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## **ELECTRONIC SUPPORTING INFORMATION**

## Identification and Validation of Novel Microtubule Suppressors with an Imidazopyridine Scaffold through Structure-Based Virtual Screening and Docking.

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**Table S1:** Chemical structure of shortlisted compounds and their binding energiesby BUDE, AutoDock, and MOE.

| Cp. | ZINC-ID      | Structure | BUDE<br>binding  | AutoDock<br>binding | MOE<br>Binding     |
|-----|--------------|-----------|------------------|---------------------|--------------------|
|     |              |           | energy<br>kJ/mol | energy<br>Kcal/mol  | Energy<br>kcal/mol |
| 1   | Zinc02843810 |           | -111.013         | -9.79               | -6.28              |
| 2   | Zinc02691641 |           | -92.82           | -10.31              | -7.2               |
| 3   | Zinc03614688 |           | -91.78           | -7.87               | -5.2               |
| 4   | Zinc49543397 |           | -104.3           | -9.14               | -8.01              |
| 5   | Zinc02690781 |           | -100.75          | -10.08              | -8.24              |
| 6   | Zinc36360243 |           | -100.9           | -8.57               | -10.88             |

| 7  | Zinc18200970 | H <sub>2</sub> N 0                      | -125.35 | -9.7   | -6.9  |
|----|--------------|---|---------|--------|-------|
|    |              |   |         |        |       |
|    |              |   |         |        |       |
|    |              | F C C C C C C C C C C C C C C C C C C C | -119.48 | -9.65  | -9.61 |
| 8  | Zinc02690805 |   |         |        |       |
|    |              | F-NH O-                                 |         |        |       |
| 9  | Zinc02690789 |   | -108.21 | -10.02 | -8.93 |
|    |              |   |         |        |       |
|    |              |   |         |        |       |
|    |              |   | -83.65  | -9.79  | -5.5  |
| 10 | Zinc07095120 | NH S                                    |         |        |       |
|    |              |   |         |        |       |
|    |              |   |         |        |       |
|    | 7inc11112052 | NH                                      | -83.66  | -9.9   | -6.32 |
| 11 | Zinc11112053 |   |         |        |       |
|    |              |   |         |        |       |
|    | 7in-22422021 | CI S                                    | -84.55  | -9.9   | -5.7  |
| 12 | Zinc23483881 |   |         |        |       |
|    |              |   |         |        |       |
| 13 | Zinc09373064 |   | 105.95  | -10.91 | -6.36 |
|    |              |   |         |        |       |
|    |              |   |         |        |       |

**Table S2:** Physical properties of shortlisted compounds calculated by MOE.

| Compounds | ZINC ID      | MwSt   | Log p<br>(O/W) | H-bond<br>donors | H-bond<br>acceptors | tPSA   | Rotatable<br>bond |
|-----------|--------------|--------|----------------|------------------|---------------------|--------|-------------------|
| 1         | Zinc02843810 | 437.49 | 2.6            | 1                | 4                   | 87.00  | 7                 |
| 2         | Zinc02691641 | 443.49 | 4.6            |                  | 5                   | 65.07  | 5                 |
| 3         | Zinc03614688 | 368.26 | 4.08           | 1                | 3                   | 47.56  | 3                 |
| 4         | Zinc49543397 | 377.37 | 3.16           | 2                | 3                   | 103.62 | 4                 |
| 5         | Zinc02690781 | 474.48 | 3.79           | 1                | 5                   | 84.94  | 8                 |
| 6         | Zinc36360243 | 403.36 | 3.9            | 2                | 5                   | 79.87  | 4                 |
| 7         | Zinc18200970 | 490.53 | 3.4            | 2                | 5                   | 116.95 | 5                 |
| 8         | Zinc02690805 | 496.44 | 3.8            | 1                | 5                   | 84.94  | 8                 |
| 9         | Zinc02690789 | 490.48 | 3.4            | 1                | 6                   | 94.17  | 9                 |
| 10        | Zinc07095120 | 411.53 | 4.05           | 2                | 4                   | 88.38  | 6                 |
| 11        | Zinc11112053 | 439.53 | 3.7            | 1                | 5                   | 80.23  | 6                 |
| 12        | Zinc23483881 | 351.85 | 3.3            | 1                | 3                   | 47.56  | 6                 |
| 13        | Zinc09373064 | 459.53 | 3.2            | 1                | 6                   | 91.16  | 5                 |

Table S3 The effect of compound 6 on the phases of the MCF7 cell cycle.

| compounds | %G0-G1 | %S    | %G2-M | %Pre G1 |
|-----------|--------|-------|-------|---------|
| Control   |        |       |       |         |
|           | 53.91  | 42.71 | 3.38  | 1.48    |
| 6         | 36.28  | 29.66 | 34.06 | 27.36   |

**Table S4** Hydrogen bonding of compounds **6,8** and **9** docked into the colchicine binding site.

| Ср | Interacting moiety<br>in compound | Amino acid involved | Distance<br>Å | Type of interaction |
|----|-----------------------------------|---------------------|---------------|---------------------|
| 6  | OH-imidazopyridine                | C=O Thra179         | 2.5           | H-bond              |
|    | OH-phenyl                         | NH Alaβ250          | 3.1           | H-bond              |
|    |                                   |                     |               |                     |
| 8  | C=O                               | SH Cysβ241          | 3.1           | H-bond              |
|    | C=O                               | OH Sera178          | 2.9           | H-bond              |
| 9  | OCH₃                              | SH Cysβ241          | 3.6           | H-bond              |
|    | C=O                               | OH Sera178          | 3.0           | H-bond              |

