

# Supporting Information

---

## Rational development of a potent 15-lipoxygenase-1 inhibitor with *in vitro* and *ex vivo* anti-inflammatory properties

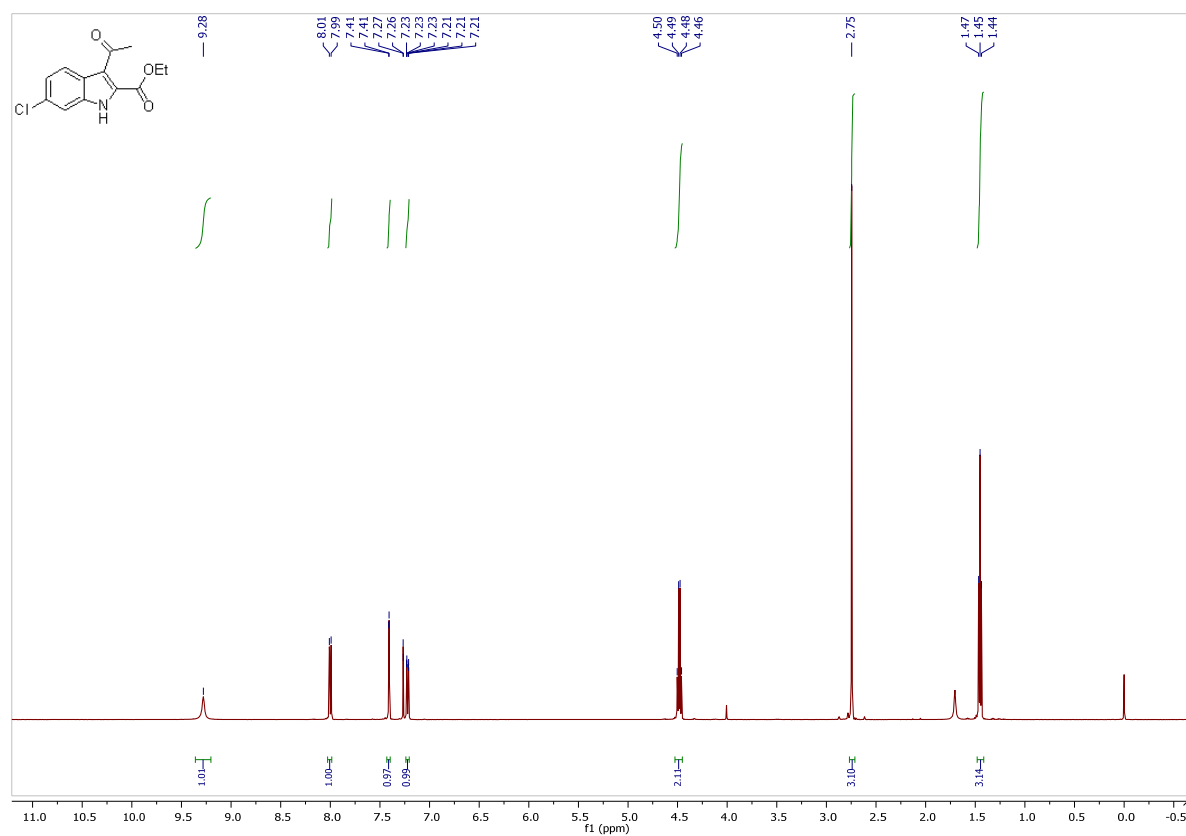
Nikolaos Eleftheriadis,<sup>†</sup> Constantinos G. Neochoritis,<sup>‡</sup> Niek G.J. Leus,<sup>†</sup> Petra E. van der Wouden,<sup>†</sup> Alexander Dömling,<sup>‡</sup> Frank J. Dekker<sup>†,\*</sup>

### Table of Contents

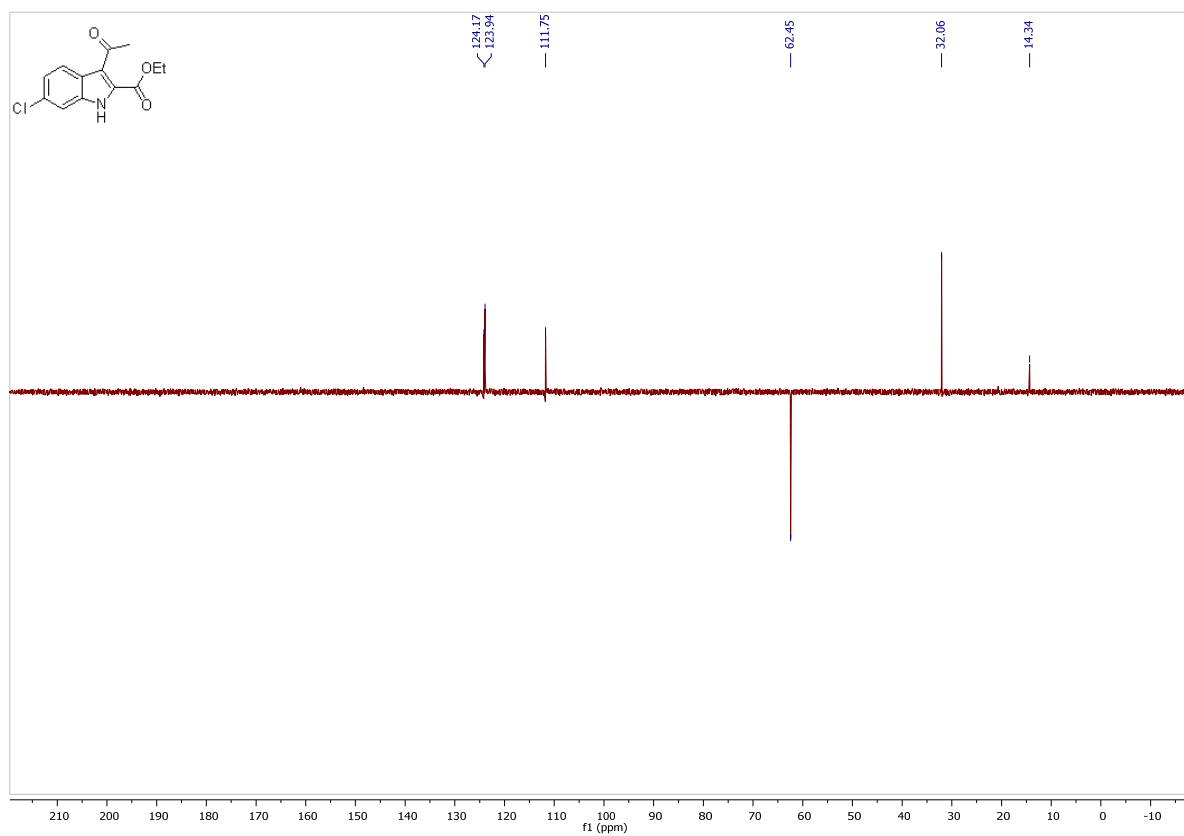
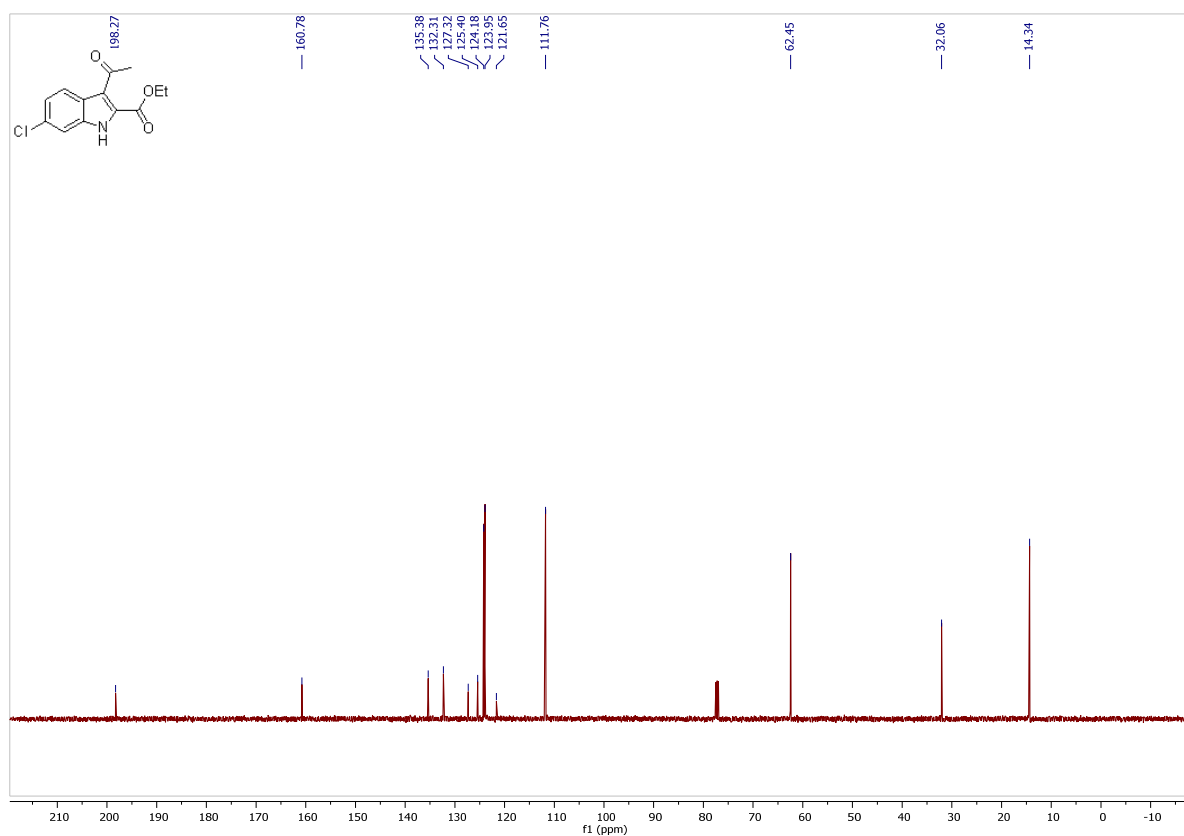
|                                         |    |
|-----------------------------------------|----|
| NMR spectra .....                       | 2  |
| Enzyme inhibition studies .....         | 32 |
| Focused library .....                   | 32 |
| Screening Results.....                  | 37 |
| Screening Results (column graphs) ..... | 38 |
| IC <sub>50</sub> graphs .....           | 40 |
| Enzyme Kinetics .....                   | 41 |
| Molecular Modeling .....                | 42 |

