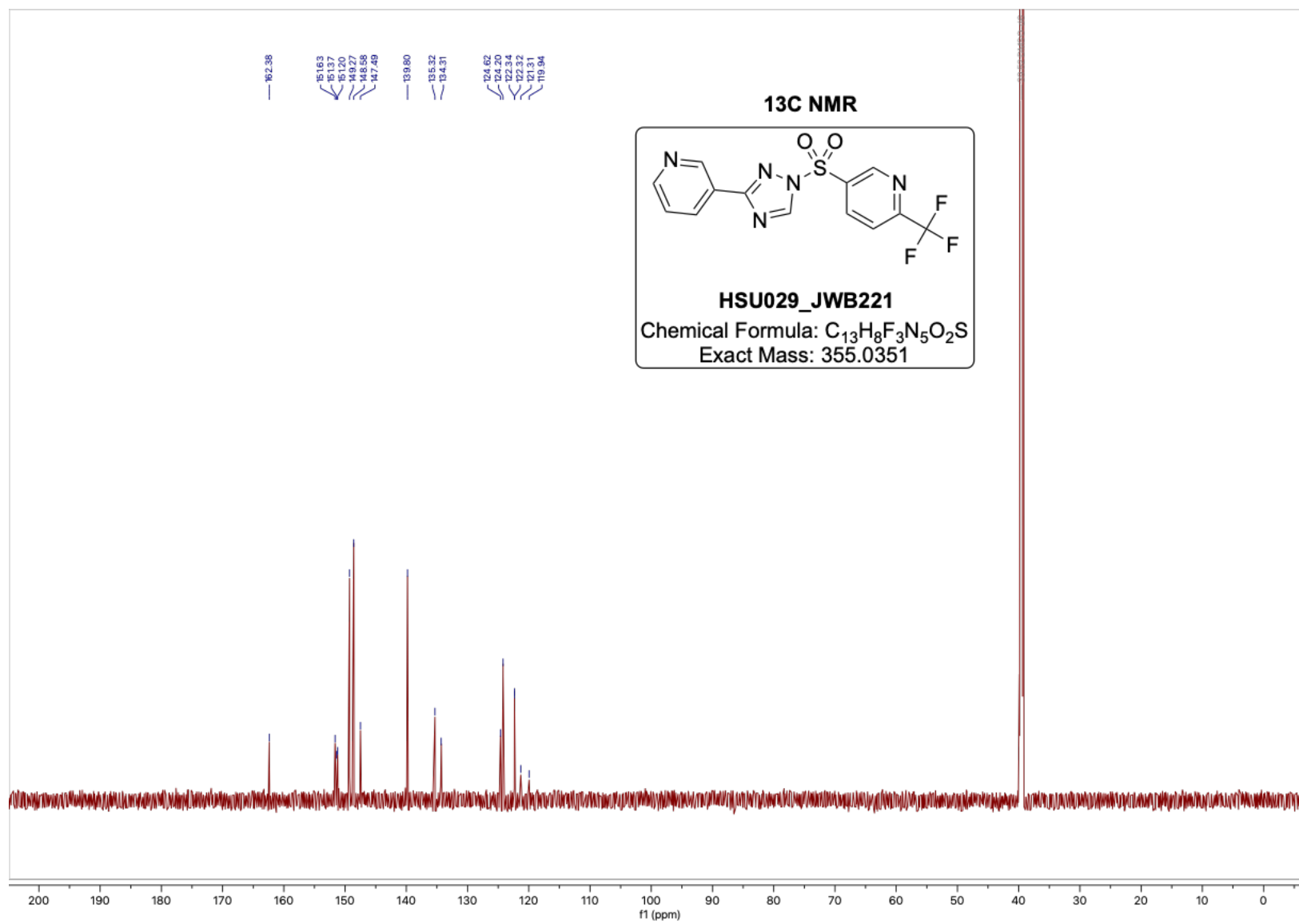


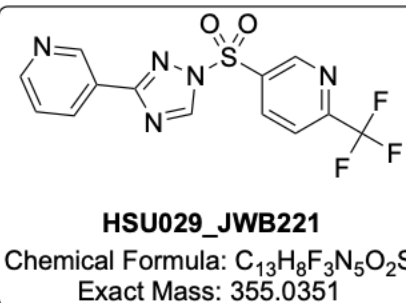
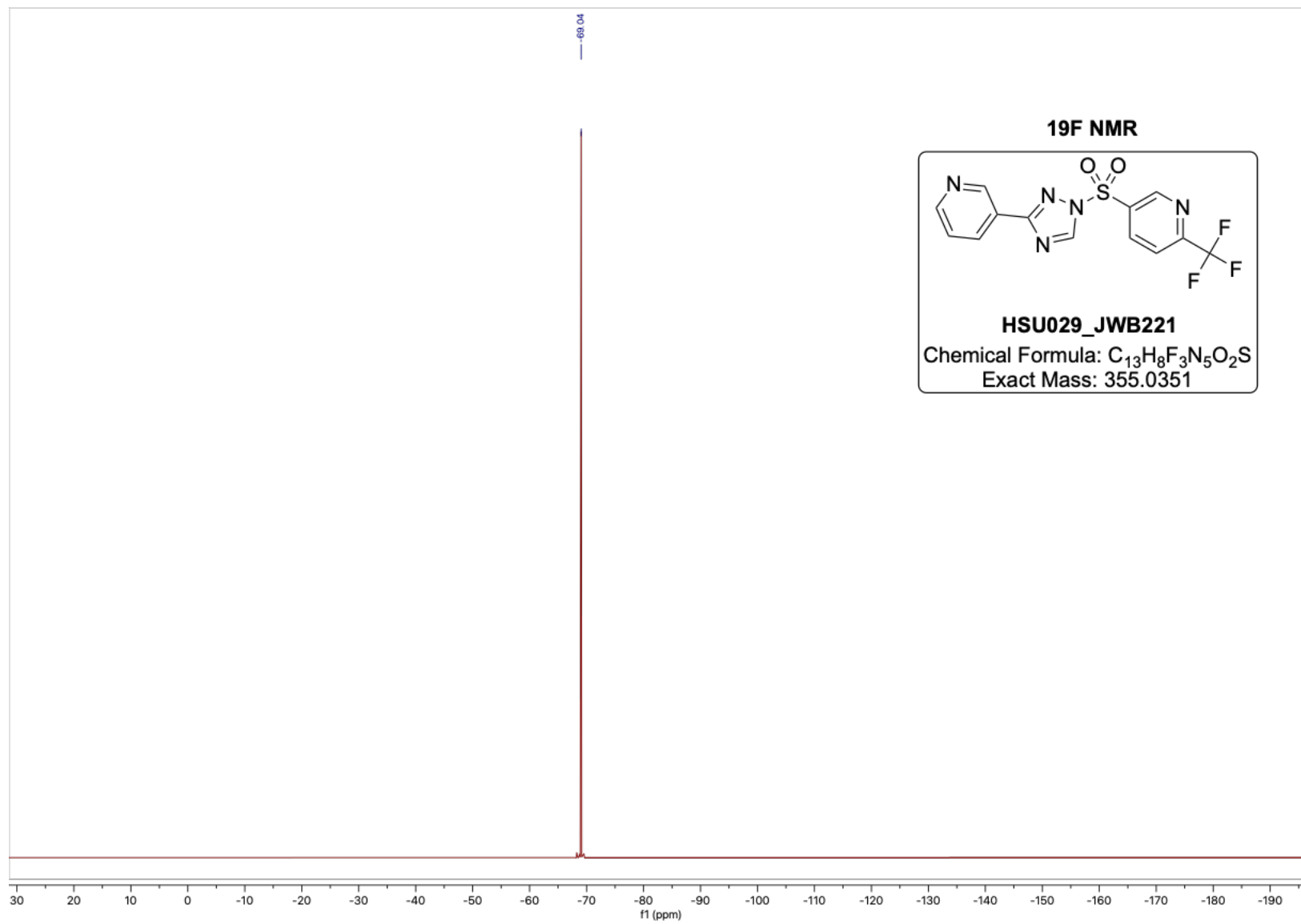