Table S5 Average RMSD values (Å) of complexes of 6, 8, 9, 14, 15 and colchicine with tubulin over 1.5  $\mu$ s of simulation.

| Å                            | 6         | 8         | 9         | 14        | 15        | colchicine |
|------------------------------|-----------|-----------|-----------|-----------|-----------|------------|
| Average RMSD<br>ligand       | 1.4 ± 0.3 | 3.4 ± 0.5 | 2.2 ± 0.8 | 2.2 ± 0.8 | 1.7 ± 0.3 | 1.5 ± 0.4  |
| Average RMSD<br>protein (AB) | 2.5 ± 0.2 | 2.5 ± 0.2 | 2.4 ± 0.2 | 2.6 ± 0.3 | 2.6 ± 0.3 | 2.5 ± 0.2  |

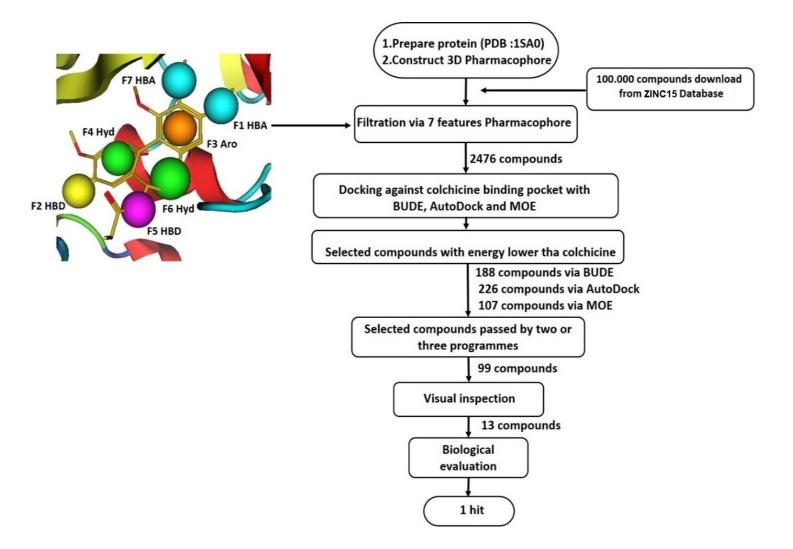
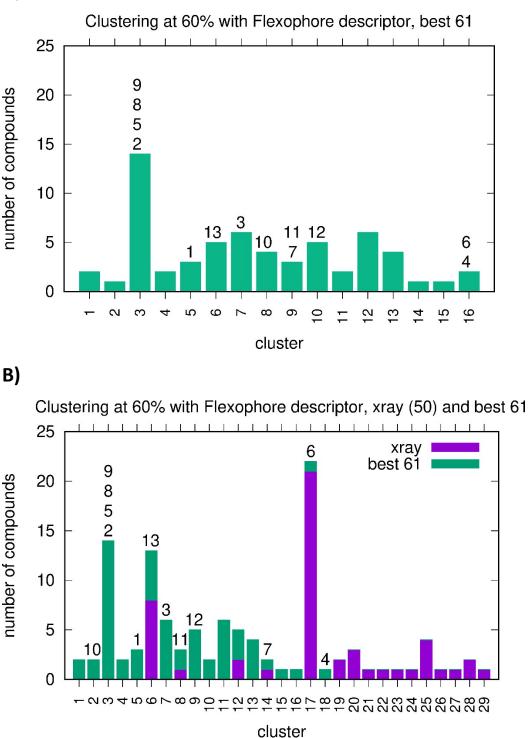
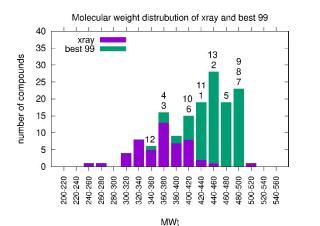


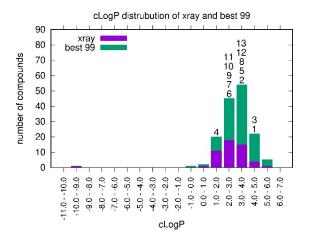
Figure S1 Schematic view of pharmacophore structure-based virtual screening.

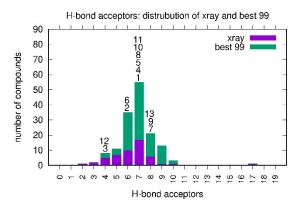


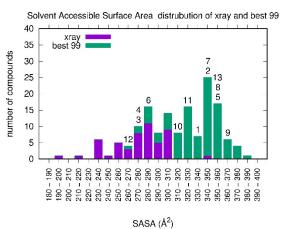


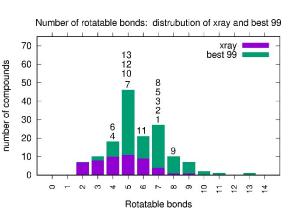
**Figure S2 A)** Clustering of the best 61 compounds with the Flexophore descriptor at 60% similarity. **B)** Clustering of the best 61 compounds with the 50 ligands in the colchicine site from crystal structures reported to date, using the Flexophore descriptor at 60% similarity. The cluster locations of **1-13** are indicated.