## NMR spectra

$^1\text{H}$  NMR spectrum of **14a**, DN397



<sup>13</sup>C NMR and DEPT-135 spectra of **14a**, DN397

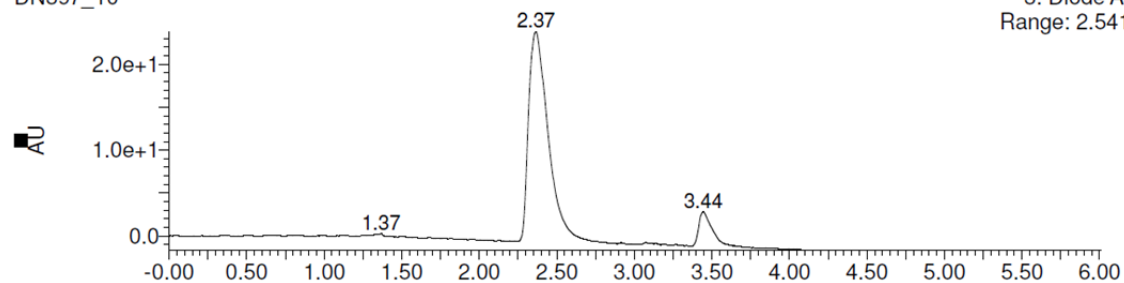


Chromatogram and MS spectra of **14a**, DN397

DN397\_10\_Col4\_sol1\_5-30%\_6min

DN397\_10

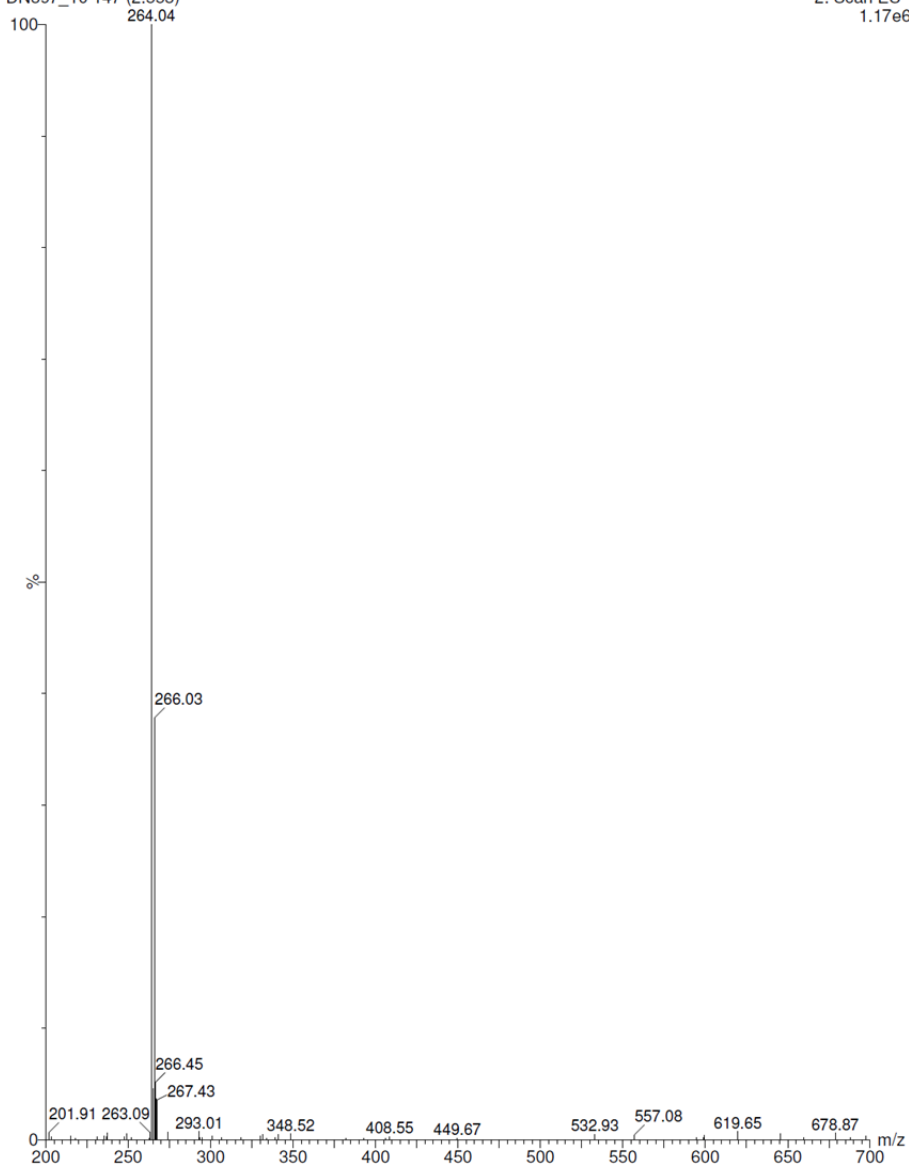
