

Supplementary Materials

Table S1. Data collection and refinement statistics of the ternary complexes of *Pf*DHFR-TS V1/S.

| | FB8 | FB6 |
|-------------------------------------|---|---|
| PDB ID code | 8YQ8 | 8YQ9 |
| <i>Data collection</i> | | |
| Wavelength (Å) | 1.5418 | 1.5418 |
| Space group | <i>P</i> 2 ₁ 2 ₁ 2 ₁ | <i>P</i> 2 ₁ 2 ₁ 2 ₁ |
| Molecules/ASU | 2 | 2 |
| Unit-Cell Parameters | | |
| <i>a b c</i> (Å) | 57.763 156.41 165.065 | 58.064 156.496 164.865 |
| $\alpha \beta \gamma$ (deg) | 90 90 90 | 90 90 90 |
| Resolution ^a (Å) | 26.1–2.1 (2.2–2.1) | 24.55–2.4 (2.5–2.4) |
| Completeness (%) | 99.7 (100) | 99.5 (100.0) |
| Redundancy | 5.9 (4.9) | 5.9 (5.6) |
| $\langle I/\sigma(I) \rangle$ | 13.5 (2.4) | 9.1 (2.0) |
| CC(1/2) | 0.998 (0.915) | 0.995 (0.873) |
| Wilson B-factor | 26.8 | 30.84 |
| R_{merge}^b | 0.077 (0.479) | 0.127 (0.483) |
| <i>Refinement</i> | | |
| $R_{\text{work}}/R_{\text{free}}^c$ | 0.1945 / 0.2404 | 0.217 / 0.2909 |
| Average B-factors (Å ²) | | |
| Chains A, B | | |
| Protein | 32.2, 38.9 | 37.8, 47.8 |
| Drg | 36.5, 66.9 | 33.3, 63.5 |
| NDP | 28.5, 64.0 | 38.0, 83.5 |
| UMP | 35.0, 40.1 | 73.4, 83.6 |
| GOL | - | 43.5, 40.5 |
| Water | 35.5 | 33.5 |
| R.m.s. deviation | | |
| Bond lengths (Å) | 0.0092 | 0.0087 |
| Bond angles (°) | 1.8340 | 1.8186 |
| Ramachadran Plot | | |
| favored (%) | 97.6 | 96.4 |
| allowed (%) | 2.4 | 3.6 |
| outlier (%) | - | - |

^aValues in parentheses are for the highest-resolution shell.

^b $R_{\text{merge}} = \frac{\sum_{hkl} \sum_i |I_i(hkl) - \langle I(hkl) \rangle|}{\sum_{hkl} \sum_i I_i(hkl)}$, where $I_i(hkl)$ is the intensity of an individual reflection and $\langle I(hkl) \rangle$ is the mean intensity of symmetry-equivalent reflections.

^c $R_{\text{work}} = \frac{\sum_{hkl} ||F_{\text{obs}}| - |F_{\text{calc}}||}{\sum_{hkl} |F_{\text{obs}}|}$, where F_{obs} and F_{calc} are the observed and calculated structure-factor amplitudes, respectively. R_{free} was calculated in the same manner as R_{work} but using only a 5% unrefined subset of the reflection data.

Table S2. Docking scores obtained for the lead compounds against QM and WT *PfDHFR* and *HsDHFR*.

| Compound | Docking score (kcal/mol) | | |
|----------|--------------------------|------------------|---------------|
| | QM <i>PfDHFR</i> | WT <i>PfDHFR</i> | <i>HsDHFR</i> |
| FB1 | -11.24 | -10.33 | -9.12 |
| FB2 | -10.51 | -11.0 | -7.34 |
| FB3 | -9.37 | -8.88 | -8.69 |
| FB4 | -10.27 | -10.37 | -7.71 |
| FB5 | -9.83 | -9.00 | -5.79 |
| FB6 | -9.913 | -9.88 | -6.53 |
| FB7 | -9.97 | -7.52 | -5.69 |
| FB8 | -9.69 | -11.33 | -7.47 |

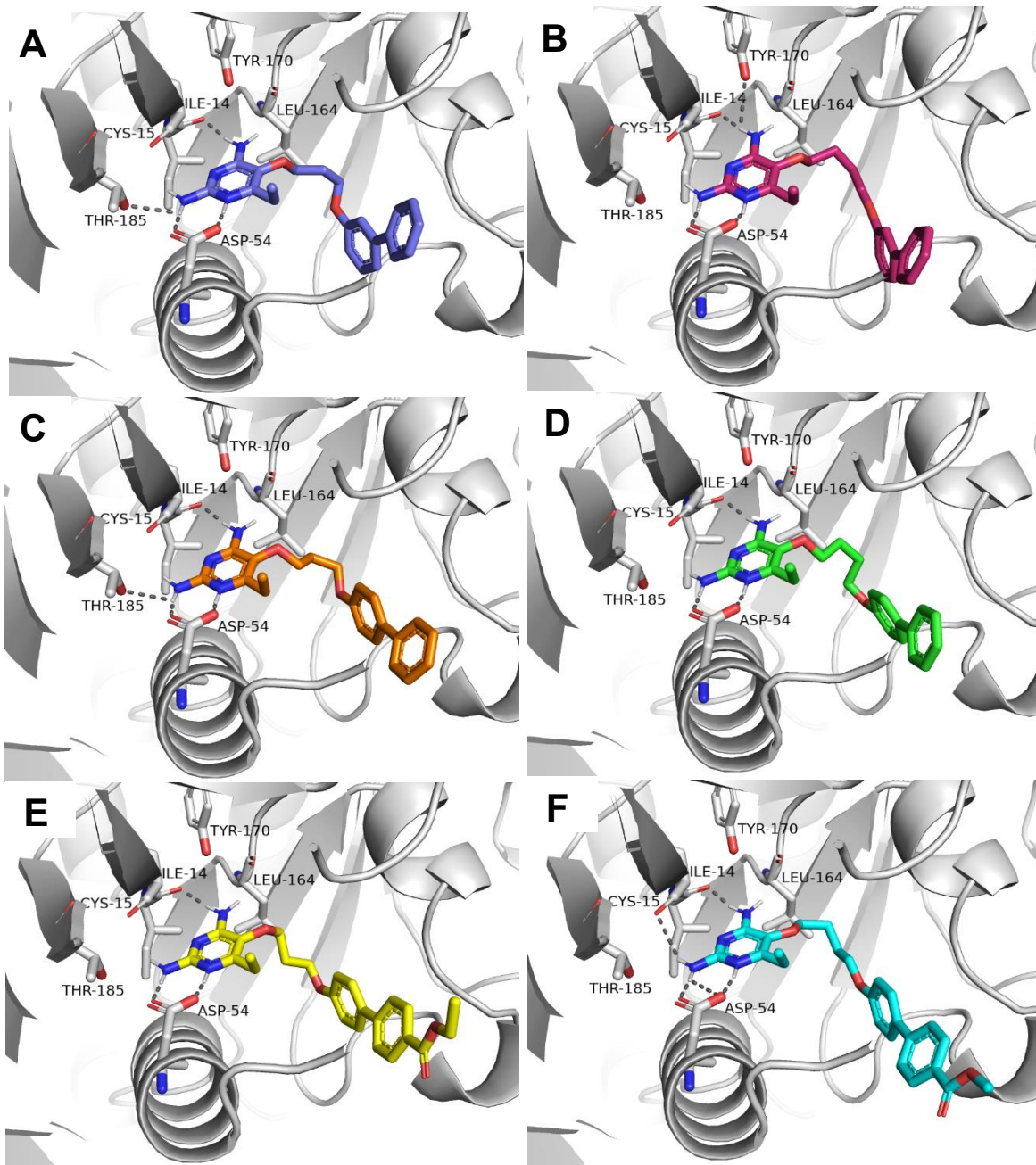


Figure S1. Individual molecular docking poses obtained for lead compounds in QM *PfdHFR*. (A) FB1, (B) FB2, (C) FB3, (D) FB4, (E) FB5, (F) FB7.

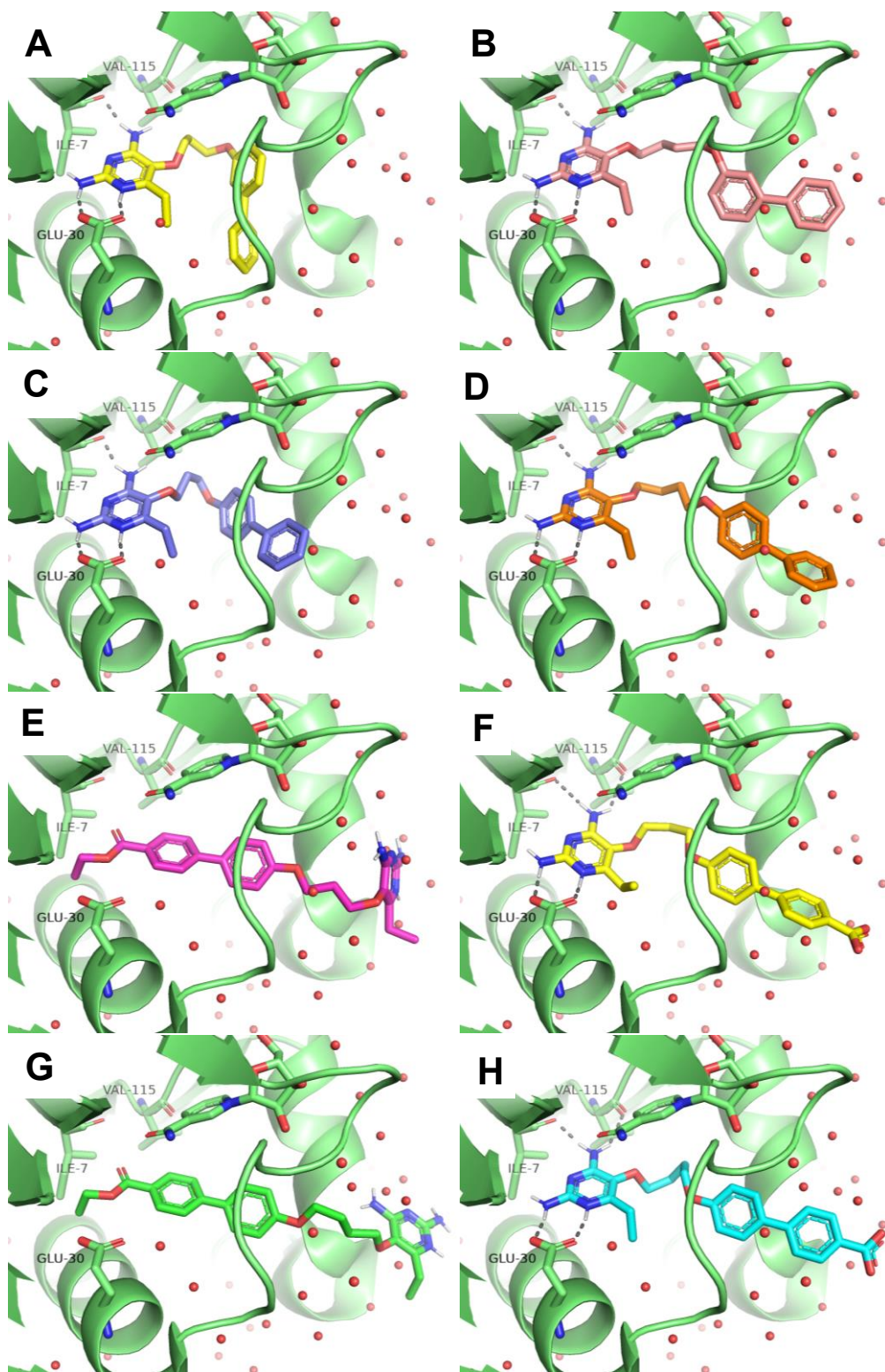


Figure S2. Individual molecular docking poses obtained for lead compounds in *HsDHFR*. (A) FB1, (B) FB2, (C) FB3, (D) FB4, (E) FB5, (F) FB6, (G) FB7, (H) FB8.