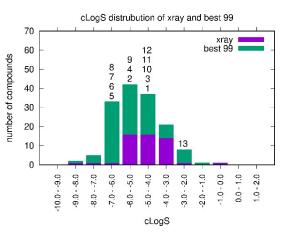


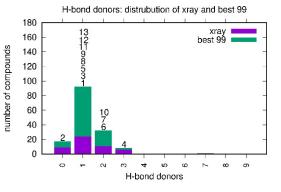




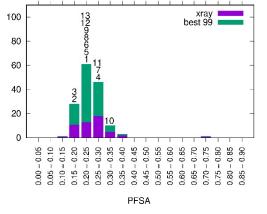








Polar Fractional Surface Area distrubution of xray and best 99



**Figure S3** Histograms of ligand properties, comparing the 99 ligands from virtual screening with the 50 ligands from known crystal structures. Locations of the shortlisted compounds are indicated by the numbers.

number of compounds

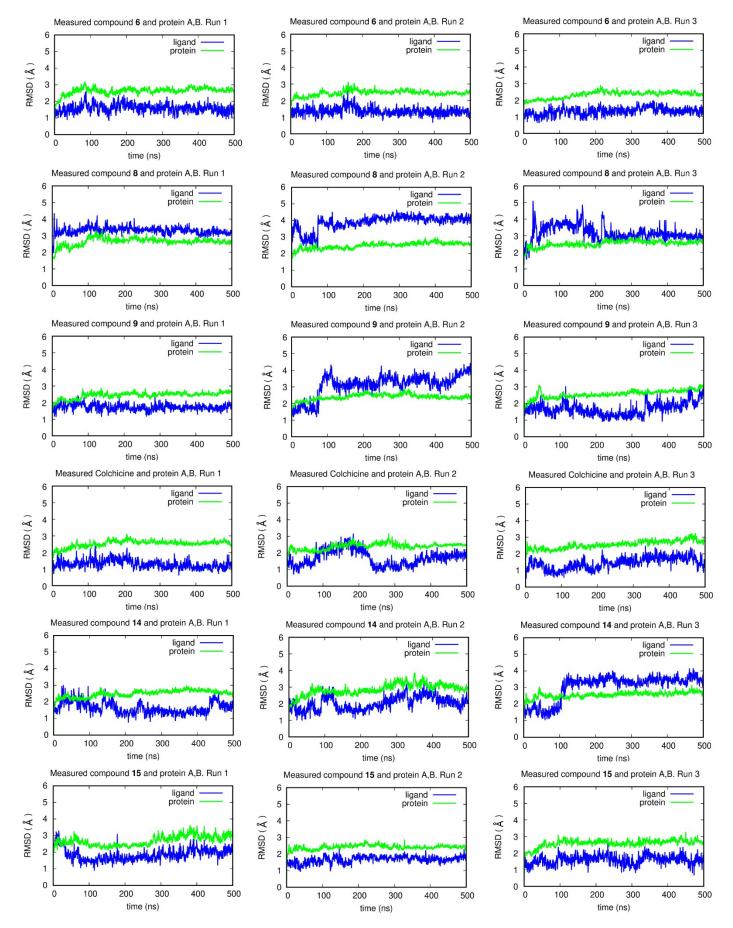
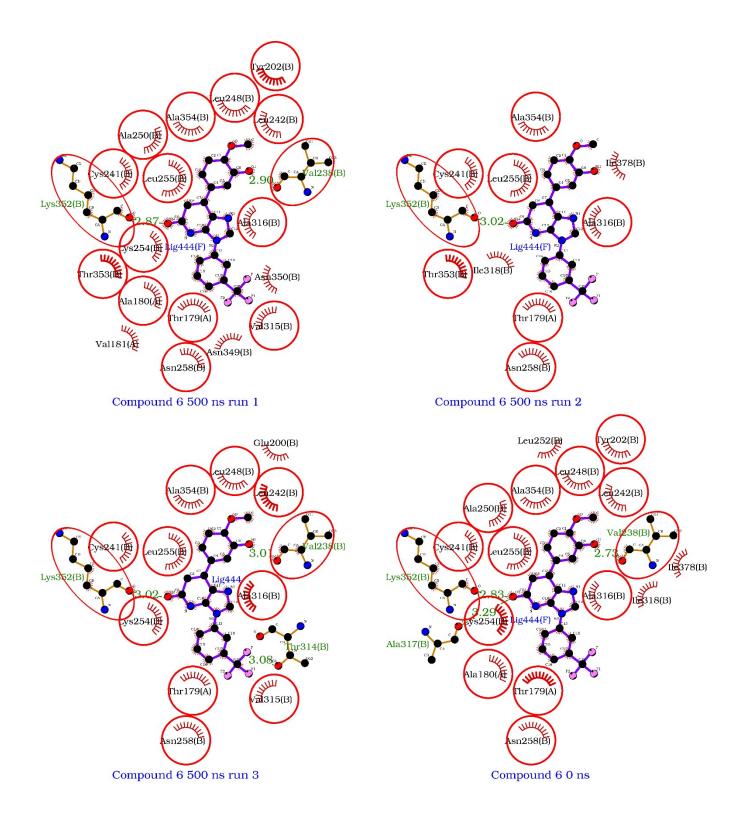
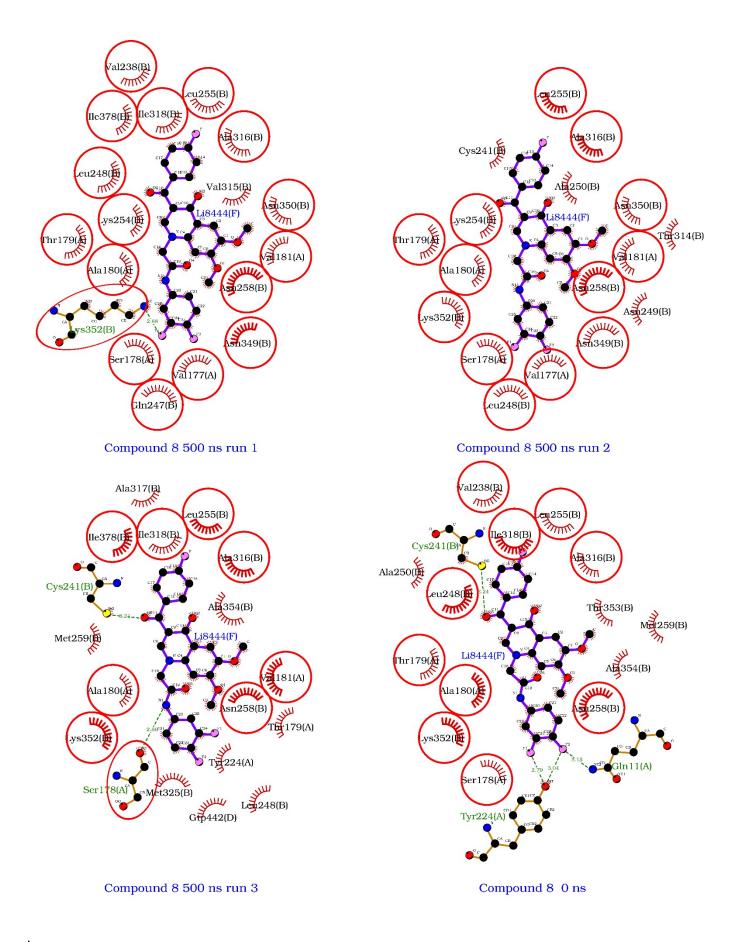