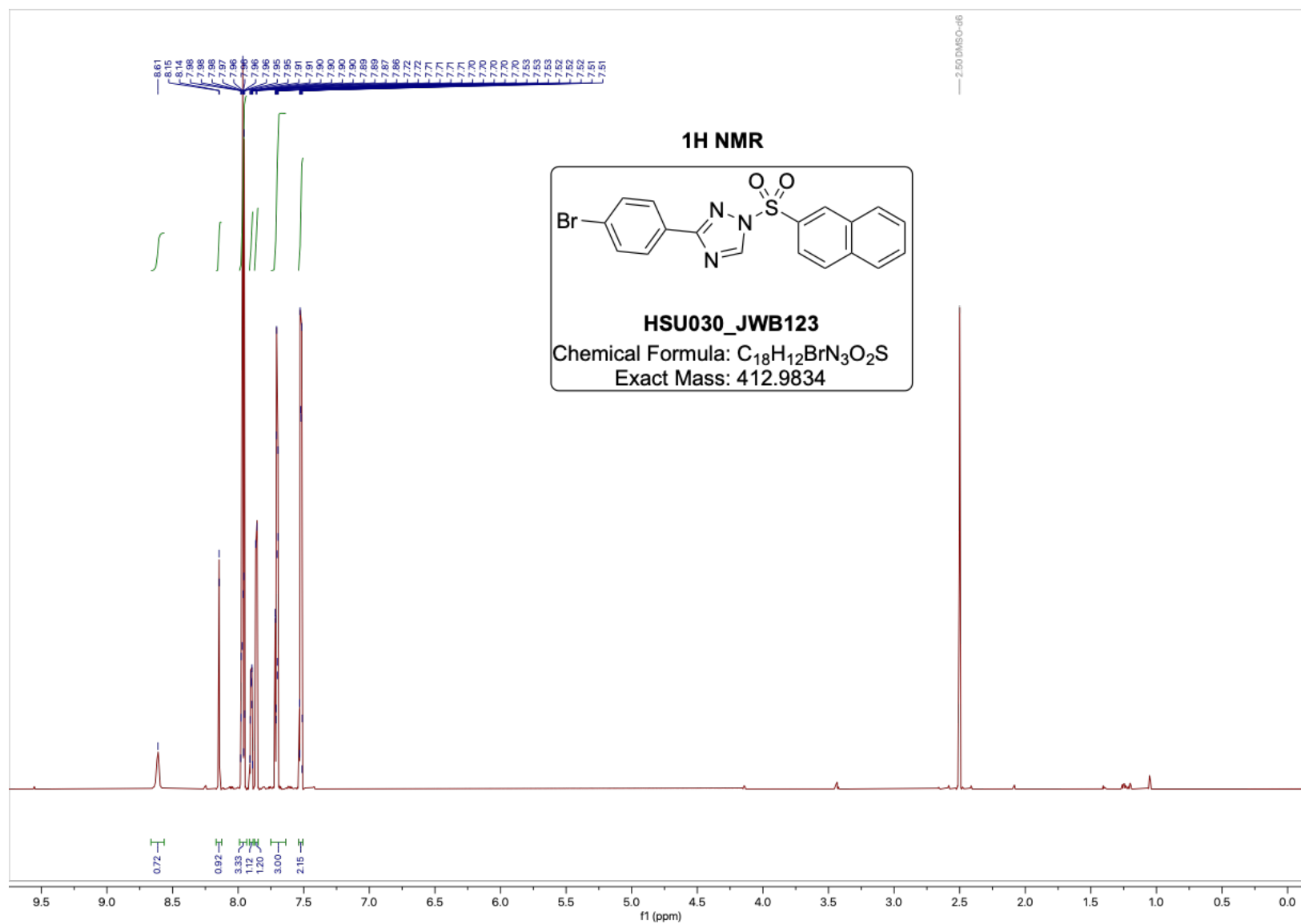


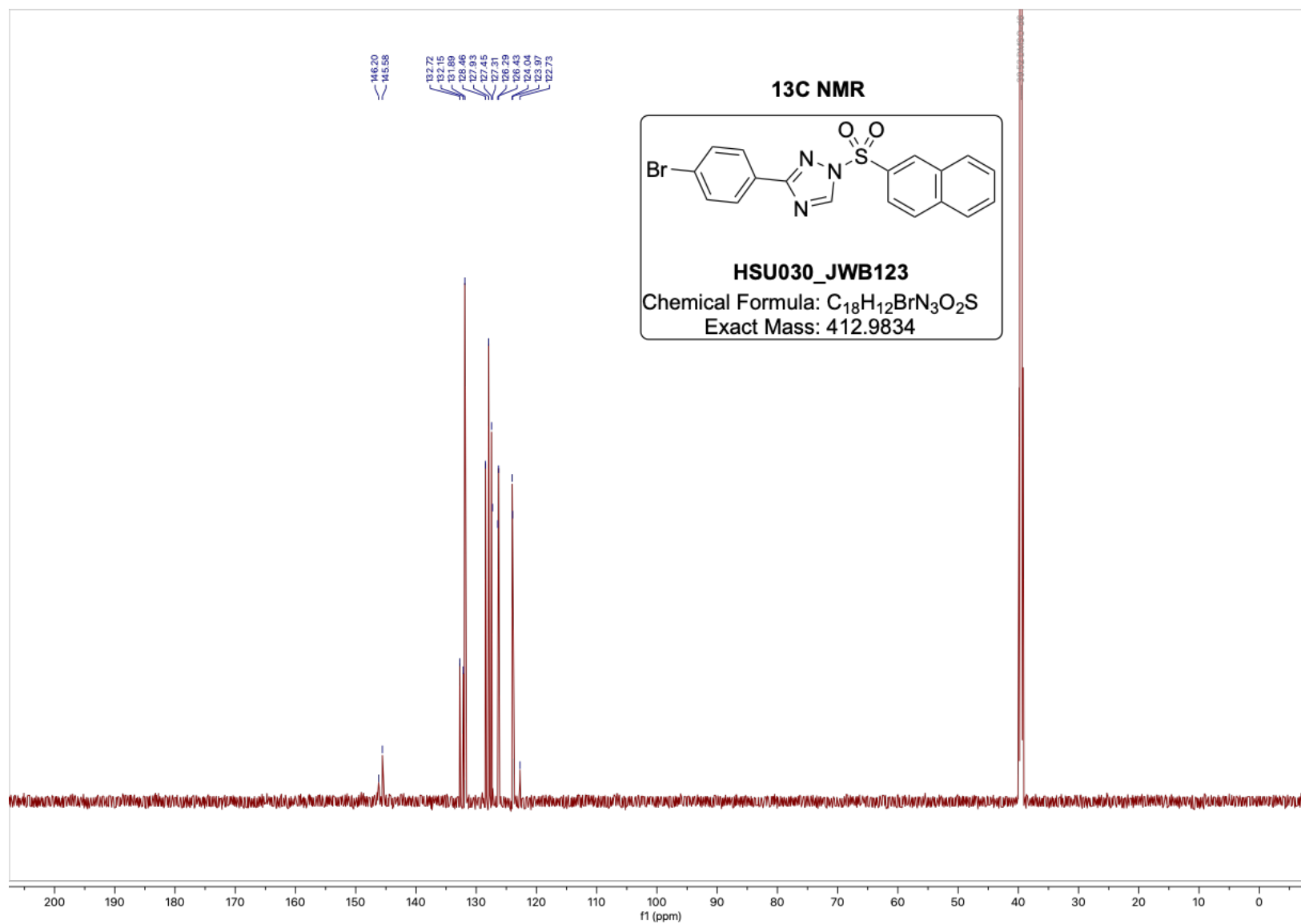


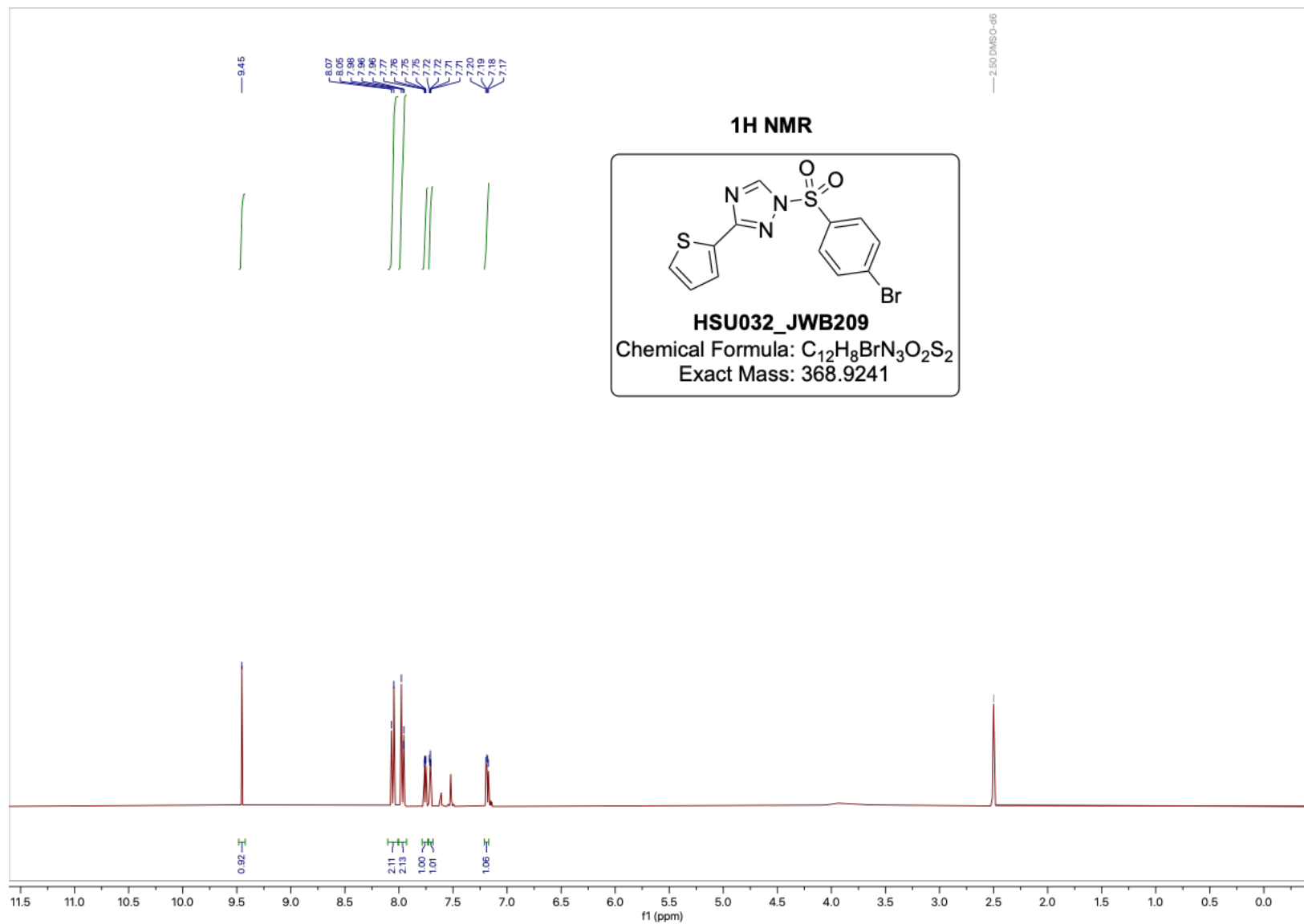


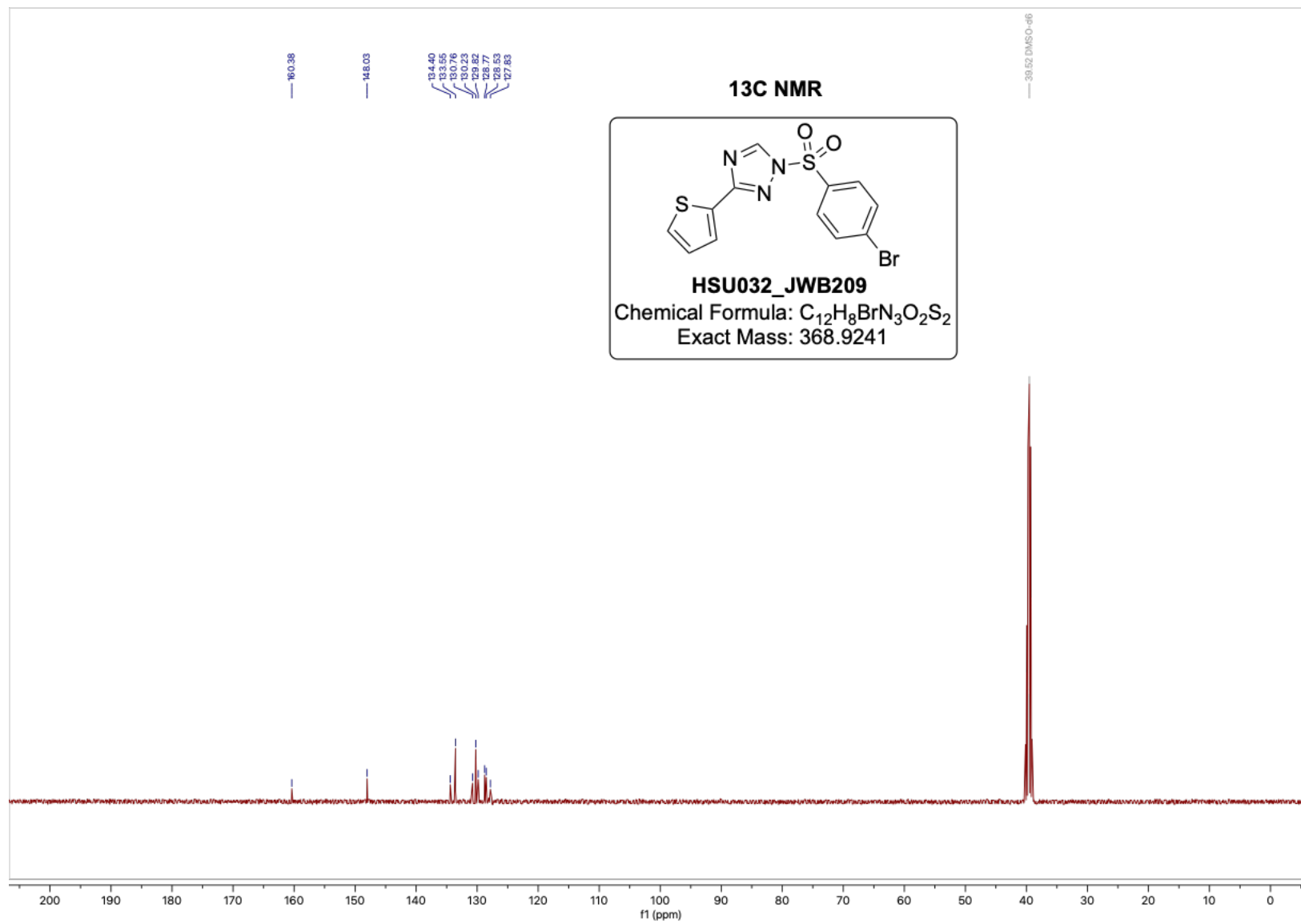


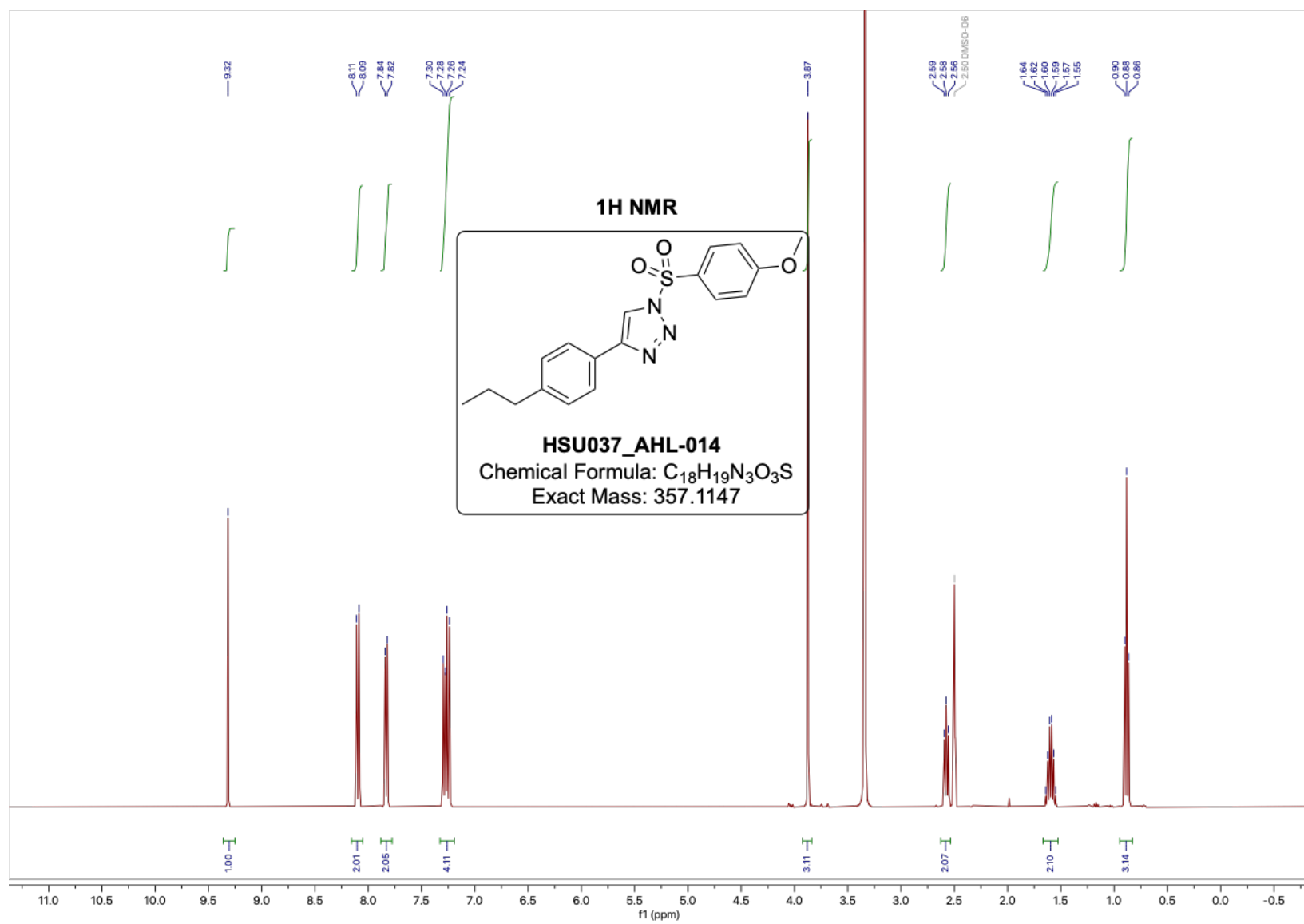


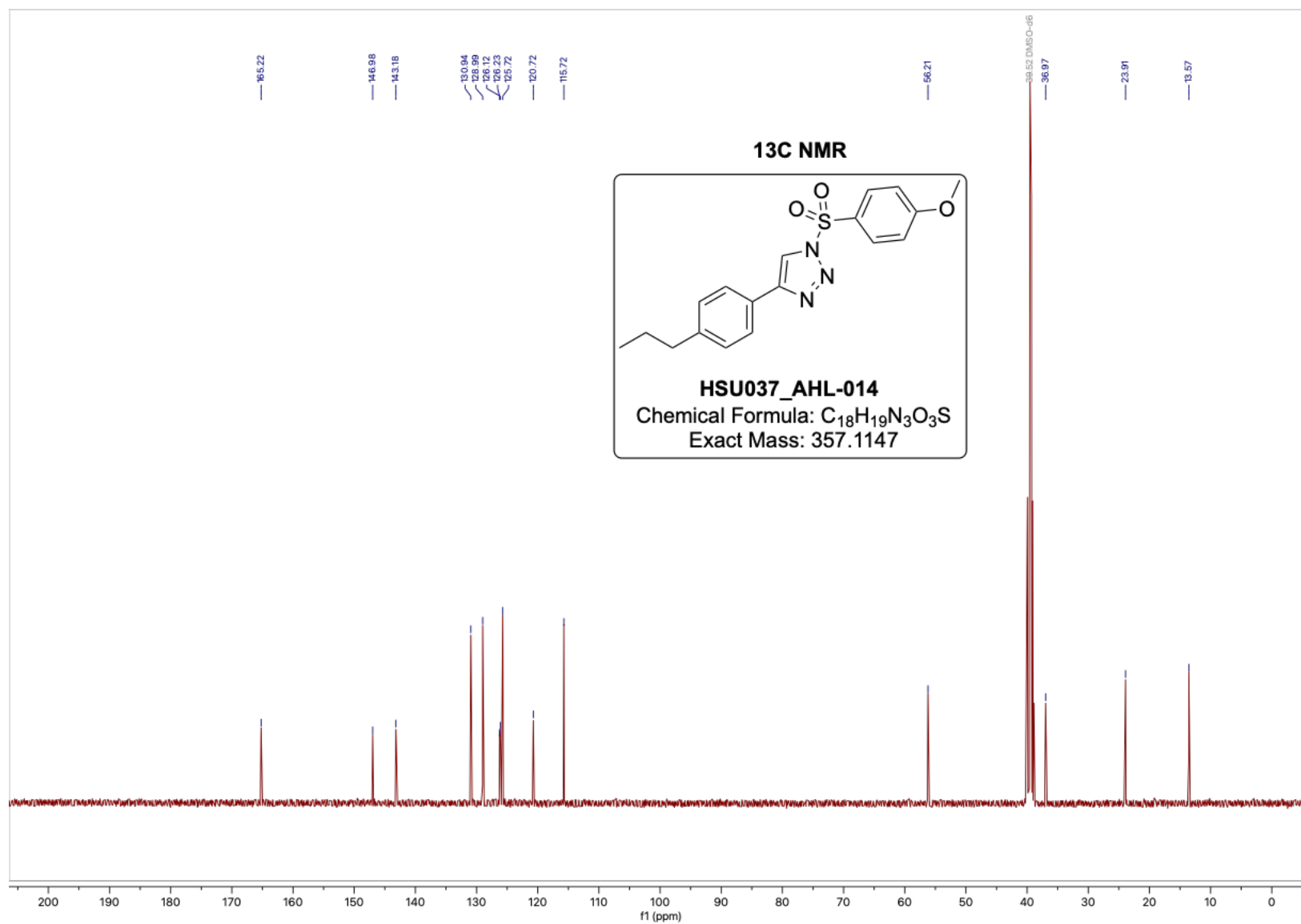




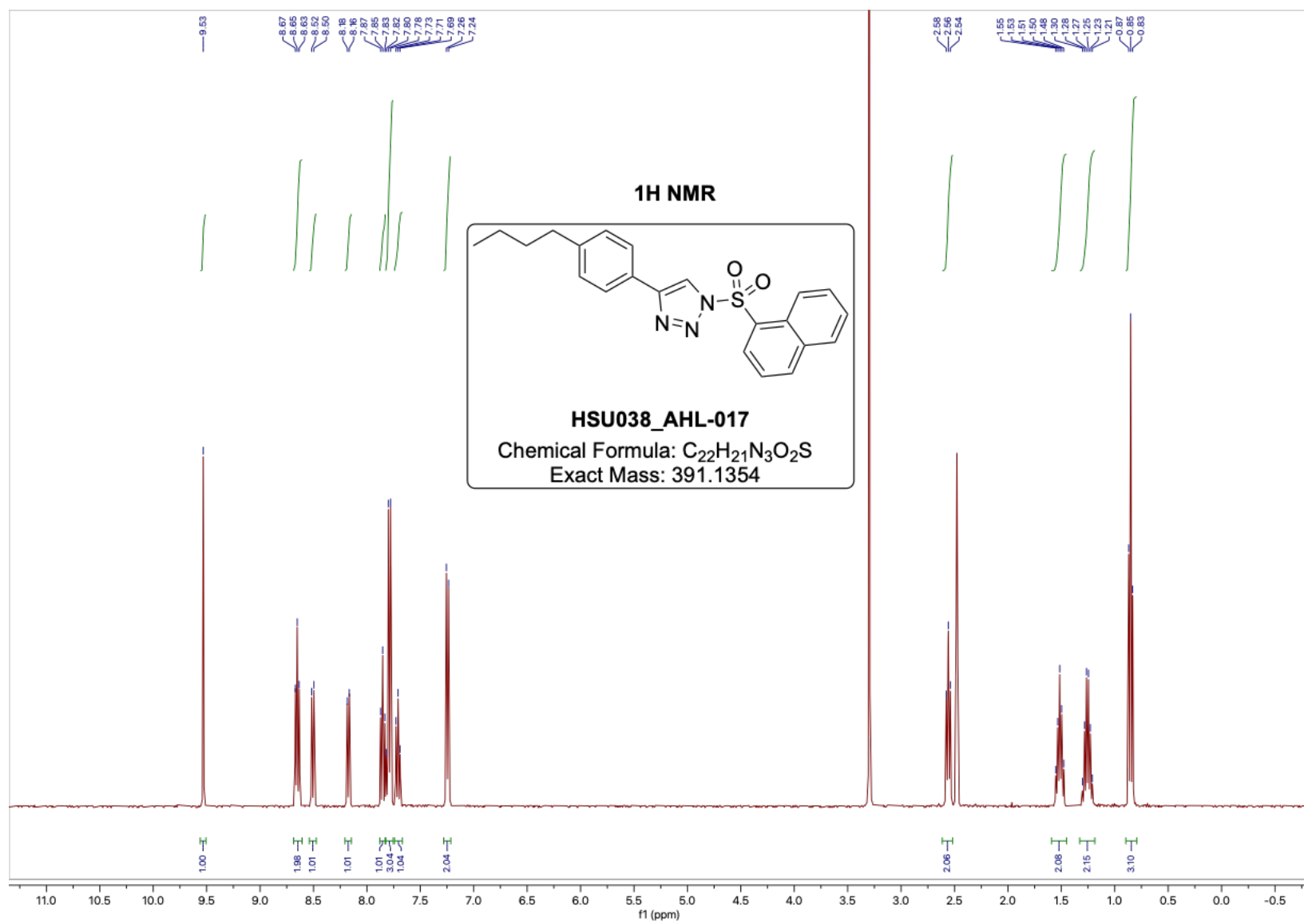


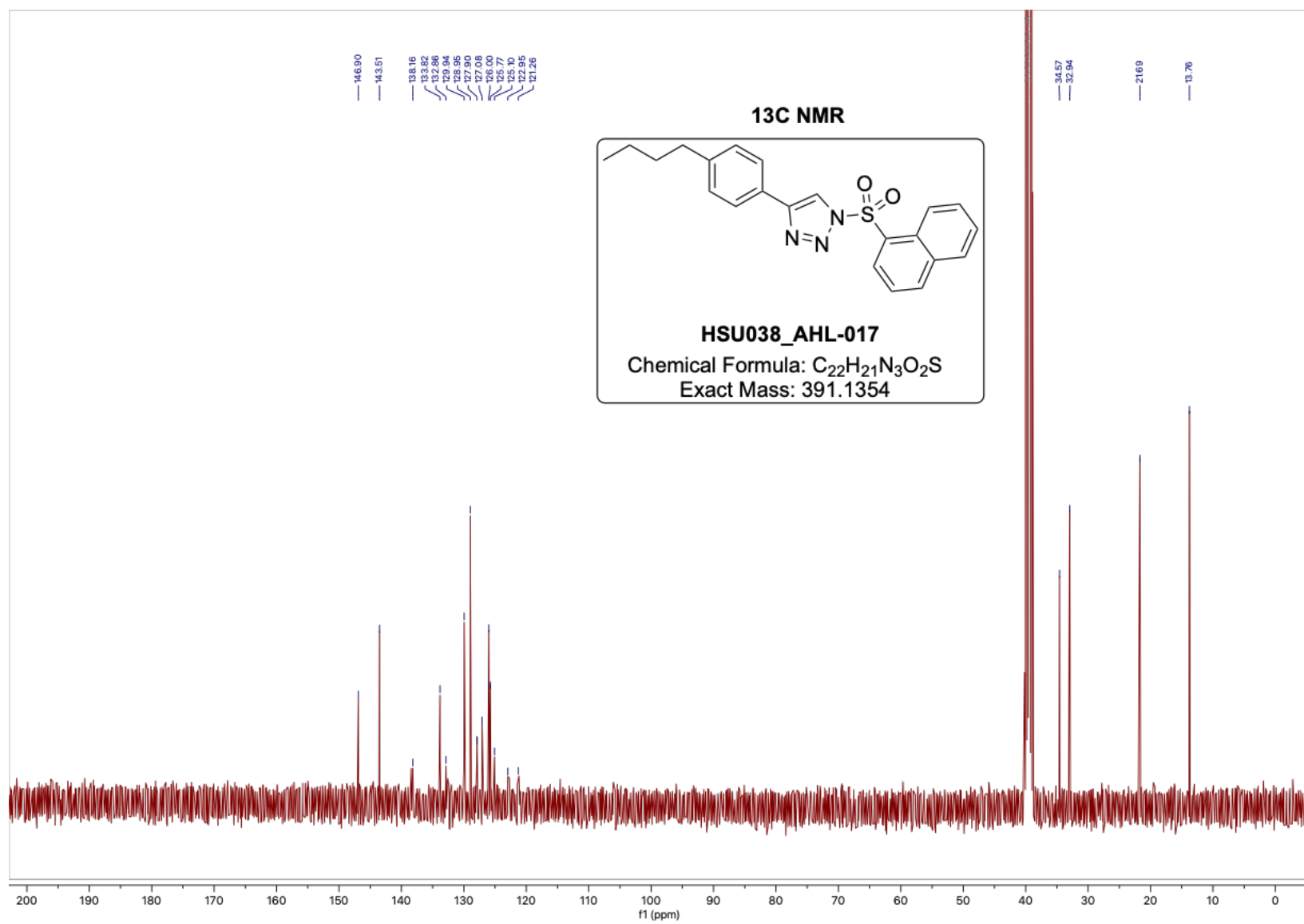


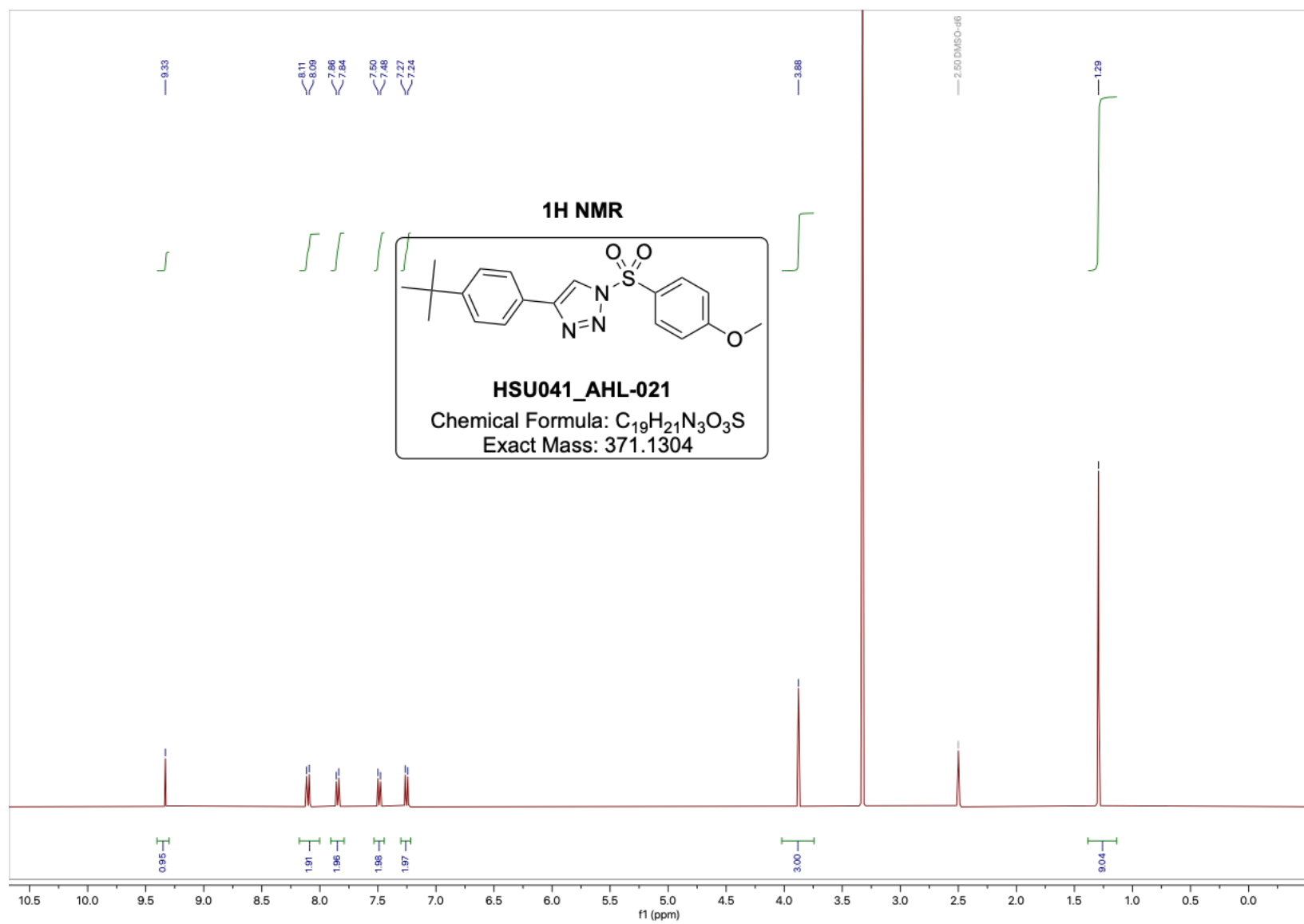


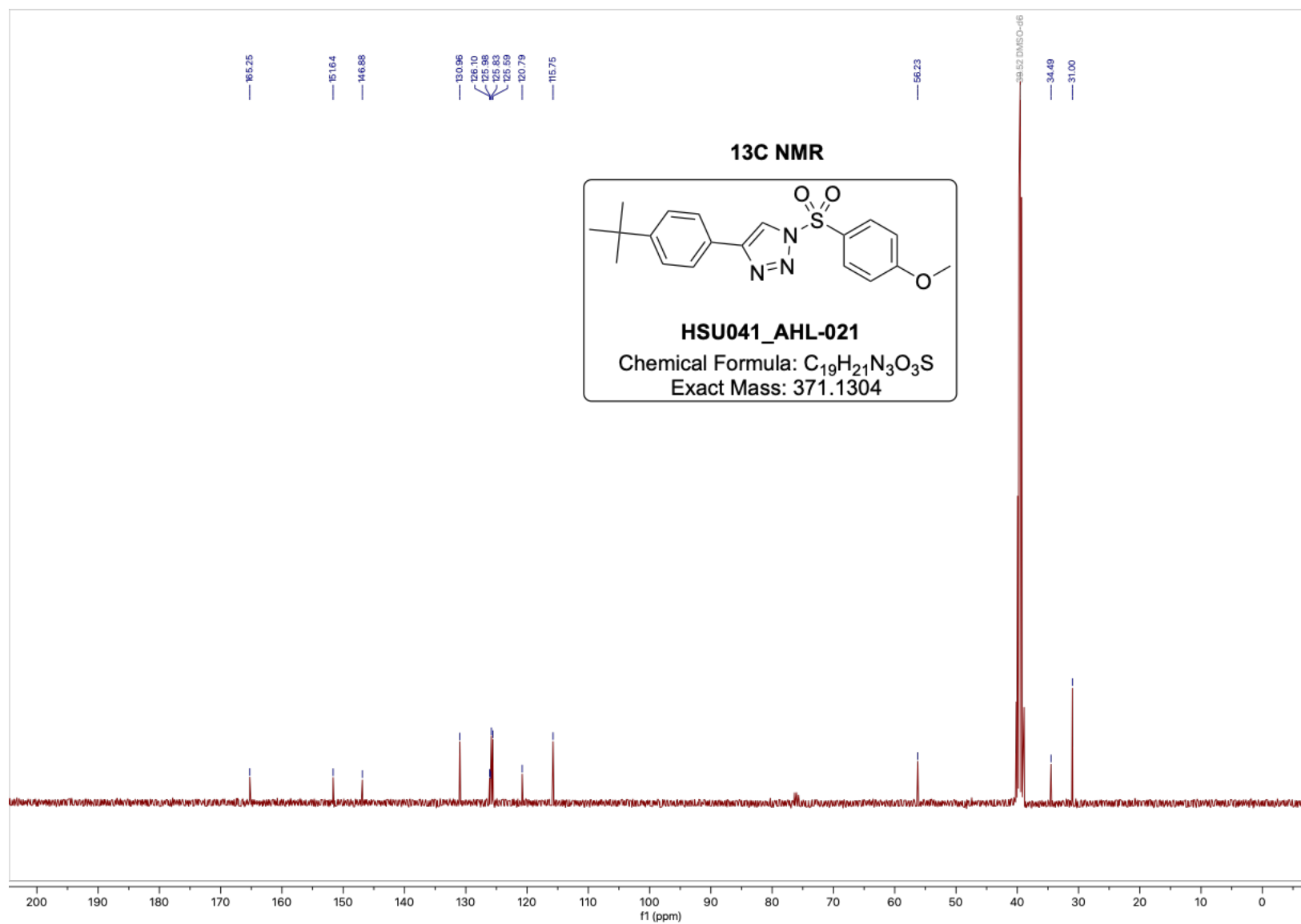


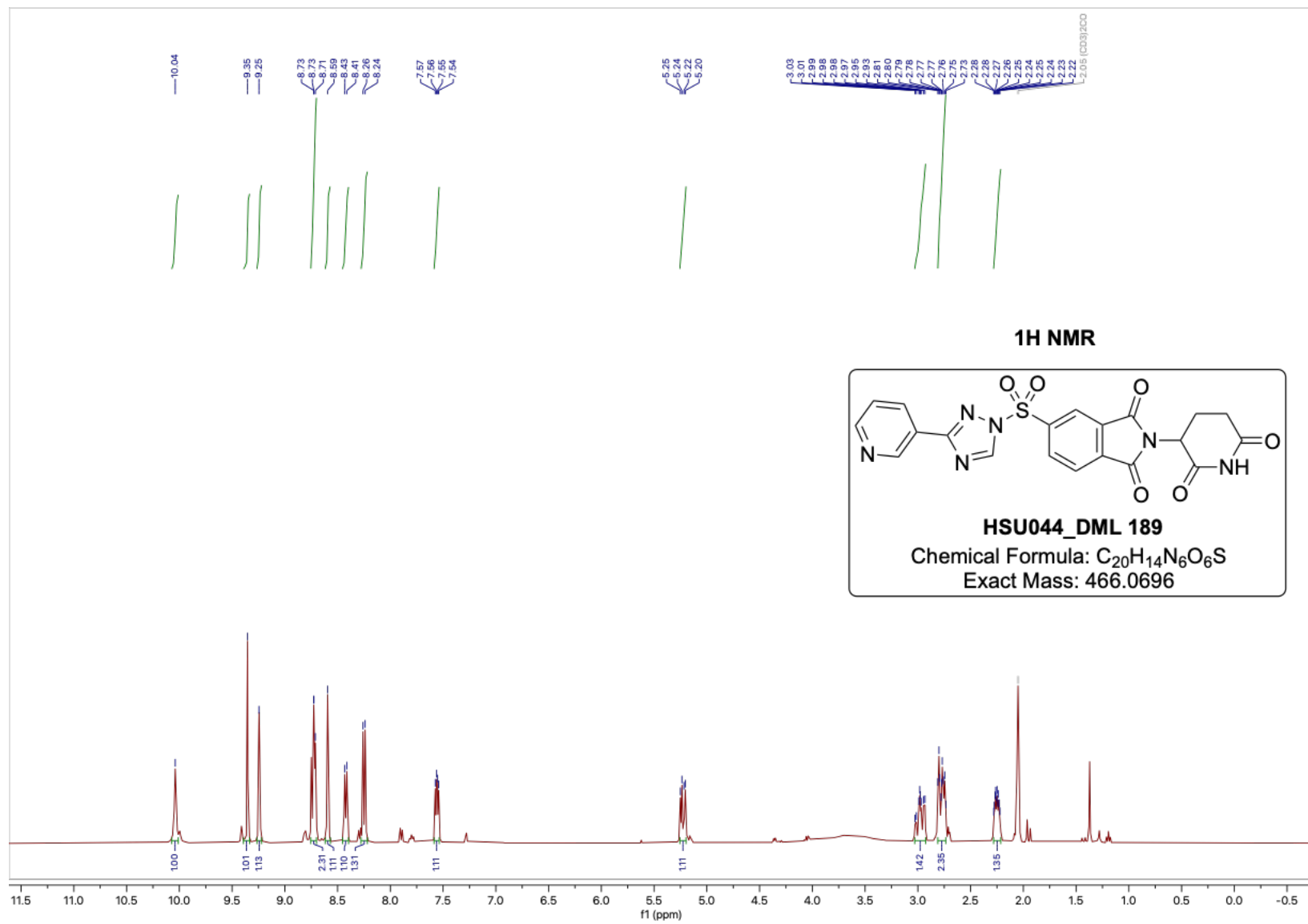


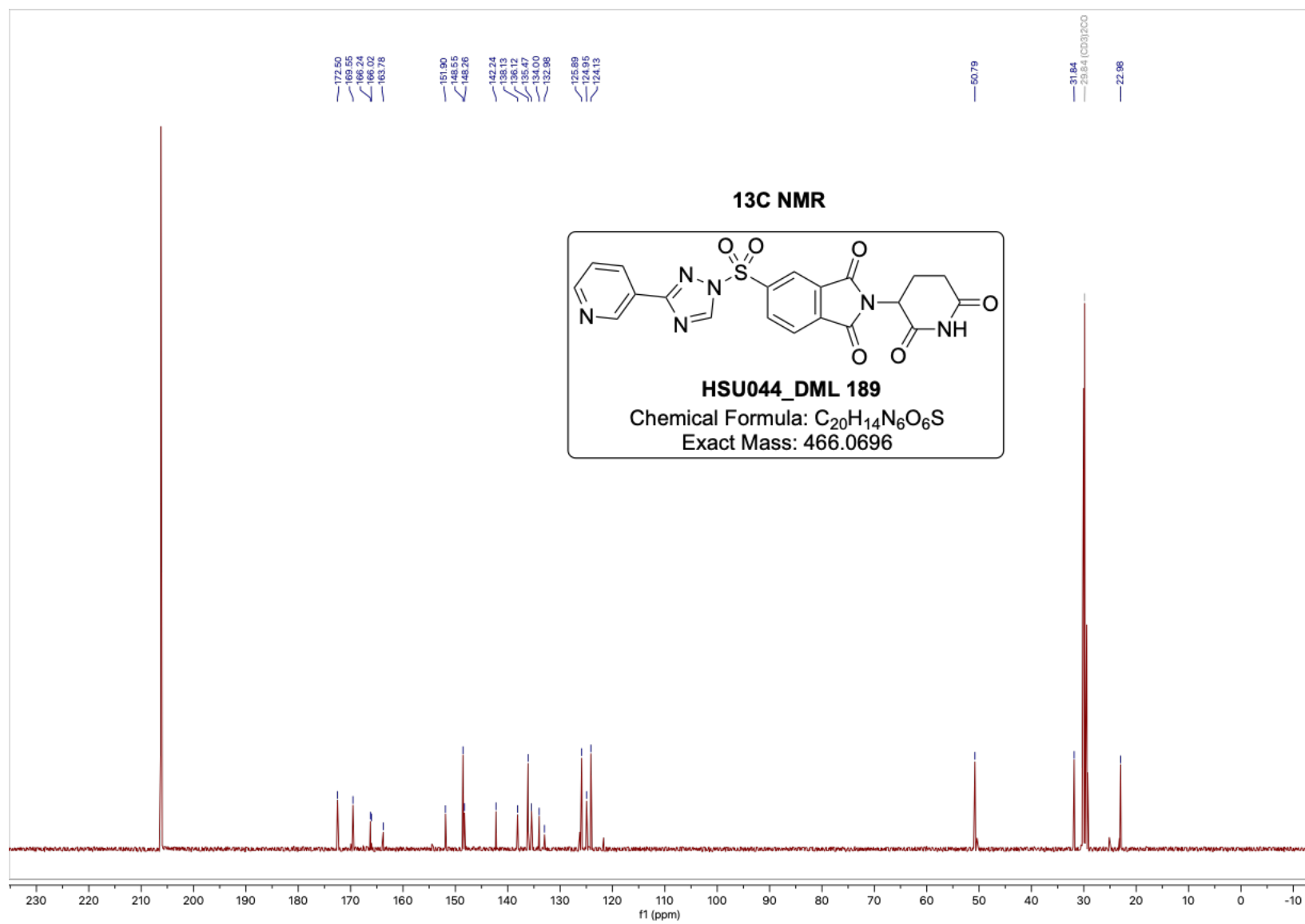


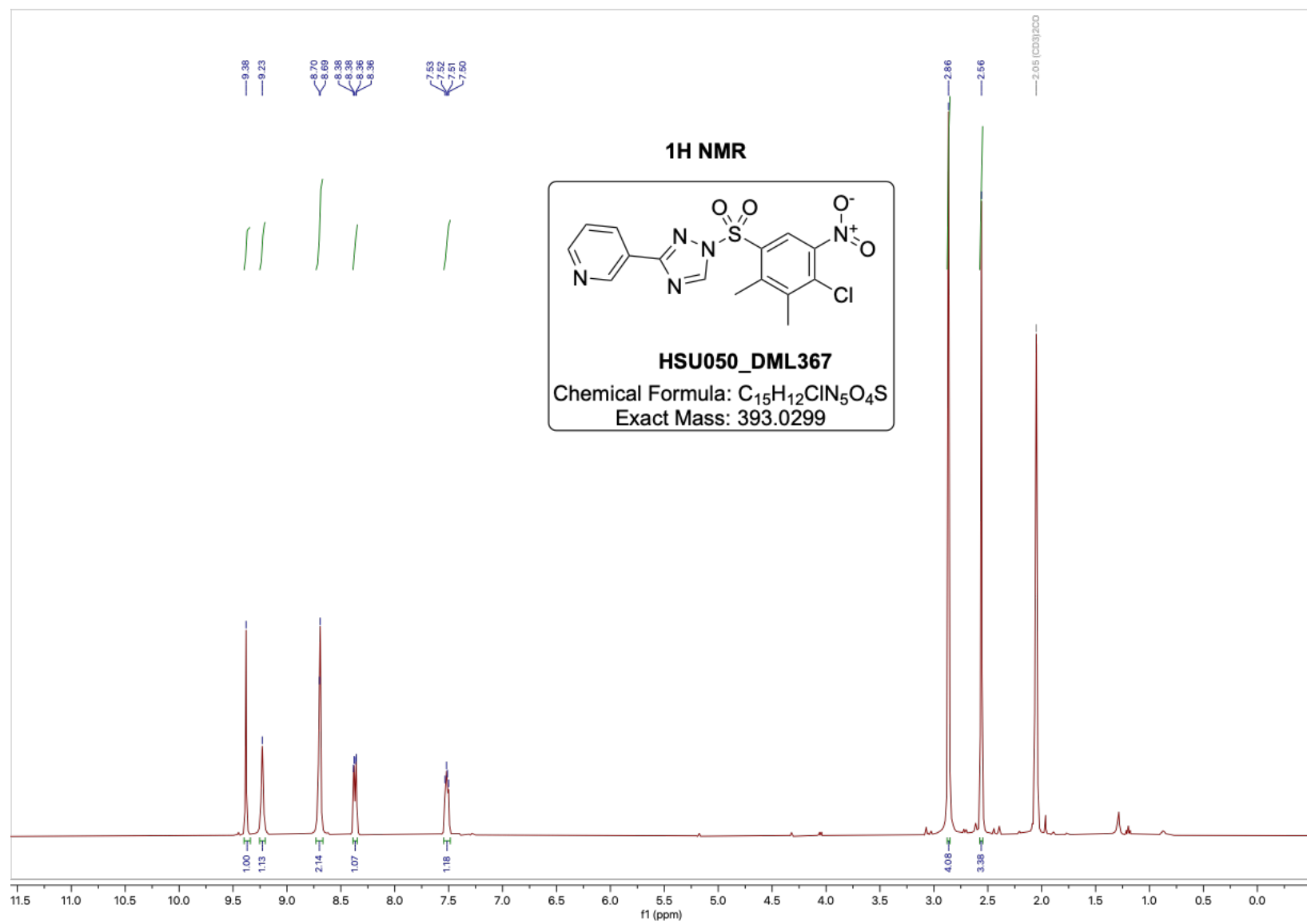


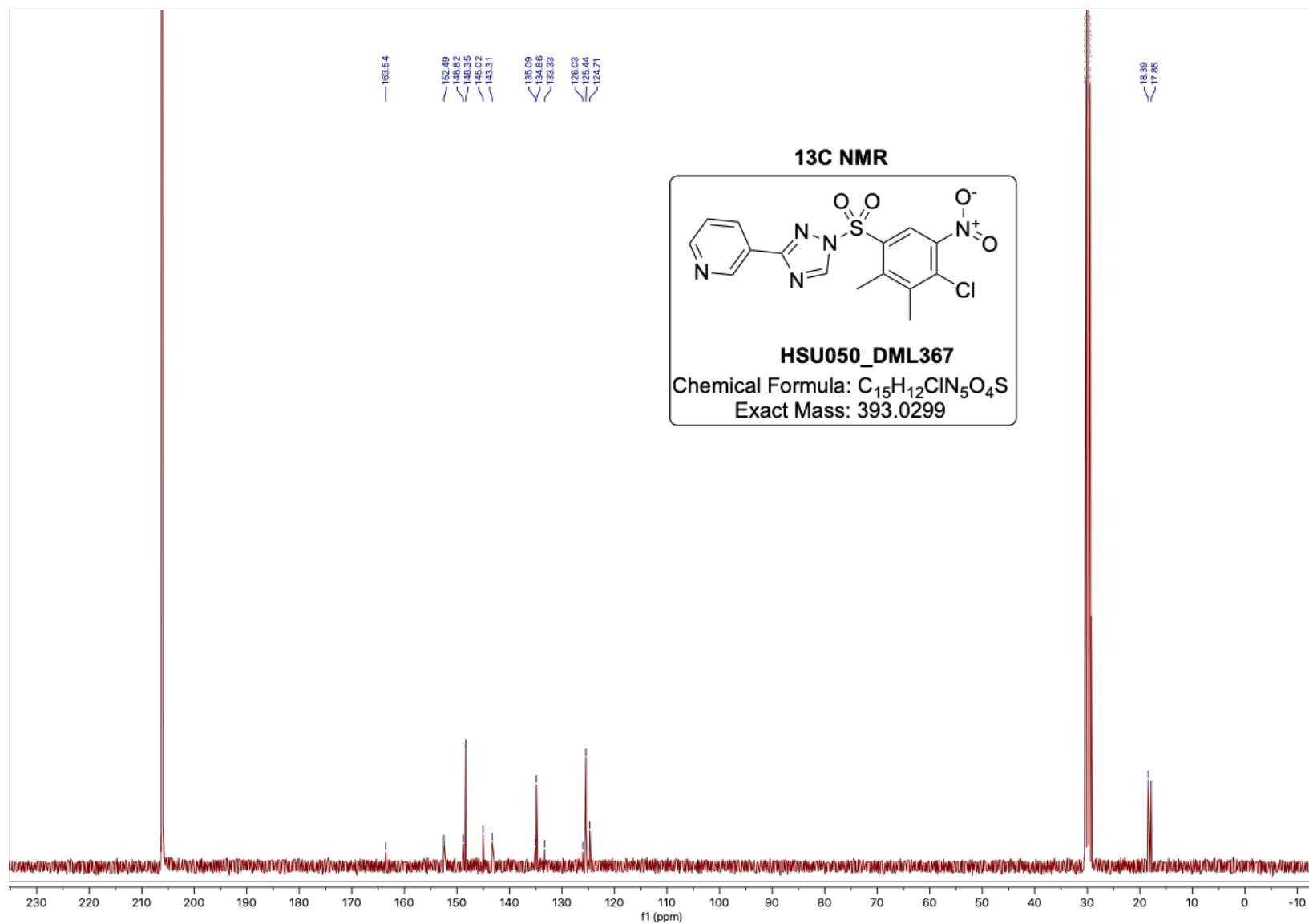




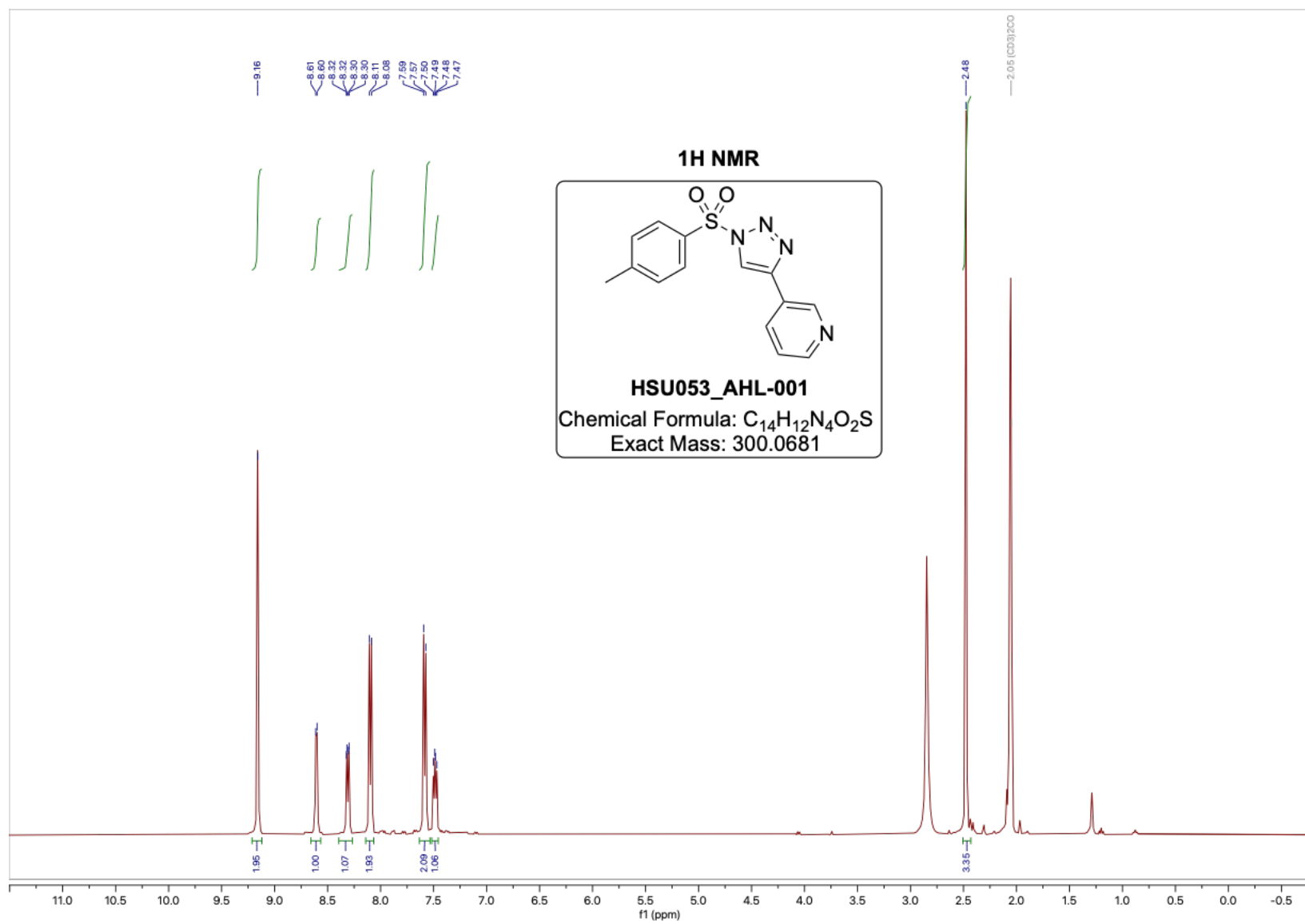


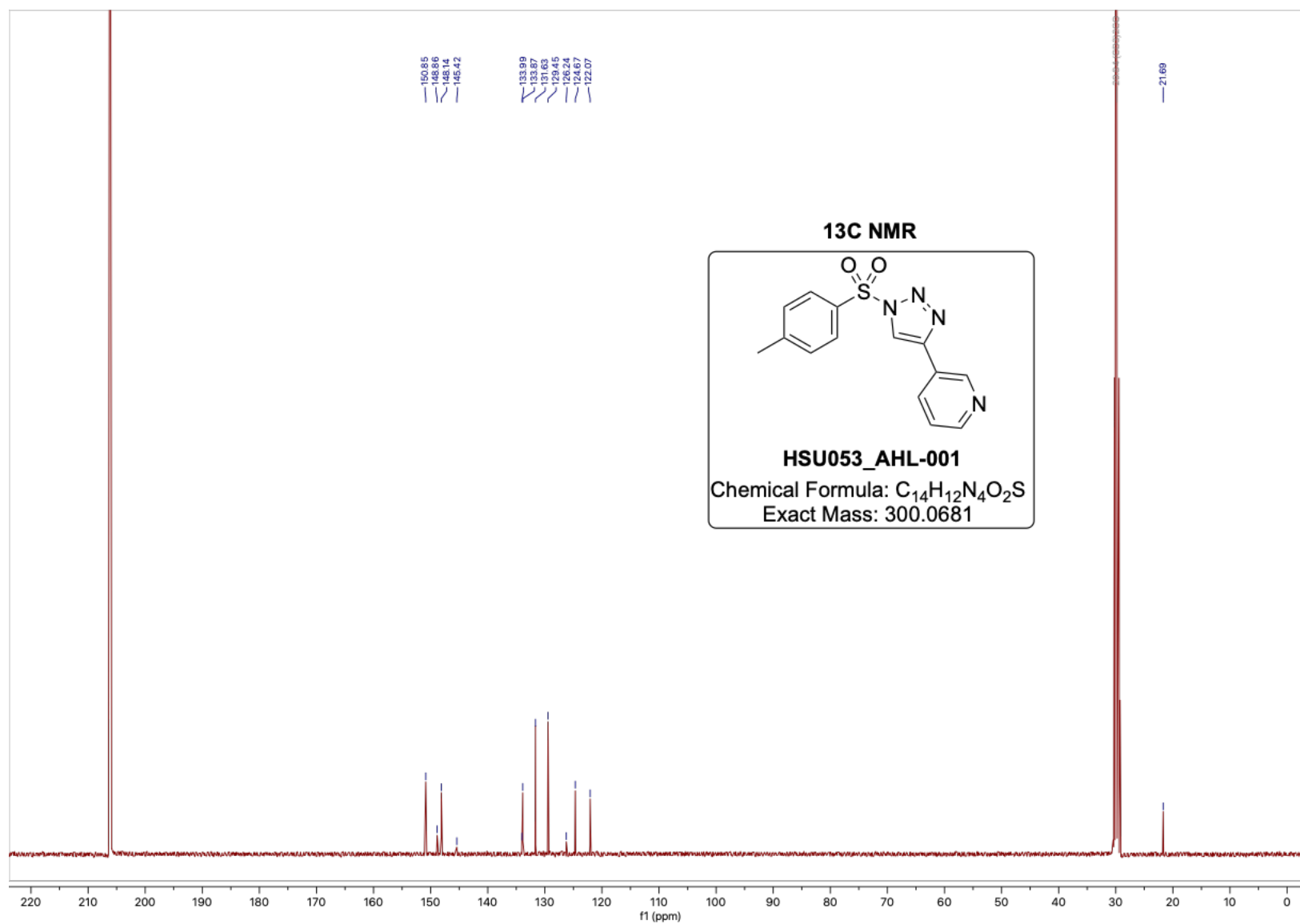


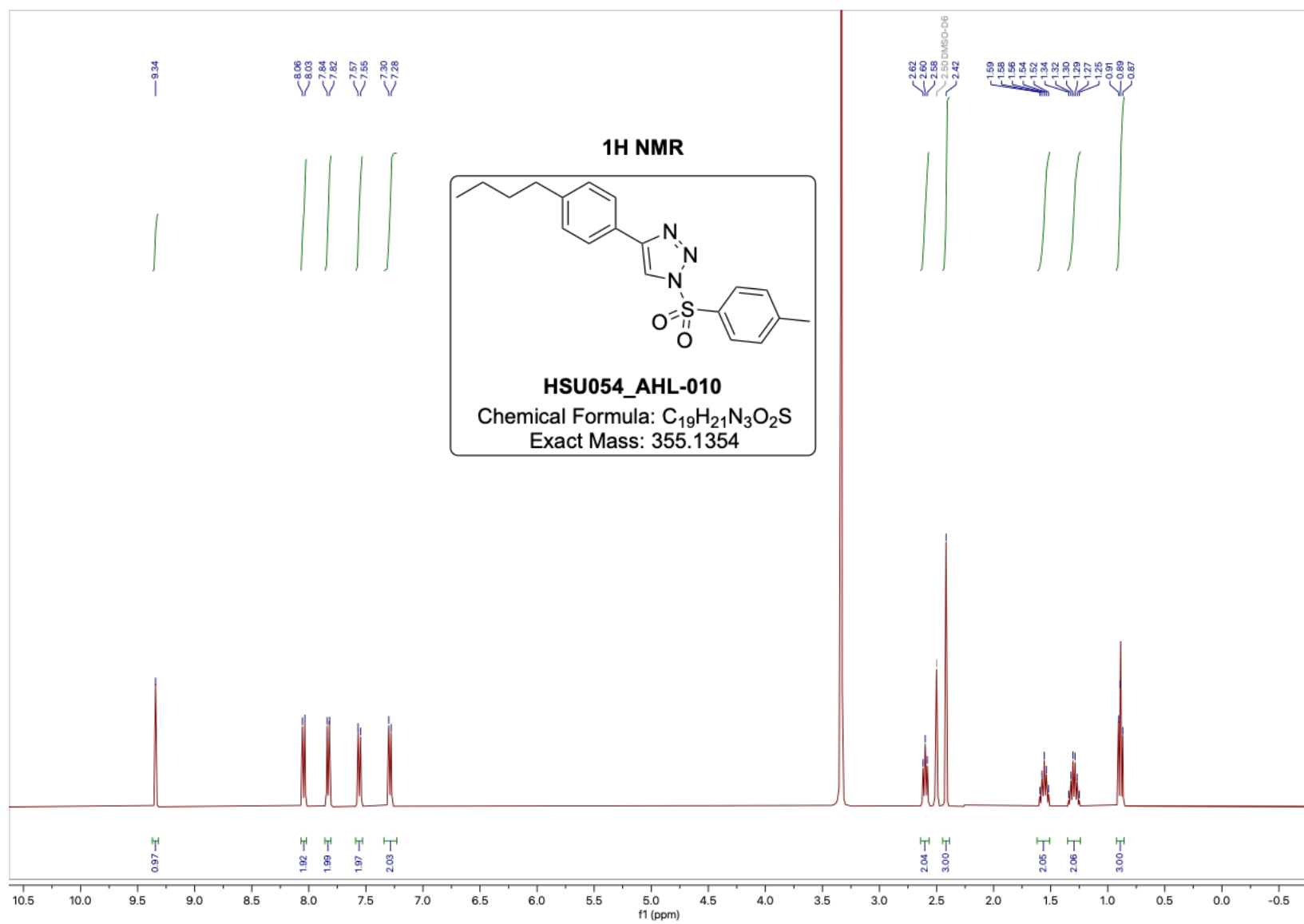


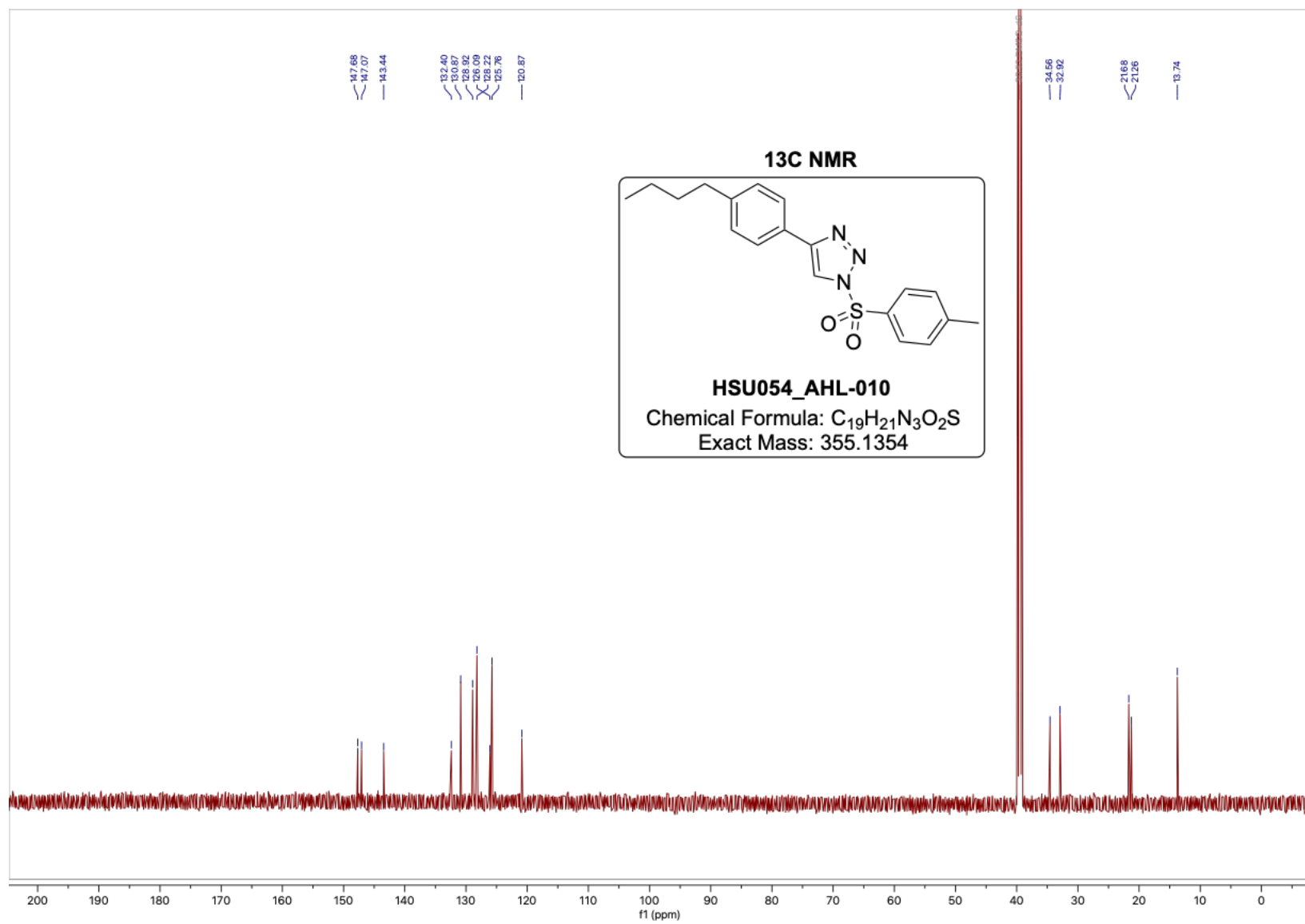


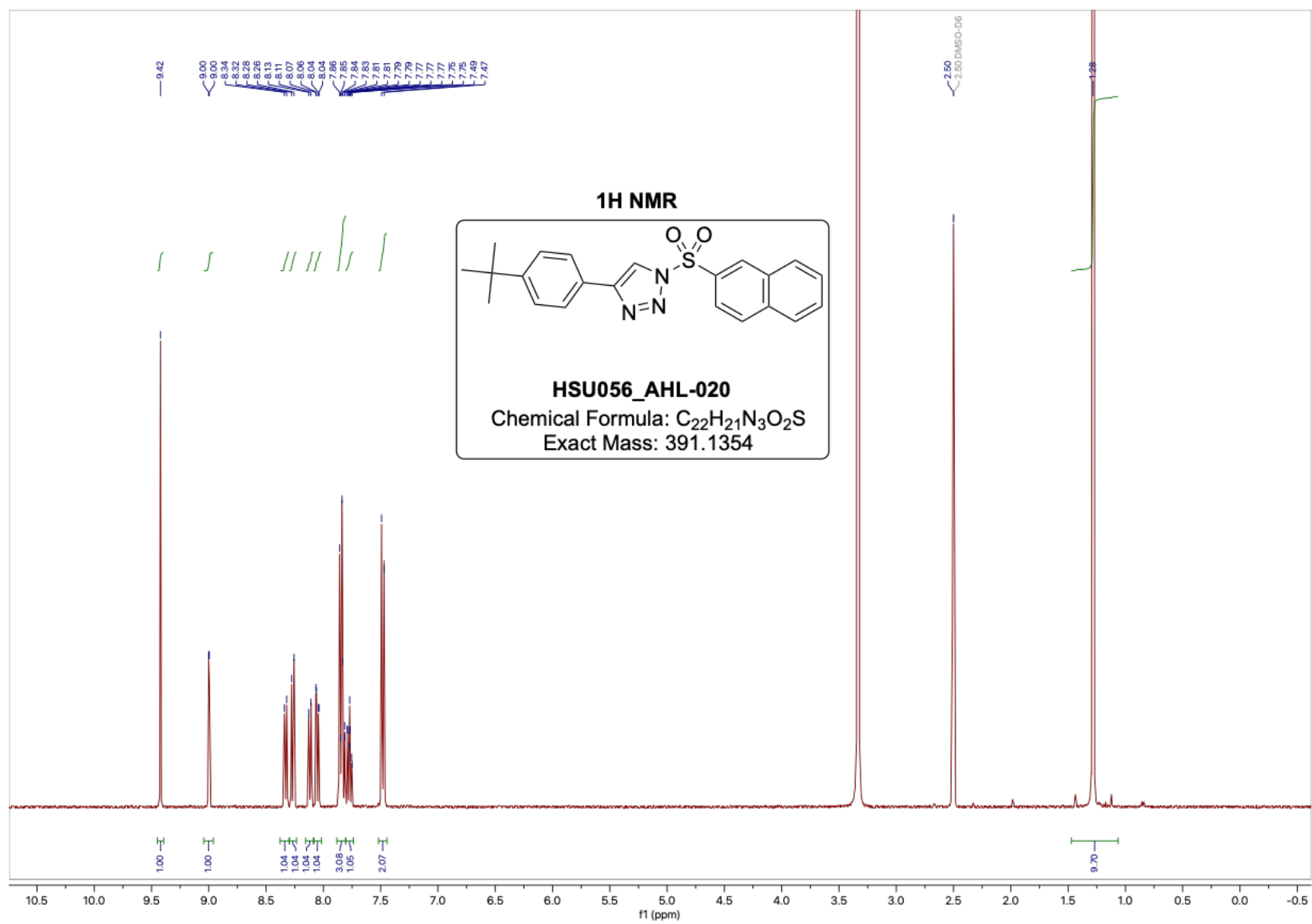


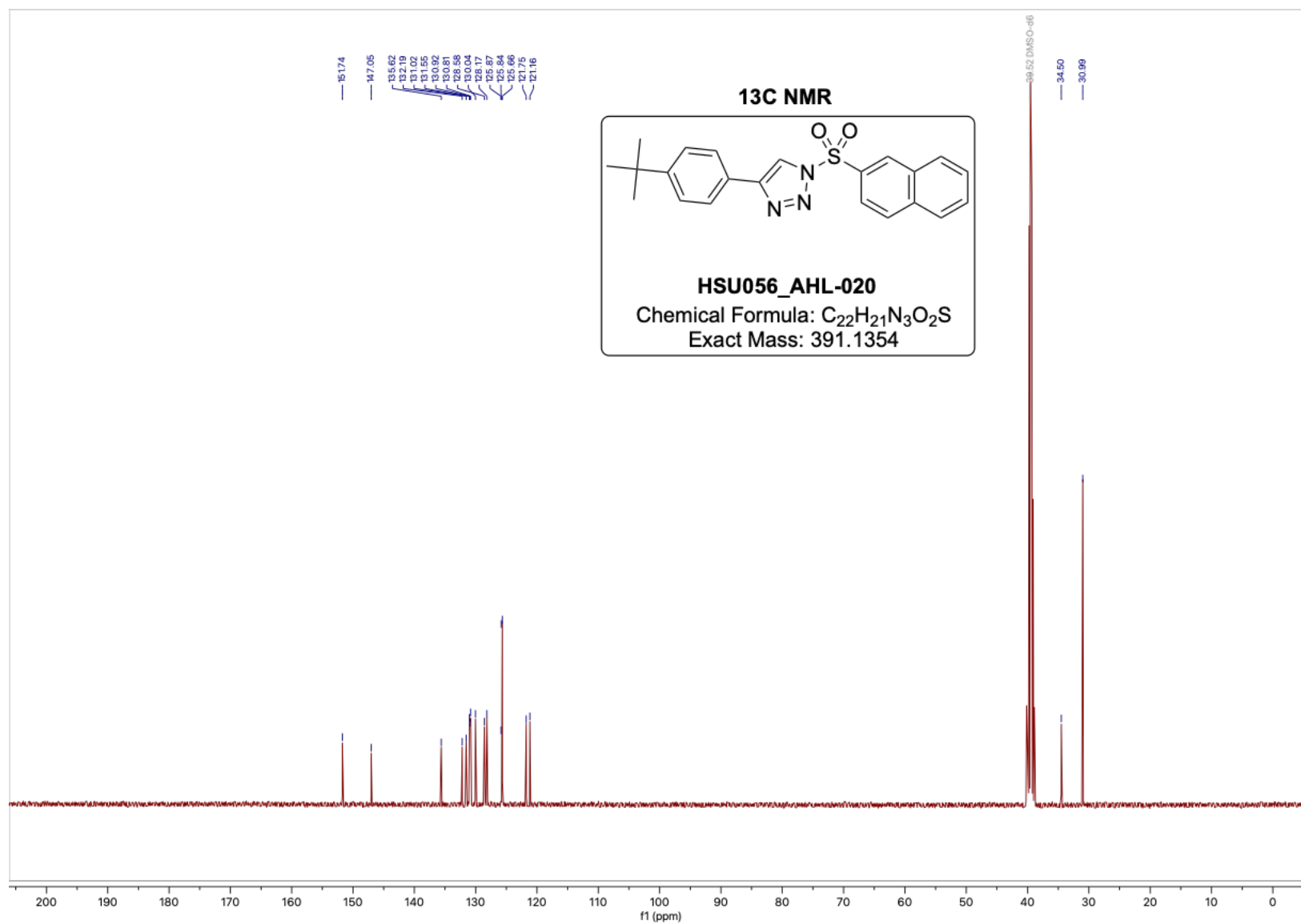


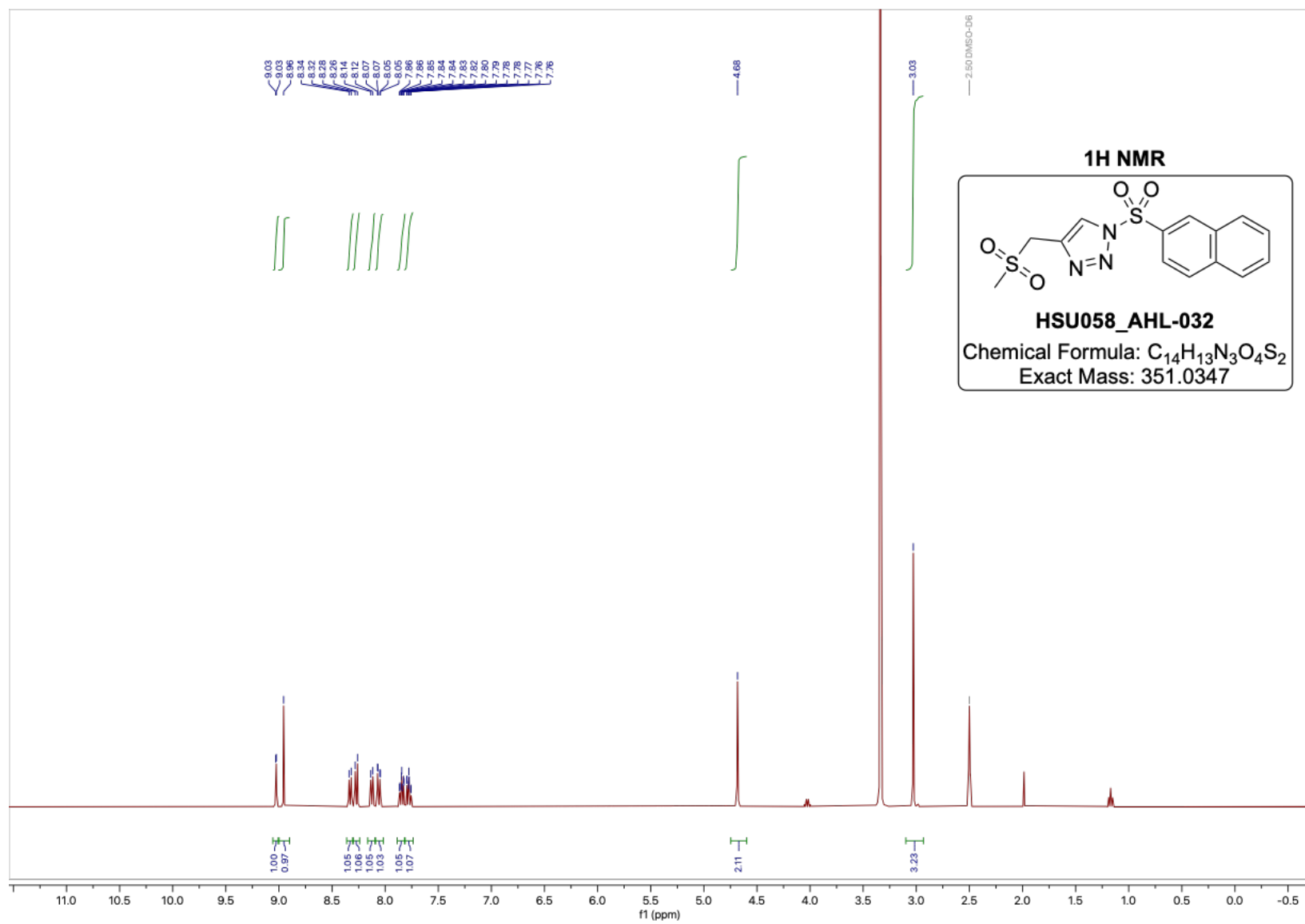


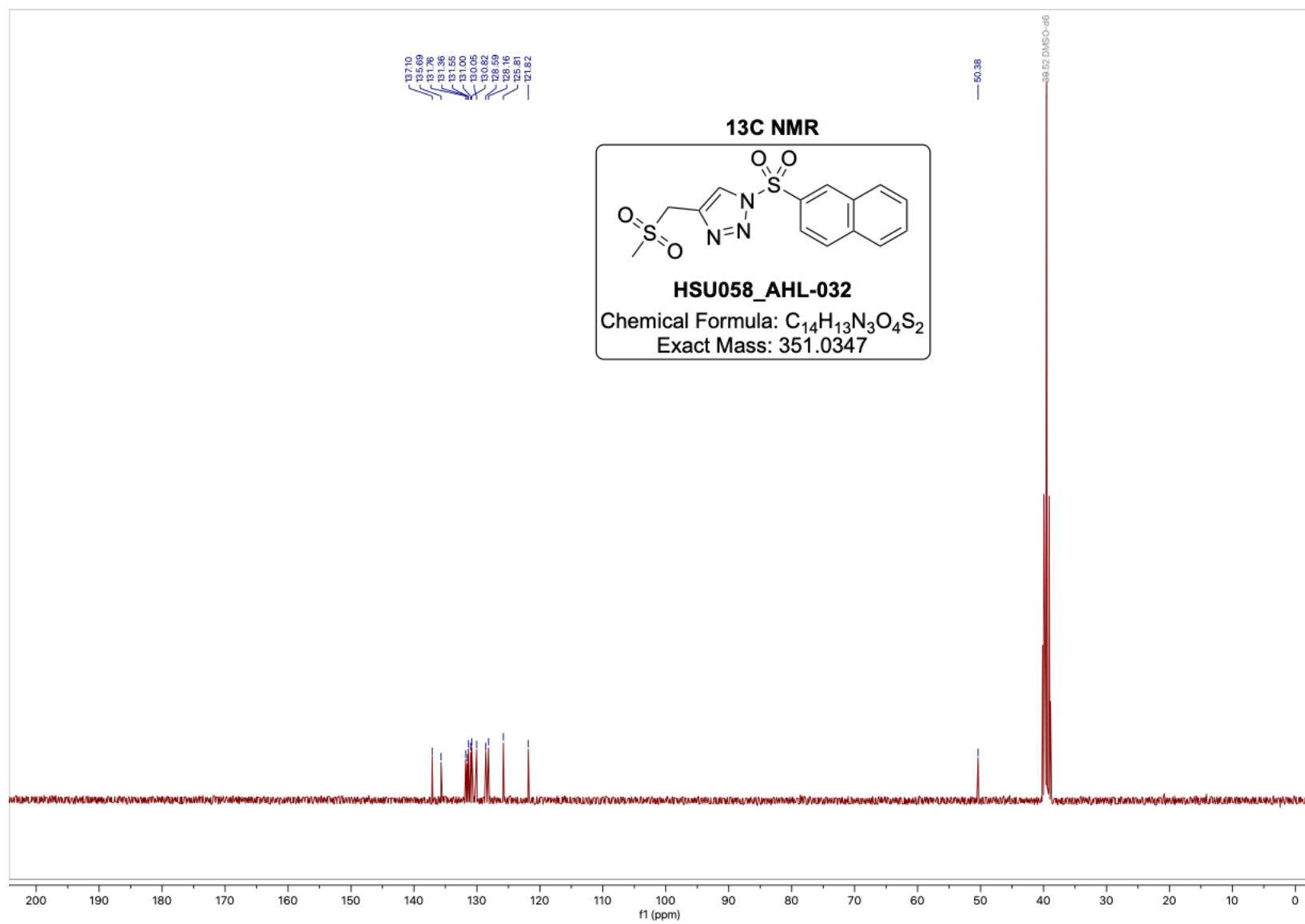