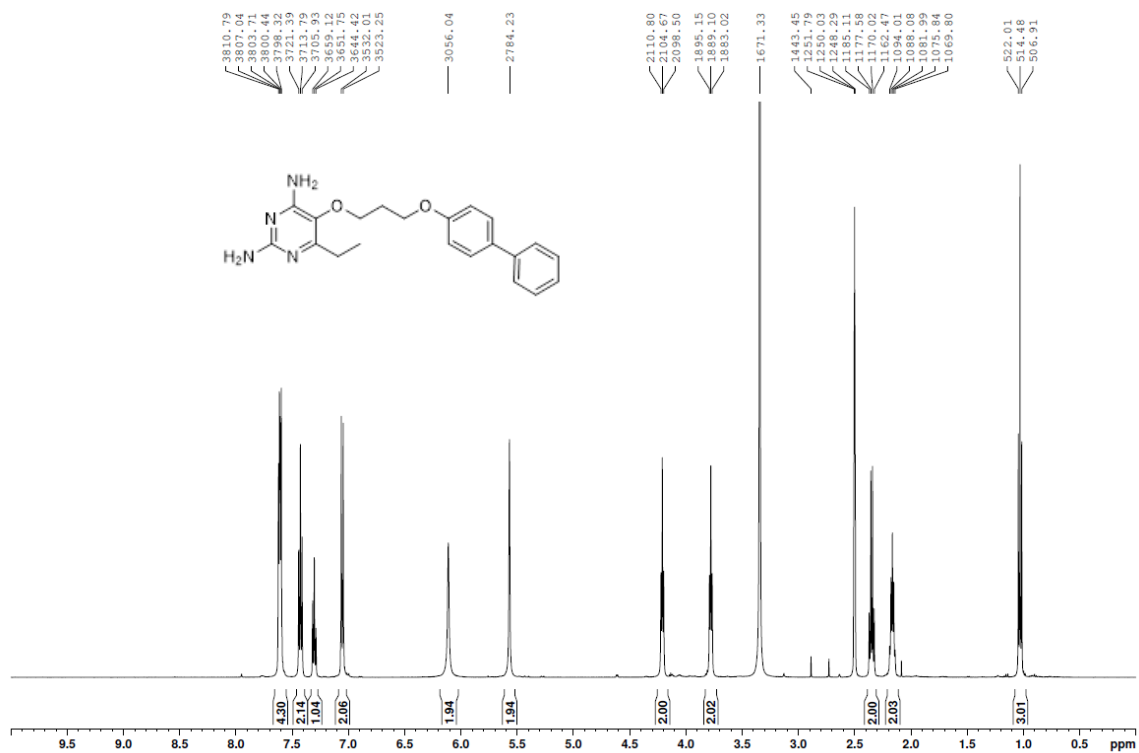


Figure S7. ¹H NMR spectrum of compound 16.

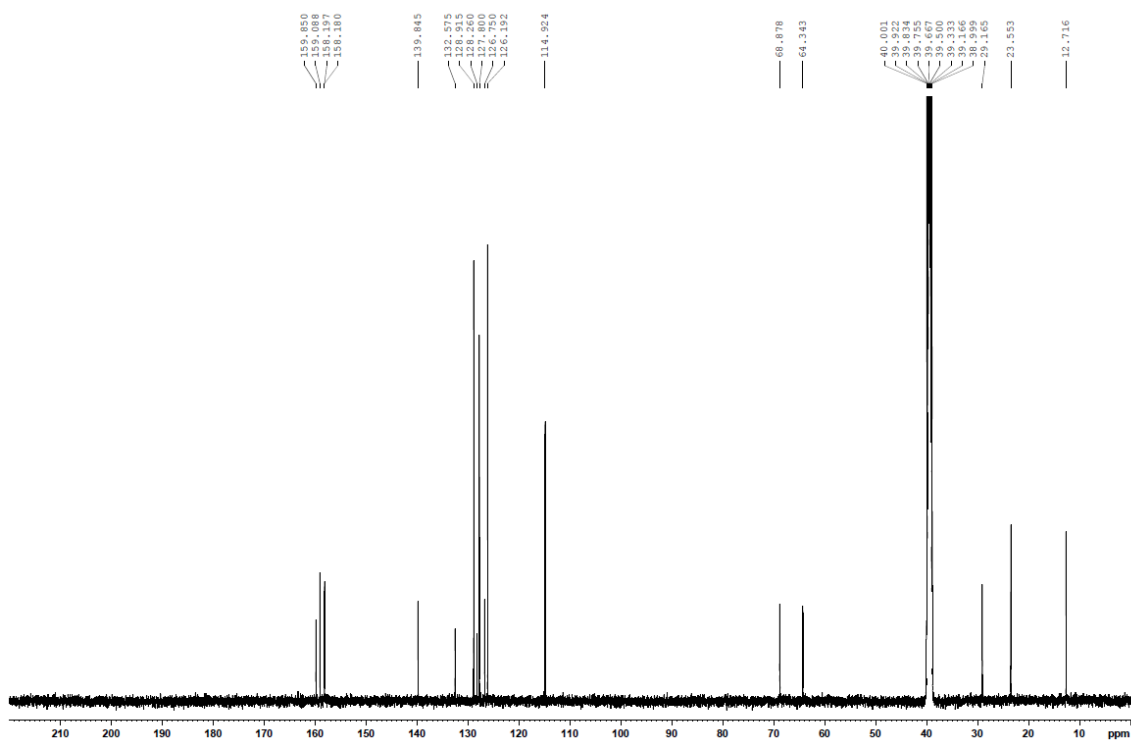


Figure S8. ¹³C NMR spectrum of compound 16.

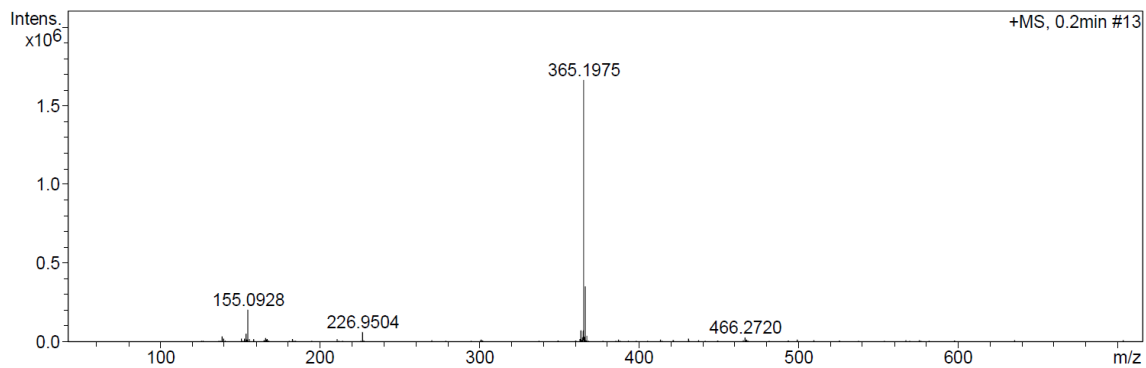


Figure S9. HRMS spectrum of compound 16.

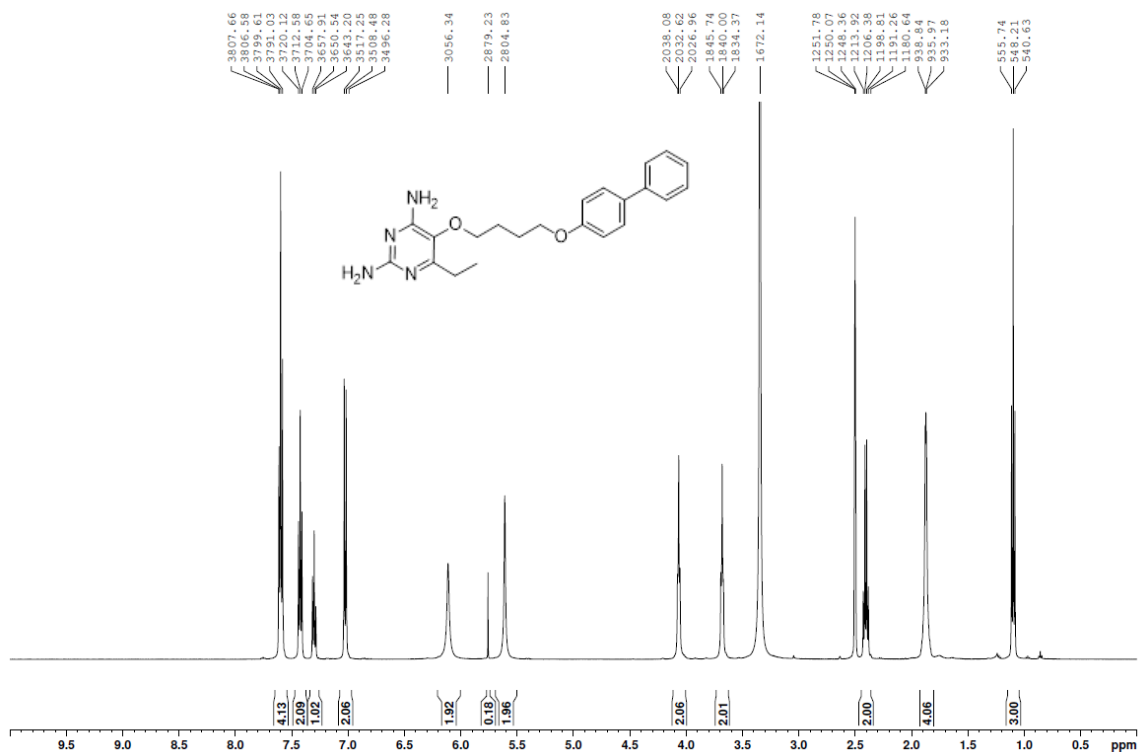


Figure S10. ^1H NMR spectrum of compound 17.