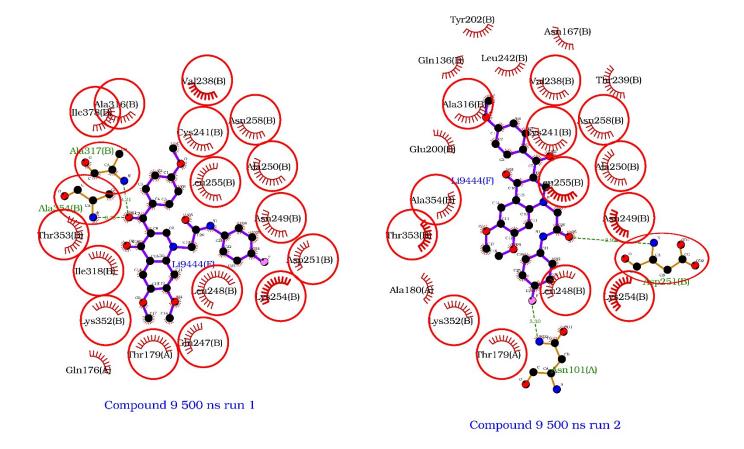
Figure S4 RMSD plots of all protein atoms of tubulin subunits A and B ( $\alpha$  and  $\beta$ ) superimposed on the time = 0 ns structure over the trajectories and measuring

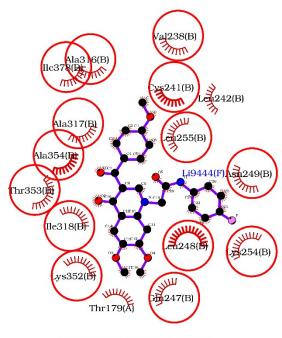
all atoms of the tubulin protein subunits A and B (green) and all atoms of the ligand (blue).



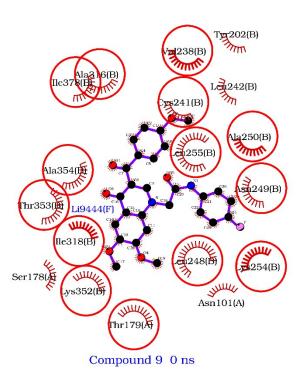
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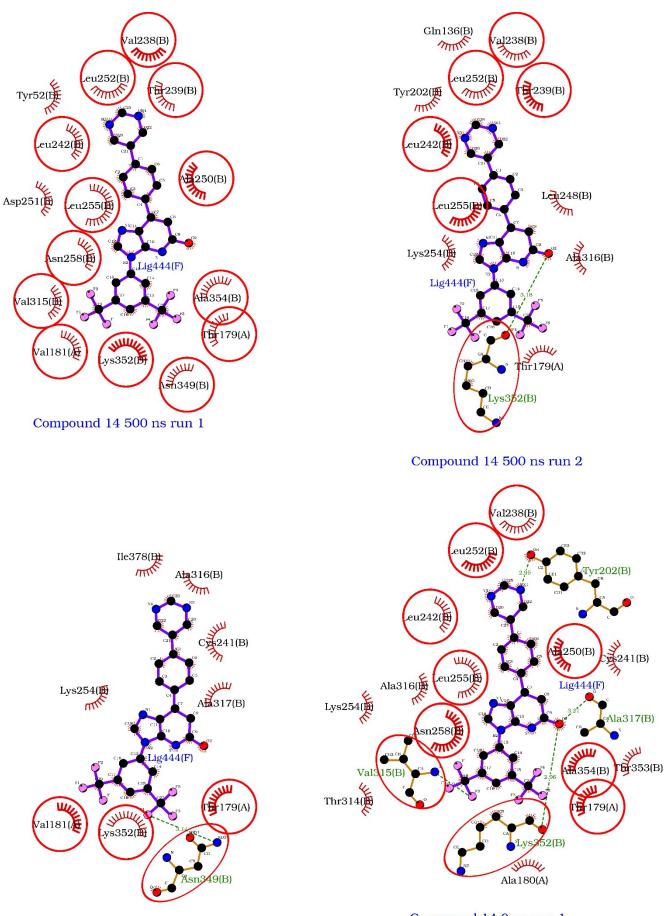






