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Appendix

A. Pharmacophore-based virtual screening

Table A1. Evaluation of the quality of SHP2 protein (PDB ID: 5EHR) by plot statistics in PROCHECK

Parameters for evaluation	No. of residues	Score
Residues in most favoured regions [A,B,L]	415	89.2%
Residues in additional allowed regions [a,b,l,p]	44	10.1%
Residues in generously allowed regions [~a,~b,~l,~p]	2	0.5%
Residues in disallowed regions	1	0.2%
Number of non-glycine and non-proline residues	436	100.00%
Number of end-residues (excl. Gly and Pro)	8	-
Number of glycine residues (shown as triangles)*	35	-
Number of proline residues	19	-
Total number of residues	498	-

*Refer to the Figure A2 for details.

A brief protocol for evaluation of protein through SAVES server

The optimized protein structure of SHP2 (PDB ID: 5EHR) was uploaded to SAVES v6.0 web server (<https://saves.mbi.ucla.edu/>). Then protein was analyzed for different parameter (ERRAT, Verify 3D, PROVE, WHATCHEK, PROCHECK and CRYST). The Ramachandran plot from the results of PROCHECK was then downloaded. The graph of the Ramachandran plot is given in Figure A2.

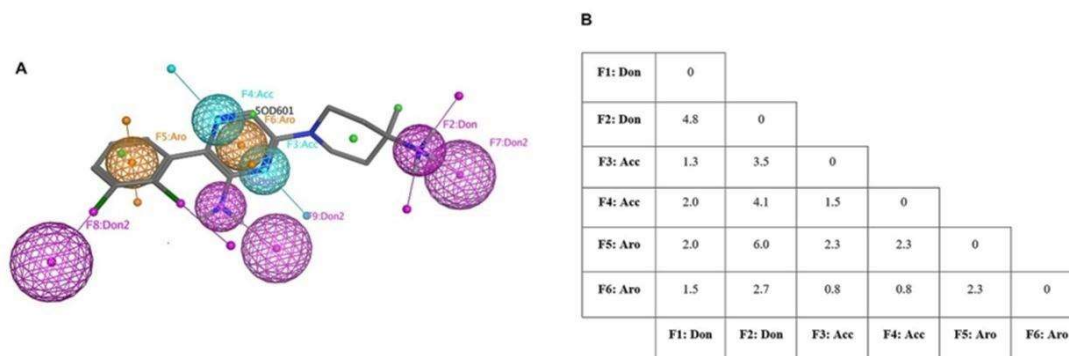


Figure A1. A) Pharmacophore model of co-crystallized **SHP099** showing the hypothetical spheres developed by MOE. Pink sphere = donor, blue sphere = acceptor, orange sphere = aromatic; B) The distance between the pharmacophoric features (in angstrom, Å) is presented in the distance matrix. Aro: Aromatic; don: donor; hyd: hydrophobic; acc: acceptor.

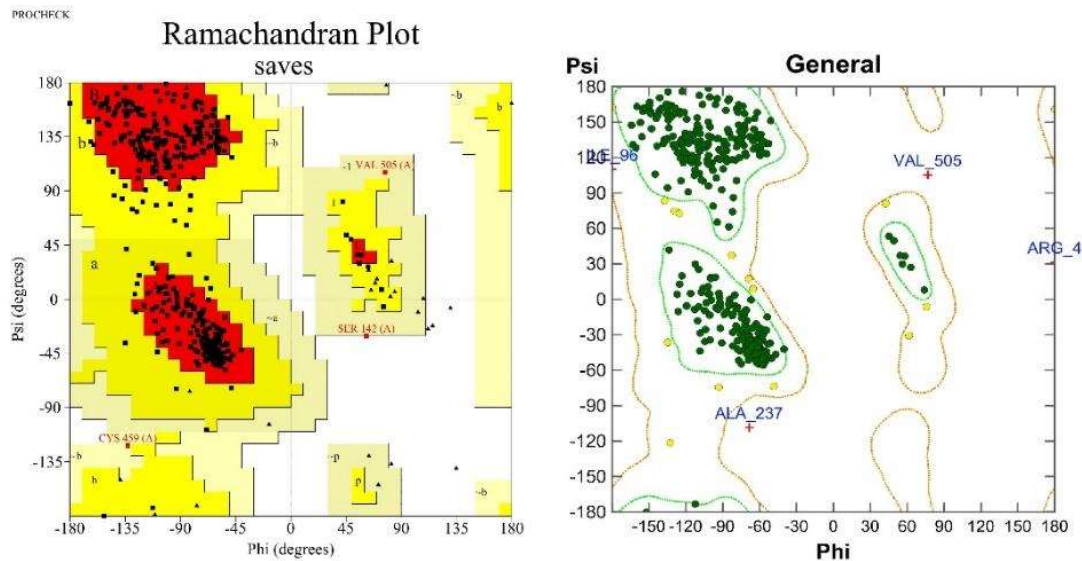


Figure A2. Ramachandran plot of SHP2 (PDB ID: 5EHR) obtained from PROCHECK (red & yellow) and phi-psi plot from MOE (black & white), respectively

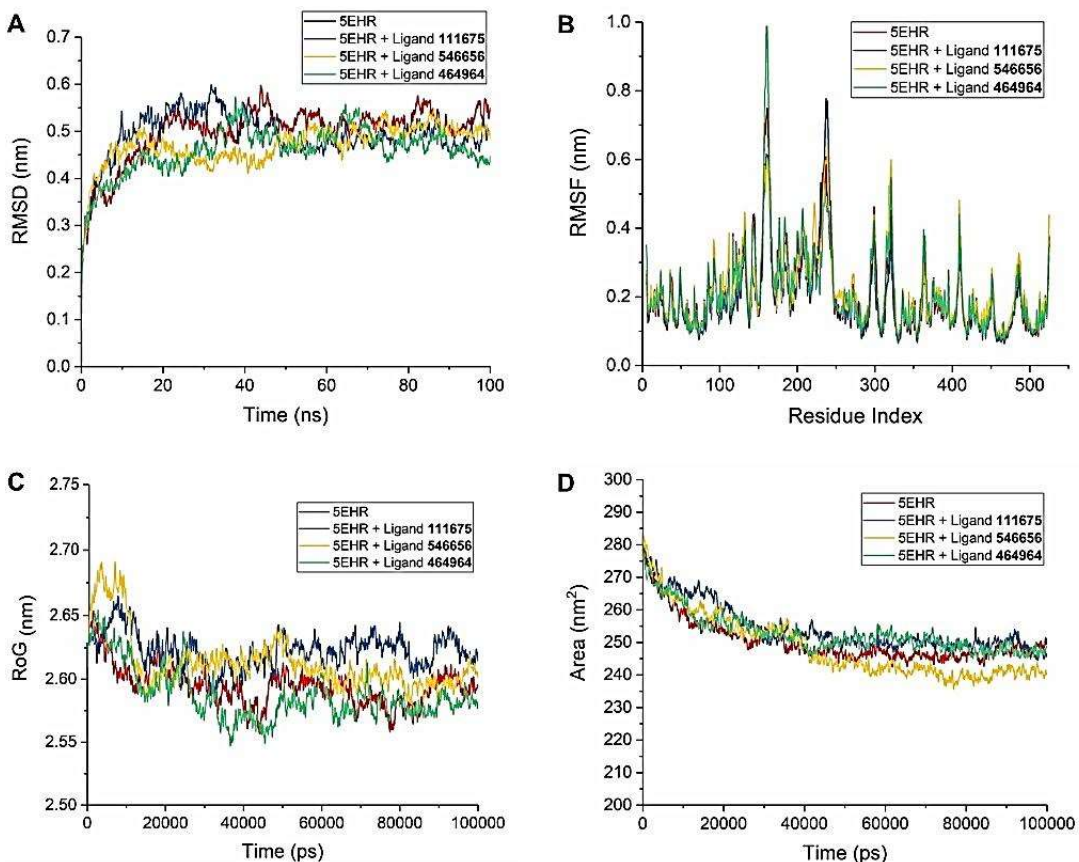


Figure A3. MD simulation results of most and least active hits; (A) RMSD of apoprotein and complex with ligands **111675**, **546656** and **464964**; (B) RMSF of apoprotein and complex with ligands **111675**, **546656** and **464964**; (C) RoG of

apoprotein and complexes of ligand **111675**, **546656** and **464964**; (D) solvent accessible surface area of apoprotein and complexes of ligand **111675**, **546656** and **464964**.

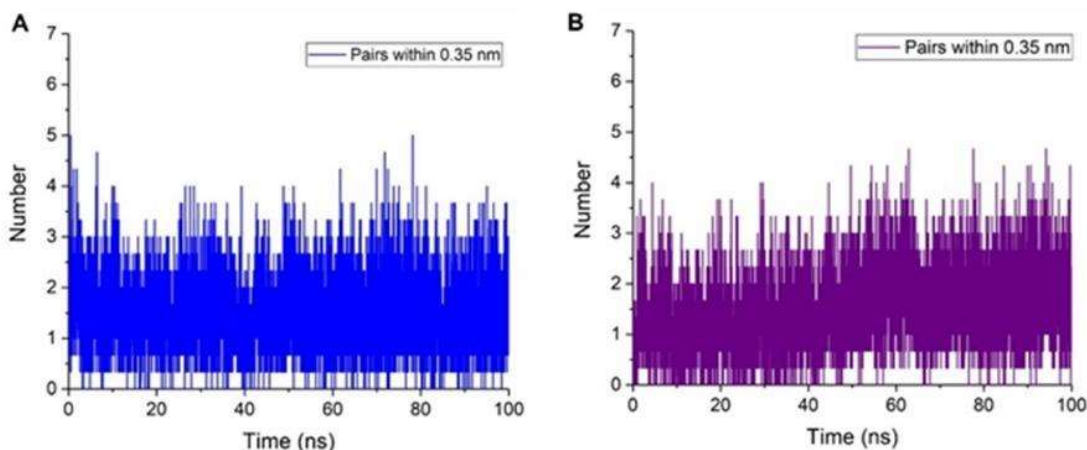


Figure A4. Hydrogen bonding analysis of 5EHR complexed with A) compound **111675** (shown in blue) and B) compound **546656** (shown in purple)

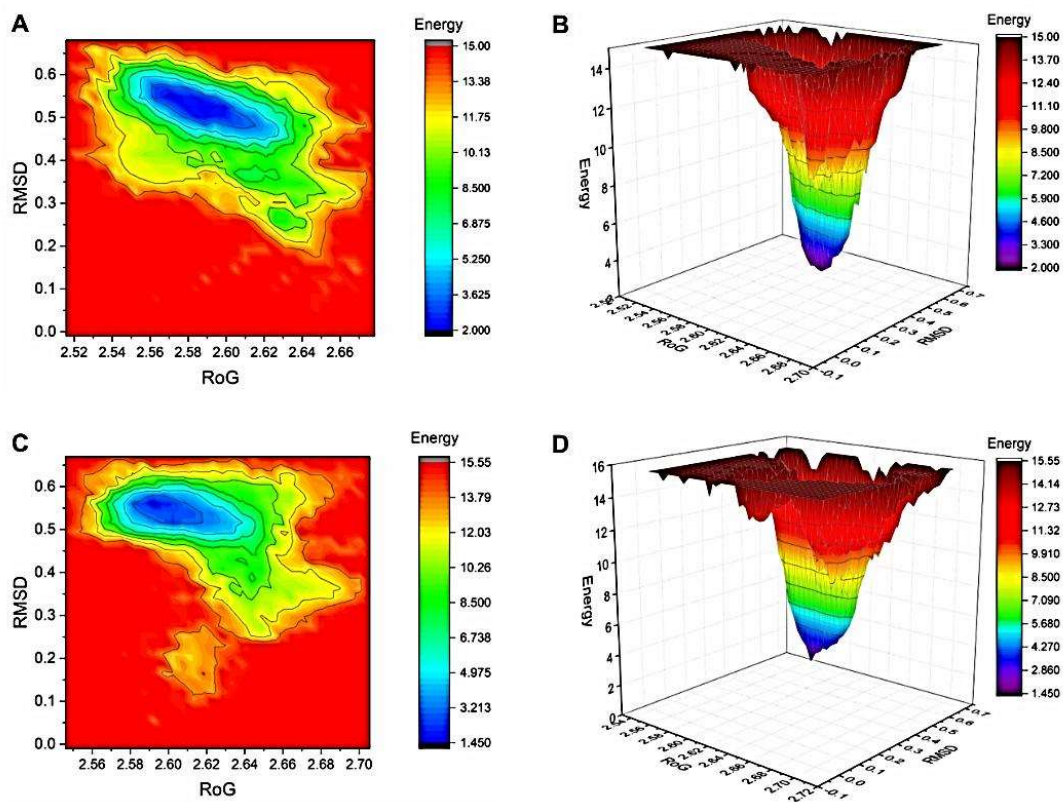


Figure A5. Free energy landscape (FEL) diagram of 5EHR apoprotein and its respective complexes; (A) 2D contour map of 5EHR; (B) funnel like 3D energy projection diagram of 5EHR; (C) 2D contour map of 5EHR and ligand **111675** complex; (D) funnel like 3D energy projection diagram of 5EHR and ligand **111675** complex

B. Spectral data of selected synthesized compounds

B1. Thioacetamide-tethered 1,3,4-thiadiazole-1,2,4-triazole hybrids (STT series)

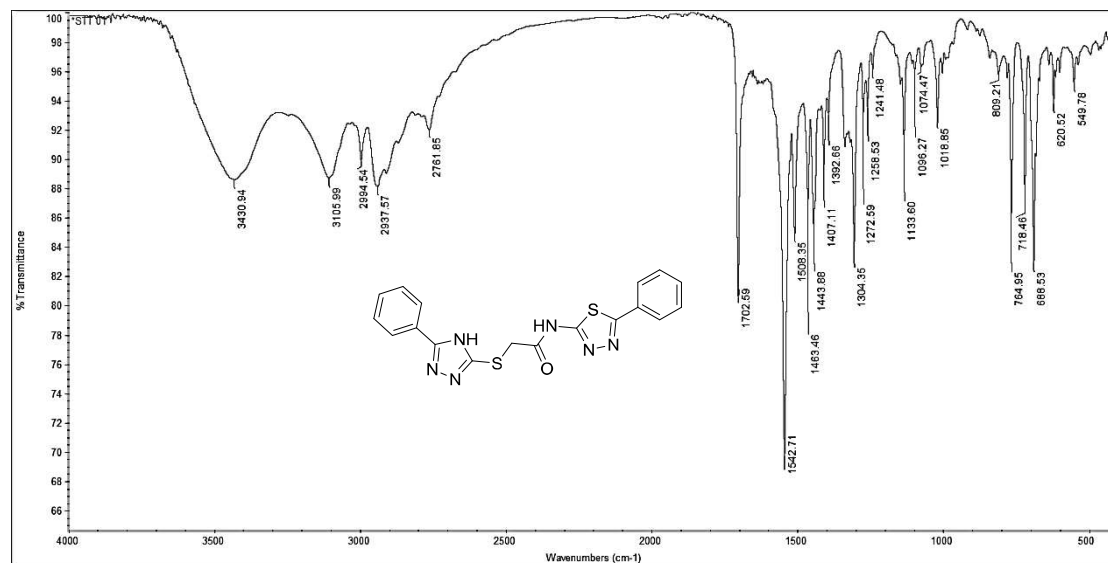
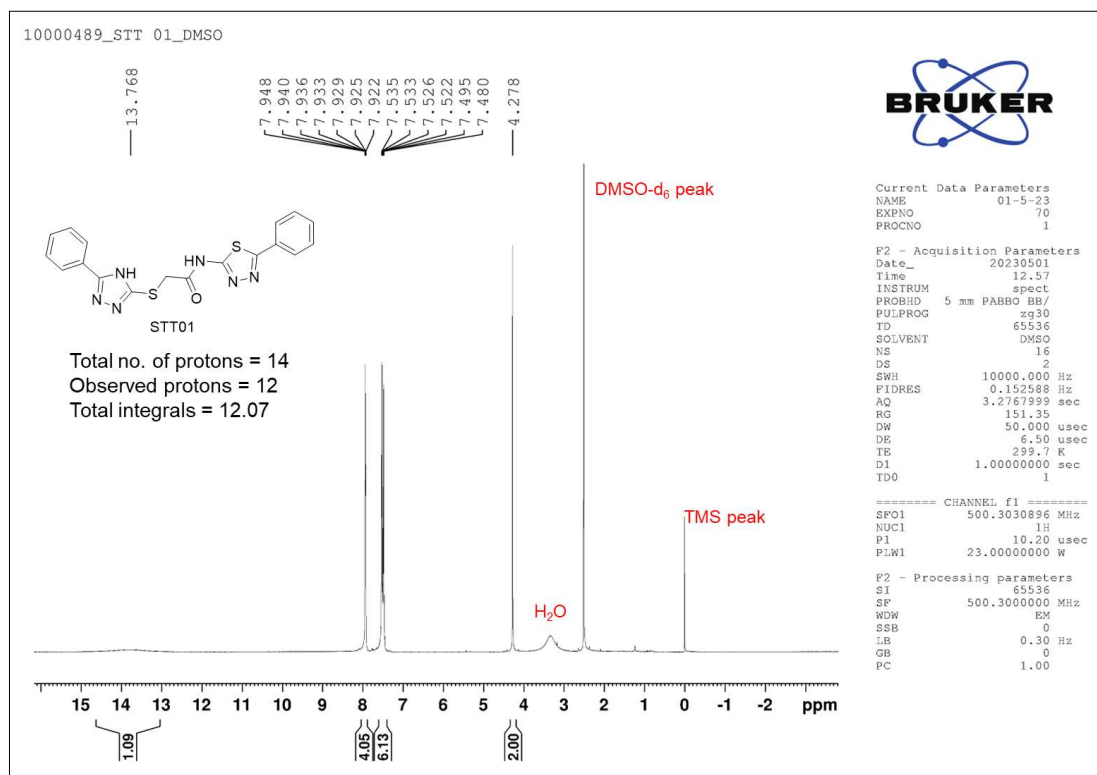