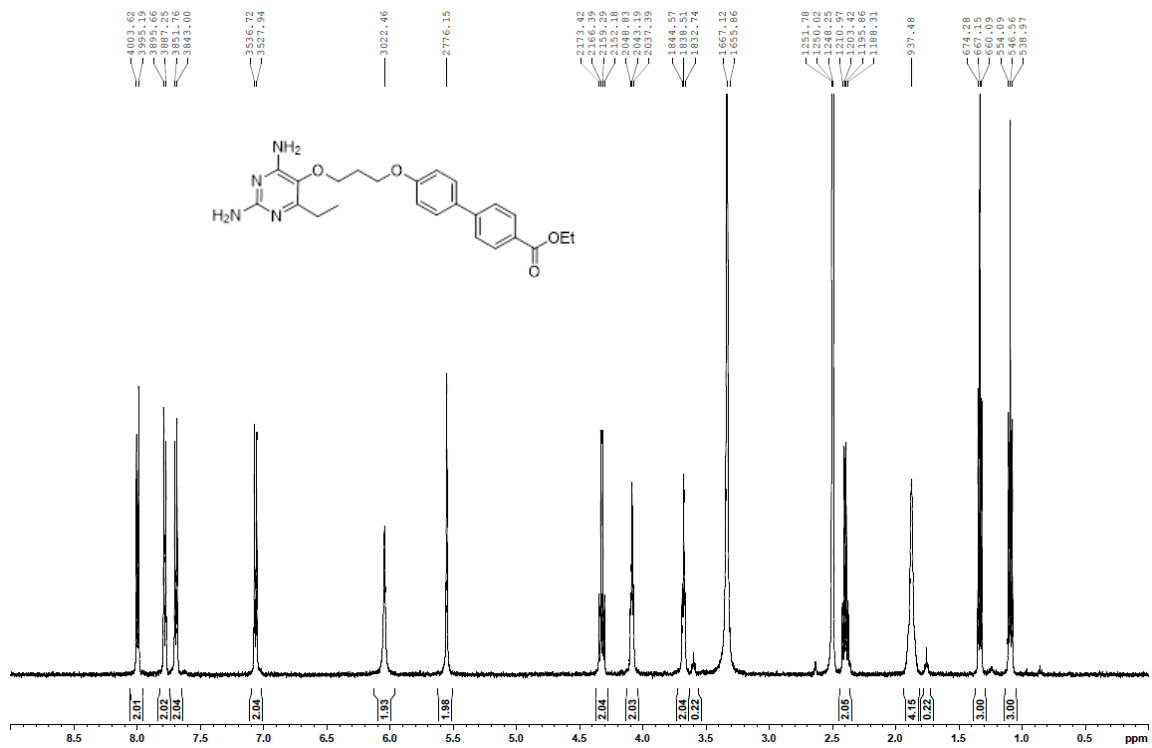


Figure S13. ¹H NMR spectrum of compound 18.

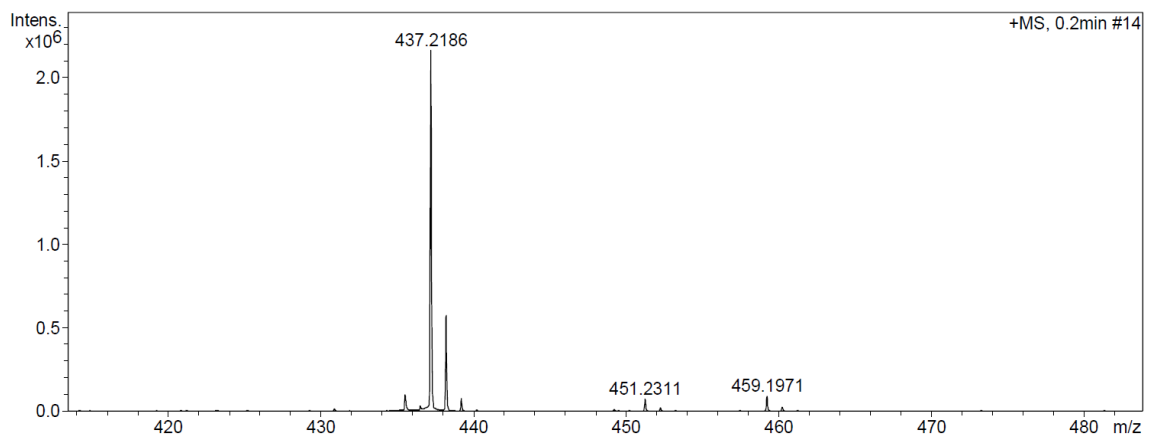


Figure S14. HRMS spectrum of compound 18.

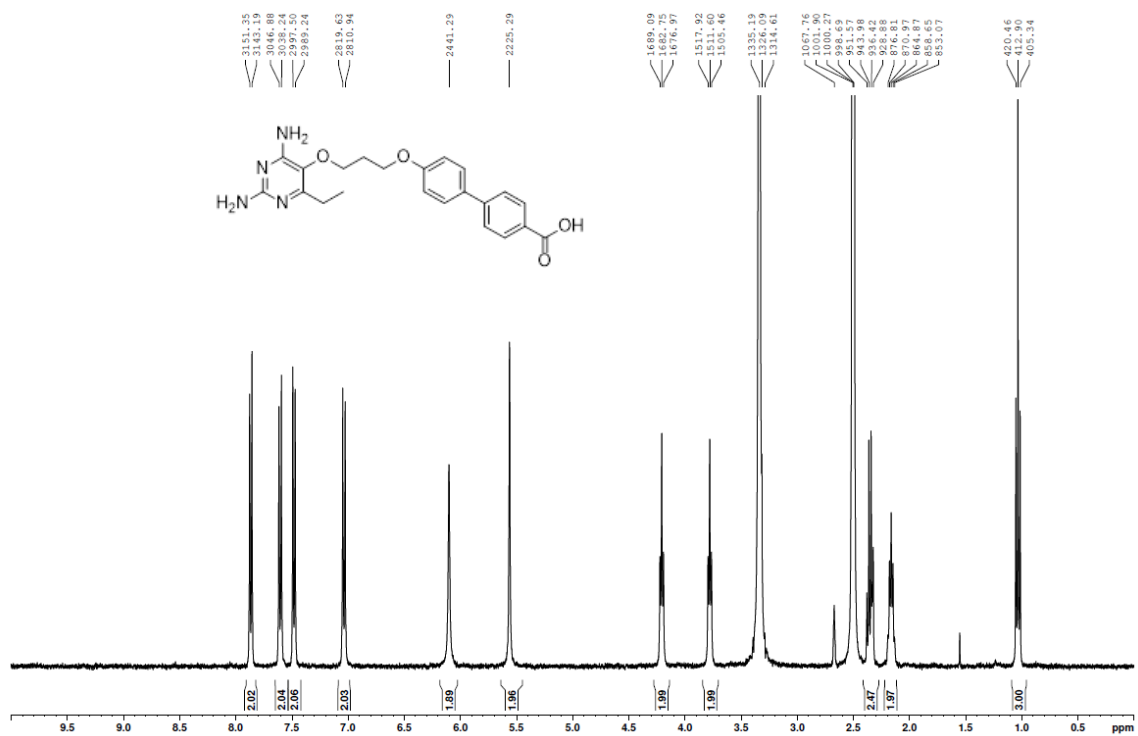


Figure S15. ¹H NMR spectrum of compound 19.

Spectrum View

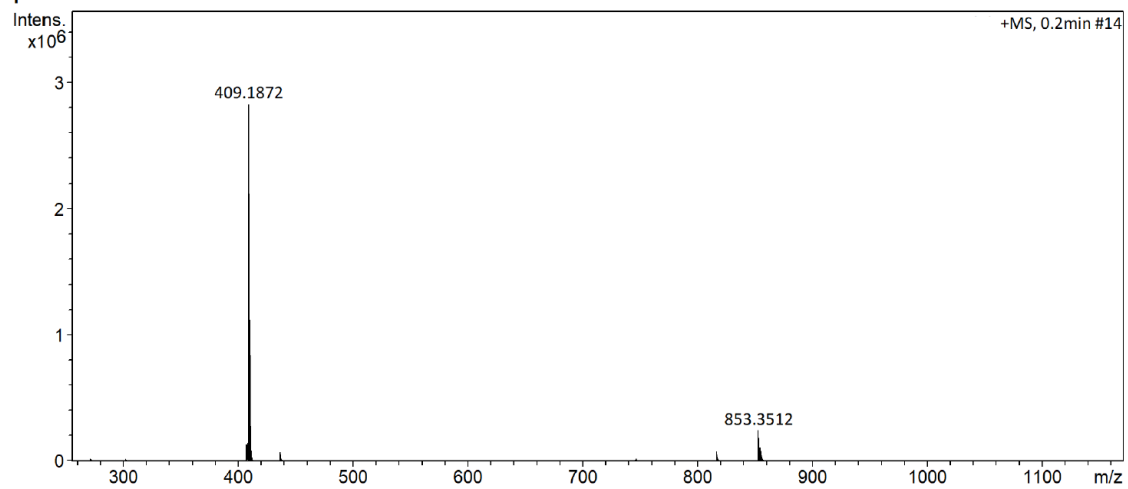


Figure S16. HRMS spectrum of compound 19.

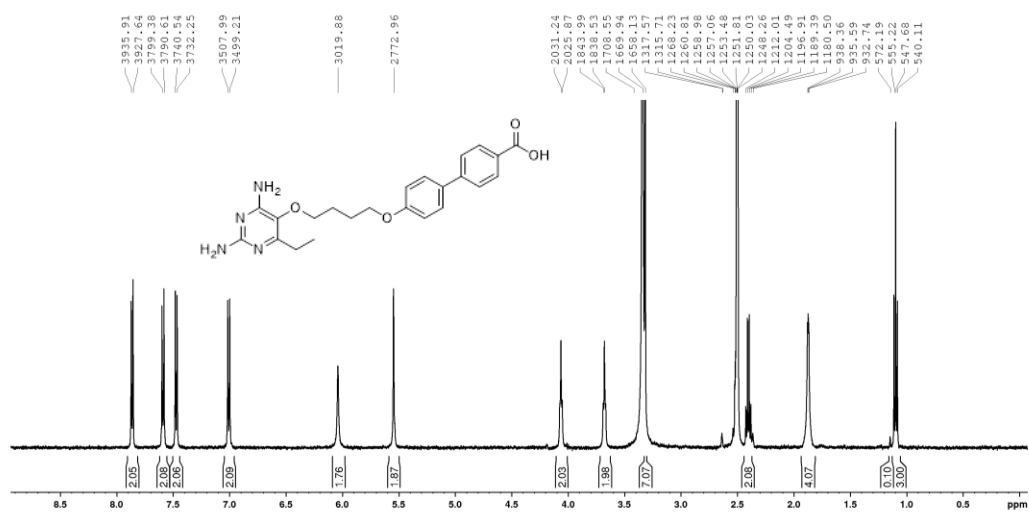


Figure S19. ¹H NMR spectrum of compound 21.