Compound 9 500 ns run 3

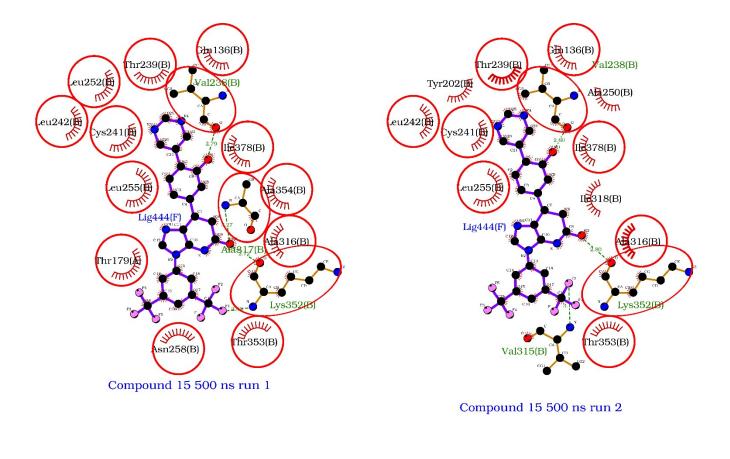


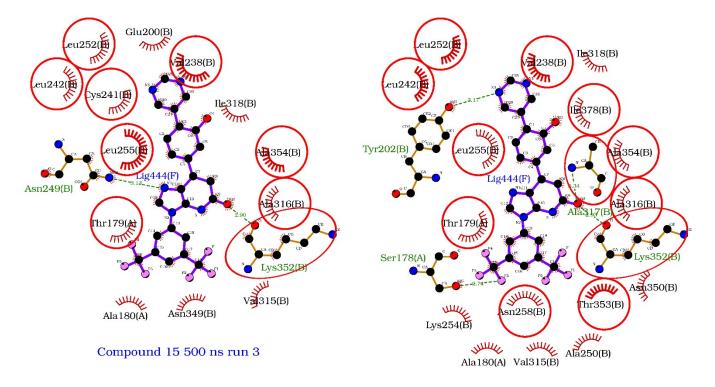


Compound 14 500 ns run 3

.

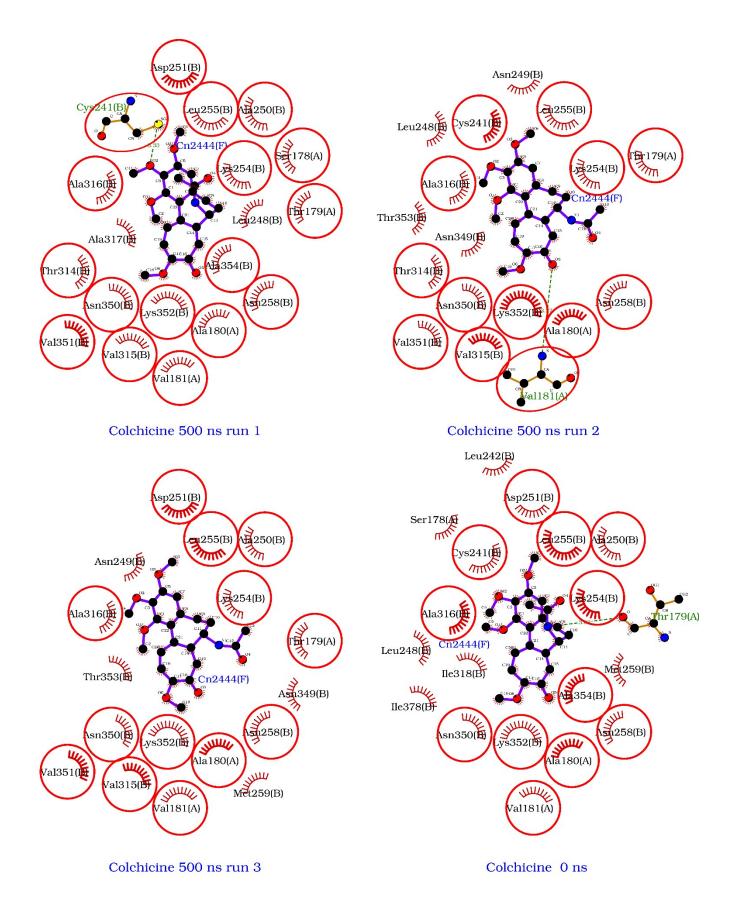
Compound 14 0 ns run 1





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Compound 15 0 ns run 1



**Figure S5** Ligplot<sup>+</sup> diagrams of the 500 ns structures for each simulation showing ligand-residue contacts at that moment, plus their respective starting structures.

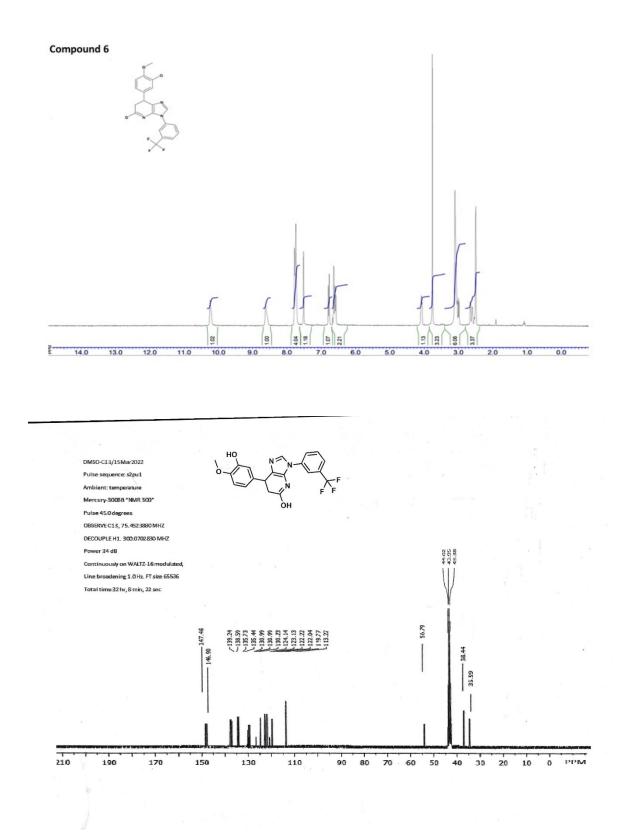


Figure S6 <sup>1</sup>H and <sup>13</sup>C NMR spectra of compound 6.