Figure A6. FTIR spectrum of compound STT01

Figure A7. ¹H NMR spectrum of compound STT01

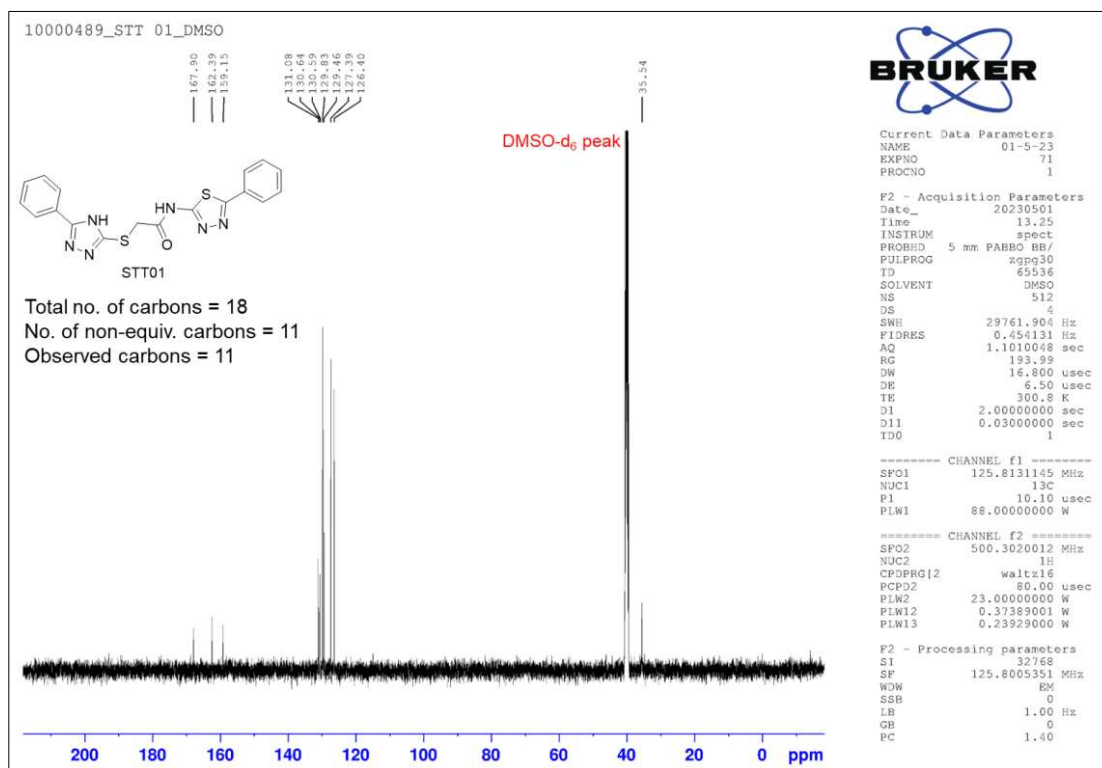
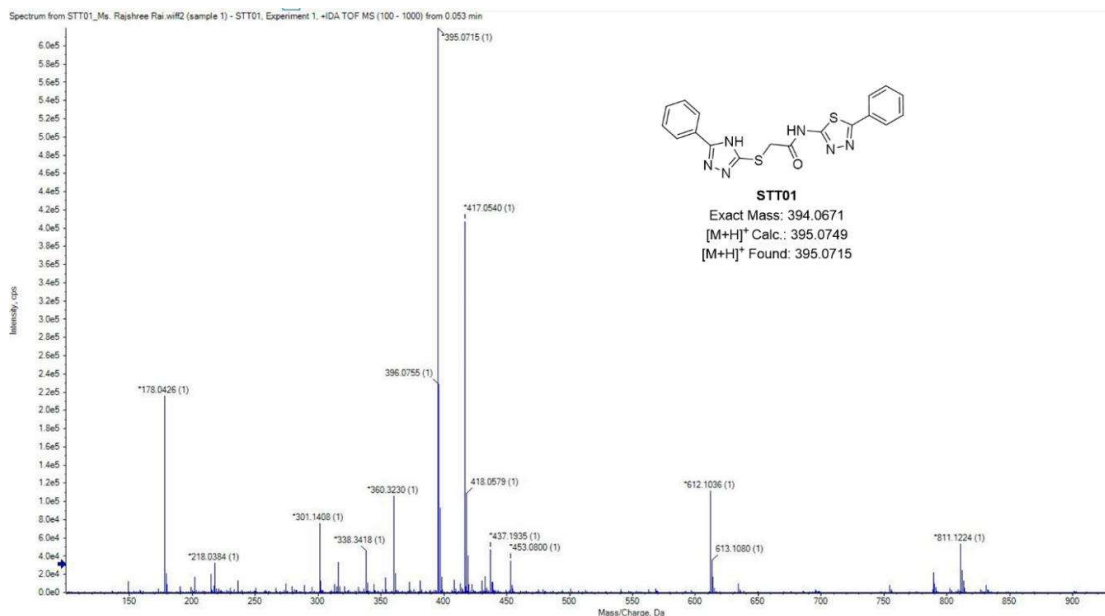
Figure A8. ¹³C NMR spectrum of compound STT01