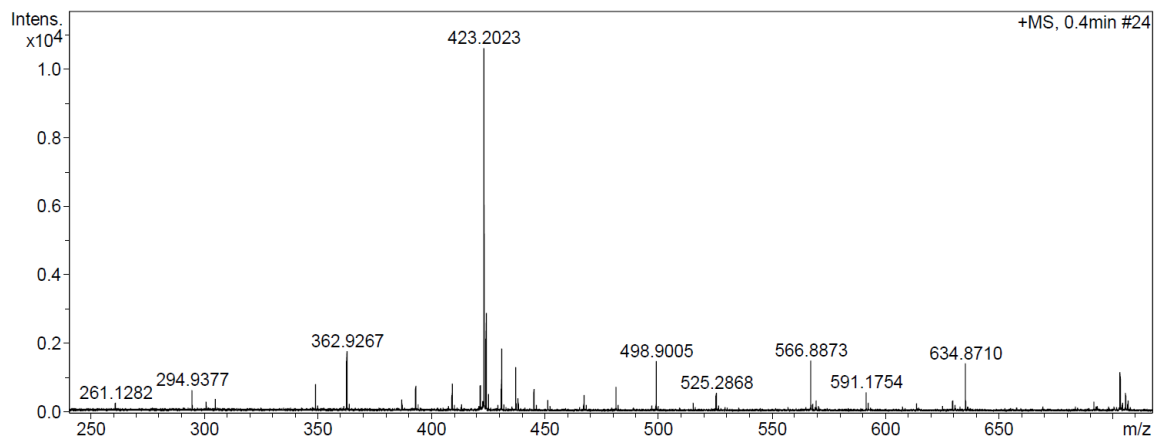


Figure S20. HRMS spectrum of compound 21.

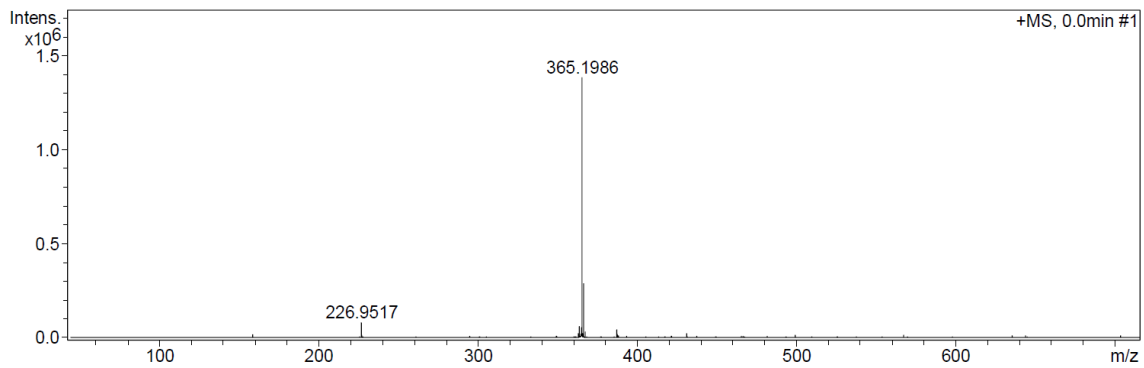


Figure S23. HRMS spectrum of compound FB1.

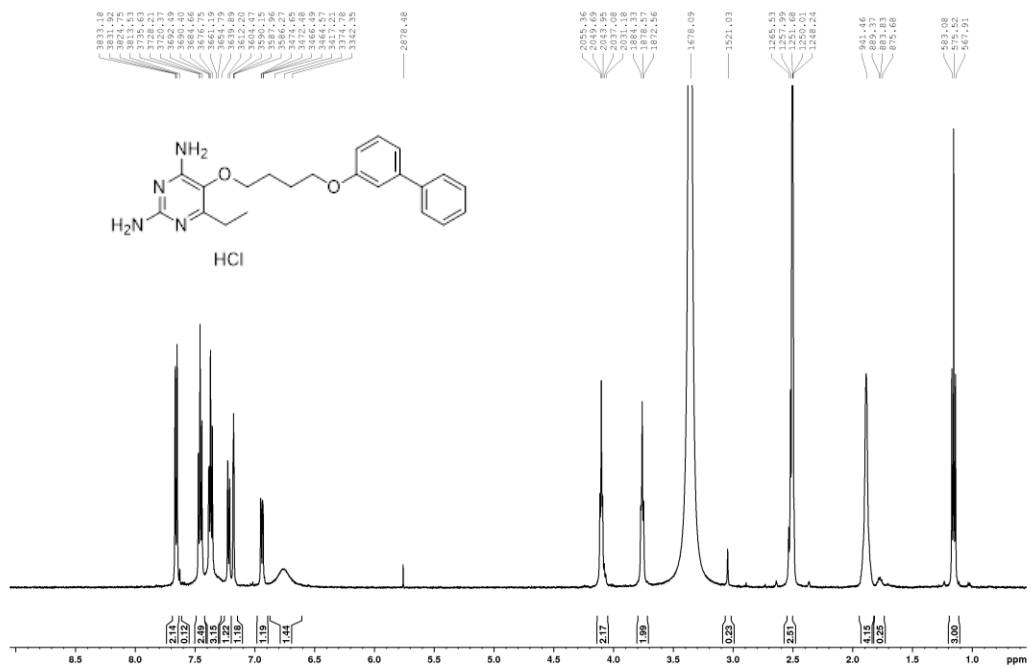


Figure S24. ¹H NMR spectrum of compound FB2.

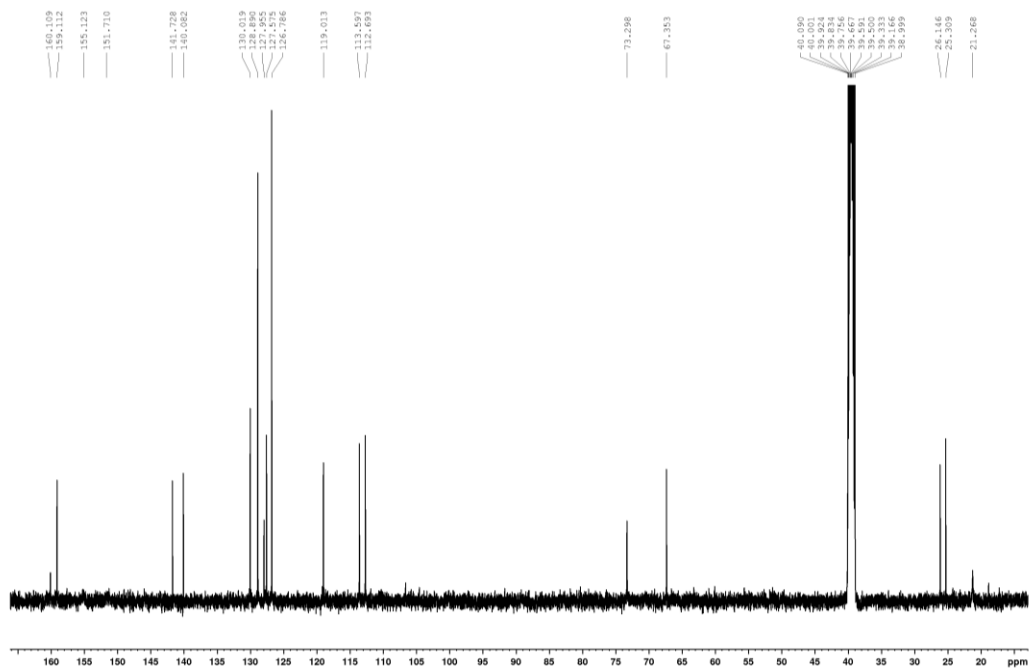


Figure S25. ^{13}C NMR spectrum of compound FB2.

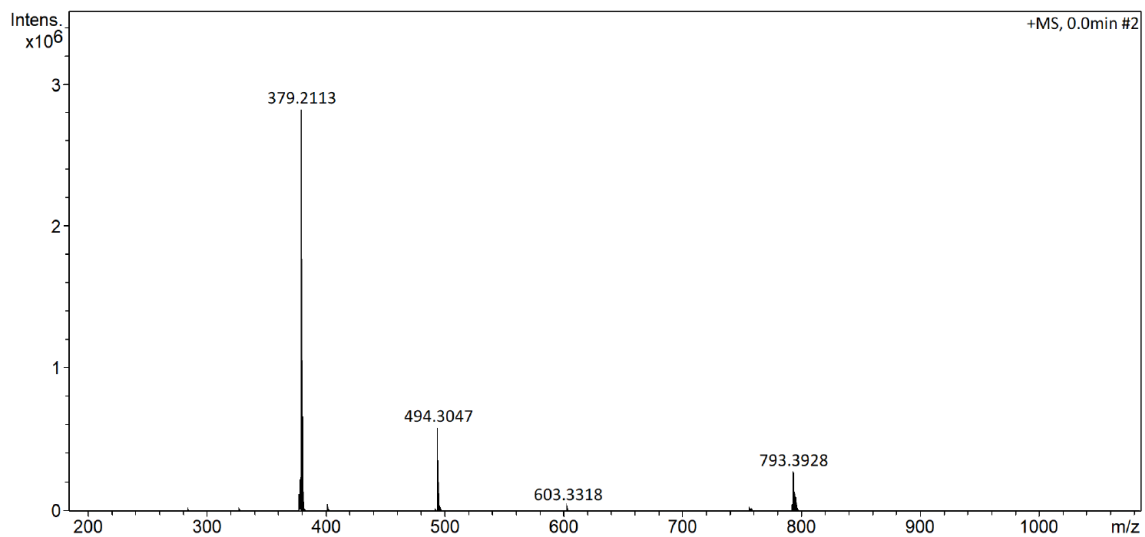


Figure S26. HRMS spectrum of compound FB2.