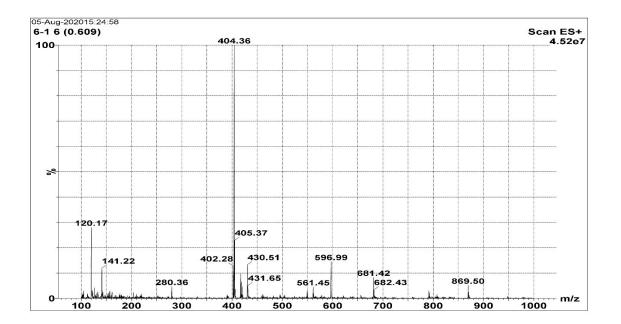
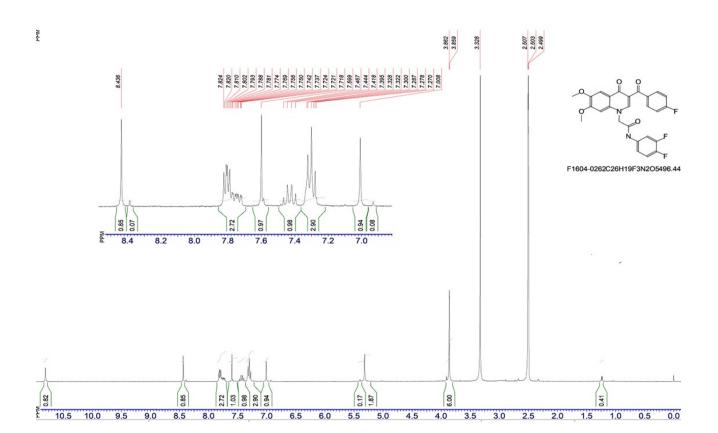


Figure S7 Mass spectrum of compound 6.



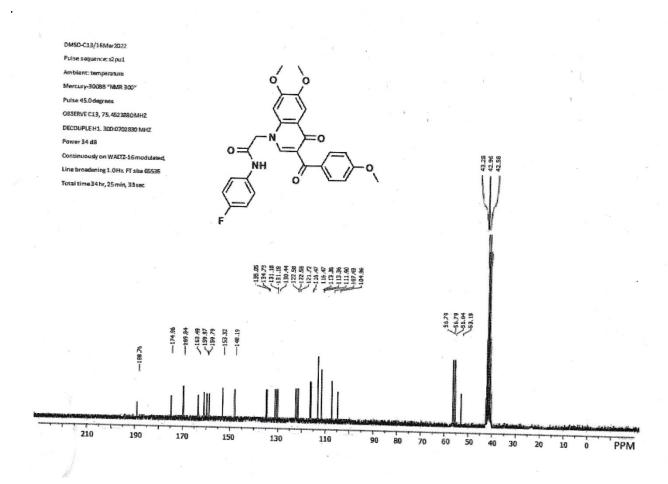


Figure S8 <sup>1</sup>H and <sup>13</sup>C NMR spectrum of compound 8

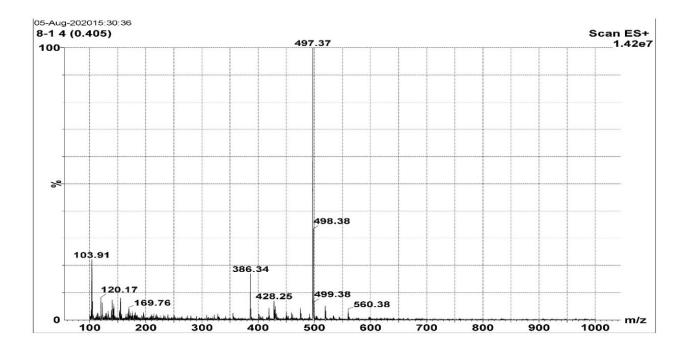


Figure S9 Mass spectrum of compound 8.