Figure A9. HRMS spectrum of compound STT01

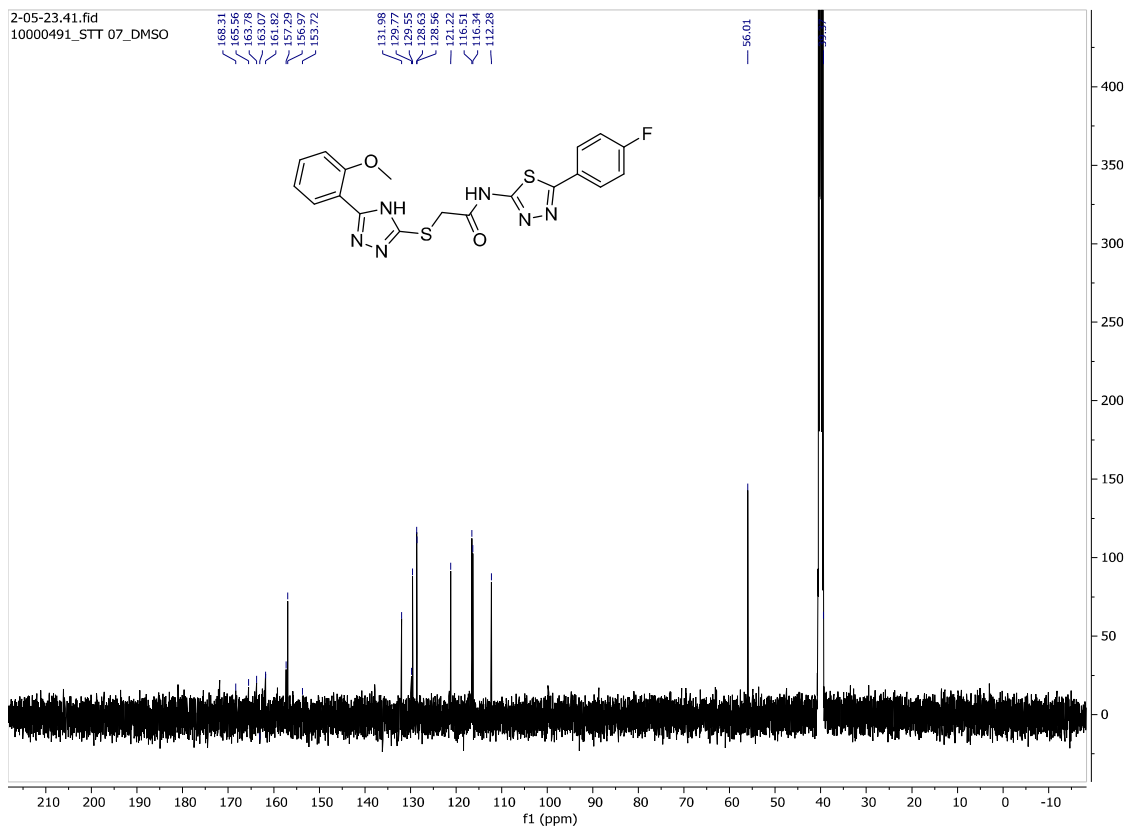


Figure A12. ^{13}C NMR spectrum of compound STT07

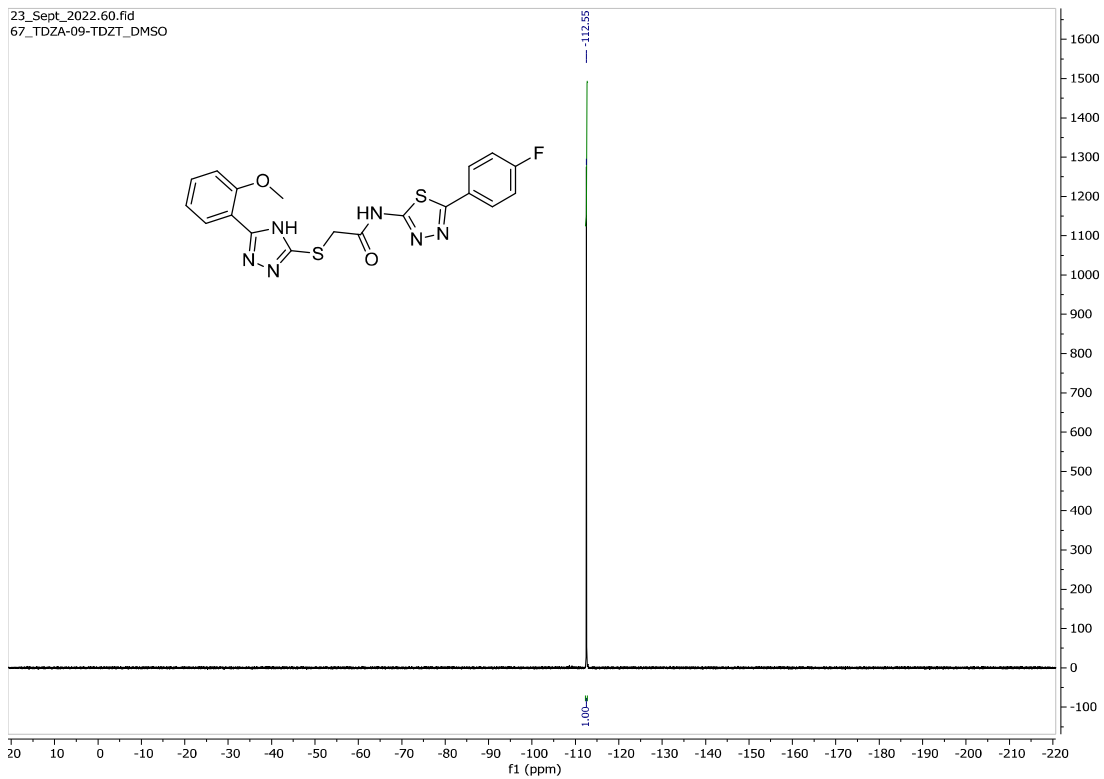


Figure A13. ^{19}F NMR spectrum of compound STT07

Appendix

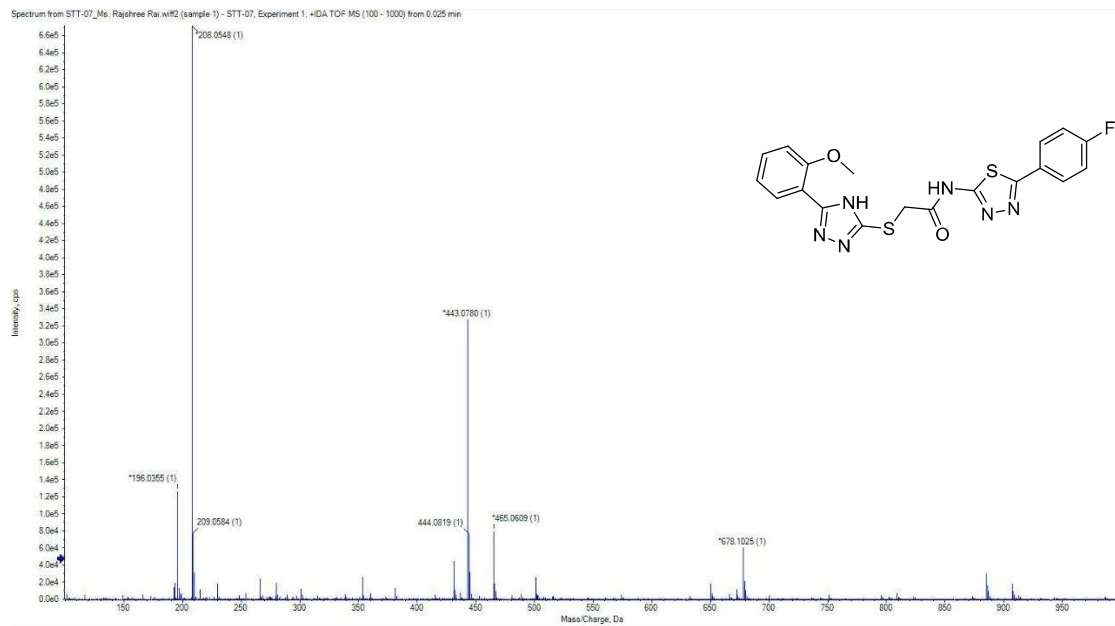


Figure A14. HRMS spectrum of compound STT07

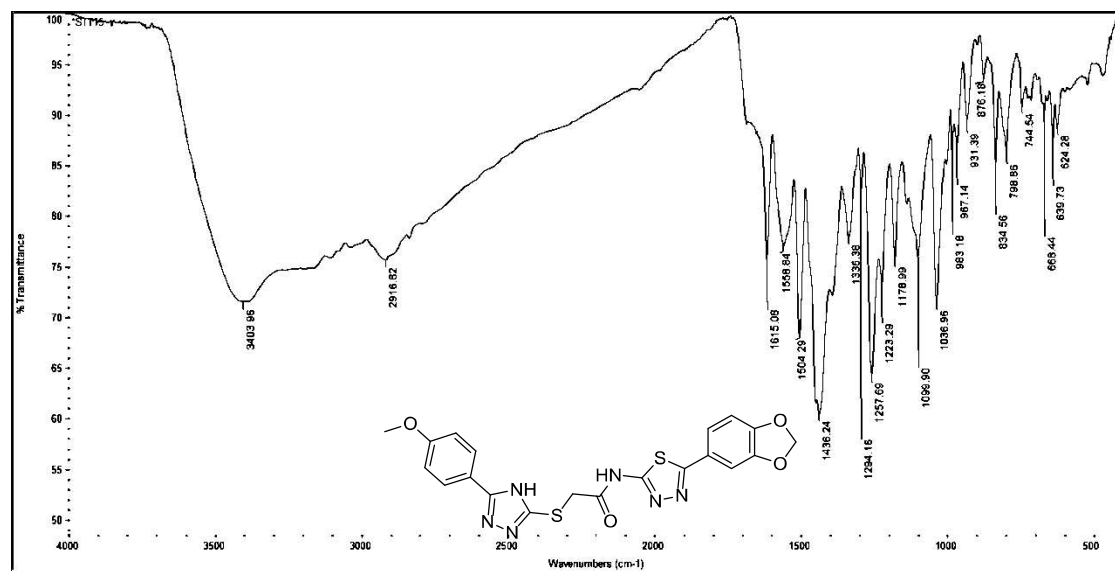


Figure A15. FTIR spectrum of compound STT13

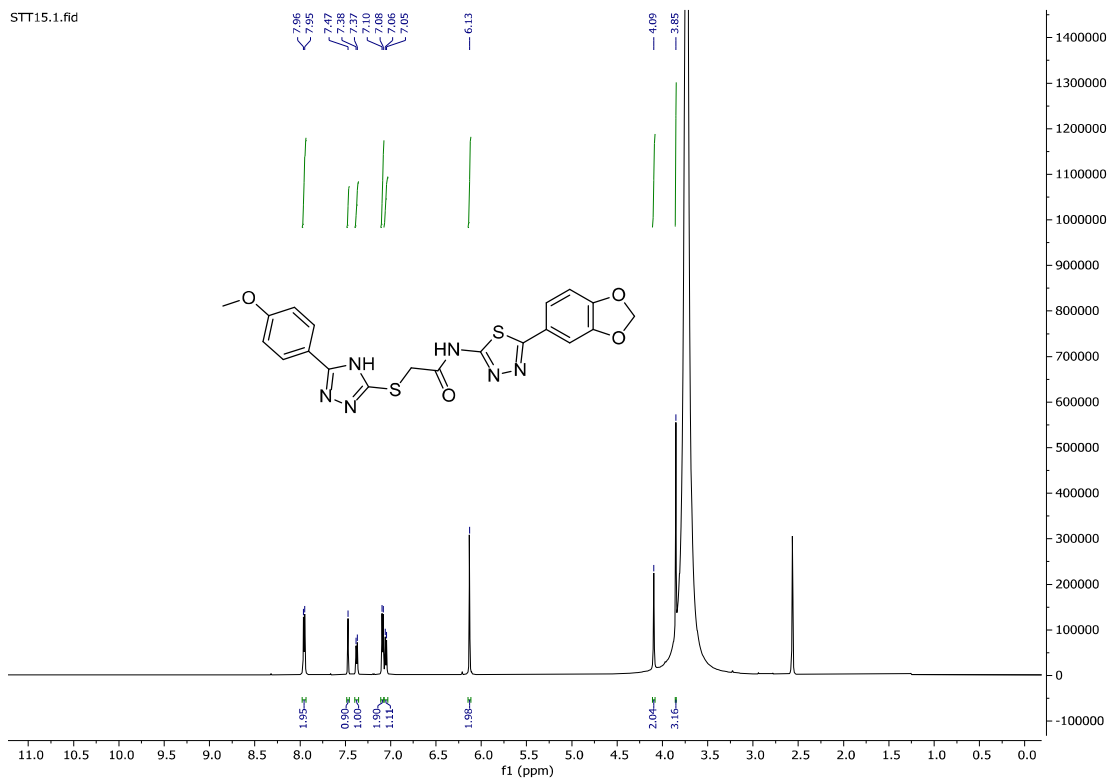


Figure A16. ^1H NMR spectrum of compound STT13

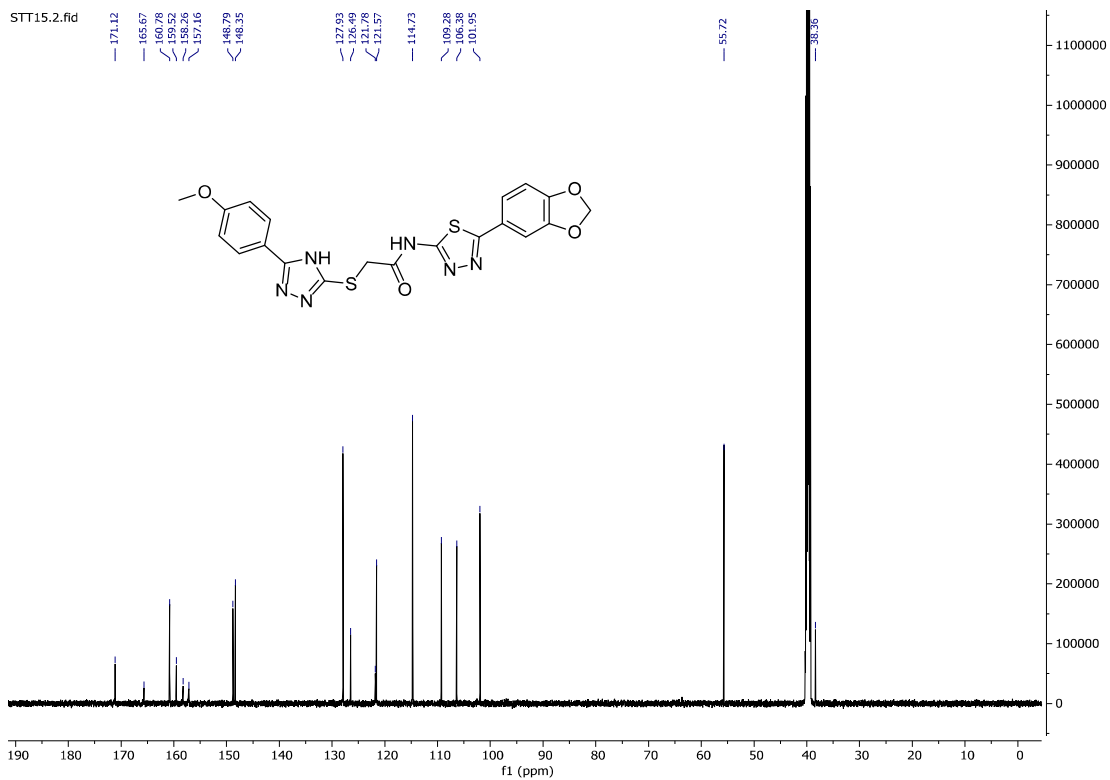


Figure A17. ^{13}C NMR spectrum of compound STT13

Appendix

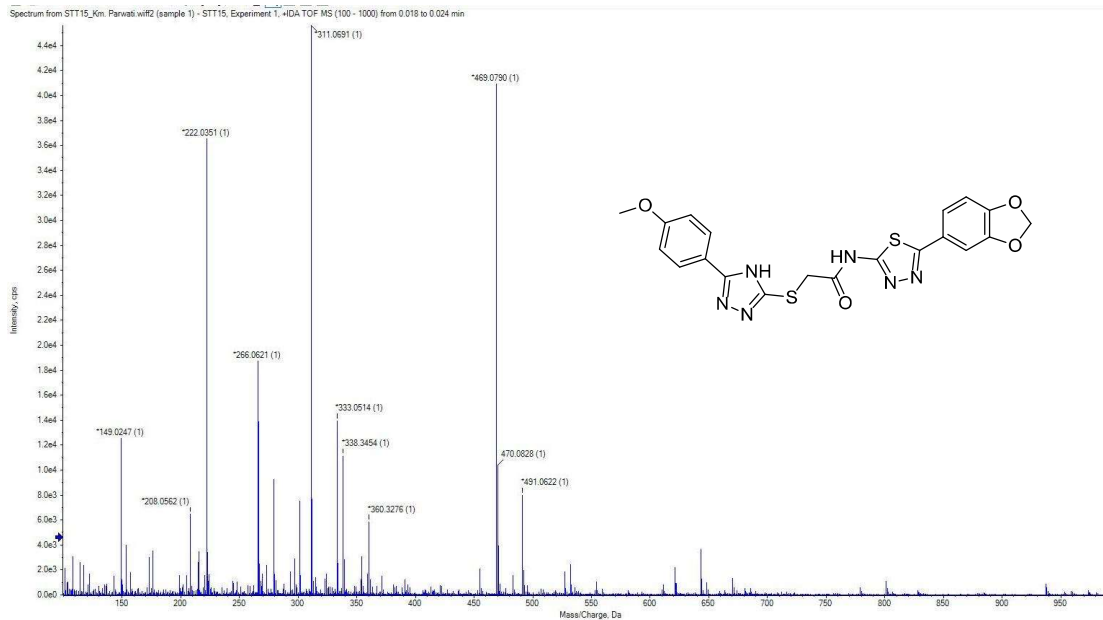


Figure A18. HRMS spectrum of compound STT13

B2. 5-(Substituted phenyl)-1,3,4-thiadiazole-2-amine derived sulphur-linked acetamides (STS series)