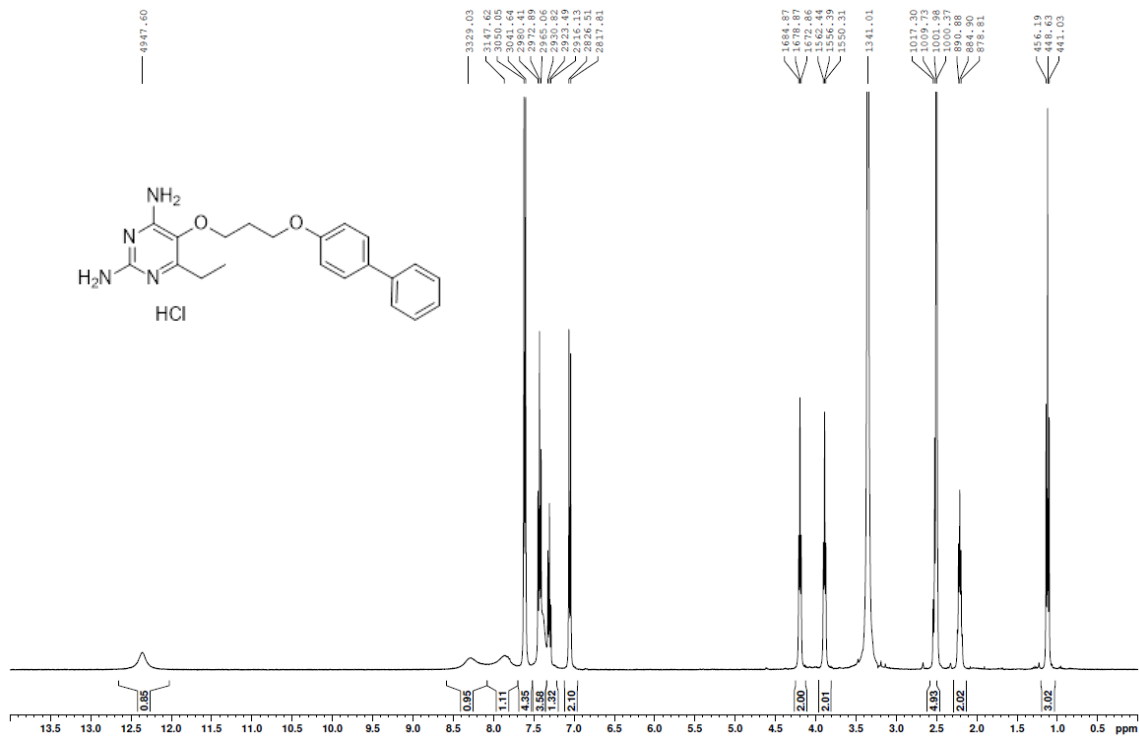


Figure S27. ¹H NMR spectrum of compound FB3.

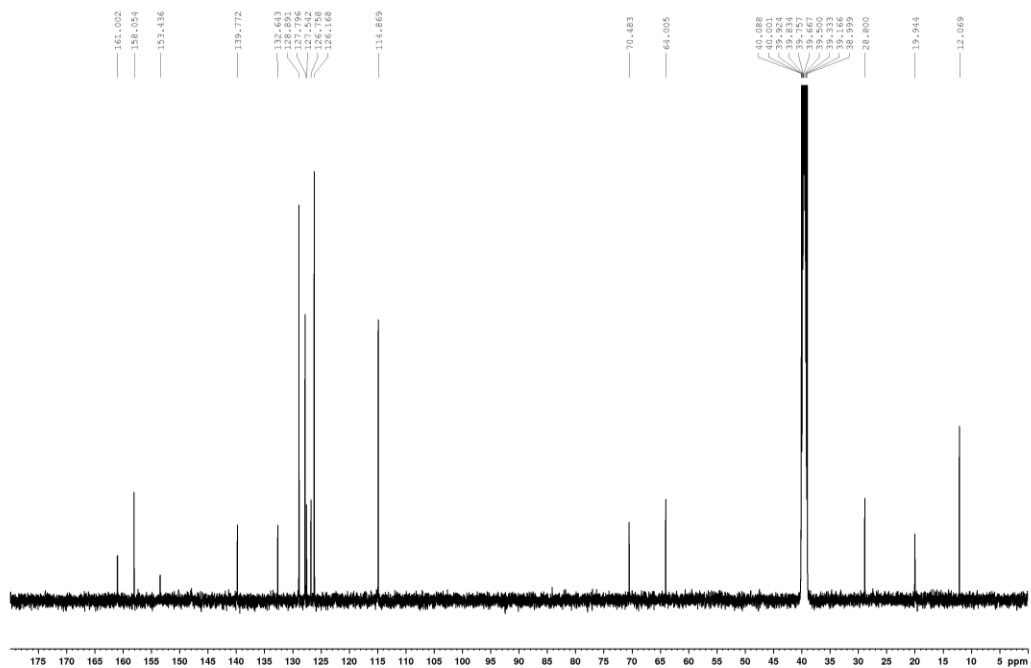


Figure S28. ¹³C NMR spectrum of compound FB3.

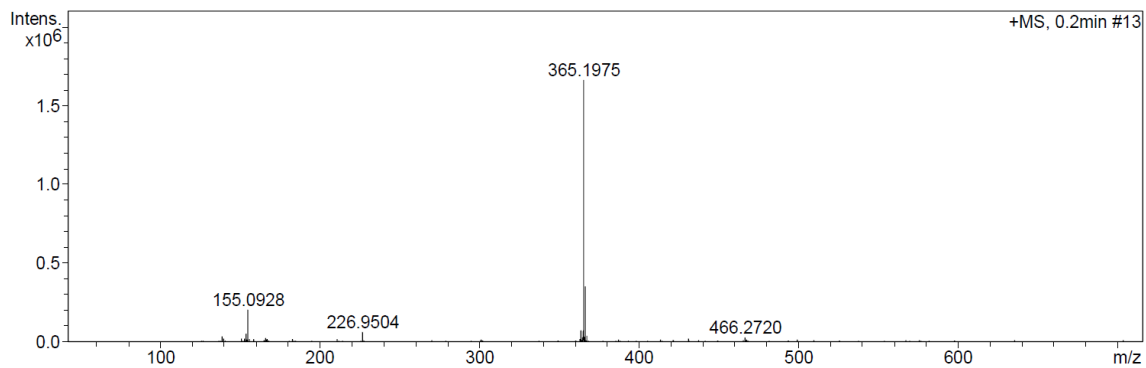


Figure S29. HRMS spectrum of compound FB3.

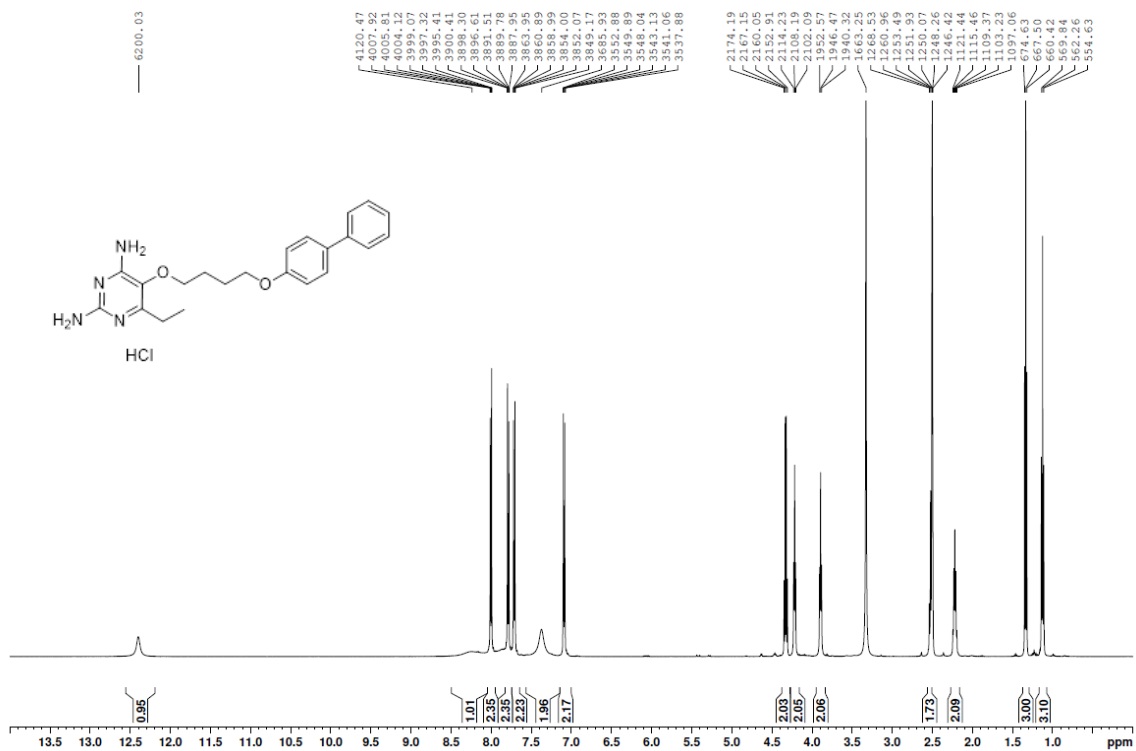


Figure S30. ¹H NMR spectrum of compound FB4.

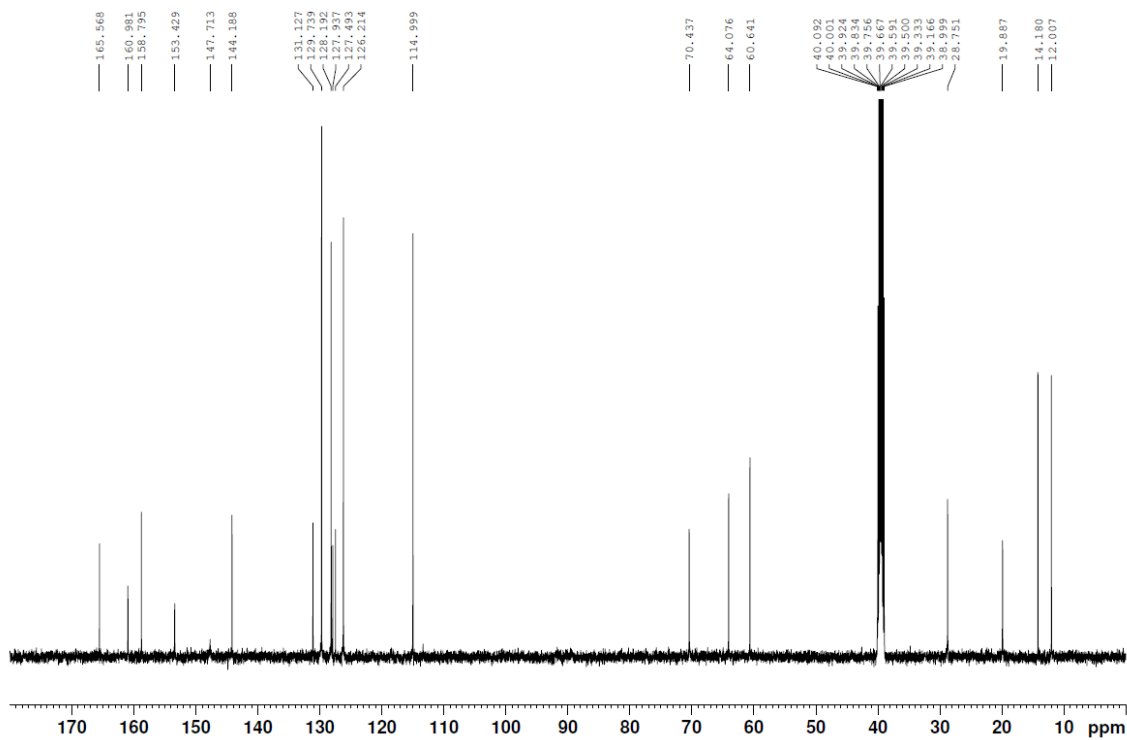


Figure S31. ^{13}C NMR spectrum of compound FB4.