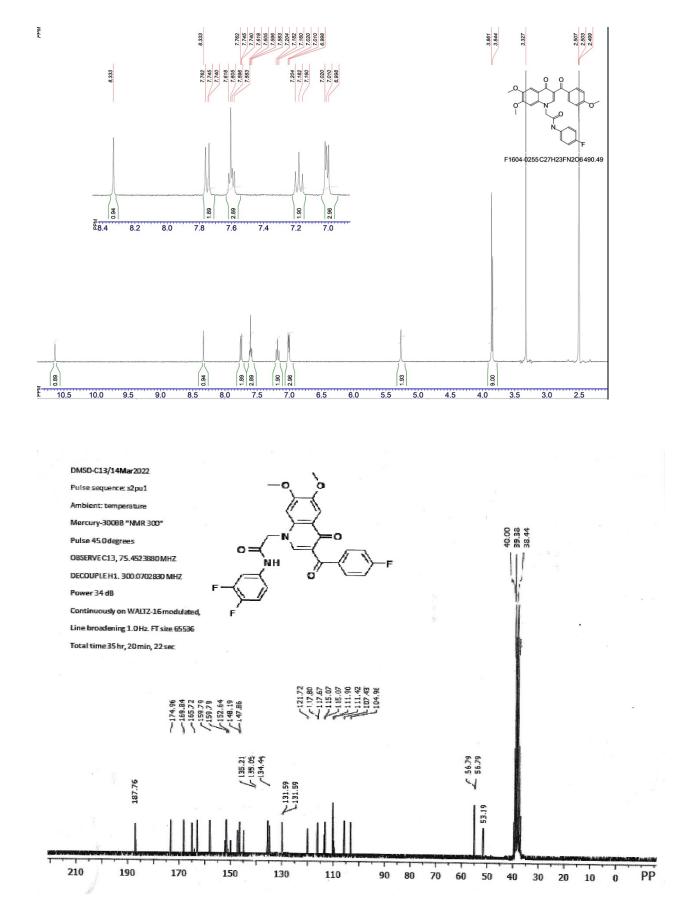
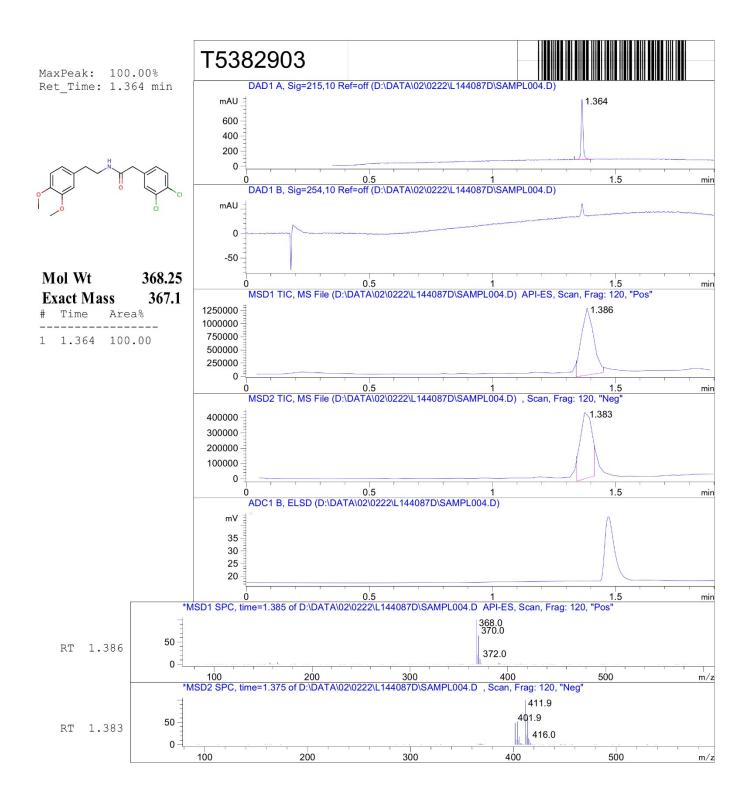
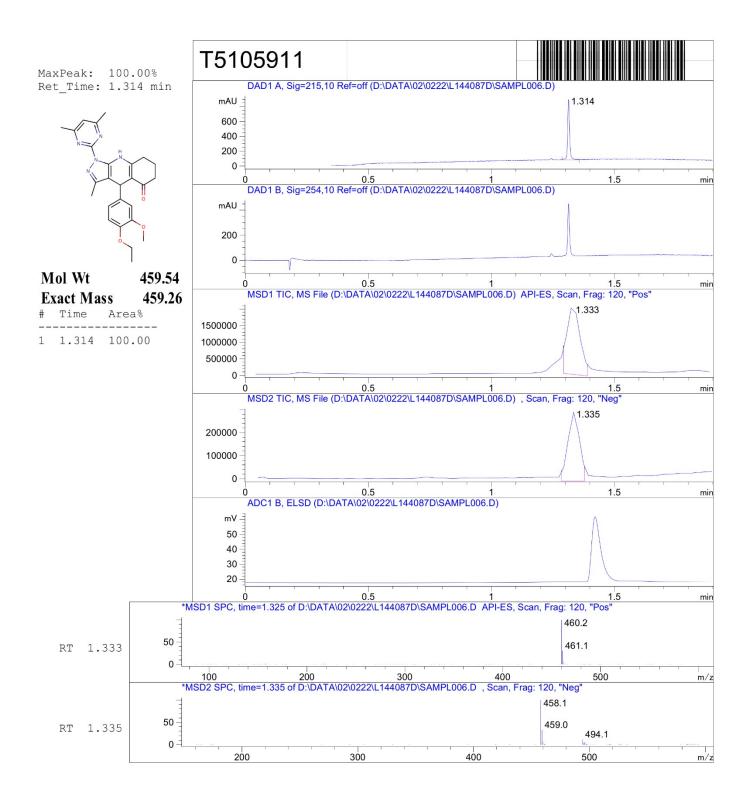