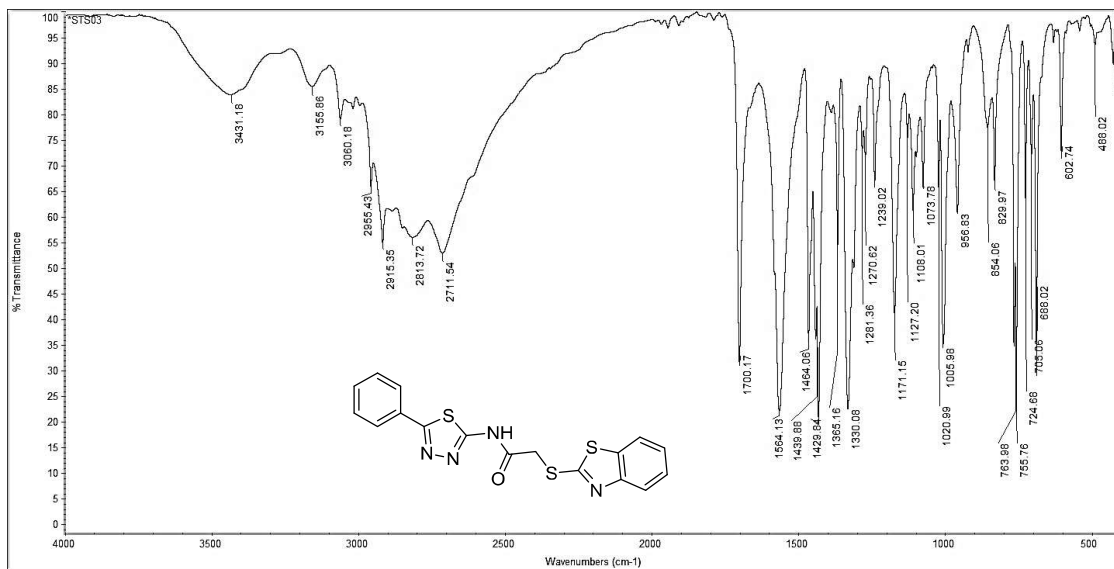


Figure A19. FTIR spectrum of compound STS03

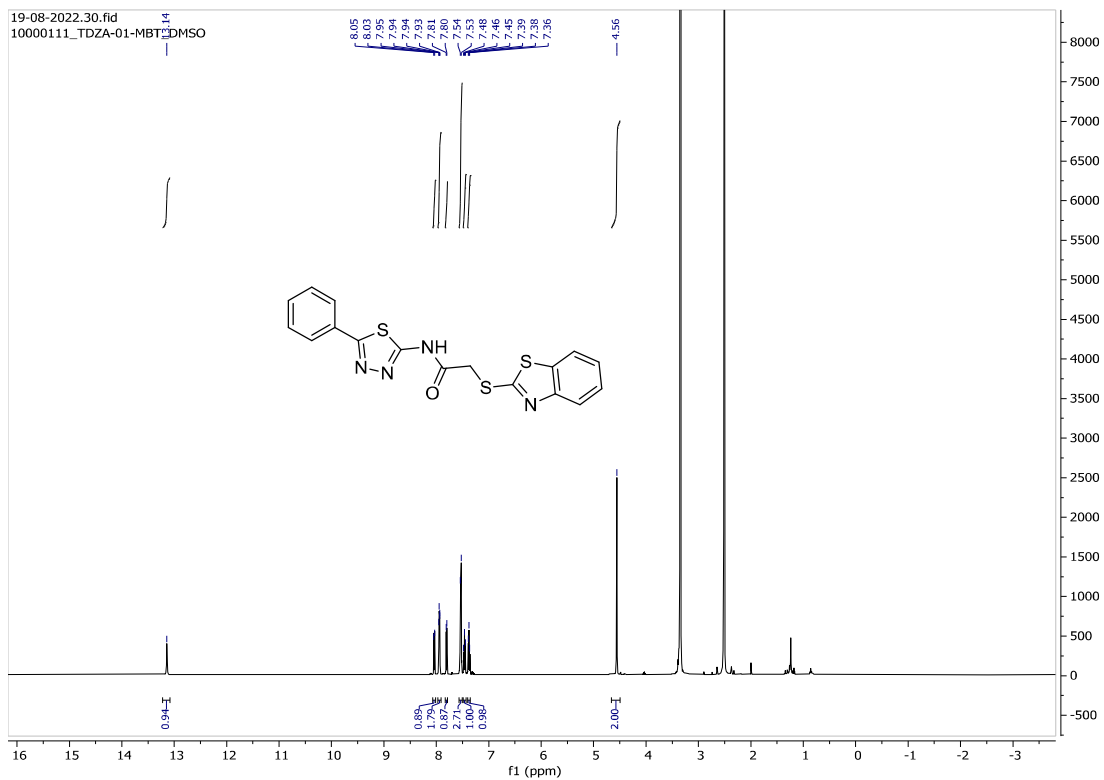


Figure A20. ^1H NMR spectrum of compound STS03

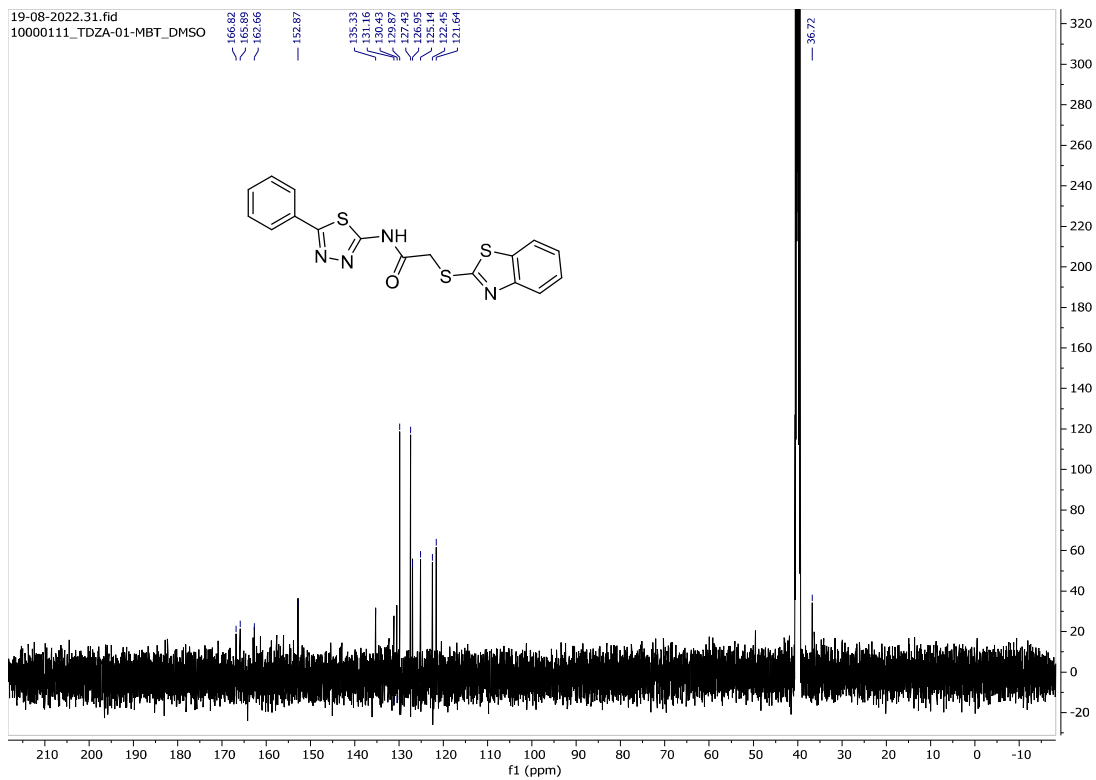


Figure A21. ^{13}C NMR spectrum of compound STS03

Appendix

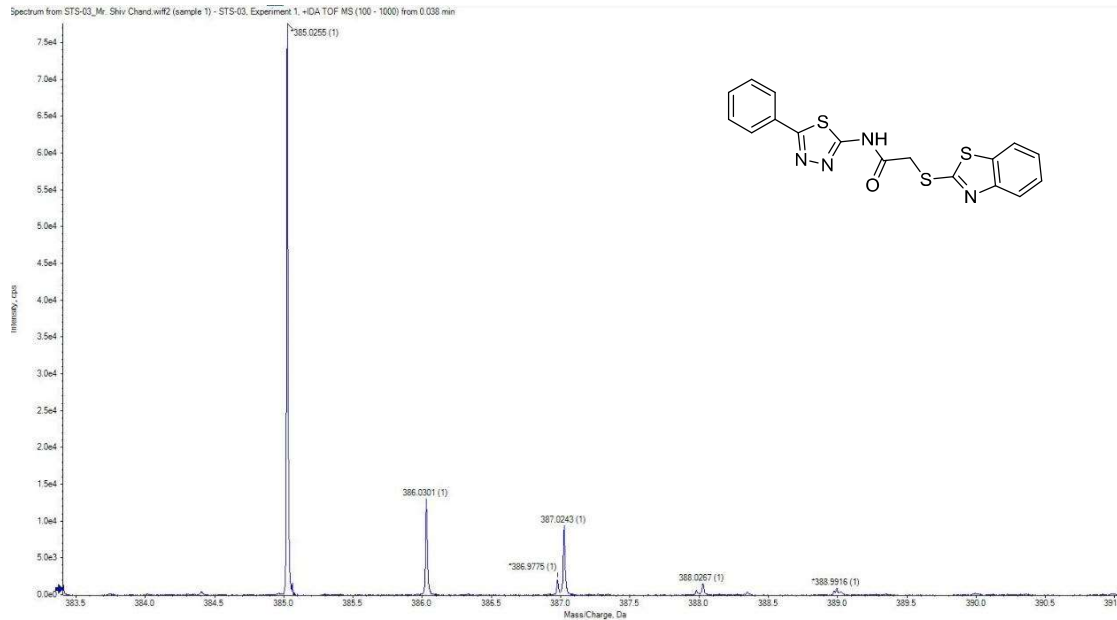


Figure A22. HRMS spectrum of compound STS03

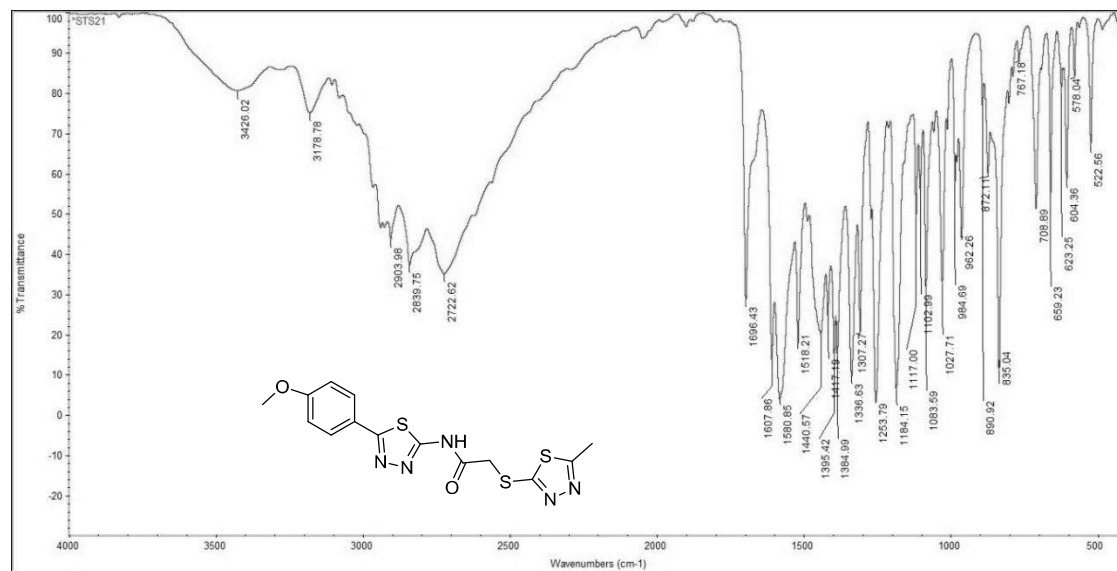


Figure A23. FTIR spectrum of compound STS11

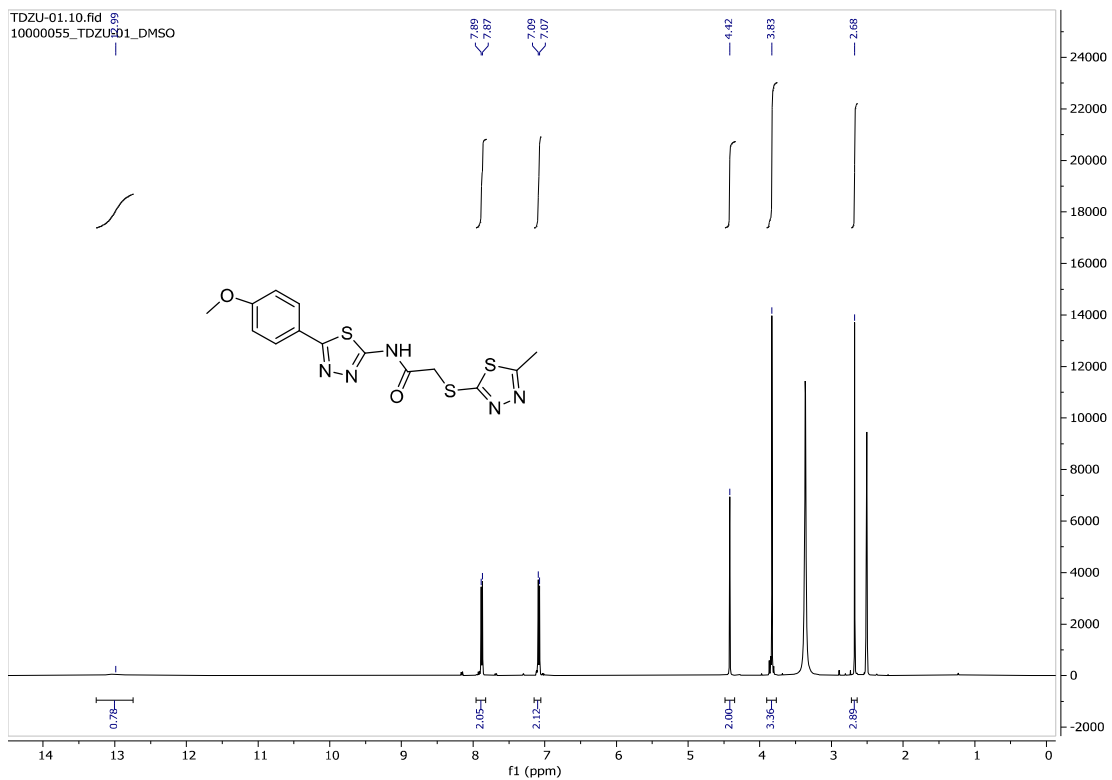


Figure A24. ^1H NMR spectrum of compound STS11

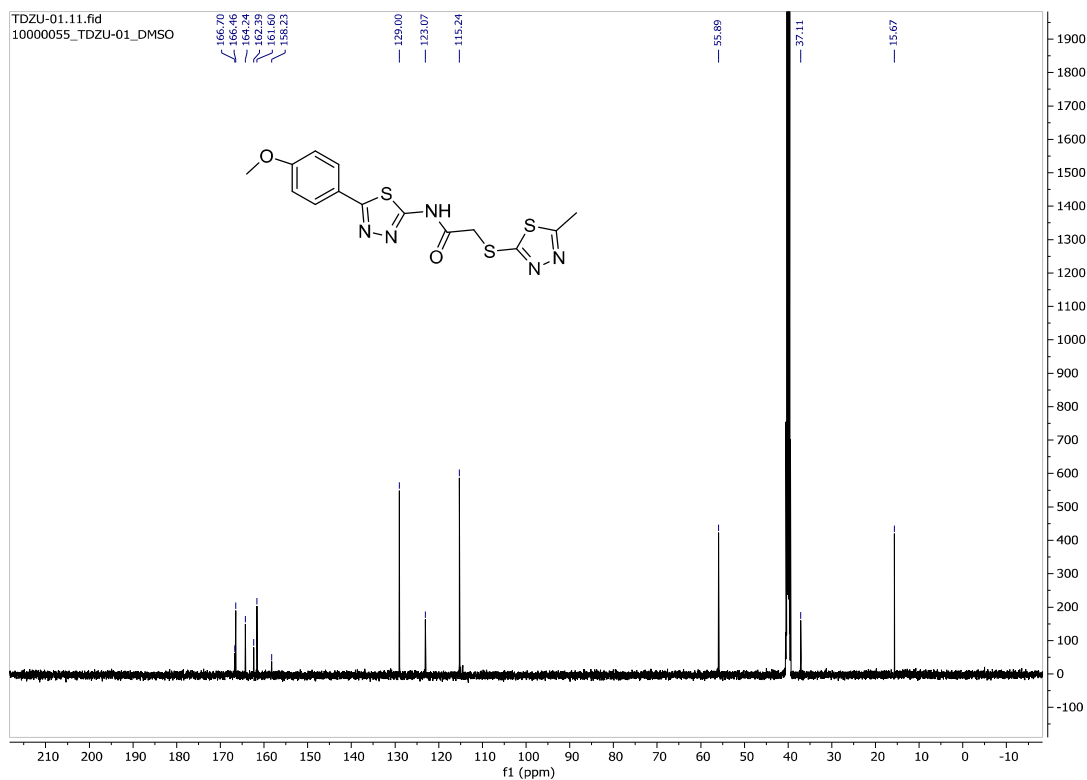