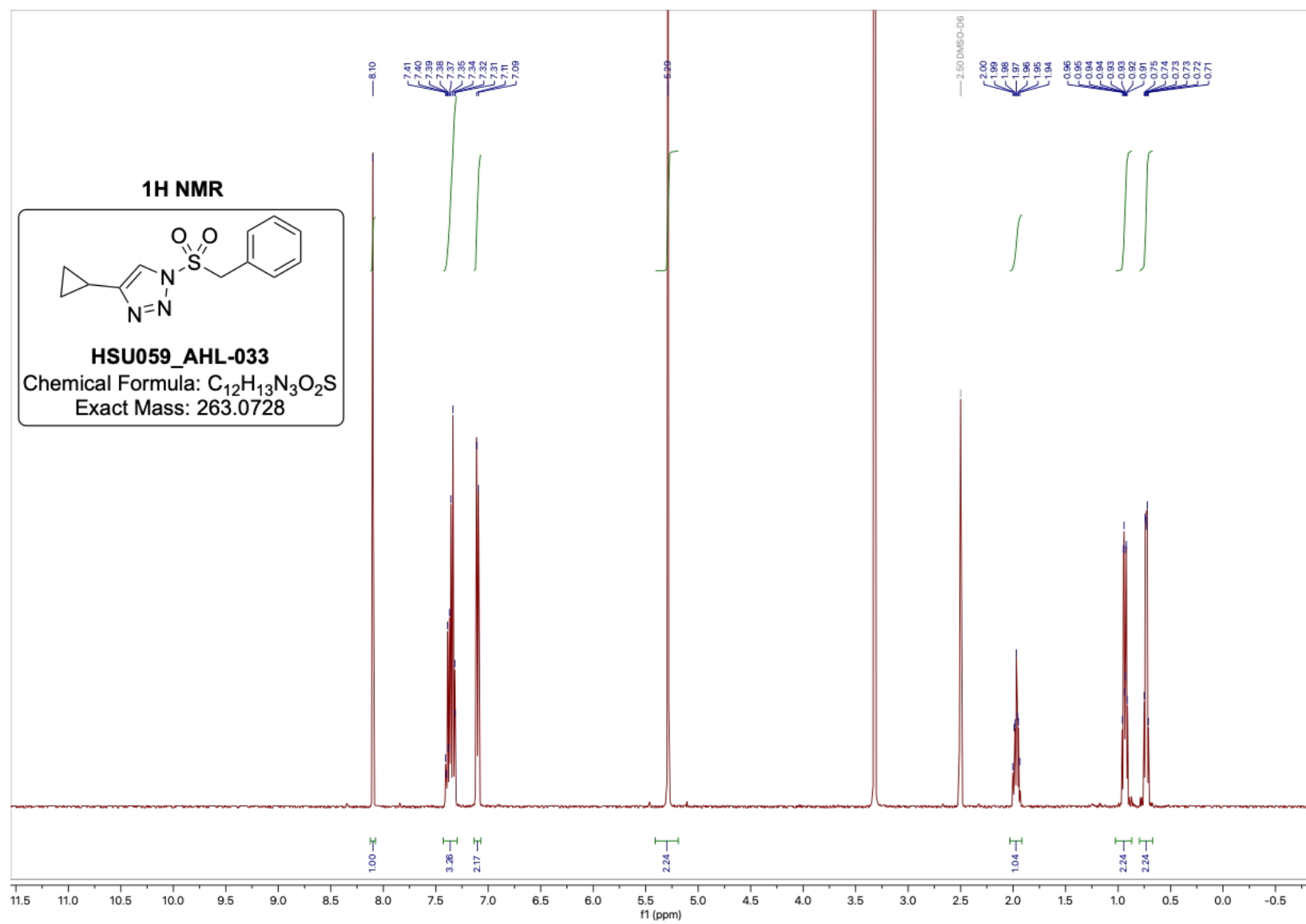


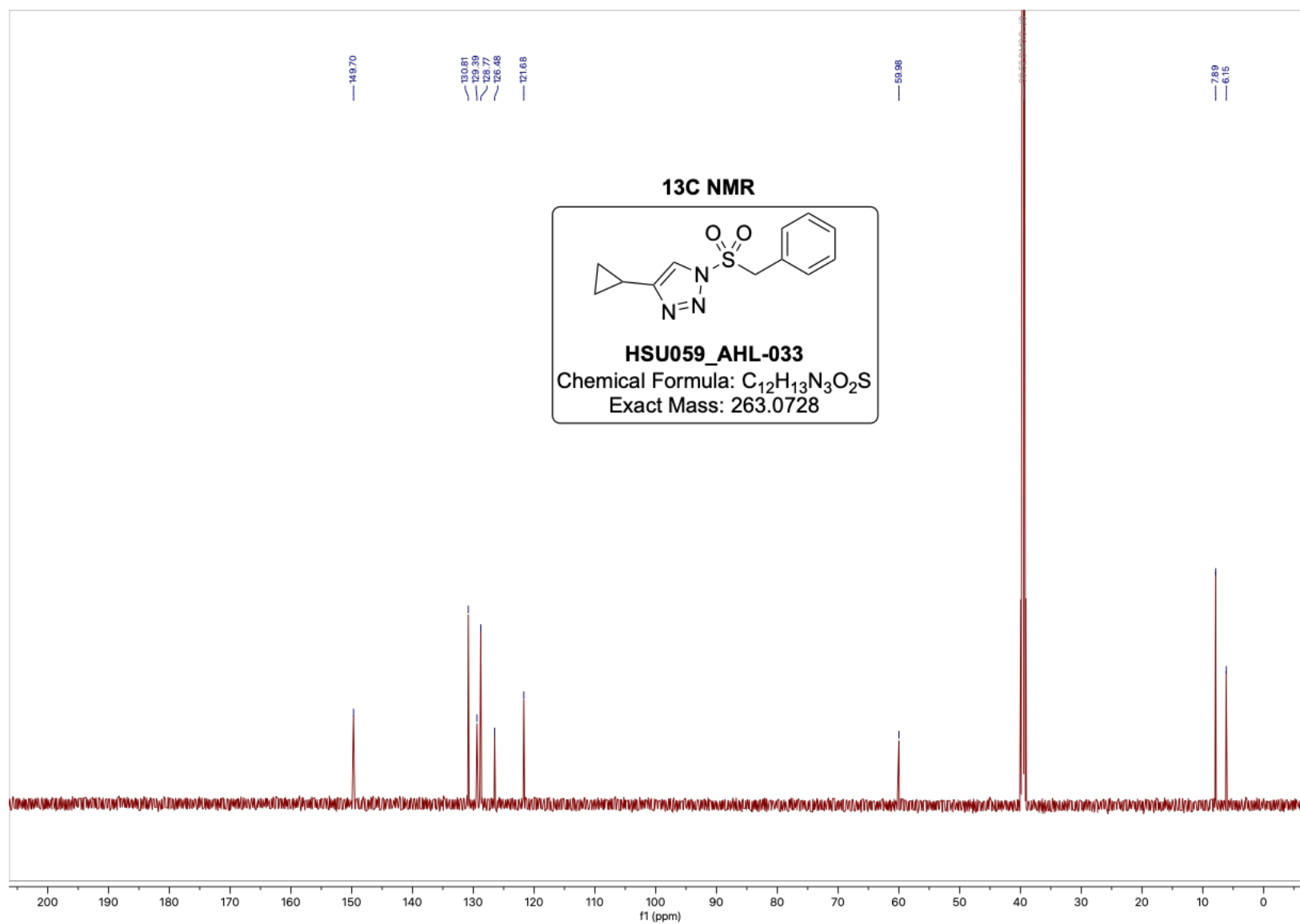


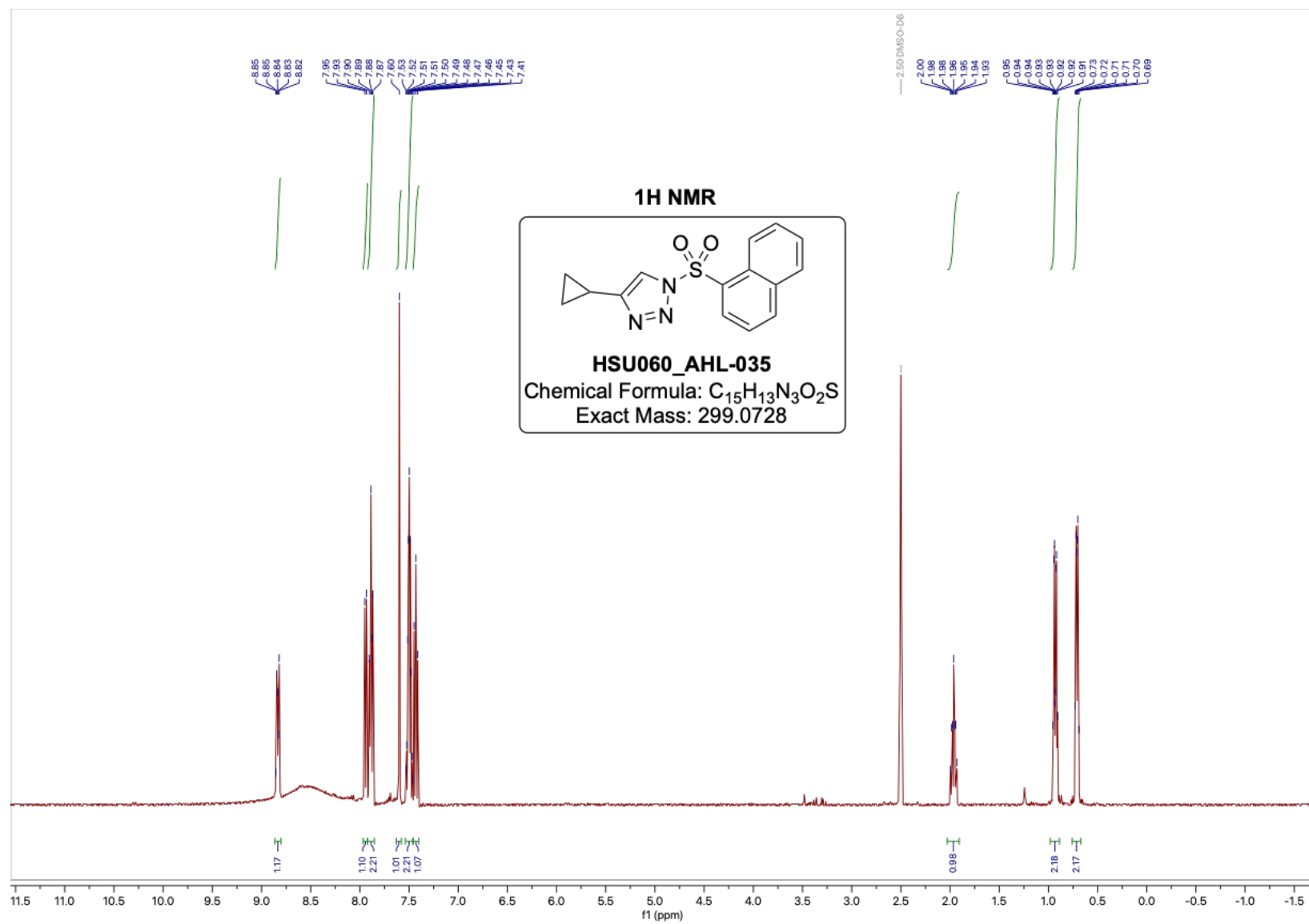


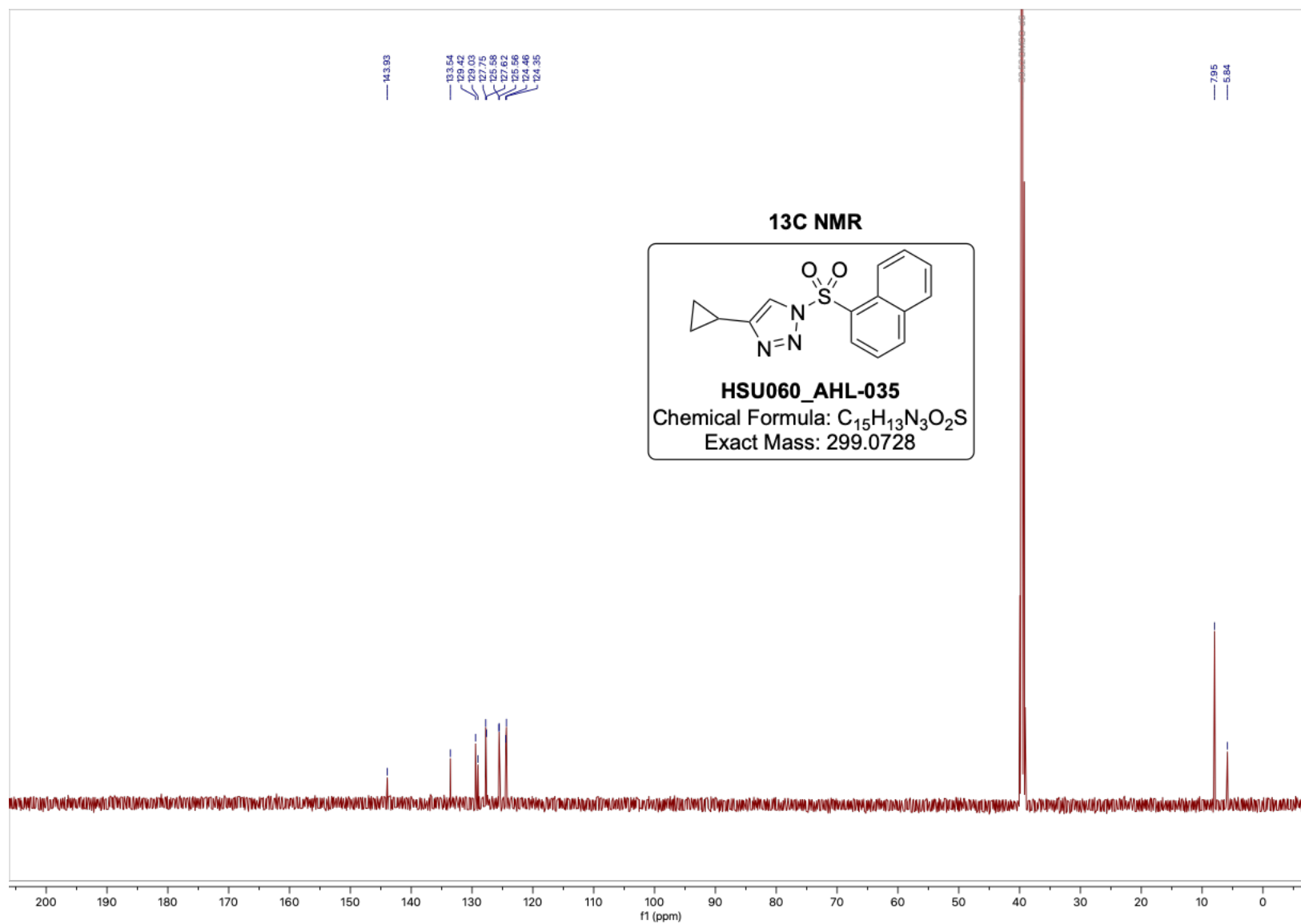


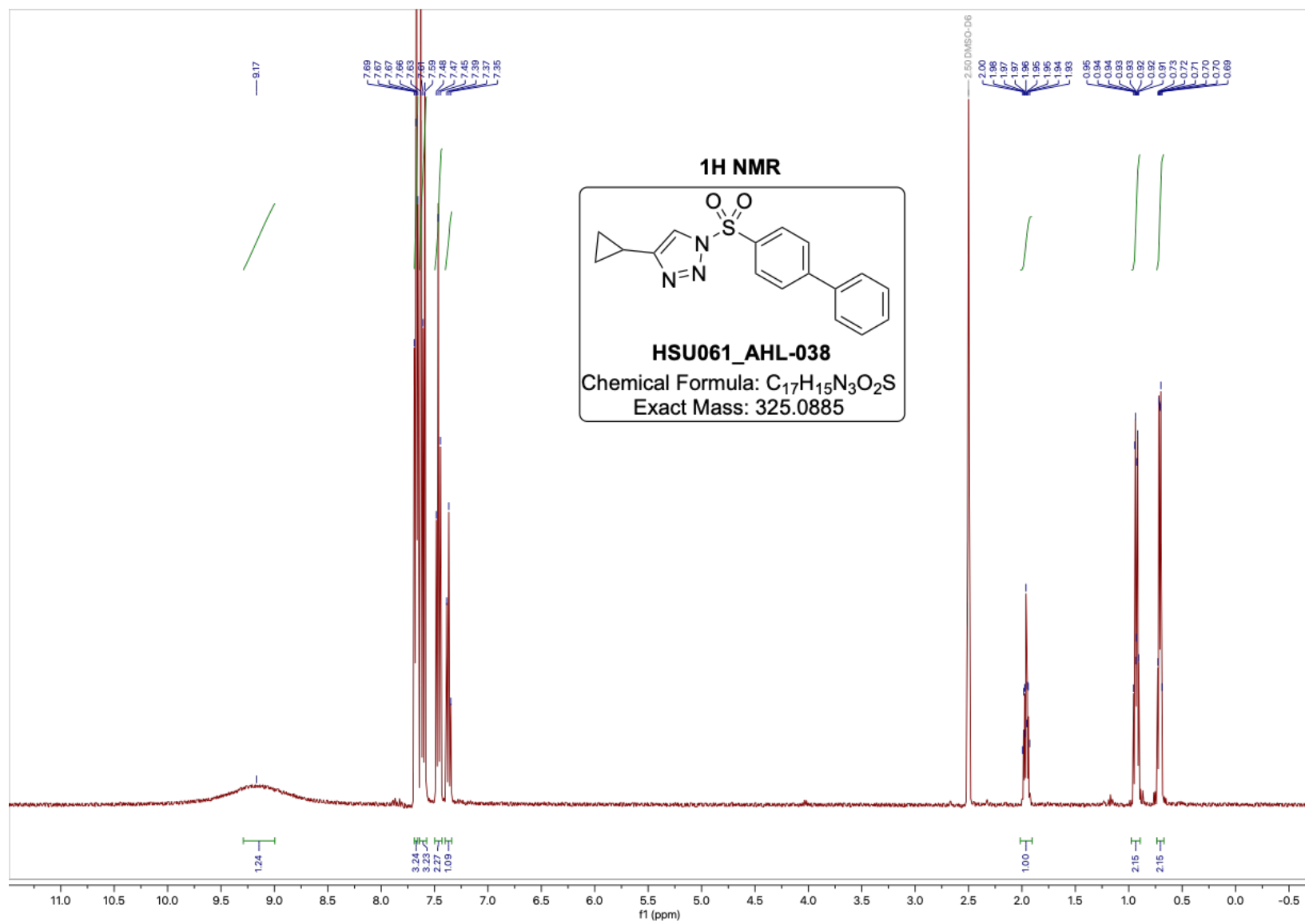


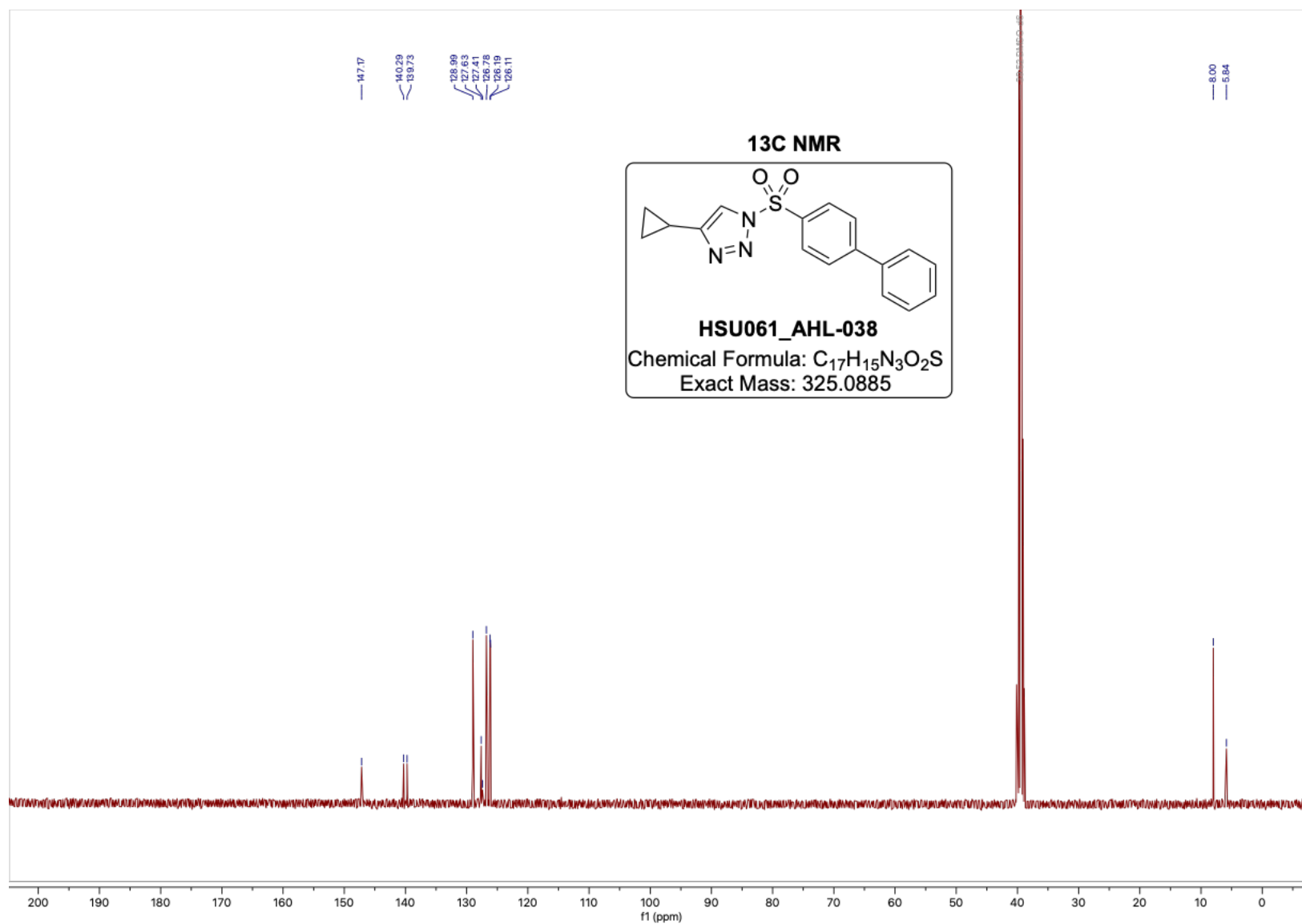


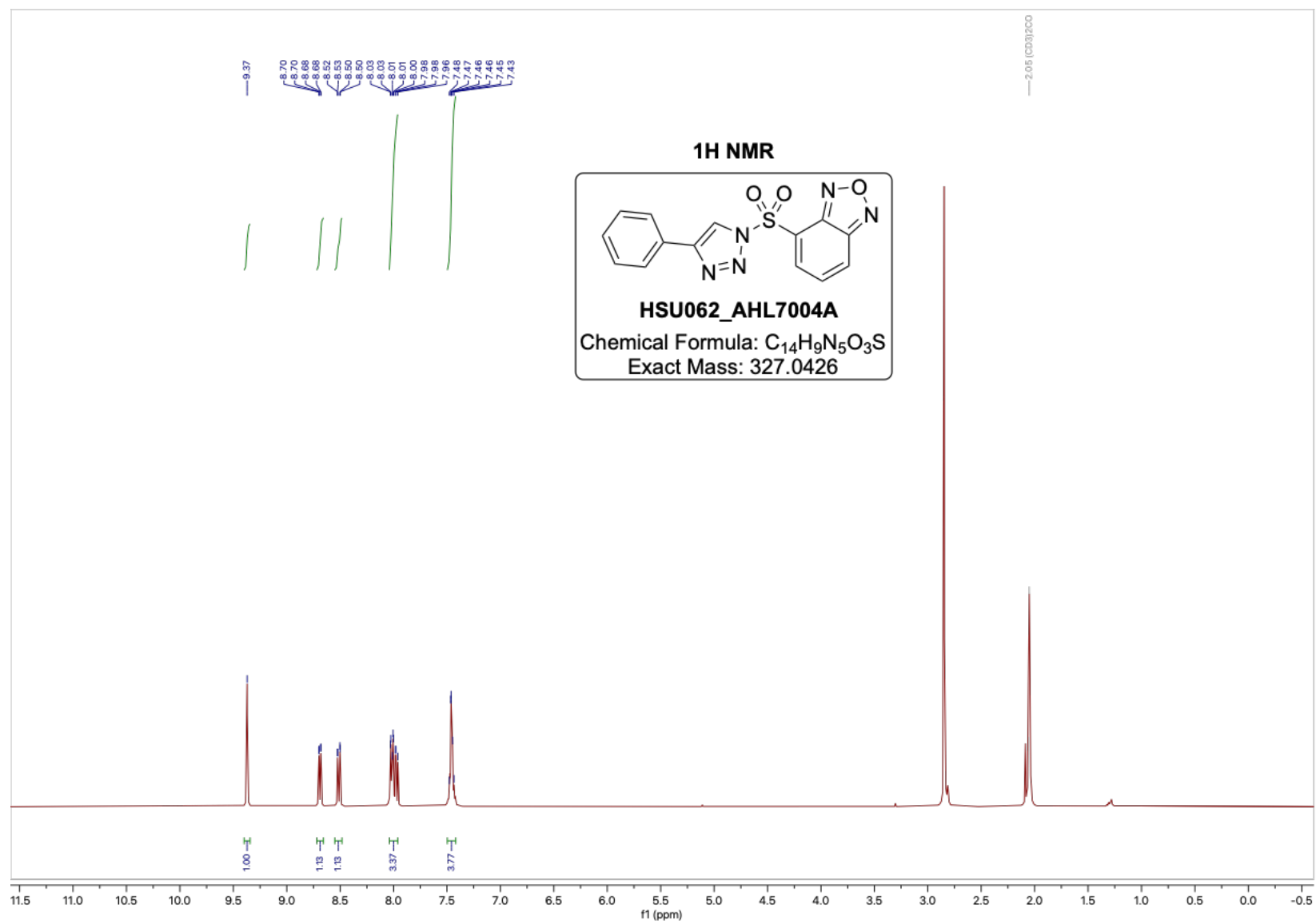


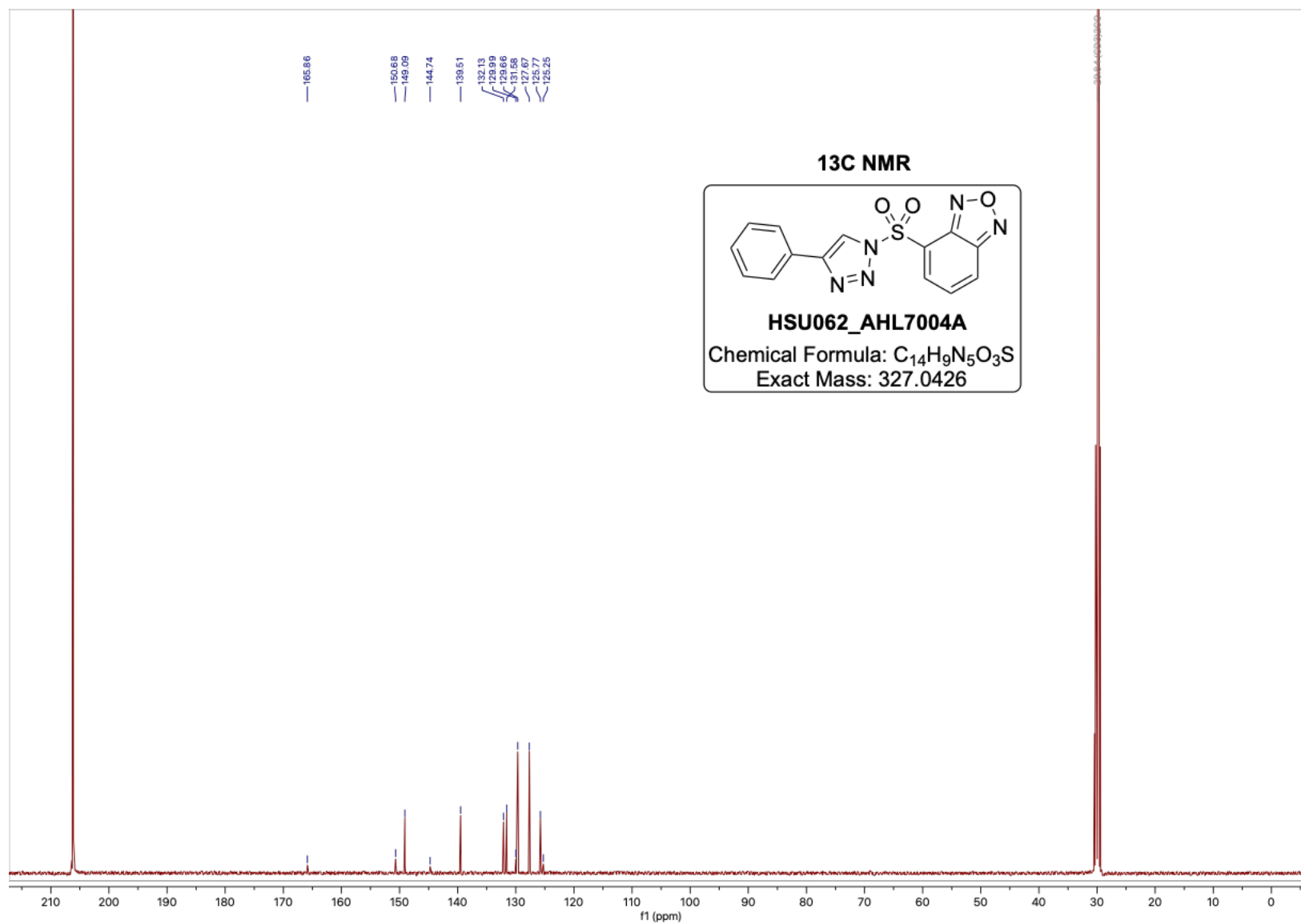




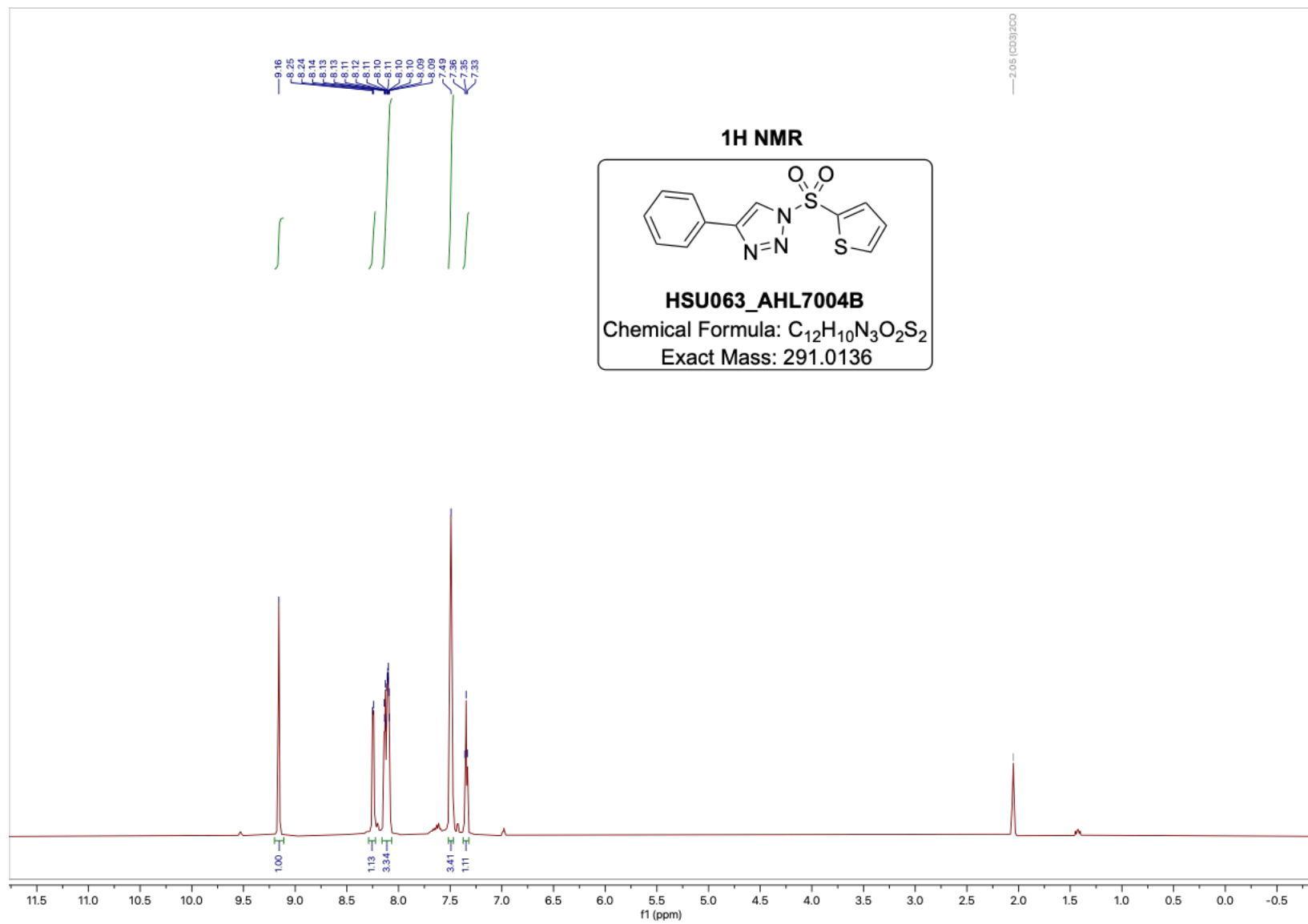


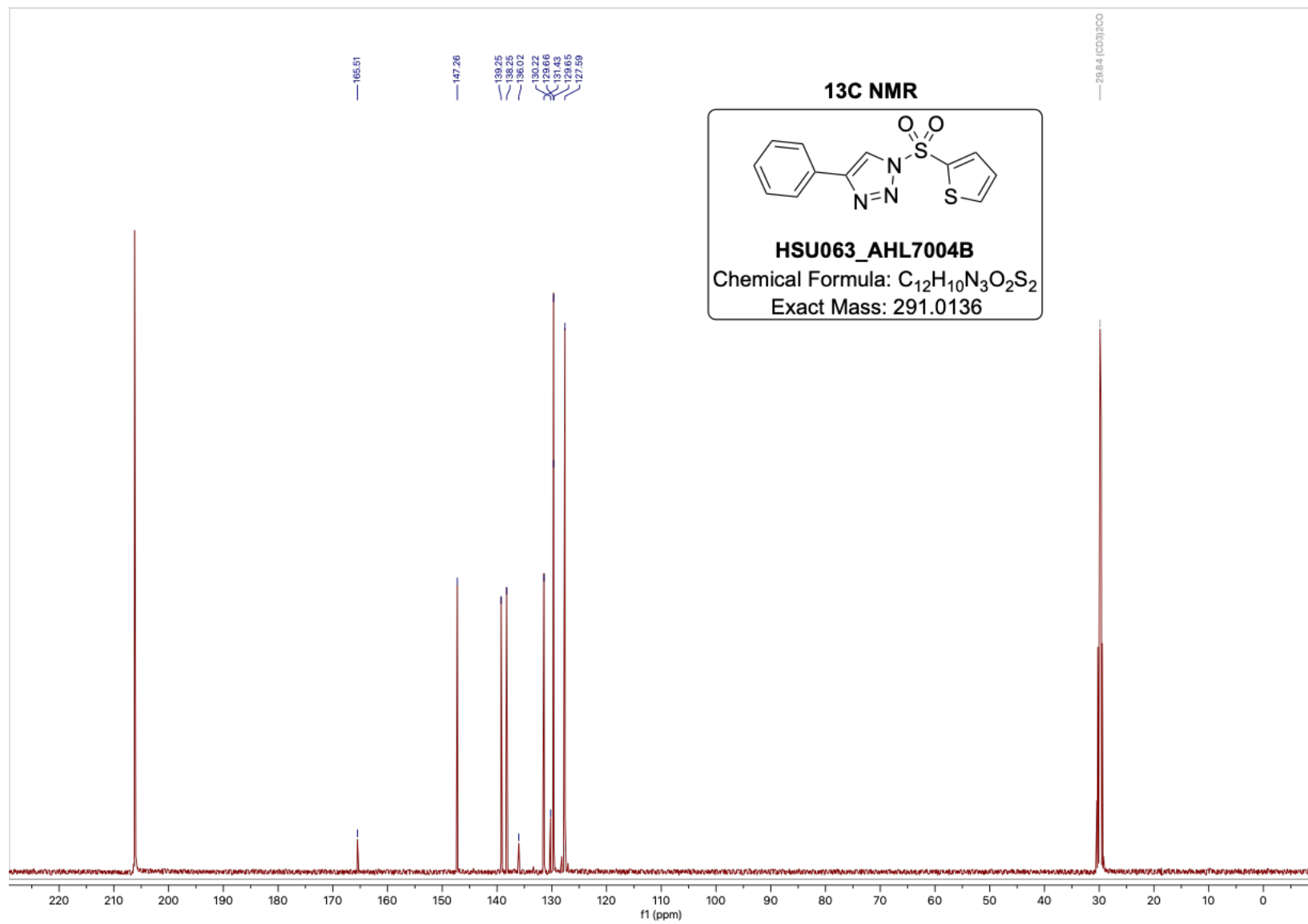


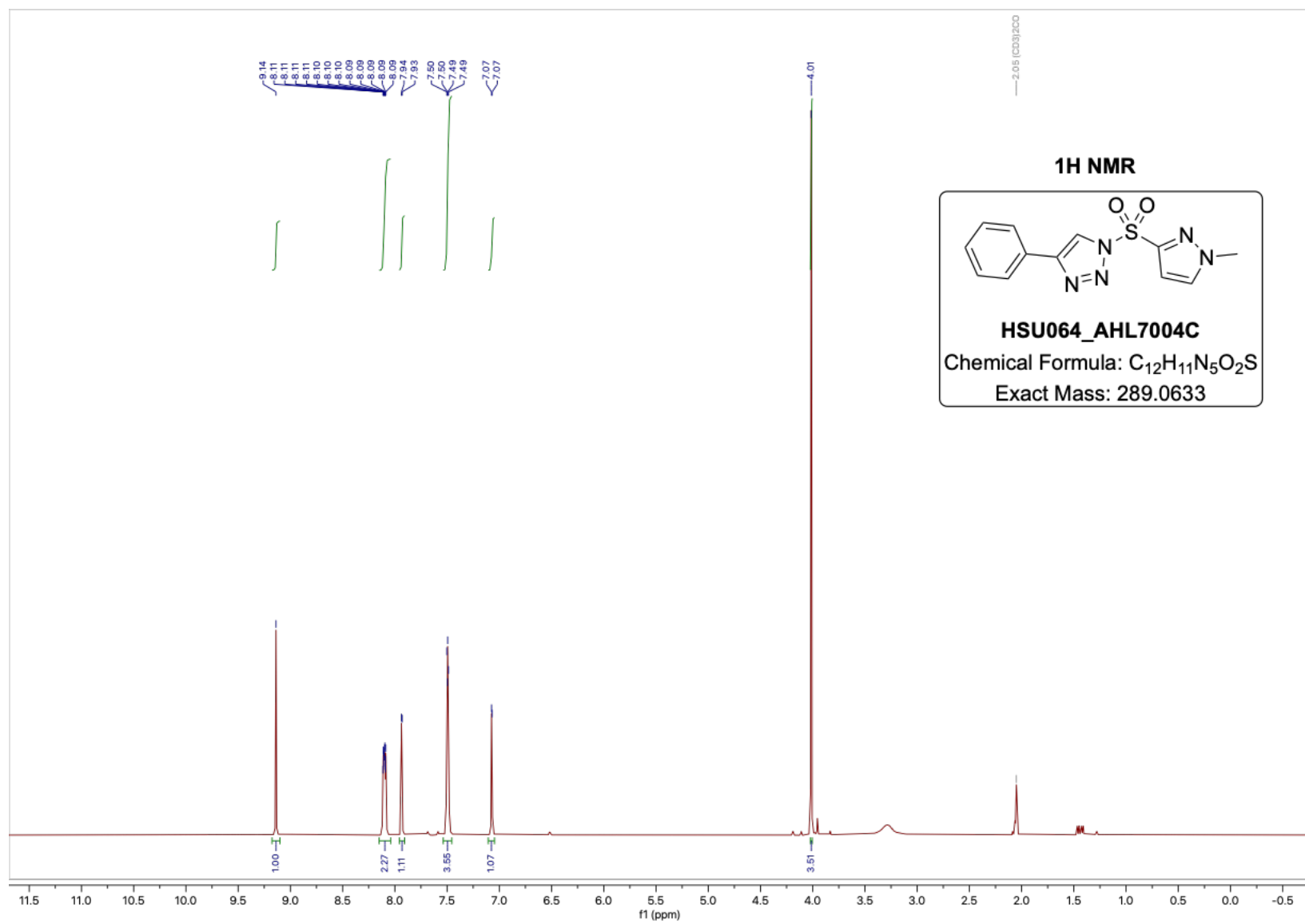


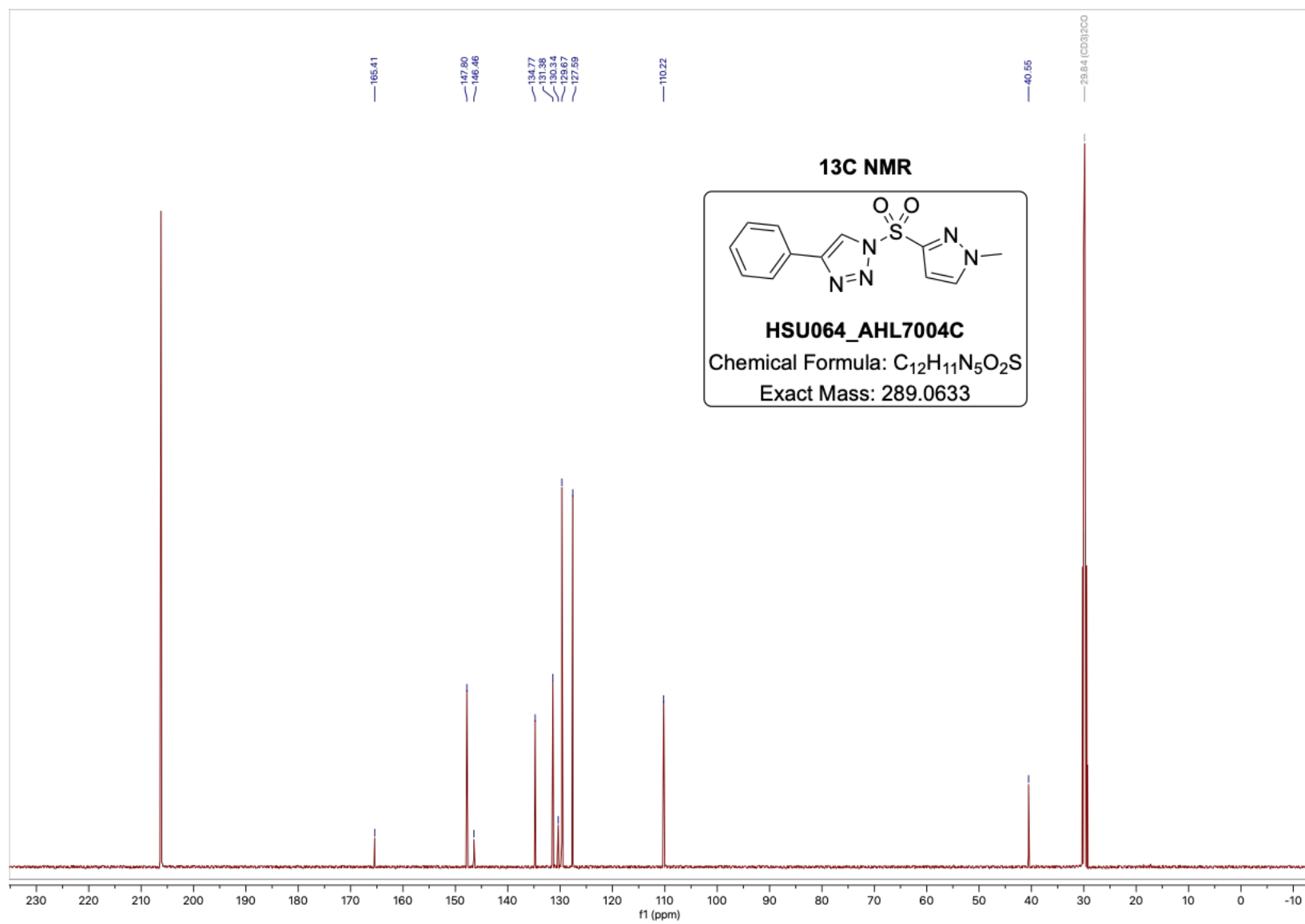


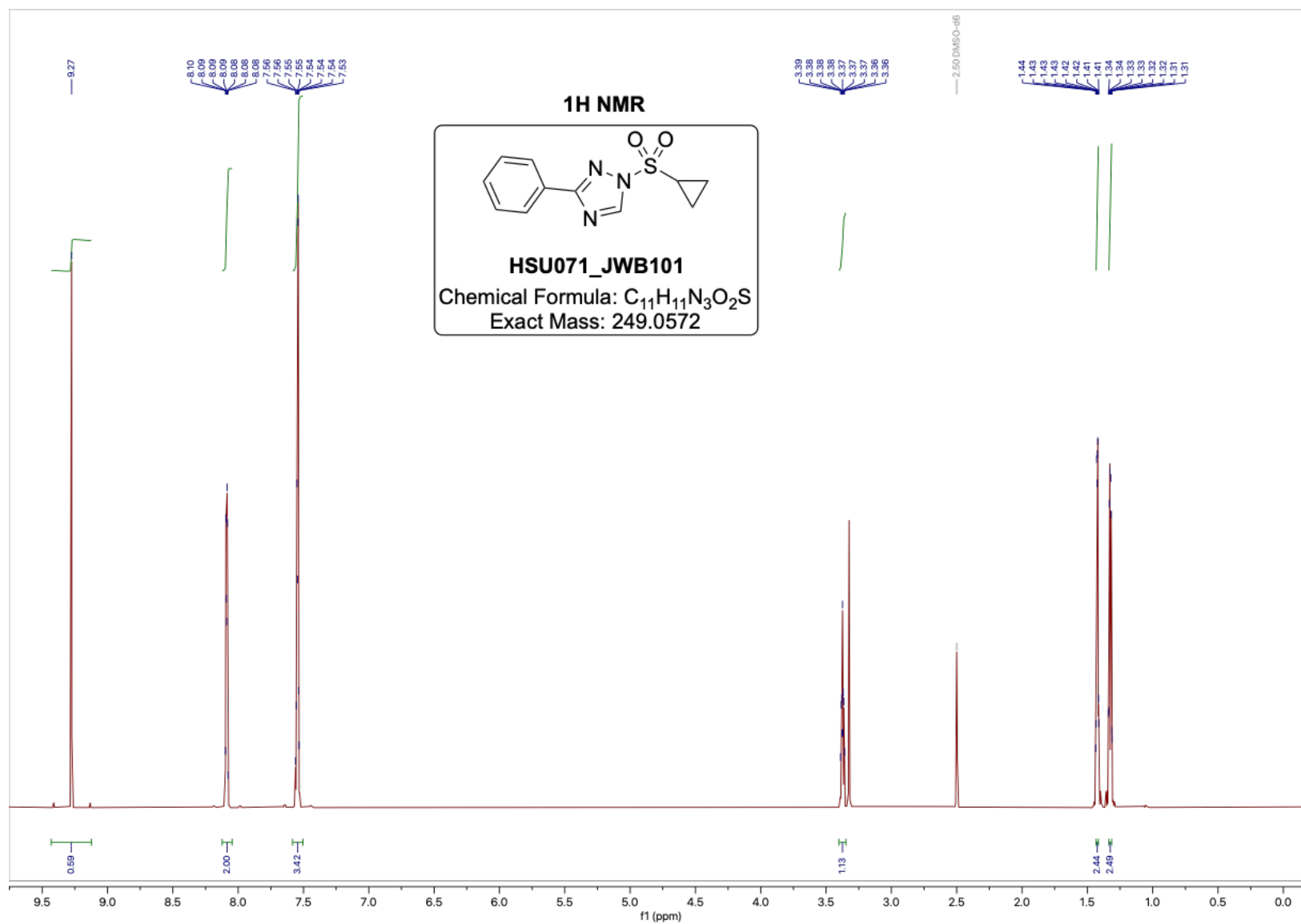


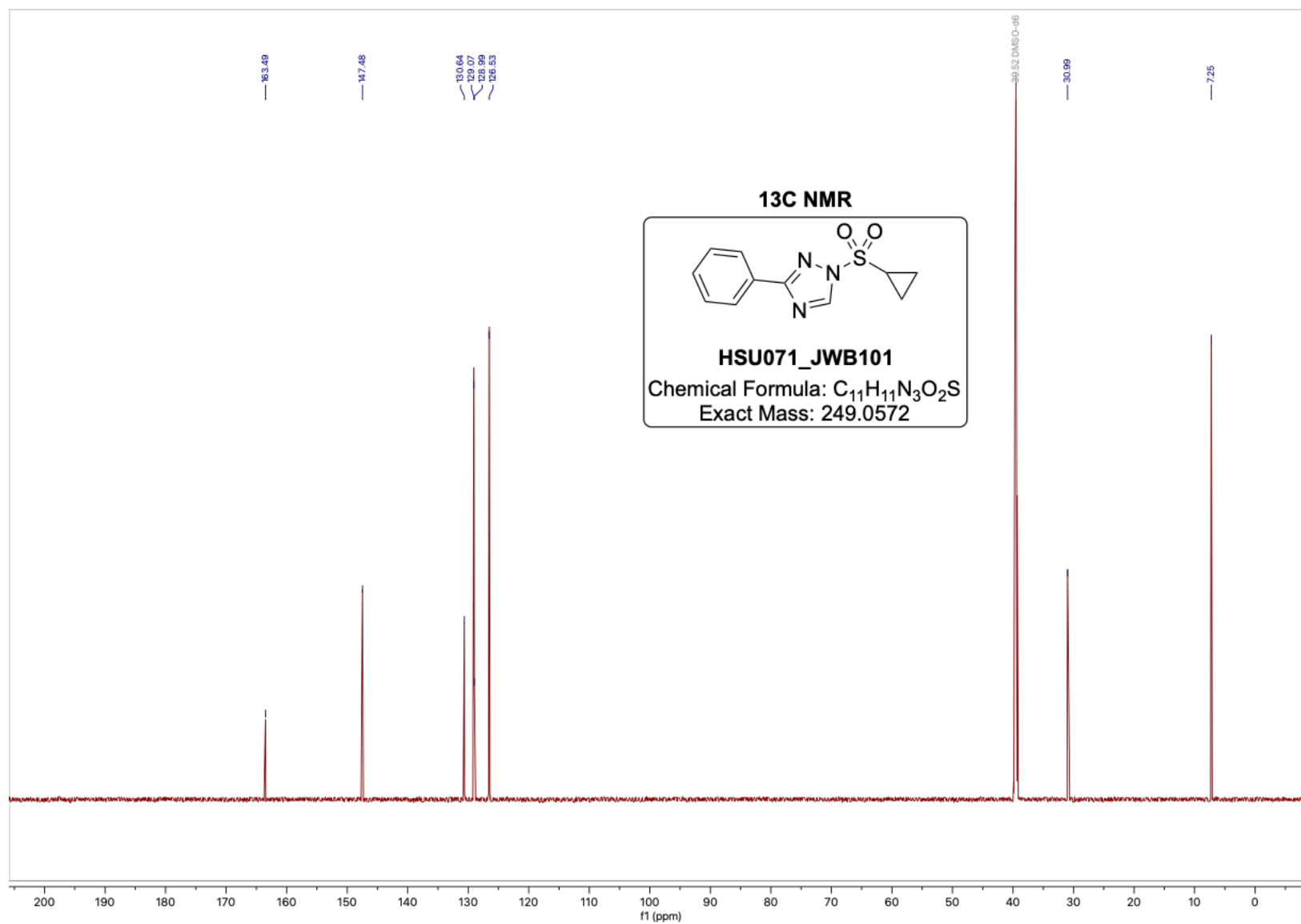


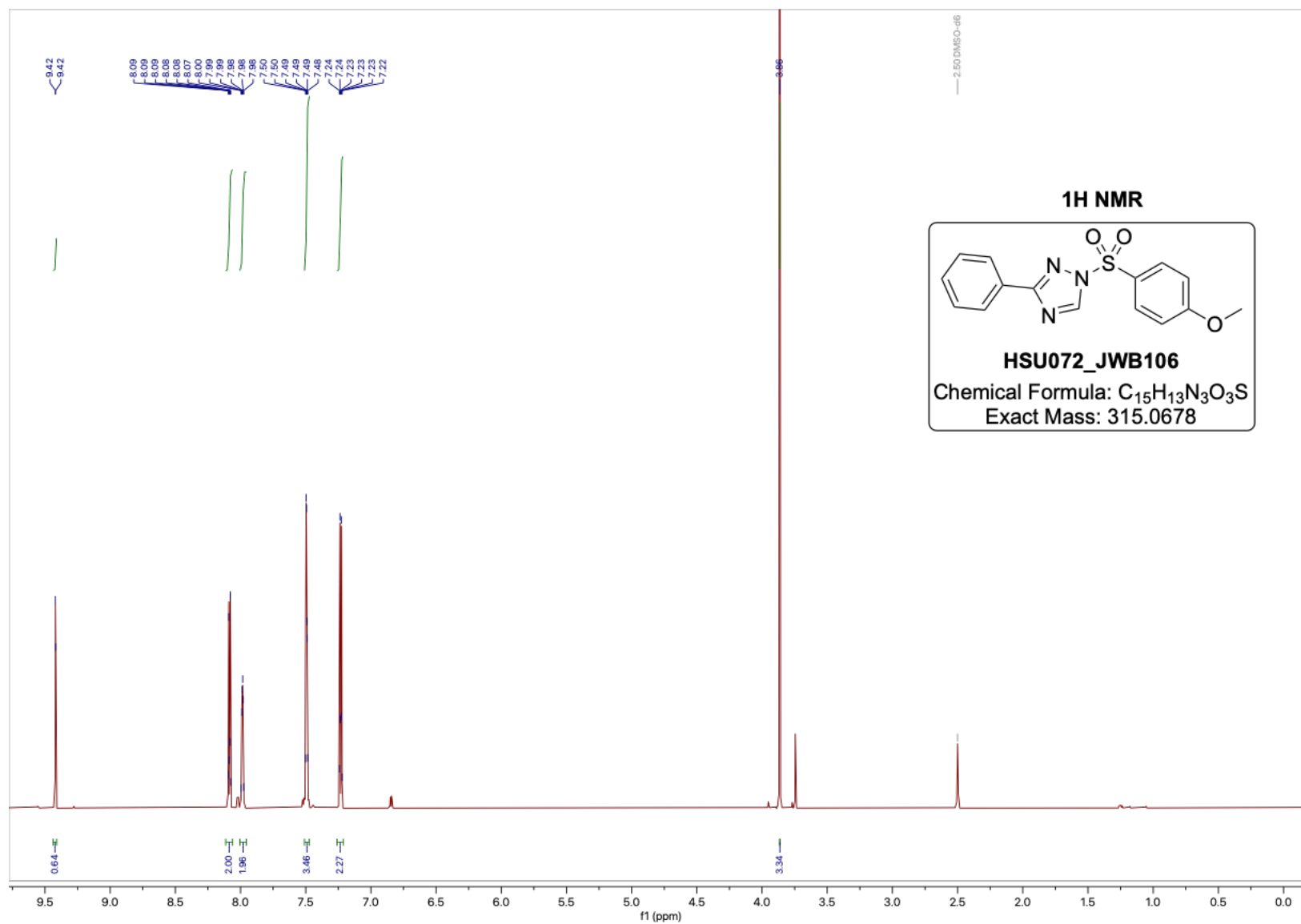


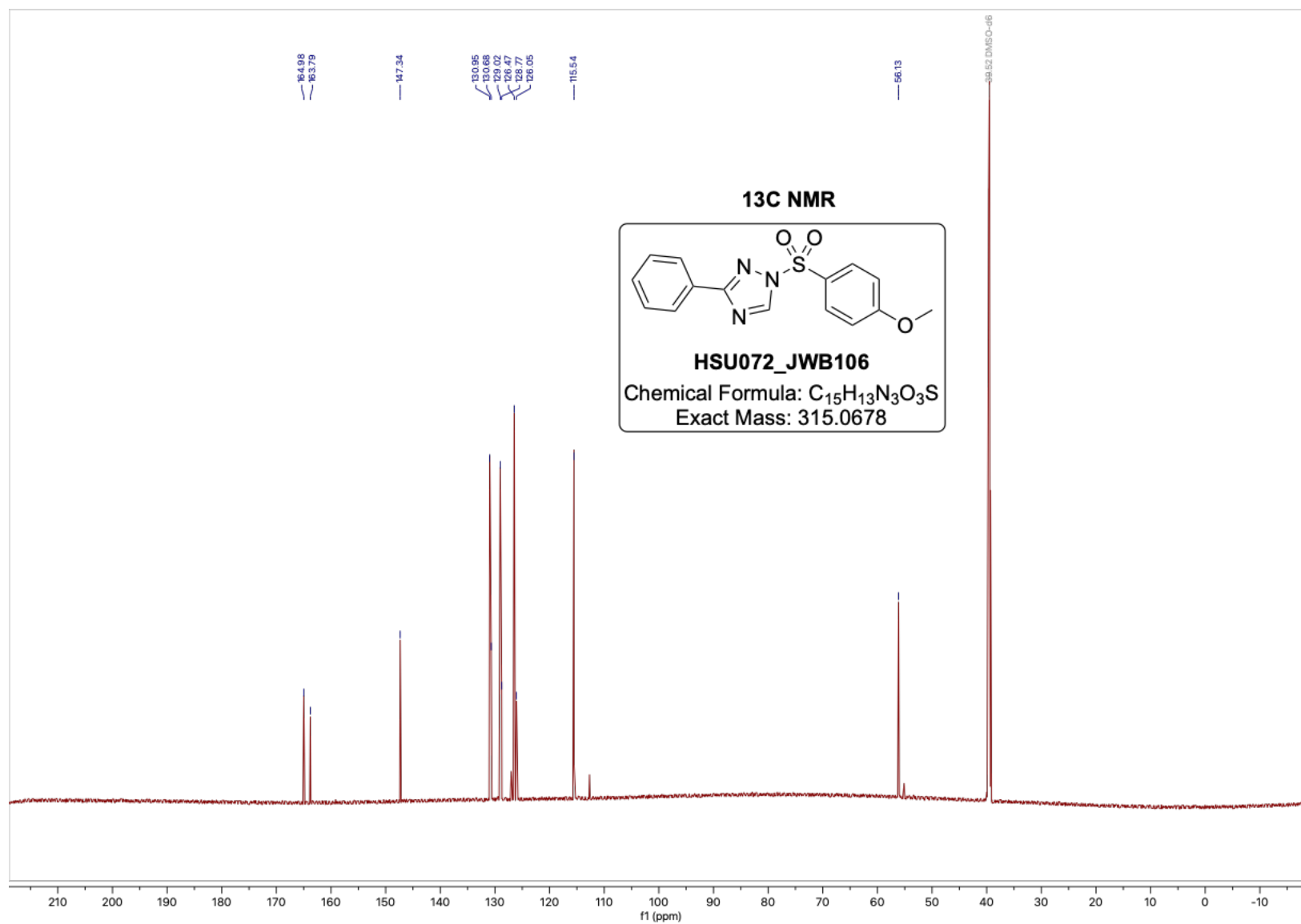




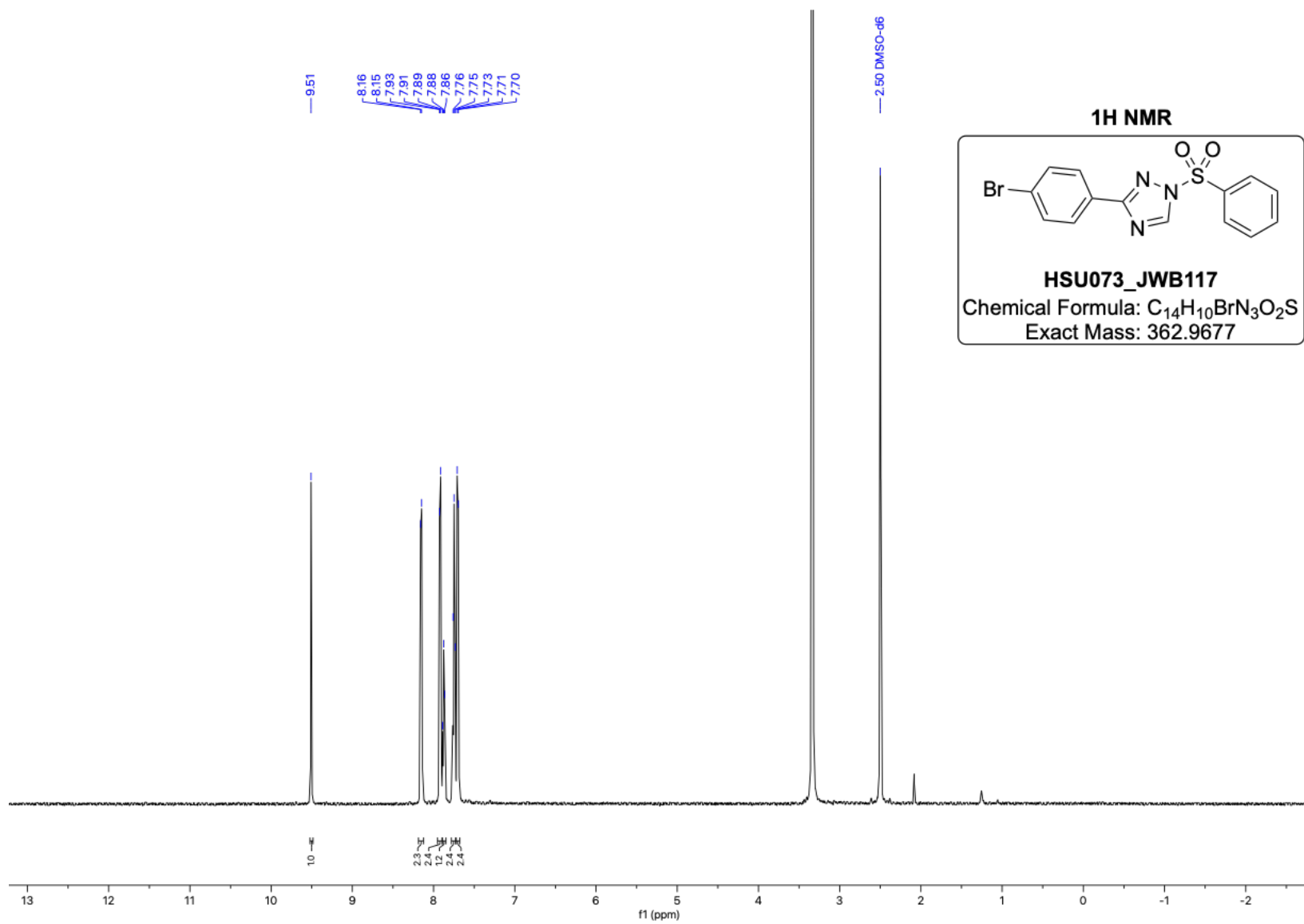


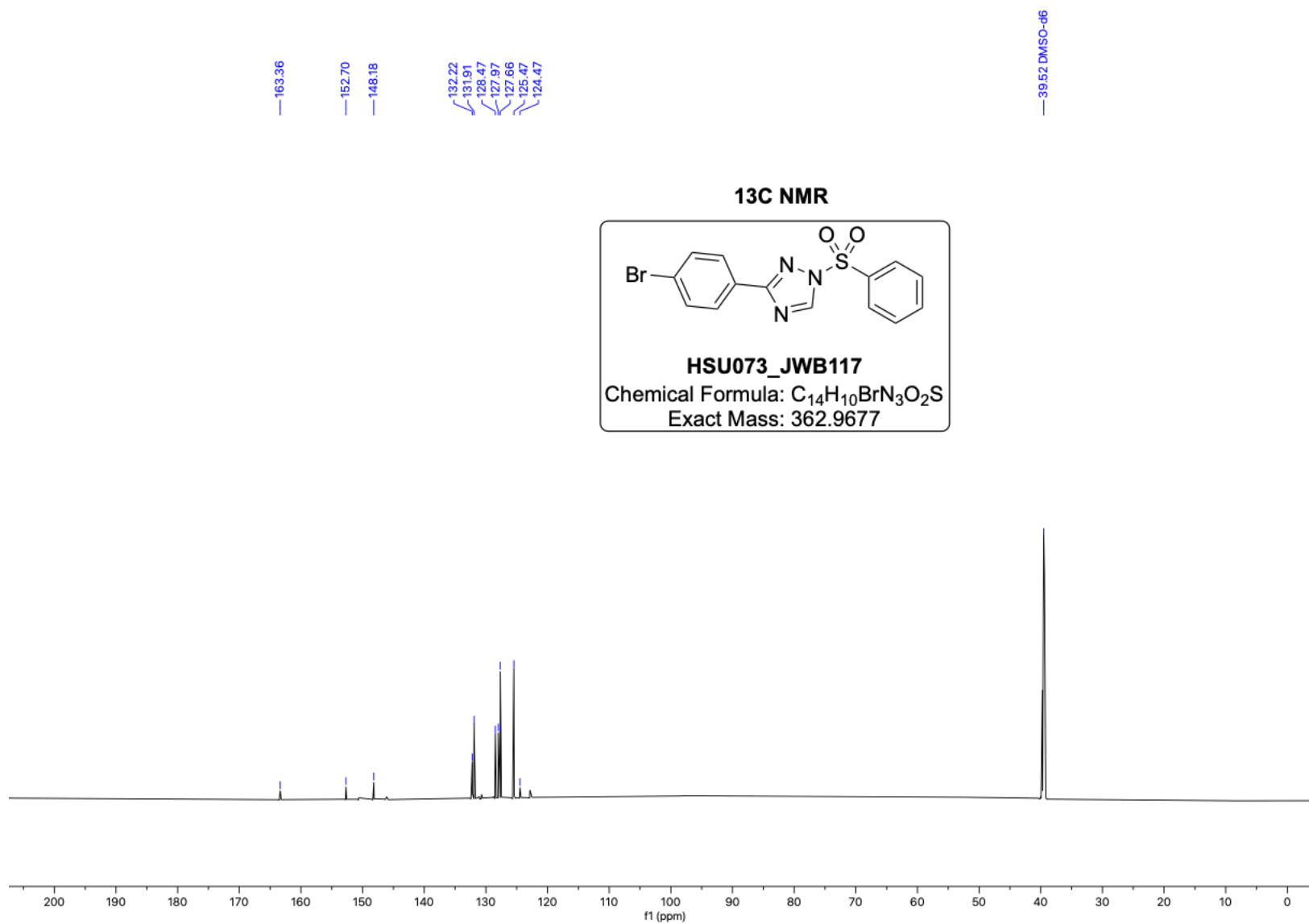


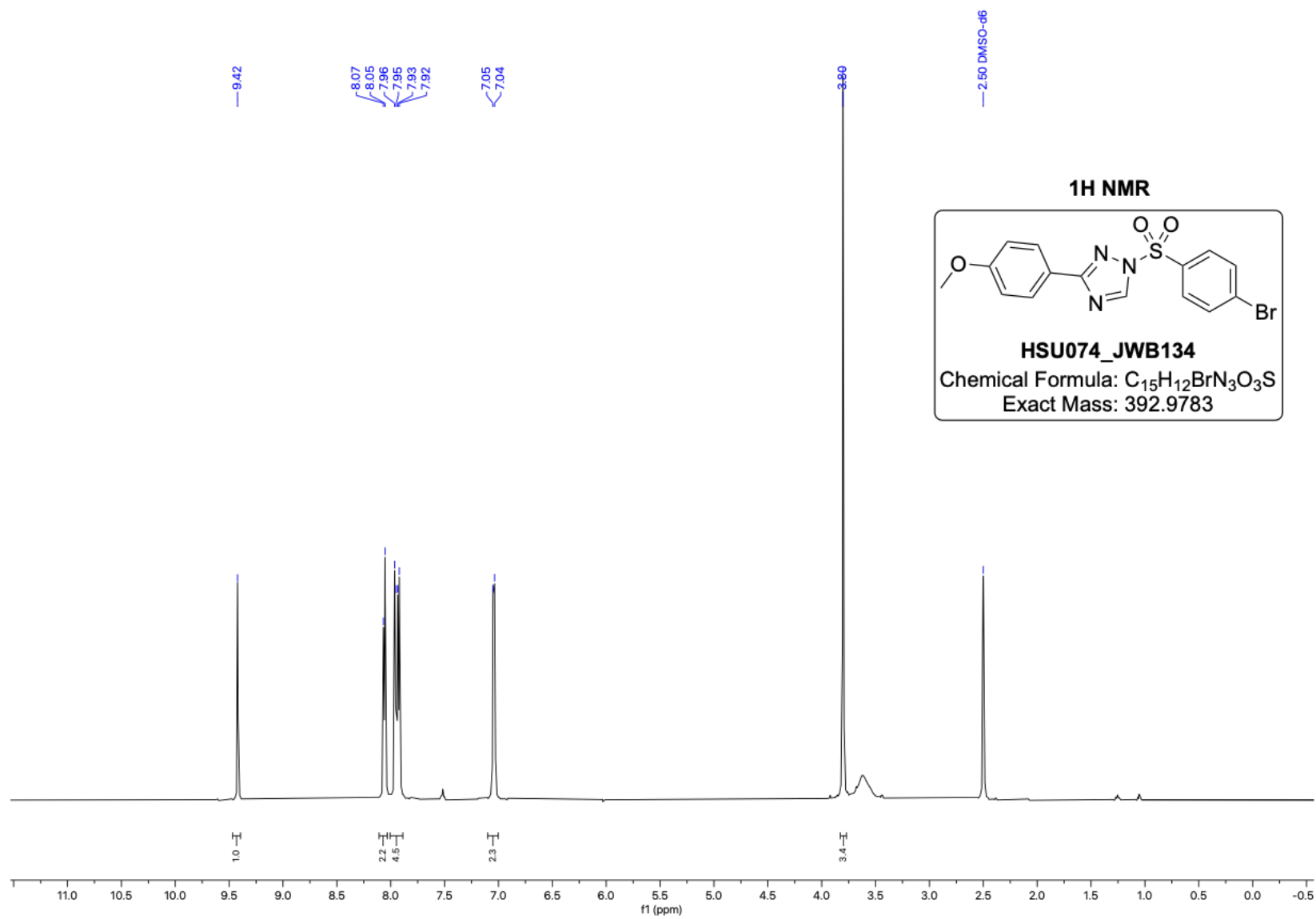


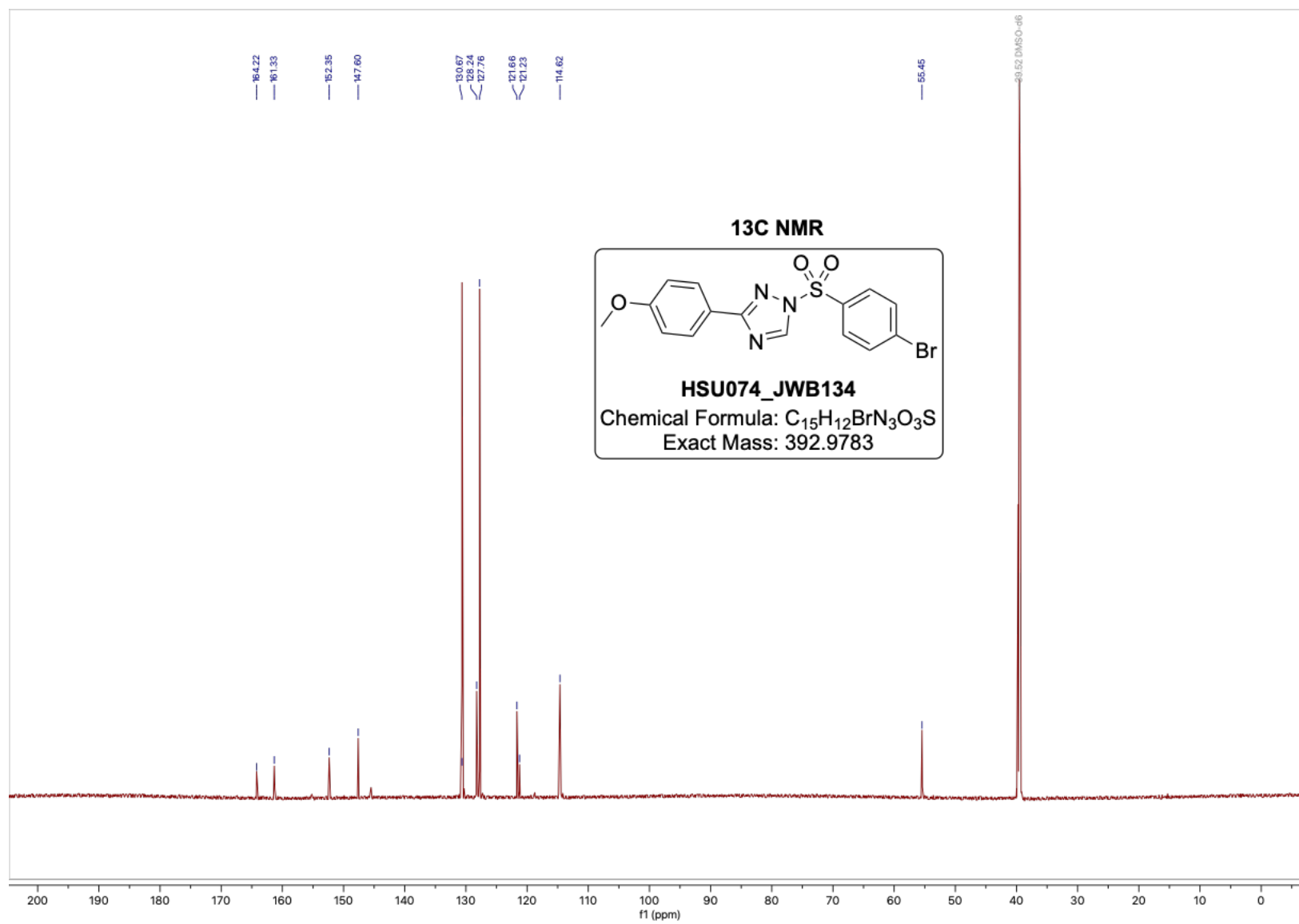




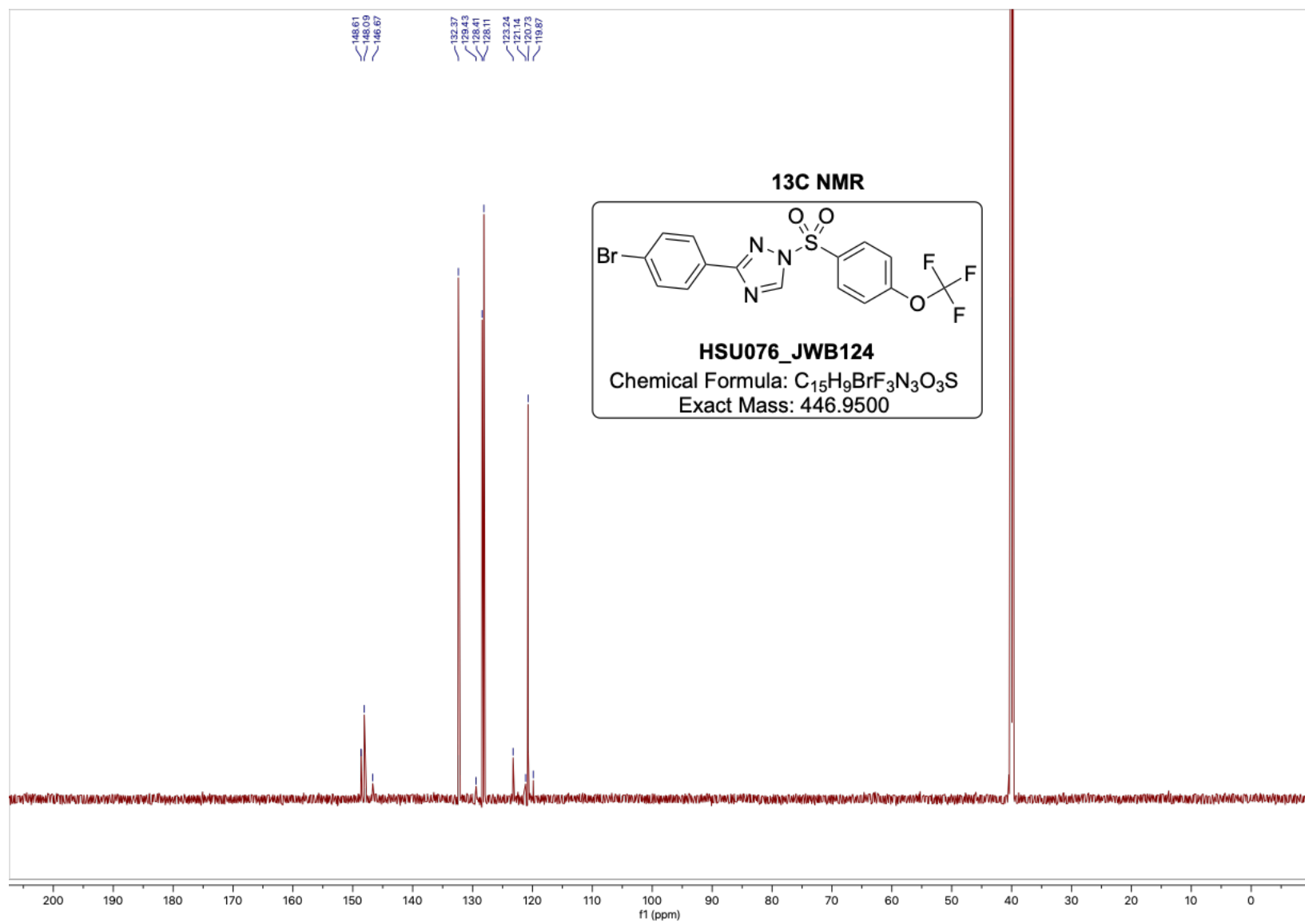


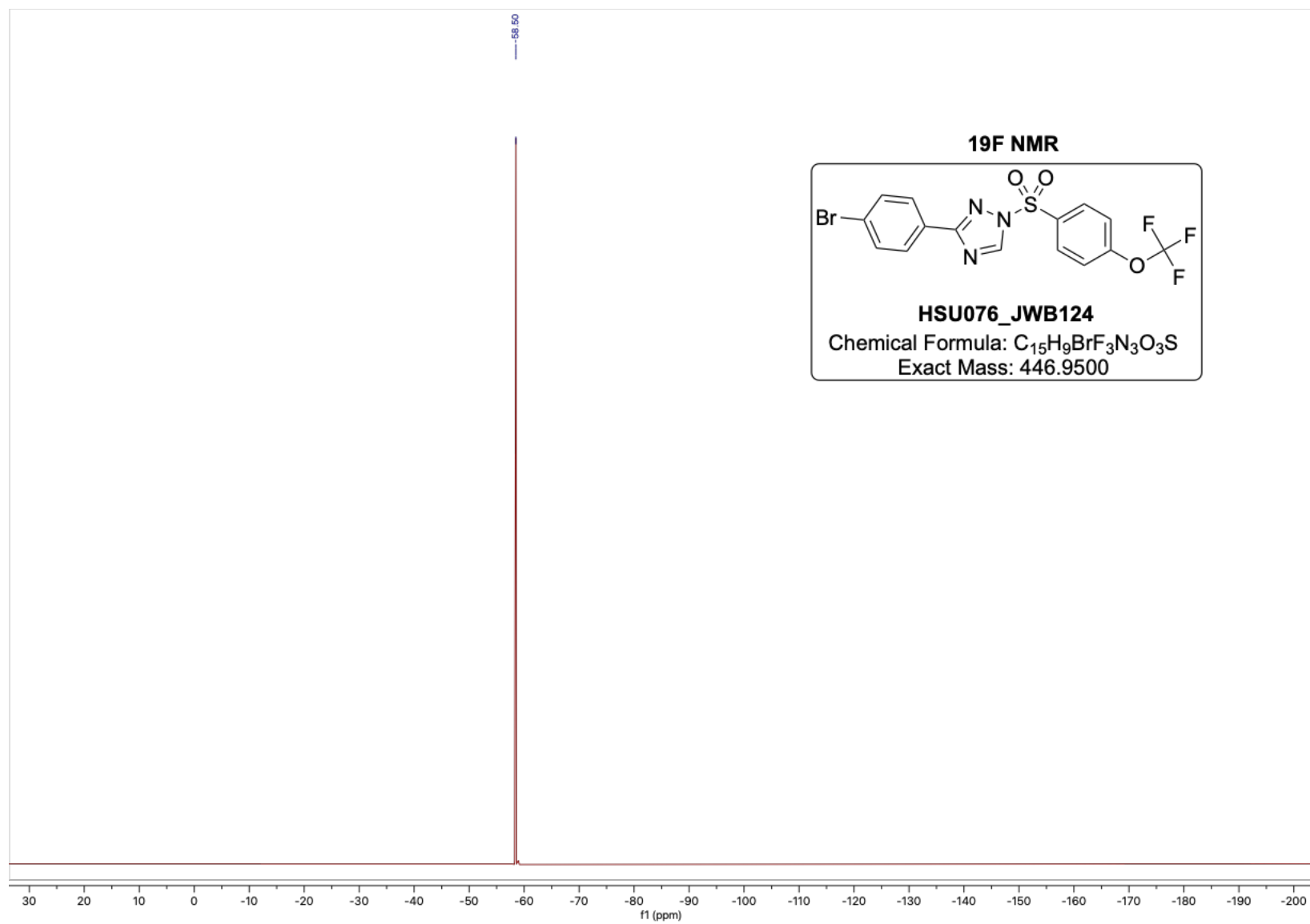


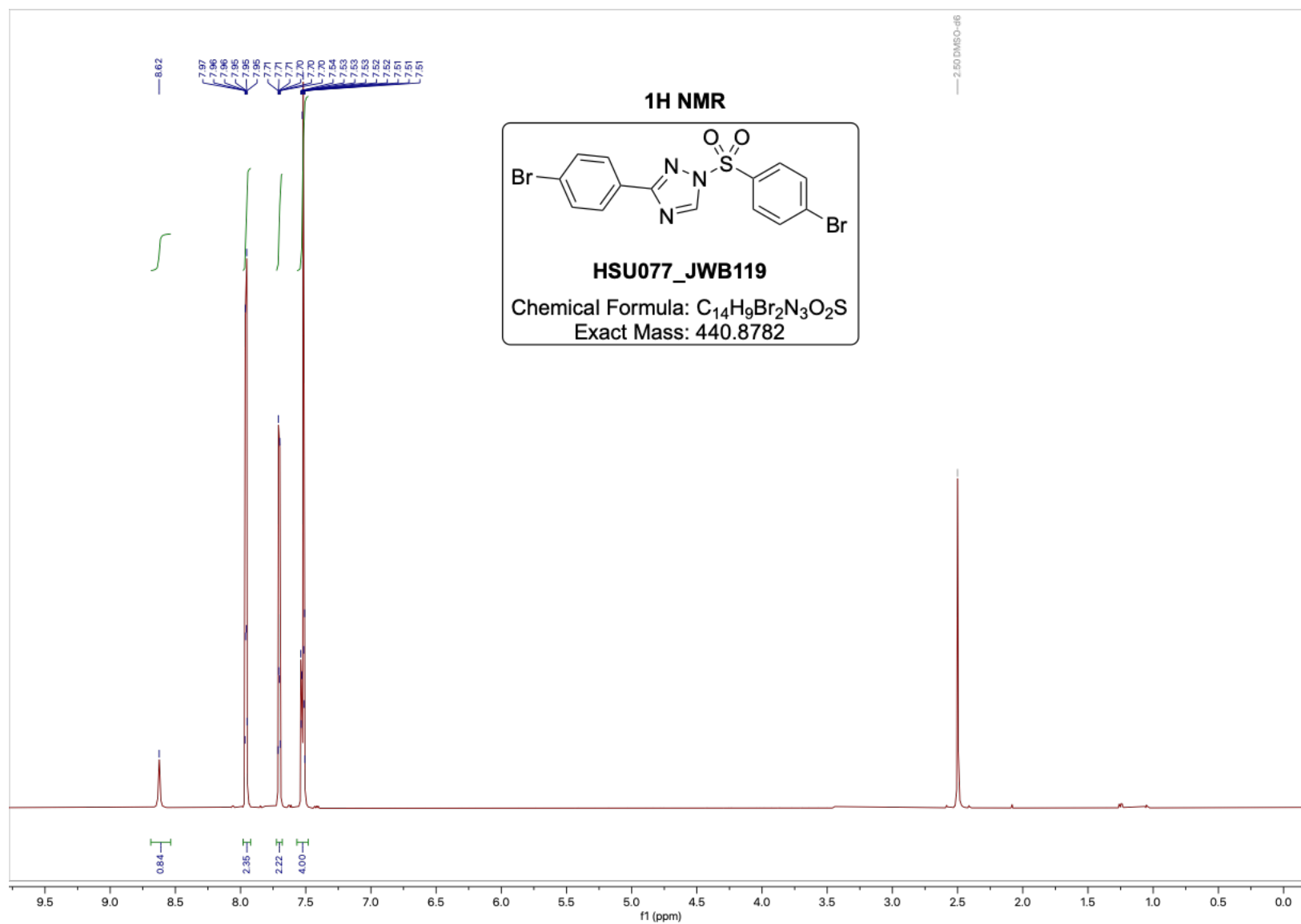