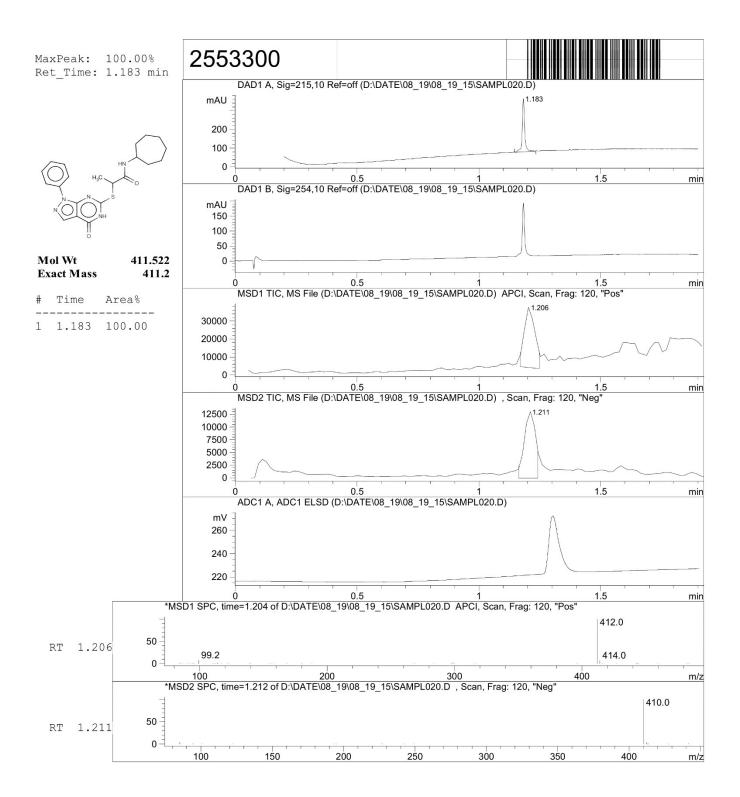
Figure S10 <sup>1</sup>H and <sup>13</sup>C NMR spectrum of compound 9



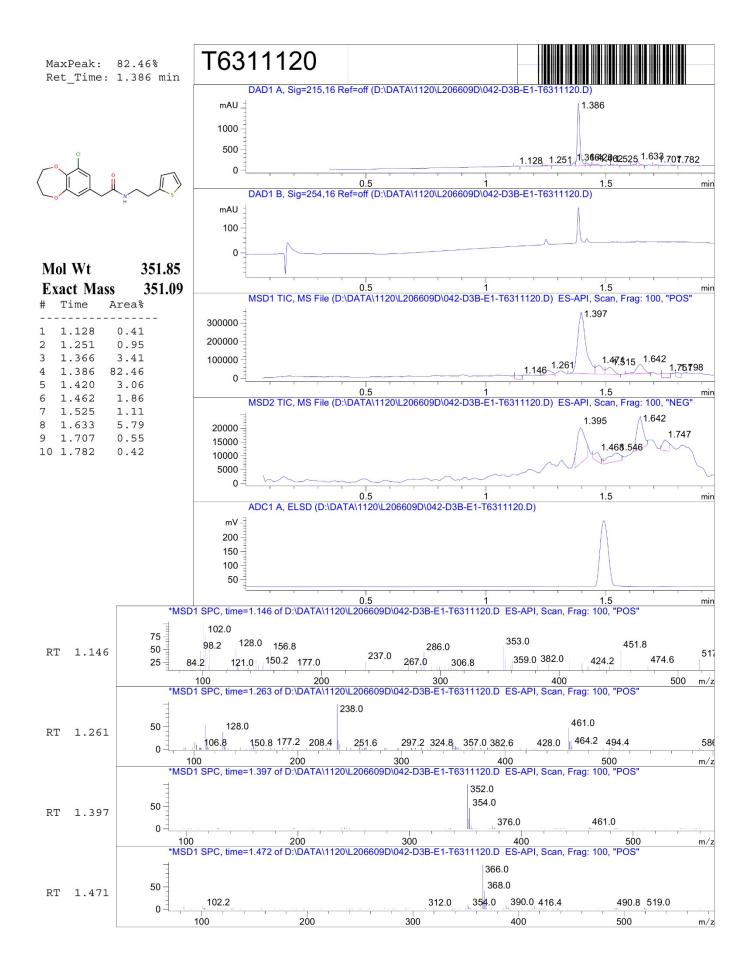
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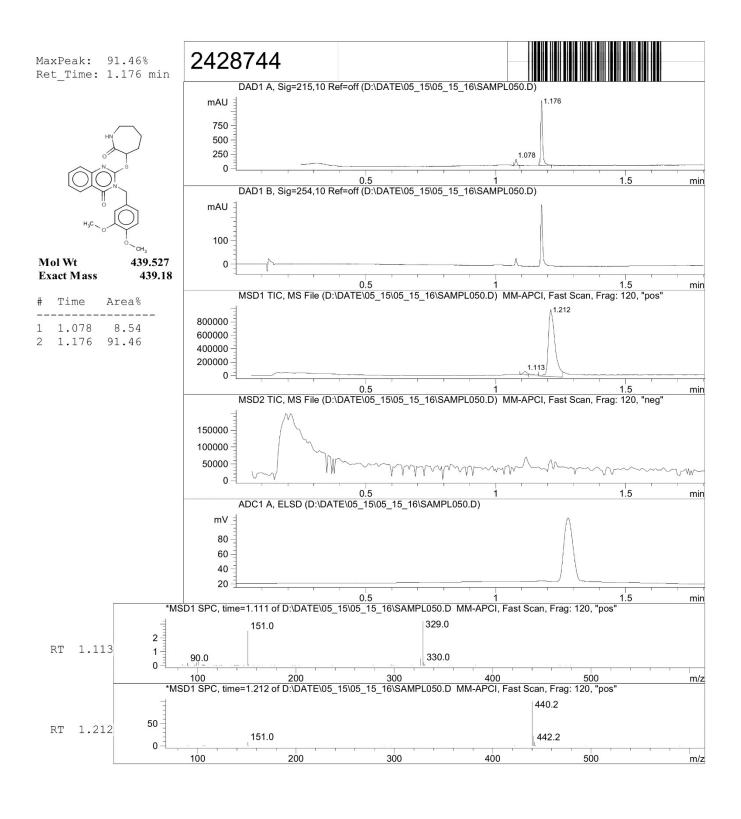


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## Figure S11 Spectra and analytical date of the short-listed compounds