Figure A25. ^{13}C NMR spectrum of compound STS11

Appendix

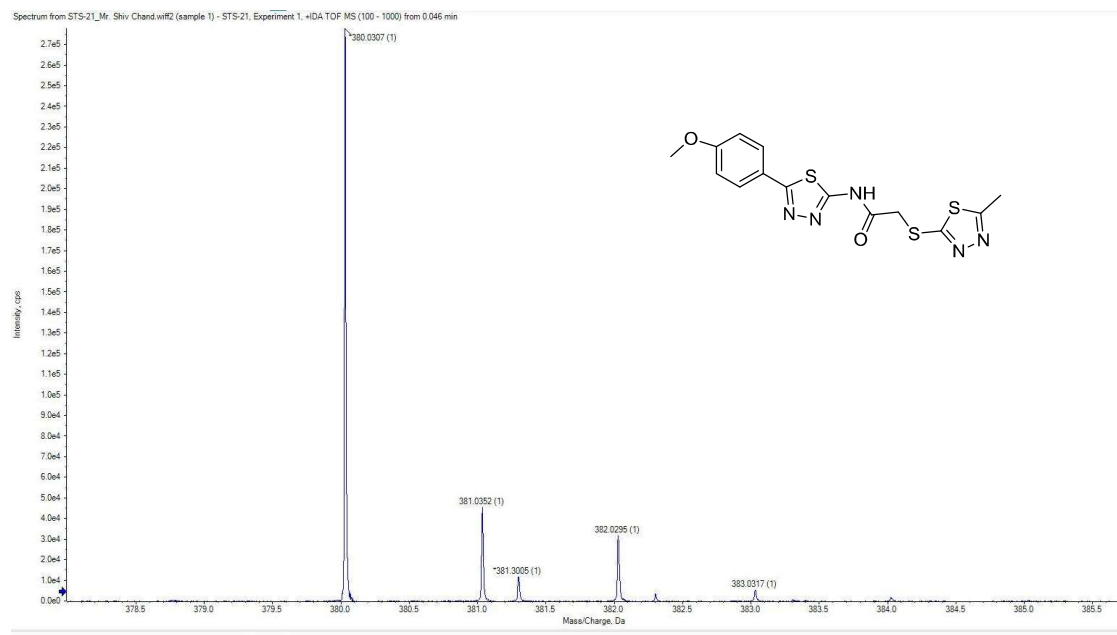


Figure A26. HRMS spectrum of compound STS11

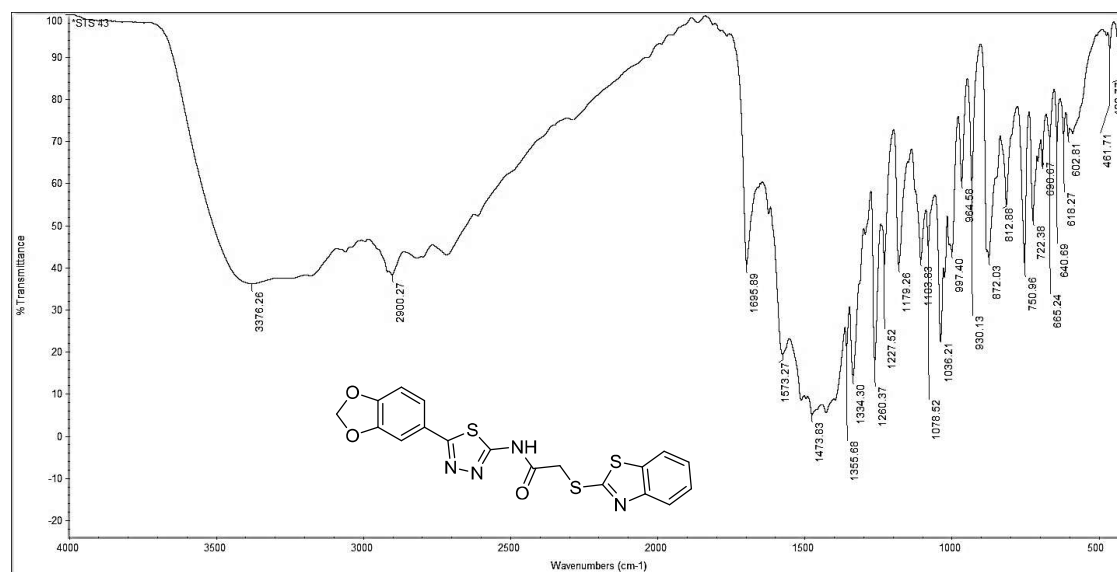


Figure A27. FTIR spectrum of compound STS23

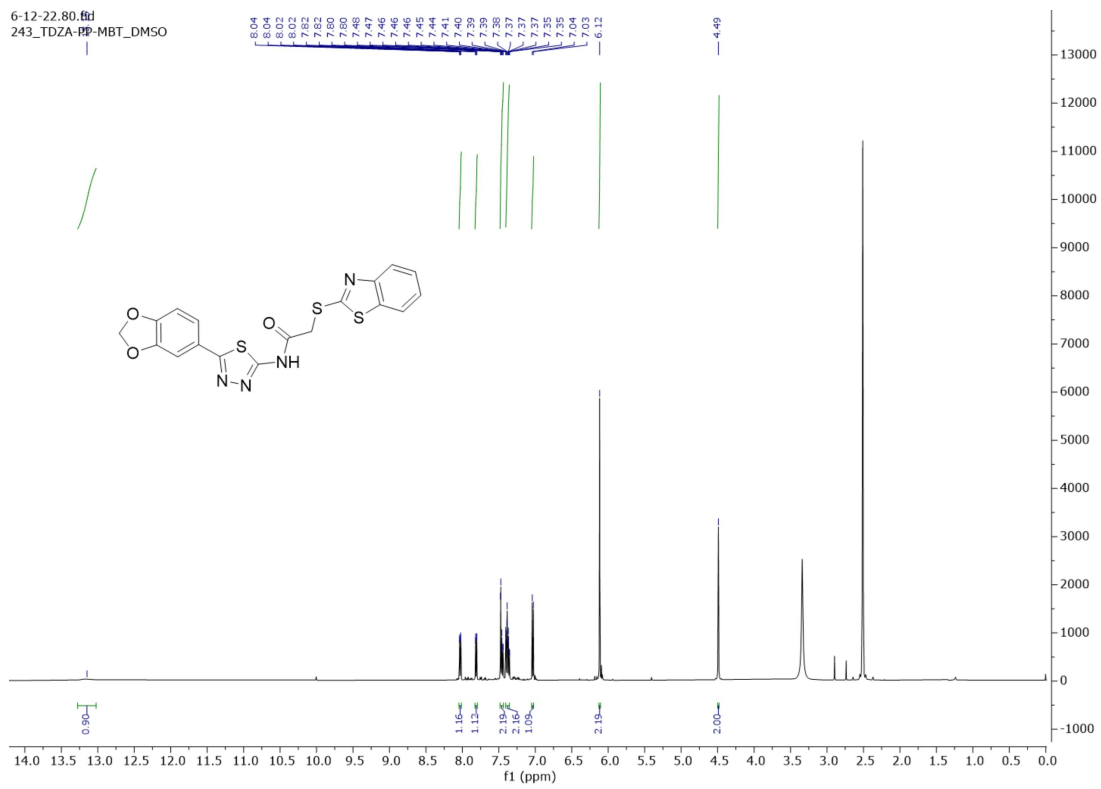


Figure A28. ^1H NMR spectrum of compound STS23

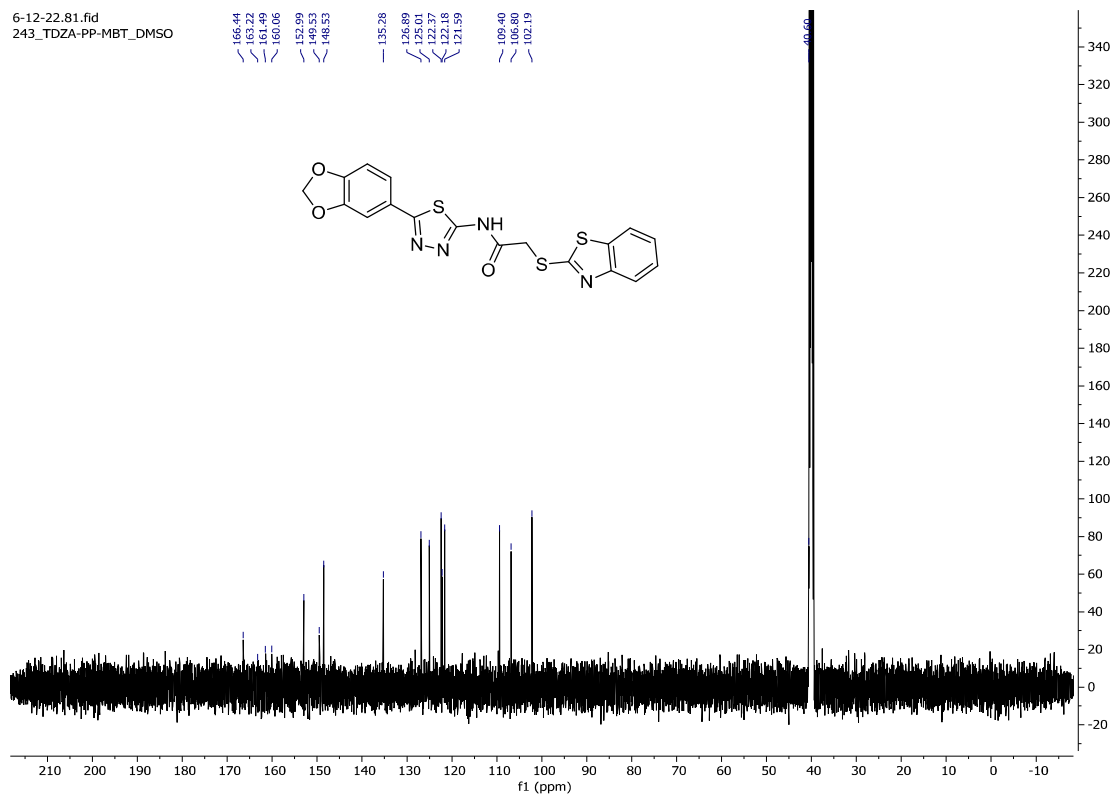


Figure A29. ^{13}C NMR spectrum of compound STS23

Appendix

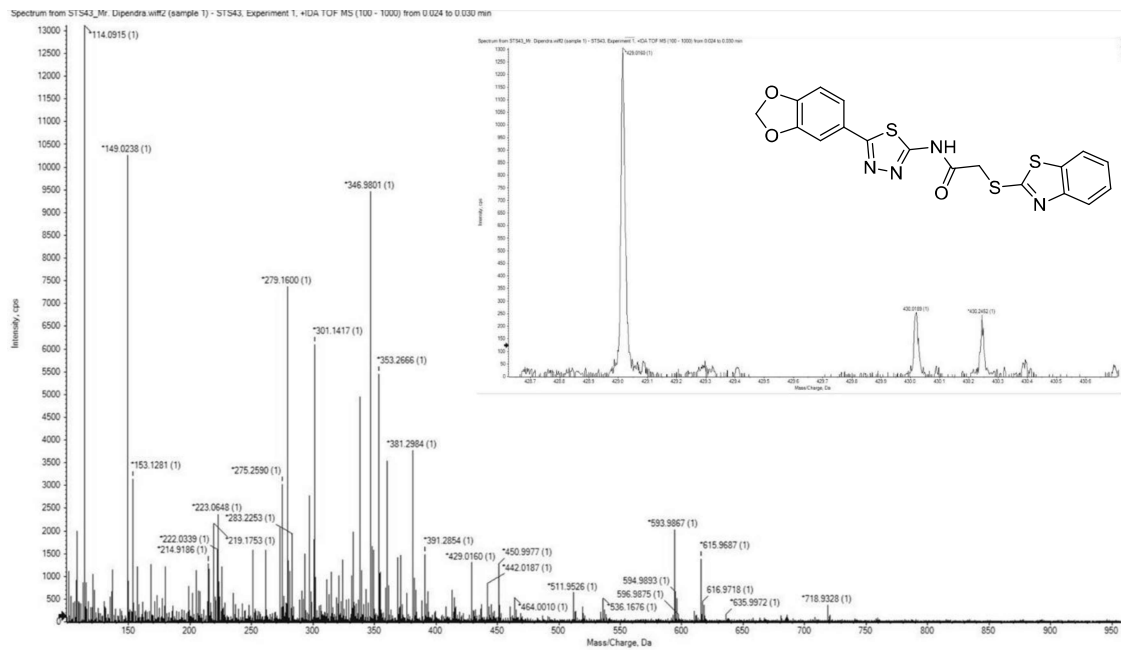


Figure A30. HRMS spectrum of compound STS23

B3. *S*-Acetohydrazone of 5-methyl-1,3,4-thiadiazole-2-thiol (TEH series)