3: Diode Array  
Range: 2.541e+1



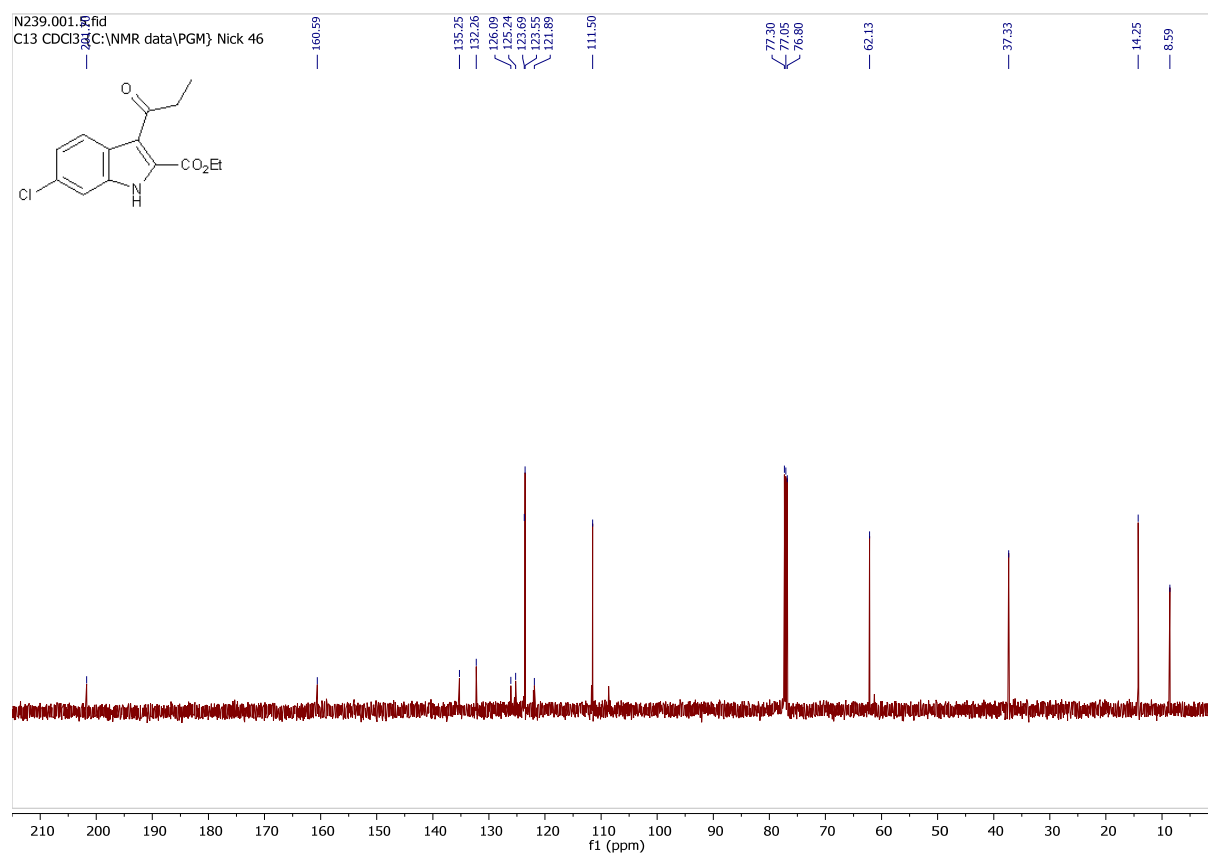
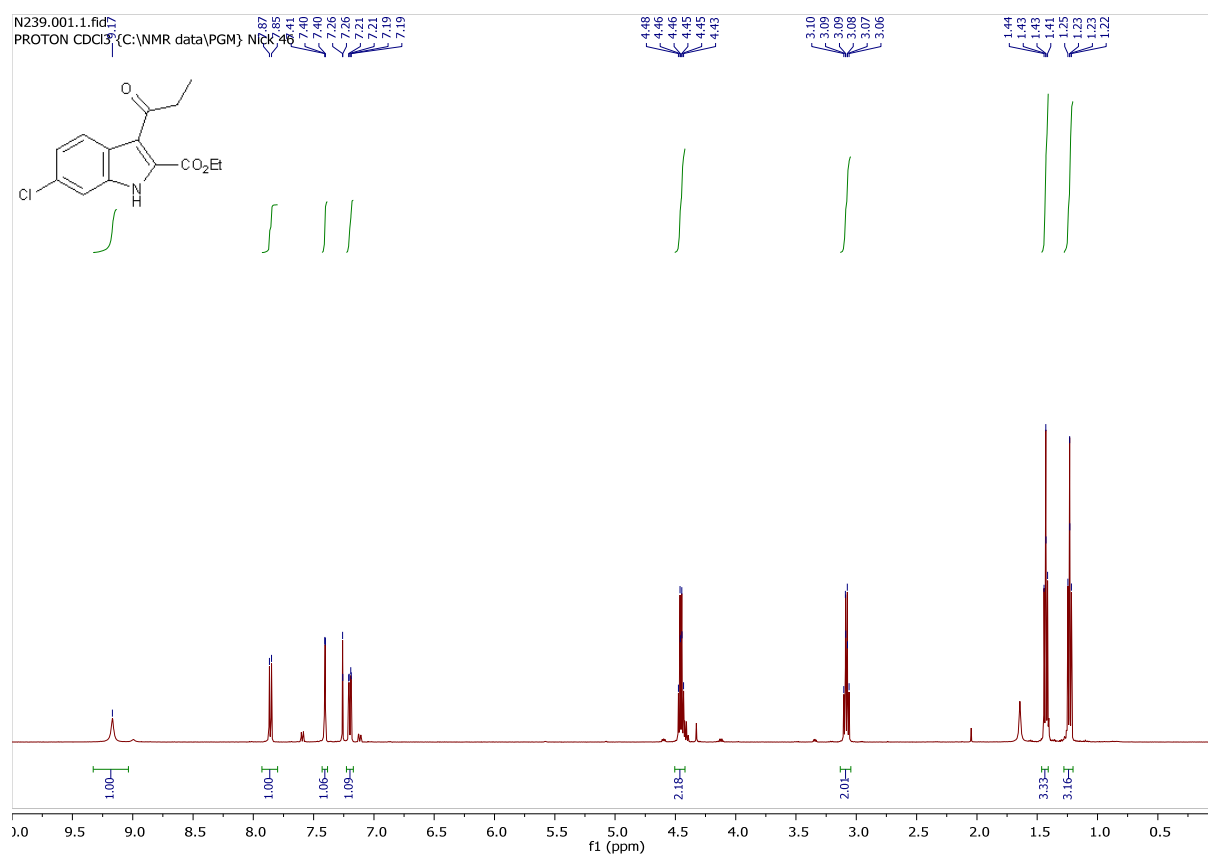
DN397\_10\_Col4\_sol1\_5-30%\_6min

DN397\_10 147 (2.553)

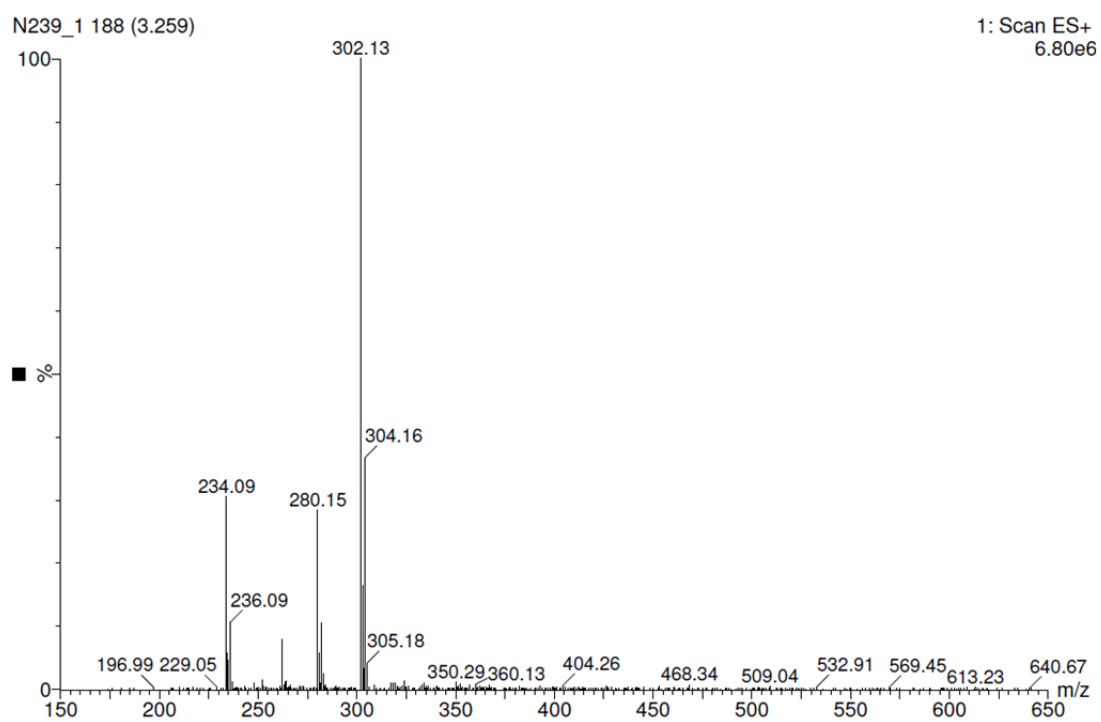
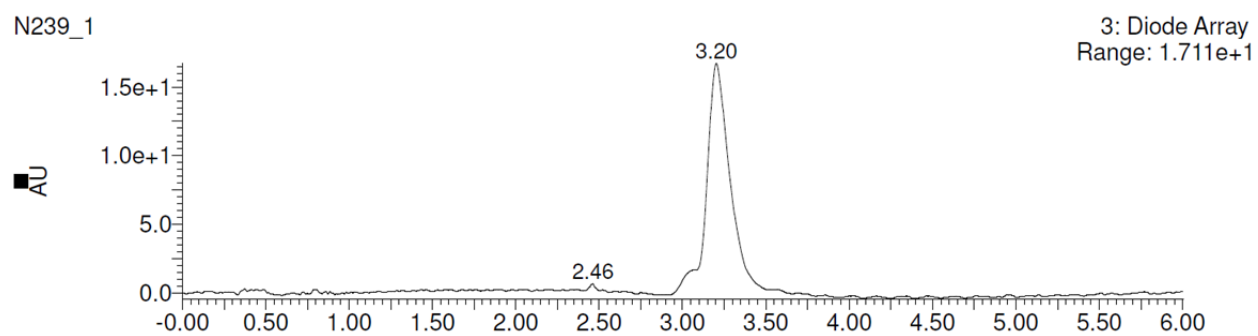
2: Scan ES-  
1.17e6