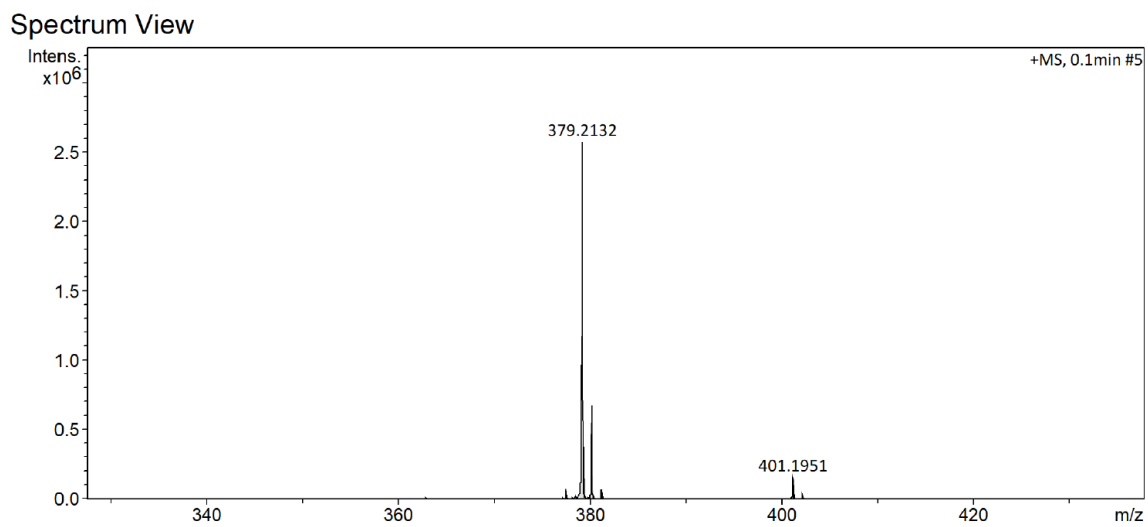


Figure S32. HRMS spectrum of compound FB4.

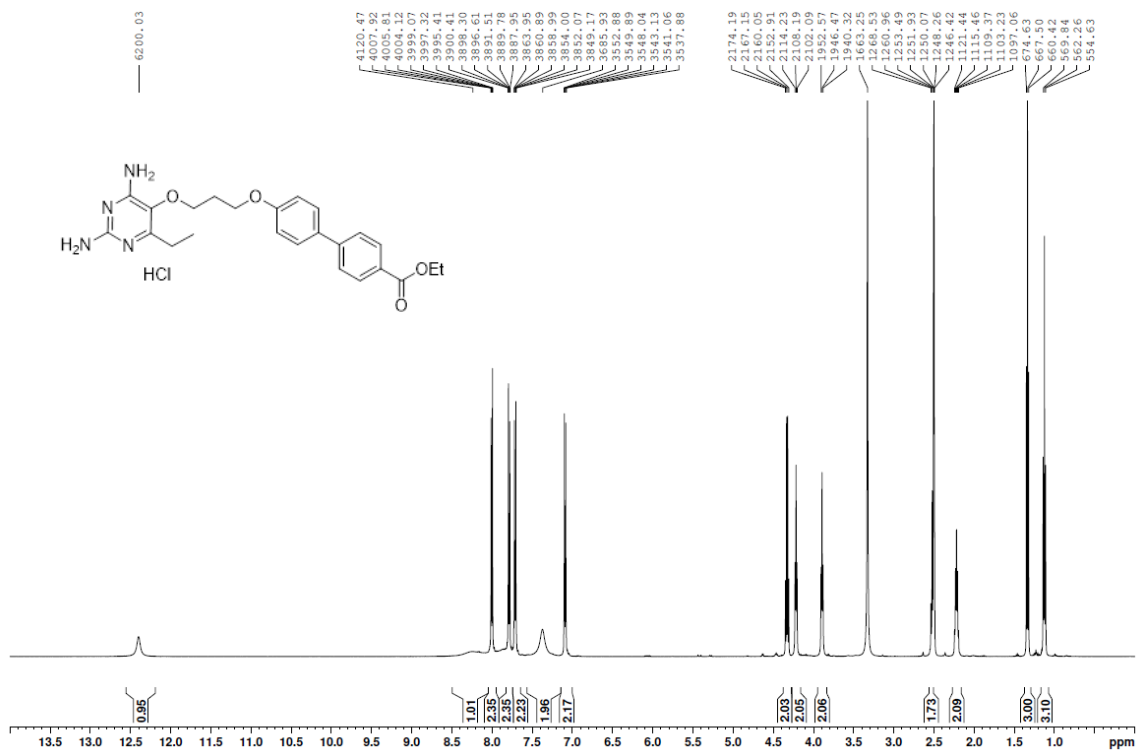


Figure S33. ¹H NMR spectrum of compound FB5.

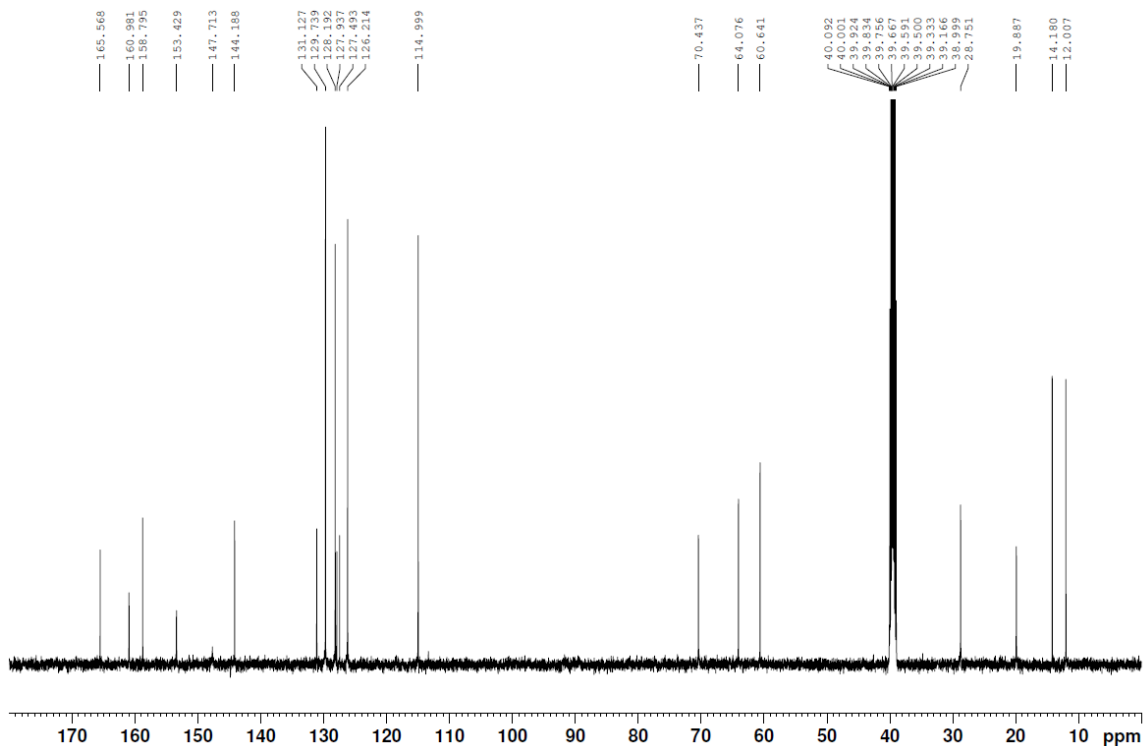


Figure S34. ¹³C NMR spectrum of compound FB5.

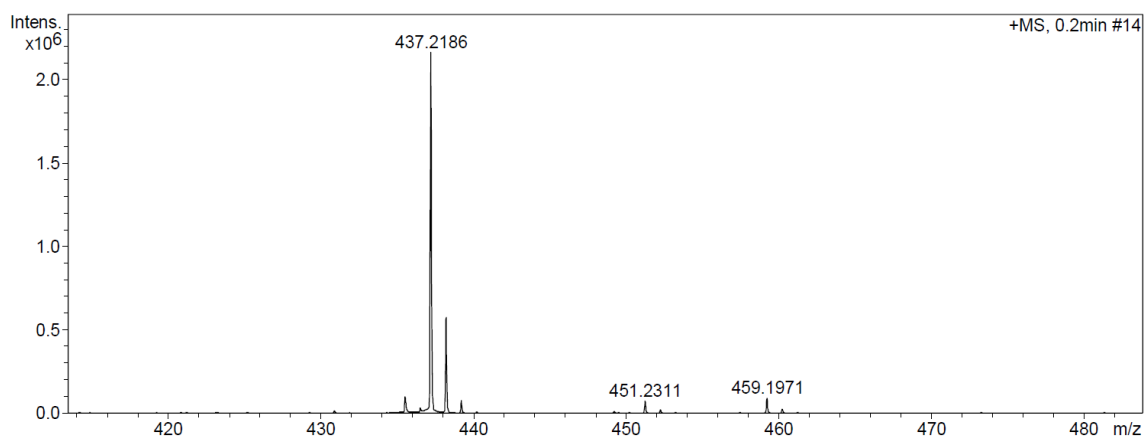


Figure S35. HRMS spectrum of compound FB5.