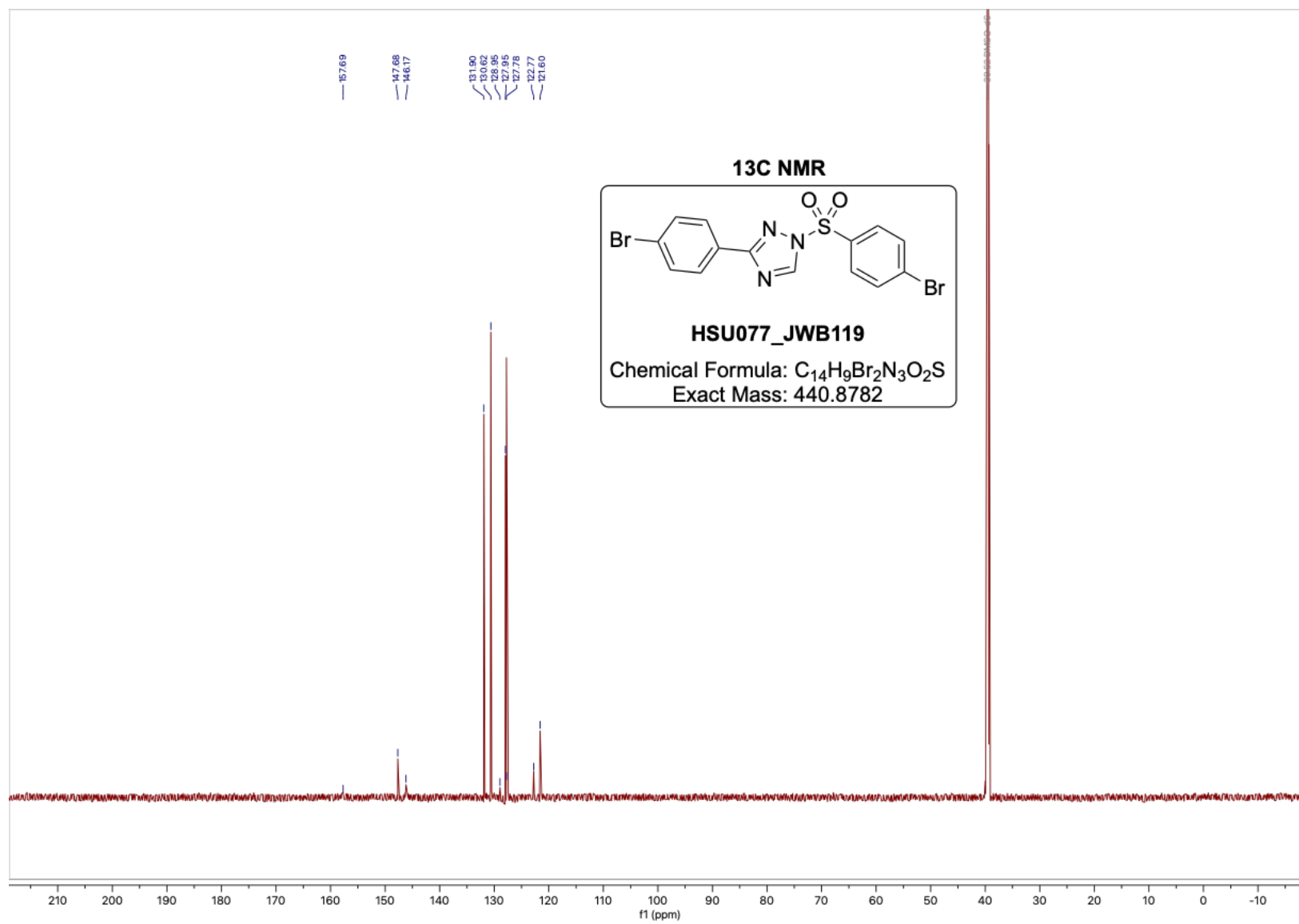


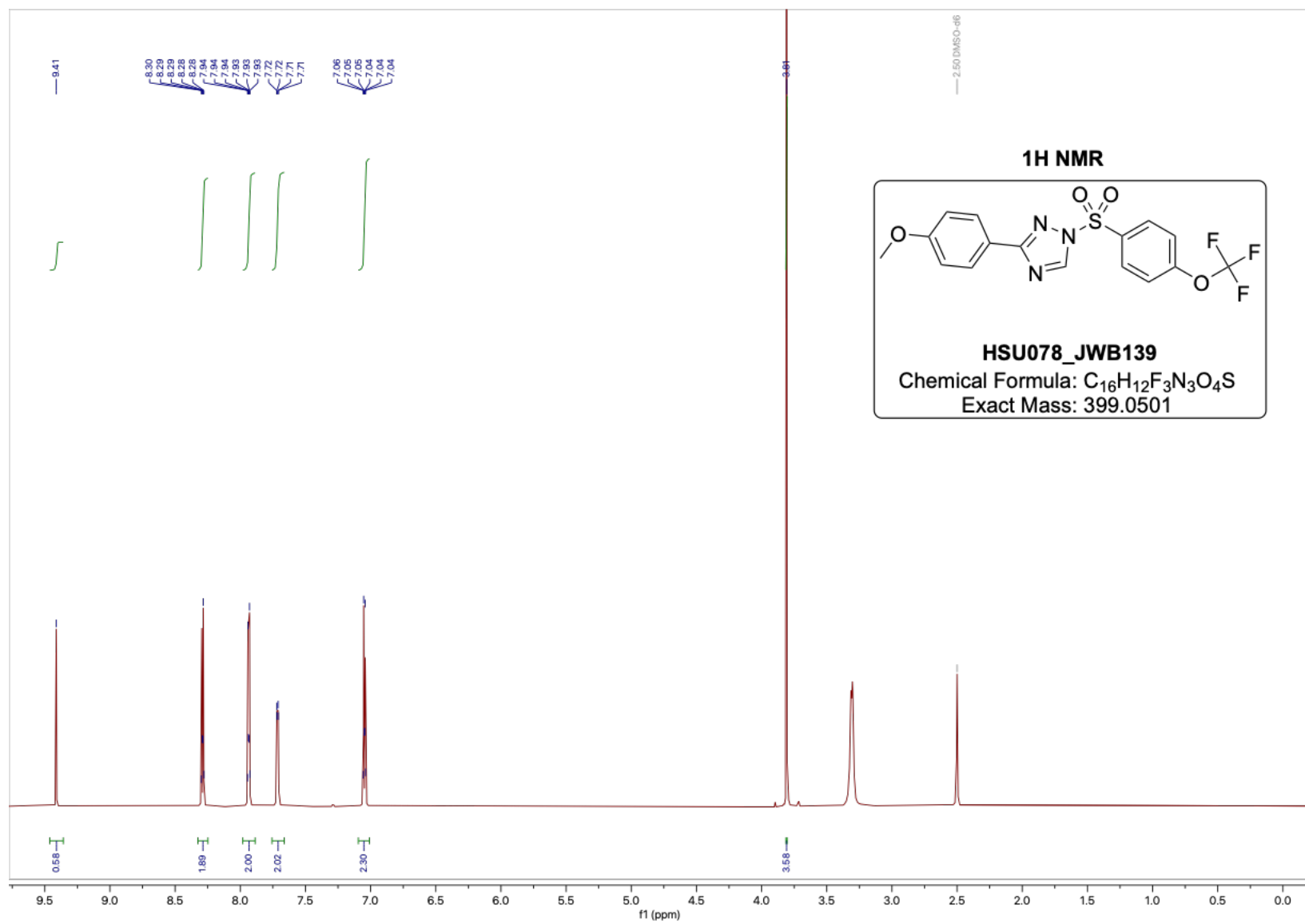


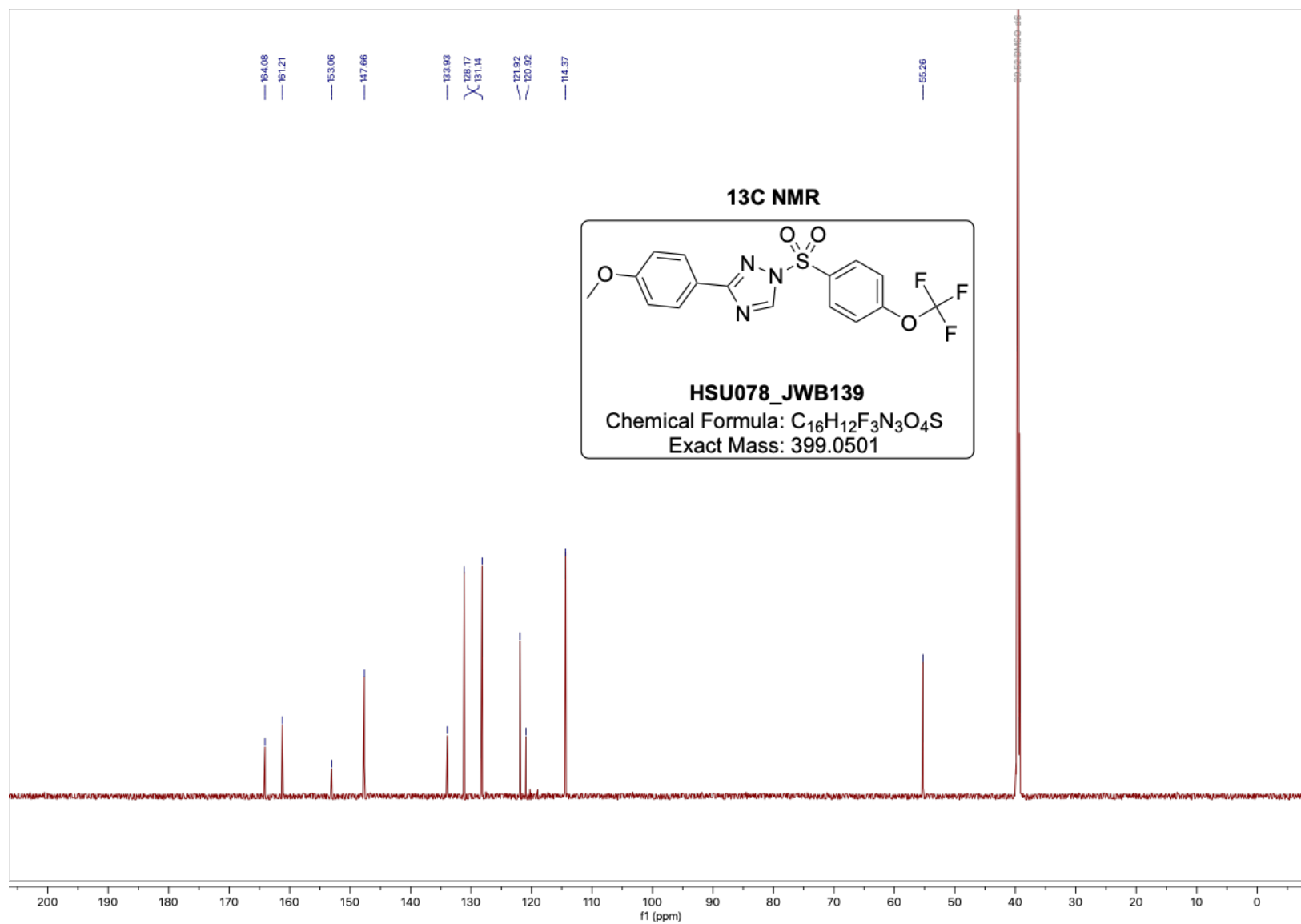


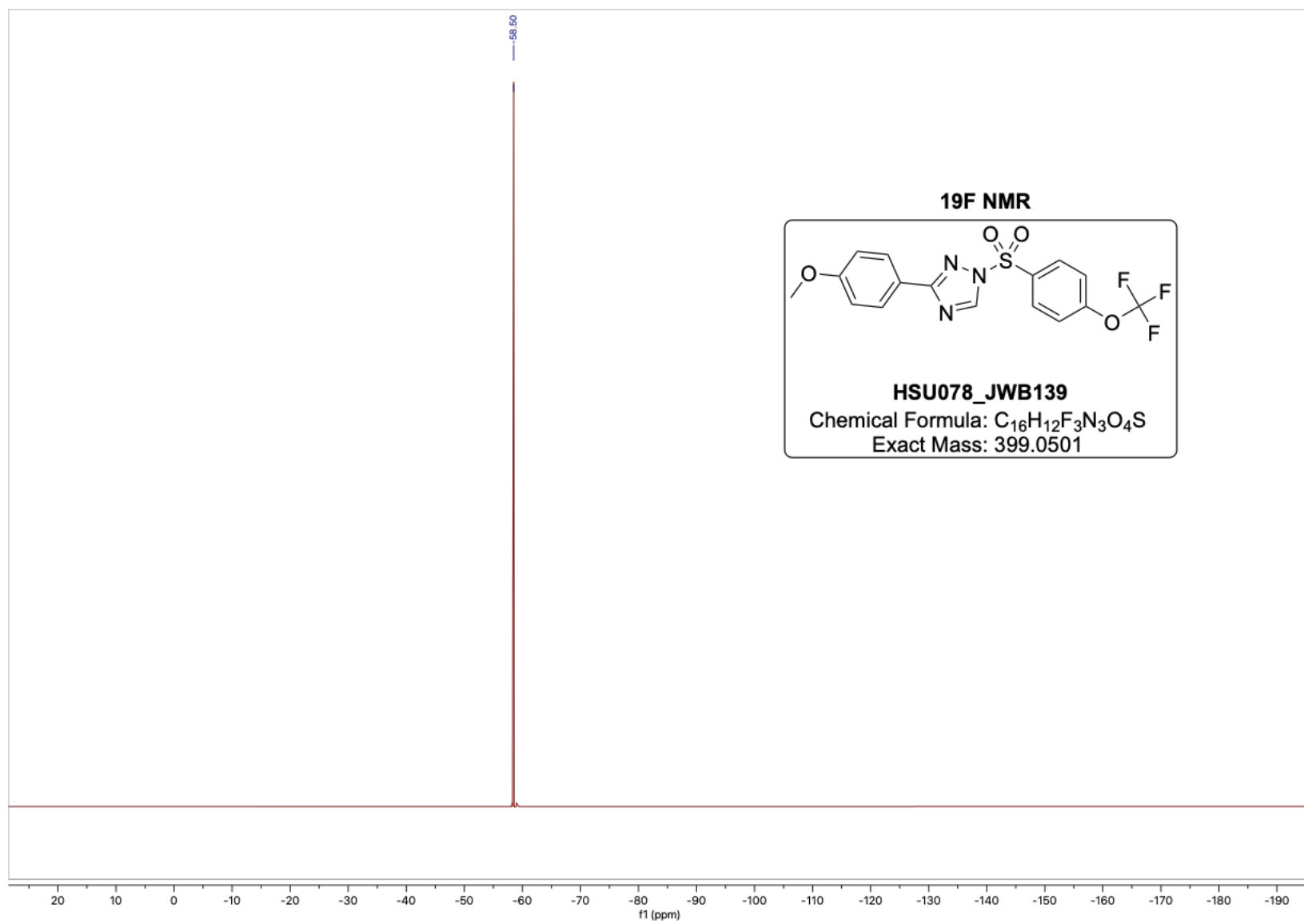


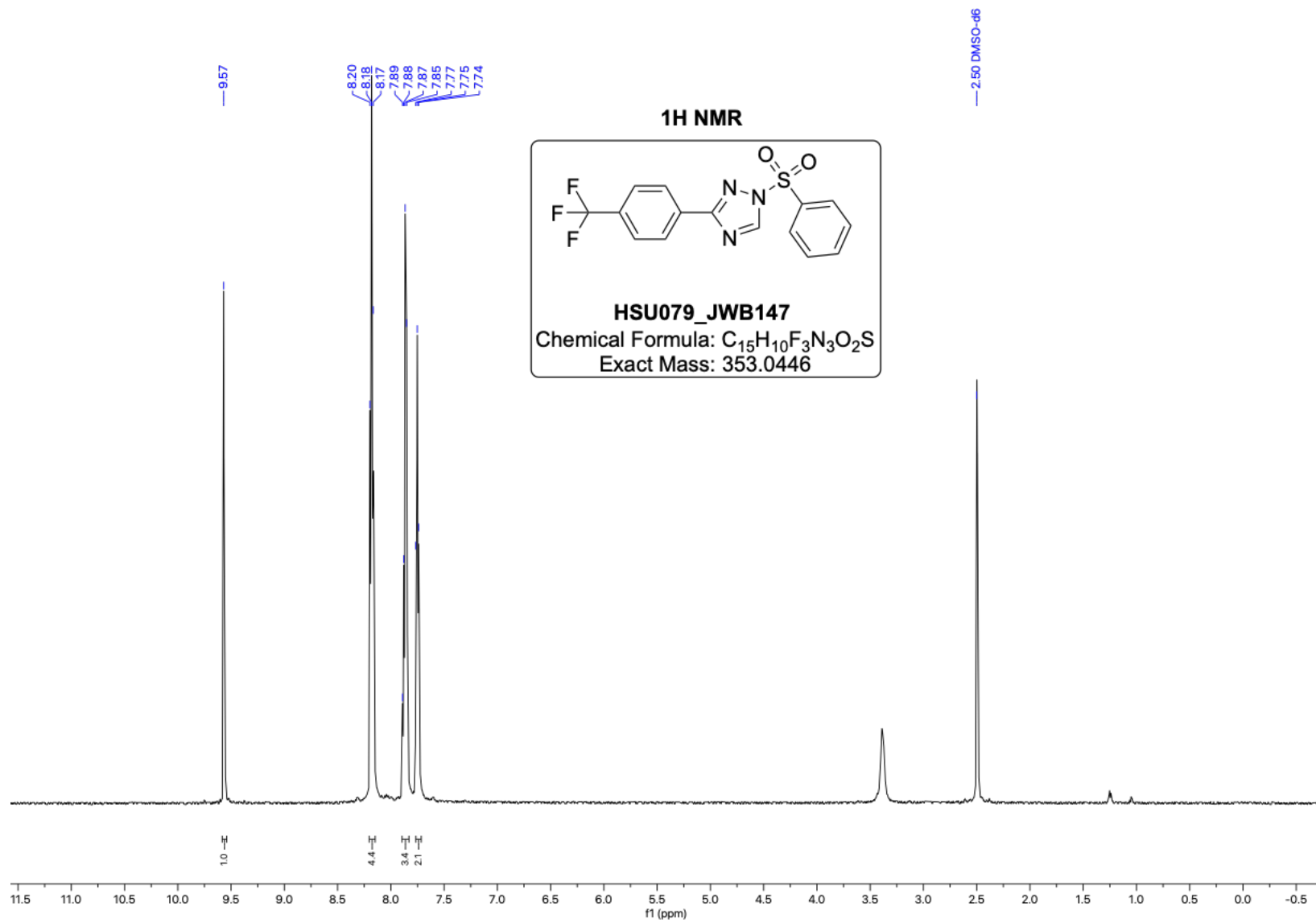


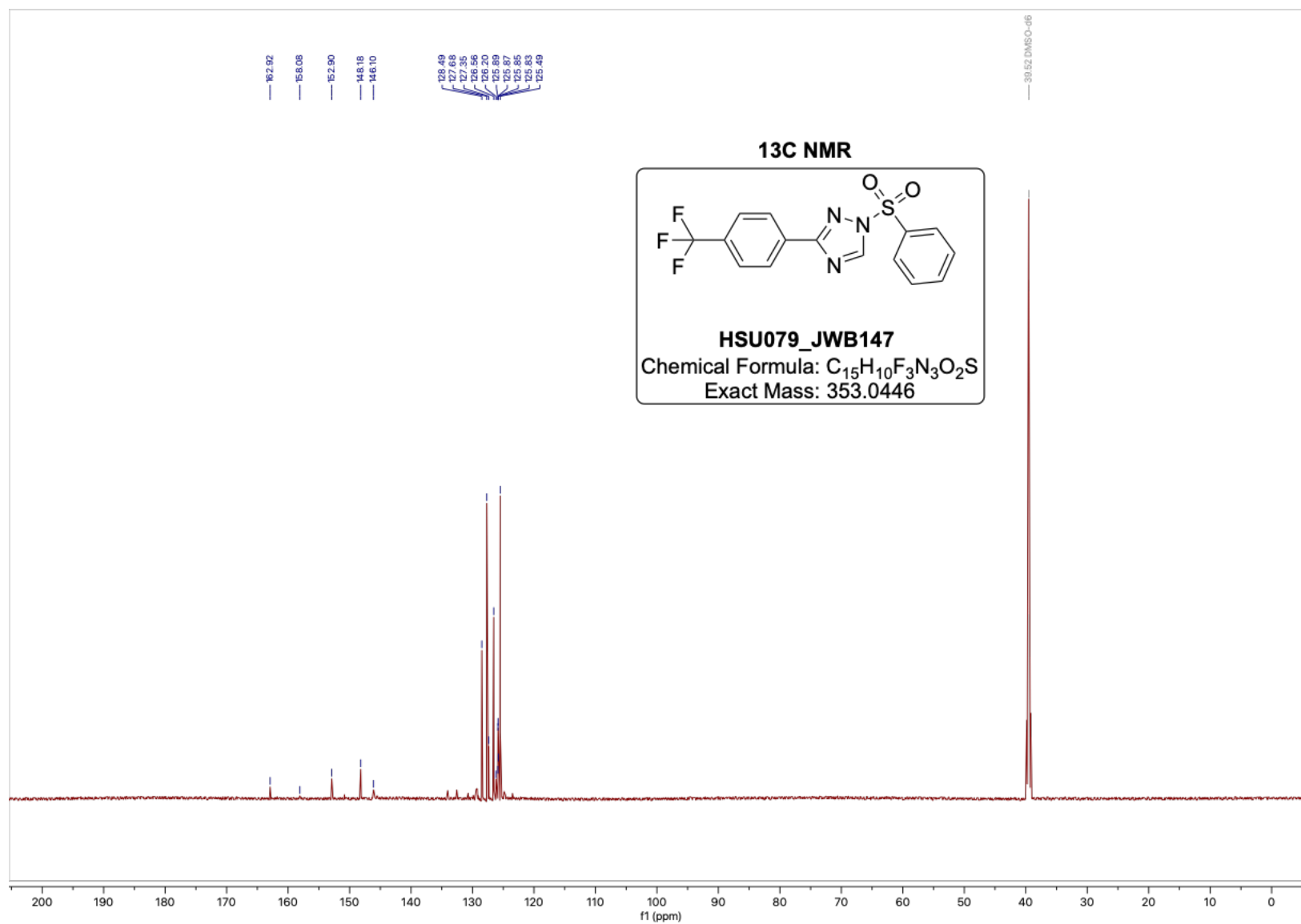


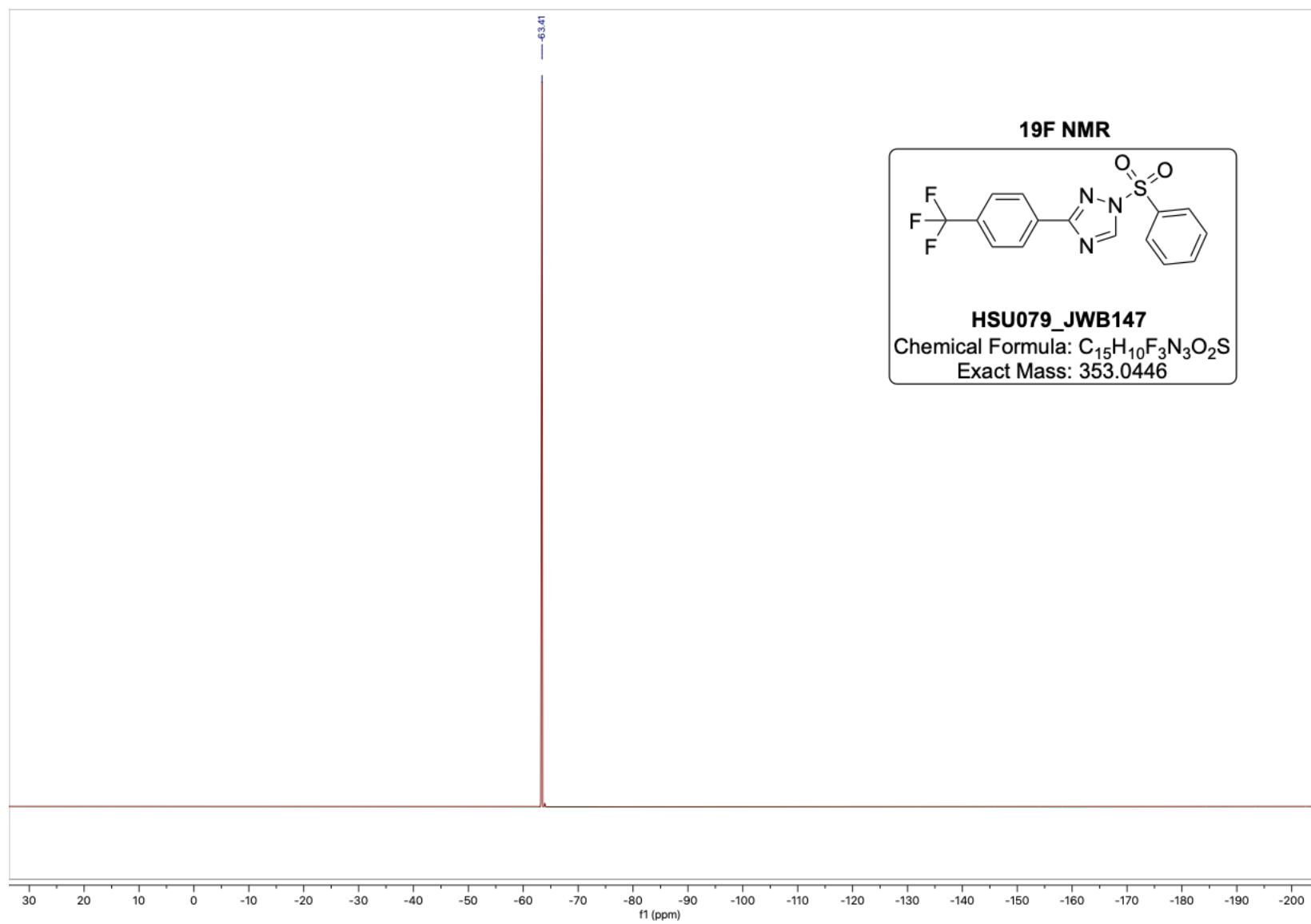


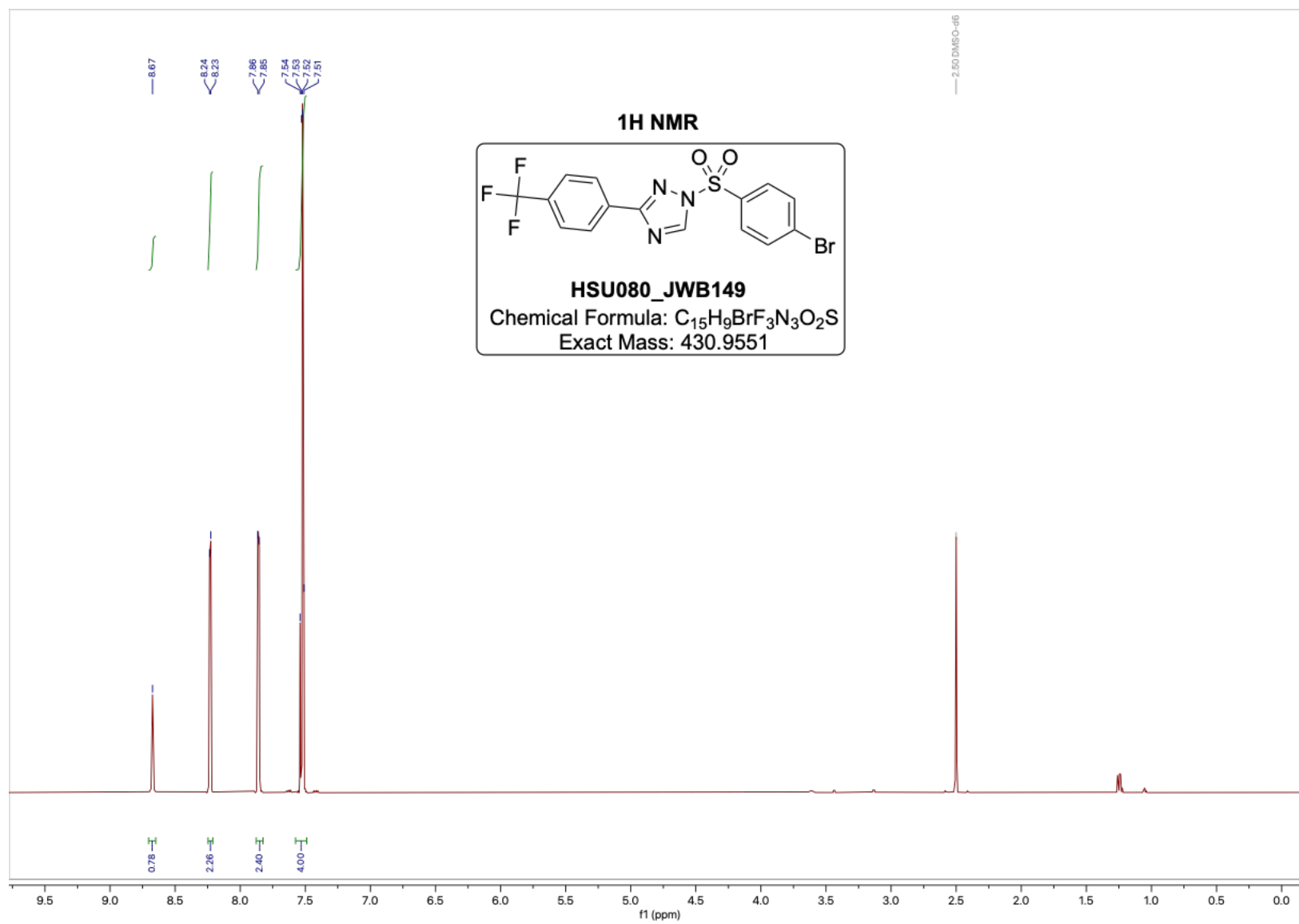




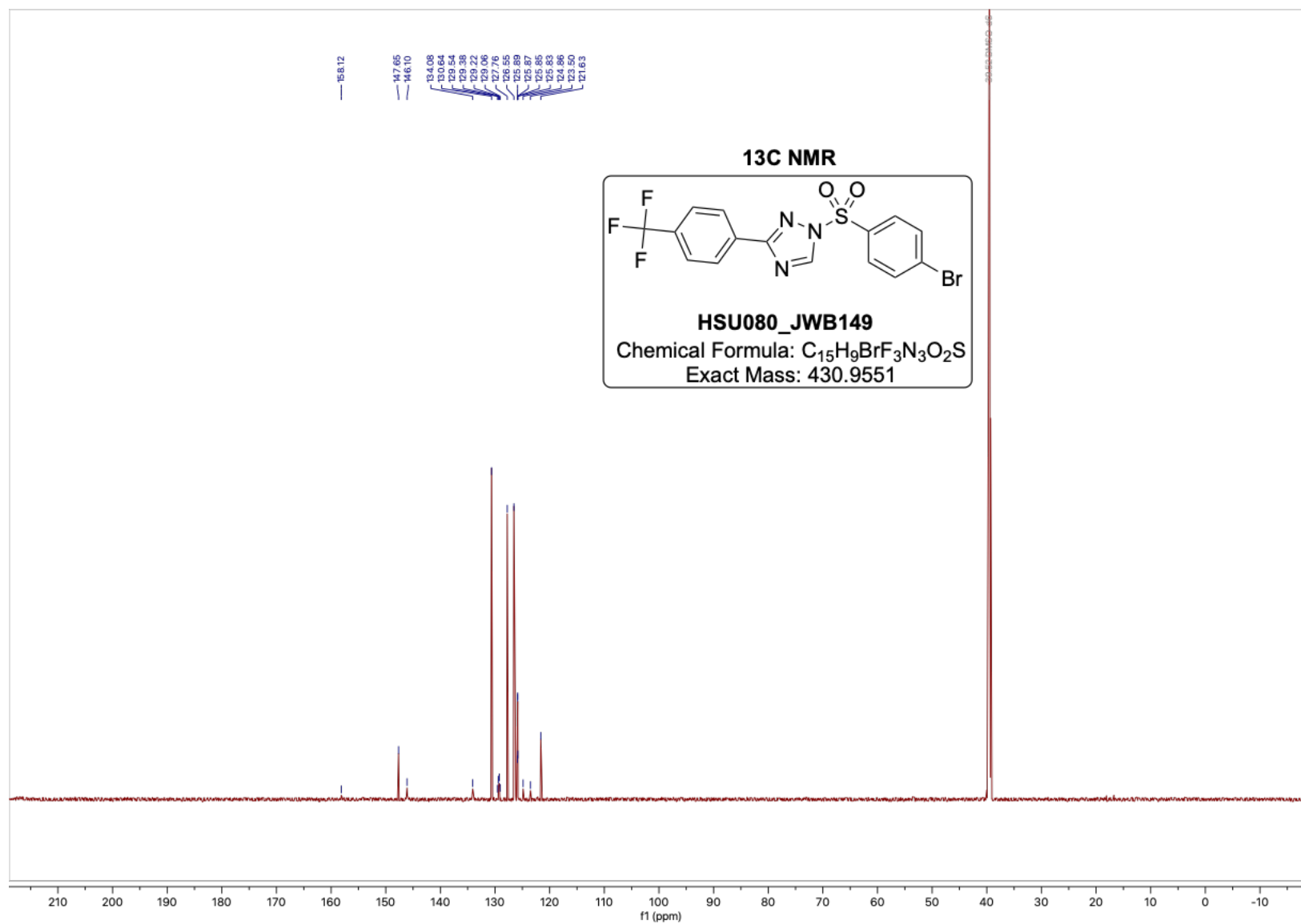


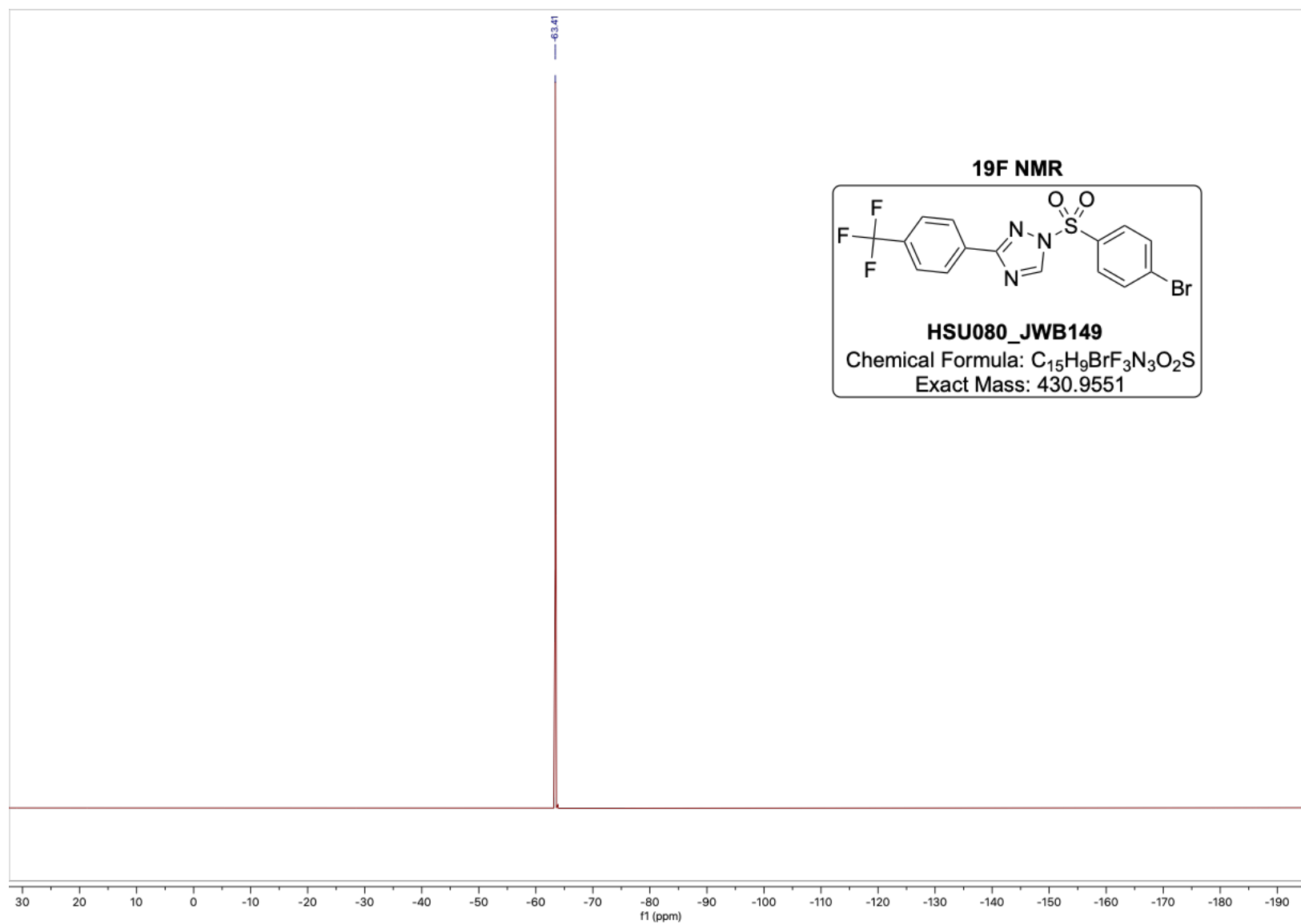


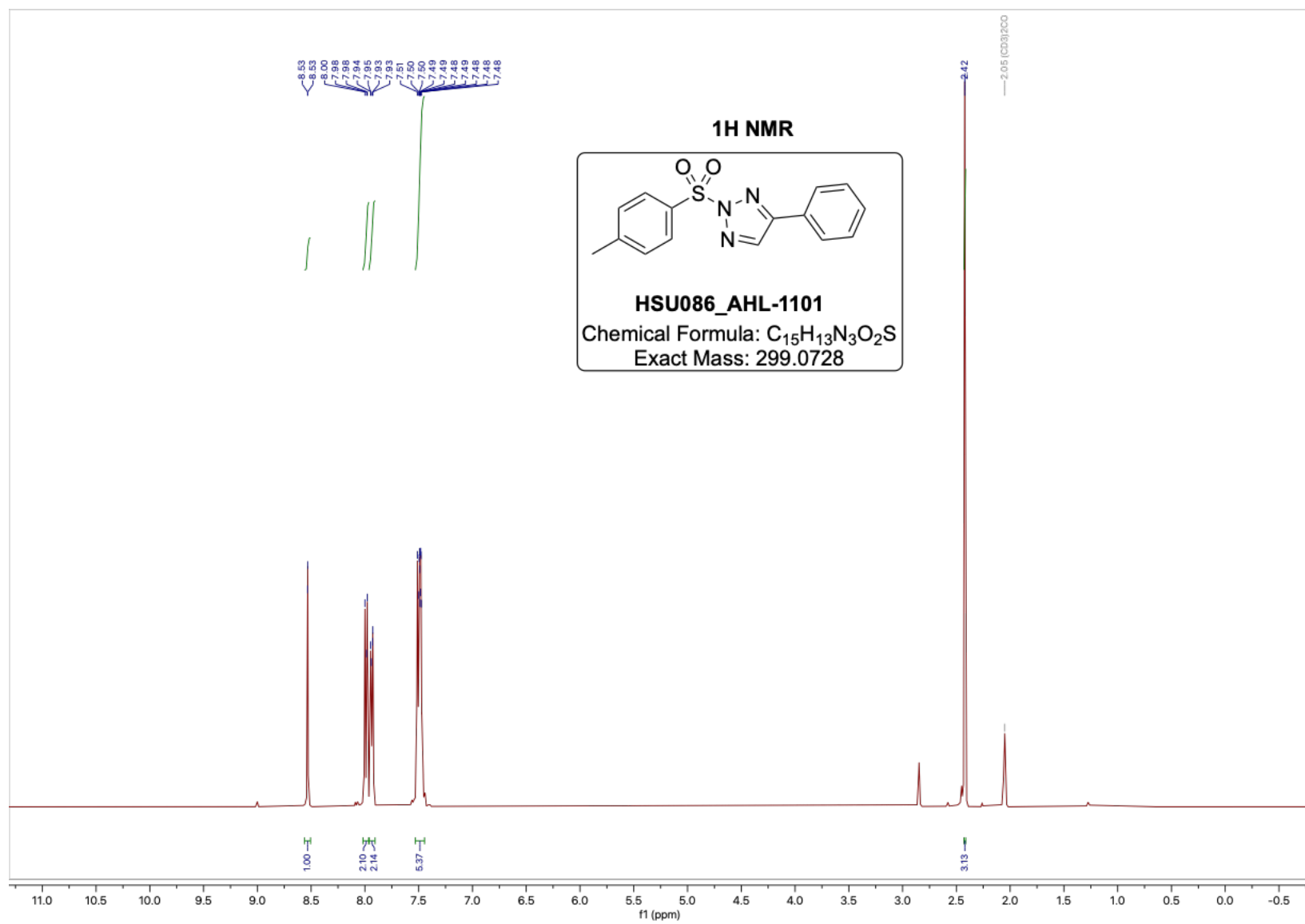


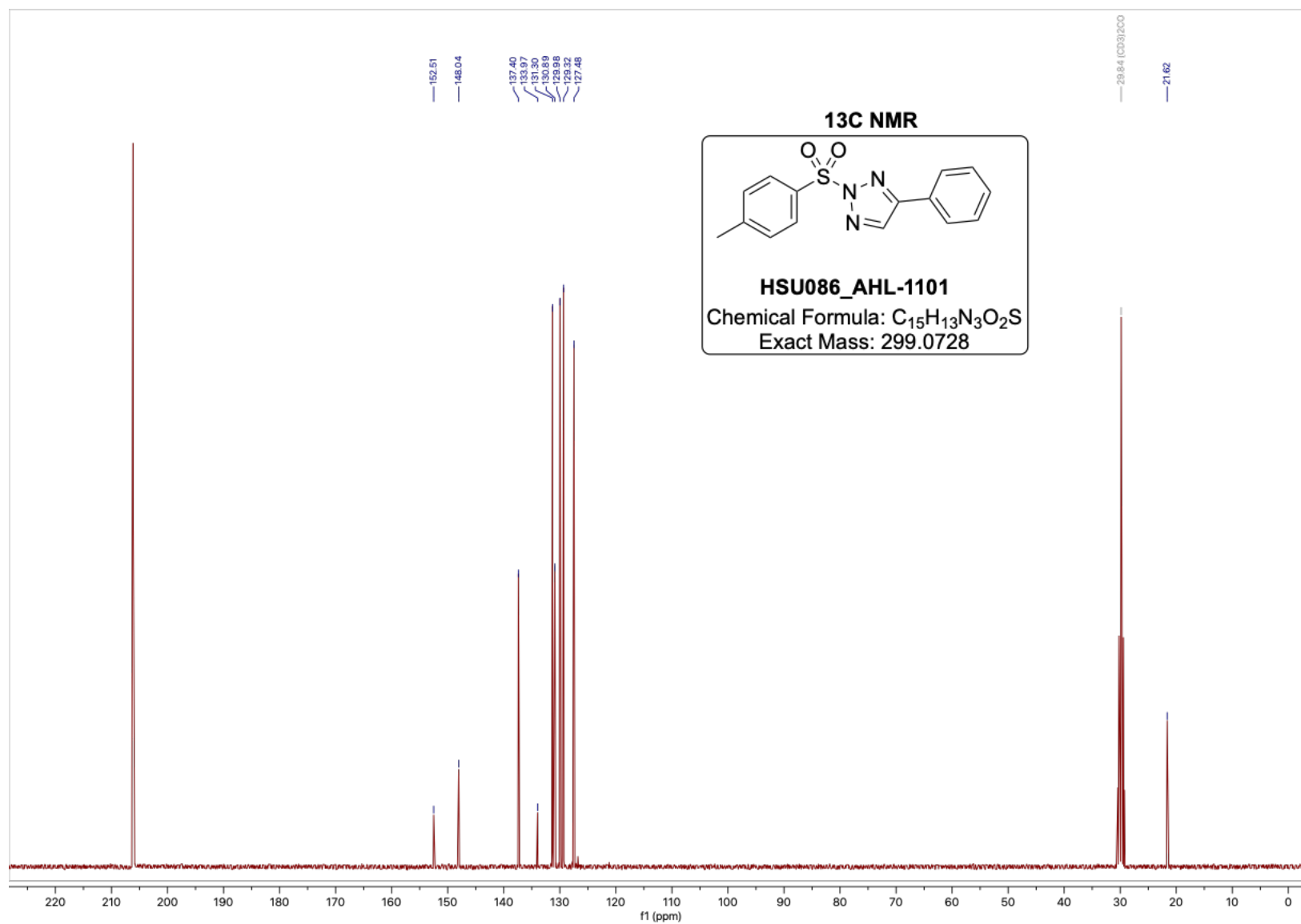




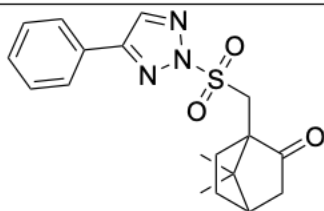




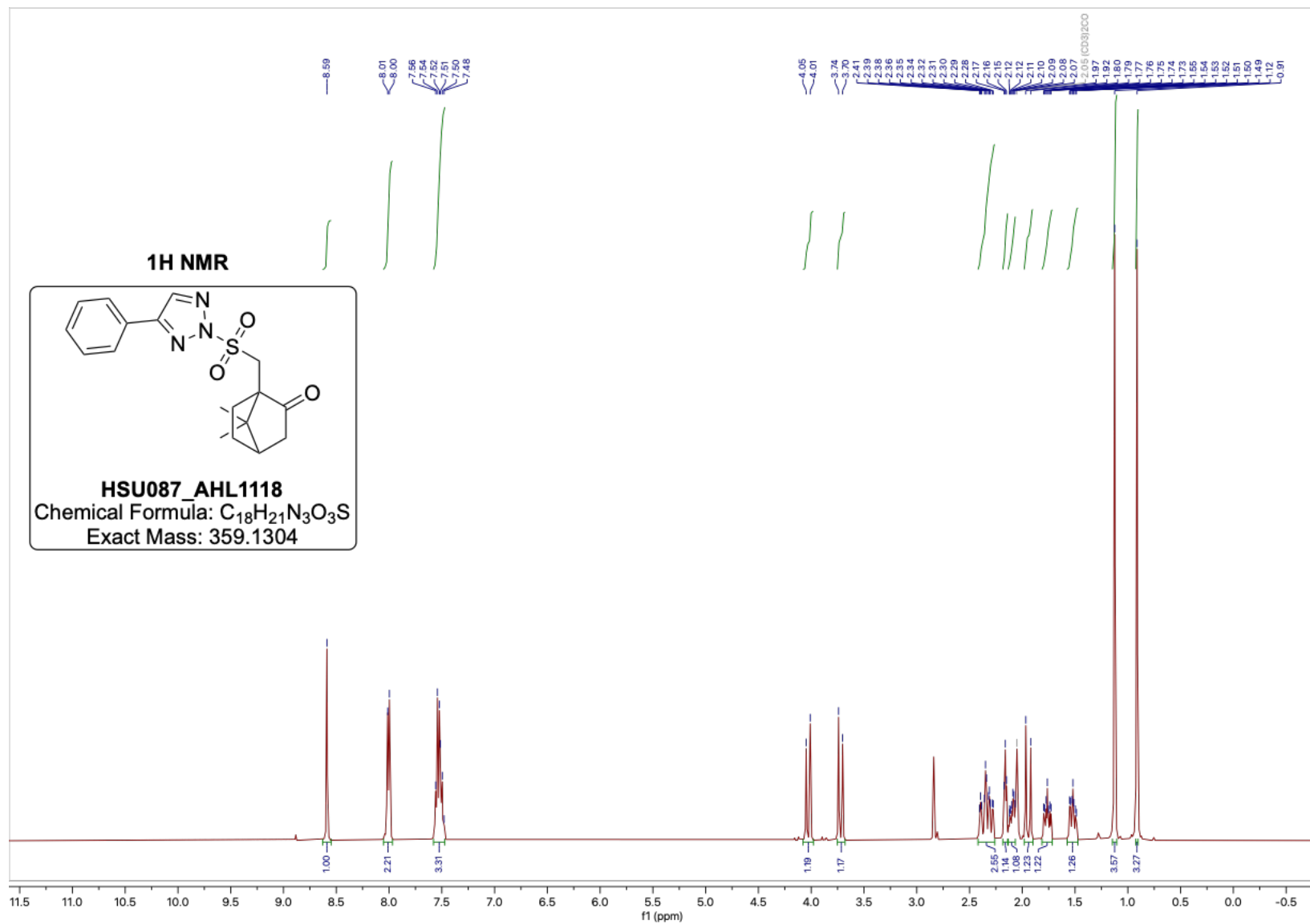


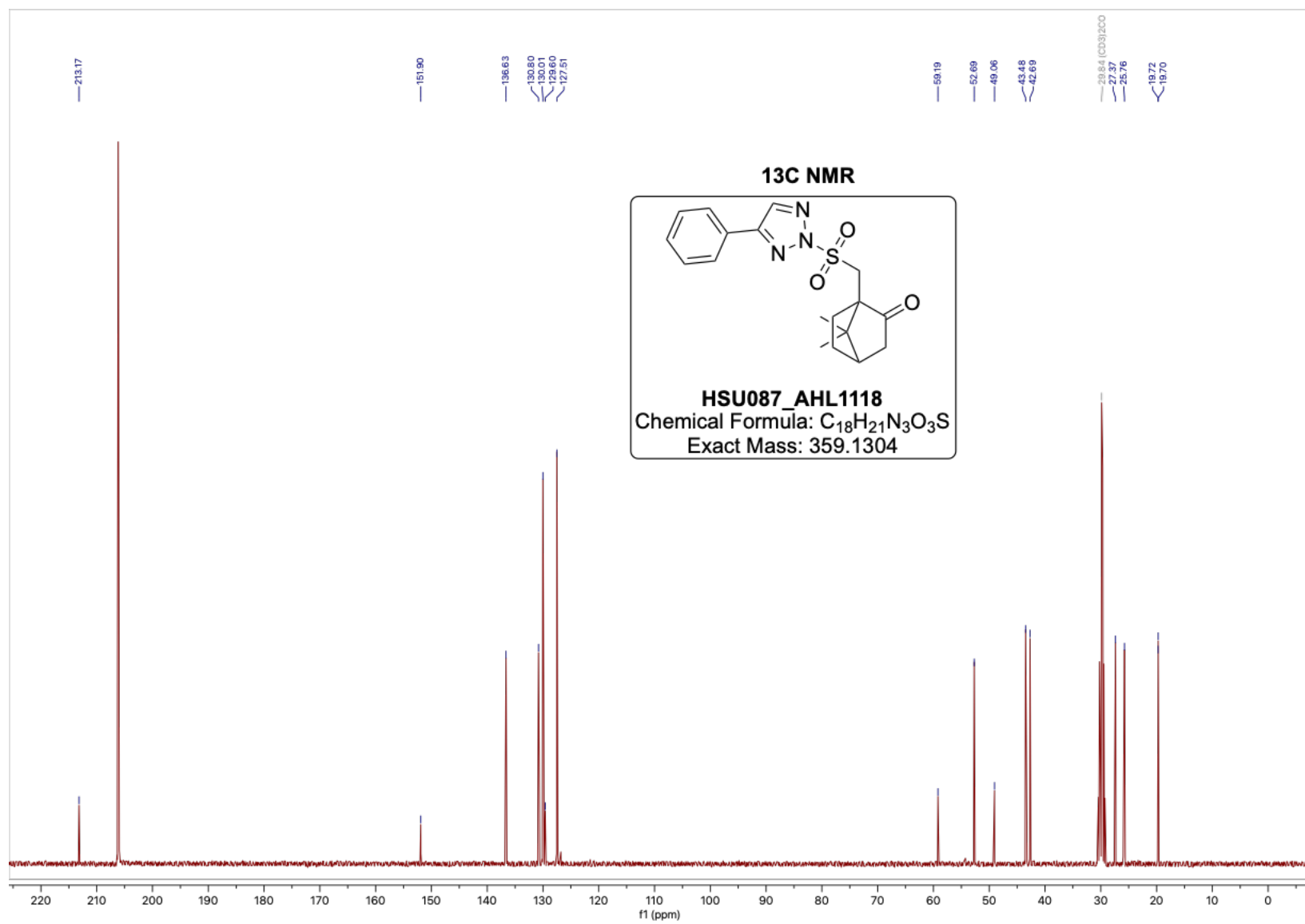


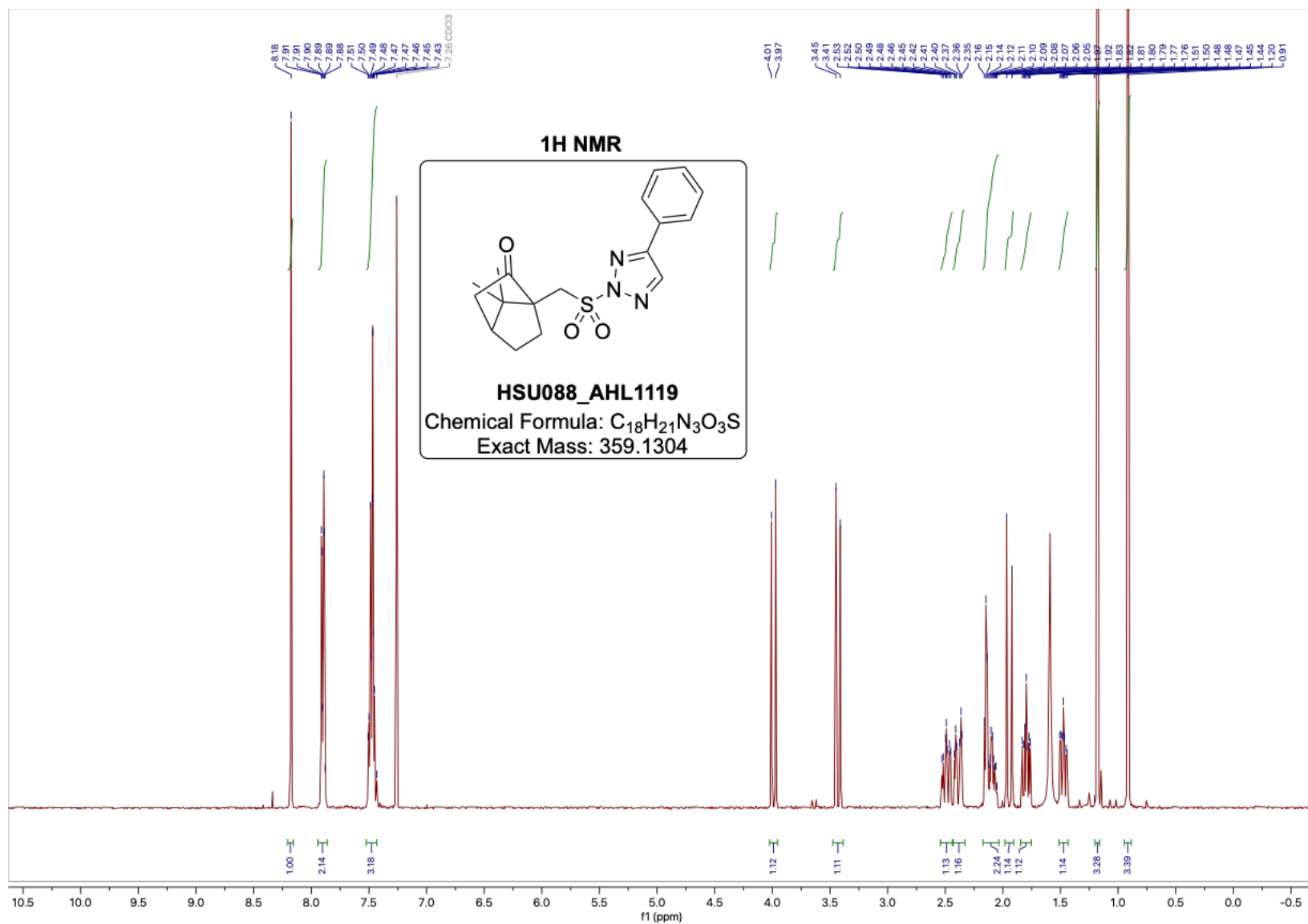
# <sup>1</sup>H NMR

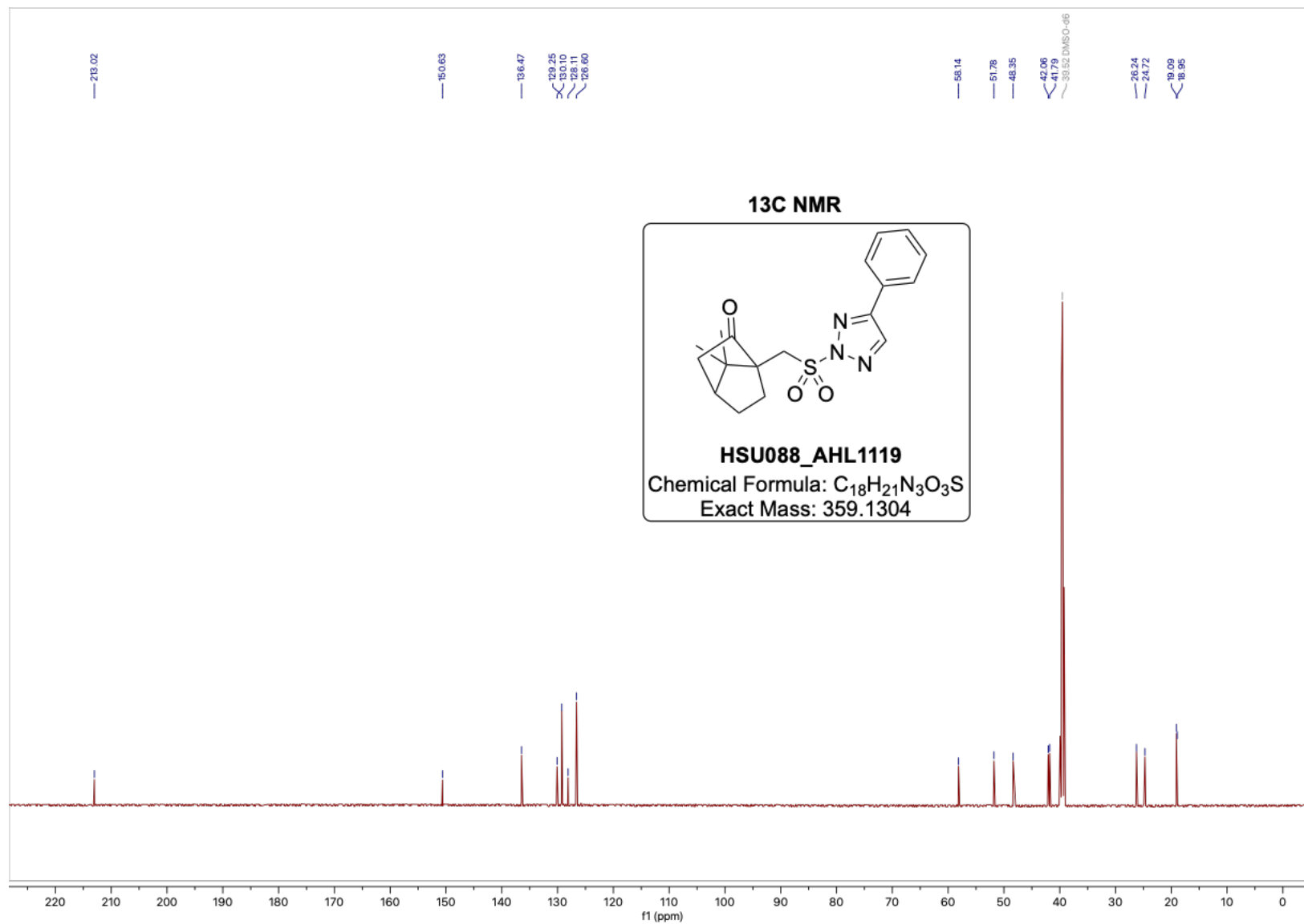


**HSU087\_AHL1118**  
Chemical Formula: C<sub>18</sub>H<sub>21</sub>N<sub>3</sub>O<sub>3</sub>S  
Exact Mass: 359.1304

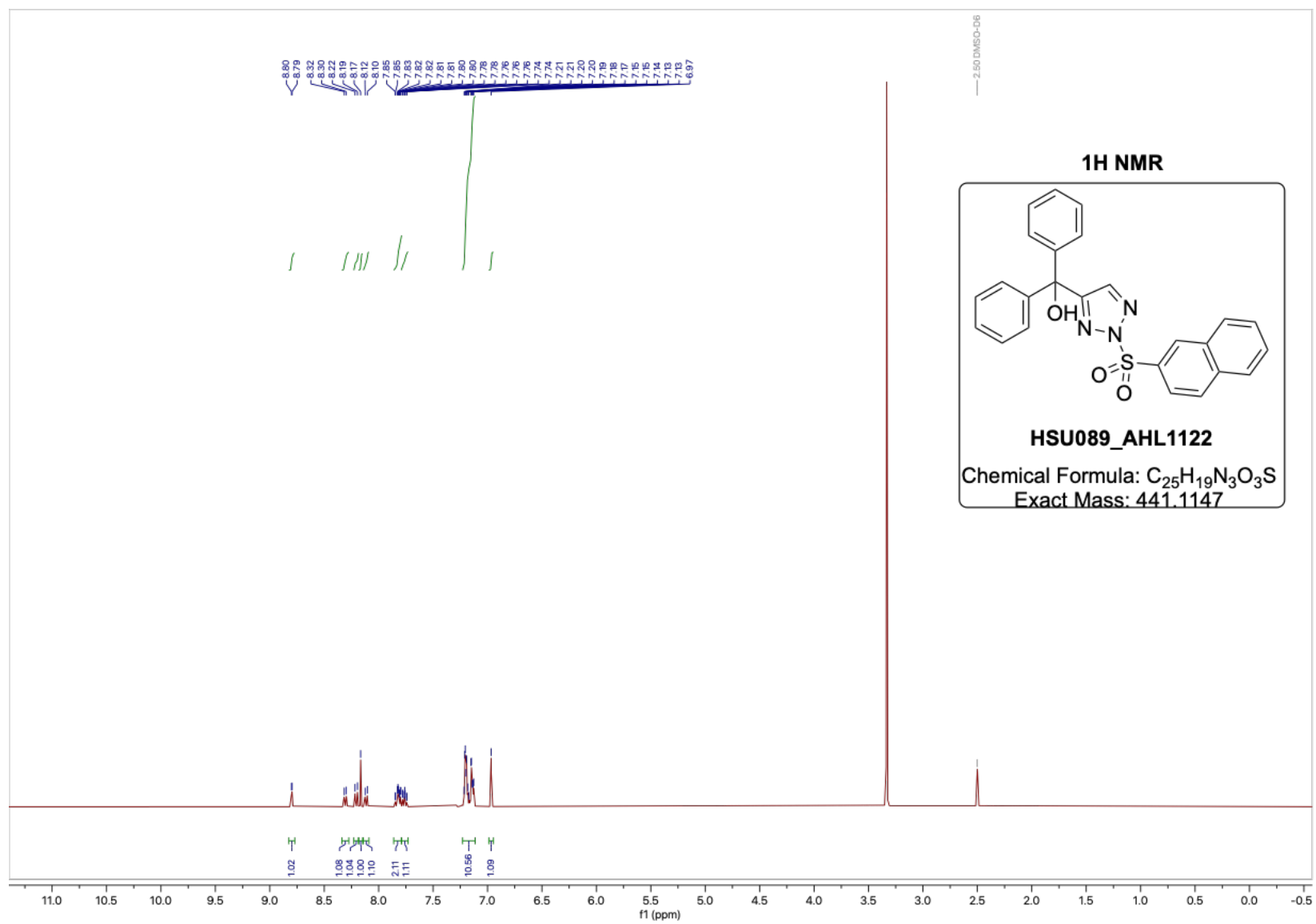




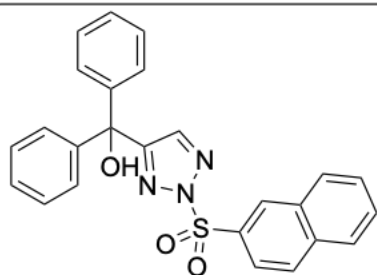