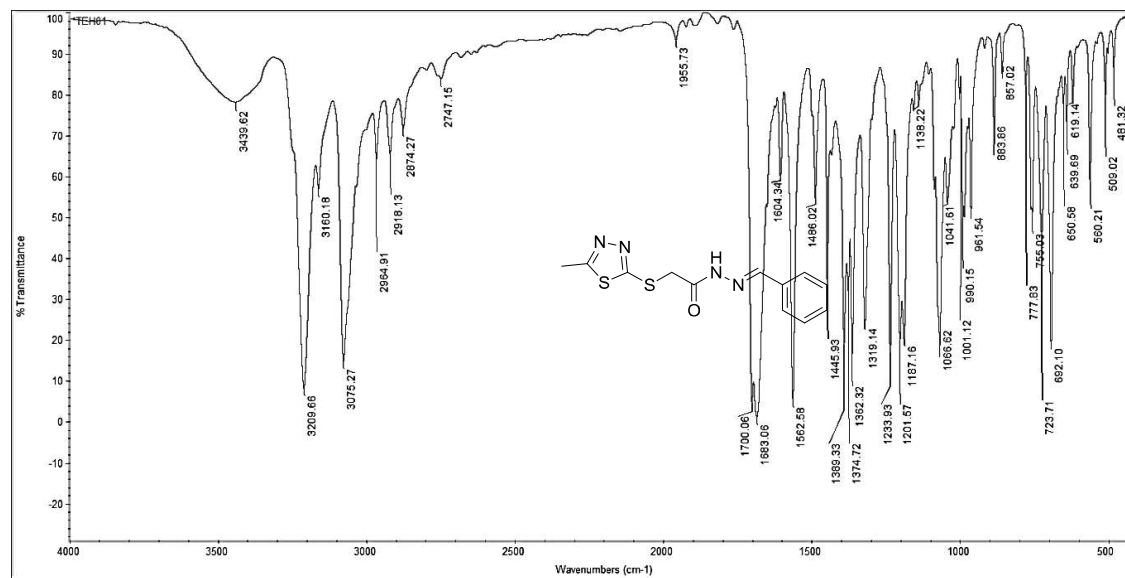


Figure A31. FTIR spectrum of compound TEH01

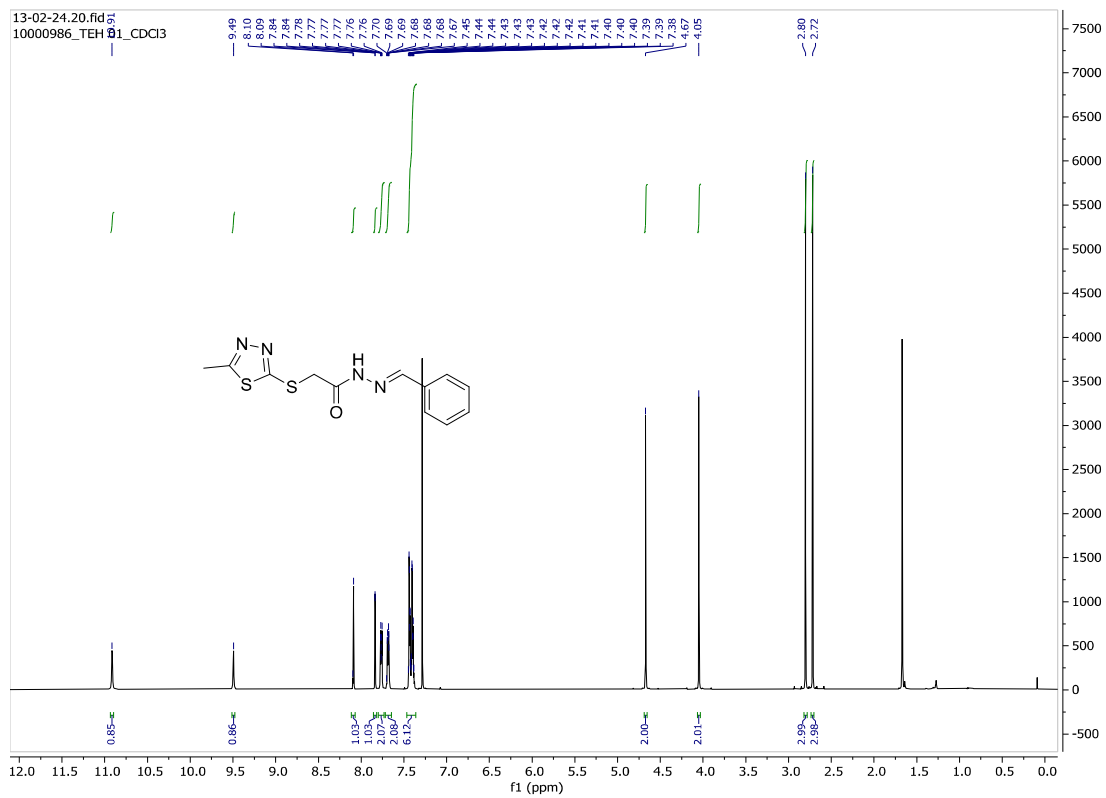


Figure A32. ^1H NMR spectrum of compound TEH01

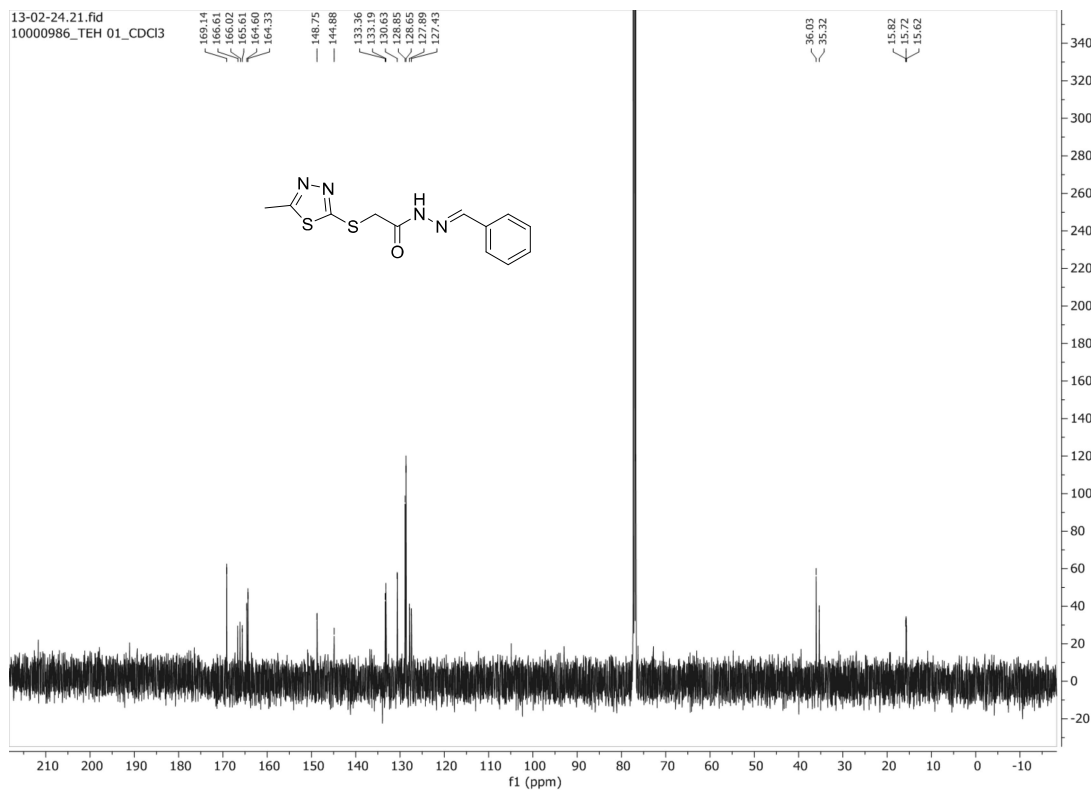


Figure A33. ^{13}C NMR spectrum of compound TEH01

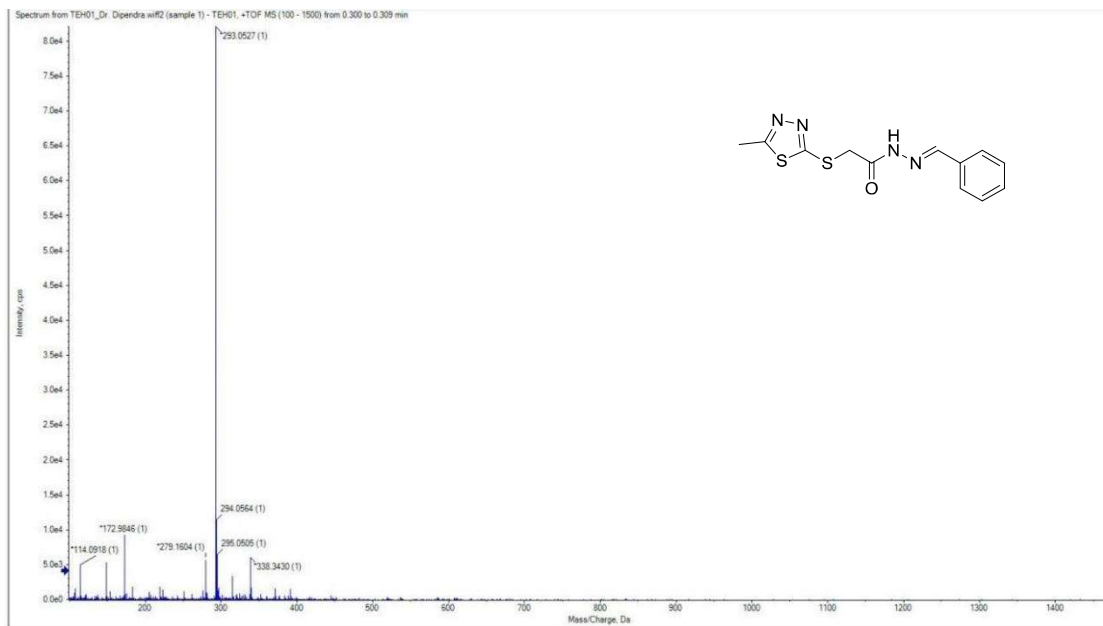


Figure A34. HRMS spectrum of compound TEH01

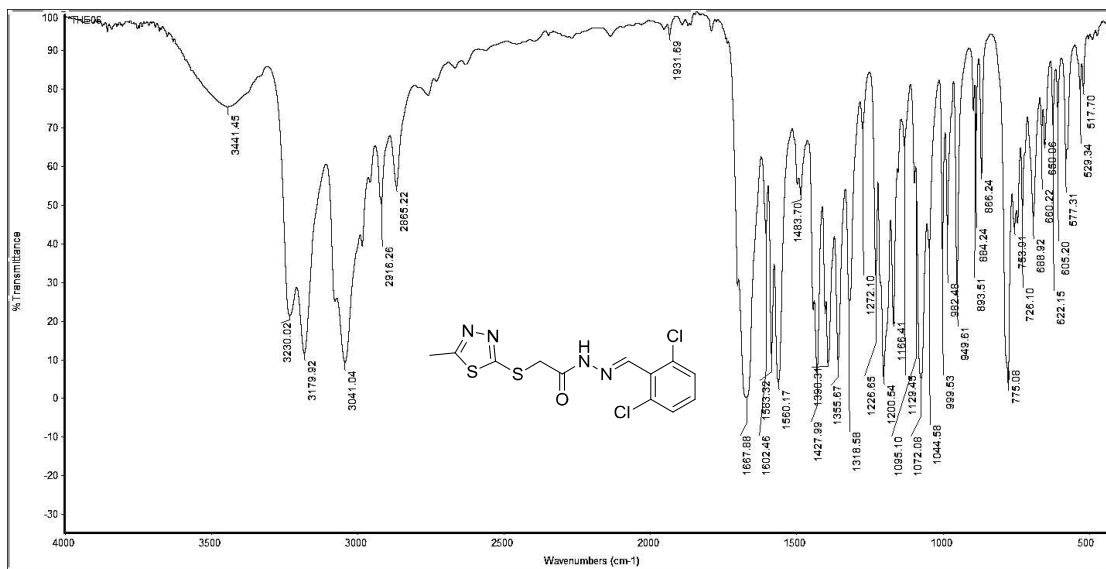
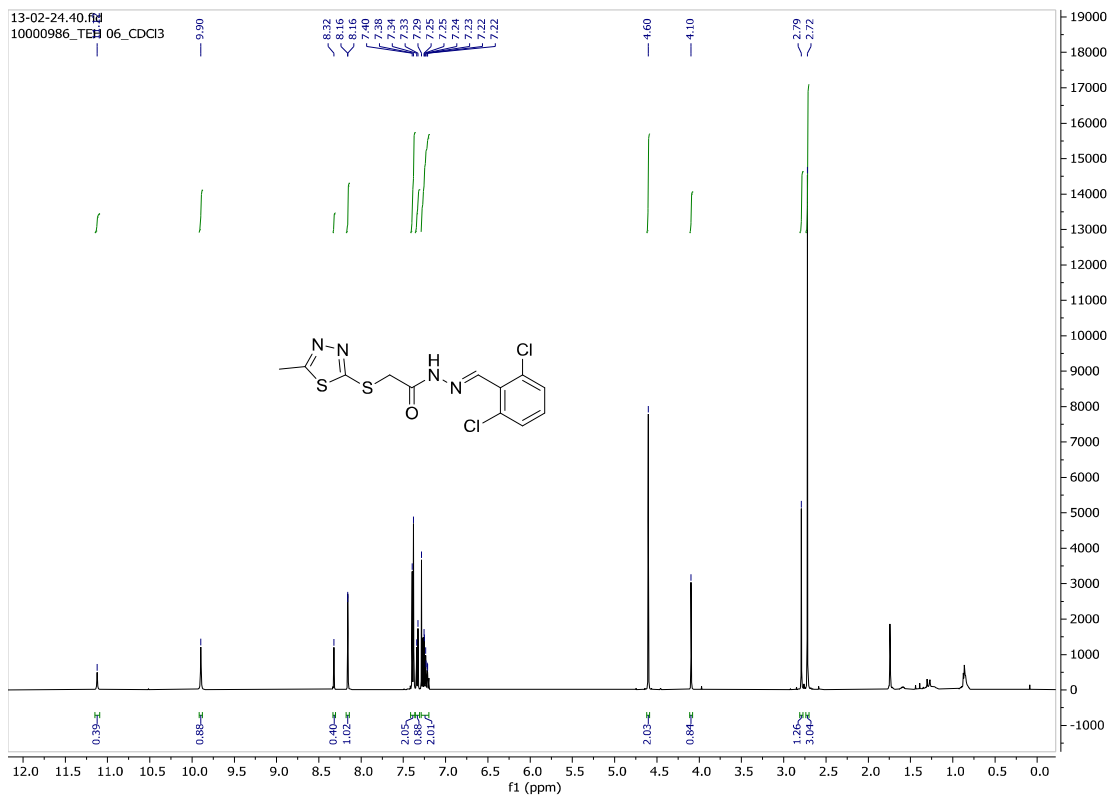
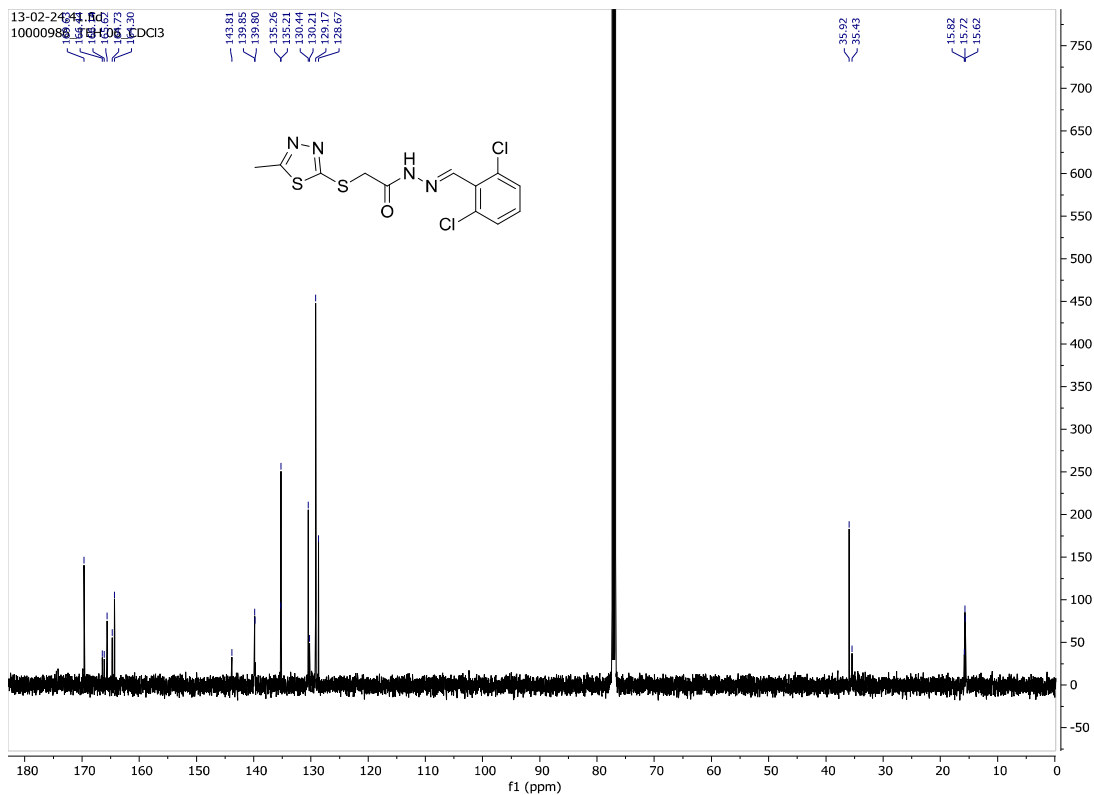


Figure A35. FTIR spectrum of compound TEH06

Figure A36. ^1H NMR spectrum of compound TEH06Figure A37. ^{13}C NMR spectrum of compound TEH06