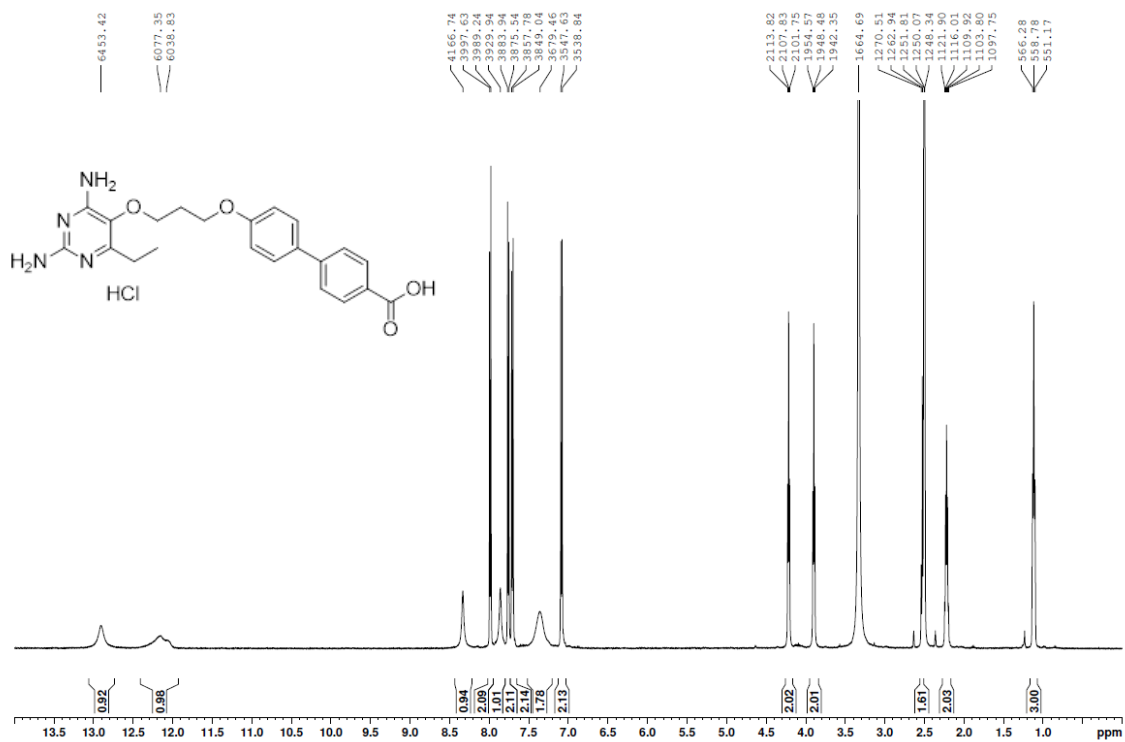


Figure S36. ¹H NMR spectrum of compound FB6.

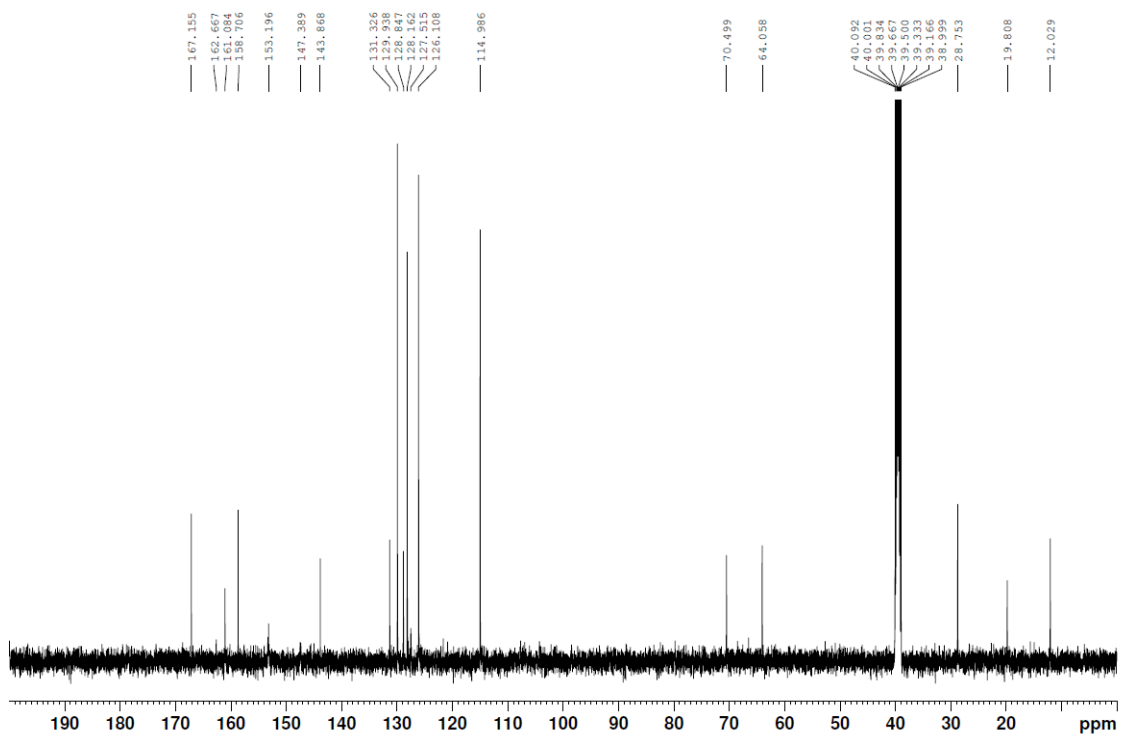


Figure S37. ^{13}C NMR spectrum of compound FB6.

Spectrum View

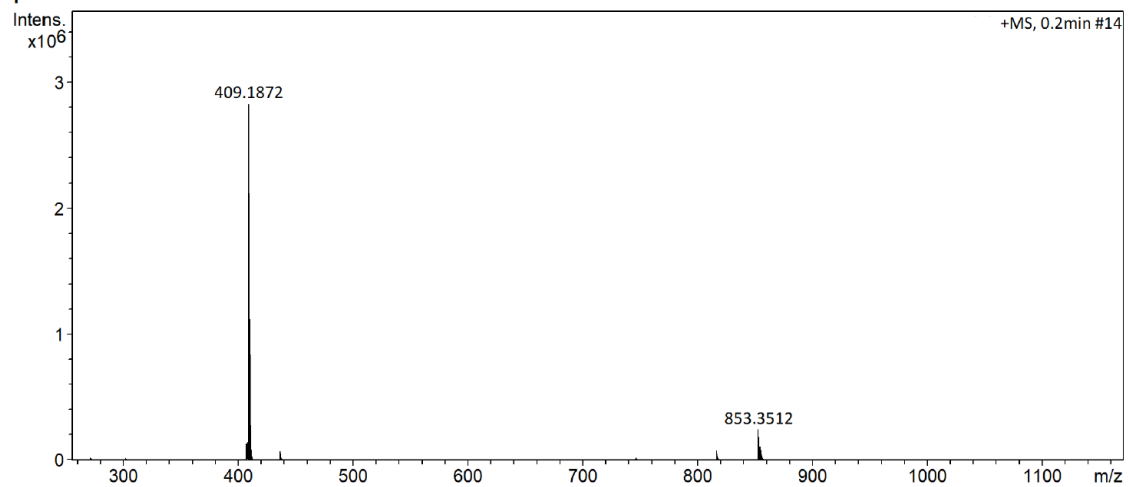


Figure S38. HRMS spectrum of compound FB6.

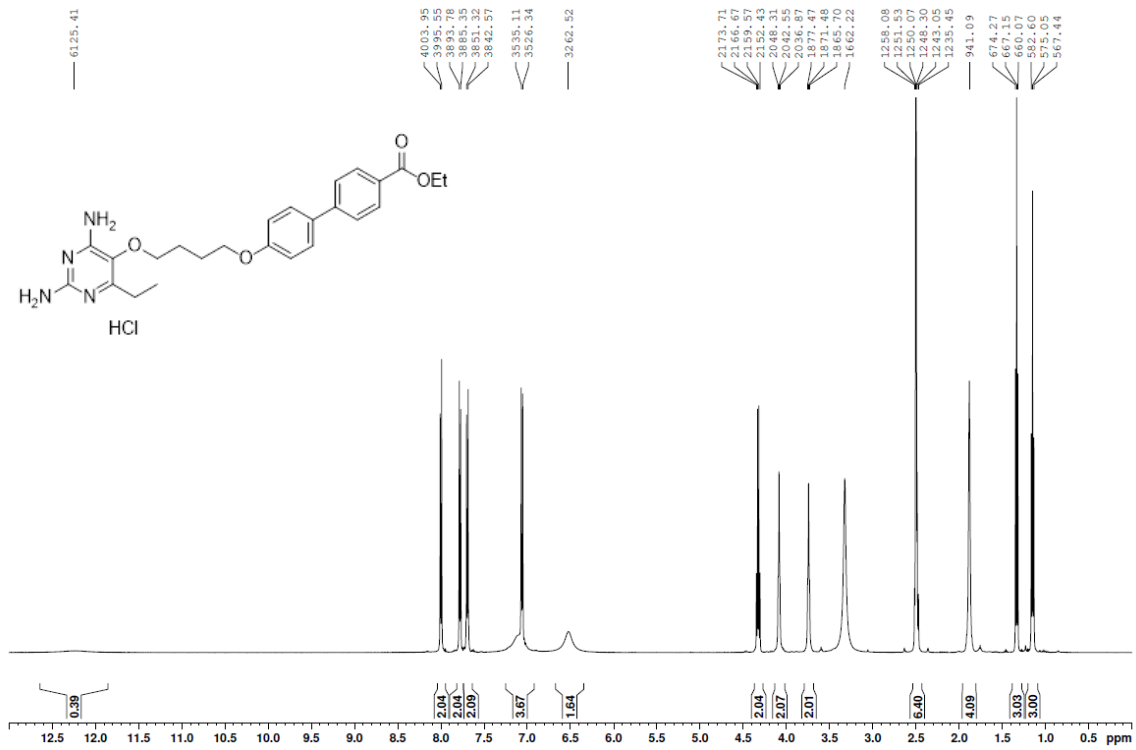


Figure S39. ¹H NMR spectrum of compound FB7.