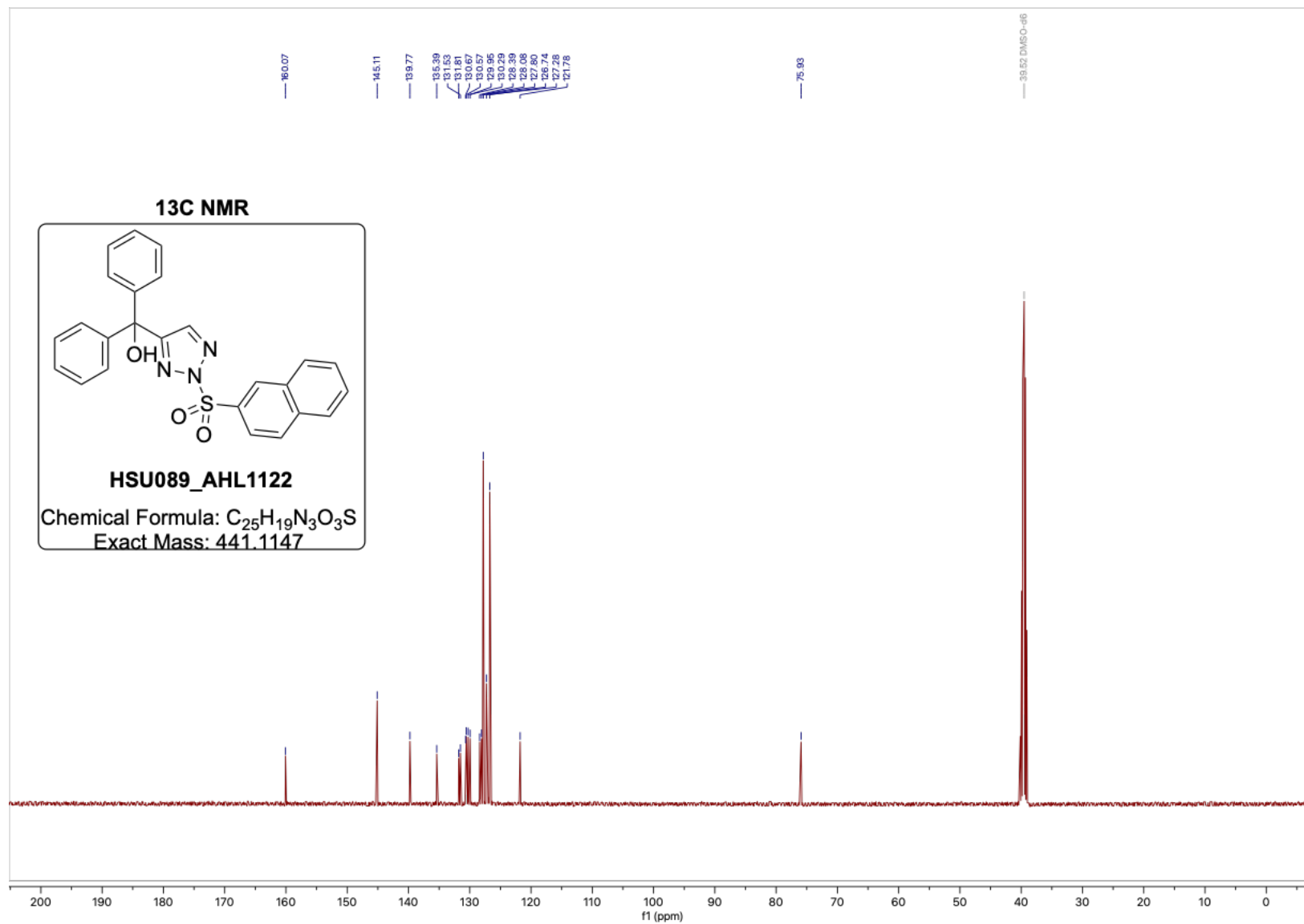


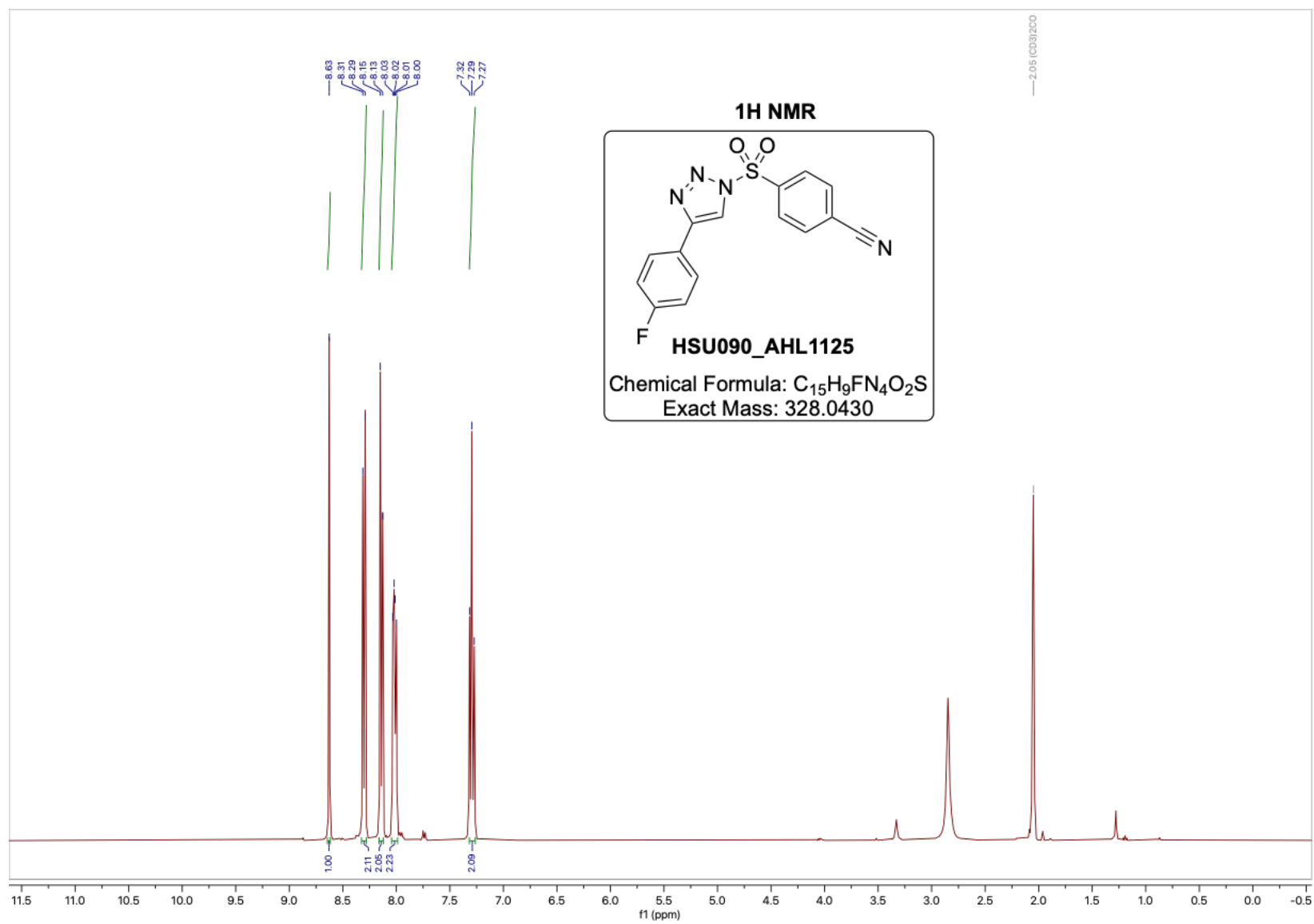
# **<sup>13</sup>C NMR**

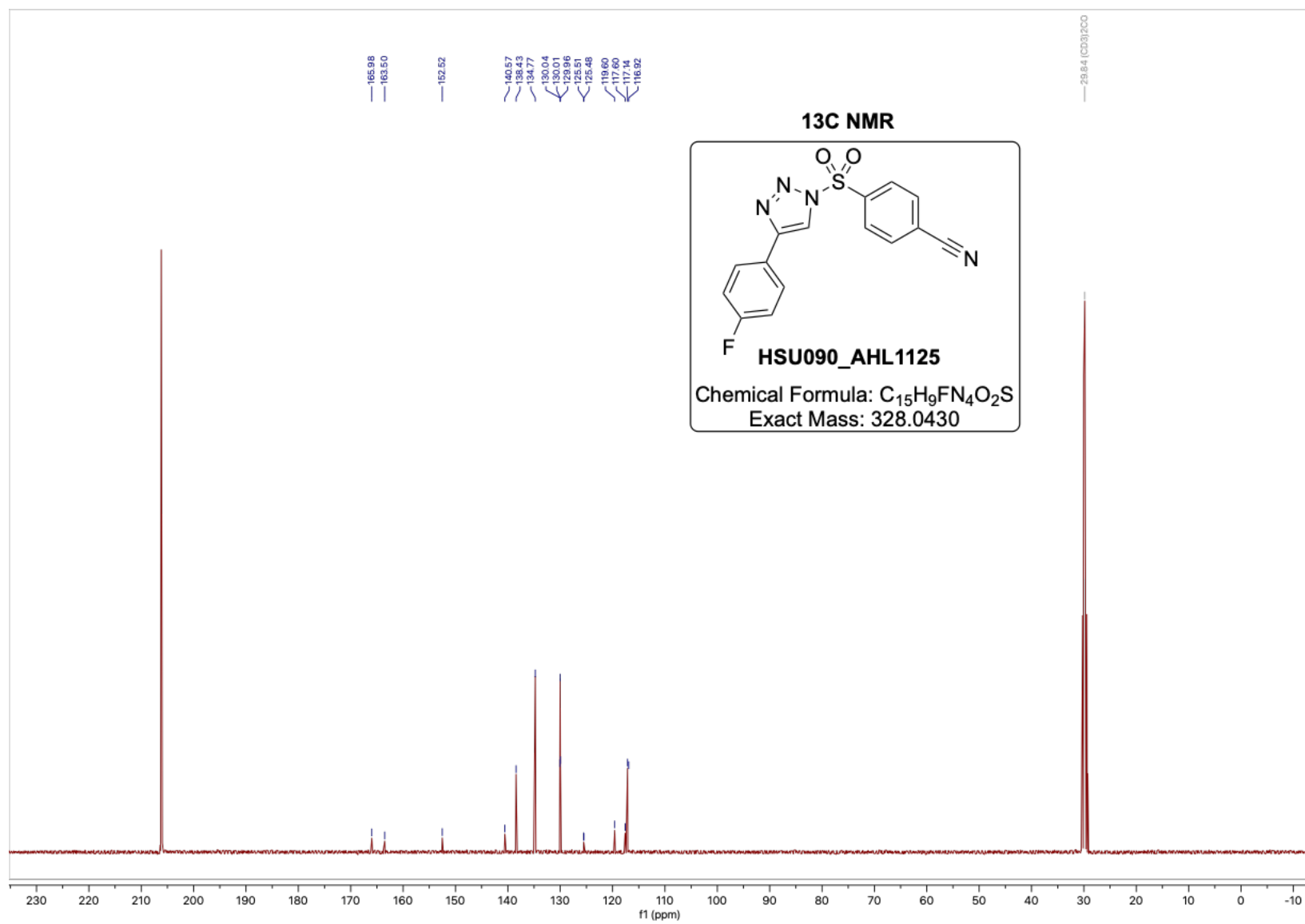


**HSU089\_AHL1122**

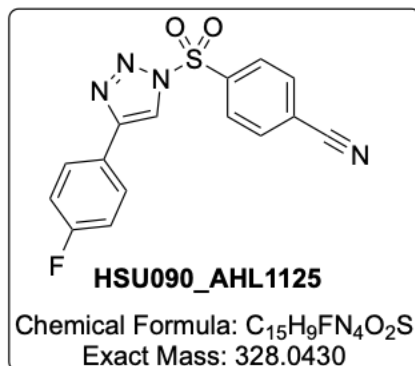
Chemical Formula: C<sub>25</sub>H<sub>19</sub>N<sub>3</sub>O<sub>3</sub>S  
Exact Mass: 441.1147



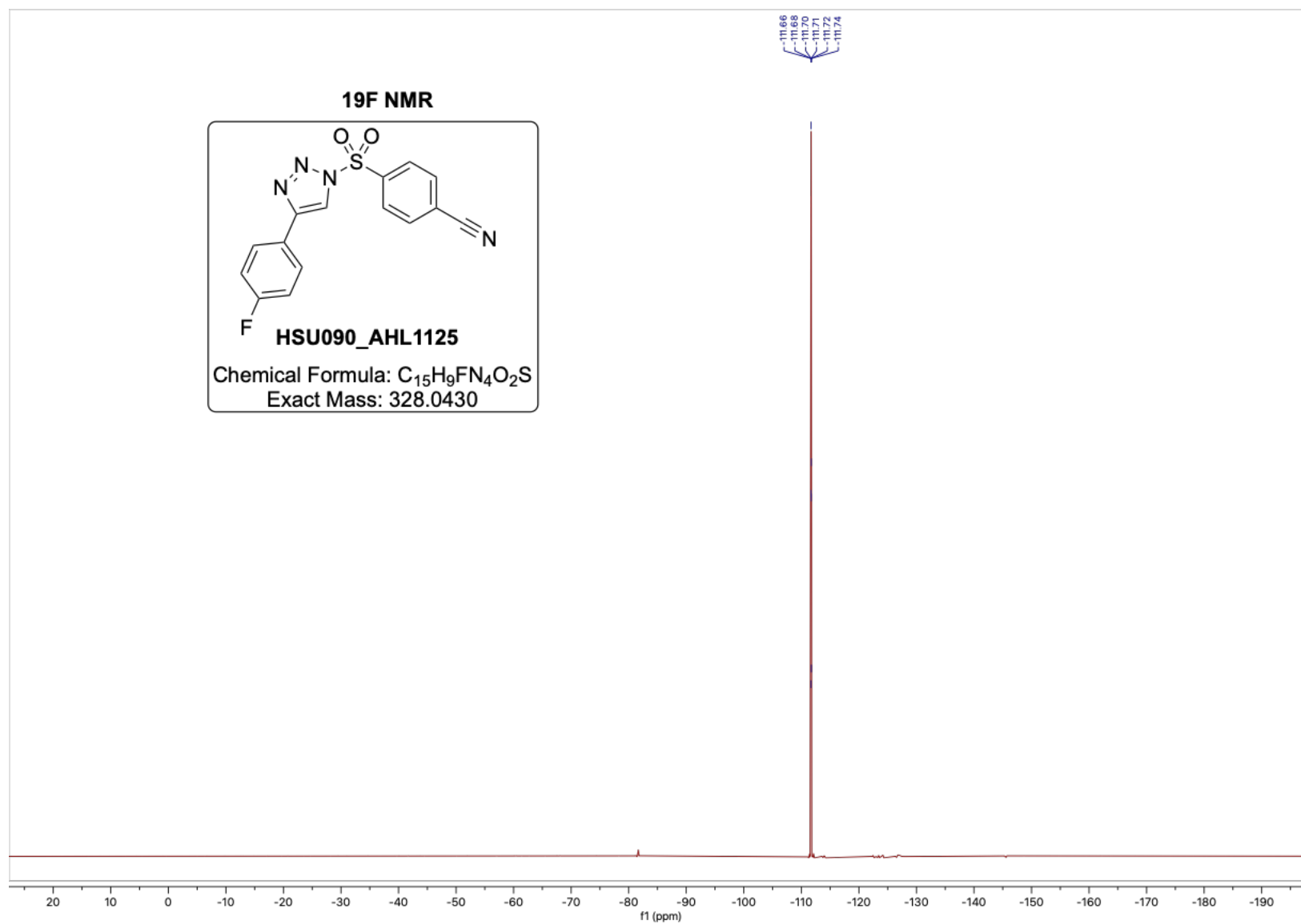


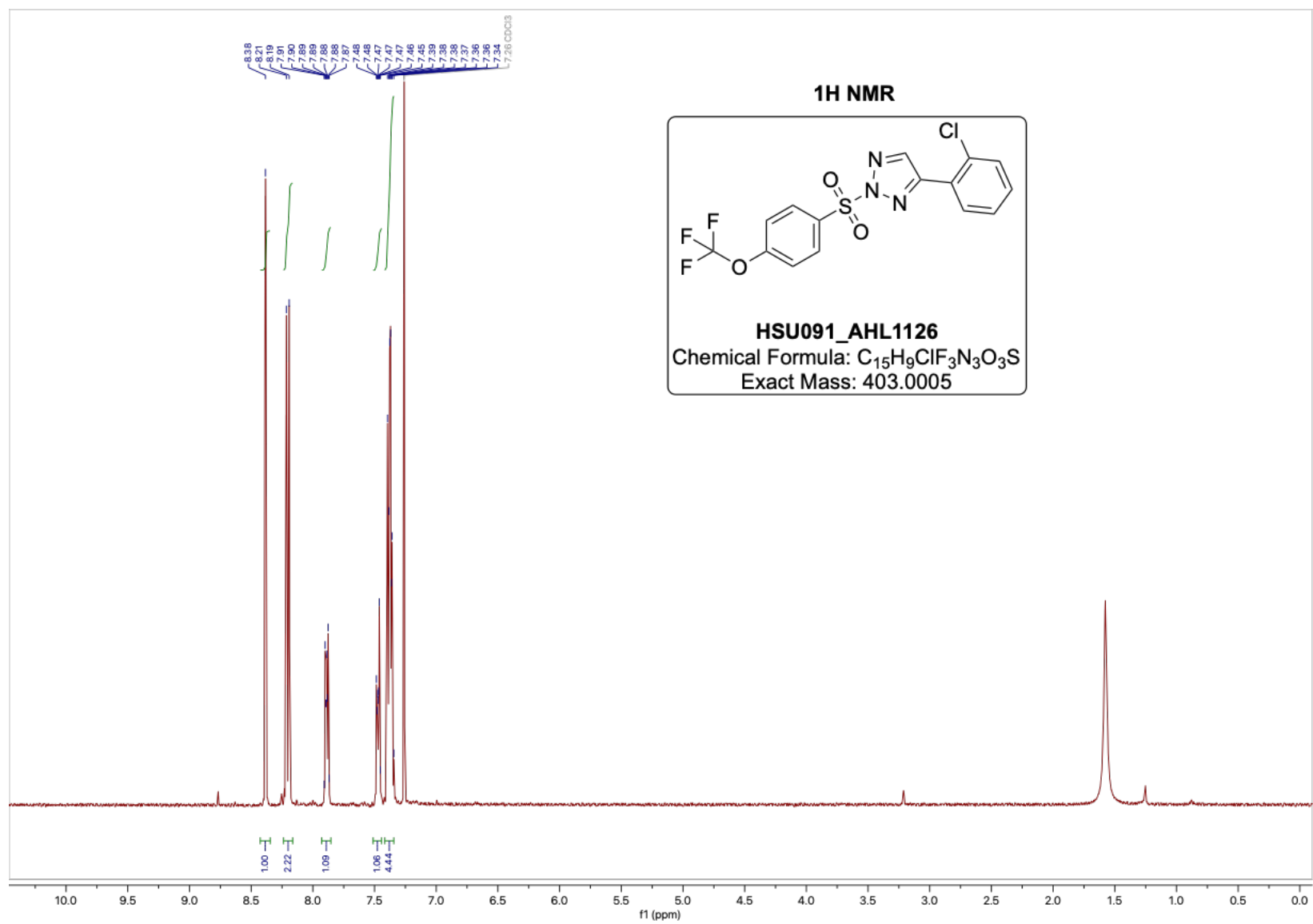


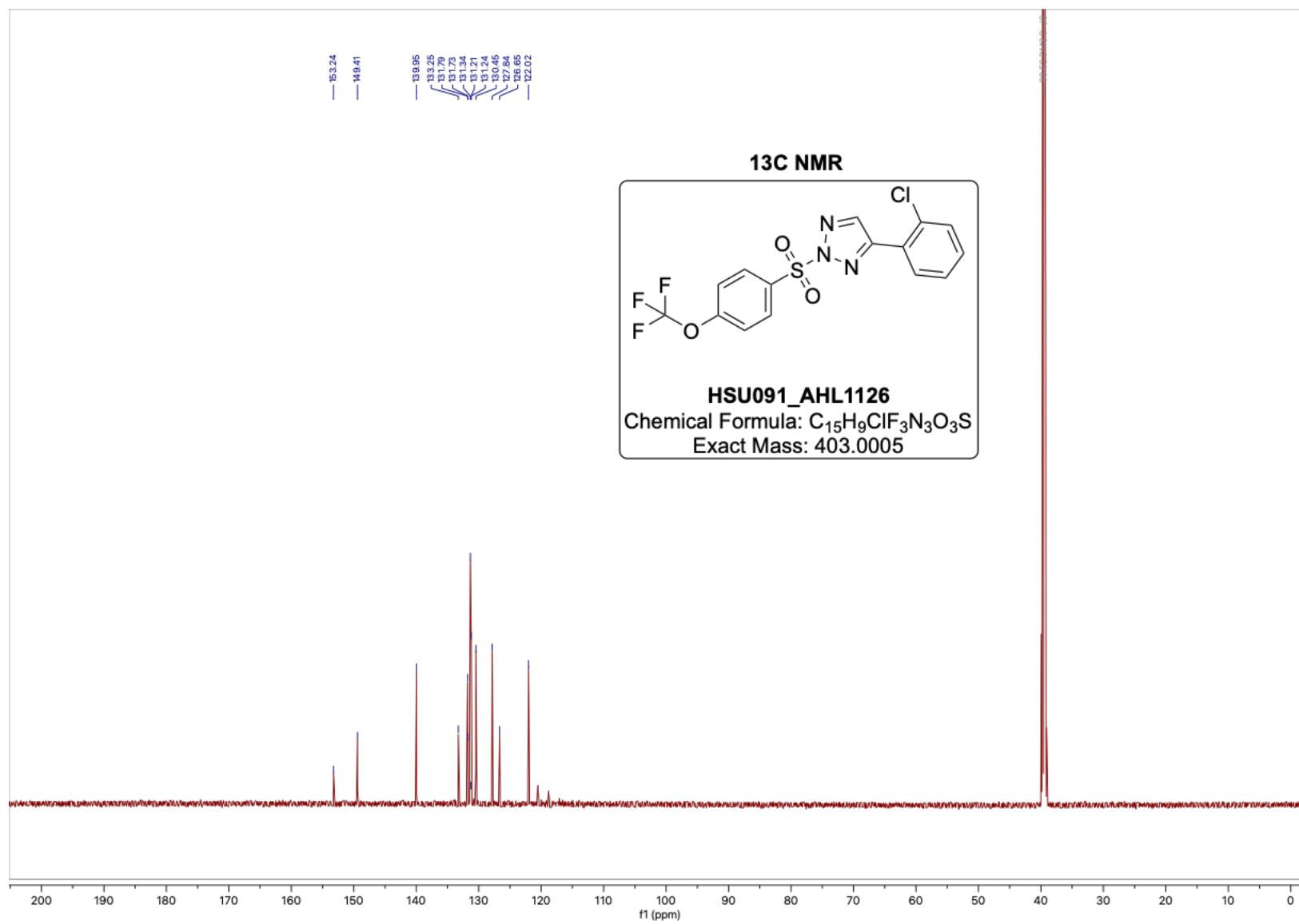
**<sup>19</sup>F NMR**

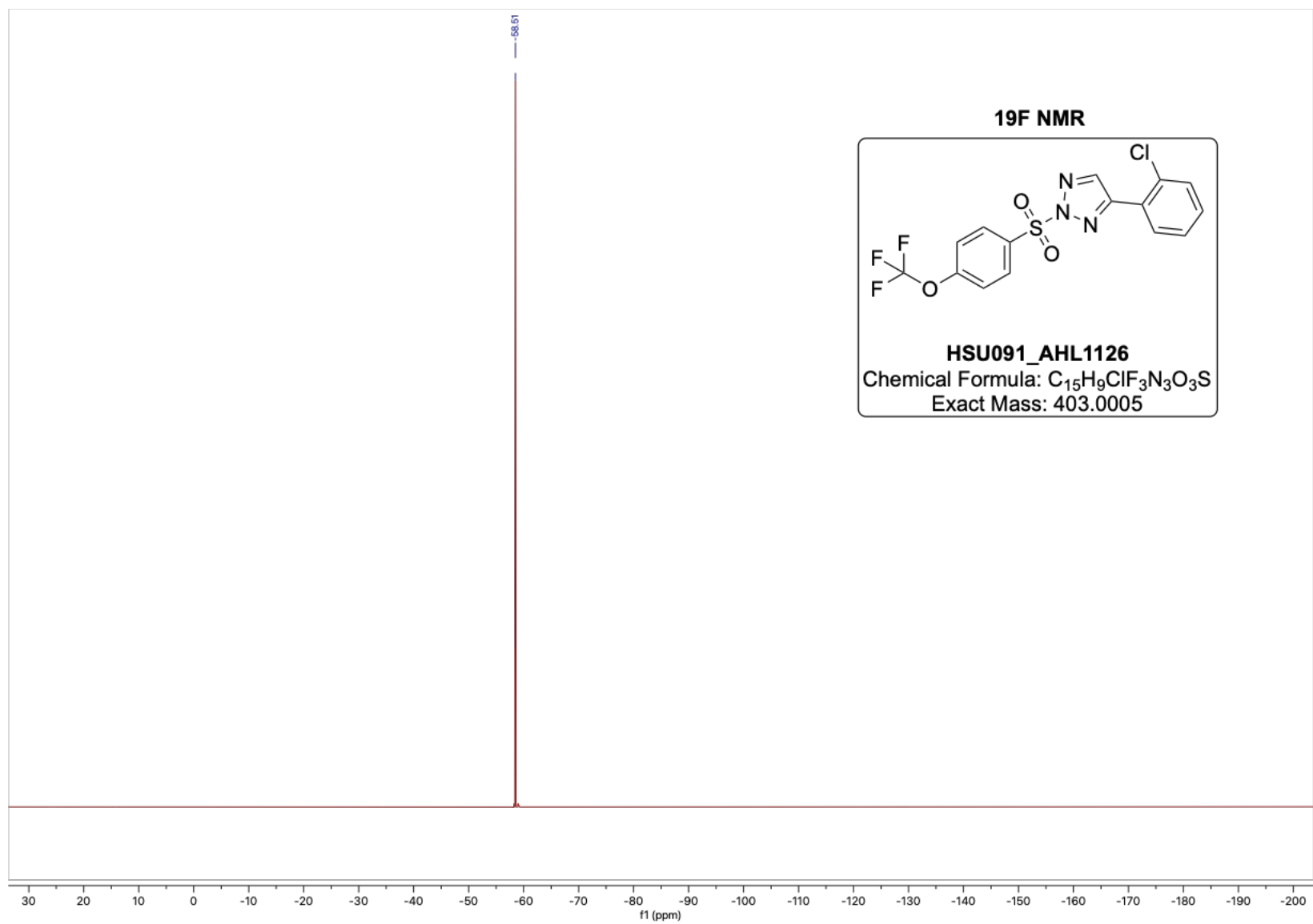


-111.66  
-111.68  
-111.70  
-111.72  
-111.74

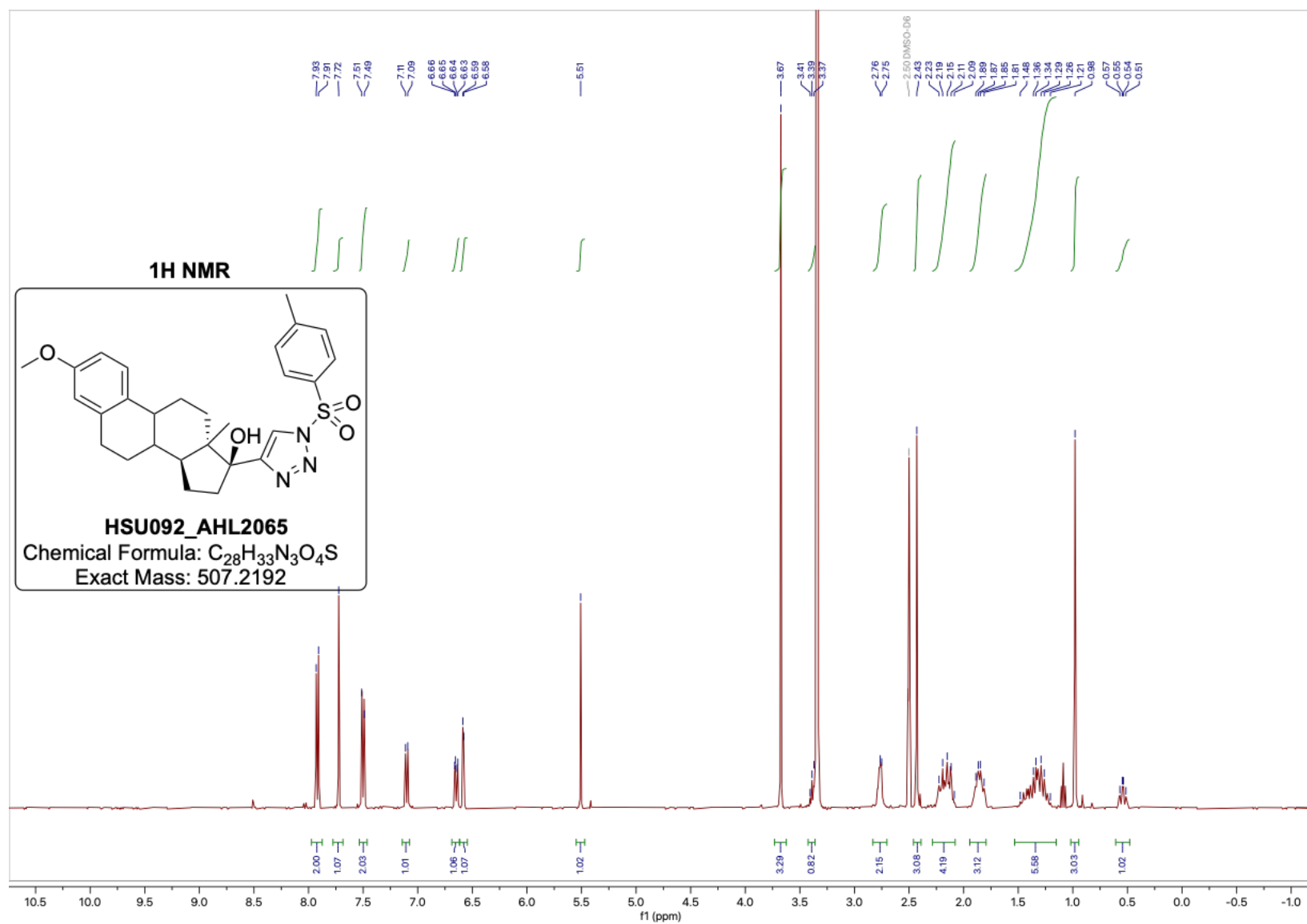




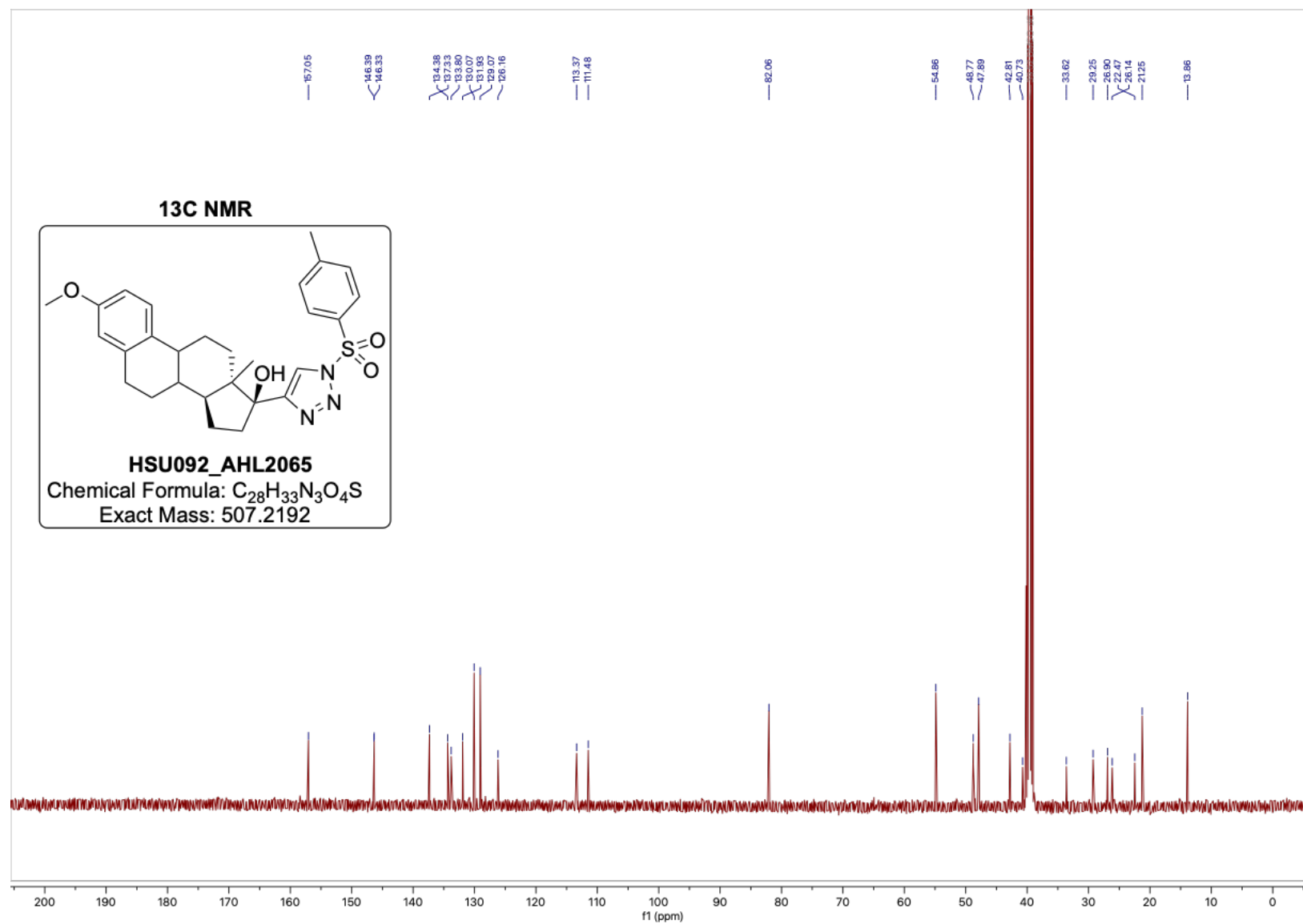
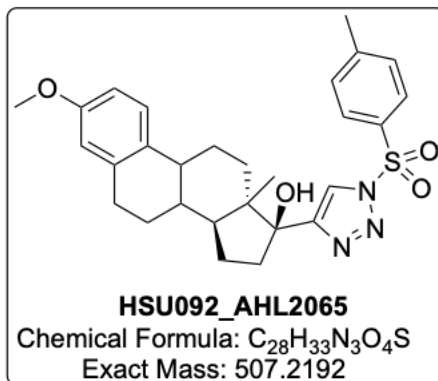


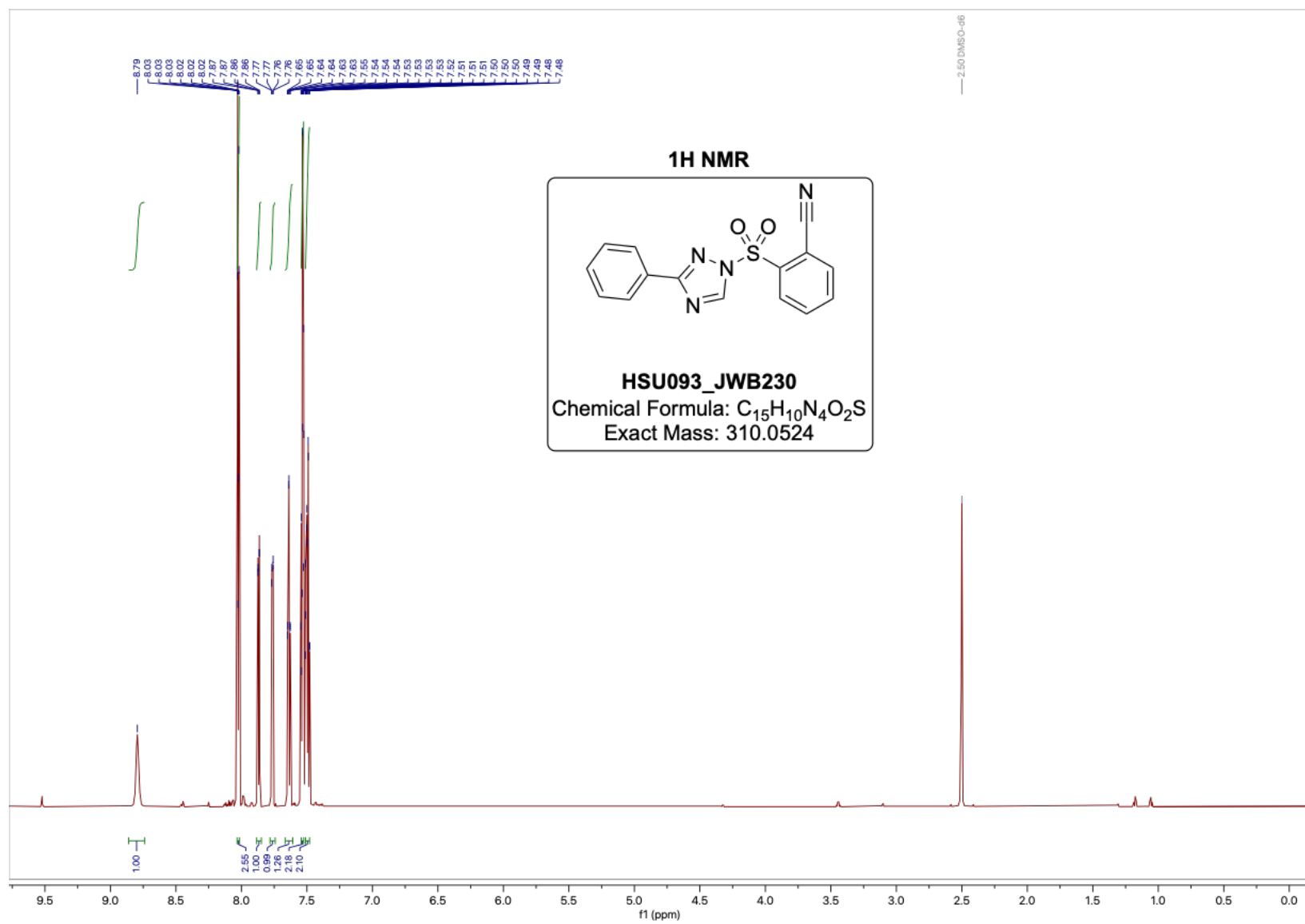


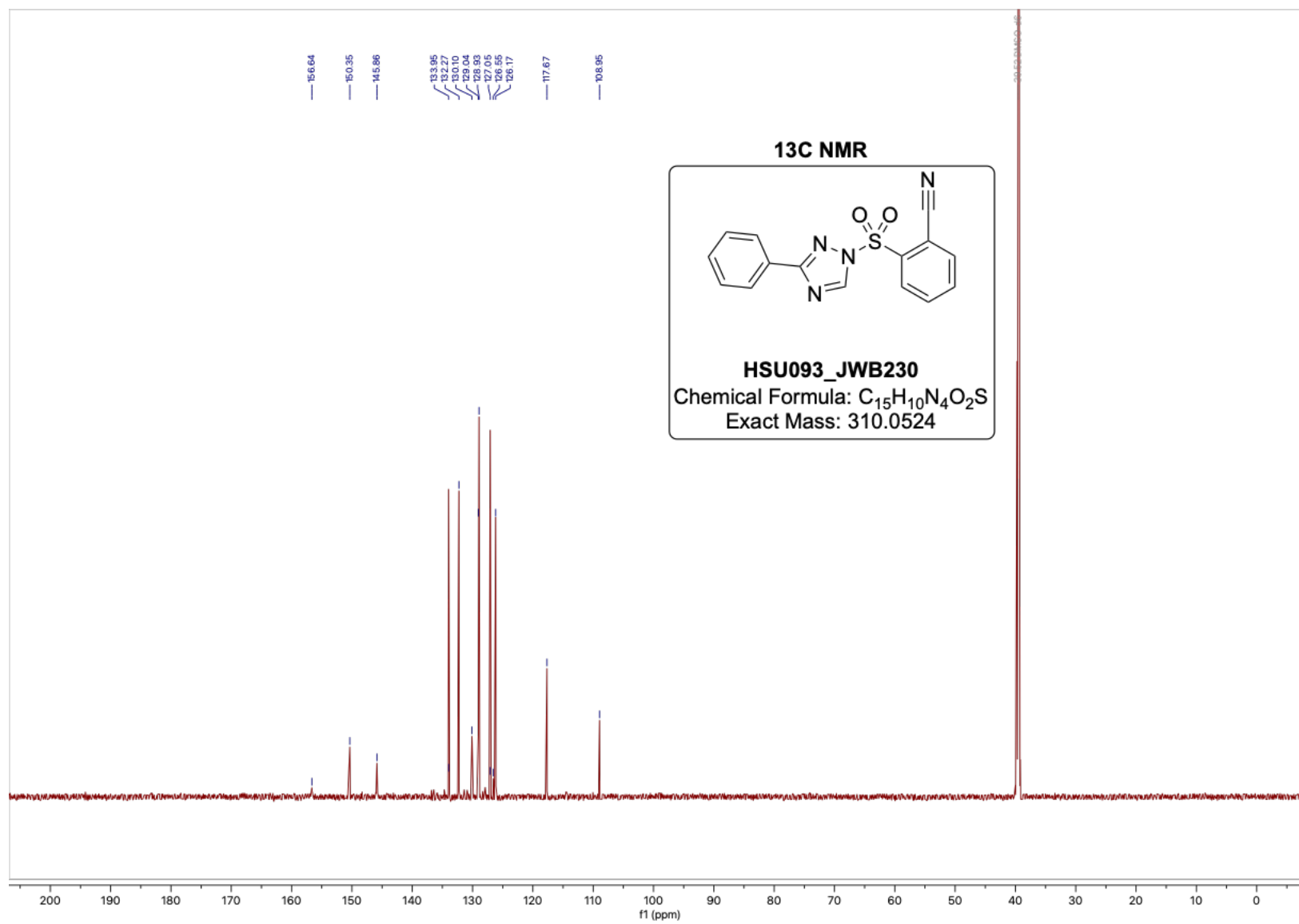


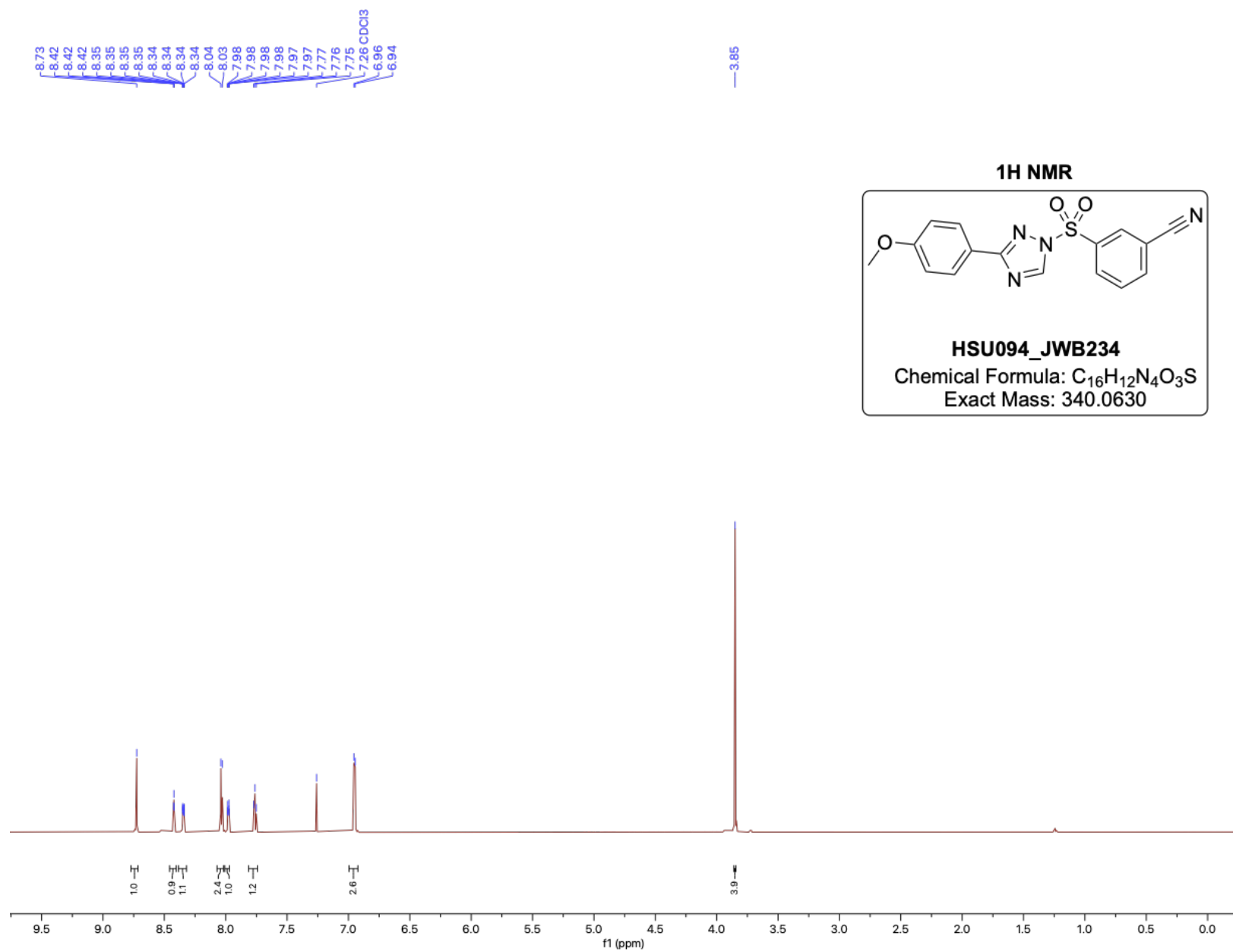


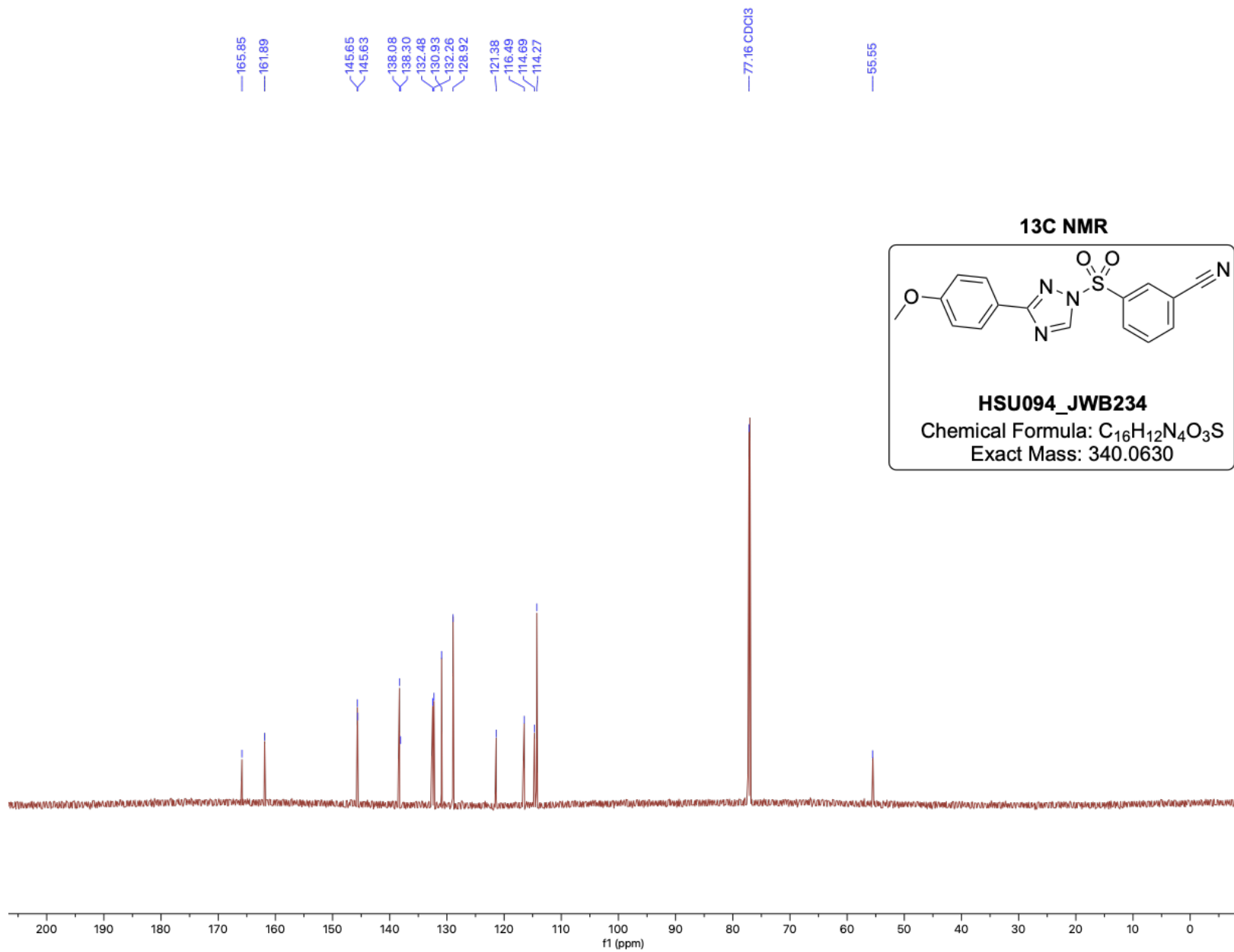
# **<sup>13</sup>C NMR**



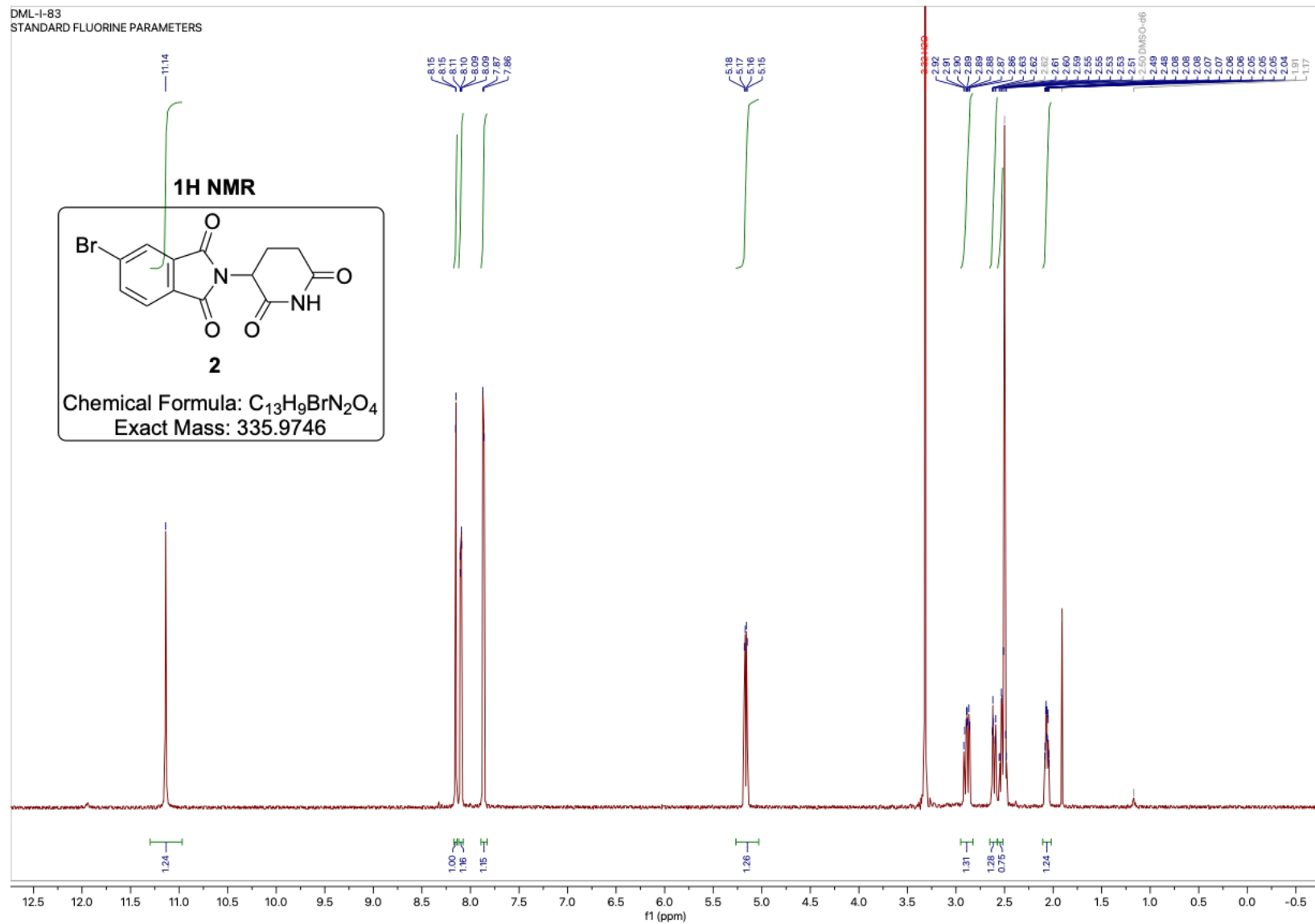
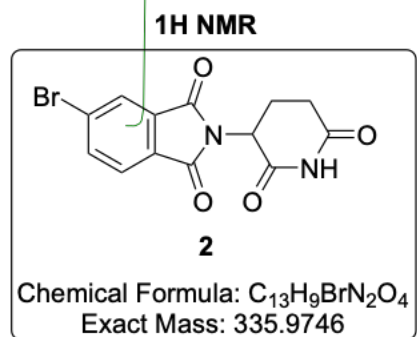