Appendix

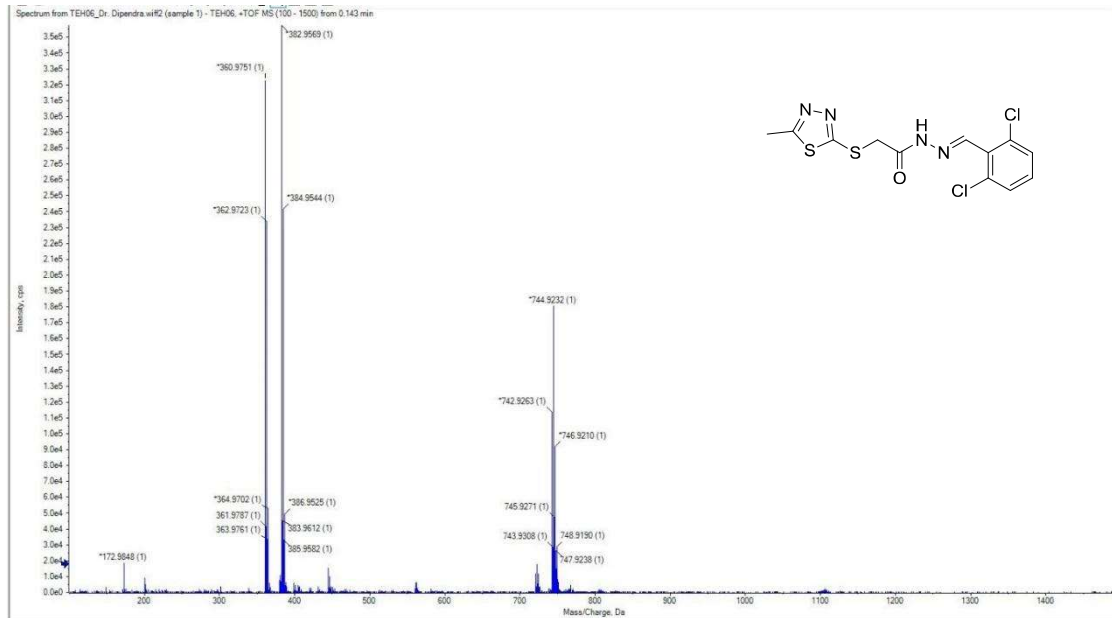


Figure A38. HRMS spectrum of compound TEH06

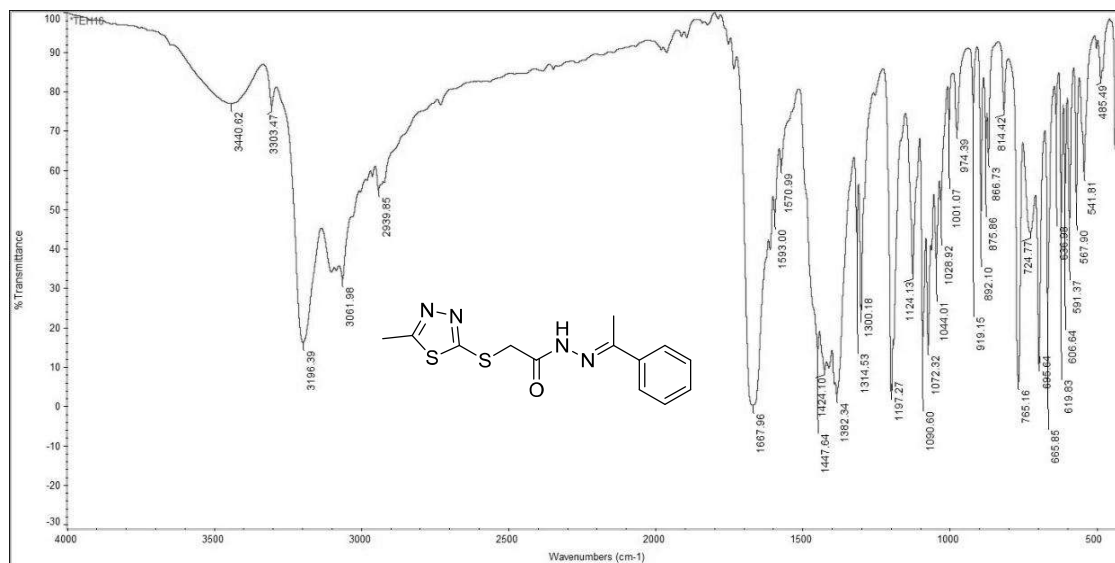


Figure A39. FTIR spectrum of compound TEH16

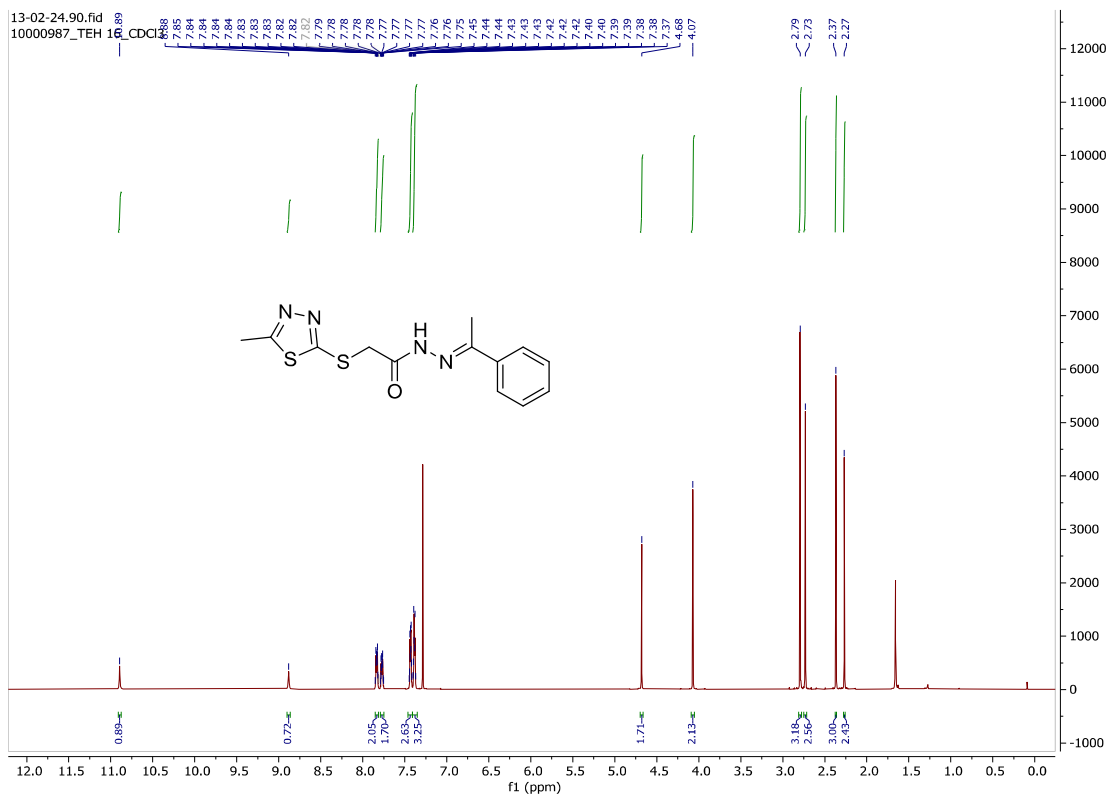


Figure A40. ^1H NMR spectrum of compound TEH16

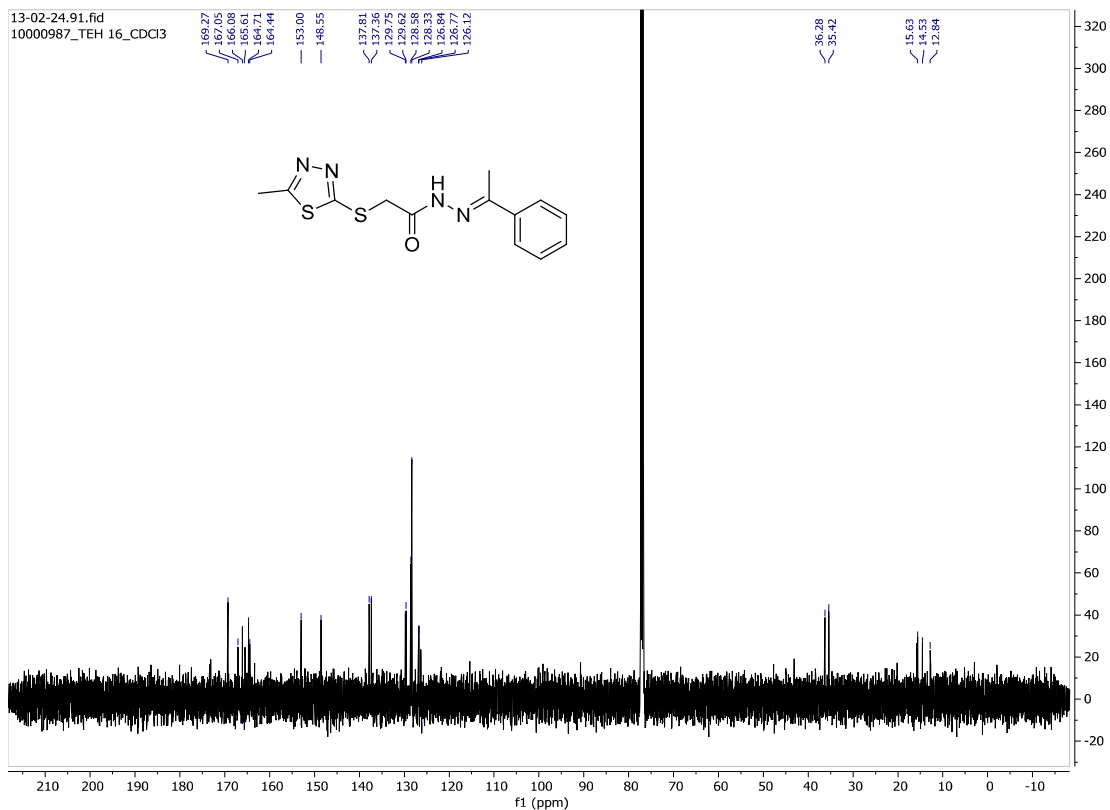


Figure A41. ^{13}C NMR spectrum of compound TEH16

Appendix

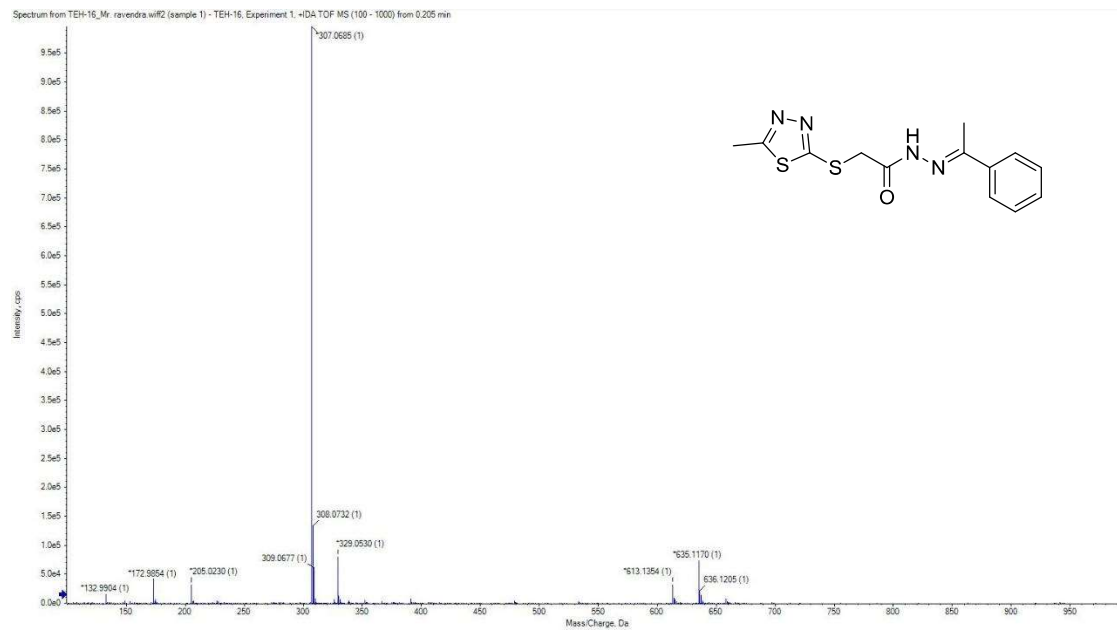


Figure A42. HRMS spectrum of compound **TEH16**

C. UV-based LogP determination

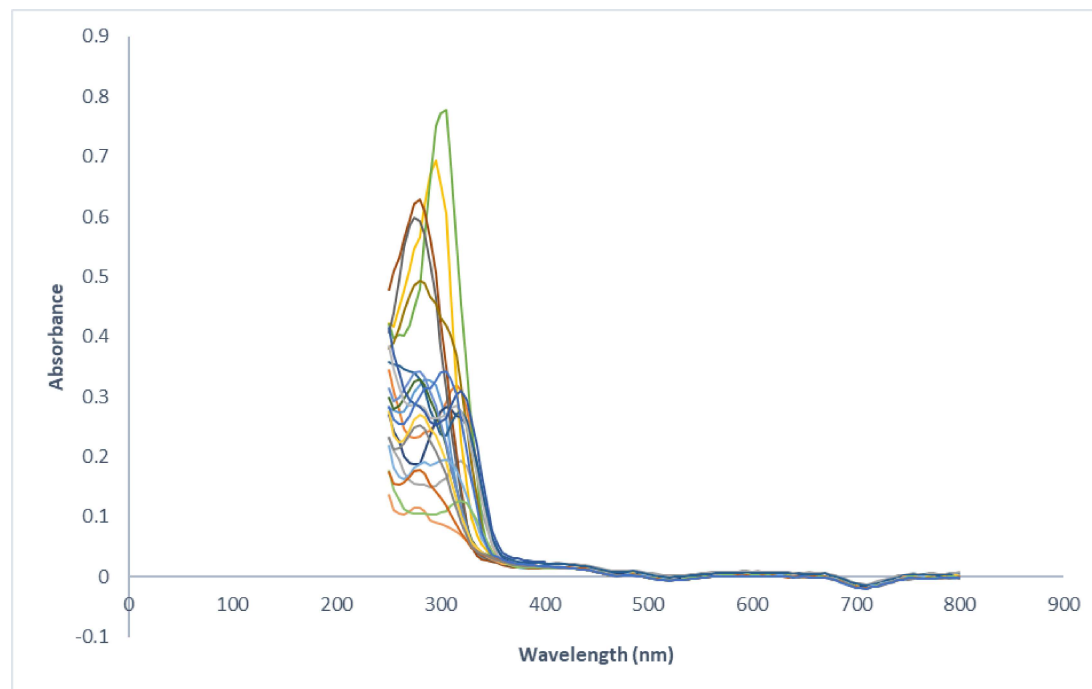


Figure A43. UV λ_{\max} spectra of **STT** series of compounds

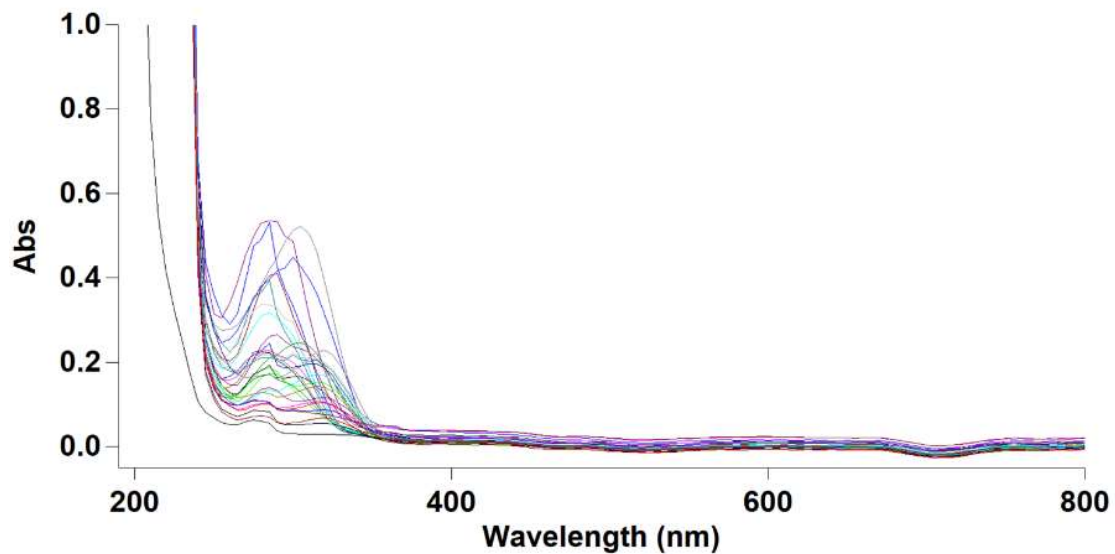


Figure A44. UV λ_{\max} spectra of STS series of compounds

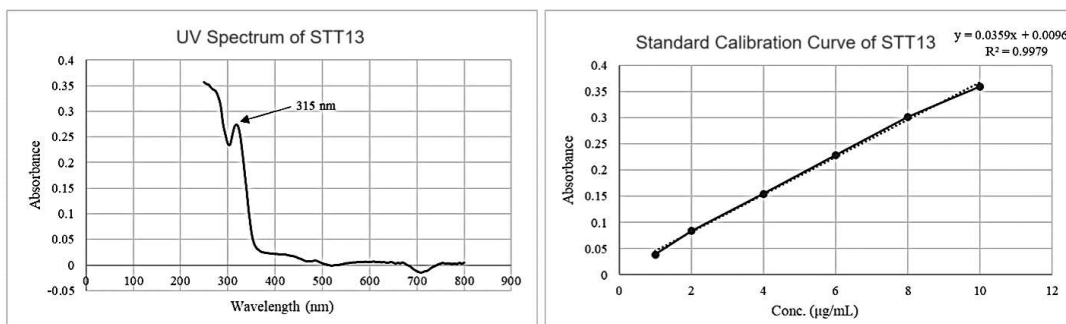


Figure A45. UV spectrum (at a concentration of 10 $\mu\text{g/mL}$ in HPLC grade methanol) and standard calibration curve (at $\lambda_{\max} = 315$ nm) of compound STT13