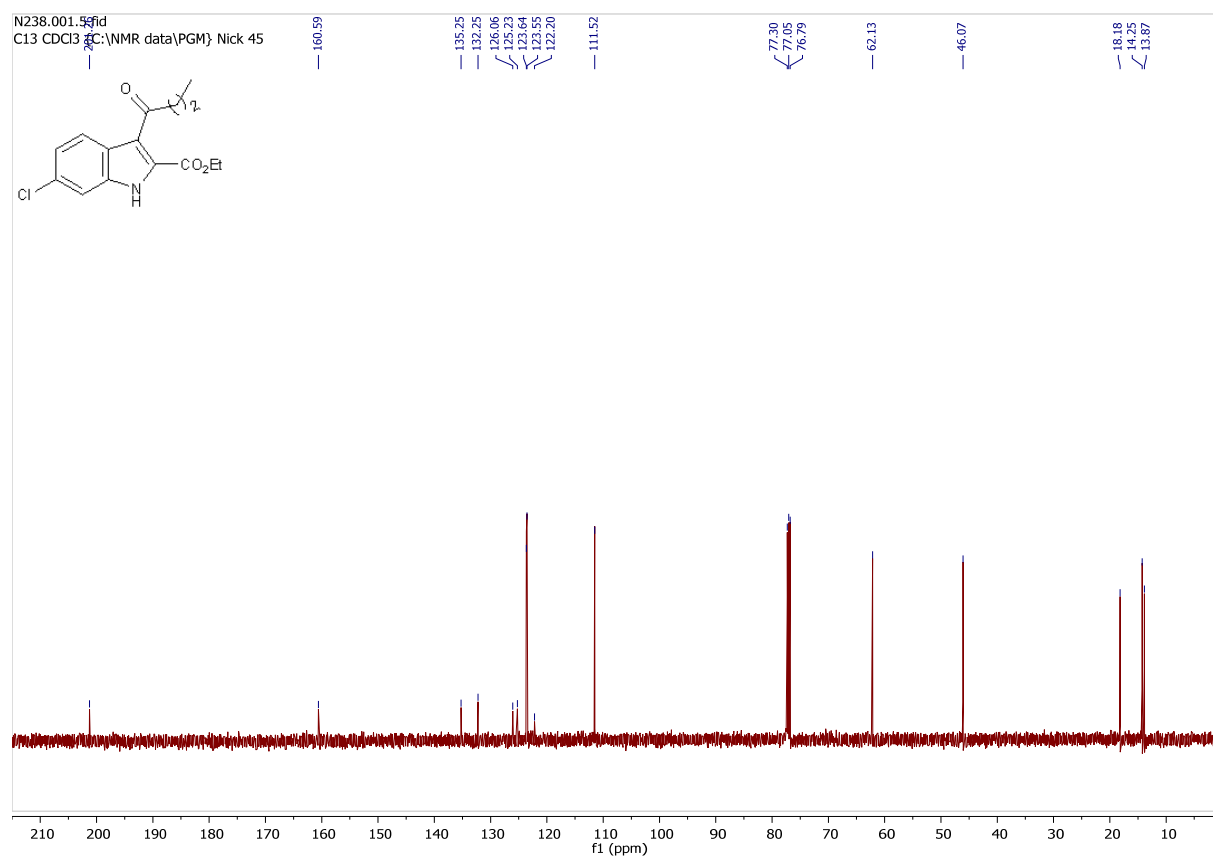
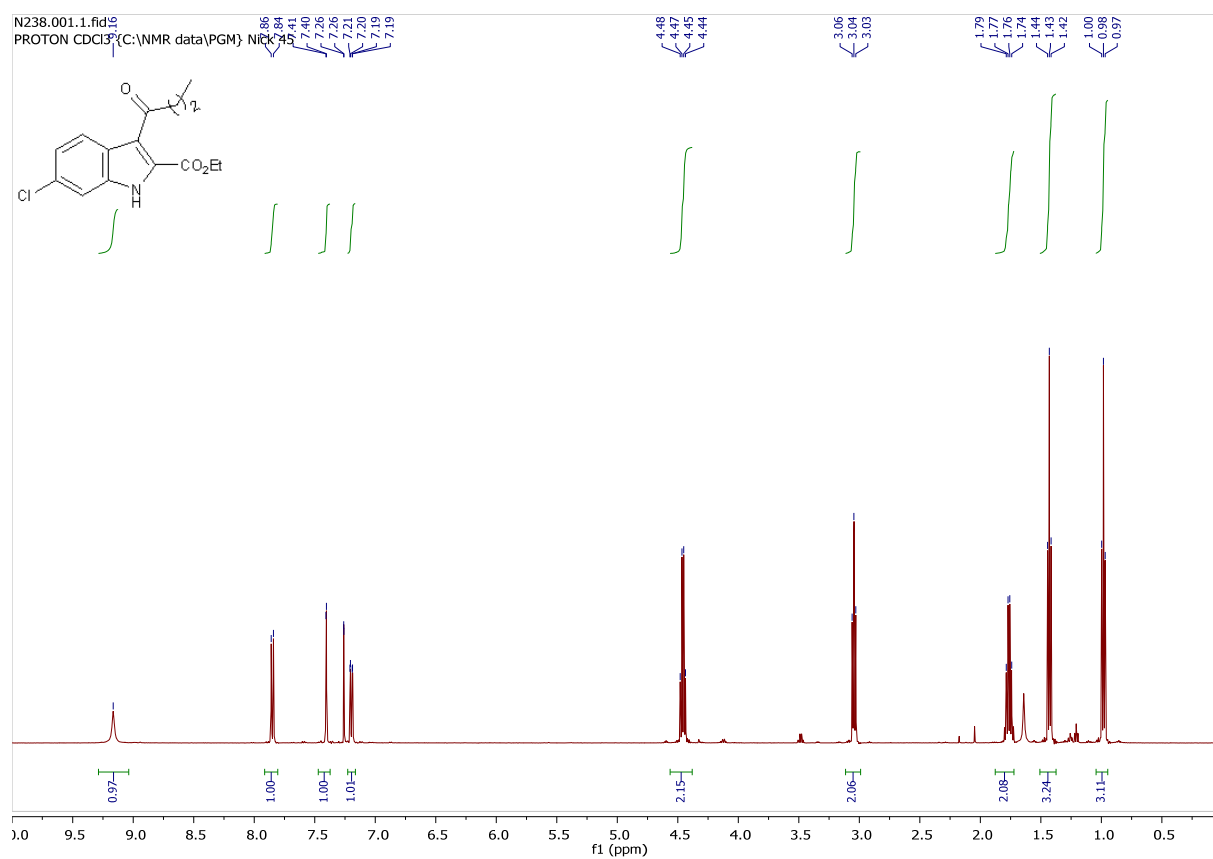
$^1\text{H}$  NMR and  $^{13}\text{C}$  NMR spectra of **14b**, N239



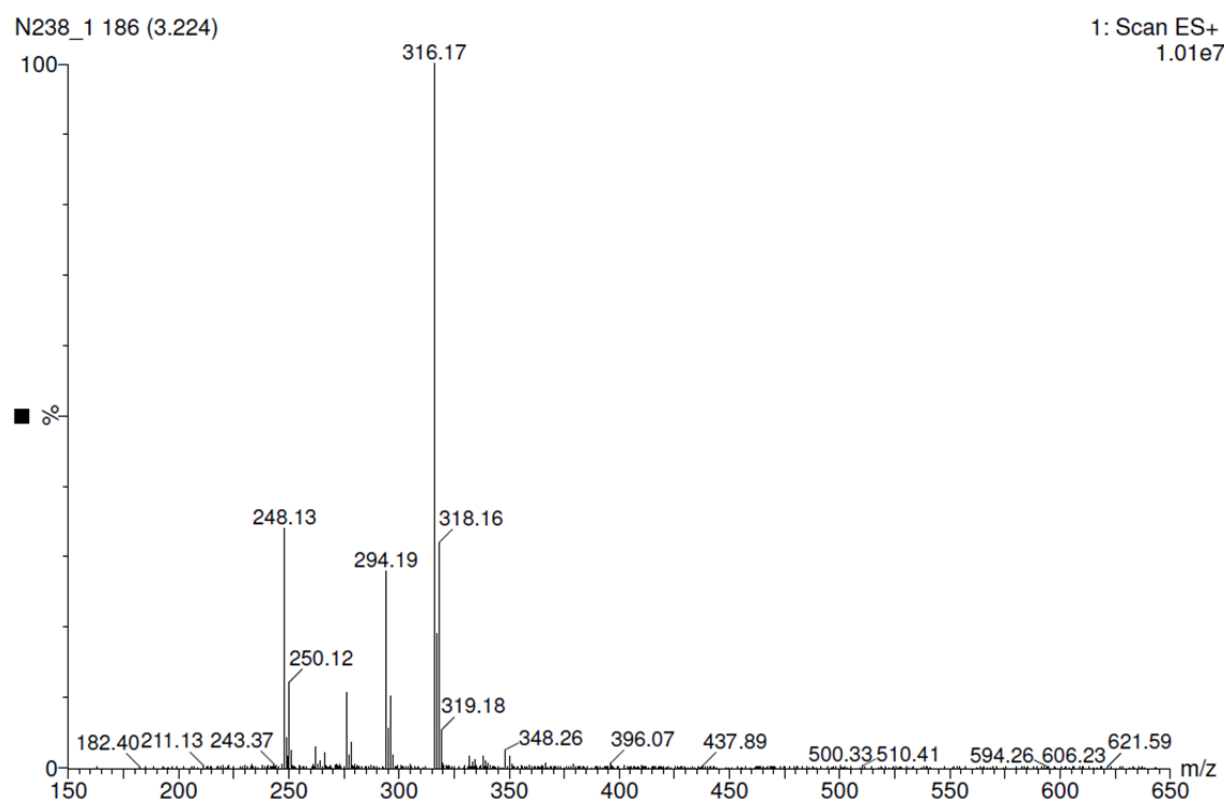
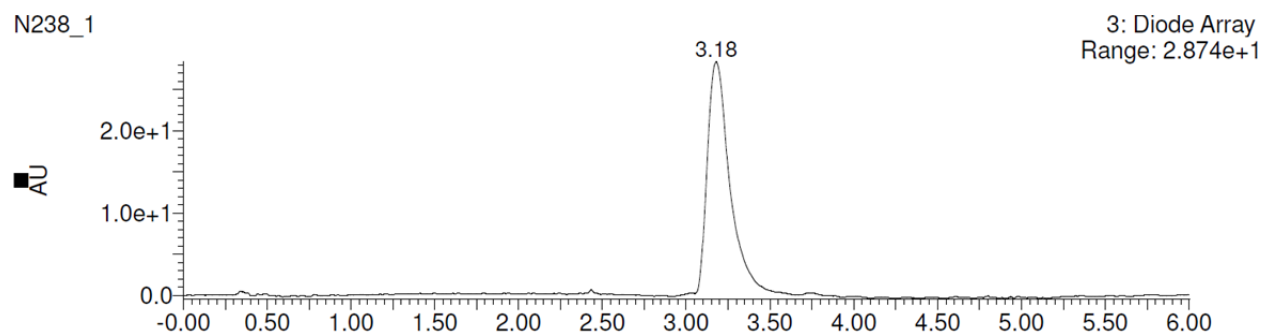
Chromatogram and MS spectra of **14b**, N239