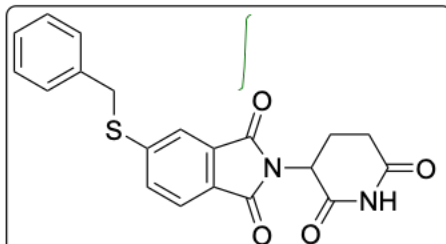


DML-I-83  
STANDARD FLUORINE PARAMETERS



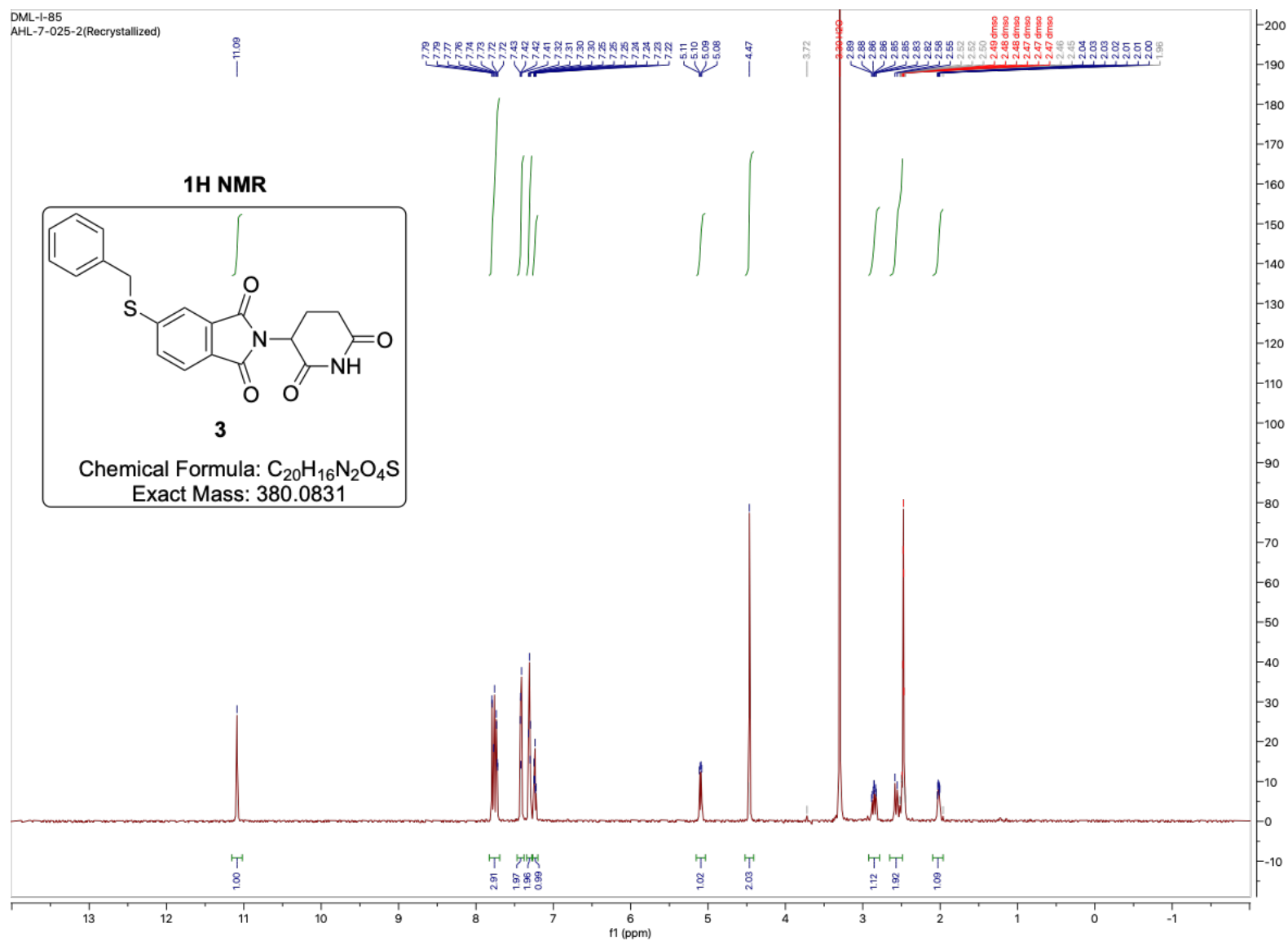
DML-I-85  
AHL-7-025-2(Recrystallized)

# 1H NMR



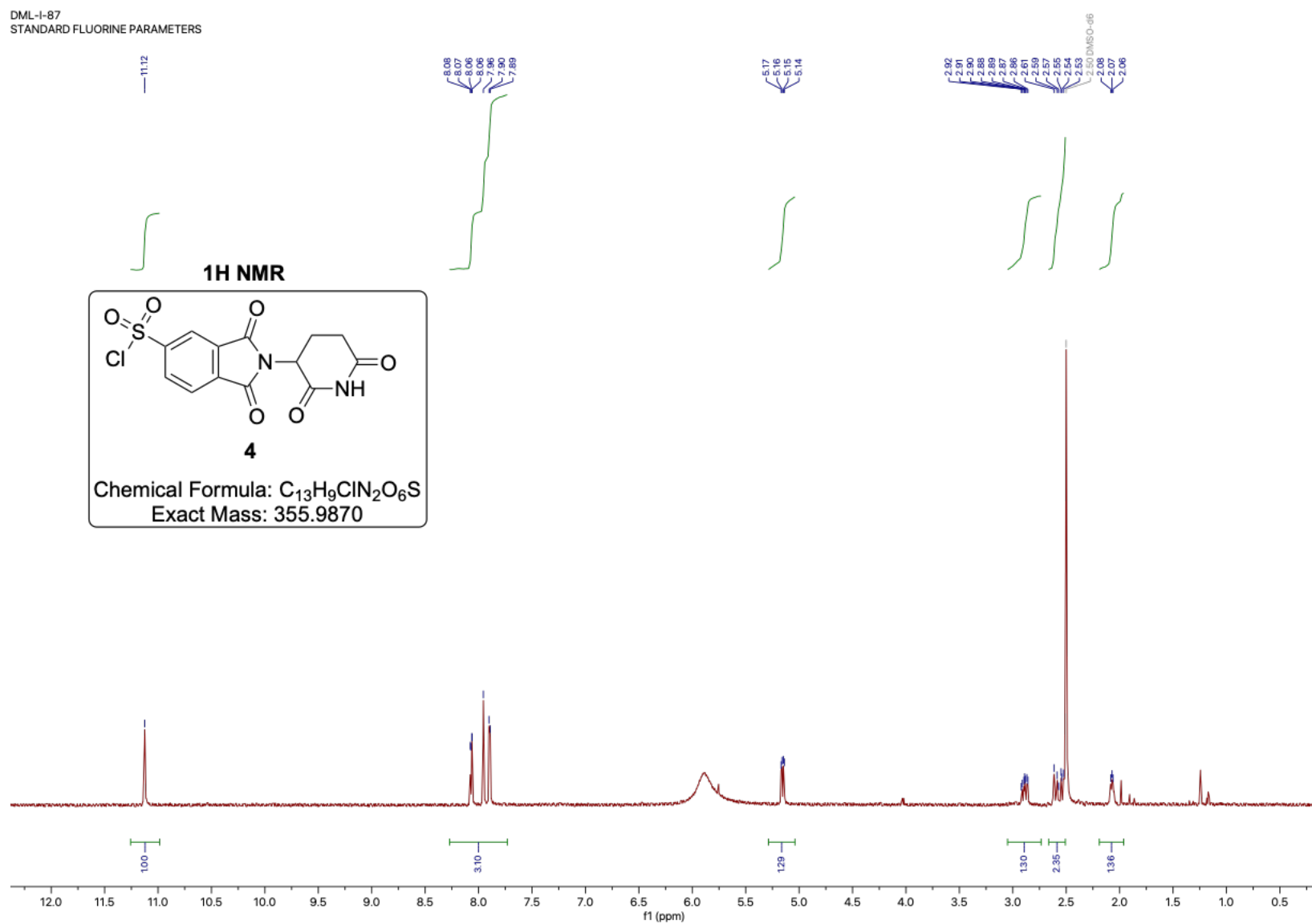
3

Chemical Formula:  $C_{20}H_{16}N_2O_4S$   
Exact Mass: 380.0831





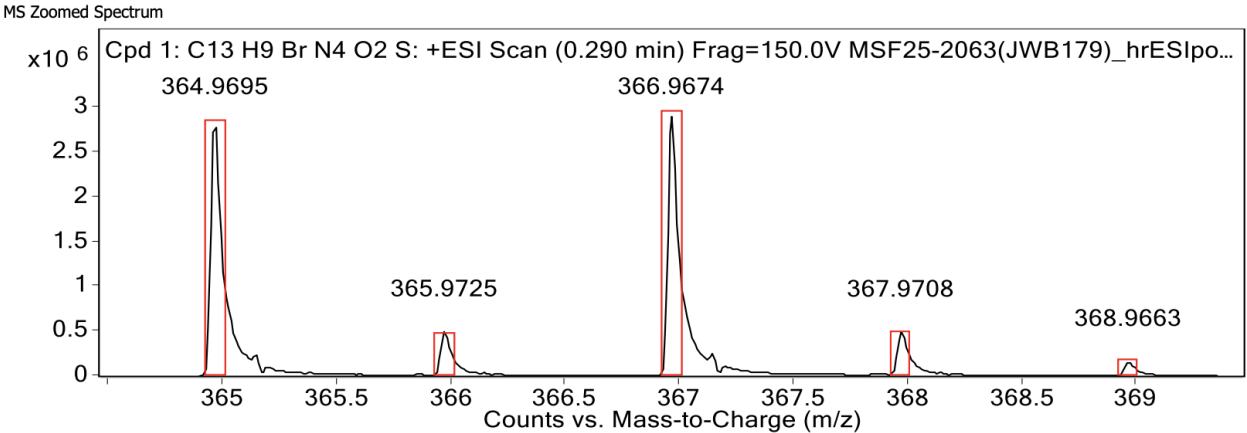
DML-I-87  
STANDARD FLUORINE PARAMETERS



# Target Compound Screening Report

## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

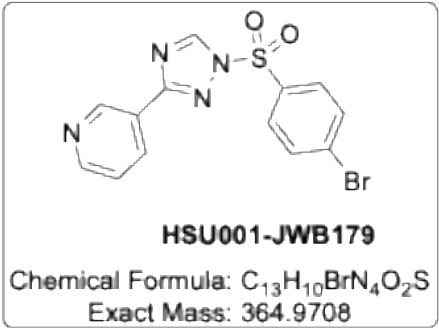
**Data File** MSF25-2063(JWB179)\_hrESIpos1.d    **Sample Name** 2063(JWB179)    **Comment** 2063(JWB179)  
**Position** P1-D1    **Instrument Name** 6530    **User Name**  
**Acq Method** FIA\_pos.m    **Acquired Time** 12/4/2025 10:43:53 AM    **DA Method** MSF.m



MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
364.9695	364.9702	1	2836650	C13H9BrN4O2S	(M+H)+	2.1
365.9725	365.9729	1	501572	C13H9BrN4O2S	(M+H)+	1.08
366.9674	366.9682	1	2928898	C13H9BrN4O2S	(M+H)+	2.13
367.9708	367.9708	1	497794	C13H9BrN4O2S	(M+H)+	0.07
368.9663	368.9666	1	170136	C13H9BrN4O2S	(M+H)+	0.66

--- End Of Report ---

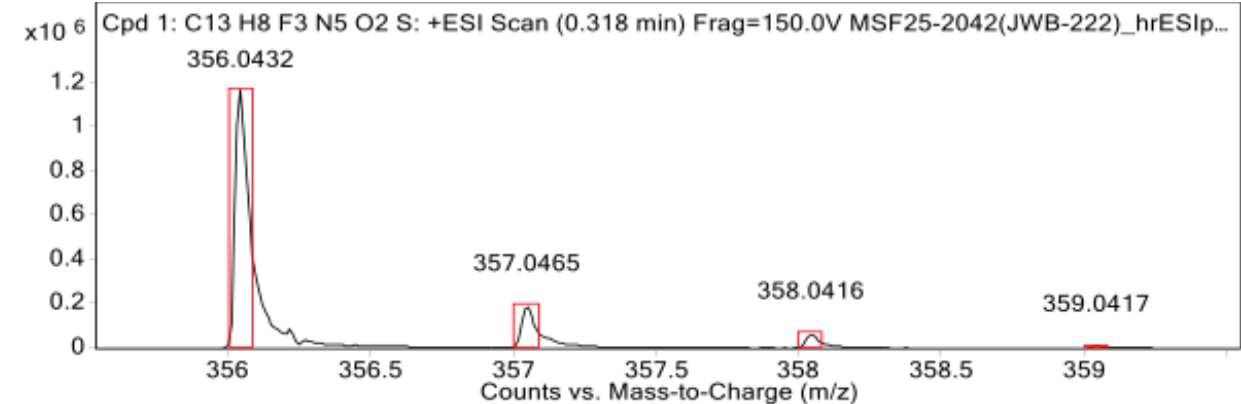


# Target Compound Screening Report

## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

Data File MSF25-2042(JWB-222)\_hrESIp0s1.d      Sample Name 2042(JWB-222)      Comment 2042(JWB-222)  
Position P1-A9      Instrument Name 6530      User Name  
Acq Method FIA\_pos.m      Acquired Time 11/24/2025 2:05:59 PM      DA Method MSF.m

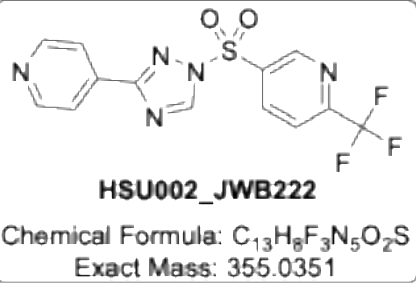
MS Zoomed Spectrum



### MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
356.0432	356.0424	1	1173525	C13H8F3N5O2S	(M+H)+	-2.26
357.0465	357.0449	1	192606	C13H8F3N5O2S	(M+H)+	-4.47
358.0416	358.0407	1	67591	C13H8F3N5O2S	(M+H)+	-2.58
359.0417	359.0422	1	9398	C13H8F3N5O2S	(M+H)+	1.33

--- End Of Report ---



Sample Name  
User Name  
Sample Type  
ACQ Method

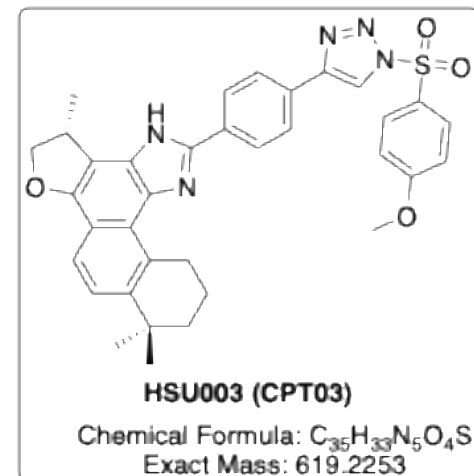
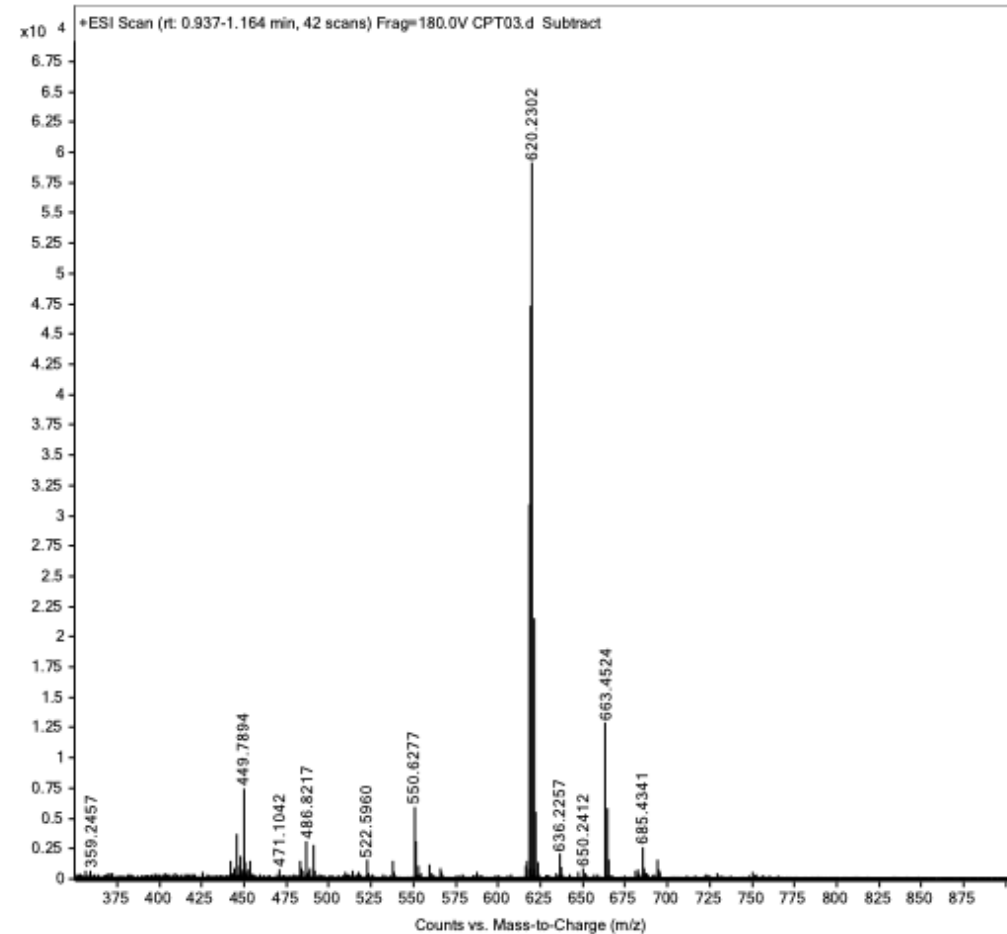
CPT03  
Sample  
Acq 100 ACN.m

Position  
Inj Vol  
IRM Calibration Status  
Comment

P1-A1  
0.1  
Success

Instrument Name  
InjPosition  
Data Filename  
Acquired Time

Instrument 1  
CPT03.d  
2/17/2022 10:19:34 AM (UTC-05:00)

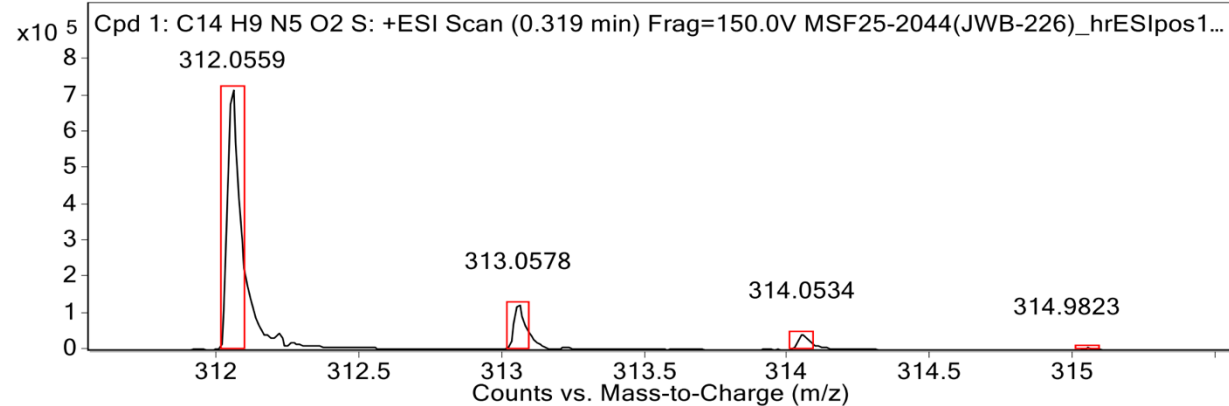


# Target Compound Screening Report

## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2044(JWB-226)\_hrESIpos1.d **Sample Name** 2044(JWB-226) **Comment** 2044(JWB-226)  
**Position** P1-B2 **Instrument Name** 6530 **User Name**  
**Acq Method** FIA\_pos.m **Acquired Time** 11/24/2025 2:10:01 PM **DA Method** MSF.m

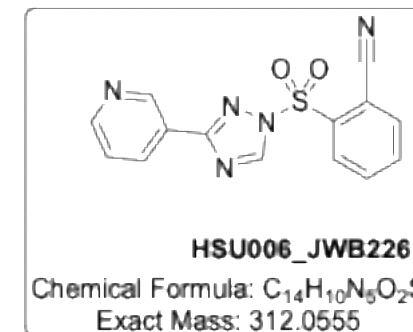
### MS Zoomed Spectrum



### MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
312.0559	312.0550	1	724030	C <sub>14</sub> H <sub>9</sub> N <sub>5</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	-2.87
313.0578	313.0575	1	126193	C <sub>14</sub> H <sub>9</sub> N <sub>5</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	-0.72
314.0534	314.0535	1	45243	C <sub>14</sub> H <sub>9</sub> N <sub>5</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	0.4
315.9266	316.0561	1	599	C <sub>14</sub> H <sub>9</sub> N <sub>5</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	409.74
316.8873	317.0575	1	484	C <sub>14</sub> H <sub>9</sub> N <sub>5</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	537.24

--- End Of Report ---

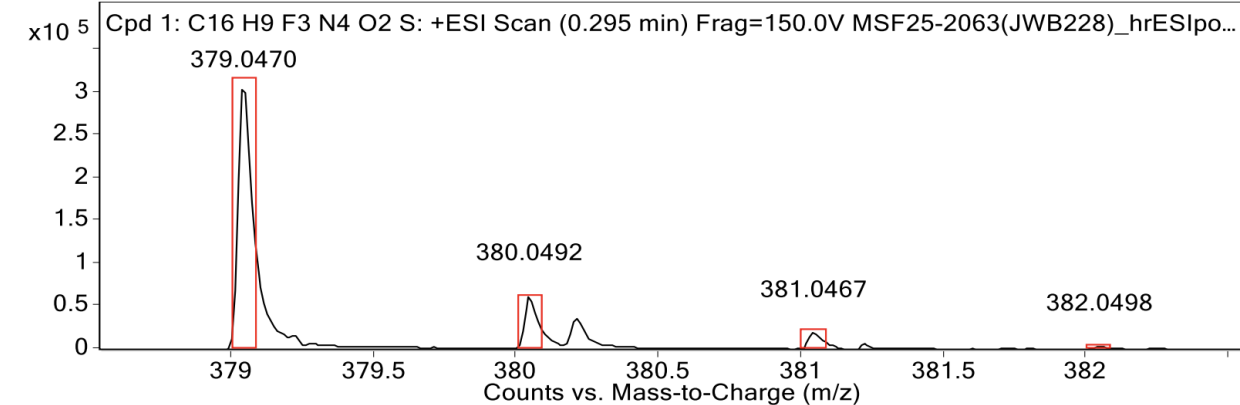


# Target Compound Screening Report

## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2063(JWB228)\_hrESIpos1.d      **Sample Name** 2063(JWB228)      **Comment** 2063(JWB228)  
**Position** P1-D3      **Instrument Name** 6530      **User Name**  
**Acq Method** FIA\_pos.m      **Acquired Time** 12/4/2025 10:47:54 AM      **DA Method** MSF.m

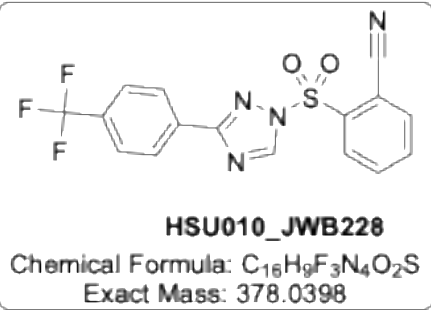
MS Zoomed Spectrum



### MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
302.1529			1704604			
379.0470	379.0471	1	315264	C <sub>16</sub> H <sub>9</sub> F <sub>3</sub> N <sub>4</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	0.19
380.0492	380.0499	1	62977	C <sub>16</sub> H <sub>9</sub> F <sub>3</sub> N <sub>4</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	1.74
381.0467	381.0461	1	19793	C <sub>16</sub> H <sub>9</sub> F <sub>3</sub> N <sub>4</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	-1.64
382.0498	382.0474	1	4104	C <sub>16</sub> H <sub>9</sub> F <sub>3</sub> N <sub>4</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	-6.17

--- End Of Report ---

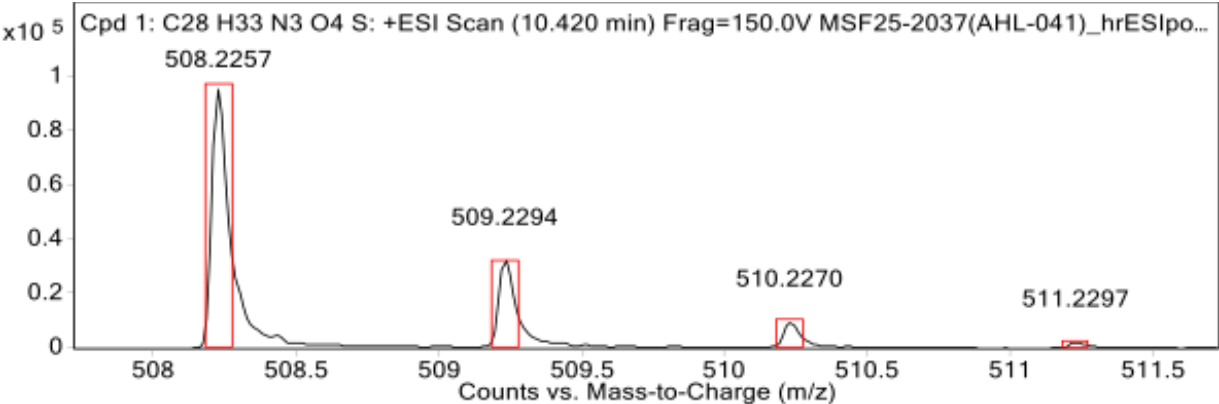


# Target Compound Screening Report

## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2037(AHL-041)\_hrESIposLC1.d      **Sample Name** 2037(AHL-041)      **Comment** 2037(AHL-041)  
**Position** P1-A4      **Instrument Name** 6530      **User Name**  
**Acq Method** LC\_C18\_pos\_jl.m      **Acquired Time** 11/24/2025 3:33:43 PM      **DA Method** MSF.m

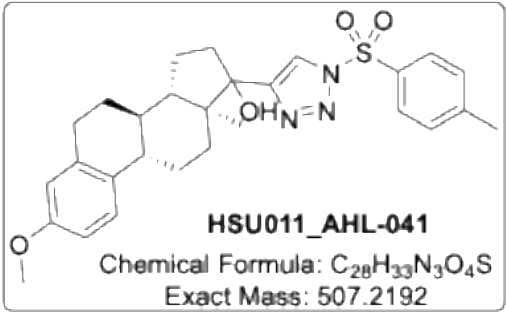
MS Zoomed Spectrum



### MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
508.2257	508.2265	1	95809	C <sub>28</sub> H <sub>33</sub> N <sub>3</sub> O <sub>4</sub> S	(M+H) <sup>+</sup>	1.44
509.2294	509.2295	1	32568	C <sub>28</sub> H <sub>33</sub> N <sub>3</sub> O <sub>4</sub> S	(M+H) <sup>+</sup>	0.2
510.2270	510.2280	1	9952	C <sub>28</sub> H <sub>33</sub> N <sub>3</sub> O <sub>4</sub> S	(M+H) <sup>+</sup>	2.08
511.2297	511.2289	1	2526	C <sub>28</sub> H <sub>33</sub> N <sub>3</sub> O <sub>4</sub> S	(M+H) <sup>+</sup>	-1.59
512.2447	512.2303	1	442	C <sub>28</sub> H <sub>33</sub> N <sub>3</sub> O <sub>4</sub> S	(M+H) <sup>+</sup>	-28.12

--- End Of Report ---

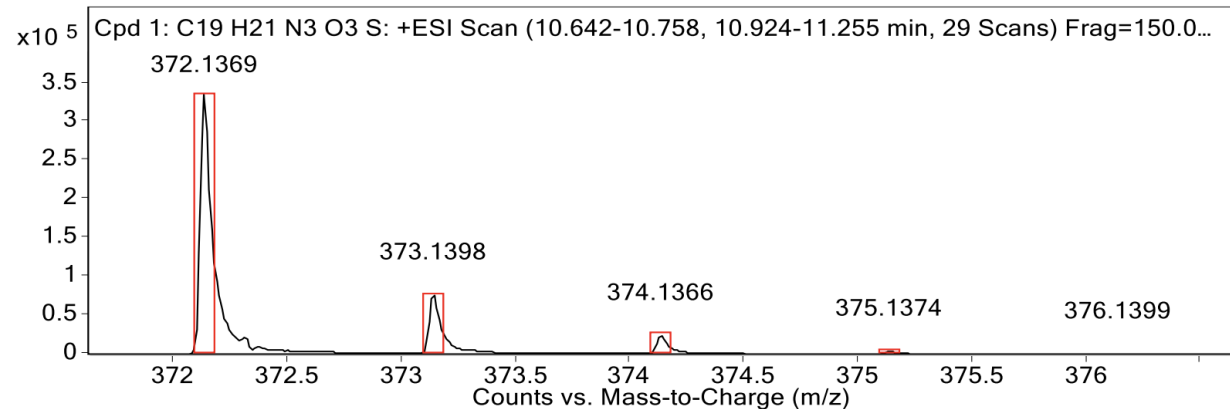


## Target Compound Screening Report

### Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2087(AHL011)\_hrESIposLC1.d **Sample Name** 2087(AHL011) **Comment** 2087(AHL011)  
**Position** P1-C5 **Instrument Name** 6530 **User Name**  
**Acq Method** LC\_C18\_pos\_jl.m **Acquired Time** 12/4/2025 5:48:51 PM **DA Method** MSF.m

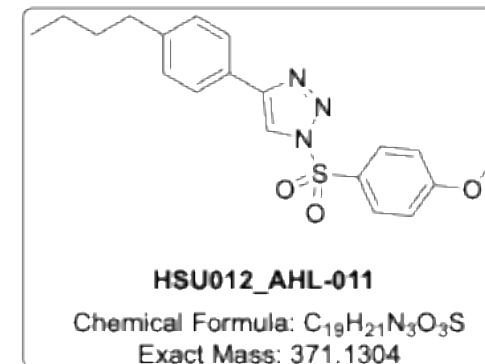
#### MS Zoomed Spectrum



#### MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
372.1369	372.1376	1	335642	C <sub>19</sub> H <sub>21</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	2.04
373.1398	373.1406	1	76837	C <sub>19</sub> H <sub>21</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	2.21
374.1366	374.1374	1	24236	C <sub>19</sub> H <sub>21</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	2.16
375.1374	375.1387	1	4089	C <sub>19</sub> H <sub>21</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	3.41
376.1399	376.1401	1	497	C <sub>19</sub> H <sub>21</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	0.66

--- End Of Report ---

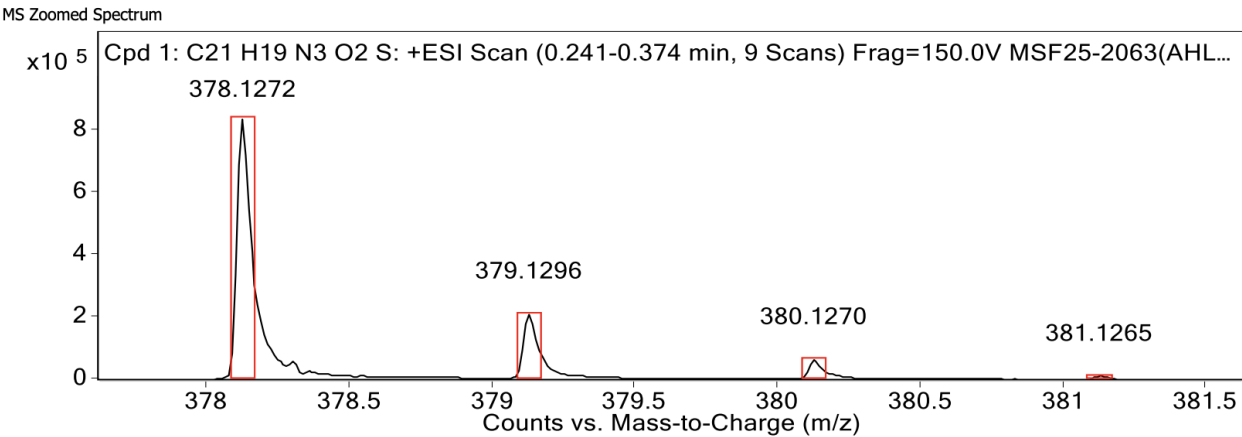




Target Compound Screening Report

Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

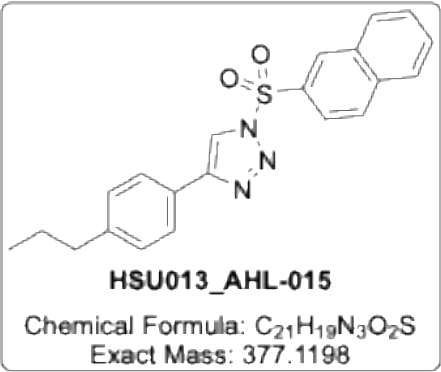
Data File MSF25-2063(AHL015)\_hrESIpos1.d      Sample Name 2063(AHL015)      Comment 2063(AHL015)  
Position P1-E2      Instrument Name 6530      User Name  
Acq Method FIA\_pos.m      Acquired Time 12/4/2025 11:04:01 AM      DA Method MSF.m



MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
372.1022			3559317			
378.1272	378.1271	1	834563	C21H19N3O2S	(M+H)+	-0.35
379.1296	379.1301	1	210880	C21H19N3O2S	(M+H)+	1.26
380.1270	380.1272	1	64437	C21H19N3O2S	(M+H)+	0.35
381.1265	381.1282	1	10923	C21H19N3O2S	(M+H)+	4.28

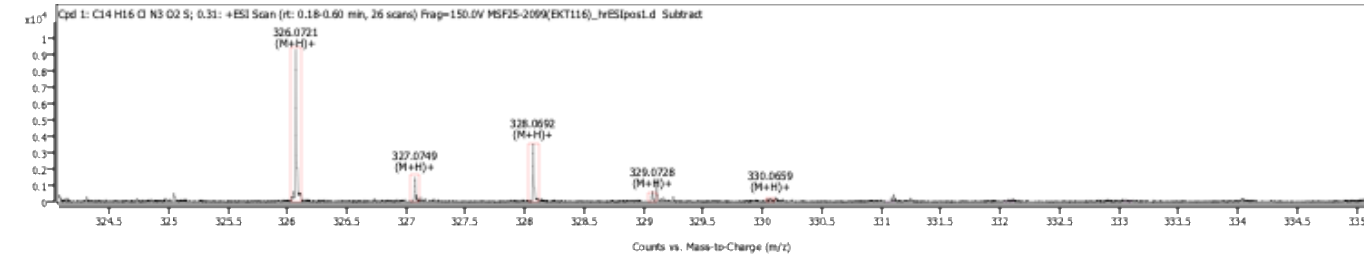
--- End Of Report ---



# Target Screening Report

## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

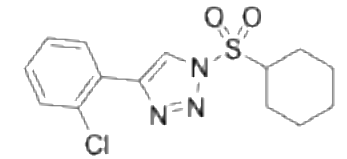
<b>Data File</b>	MSF25-2099(EKT116).hrESIspos1.d	<b>Sample Name</b>	2099(EKT116)	<b>Comment</b>	2099(EKT116)
<b>Position</b>	P1-P3	<b>Instrument Name</b>	Instrument 1		
<b>Acquisition Method</b>	FIA_Pos.m	<b>Acquired Time</b>	12/15/2025 5:16:48 PM (UTC-06:00)	<b>DA Method</b>	MSF.m



### Spectrum Peaks

Obs. m/z	Calc. m/z	Charge	Abund	Formula	Ion Species	Tgt Mass Error (PPM)
326.0721	326.0725	1	9354	C14H16ClN3O2S	(M+H)+	-1.17
327.0749	327.0753	1	1485	C14H16ClN3O2S	(M+H)+	-1.26
328.0692	328.0697	1	3574	C14H16ClN3O2S	(M+H)+	-1.57
329.0728	329.0724	1	632	C14H16ClN3O2S	(M+H)+	1.39
330.0659	330.0683	1	197	C14H16ClN3O2S	(M+H)+	-7.27
130.1590			8001873			

MassHunter Qual 10.0  
(End of Report)



**HSU014\_EKT116**

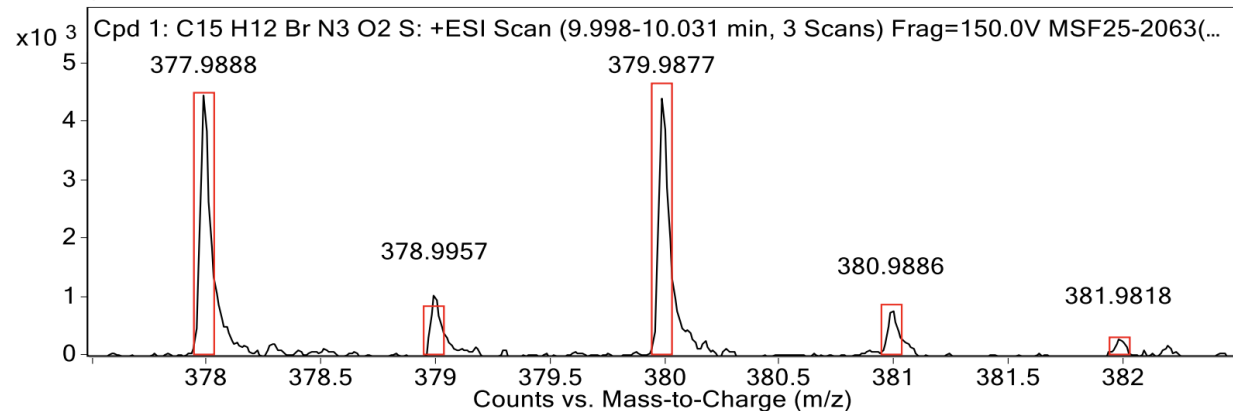
Chemical Formula: C<sub>14</sub>H<sub>16</sub>ClN<sub>3</sub>O<sub>2</sub>S  
Exact Mass: 325.0652

## Target Compound Screening Report

### Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2063(EKT151)\_hrESIposLC1.d  
**Position** P1-D8  
**Acq Method** LC\_C18\_pos\_jl.m  
**Sample Name** 2063(EKT151)  
**Instrument Name** 6530  
**Acquired Time** 12/4/2025 10:01:33 PM  
**Comment** 2063(EKT151)  
**User Name**  
**DA Method** MSF.m

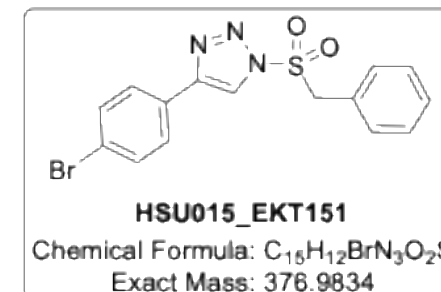
MS Zoomed Spectrum



MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
377.9888	377.9906	1	4475	C <sub>15</sub> H <sub>12</sub> BrN <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	4.78
378.9957	378.9935	1	1048	C <sub>15</sub> H <sub>12</sub> BrN <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	-5.96
379.9877	379.9886	1	4426	C <sub>15</sub> H <sub>12</sub> BrN <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	2.36
380.9886	380.9914	1	794	C <sub>15</sub> H <sub>12</sub> BrN <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	7.42
381.9818	381.9874	1	287	C <sub>15</sub> H <sub>12</sub> BrN <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	14.61
922.0021			59582			

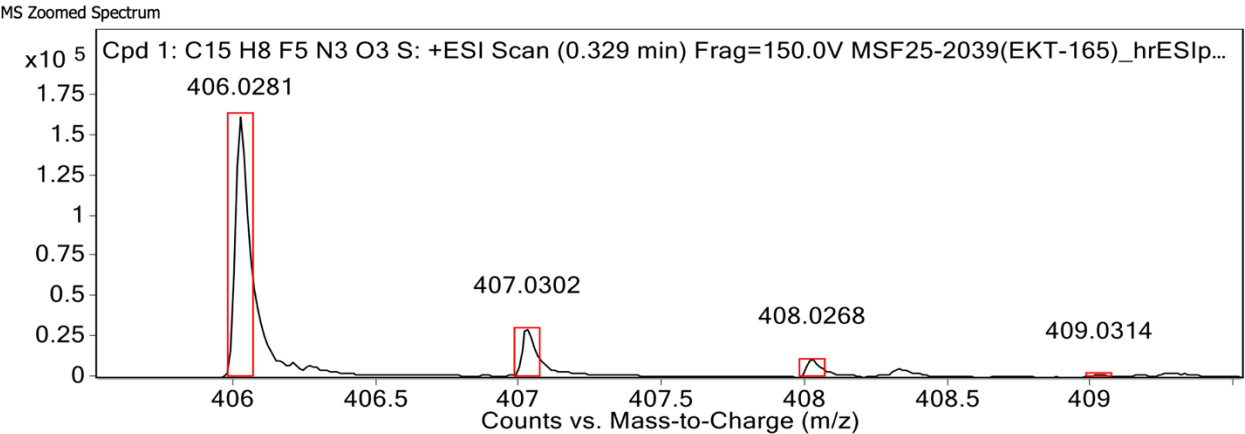
--- End Of Report ---



# Target Compound Screening Report

## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

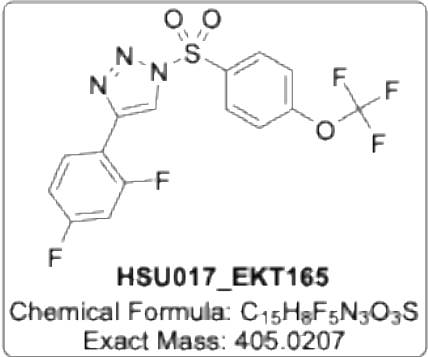
**Data File** MSF25-2039(EKT-165)\_hrESIpos1.d      **Sample Name** 2039(EKT-165)      **Comment** 2039(EKT-165)  
**Position** P1-A6      **Instrument Name** 6530      **User Name**  
**Acq Method** FIA\_pos.m      **Acquired Time** 11/24/2025 1:59:51 PM      **DA Method** MSF.m



### MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
406.0281	406.0279	1	161689	C <sub>15</sub> H <sub>8</sub> F <sub>5</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	-0.33
407.0302	407.0308	1	30496	C <sub>15</sub> H <sub>8</sub> F <sub>5</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	1.44
408.0268	408.0268	1	11438	C <sub>15</sub> H <sub>8</sub> F <sub>5</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	0.17
409.0314	409.0285	1	2202	C <sub>15</sub> H <sub>8</sub> F <sub>5</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	-7.02
409.9521	410.0297	1	261	C <sub>15</sub> H <sub>8</sub> F <sub>5</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	189.15

--- End Of Report ---

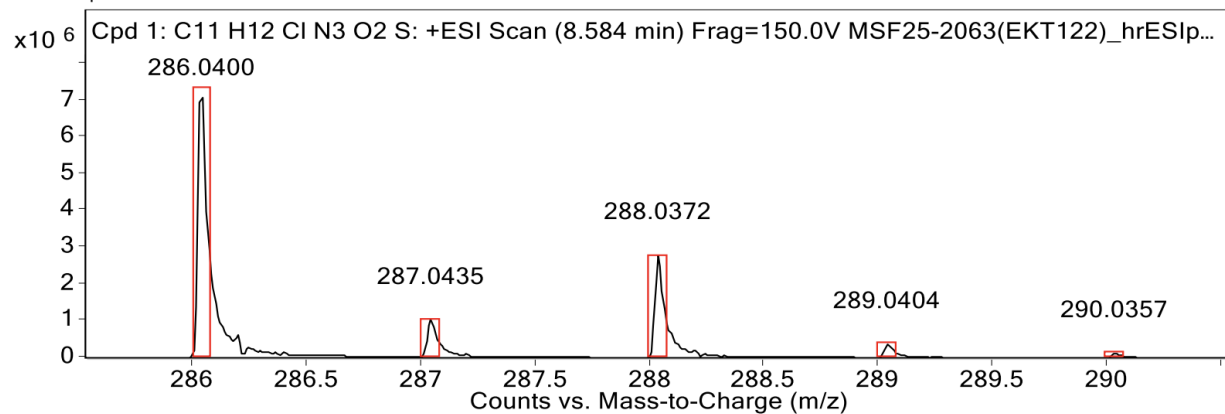


## Target Compound Screening Report

### Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2063(EKT122)\_hrESIposLC1.d **Sample Name** 2063(EKT122) **Comment** 2063(EKT122)  
**Position** P1-D7 **Instrument Name** 6530 **User Name**  
**Acq Method** LC\_C18\_pos\_jl.m **Acquired Time** 12/4/2025 1:36:37 PM **DA Method** MSF.m

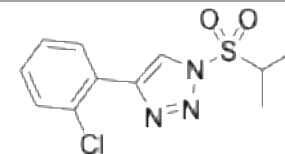
MS Zoomed Spectrum



MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
180.0313			9057772			
286.0400	286.0412	1	7222646	C <sub>11</sub> H <sub>12</sub> ClN <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	3.99
287.0435	287.0438	1	1044582	C <sub>11</sub> H <sub>12</sub> ClN <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	1.01
288.0372	288.0383	1	2811036	C <sub>11</sub> H <sub>12</sub> ClN <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	4.1
289.0404	289.0409	1	388823	C <sub>11</sub> H <sub>12</sub> ClN <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	1.73
290.0357	290.0363	1	141438	C <sub>11</sub> H <sub>12</sub> ClN <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	1.97

--- End Of Report ---



**HSU019\_EKT122**

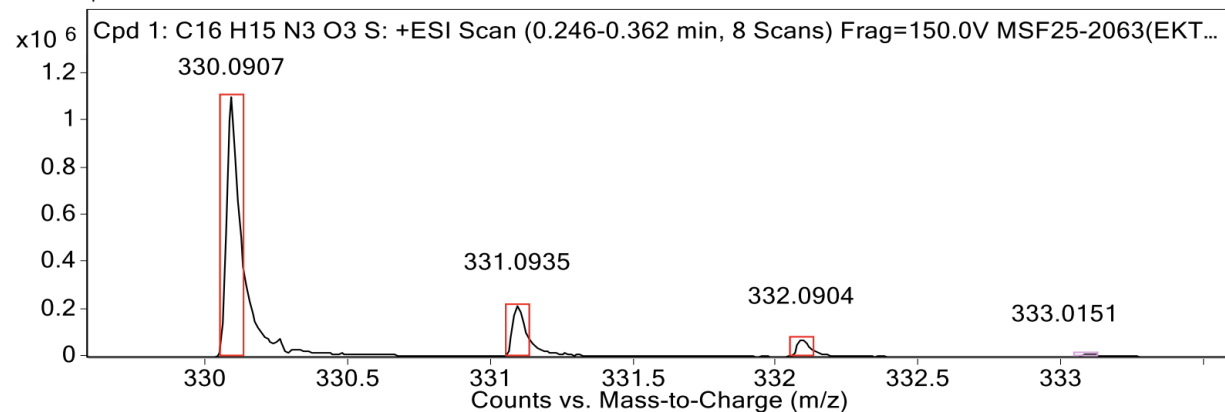
Chemical Formula: C<sub>11</sub>H<sub>12</sub>ClN<sub>3</sub>O<sub>2</sub>S  
Exact Mass: 285.0339

# Target Compound Screening Report

## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2063(EKT080)\_hrESIpos1.d **Sample Name** 2063(EKT080) **Comment** 2063(EKT080)  
**Position** P1-D6 **Instrument Name** 6530 **User Name**  
**Acq Method** FIA\_pos.m **Acquired Time** 12/4/2025 10:53:57 AM **DA Method** MSF.m

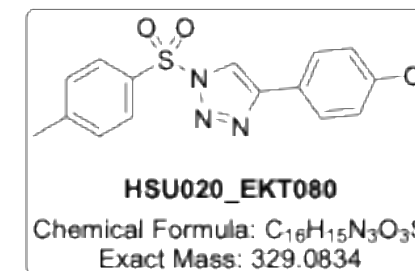
MS Zoomed Spectrum



MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
330.0907	330.0907	1	1106115	C <sub>16</sub> H <sub>15</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	-0.07
331.0935	331.0936	1	220098	C <sub>16</sub> H <sub>15</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	0.11
332.0904	332.0898	1	75988	C <sub>16</sub> H <sub>15</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	-1.58
333.0151	333.0914	1	825	C <sub>16</sub> H <sub>15</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	228.98

--- End Of Report ---

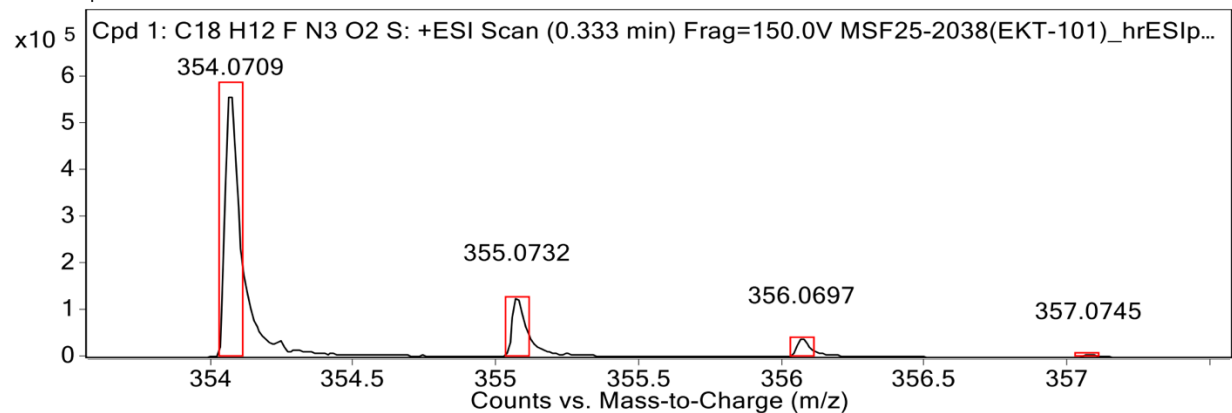


## Target Compound Screening Report

### Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

Data File MSF25-2038(EKT-101)\_hrESIpos1.d Sample Name 2038(EKT-101) Comment 2038(EKT-101)  
Position P1-A5 Instrument Name 6530 User Name  
Acq Method FIA\_pos.m Acquired Time 11/24/2025 1:57:48 PM DA Method MSF.m

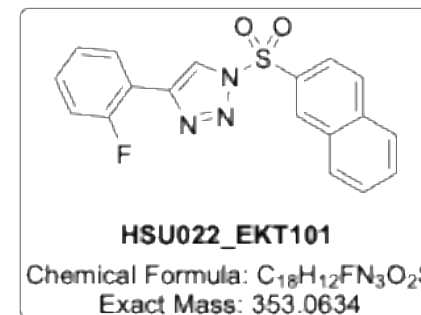
MS Zoomed Spectrum



#### MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
354.0709	354.0707	1	581540	C <sub>18</sub> H <sub>12</sub> FN <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	-0.48
355.0732	355.0736	1	131547	C <sub>18</sub> H <sub>12</sub> FN <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	1.07
356.0697	356.0701	1	40099	C <sub>18</sub> H <sub>12</sub> FN <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	1.19
357.0745	357.0714	1	6318	C <sub>18</sub> H <sub>12</sub> FN <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	-8.55
364.0203			1881949			

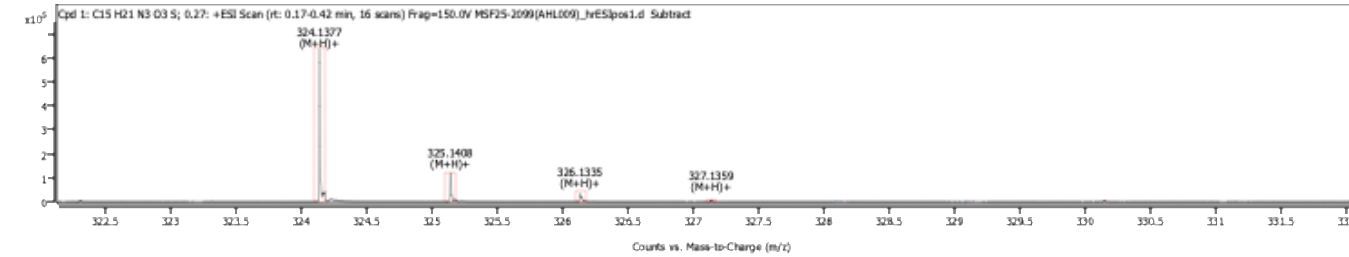
--- End Of Report ---



# Target Screening Report

## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

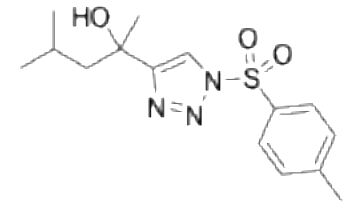
<b>Data File</b>	MSF25-2099(AHL009)_hrES(pos1.d	<b>Sample Name</b>	2099(AHL009)	<b>Comment</b>	2099(AHL009)
<b>Position</b>	P1-E6	<b>Instrument Name</b>	Instrument 1		
<b>Acquisition Method</b>	FIA_Pos.m	<b>Acquired Time</b>	12/15/2025 5:07:02 PM (UTC-06:00)	<b>DA Method</b>	MSF.m



### Spectrum Peaks

Obs. m/z	Calc. m/z	Charge	Abund	Formula	Ion Species	Tgt Mass Error (PPM)
324.1377	324.1376	1	940805	C15H21N3O3S	(M+H)+	0.15
325.1408	325.1405	1	122866	C15H21N3O3S	(M+H)+	0.81
326.1335	326.1366	1	34437	C15H21N3O3S	(M+H)+	-9.34
327.1359	327.1383	1	5528	C15H21N3O3S	(M+H)+	-7.10
130.1590			1107853			

MassHunter Qual 10.0  
(End of Report)



**HSU024\_AHL-009**  
Chemical Formula: C<sub>15</sub>H<sub>21</sub>N<sub>3</sub>O<sub>3</sub>S  
Exact Mass: 323.1304

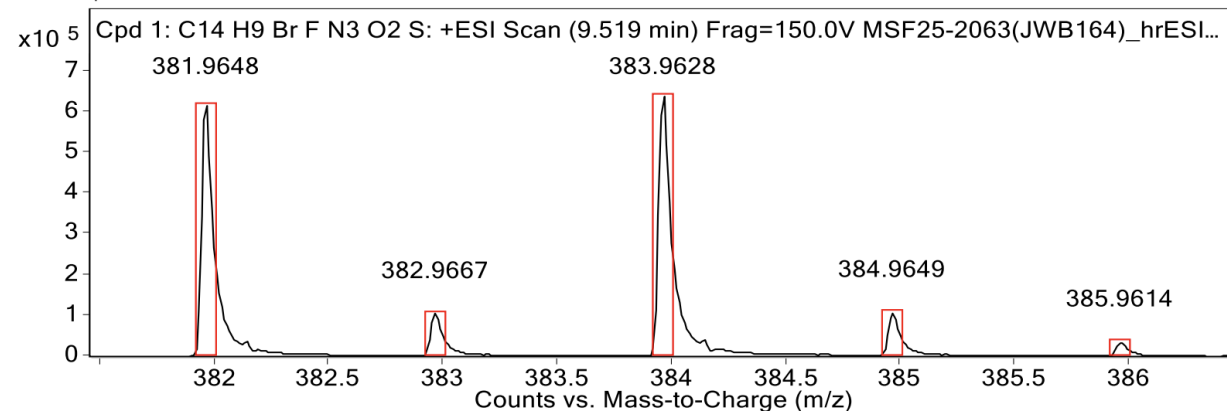


# Target Compound Screening Report

## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2063(JWB164)\_hrESIposLC1.d **Sample Name** 2063(JWB164) **Comment** 2063(JWB164)  
**Position** P1-C9 **Instrument Name** 6530 **User Name**  
**Acq Method** LC\_C18\_pos\_jl.m **Acquired Time** 12/4/2025 12:50:41 PM **DA Method** MSF.m