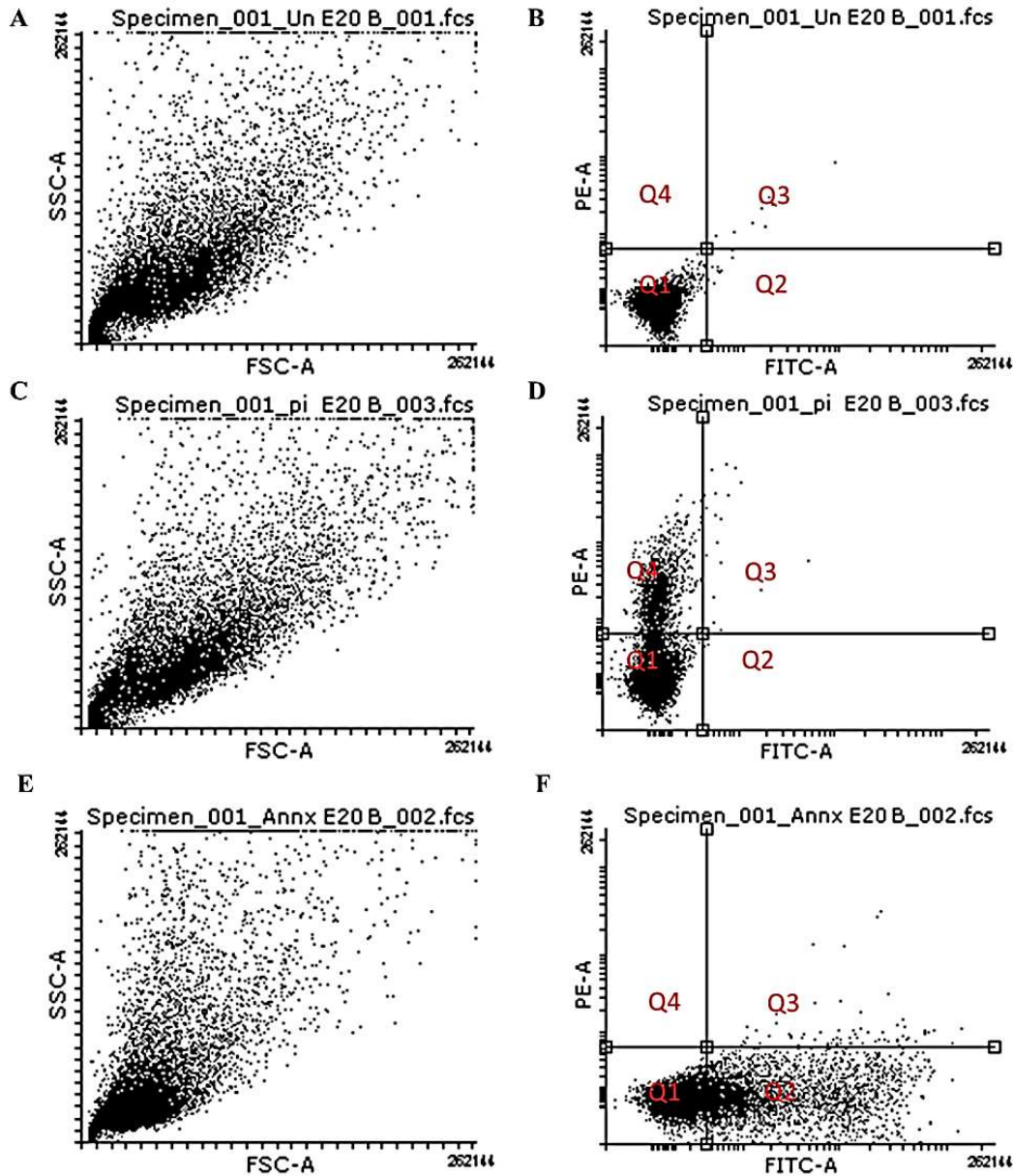
D. Cellular assays for estimation of antiproliferation activity

Figure A46. Flow cytometric analysis of apoptosis of U87MG cells. A) Cell scatter plot indicating side scatter (SSC) parameter versus forward scatter (FSC) parameter and B) propidium iodide (PI) fluorescence versus Annexin V-FITC fluorescence dot plot for unstained U87MG cells showing live (Q1), early apoptotic (Q2), late apoptotic (Q3) and necrotic (Q4) cell populations. C) Cell scatter plot indicating side scatter (SSC) parameter versus forward scatter (FSC) parameter and D) propidium iodide (PI) fluorescence versus Annexin V-FITC fluorescence dot plot for U87MG cells treated with only PI. E) Cell scatter plot indicating side scatter (SSC) parameter versus forward scatter (FSC) parameter and F) propidium iodide (PI) fluorescence versus Annexin V-FITC fluorescence dot plot for U87MG cells treated with only Annexin-FITC.

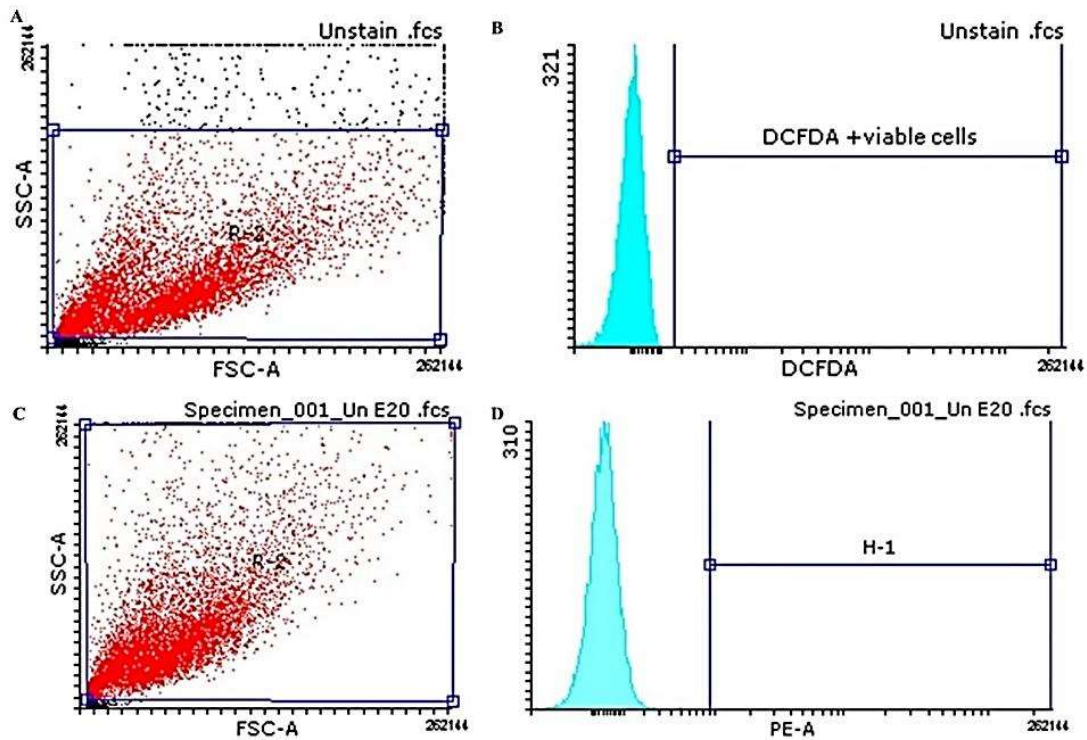


Figure A47. Flow cytometric estimation of reactive oxygen species (ROS) and mitochondrial membrane potential (MMP) of U87MG cells. A) Cell scatter plot indicating side scatter (SSC) parameter versus forward scatter (FSC) parameter and B) histogram of unstained U87MG cells showing the cell counts in Y-axis versus the H₂DCFDA intensity in X-axis. C) Cell scatter plot indicating side scatter (SSC) parameter versus forward scatter (FSC) parameter and D) histogram of unstained U87MG cells showing the cell counts in Y-axis versus the J-aggregate intensity in X-axis.

E. RP-HPLC Method Development and Validation for compound STT13

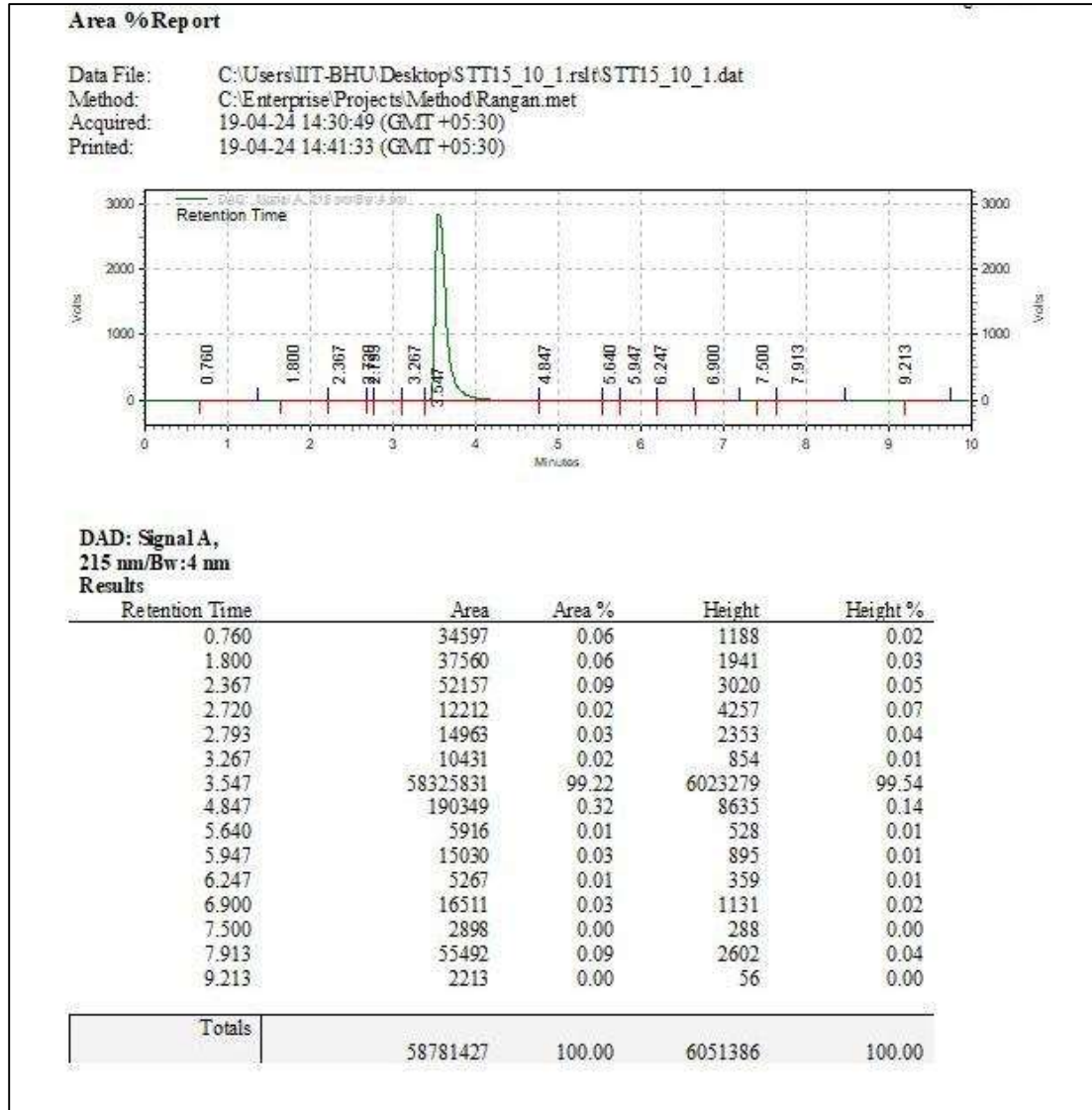


Figure A48. HPLC chromatogram of compound STT13 at a concentration of 10 µM

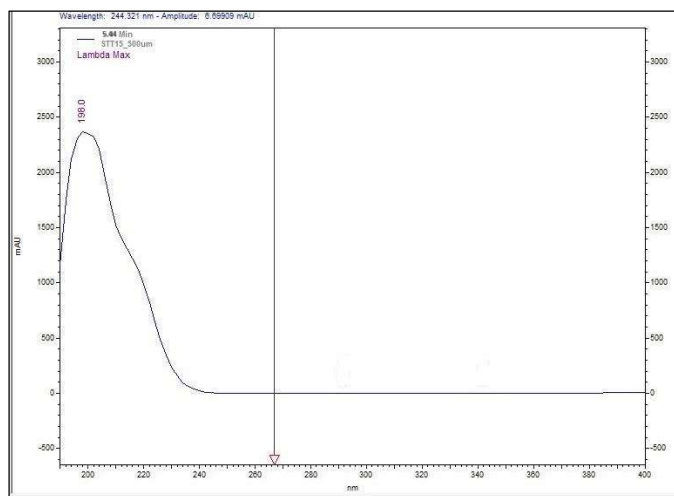


Figure A49. Spectrum of compound **STT13** in 100% acetonitrile at 5.44 min

Table A2. System Parameters for RP-HPLC Method of compound **STT13**

System Parameters	STT13	Formulae
Average % Purity (AUC upto LOD)	98.54 ± 1.63	
Retention time (t_R) [min]	5.440	
Dead time (t_0) [DMSO marker] [min]	0.960	
Adjusted retention time (t) [min]	4.48	$t = t_R - t_0$
Retention factor (k)	4.666	$k = (t_R - t_0) / t_0$
Volumetric flow rate (F_R) [mL/min]	0.7	$V = F_R \times t_R$
Retention volume (V) [mL]	3.808	$S = W_{0.05H} / 2f$
Peak width (W) [min]	0.50	$N = 16(t_R / W)^2$
Tailing (or, Symmetry) factor (S)	0.98	$H = L / N$
Theoretical plates (N)	2002	
HETP (H) [mm]	8.01	

$W_{0.05H}$ = Peak width at 5% peak height above peak baseline, f = length where vertical plumb line from the peak summit intersects a horizontal line at 5% peak height above peak baseline, LOD: limit of detection. HETP: height equivalent of a theoretical plate.

Table A3. System suitability parameters for RP-HPLC method developed for compound **STT13** (n=6)

System Suitability Parameters	Value obtained	SD	%RSD	Acceptance Criteria
Retention (or, Capacity) factor (k)	4.620	0.091	1.954	$k \geq 2.0$
Tailing (or, Symmetry) factor (S)	0.98	0.002	0.178	$S \leq 2.0$
Theoretical plates (N)	2002	19.112	0.954	$N \geq 2000$
HETP (H) [mm]	8.01	--	--	--
Retention time (t_R) [min]	5.44	0.043	0.792	$RSD \leq 1.0\%$
Peak purity	1	0.000	0.000	$RSD \leq 1.0\%$

LIST OF PUBLICATIONS

1. **Mitra, R.**, and Ayyannan, S. R. (2021). Small-Molecule Inhibitors of Shp2 Phosphatase as Potential Chemotherapeutic Agents for Glioblastoma: A Minireview. *ChemMedChem*, 16(5), 777-787. doi: 10.1002/cmdc.202000706.
2. **Mitra, R.**, and Ayyannan, S. R. (2022). Role of Lysine-specific Demethylase 1 and Its Small Molecule Inhibitors in Glioblastoma Multiforme Therapy. *Anticancer Agents Med .Chem.*, 22(18), 3062-3085. doi: 10.2174/1871520622666220421092414.
3. **Mitra, R.**, Singh, D., and Ayyannan, S. R. (2022). Design, synthesis and evaluation of 5-substituted-2-amino-1,3,4-thiadiazole derivatives as anticancer agents. *J. Pharm. Chem.*, 8(Supplement). 150. Retrieved from <https://pubs.vensel.org/index.php/jphchem/article/view/247>.
4. **Mitra, R.**, Kumar, S., and Ayyannan, S. R. (2023). Identification of new small molecule allosteric SHP2 inhibitor through pharmacophore-based virtual screening, molecular docking, molecular dynamics simulation studies, synthesis and in vitro evaluation. *J. Biomol. Struct. Dyn.*, 1-20. doi: 10.1080/07391102.2023.2291733.
5. **Mitra, R.**, Kumar, S., Chaudhuri, A., Agrawal, A. K., and Ayyannan, S. R. (2024). Design, Synthesis and Evaluation of Thioacetamide-Tethered Thiadiazole-1,2,4-Triazole Hybrids as SHP2 Inhibitors for Cancer Therapy. *ChemSelect*, 9(27), e202401602. doi: 10.1002/slct.202401602.
6. **Mitra, R.**, Kumar, S., Chaudhuri, A., Majumdar, S., Agrawal, A. K., Sairam, K., Ayyannan, S. R. (2024). *S*-Acetohydrazones of 5-Methyl-1,3,4-thiadiazole-2-thiol: A New Class of SHP2 Inhibitory Scaffold Possessing Potential Anti-Breast Cancer Activity. (*In communication*)
7. **Mitra, R.**, Kumar, S., Chaudhuri, A., Agrawal, A. K., and Ayyannan, S. R. A HETEROCYCLIC THIOACETAMIDE DERIVATIVE AND A METHOD OF SYNTHESIS THEREOF. (2024). Indian Patent Application No. 202411046345.