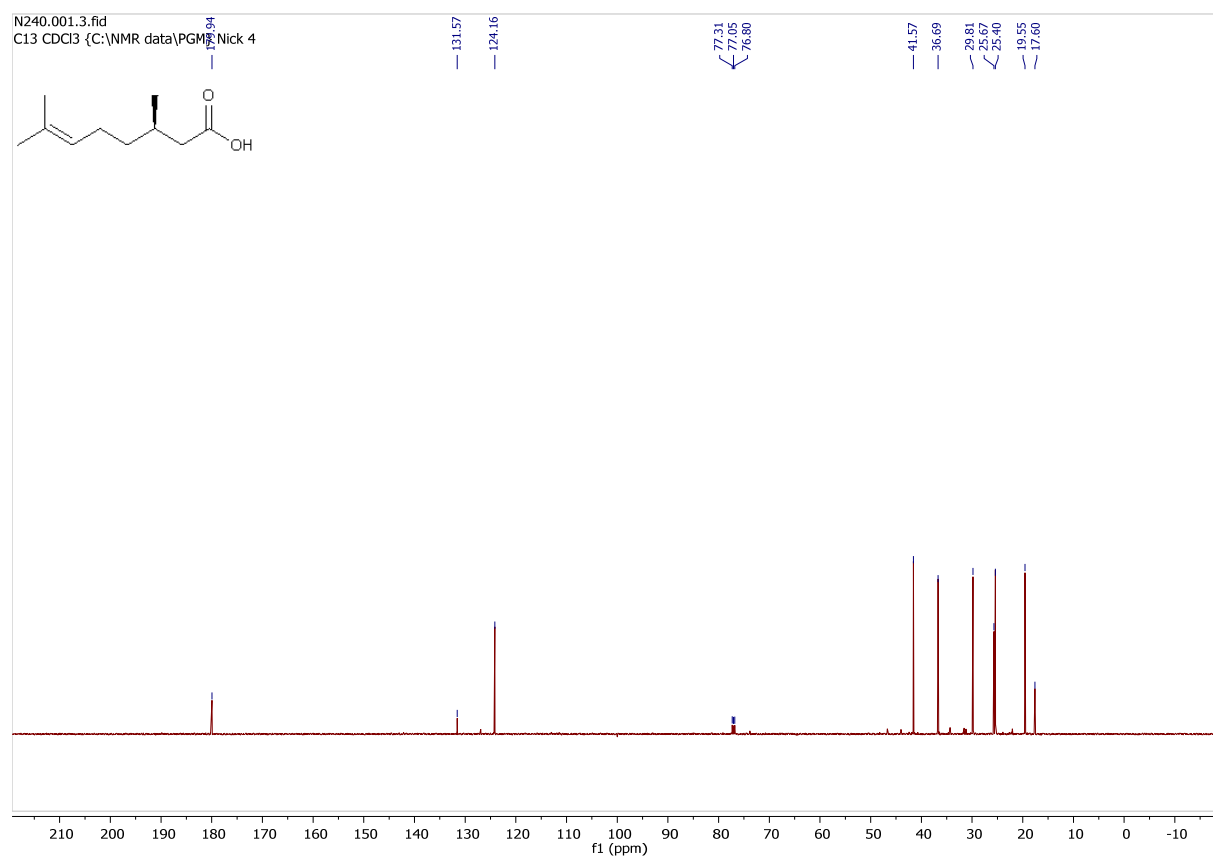
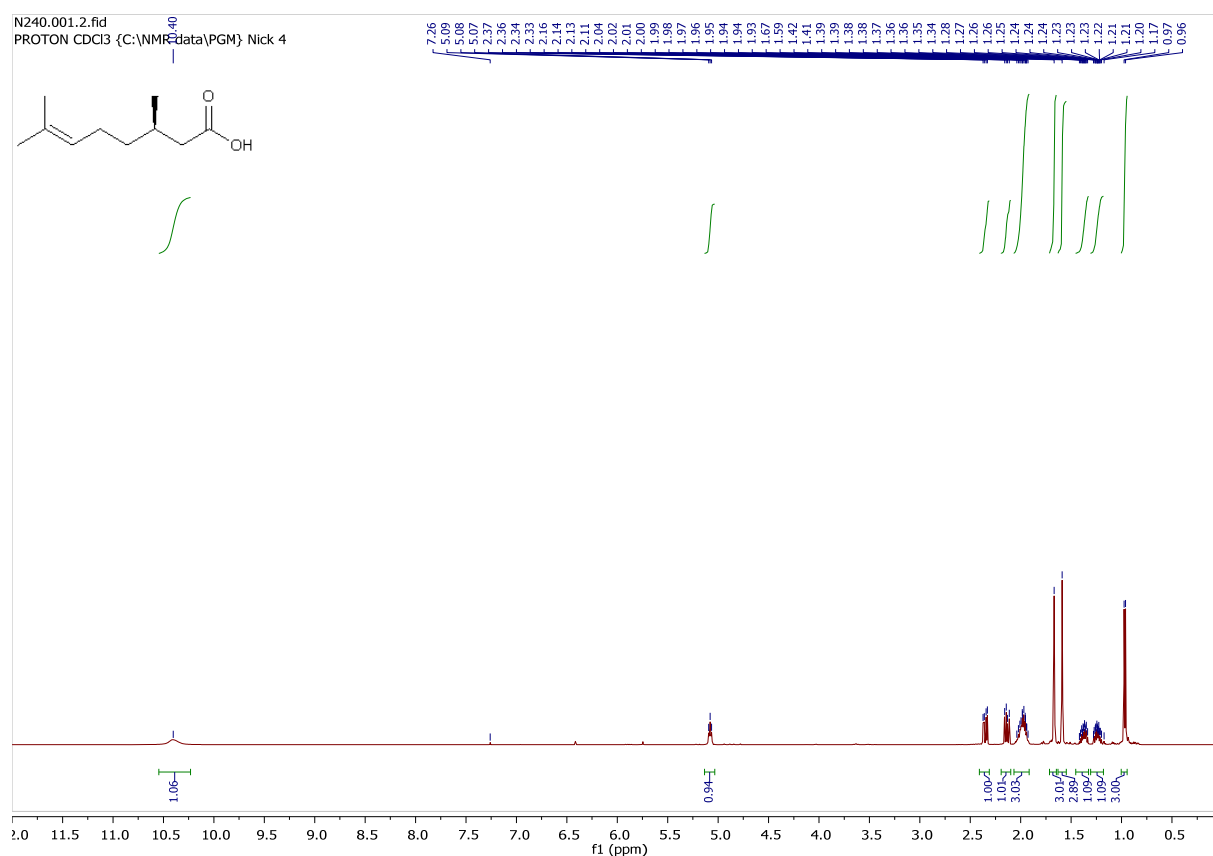
$^1\text{H}$  NMR and  $^{13}\text{C}$  NMR spectra of **14c**, N238