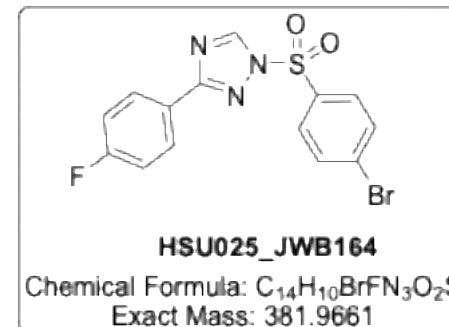
### MS Zoomed Spectrum



### MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
381.9648	381.9656	1	619445	C14H9BrFN3O2S	(M+H)+	1.99
382.9667	382.9684	1	105565	C14H9BrFN3O2S	(M+H)+	4.22
383.9628	383.9636	1	642132	C14H9BrFN3O2S	(M+H)+	1.92
384.9649	384.9663	1	108586	C14H9BrFN3O2S	(M+H)+	3.5
385.9614	385.9621	1	36185	C14H9BrFN3O2S	(M+H)+	1.79
386.9641	386.9638	1	4732	C14H9BrFN3O2S	(M+H)+	-0.84

--- End Of Report ---

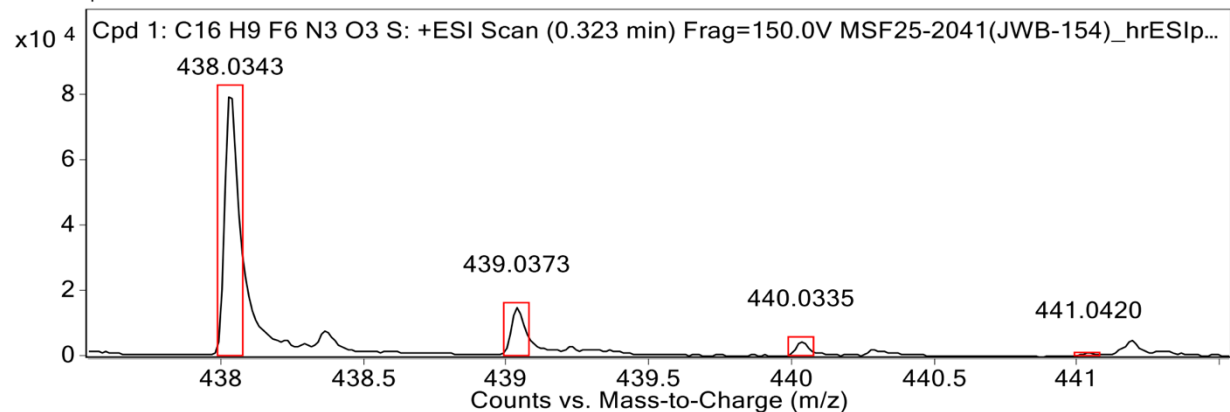


## Target Compound Screening Report

### Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

Data File MSF25-2041(JWB-154)\_hrESIpos1.d Sample Name 2041(JWB-154) Comment 2041(JWB-154)  
Position P1-A8 Instrument Name 6530 User Name  
Acq Method FIA\_pos.m Acquired Time 11/24/2025 2:03:59 PM DA Method MSF.m

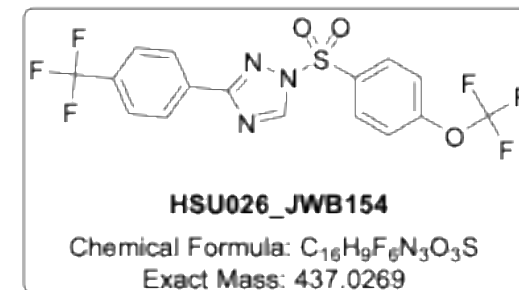
#### MS Zoomed Spectrum



#### MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
436.3627			98431			
438.0343	438.0342	1	82450	C <sub>16</sub> H <sub>9</sub> F <sub>6</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	-0.35
439.0373	439.0370	1	15163	C <sub>16</sub> H <sub>9</sub> F <sub>6</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	-0.66
440.0335	440.0333	1	4733	C <sub>16</sub> H <sub>9</sub> F <sub>6</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	-0.41
441.0420	441.0348	1	1091	C <sub>16</sub> H <sub>9</sub> F <sub>6</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	-16.3

--- End Of Report ---

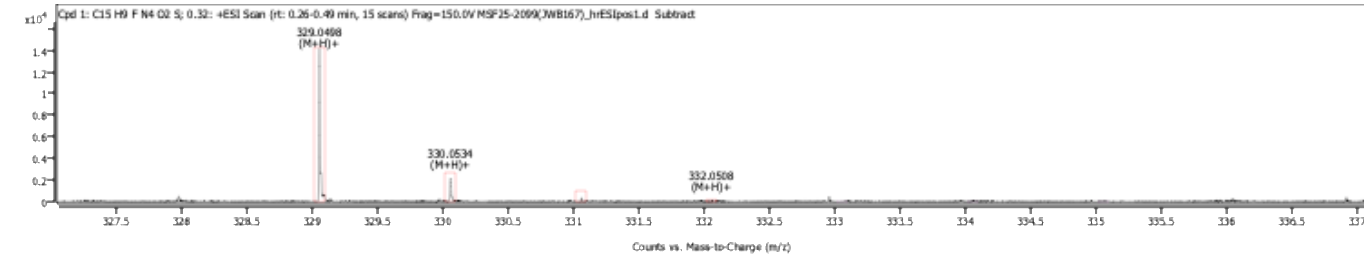


# Target Screening Report



## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

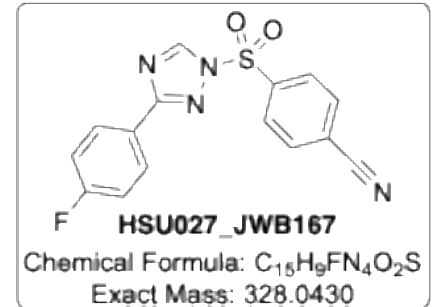
Data File	MSF25-2099_JWB167_hrESIspos1.d	Sample Name	2099_JWB167	Comment	2099_JWB167
Position	P1-F5	Instrument Name	Instrument 1		
Acquisition Method	FIA_Pos.m	Acquired Time	12/15/2025 5:20:03 PM (UTC-06:00)	DA Method	MSF.m



### Spectrum Peaks

Obs. m/z	Calc. m/z	Charge	Abund	Formula	Ion Species	Tgt Mass Error (PPM)
329.0498	329.0503	1	14190	C15H9FN4O2S	(M+H)+	-1.52
330.0534	330.0530	1	2114	C15H9FN4O2S	(M+H)+	1.16
332.0508	332.0505	1	51	C15H9FN4O2S	(M+H)+	0.93
130.1990			6756261			

MassHunter Qual 10.0  
(End of Report)

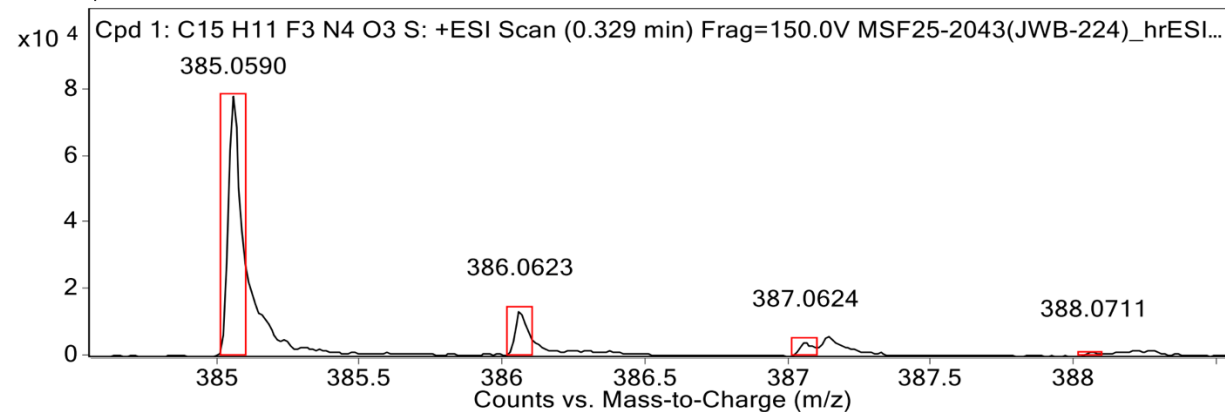


## Target Compound Screening Report

### Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2043(JWB-224)\_hrESIpos1.d **Sample Name** 2043(JWB-224) **Comment** 2043(JWB-224)  
**Position** P1-B1 **Instrument Name** 6530 **User Name**  
**Acq Method** FIA\_pos.m **Acquired Time** 11/24/2025 2:08:00 PM **DA Method** MSF.m

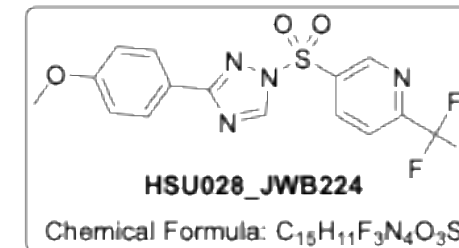
#### MS Zoomed Spectrum



#### MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
385.0590	385.0577	1	78326	C <sub>15</sub> H <sub>11</sub> F <sub>3</sub> N <sub>4</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	-3.57
386.0623	386.0604	1	13764	C <sub>15</sub> H <sub>11</sub> F <sub>3</sub> N <sub>4</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	-4.87
387.0624	387.0566	1	4353	C <sub>15</sub> H <sub>11</sub> F <sub>3</sub> N <sub>4</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	-15.07
388.0711	388.0581	1	1326	C <sub>15</sub> H <sub>11</sub> F <sub>3</sub> N <sub>4</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	-33.48
389.1370	389.0593	1	1675	C <sub>15</sub> H <sub>11</sub> F <sub>3</sub> N <sub>4</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	-199.73

--- End Of Report ---

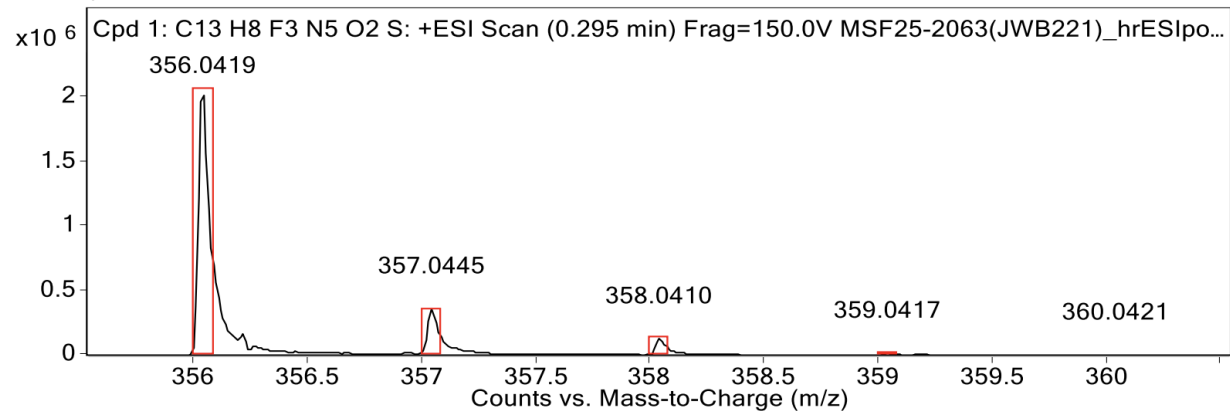


## Target Compound Screening Report

### Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2063(JWB221)\_hrESIpos1.d **Sample Name** 2063(JWB221) **Comment** 2063(JWB221)  
**Position** P1-D2 **Instrument Name** 6530 **User Name**  
**Acq Method** FIA\_pos.m **Acquired Time** 12/4/2025 10:45:54 AM **DA Method** MSF.m

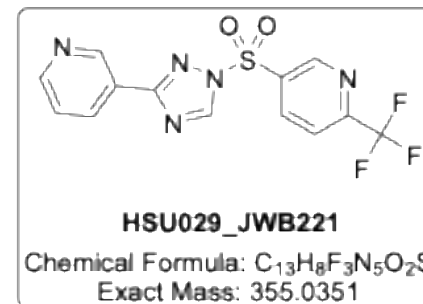
MS Zoomed Spectrum



MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
356.0419	356.0424	1	2054974	C <sub>13</sub> H <sub>8</sub> F <sub>3</sub> N <sub>5</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	1.31
357.0445	357.0449	1	360844	C <sub>13</sub> H <sub>8</sub> F <sub>3</sub> N <sub>5</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	0.89
358.0410	358.0407	1	127975	C <sub>13</sub> H <sub>8</sub> F <sub>3</sub> N <sub>5</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	-0.94
359.0417	359.0422	1	16257	C <sub>13</sub> H <sub>8</sub> F <sub>3</sub> N <sub>5</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	1.35
360.0421	360.0433	1	2193	C <sub>13</sub> H <sub>8</sub> F <sub>3</sub> N <sub>5</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	3.32

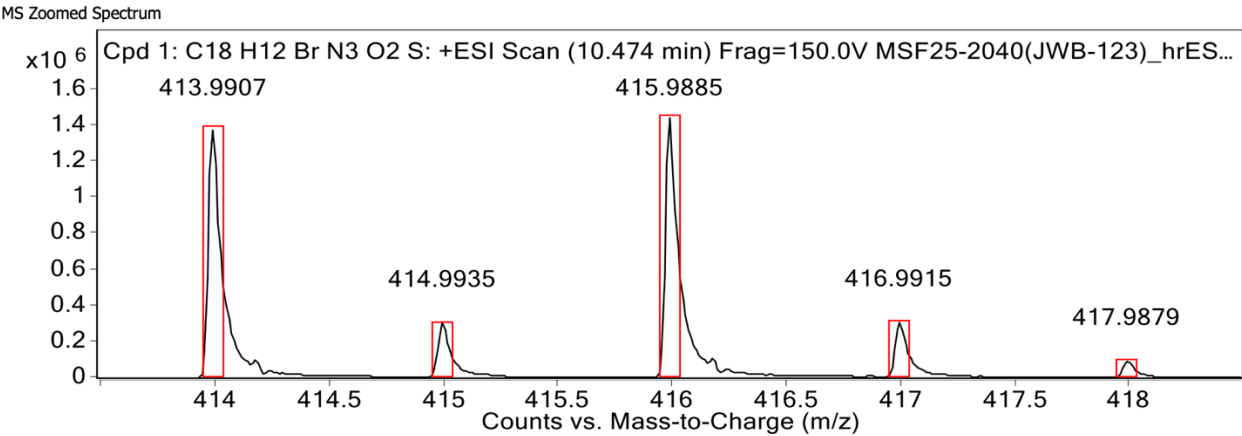
--- End Of Report ---



# Target Compound Screening Report

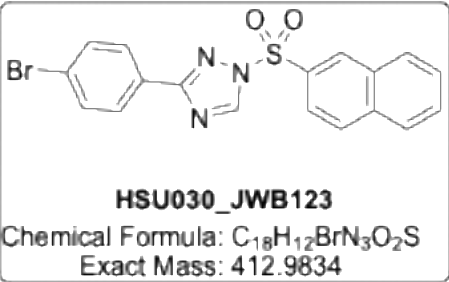
## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

<b>Data File</b>	MSF25-2040(JWB-123)_hrESIposLC1.d	<b>Sample Name</b>	Unavailable	<b>Comment</b>	Sample information is unavailable
<b>Position</b>	Unavailable	<b>Instrument Name</b>	Unavailable	<b>User Name</b>	Unavailable
<b>Acq Method</b>		<b>Acquired Time</b>	Unavailable	<b>DA Method</b>	MSF.m



MS Spectrum Peak List						
Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
413.9907	413.9906	1	1377179	C <sub>18</sub> H <sub>12</sub> BrN <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	-0.06
414.9935	414.9935	1	311013	C <sub>18</sub> H <sub>12</sub> BrN <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	0.14
415.9885	415.9887	1	1446529	C <sub>18</sub> H <sub>12</sub> BrN <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	0.37
416.9915	416.9915	1	314084	C <sub>18</sub> H <sub>12</sub> BrN <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	0.05
417.9879	417.9881	1	99280	C <sub>18</sub> H <sub>12</sub> BrN <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	0.58

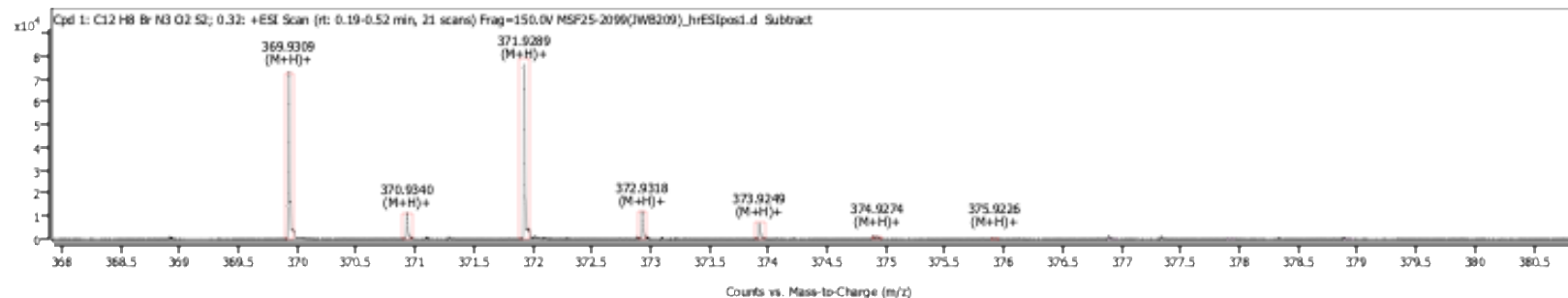
--- End Of Report ---



# Target Screening Report

## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

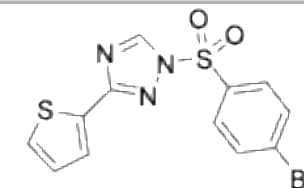
<b>Data File</b>	MSF25-2099(JWB209)_hrESIpos1.d	<b>Sample Name</b>	2099(JWB209)	<b>Comment</b>	2099(JWB209)
<b>Position</b>	P1-F6	<b>Instrument Name</b>	Instrument 1		
<b>Acquisition Method</b>	FIA_Pos.m	<b>Acquired Time</b>	12/15/2025 5:21:40 PM (UTC-06:00)	<b>DA Method</b>	MSF.m



### Spectrum Peaks

Obs. m/z	Calc. m/z	Charge	Abund	Formula	Ion Species	Tgt Mass Error (PPM)
369.9309	369.9314	1	74221	C12H8BrN3O2S2	(M+H)+	-1.48
370.9340	370.9340	1	11803	C12H8BrN3O2S2	(M+H)+	0.05
371.9289	371.9293	1	77349	C12H8BrN3O2S2	(M+H)+	-1.03
372.9318	372.9318	1	12417	C12H8BrN3O2S2	(M+H)+	-0.07
373.9249	373.9255	1	6098	C12H8BrN3O2S2	(M+H)+	-4.35
374.9274	374.9286	1	1213	C12H8BrN3O2S2	(M+H)+	-3.03
375.9226	375.9249	1	216	C12H8BrN3O2S2	(M+H)+	-6.13
130.1589			10468549			

MassHunter Qual 10.0  
(End of Report)



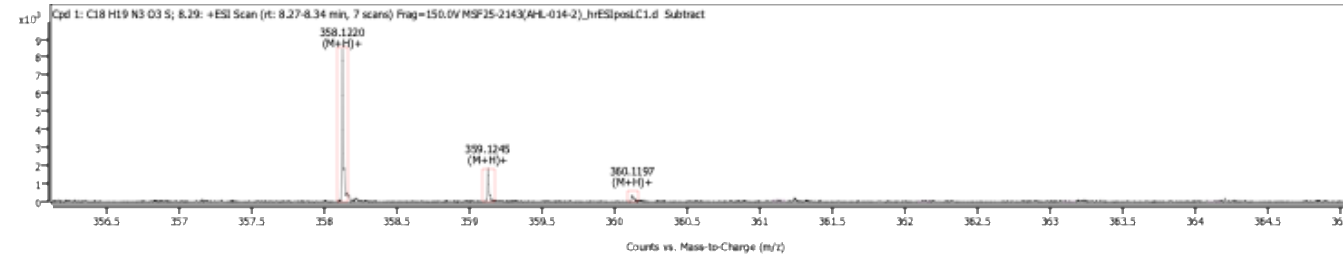
**HSU032\_JWB209**

Chemical Formula: C<sub>12</sub>H<sub>8</sub>BrN<sub>3</sub>O<sub>2</sub>S<sub>2</sub>  
Exact Mass: 368.9241

# Target Screening Report

## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

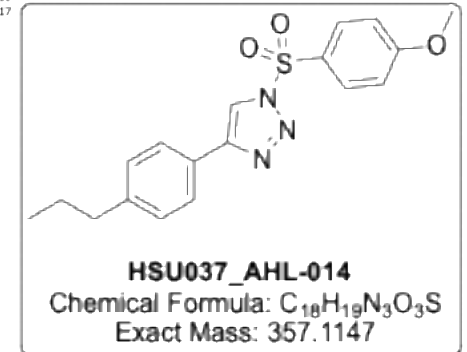
Data File: MSP25-2143/AHL-014-2)\_hrESIPosLC1.d  
 Sample Name:   
 Position:   
 Instrument Name:   
 Acquired Time: 12/19/2025 2:56:13 PM (UTC-06:00)  
 DA Method: MSP.m  
 Acquisition Method:   
 Comment:   
 Cpd 1: C18 H19 N3 O3 S; 8.29: +ESI Scan (rt: 8.27-8.34 min, 7 scans) Frag=150.0V MSP25-2143/AHL-014-2)\_hrESIPosLC1.d Subtract



### Spectrum Peaks

Obs. m/z	Calc. m/z	Charge	Abund	Formula	Ion Species	Tgt Mass Error (PPM)
358.1220	358.1220	1	8499	C18H19NO3S	(M+H)+	0.01
359.1245	359.1245	1	1859	C18H19NO3S	(M+H)+	-1.18
360.1197	360.1216	1	377	C18H19NO3S	(M+H)+	-5.17

MassHunter Qual 10.0  
 (End of Report)



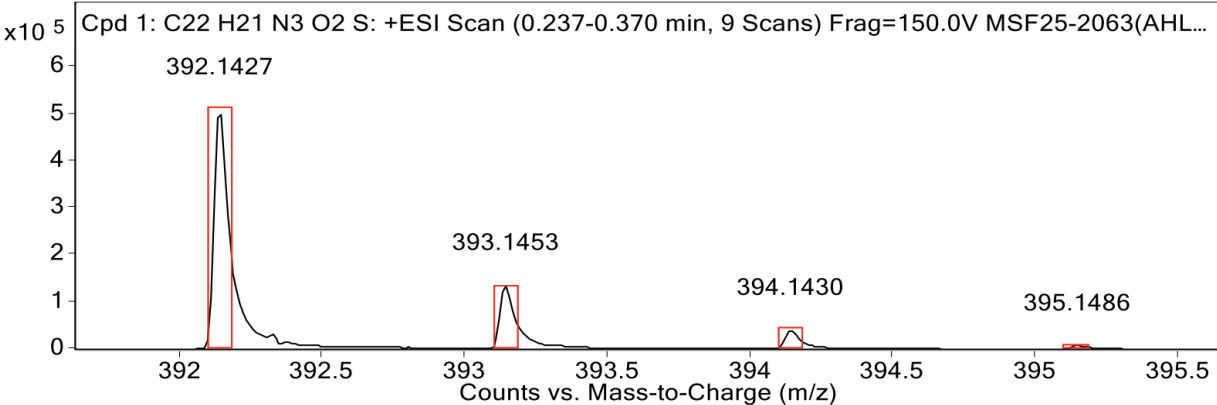


# Target Compound Screening Report

## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2063(AHL017)\_hrESIpos1.d      **Sample Name** 2063(AHL017)      **Comment** 2063(AHL017)  
**Position** P1-E3      **Instrument Name** 6530      **User Name**  
**Acq Method** FIA\_pos.m      **Acquired Time** 12/4/2025 11:06:02 AM      **DA Method** MSF.m

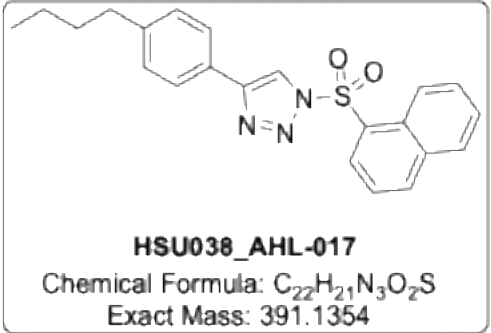
MS Zoomed Spectrum



MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
386.1182			1176102			
392.1427	392.1427	1	510087	C <sub>22</sub> H <sub>21</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	0.15
393.1453	393.1457	1	136366	C <sub>22</sub> H <sub>21</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	1.18
394.1430	394.1430	1	40763	C <sub>22</sub> H <sub>21</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	0
395.1486	395.1440	1	7586	C <sub>22</sub> H <sub>21</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	-11.75

--- End Of Report ---

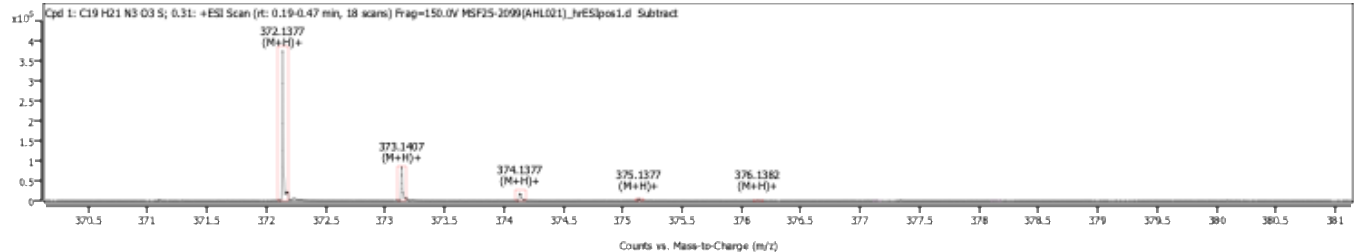


Target Screening Report



Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

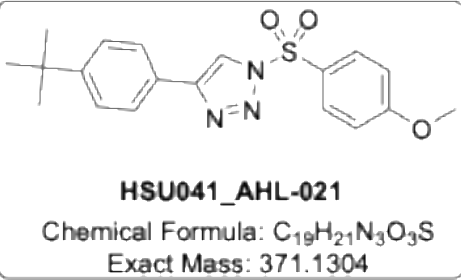
Data File	MSF25-2099(AHL021)_hrES(pos1.d	Sample Name	2099(AHL021)	Comment	2099(AHL021)
Position	P1-E7	Instrument Name	Instrument 1		
Acquisition Method	FIA_Pos.m	Acquired Time	12/15/2025 5:08:39 PM (UTC-06:00)	DA Method	MSF.m



Spectrum Peaks

Obs. m/z	Calc. m/z	Charge	Abund	Formula	Ion Species	Tgt Mass Error (PPM)
372.1377	372.1376	1	385128	C19H21N3O3S	(M+H)+	0.05
373.1407	373.1406	1	86521	C19H21N3O3S	(M+H)+	0.36
374.1377	374.1374	1	19401	C19H21N3O3S	(M+H)+	0.61
375.1377	375.1387	1	4561	C19H21N3O3S	(M+H)+	-2.62
376.1382	376.1401	1	749	C19H21N3O3S	(M+H)+	-5.04
130.1990			9895309			

MassHunter Qual 10.0  
(End of Report)

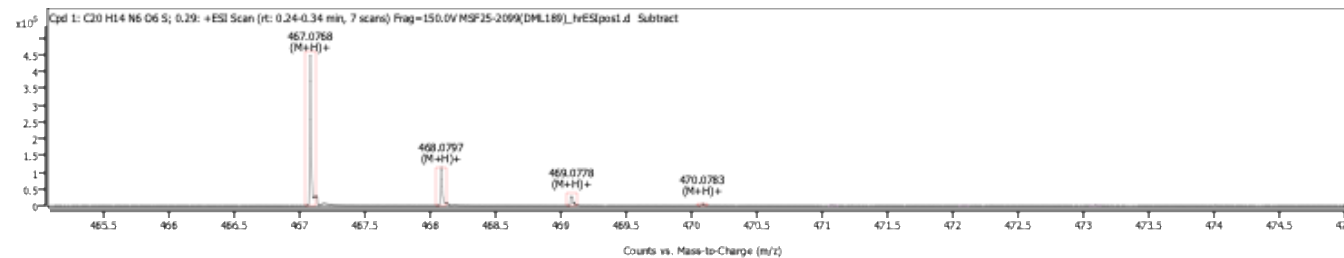


# Target Screening Report



## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

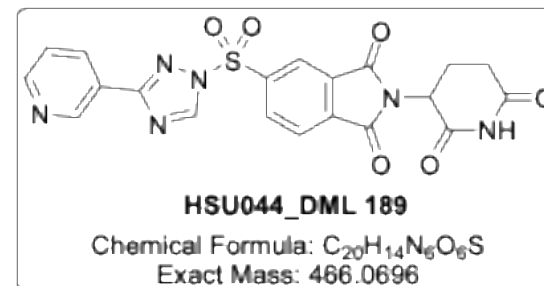
<b>Data File</b>	MSF25-2099(DML189)_hrESI.post.d	<b>Sample Name</b>	2099(DML189)	<b>Comment</b>	2099(DML189)
<b>Position</b>	P1-F4	<b>Instrument Name</b>	Instrument 1		
<b>Acquisition Method</b>	FIA_Pos.m	<b>Acquired Time</b>	12/15/2023 5:18:26 PM (UTC-06:00)	<b>DA Method</b>	MSF.m



### Spectrum Peaks

Obs. m/z	Calc. m/z	Charge	Abund	Formula	Ion Species	Tgt Mass Error (PPM)
467.0768	467.0768	1	456293	C20H14N6O6S	(M+H)+	-0.09
468.0797	468.0795	1	116254	C20H14N6O6S	(M+H)+	0.40
469.0778	469.0771	1	26809	C20H14N6O6S	(M+H)+	1.42
470.0783	470.0784	1	5894	C20H14N6O6S	(M+H)+	-0.09
130.1589			11053639			

MassHunter Qual 10.0  
(End of Report)

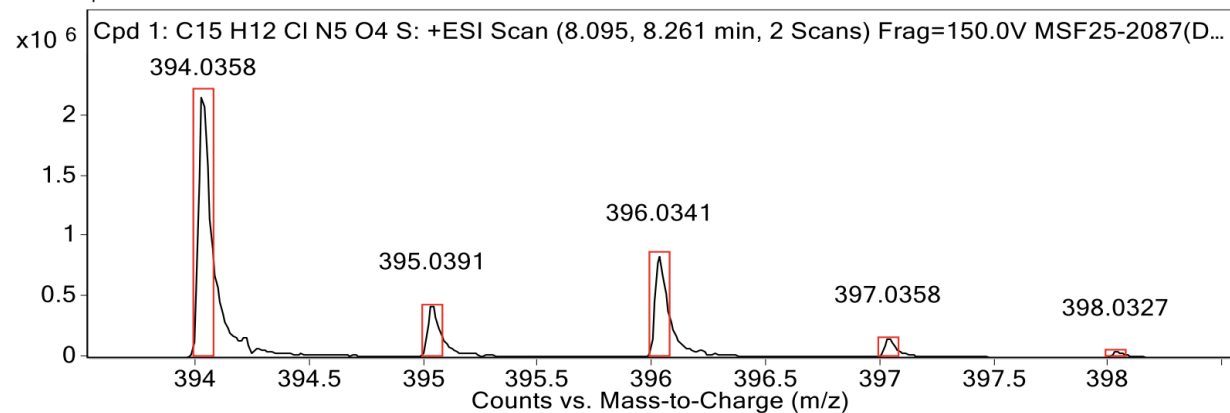


## Target Compound Screening Report

### Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2087(DML367)\_hrESIposLC1.d **Sample Name** 2087(DML367) **Comment** 2087(DML367)  
**Position** P1-D5 **Instrument Name** 6530 **User Name**  
**Acq Method** LC\_C18\_pos\_jl.m **Acquired Time** 12/4/2025 9:15:35 PM **DA Method** MSF.m

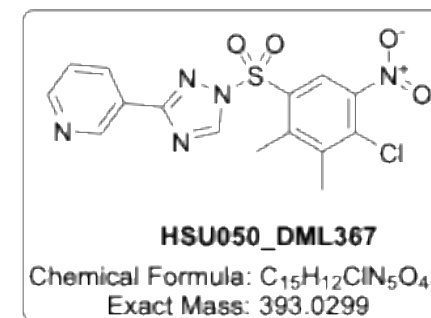
MS Zoomed Spectrum



MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
394.0358	394.0371	1	2211590	C <sub>15</sub> H <sub>12</sub> ClN <sub>5</sub> O <sub>4</sub> S	(M+H) <sup>+</sup>	3.32
395.0391	395.0397	1	437564	C <sub>15</sub> H <sub>12</sub> ClN <sub>5</sub> O <sub>4</sub> S	(M+H) <sup>+</sup>	1.76
396.0341	396.0345	1	840389	C <sub>15</sub> H <sub>12</sub> ClN <sub>5</sub> O <sub>4</sub> S	(M+H) <sup>+</sup>	1.19
397.0358	397.0369	1	162511	C <sub>15</sub> H <sub>12</sub> ClN <sub>5</sub> O <sub>4</sub> S	(M+H) <sup>+</sup>	2.95
398.0327	398.0336	1	53848	C <sub>15</sub> H <sub>12</sub> ClN <sub>5</sub> O <sub>4</sub> S	(M+H) <sup>+</sup>	2.23

--- End Of Report ---

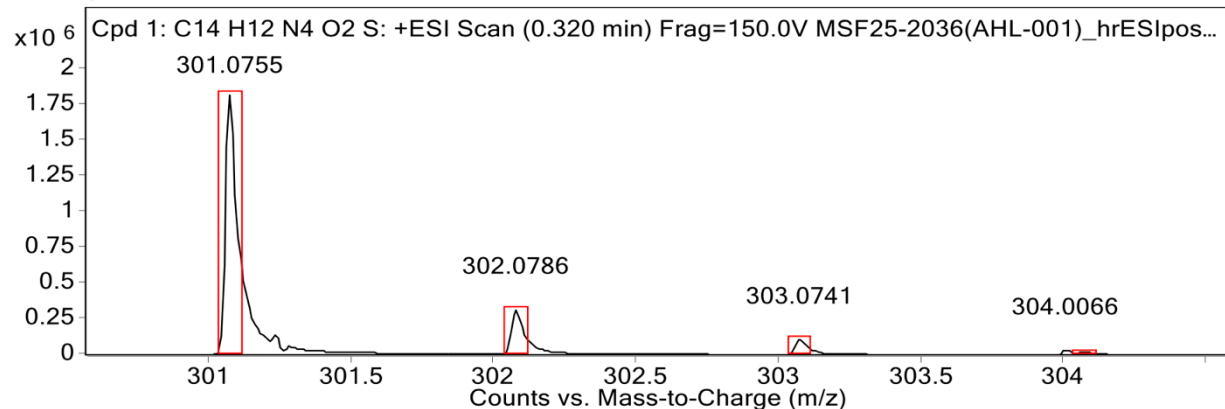


## Target Compound Screening Report

### Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2036(AHL-001)\_hrESIpos1.d **Sample Name** 2036(AHL-001) **Comment** 2036(AHL-001)  
**Position** P1-A3 **Instrument Name** 6530 **User Name**  
**Acq Method** FIA\_pos.m **Acquired Time** 11/24/2025 1:53:43 PM **DA Method** MSF.m

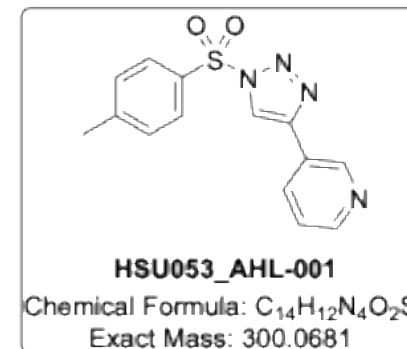
#### MS Zoomed Spectrum



#### MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
301.0755	301.0754	1	1815759	C <sub>14</sub> H <sub>12</sub> N <sub>4</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	-0.28
302.0786	302.0781	1	320167	C <sub>14</sub> H <sub>12</sub> N <sub>4</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	-1.76
303.0741	303.0739	1	110800	C <sub>14</sub> H <sub>12</sub> N <sub>4</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	-0.62
304.0066	304.0755	1	35131	C <sub>14</sub> H <sub>12</sub> N <sub>4</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	226.51
305.0077	305.0766	1	6387	C <sub>14</sub> H <sub>12</sub> N <sub>4</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	225.89

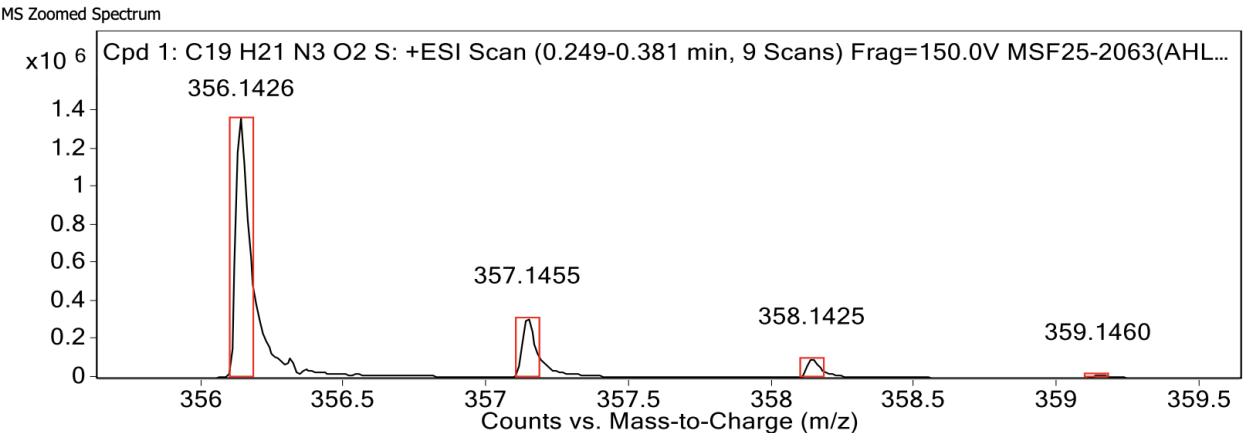
--- End Of Report ---



Target Compound Screening Report

Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

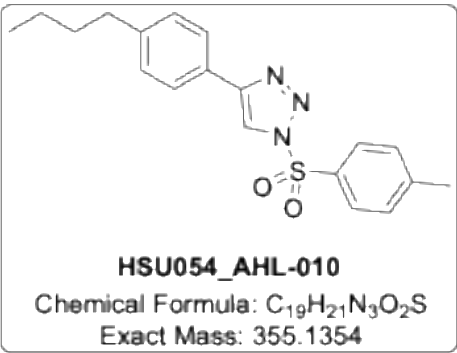
Data File MSF25-2063(AHL010)\_hrESIpos1.d      Sample Name 2063(AHL010)      Comment 2063(AHL010)  
Position P1-D9      Instrument Name 6530      User Name  
Acq Method FIA\_pos.m      Acquired Time 12/4/2025 11:00:01 AM      DA Method MSF.m



MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
350.1180			2843358			
356.1426	356.1427	1	1362227	C19H21N3O2S	(M+H)+	0.32
357.1455	357.1457	1	311766	C19H21N3O2S	(M+H)+	0.45
358.1425	358.1424	1	97903	C19H21N3O2S	(M+H)+	-0.37
359.1460	359.1436	1	15085	C19H21N3O2S	(M+H)+	-6.87

--- End Of Report ---

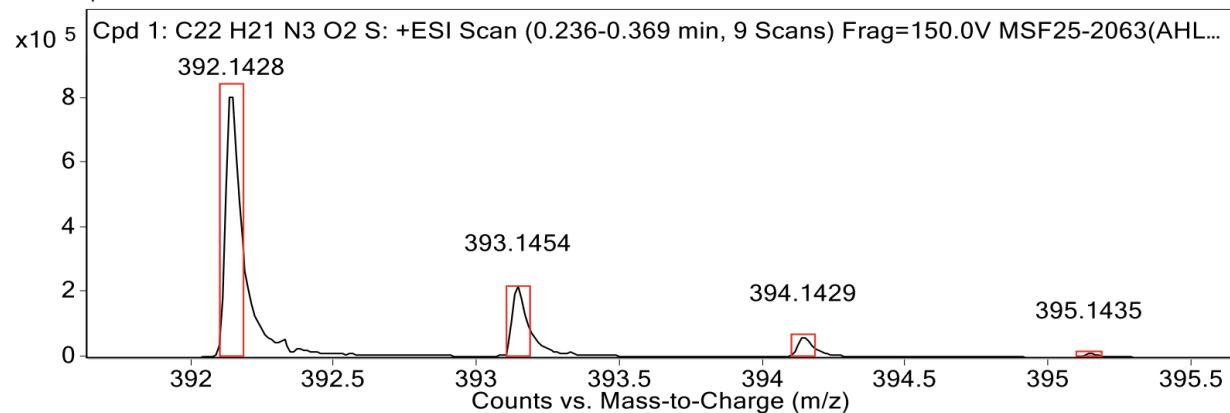


## Target Compound Screening Report

### Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

<b>Data File</b>	MSF25-2063(AHL020)_hrESIpos1.d	<b>Sample Name</b>	2063(AHL020)	<b>Comment</b>	2063(AHL020)
<b>Position</b>	P1-E4	<b>Instrument Name</b>	6530	<b>User Name</b>	
<b>Acq Method</b>	FIA_pos.m	<b>Acquired Time</b>	12/4/2025 11:08:03 AM	<b>DA Method</b>	MSF.m

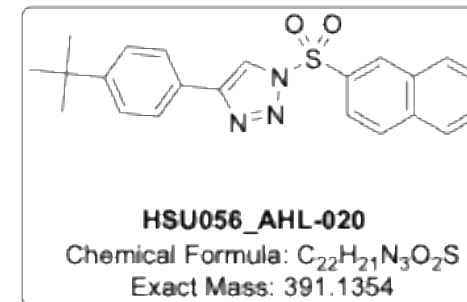
MS Zoomed Spectrum



MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
386.1178			2835460			
392.1428	392.1427	1	839543	C <sub>22</sub> H <sub>21</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	-0.2
393.1454	393.1457	1	220208	C <sub>22</sub> H <sub>21</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	0.83
394.1429	394.1430	1	65407	C <sub>22</sub> H <sub>21</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	0.34
395.1435	395.1440	1	11697	C <sub>22</sub> H <sub>21</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	1.21

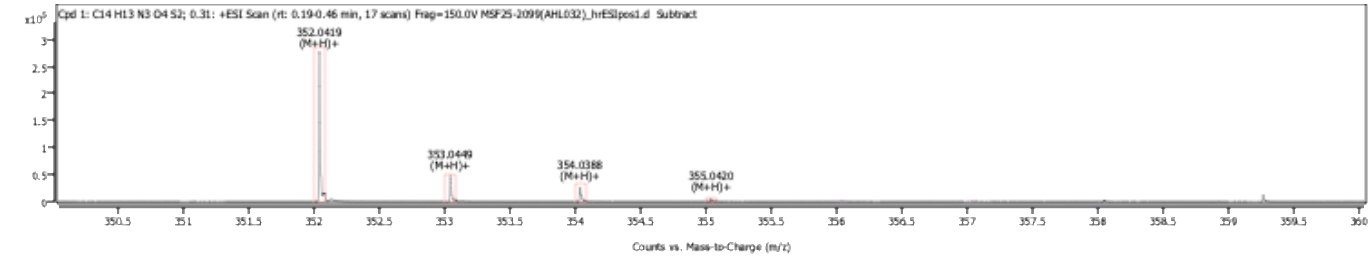
--- End Of Report ---



# Target Screening Report

## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

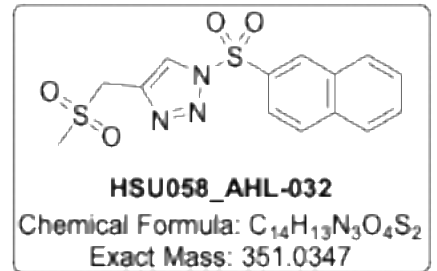
Data File	MSF25-2099(AHL032)_hrESIpos1.d	Sample Name	2099(AHL032)	Comment	2099(AHL032)
Position	P1-E8	Instrument Name	Instrument 1	DA Method	MSF.m
Acquisition Method	FIA_Pos.m	Acquired Time	12/15/2025 5:10:18 PM (UTC-06:00)		



### Spectrum Peaks

Obs. m/z	Calc. m/z	Charge	Abund	Formula	Ion Species	Tgt Mass Error (PPM)
352.0419	352.0420	1	284062	C14H13N3O4S2	(M+H)+	-0.33
353.0449	353.0447	1	49068	C14H13N3O4S2	(M+H)+	0.55
354.0388	354.0397	1	27153	C14H13N3O4S2	(M+H)+	-2.72
355.0420	355.0418	1	3876	C14H13N3O4S2	(M+H)+	0.77
130.1590			9651942			

MassHunter Qual 10.0  
(End of Report)



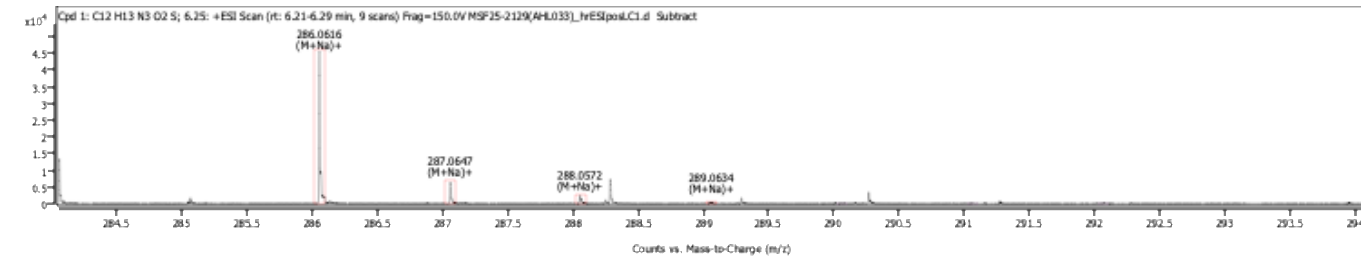


# Target Screening Report



## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

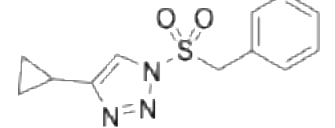
Data File	MSF25-2129(AHL033)_hrESI(pos).CL.d	Sample Name	2129(AHL033)	Comment	2129(AHL033)
Position	P1-C3	Instrument Name	Instrument 1		
Acquisition Method	LC C18 OBO4 .JLm	Acquired Time	12/18/2025 8:37:03 PM (UTC-06:00)	DA Method	MSF.m



### Spectrum Peaks

Obs. m/z	Calc. m/z	Charge	Abund	Formula	Ion Species	Tgt Mass Error (PPM)
286.0616	286.0621	1	45887	C12H13NO2S	(M+Na)+	-1.75
287.0647	287.0648	1	6431	C12H13NO2S	(M+Na)+	-0.36
288.0572	288.0602	1	2512	C12H13NO2S	(M+Na)+	-10.44
289.0634	289.0620	1	351	C12H13NO2S	(M+Na)+	4.81
260.1182			5713466			

MassHunter Qual 10.0  
(End of Report)



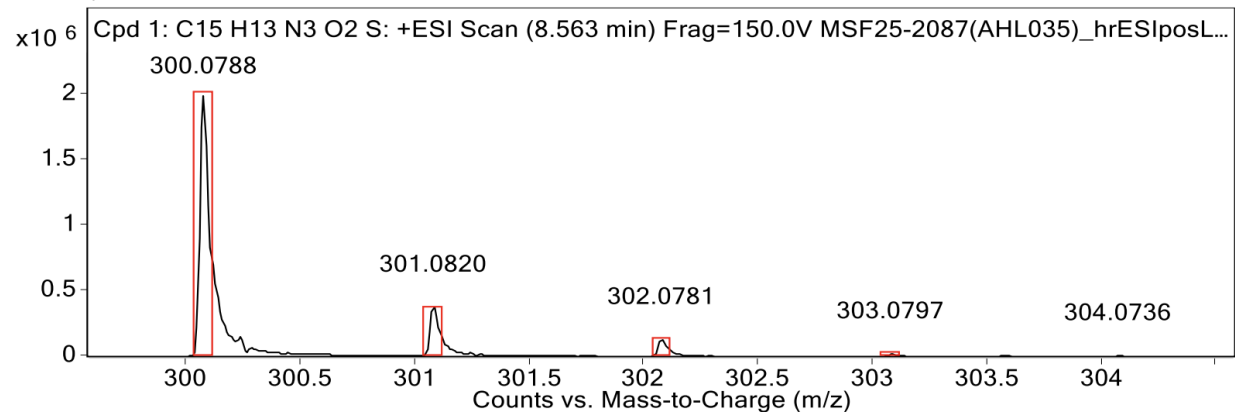
**HSU059\_AHL-033**  
Chemical Formula: C<sub>12</sub>H<sub>13</sub>N<sub>3</sub>O<sub>2</sub>S  
Exact Mass: 263.0728

## Target Compound Screening Report

### Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2087(AHL035)\_hrESIposLC1.d **Sample Name** 2087(AHL035) **Comment** 2087(AHL035)  
**Position** P1-C6 **Instrument Name** 6530 **User Name**  
**Acq Method** LC\_C18\_pos\_jl.m **Acquired Time** 12/4/2025 6:11:50 PM **DA Method** MSF.m

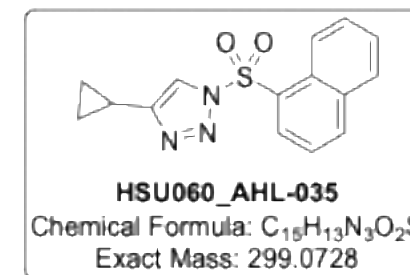
MS Zoomed Spectrum



MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
300.0788	300.0801	1	1995820	C <sub>15</sub> H <sub>13</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	4.34
301.0820	301.0830	1	378066	C <sub>15</sub> H <sub>13</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	3.25
302.0781	302.0789	1	130701	C <sub>15</sub> H <sub>13</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	2.68
303.0797	303.0804	1	21773	C <sub>15</sub> H <sub>13</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	2.34
304.0736	304.0817	1	7650	C <sub>15</sub> H <sub>13</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	26.61
305.0780	305.0833	1	1398	C <sub>15</sub> H <sub>13</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	17.4

--- End Of Report ---



## Target Compound Screening Report

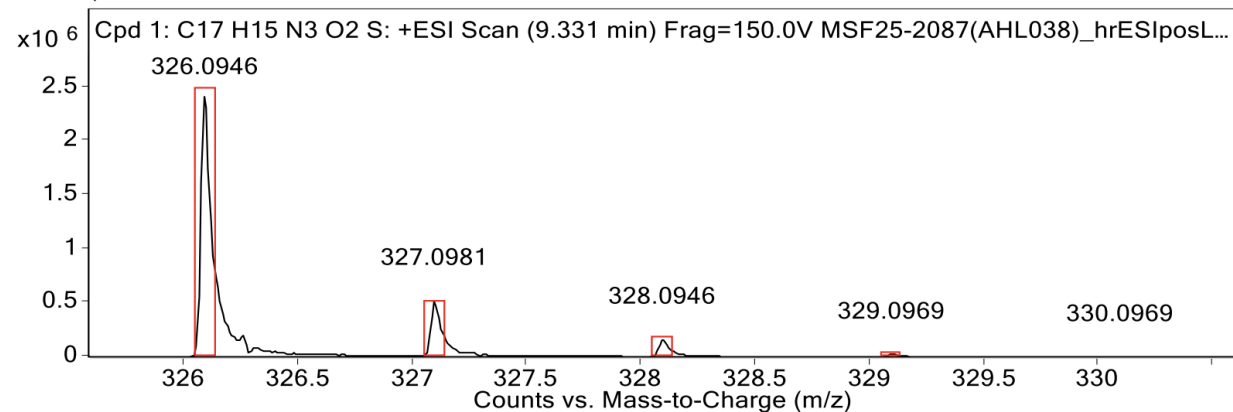
### Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2087(AHL038)\_hrESIposLC1.d  
**Position** P1-C7  
**Acq Method** LC\_C18\_pos\_jl.m

**Sample Name** 2087(AHL038)  
**Instrument Name** 6530  
**Acquired Time** 12/4/2025 6:34:49 PM

**Comment** 2087(AHL038)  
**User Name**  
**DA Method** MSF.m

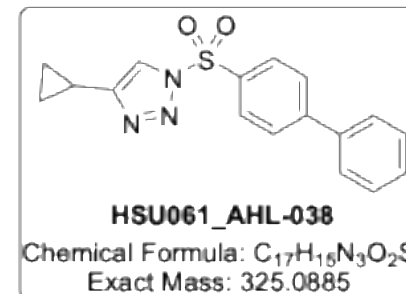
MS Zoomed Spectrum



#### MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
326.0946	326.0958	1	2479350	C <sub>17</sub> H <sub>15</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	3.72
327.0981	327.0987	1	524298	C <sub>17</sub> H <sub>15</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	1.68
328.0946	328.0950	1	164961	C <sub>17</sub> H <sub>15</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	1.03
329.0969	329.0964	1	25706	C <sub>17</sub> H <sub>15</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	-1.78
330.0969	330.0977	1	2744	C <sub>17</sub> H <sub>15</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	2.54

--- End Of Report ---

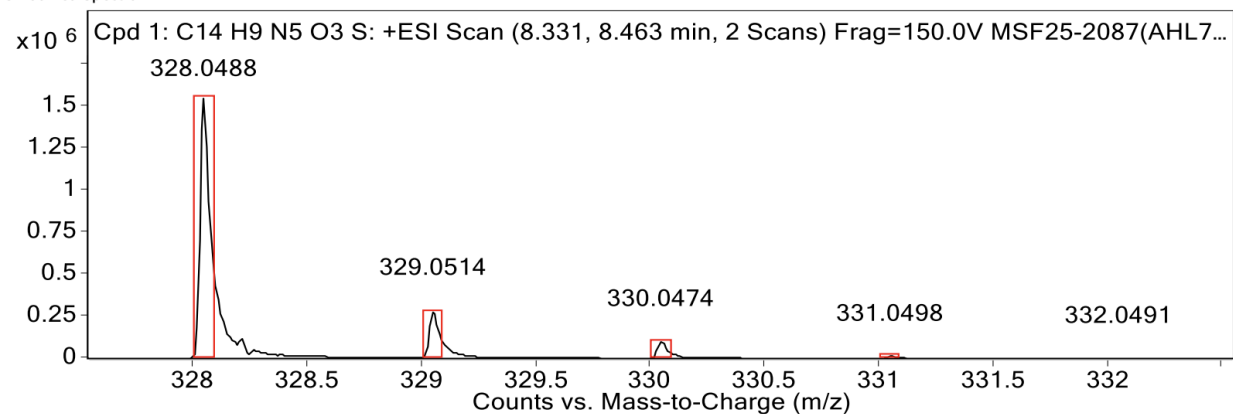


## Target Compound Screening Report

### Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

<b>Data File</b>	MSF25-2087(AHL7004A)_hrESIposLC1.d	<b>Sample Name</b>	2087(AHL7004A)	<b>Comment</b>	2087(AHL7004A)
<b>Position</b>	P1-D2	<b>Instrument Name</b>	6530	<b>User Name</b>	
<b>Acq Method</b>	LC_C18_pos_jl.m	<b>Acquired Time</b>	12/4/2025 8:06:41 PM	<b>DA Method</b>	MSF.m

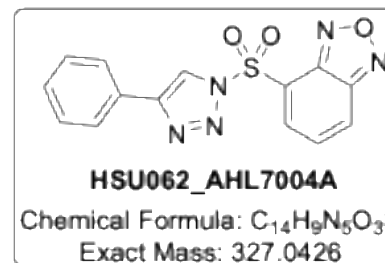
#### MS Zoomed Spectrum



#### MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
328.0488	328.0499	1	1550059	C <sub>14</sub> H <sub>9</sub> N <sub>5</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	3.46
329.0514	329.0524	1	287820	C <sub>14</sub> H <sub>9</sub> N <sub>5</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	3.25
330.0474	330.0486	1	102204	C <sub>14</sub> H <sub>9</sub> N <sub>5</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	3.74
331.0498	331.0501	1	14026	C <sub>14</sub> H <sub>9</sub> N <sub>5</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	1.05
332.0491	332.0512	1	1889	C <sub>14</sub> H <sub>9</sub> N <sub>5</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	6.3

--- End Of Report ---

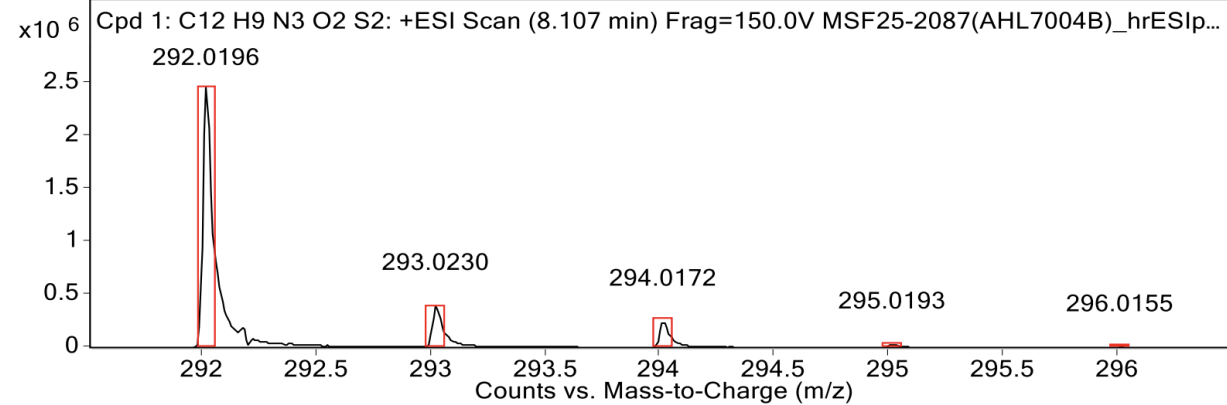


## Target Compound Screening Report

### Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2087(AHL7004B)\_hrESIposLC1.d **Sample Name** 2087(AHL7004B) **Comment** 2087(AHL7004B)  
**Position** P1-D3 **Instrument Name** 6530 **User Name**  
**Acq Method** LC\_C18\_pos\_jl.m **Acquired Time** 12/4/2025 8:29:36 PM **DA Method** MSF.m