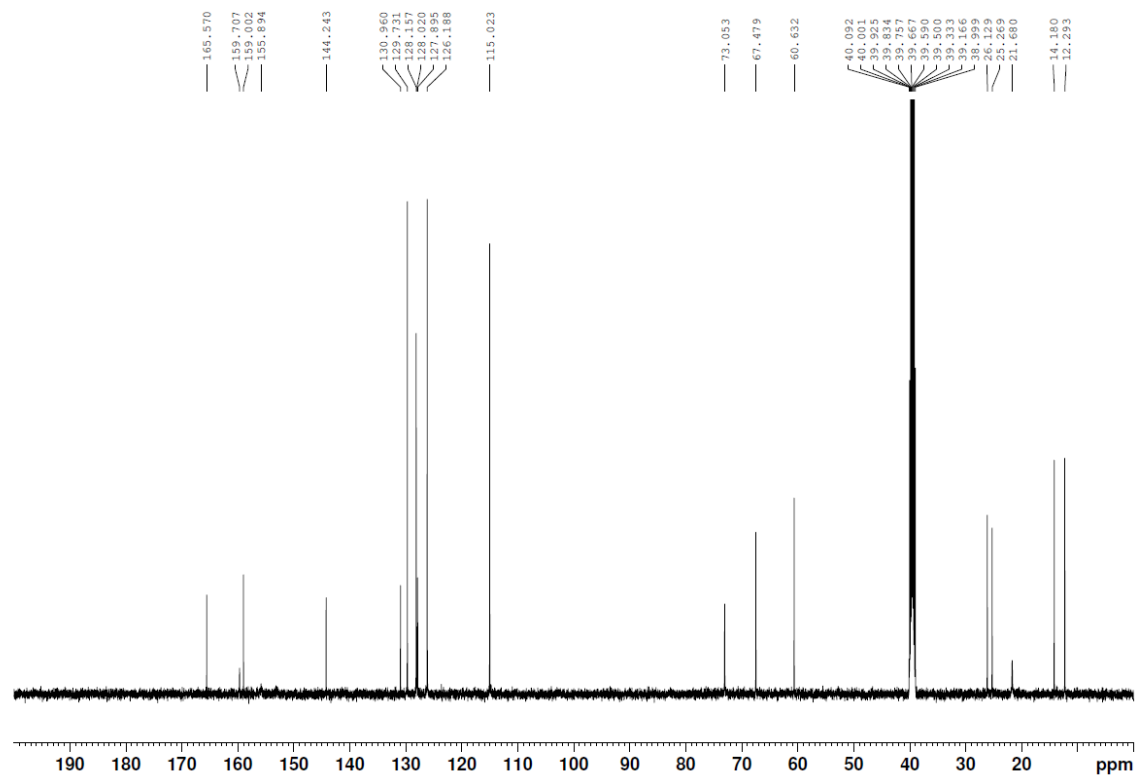


Figure S40. ¹³C NMR spectrum of compound FB7.

Spectrum View

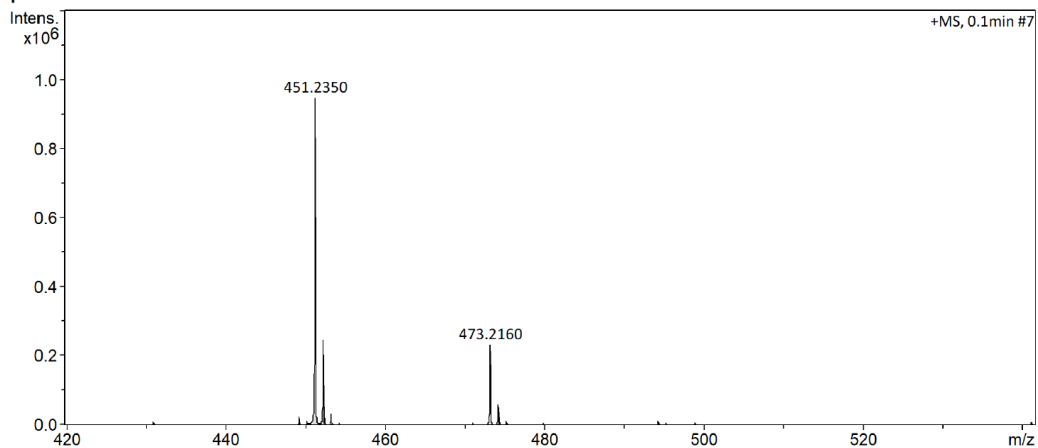


Figure S41. HRMS spectrum of compound FB7.

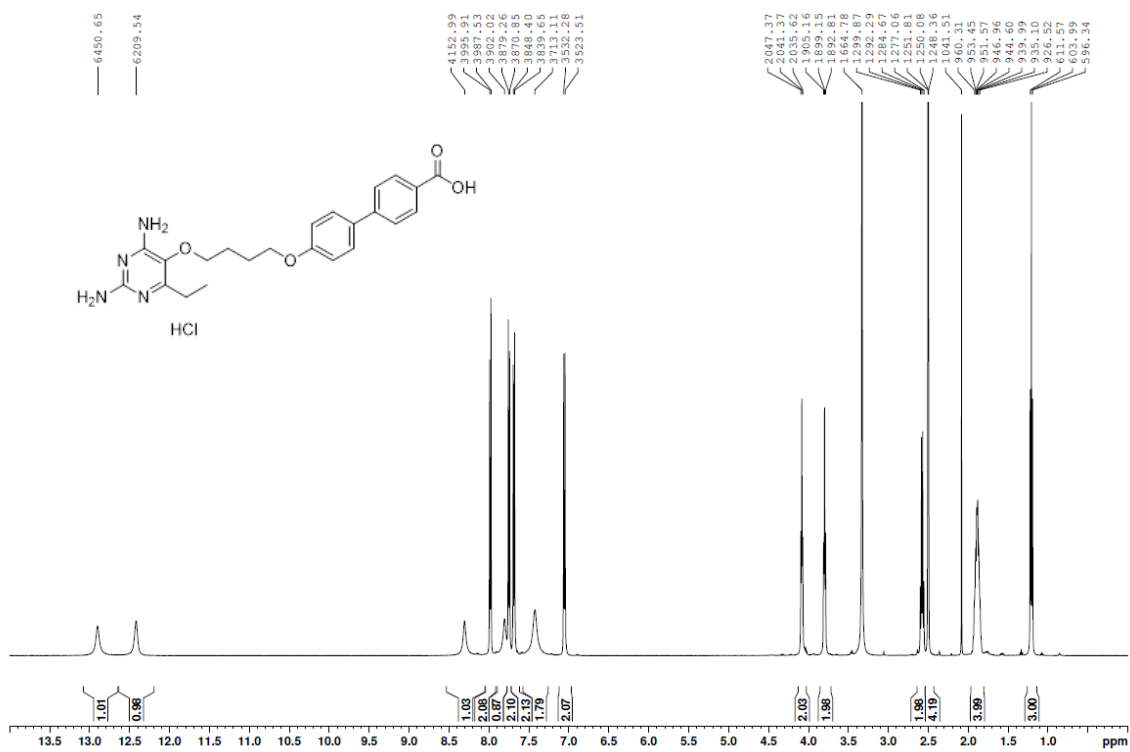


Figure S42. ^1H NMR spectrum of compound FB8.

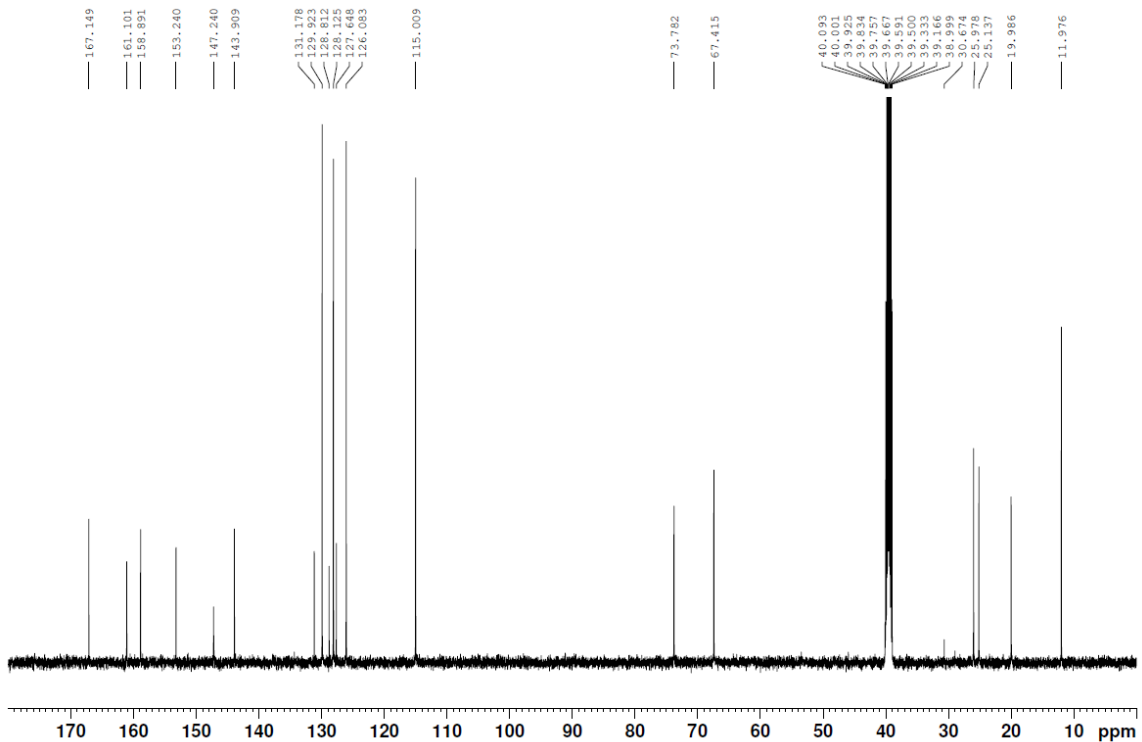


Figure S43. ^{13}C NMR spectrum of compound FB8.

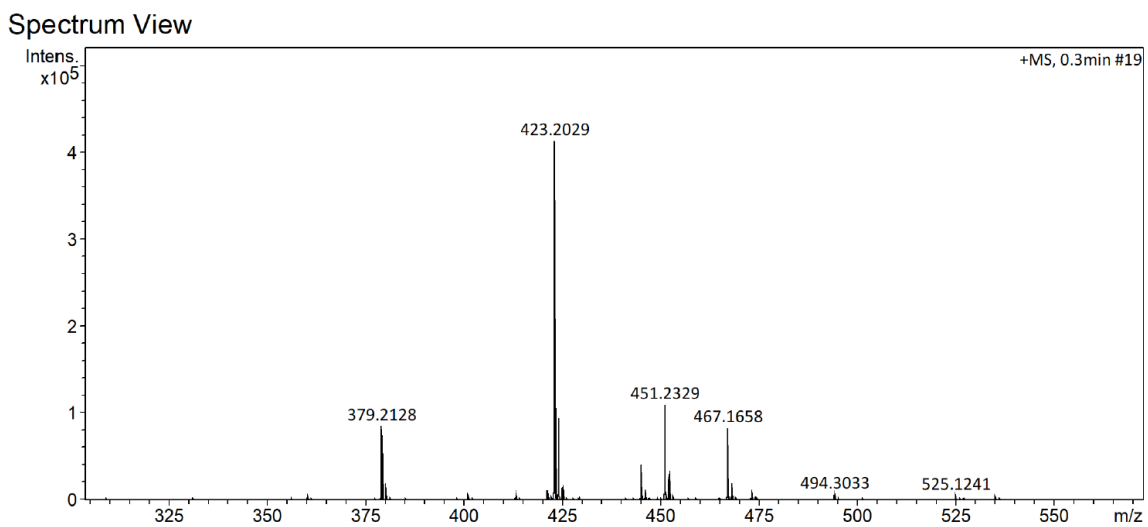


Figure S44. HRMS spectrum of compound FB8.