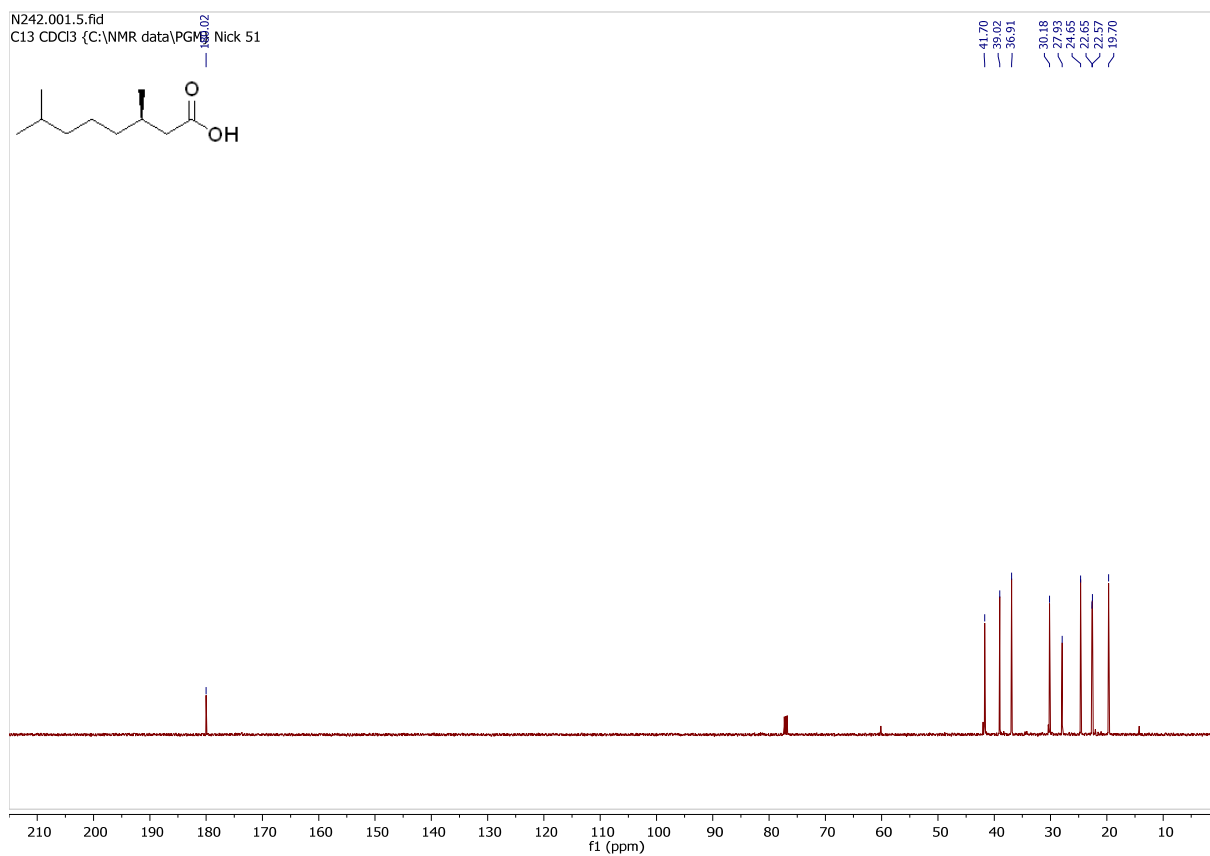
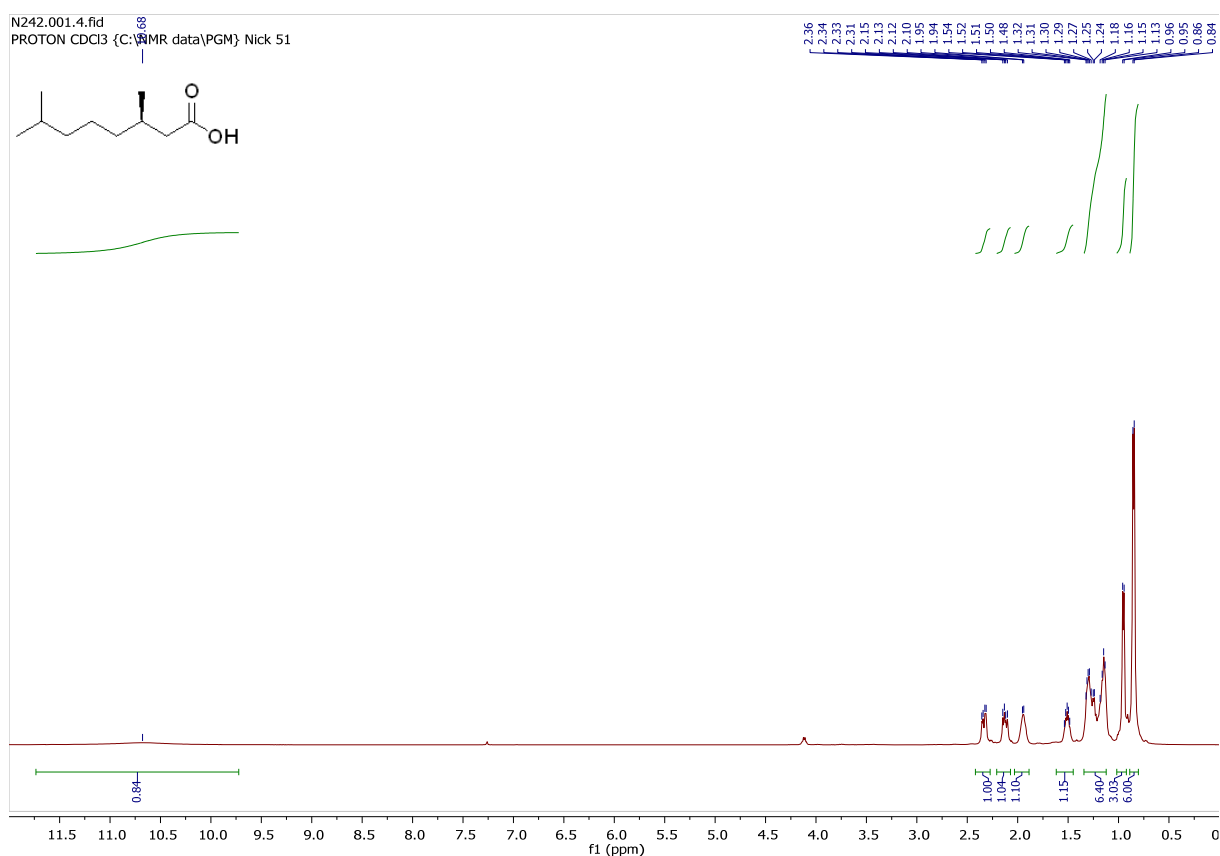
Chromatogram and MS spectra of **14c**, N238