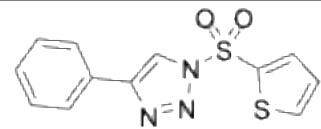
#### MS Zoomed Spectrum



#### MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
292.0196	292.0209	1	2463582	C <sub>12</sub> H <sub>9</sub> N <sub>3</sub> O <sub>2</sub> S <sub>2</sub>	(M+H) <sup>+</sup>	4.5
293.0230	293.0234	1	400507	C <sub>12</sub> H <sub>9</sub> N <sub>3</sub> O <sub>2</sub> S <sub>2</sub>	(M+H) <sup>+</sup>	1.52
294.0172	294.0181	1	251014	C <sub>12</sub> H <sub>9</sub> N <sub>3</sub> O <sub>2</sub> S <sub>2</sub>	(M+H) <sup>+</sup>	2.88
295.0193	295.0201	1	34946	C <sub>12</sub> H <sub>9</sub> N <sub>3</sub> O <sub>2</sub> S <sub>2</sub>	(M+H) <sup>+</sup>	2.81
296.0155	296.0164	1	8654	C <sub>12</sub> H <sub>9</sub> N <sub>3</sub> O <sub>2</sub> S <sub>2</sub>	(M+H) <sup>+</sup>	2.96

--- End Of Report ---



**HSU063\_AHL7004B**

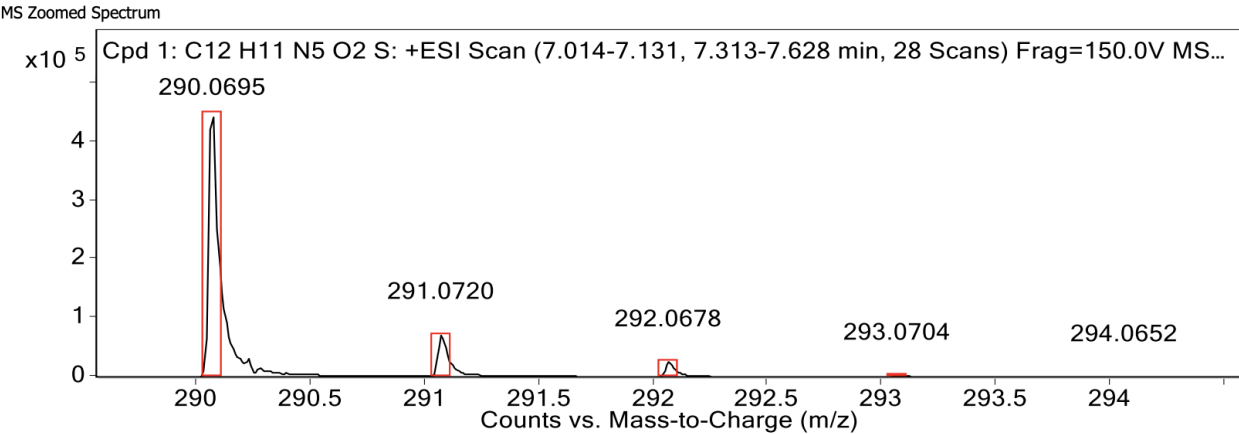
Chemical Formula: C<sub>12</sub>H<sub>10</sub>N<sub>3</sub>O<sub>2</sub>S<sub>2</sub>

Exact Mass: 291.0136

Target Compound Screening Report

Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

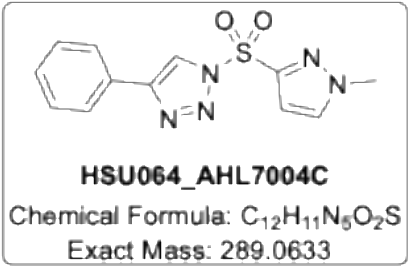
Data File MSF25-2087(AHL7004C)\_hrESIposLC1.d Sample Name 2087(AHL7004C) Comment 2087(AHL7004C)  
Position P1-D4 Instrument Name 6530 User Name  
Acq Method LC\_C18\_pos\_jl.m Acquired Time 12/4/2025 8:52:35 PM DA Method MSF.m



MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
290.0695	290.0706	1	449080	C12H11N5O2S	(M+H)+	3.72
291.0720	291.0731	1	72894	C12H11N5O2S	(M+H)+	3.55
292.0678	292.0687	1	26001	C12H11N5O2S	(M+H)+	3.26
293.0704	293.0703	1	3242	C12H11N5O2S	(M+H)+	-0.45
294.0652	294.0713	1	487	C12H11N5O2S	(M+H)+	20.67

--- End Of Report ---

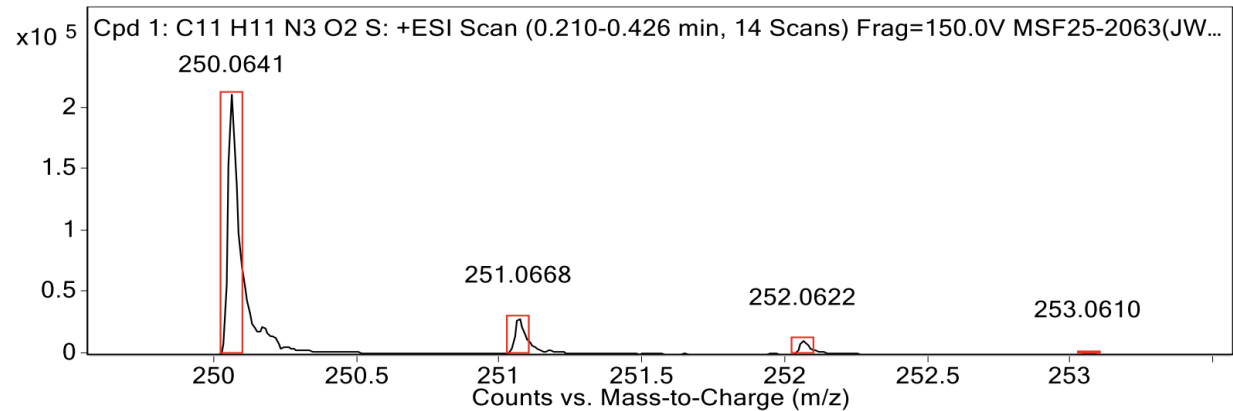


## Target Compound Screening Report

### Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2063(JWB101)\_hrESIpos1.d **Sample Name** 2063(JWB101) **Comment** 2063(JWB101)  
**Position** P1-C1 **Instrument Name** 6530 **User Name**  
**Acq Method** FIA\_pos.m **Acquired Time** 12/4/2025 10:25:37 AM **DA Method** MSF.m

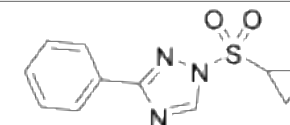
MS Zoomed Spectrum



#### MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
250.0641	250.0645	1	212953	C <sub>11</sub> H <sub>11</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	1.55
251.0668	251.0671	1	29722	C <sub>11</sub> H <sub>11</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	1.31
252.0622	252.0624	1	11501	C <sub>11</sub> H <sub>11</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	0.54
253.0610	253.0643	1	1470	C <sub>11</sub> H <sub>11</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	13.2
867.0737			212500			

--- End Of Report ---



**HSU071\_JWB101**

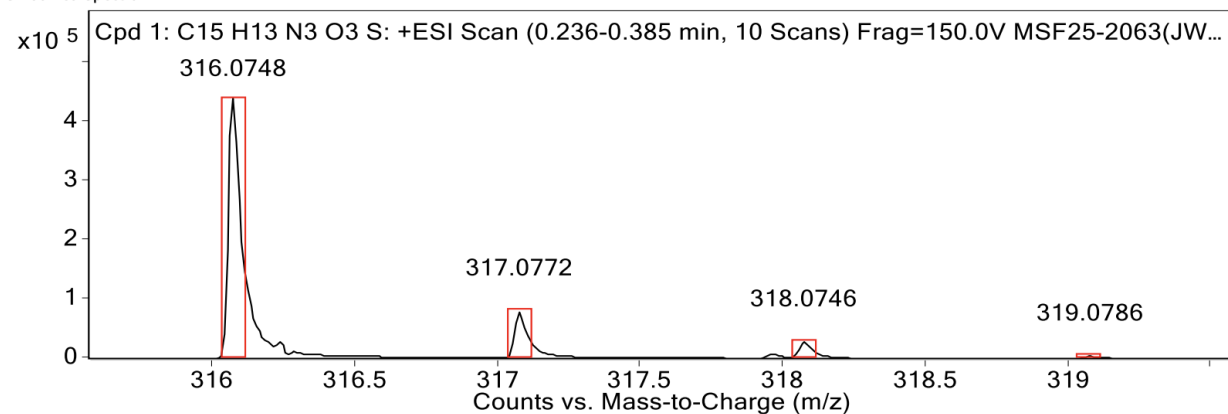
Chemical Formula: C<sub>11</sub>H<sub>11</sub>N<sub>3</sub>O<sub>2</sub>S  
Exact Mass: 249.0572

## Target Compound Screening Report

### Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2063(JWB106)\_hrESIpos1.d **Sample Name** 2063(JWB106) **Comment** 2063(JWB106)  
**Position** P1-C2 **Instrument Name** 6530 **User Name**  
**Acq Method** FIA\_pos.m **Acquired Time** 12/4/2025 10:27:38 AM **DA Method** MSF.m

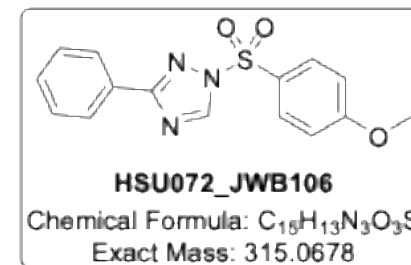
MS Zoomed Spectrum



#### MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
316.0748	316.0750	1	440410	C <sub>15</sub> H <sub>13</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	0.81
317.0772	317.0779	1	80937	C <sub>15</sub> H <sub>13</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	2.12
318.0746	318.0740	1	28563	C <sub>15</sub> H <sub>13</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	-1.94
319.0786	319.0756	1	4504	C <sub>15</sub> H <sub>13</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	-9.46
320.0772	320.0768	1	811	C <sub>15</sub> H <sub>13</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	-1.29

--- End Of Report ---

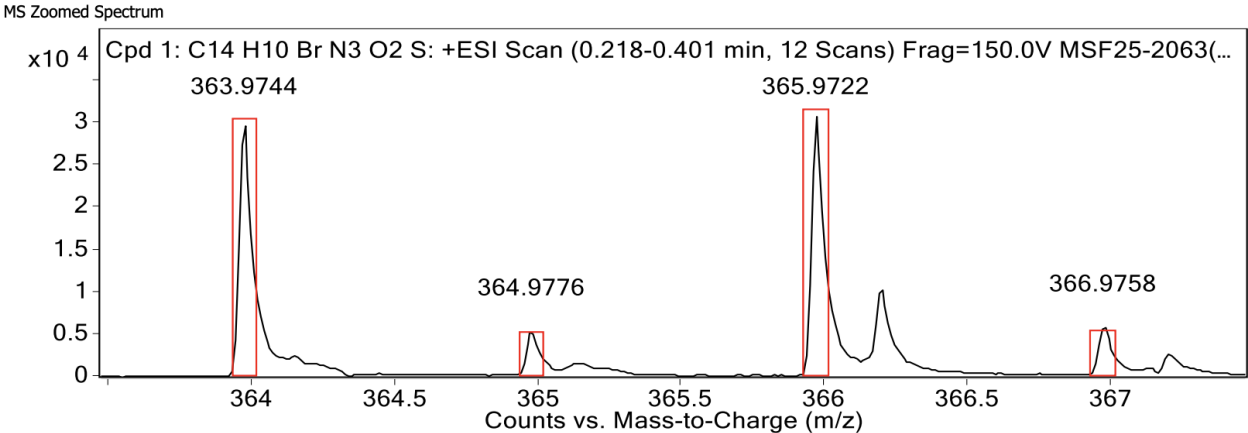




# Target Compound Screening Report

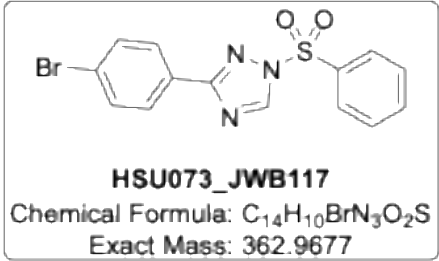
## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2063(JWB117)\_hrESIpos1.d      **Sample Name** 2063(JWB117)      **Comment** 2063(JWB117)  
**Position** P1-C3      **Instrument Name** 6530      **User Name**  
**Acq Method** FIA\_pos.m      **Acquired Time** 12/4/2025 10:29:40 AM      **DA Method** MSF.m



MS Spectrum Peak List						
Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
130.1591			422236			
363.9744	363.9750	1	29876	C14H10BrN3O2S	(M+H)+	1.66
364.9776	364.9778	1	5489	C14H10BrN3O2S	(M+H)+	0.47
365.9722	365.9730	1	30849	C14H10BrN3O2S	(M+H)+	2.12
366.9758	366.9757	1	5953	C14H10BrN3O2S	(M+H)+	-0.37

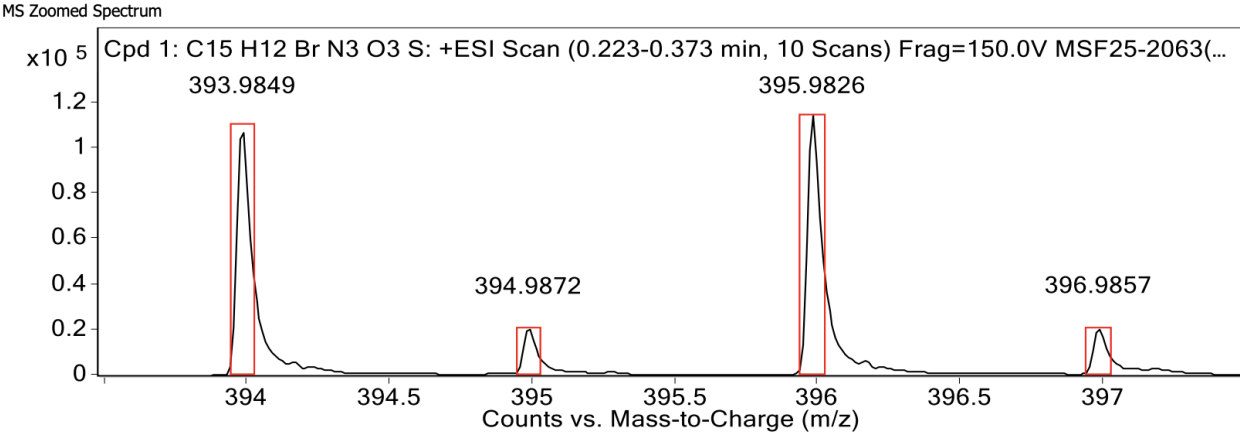
--- End Of Report ---



# Target Compound Screening Report

## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

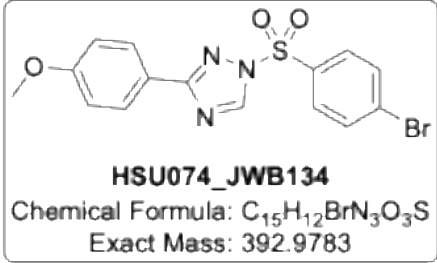
**Data File** MSF25-2063(JWB134)\_hrESIpos1.d      **Sample Name** 2063(JWB134)      **Comment** 2063(JWB134)  
**Position** P1-C6      **Instrument Name** 6530      **User Name**  
**Acq Method** FIA\_pos.m      **Acquired Time** 12/4/2025 10:35:47 AM      **DA Method** MSF.m



MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
393.9849	393.9856	1	109046	C15H12BrN3O3S	(M+H)+	1.69
394.9872	394.9884	1	20545	C15H12BrN3O3S	(M+H)+	2.99
395.9826	395.9836	1	114568	C15H12BrN3O3S	(M+H)+	2.43
396.9857	396.9863	1	20906	C15H12BrN3O3S	(M+H)+	1.67
867.0743			319119			

--- End Of Report ---

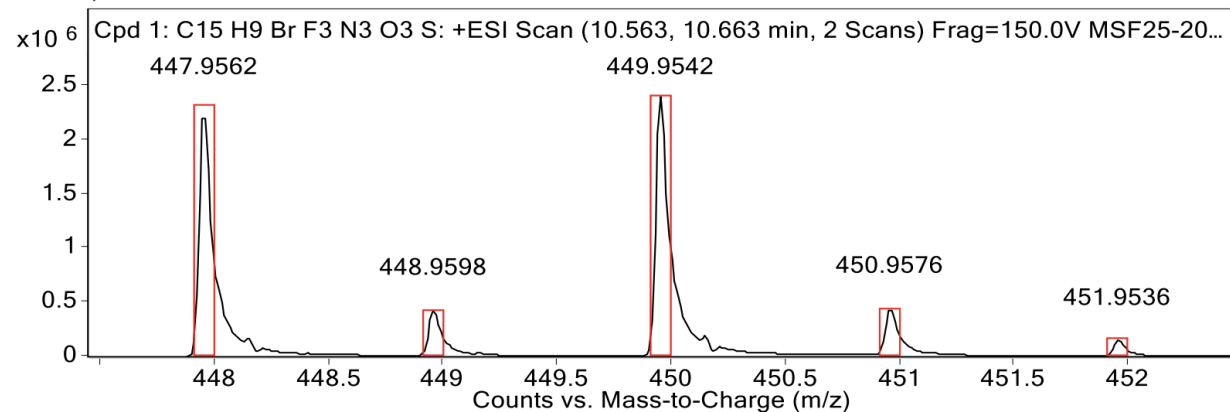


# Target Compound Screening Report

## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2063(JWB124)\_hrESIposLC1.d **Sample Name** 2063(JWB124) **Comment** 2063(JWB124)  
**Position** P1-C5 **Instrument Name** 6530 **User Name**  
**Acq Method** LC\_C18\_pos\_jl.m **Acquired Time** 12/4/2025 12:04:43 PM **DA Method** MSF.m

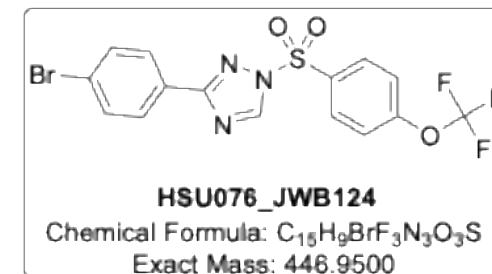
MS Zoomed Spectrum



MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
447.9562	447.9573	1	2289222	C <sub>15</sub> H <sub>9</sub> BrF <sub>3</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	2.33
448.9598	448.9601	1	430820	C <sub>15</sub> H <sub>9</sub> BrF <sub>3</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	0.73
449.9542	449.9553	1	2400483	C <sub>15</sub> H <sub>9</sub> BrF <sub>3</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	2.36
450.9576	450.9581	1	449016	C <sub>15</sub> H <sub>9</sub> BrF <sub>3</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	1.07
451.9536	451.9542	1	152213	C <sub>15</sub> H <sub>9</sub> BrF <sub>3</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	1.46

--- End Of Report ---

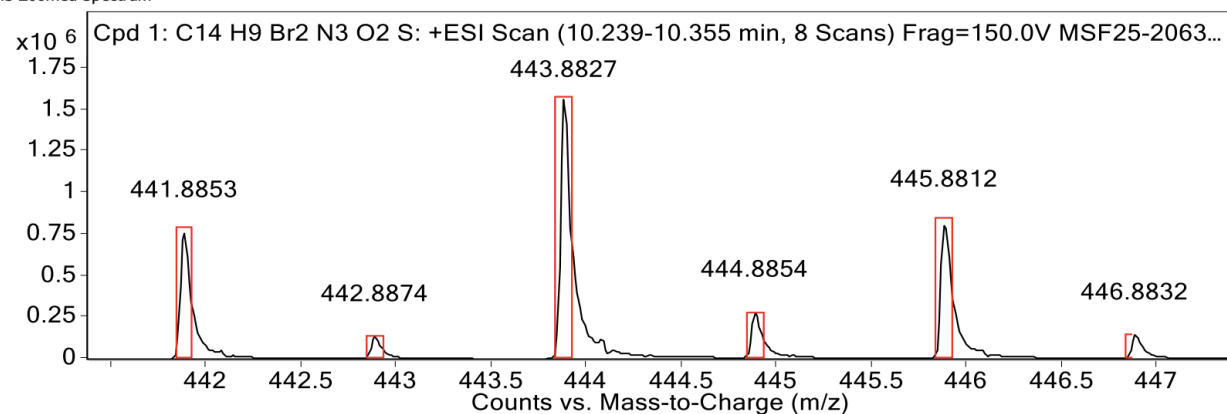


## Target Compound Screening Report

### Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

<b>Data File</b>	MSF25-2063(JWB119)_hrESIposLC1.d	<b>Sample Name</b>	2063(JWB119)	<b>Comment</b>	2063(JWB119)
<b>Position</b>	P1-C4	<b>Instrument Name</b>	6530	<b>User Name</b>	
<b>Acq Method</b>	LC_C18_pos_jl.m	<b>Acquired Time</b>	12/4/2025 5:25:51 PM	<b>DA Method</b>	MSF.m

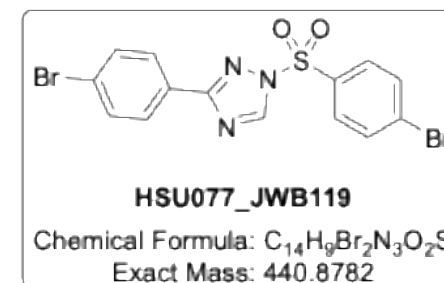
MS Zoomed Spectrum



#### MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
441.8853	441.8855	1	763367	C <sub>14</sub> H <sub>9</sub> Br <sub>2</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	0.55
442.8874	442.8883	1	136331	C <sub>14</sub> H <sub>9</sub> Br <sub>2</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	2.04
443.8827	443.8835	1	1575727	C <sub>14</sub> H <sub>9</sub> Br <sub>2</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	1.84
444.8854	444.8862	1	284008	C <sub>14</sub> H <sub>9</sub> Br <sub>2</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	1.83
445.8812	445.8815	1	826850	C <sub>14</sub> H <sub>9</sub> Br <sub>2</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	0.64
446.8832	446.8841	1	146646	C <sub>14</sub> H <sub>9</sub> Br <sub>2</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	2.22

--- End Of Report ---

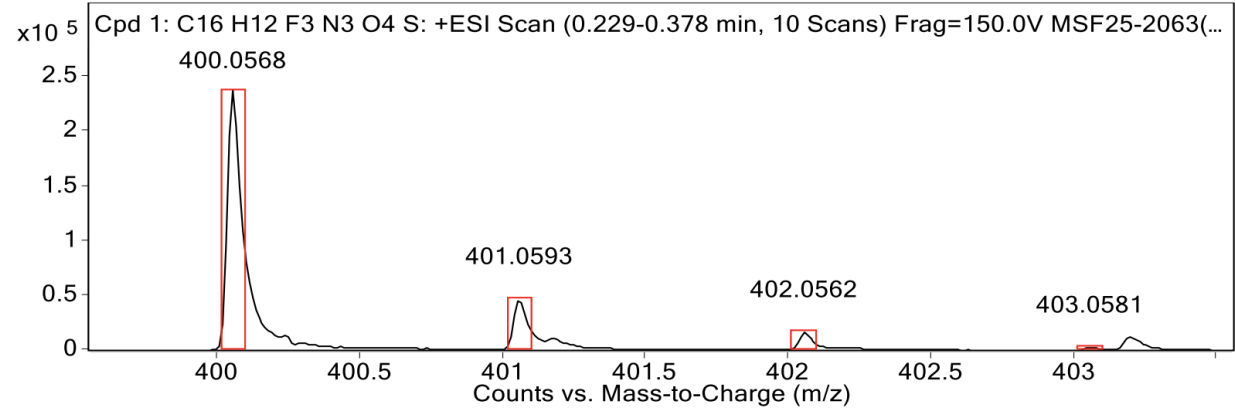


## Target Compound Screening Report

### Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2063(JWB139)\_hrESIpos1.d **Sample Name** 2063(JWB139) **Comment** 2063(JWB139)  
**Position** P1-C7 **Instrument Name** 6530 **User Name**  
**Acq Method** FIA\_pos.m **Acquired Time** 12/4/2025 10:37:50 AM **DA Method** MSF.m

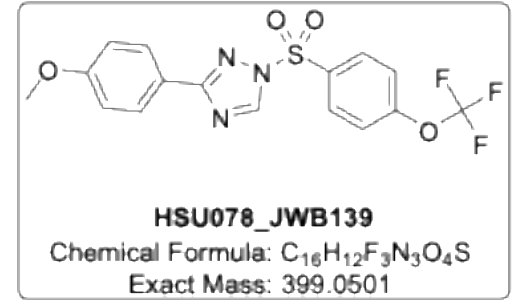
MS Zoomed Spectrum



#### MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
400.0568	400.0573	1	237845	C <sub>16</sub> H <sub>12</sub> F <sub>3</sub> N <sub>3</sub> O <sub>4</sub> S	(M+H) <sup>+</sup>	1.23
401.0593	401.0602	1	46701	C <sub>16</sub> H <sub>12</sub> F <sub>3</sub> N <sub>3</sub> O <sub>4</sub> S	(M+H) <sup>+</sup>	2.2
402.0562	402.0566	1	16432	C <sub>16</sub> H <sub>12</sub> F <sub>3</sub> N <sub>3</sub> O <sub>4</sub> S	(M+H) <sup>+</sup>	1.04
403.0581	403.0583	1	2353	C <sub>16</sub> H <sub>12</sub> F <sub>3</sub> N <sub>3</sub> O <sub>4</sub> S	(M+H) <sup>+</sup>	0.35

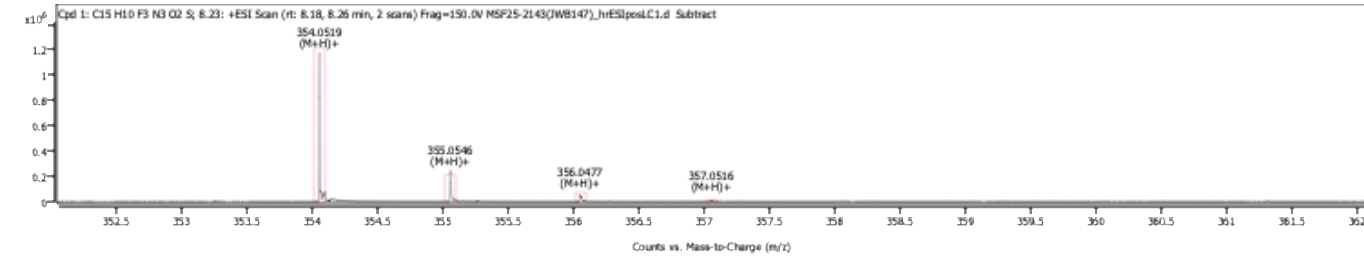
--- End Of Report ---



# Target Screening Report

## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

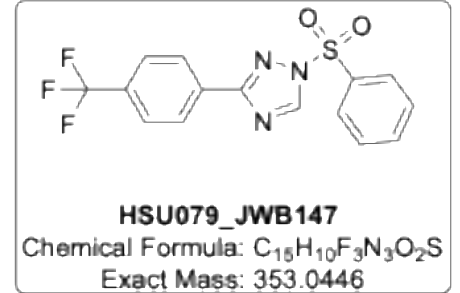
Data File	MSF25-2143_JWB147_hrESIposLC1.d	Sample Name	2143_JWB147	Comment	2143(JWB147)
Position	P1-C8	Instrument Name	Instrument 1		
Acquisition Method	LC C18 OBDN 3.0um	Acquired Time	12/19/2025 2:34:05 PM (UTC-06:00)	DA Method	MSF.m



### Spectrum Peaks

Obs. m/z	Calc. m/z	Charge	Abund	Formula	Ion Species	Tgt Mass Error (PPM)
354.0519	354.0519	1	1207995	C15H10F3N3O2S	(M+H)+	0.04
355.0546	355.0547	1	251557	C15H10F3N3O2S	(M+H)+	-0.12
356.0477	356.0506	1	55689	C15H10F3N3O2S	(M+H)+	-8.13
357.0516	357.0522	1	11552	C15H10F3N3O2S	(M+H)+	-1.49

MassHunter Qual 10.0  
(End of Report)

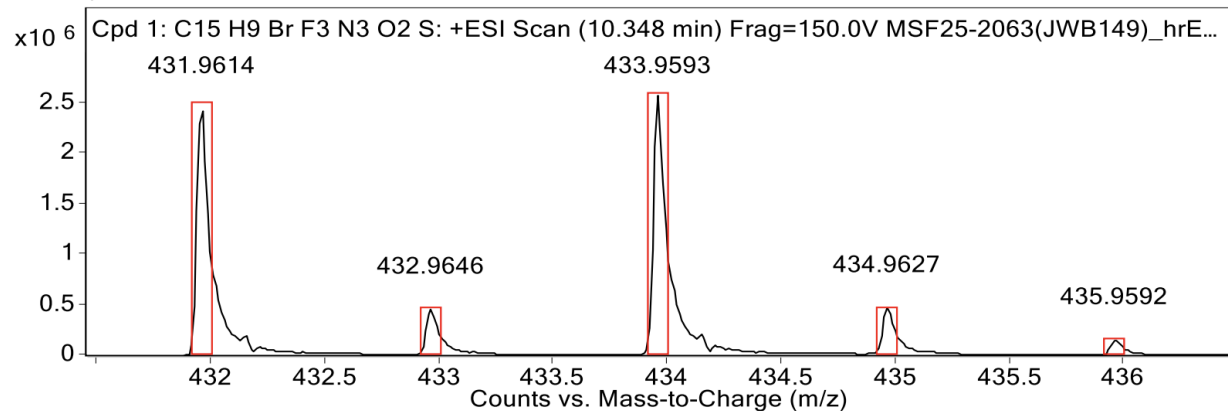


## Target Compound Screening Report

### Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2063(JWB149)\_hrESIposLC1.d **Sample Name** 2063(JWB149) **Comment** 2063(JWB149)  
**Position** P1-C8 **Instrument Name** 6530 **User Name**  
**Acq Method** LC\_C18\_pos\_jl.m **Acquired Time** 12/4/2025 12:27:42 PM **DA Method** MSF.m

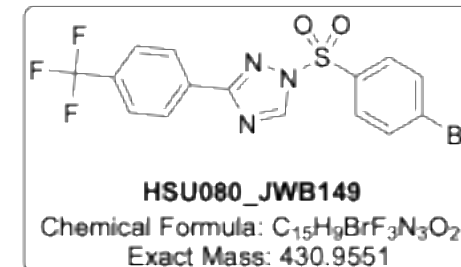
MS Zoomed Spectrum



#### MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
431.9614	431.9624	1	2447313	C <sub>15</sub> H <sub>9</sub> BrF <sub>3</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	2.28
432.9646	432.9652	1	463283	C <sub>15</sub> H <sub>9</sub> BrF <sub>3</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	1.42
433.9593	433.9604	1	2586893	C <sub>15</sub> H <sub>9</sub> BrF <sub>3</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	2.53
434.9627	434.9631	1	478074	C <sub>15</sub> H <sub>9</sub> BrF <sub>3</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	0.97
435.9592	435.9591	1	160057	C <sub>15</sub> H <sub>9</sub> BrF <sub>3</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	-0.19
436.9607	436.9607	1	20499	C <sub>15</sub> H <sub>9</sub> BrF <sub>3</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	-0.01

--- End Of Report ---

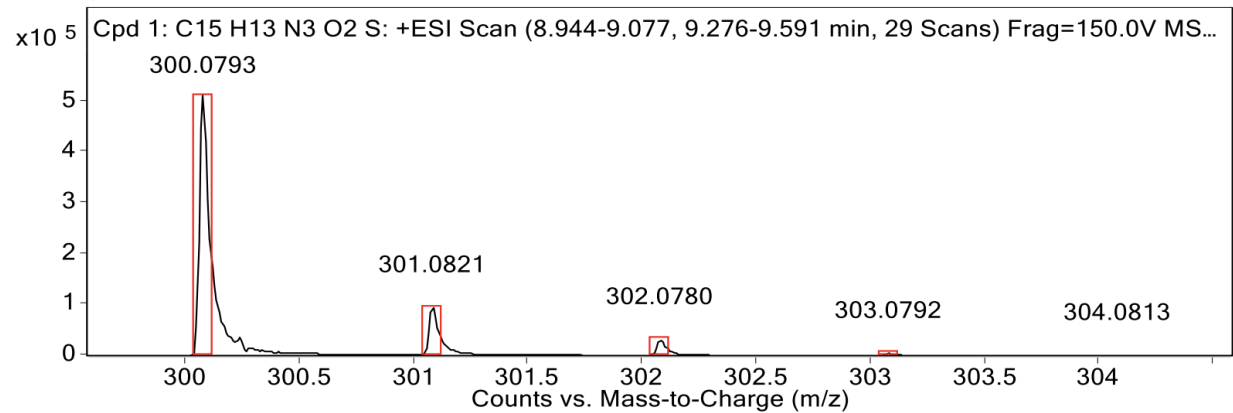


## Target Compound Screening Report

### Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2087(AHL1101)\_hrESIposLC1.d **Sample Name** 2087(AHL1101) **Comment** 2087(AHL1101)  
**Position** P1-C8 **Instrument Name** 6530 **User Name**  
**Acq Method** LC\_C18\_pos\_jl.m **Acquired Time** 12/4/2025 6:57:49 PM **DA Method** MSF.m

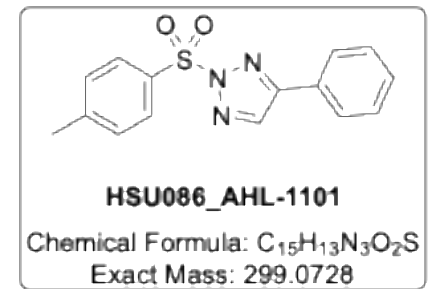
MS Zoomed Spectrum



MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
300.0793	300.0801	1	512406	C <sub>15</sub> H <sub>13</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	2.81
301.0821	301.0830	1	94533	C <sub>15</sub> H <sub>13</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	2.8
302.0780	302.0789	1	31690	C <sub>15</sub> H <sub>13</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	2.79
303.0792	303.0804	1	4518	C <sub>15</sub> H <sub>13</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	4.28
304.0813	304.0817	1	560	C <sub>15</sub> H <sub>13</sub> N <sub>3</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	1.48

--- End Of Report ---



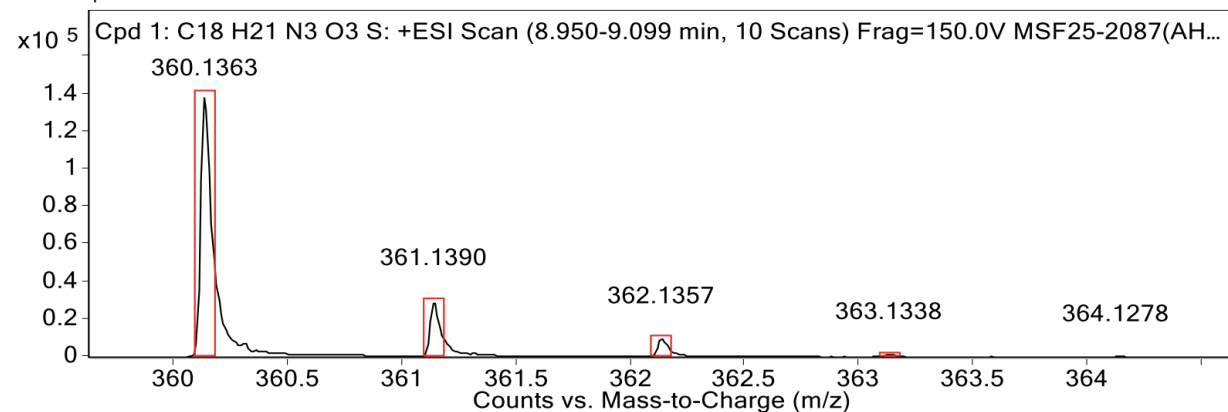


## Target Compound Screening Report

### Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2087(AHL1118)\_hrESIposLC1.d **Sample Name** 2087(AHL1118) **Comment** 2087(AHL1118)  
**Position** P1-C9 **Instrument Name** 6530 **User Name**  
**Acq Method** LC\_C18\_pos\_jl.m **Acquired Time** 12/4/2025 7:20:48 PM **DA Method** MSF.m

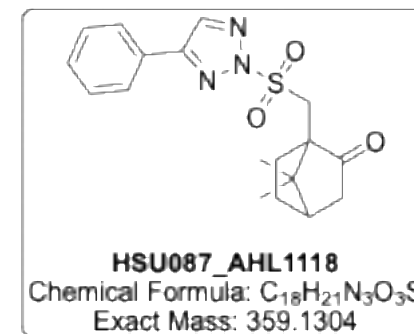
MS Zoomed Spectrum



#### MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
146.0701			4780333			
360.1363	360.1376	1	141517	C18H21N3O3S	(M+H)+	3.69
361.1390	361.1406	1	30287	C18H21N3O3S	(M+H)+	4.26
362.1357	362.1372	1	9962	C18H21N3O3S	(M+H)+	4.17
363.1338	363.1386	1	1560	C18H21N3O3S	(M+H)+	13.17

--- End Of Report ---

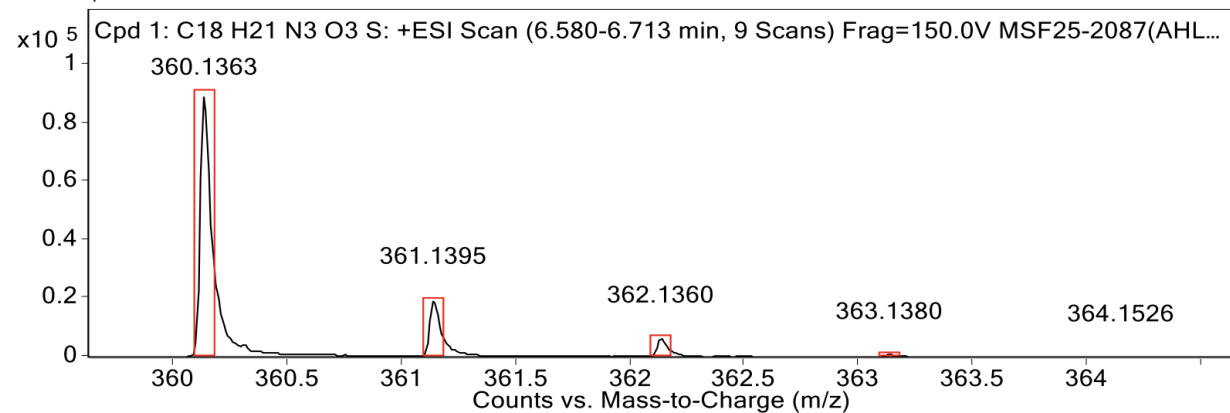


## Target Compound Screening Report

### Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2087(AHL1119)\_hrESIposLC1.d **Sample Name** 2087(AHL1119) **Comment** 2087(AHL1119)  
**Position** P1-D1 **Instrument Name** 6530 **User Name**  
**Acq Method** LC\_C18\_pos\_jl.m **Acquired Time** 12/4/2025 7:43:46 PM **DA Method** MSF.m

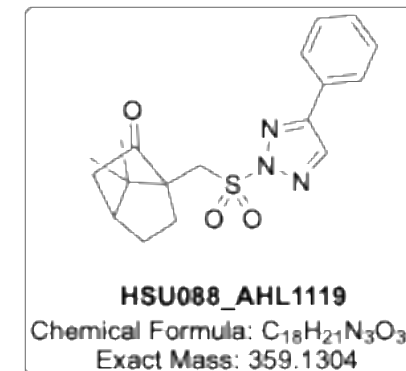
#### MS Zoomed Spectrum



#### MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
360.1363	360.1376	1	91013	C <sub>18</sub> H <sub>21</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	3.61
361.1395	361.1406	1	19815	C <sub>18</sub> H <sub>21</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	2.98
362.1360	362.1372	1	6552	C <sub>18</sub> H <sub>21</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	3.29
363.1380	363.1386	1	929	C <sub>18</sub> H <sub>21</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	1.57
364.1526	364.1400	1	101	C <sub>18</sub> H <sub>21</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	-34.75

--- End Of Report ---

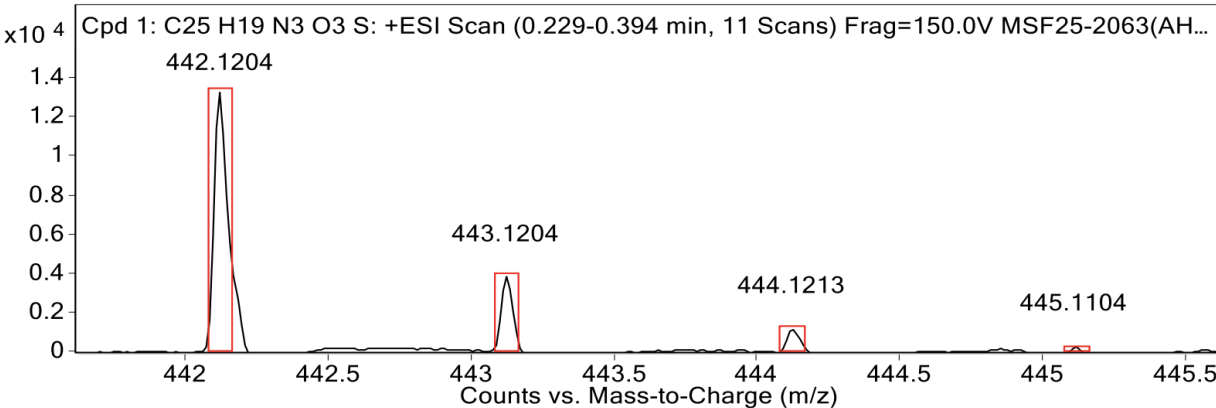


# Target Compound Screening Report

## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2063(AHL1122)\_hrESIpos1.d      **Sample Name** 2063(AHL1122)      **Comment** 2063(AHL1122)  
**Position** P1-E6      **Instrument Name** 6530      **User Name**  
**Acq Method** FIA\_pos.m      **Acquired Time** 12/4/2025 11:12:08 AM      **DA Method** MSF.m

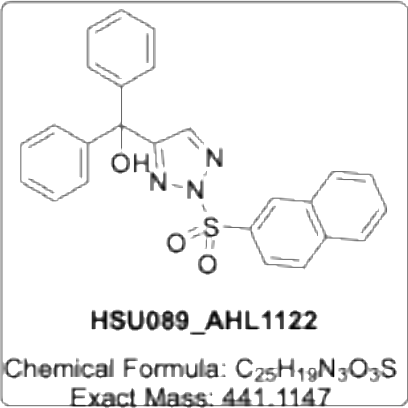
MS Zoomed Spectrum



MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
442.1204	442.1220	1	13323	C <sub>25</sub> H <sub>19</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	3.7
443.1204	443.1250	1	3886	C <sub>25</sub> H <sub>19</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	10.41
444.1213	444.1229	1	1240	C <sub>25</sub> H <sub>19</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	3.65
445.1104	445.1238	1	344	C <sub>25</sub> H <sub>19</sub> N <sub>3</sub> O <sub>3</sub> S	(M+H) <sup>+</sup>	30.11
905.2183			6399613			

--- End Of Report ---

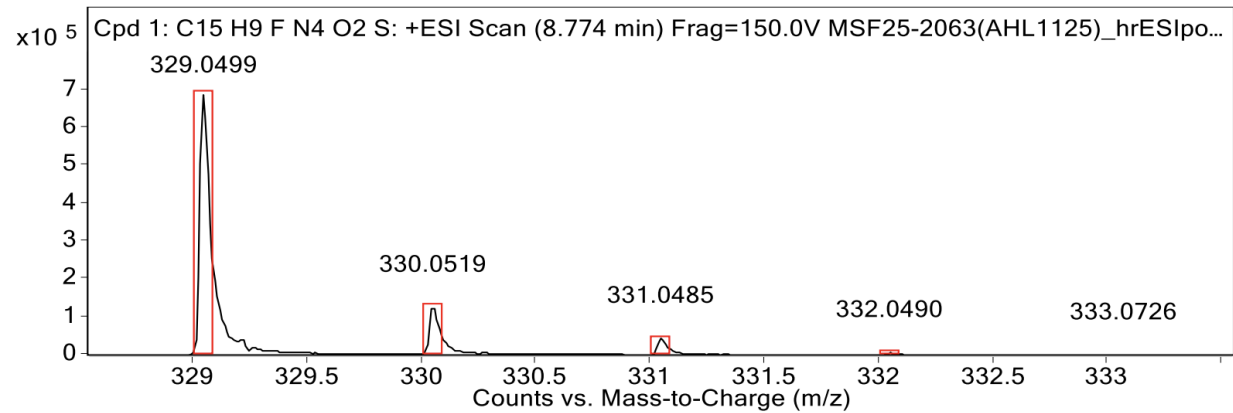


# Target Compound Screening Report

## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2063(AHL1125)\_hrESIposLC1.d **Sample Name** Unavailable **Comment** Sample information is unavailable  
**Position** Unavailable **Instrument Name** Unavailable **User Name** Unavailable  
**Acq Method** **Acquired Time** Unavailable **DA Method** MSF.m

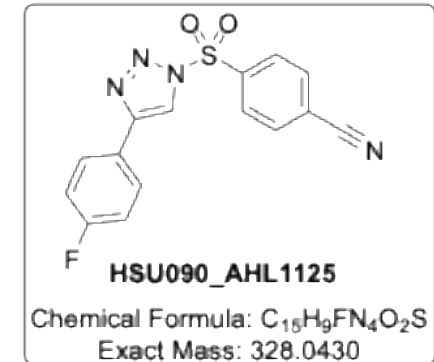
MS Zoomed Spectrum



MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
329.0499	329.0503	1	695013	C <sub>15</sub> H <sub>9</sub> FN <sub>4</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	1.19
330.0519	330.0530	1	126472	C <sub>15</sub> H <sub>9</sub> FN <sub>4</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	3.49
331.0485	331.0490	1	44544	C <sub>15</sub> H <sub>9</sub> FN <sub>4</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	1.68
332.0490	332.0505	1	6993	C <sub>15</sub> H <sub>9</sub> FN <sub>4</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	4.49
333.0726	333.0517	1	643	C <sub>15</sub> H <sub>9</sub> FN <sub>4</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	-62.64
334.0612	334.0533	1	495	C <sub>15</sub> H <sub>9</sub> FN <sub>4</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	-23.8

--- End Of Report ---

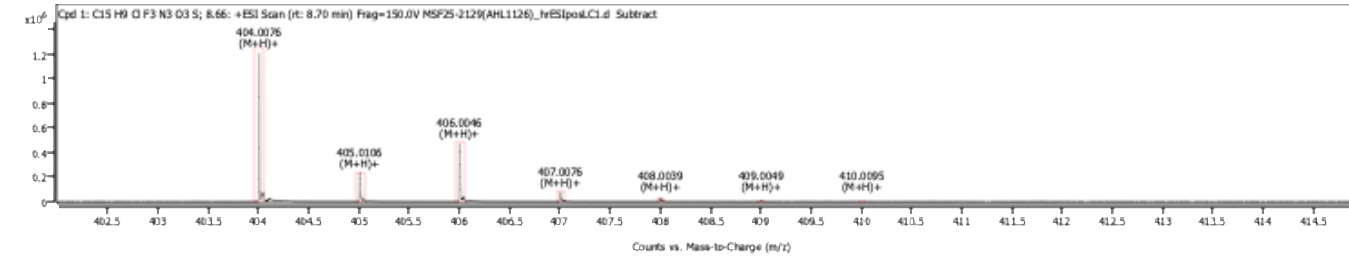


# Target Screening Report



## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

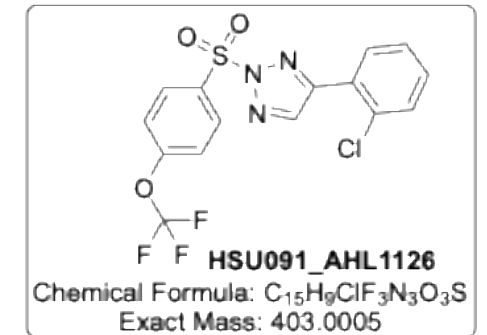
<b>Data File</b>	MSF25-2129(AHL1126)_hrESIsposLC1.d	<b>Sample Name</b>	2129(AHL1126)	<b>Comment</b>	2129(AHL1126)
<b>Position</b>	P1-C4	<b>Instrument Name</b>	Instrument 1		
<b>Acquisition Method</b>	LC C18 O-DB4 .Lum	<b>Acquired Time</b>	12/18/2025 8:59:02 PM (UTC-06:00)	<b>DA Method</b>	MSF.m



### Spectrum Peaks

Obs. m/z	Calc. m/z	Charge	Abund	Formula	Ion Species	Tgt Mass Error (PPM)
404.0076	404.0078	1	1250670	C15H9ClF3N3O3S	(M+H)+	-0.56
405.0106	405.0106	1	240559	C15H9ClF3N3O3S	(M+H)+	-0.12
406.0046	406.0052	1	477045	C15H9ClF3N3O3S	(M+H)+	-1.49
407.0076	407.0078	1	78114	C15H9ClF3N3O3S	(M+H)+	-0.46
408.0039	408.0041	1	19458	C15H9ClF3N3O3S	(M+H)+	-0.30
409.0049	409.0056	1	4511	C15H9ClF3N3O3S	(M+H)+	-1.71
410.0095	410.0097	1	517	C15H9ClF3N3O3S	(M+H)+	6.74

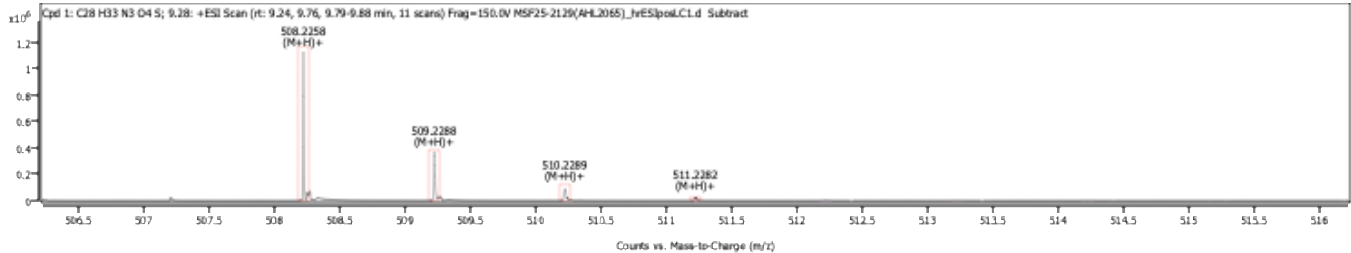
MassHunter Qual 10.0  
(End of Report)



# Target Screening Report

## Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

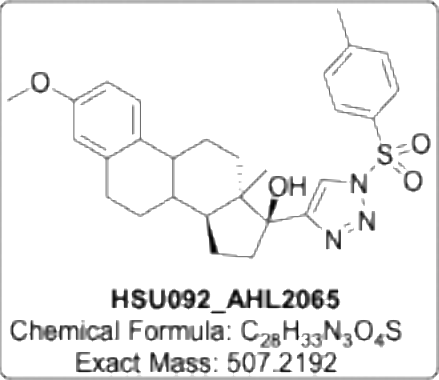
Data File	MSF25-2129(AHL2065)_hrESIsolCL1.d	Sample Name	2129(AHL2065)	Comment	2129(AHL2065)
Position	P1-C5	Instrument Name	Instrument 1		
Acquisition Method	LC C18 OBD4 .JLm	Acquired Time	12/18/2025 9:21:00 PM (UTC-06:00)	DA Method	MSF.m



### Spectrum Peaks

Obs. m/z	Calc. m/z	Charge	Abund	Formula	Ion Species	Tgt Mass Error (PPM)
508.2258	508.2265	1	1165402	C28H33NO4S	(M+H)+	-1.25
509.2288	509.2285	1	369669	C28H33NO4S	(M+H)+	-1.55
510.2289	510.2280	1	88122	C28H33NO4S	(M+H)+	1.77
511.2282	511.2289	1	20306	C28H33NO4S	(M+H)+	-1.32
536.2096			2465220			

MassHunter Qual 10.0  
(End of Report)

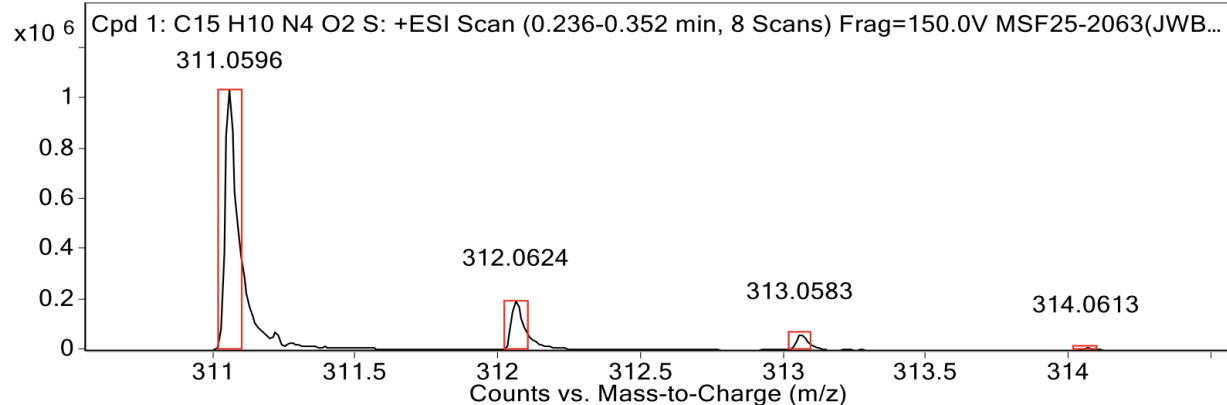


## Target Compound Screening Report

### Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2063(JWB230)\_hrESIpos1.d **Sample Name** 2063(JWB230) **Comment** 2063(JWB230)  
**Position** P1-D4 **Instrument Name** 6530 **User Name**  
**Acq Method** FIA\_pos.m **Acquired Time** 12/4/2025 10:49:55 AM **DA Method** MSF.m

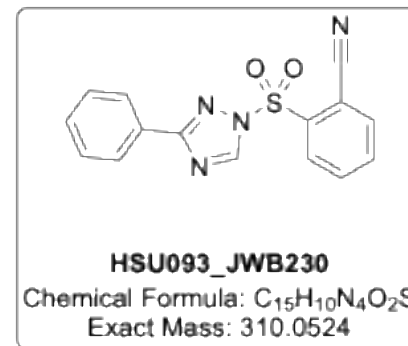
MS Zoomed Spectrum



#### MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
311.0596	311.0597	1	1034641	C <sub>15</sub> H <sub>10</sub> N <sub>4</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	0.28
312.0624	312.0624	1	196895	C <sub>15</sub> H <sub>10</sub> N <sub>4</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	0.21
313.0583	313.0585	1	64033	C <sub>15</sub> H <sub>10</sub> N <sub>4</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	0.69
314.0613	314.0599	1	10785	C <sub>15</sub> H <sub>10</sub> N <sub>4</sub> O <sub>2</sub> S	(M+H) <sup>+</sup>	-4.5
333.0414			1629377			

--- End Of Report ---

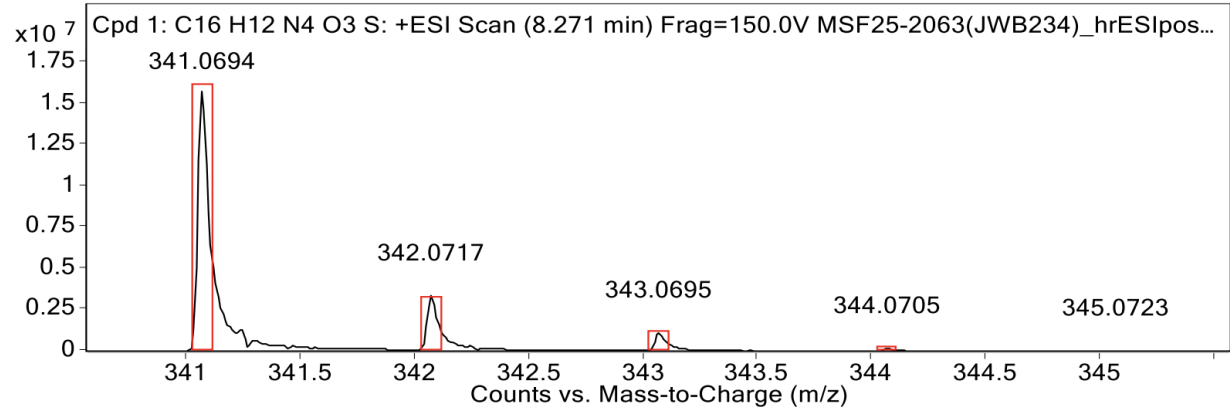


## Target Compound Screening Report

### Results Acquired by The University of Texas at Austin Mass Spectrometry Facility

**Data File** MSF25-2063(JWB234)\_hrESIposLC1.d **Sample Name** 2063(JWB234) **Comment** 2063(JWB234)  
**Position** P1-D5 **Instrument Name** 6530 **User Name**  
**Acq Method** LC\_C18\_pos\_jl.m **Acquired Time** 12/4/2025 1:13:39 PM **DA Method** MSF.m

#### MS Zoomed Spectrum



#### MS Spectrum Peak List

Obs. m/z	Calc. m/z	Charge	Abundance	Formula	Ion Species	Tgt Mass Error (ppm)
341.0694	341.0703	1	16007025	C16H12N4O3S	(M+H)+	2.53
342.0717	342.0730	1	3361506	C16H12N4O3S	(M+H)+	4.05
343.0695	343.0694	1	1111244	C16H12N4O3S	(M+H)+	-0.08
344.0705	344.0709	1	165984	C16H12N4O3S	(M+H)+	0.99
345.0723	345.0721	1	20594	C16H12N4O3S	(M+H)+	-0.61

--- End Of Report ---