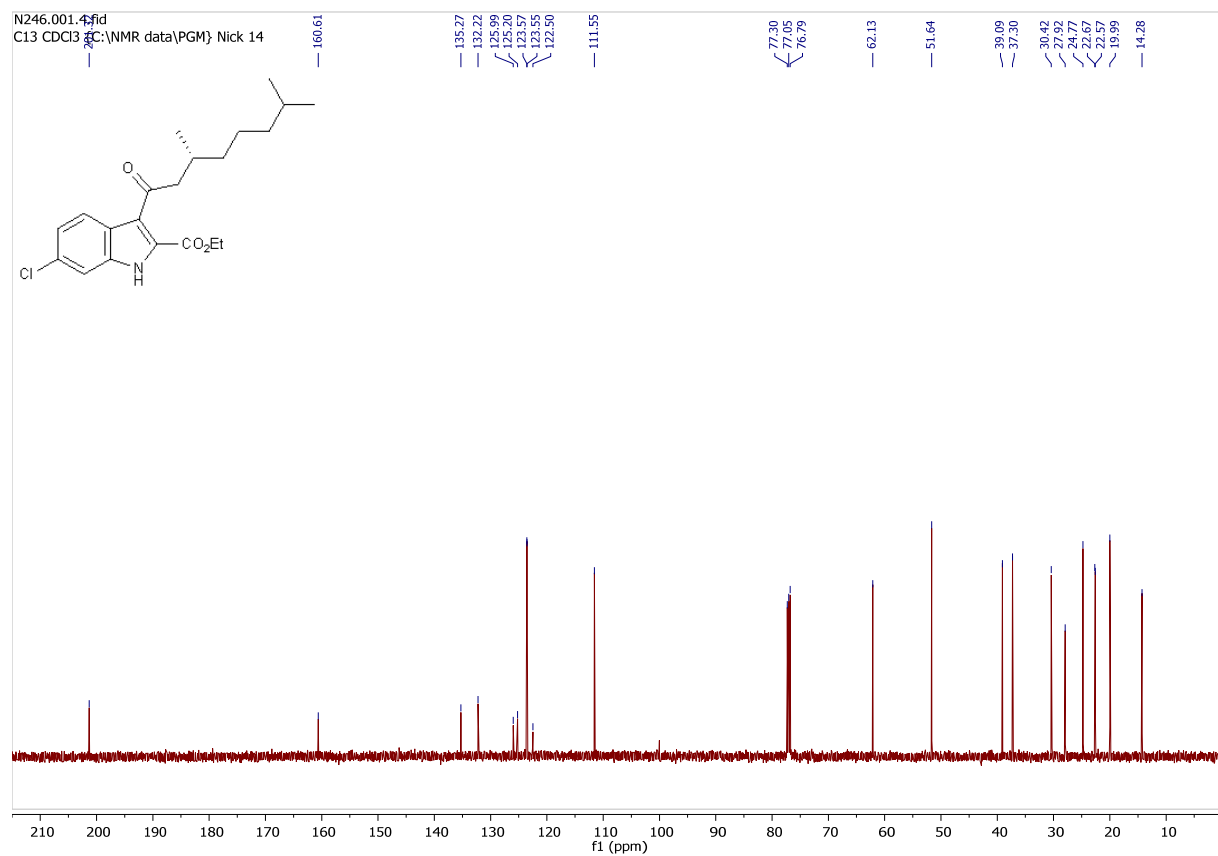
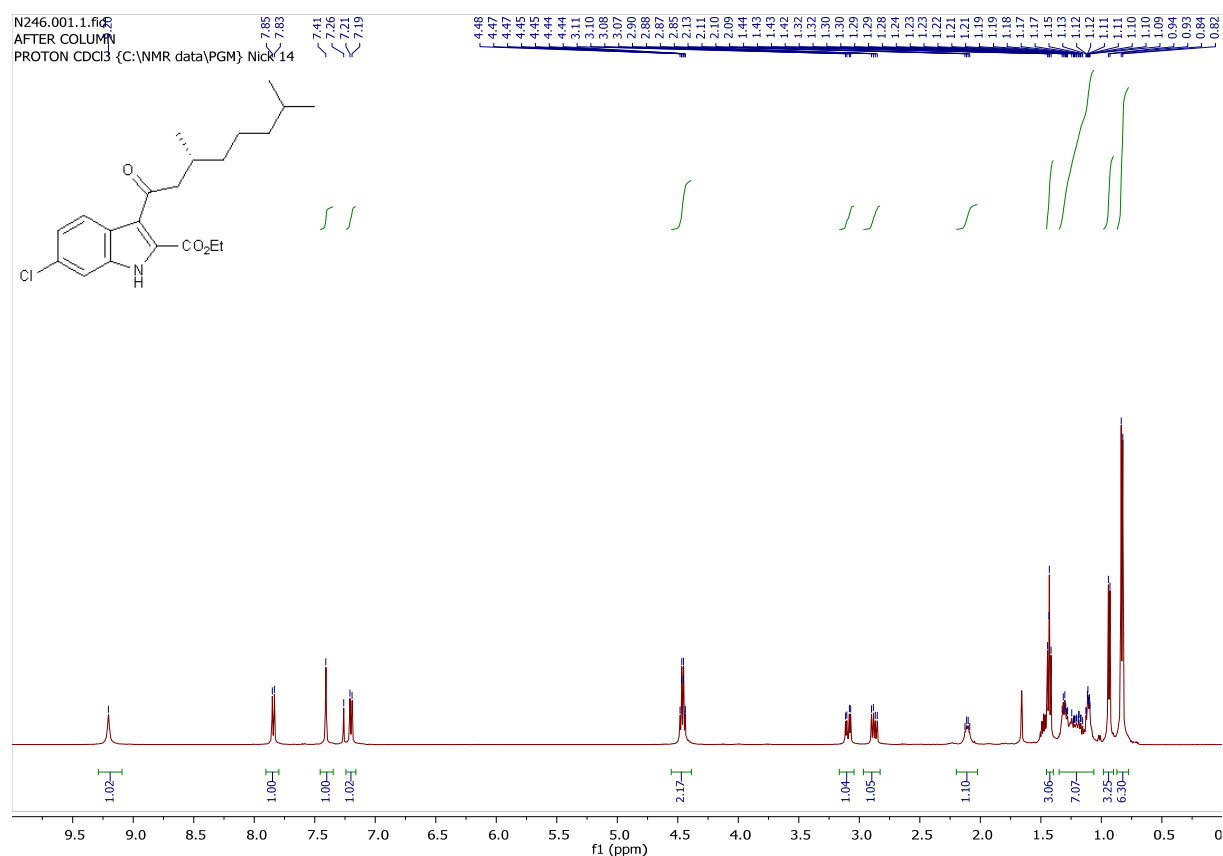
$^1\text{H}$  NMR and  $^{13}\text{C}$  NMR spectra of **11d**, N240



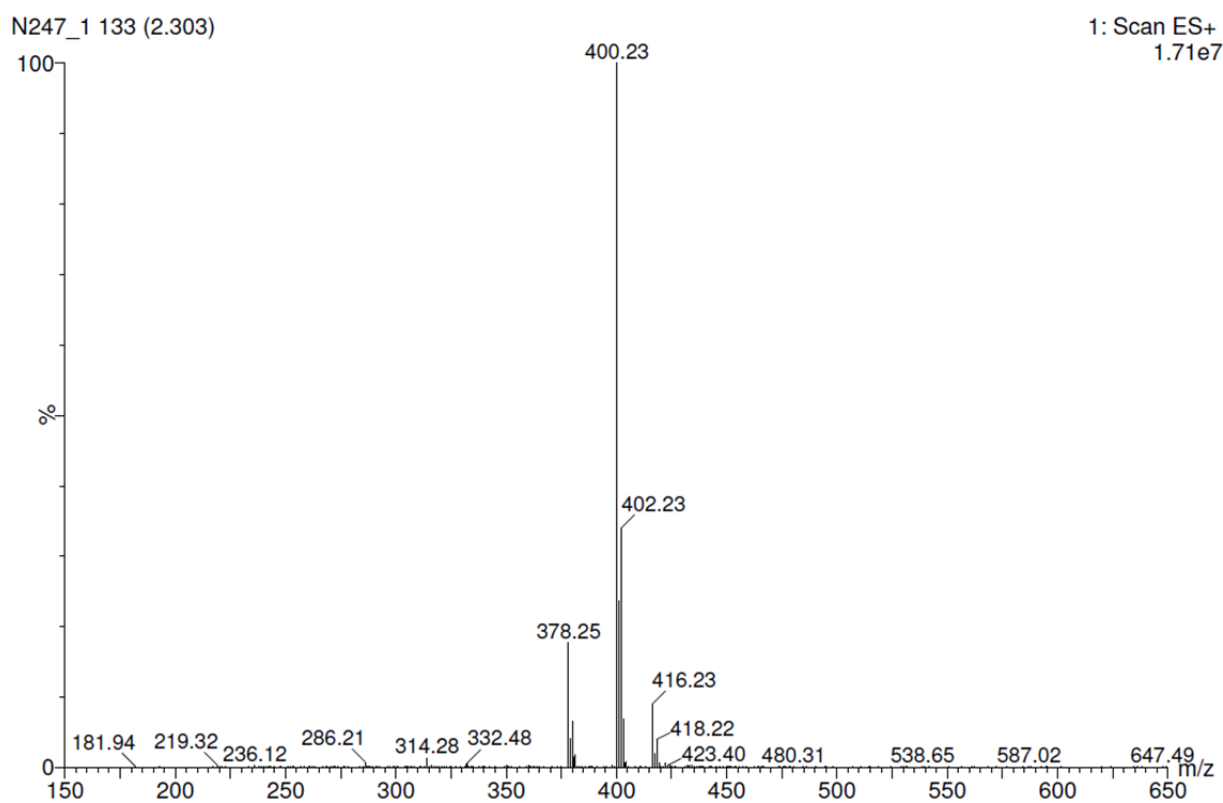
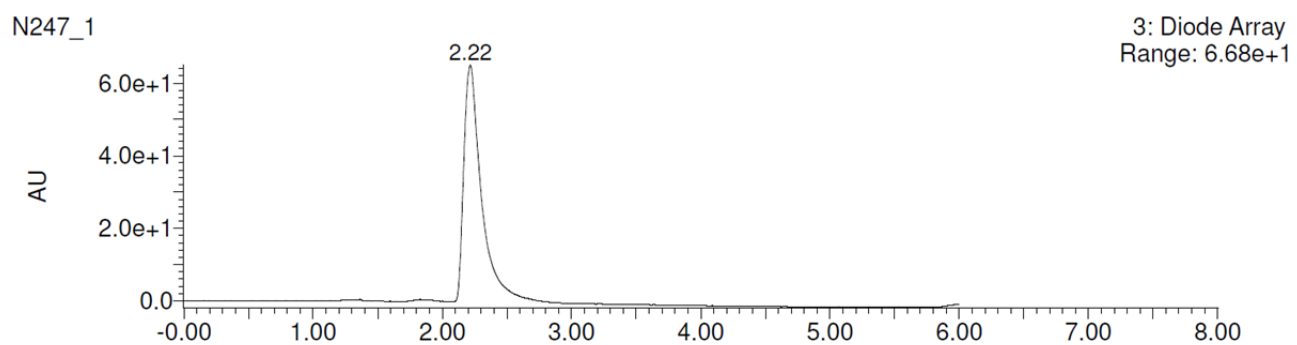
$^1\text{H}$  NMR and  $^{13}\text{C}$  NMR spectra of **12d**, N242



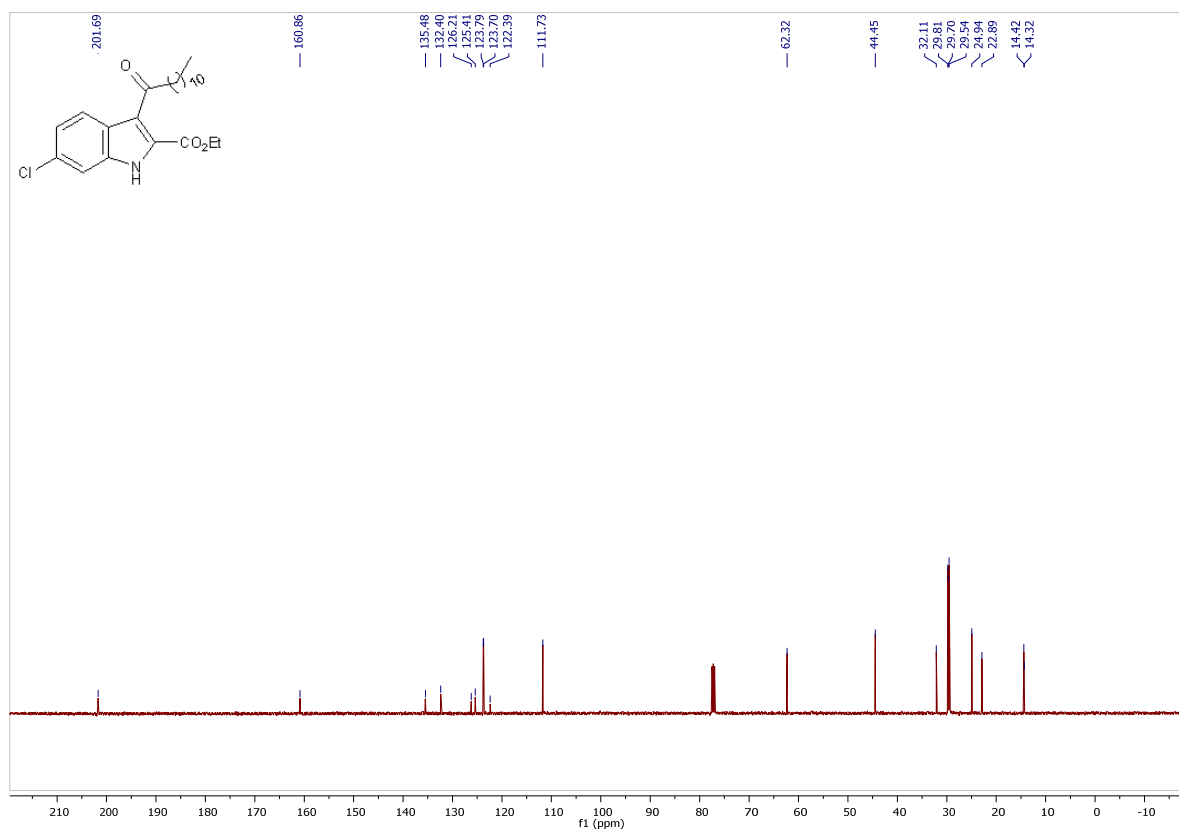
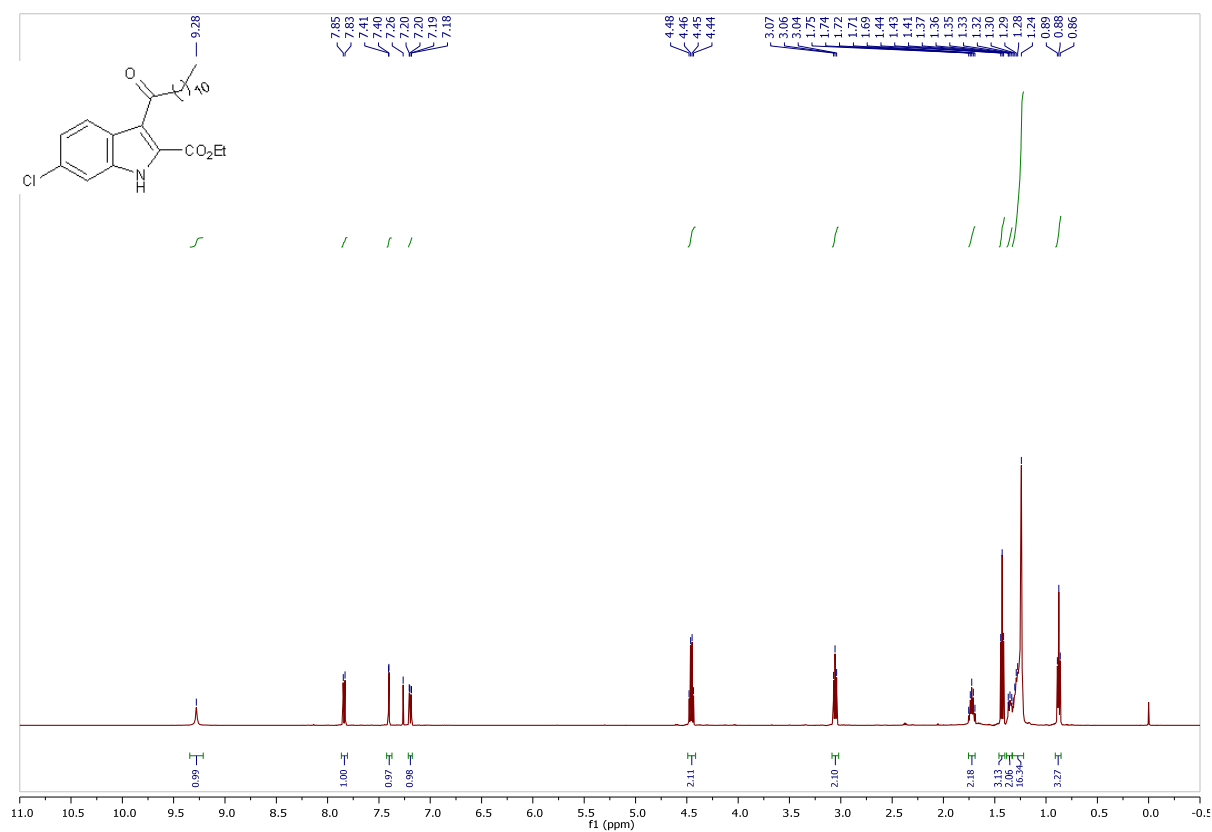
$^1\text{H}$  NMR and  $^{13}\text{C}$  NMR spectra of **14e**, N246



Chromatogram and MS spectra of **14d**, N247

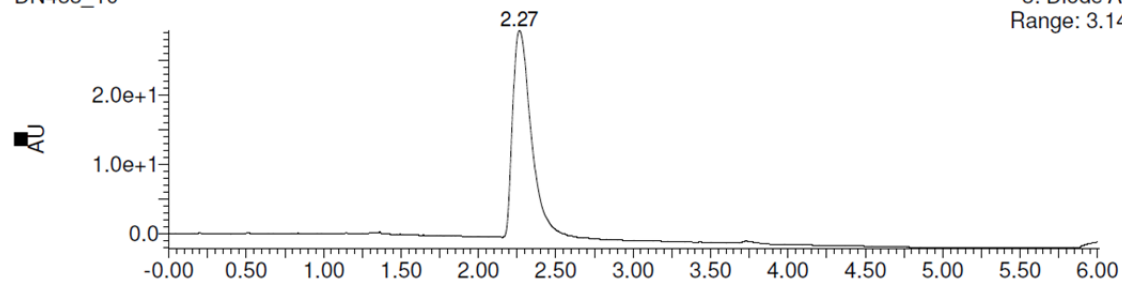


$^1\text{H}$  NMR and  $^{13}\text{C}$  NMR spectra of **14f**, DN433

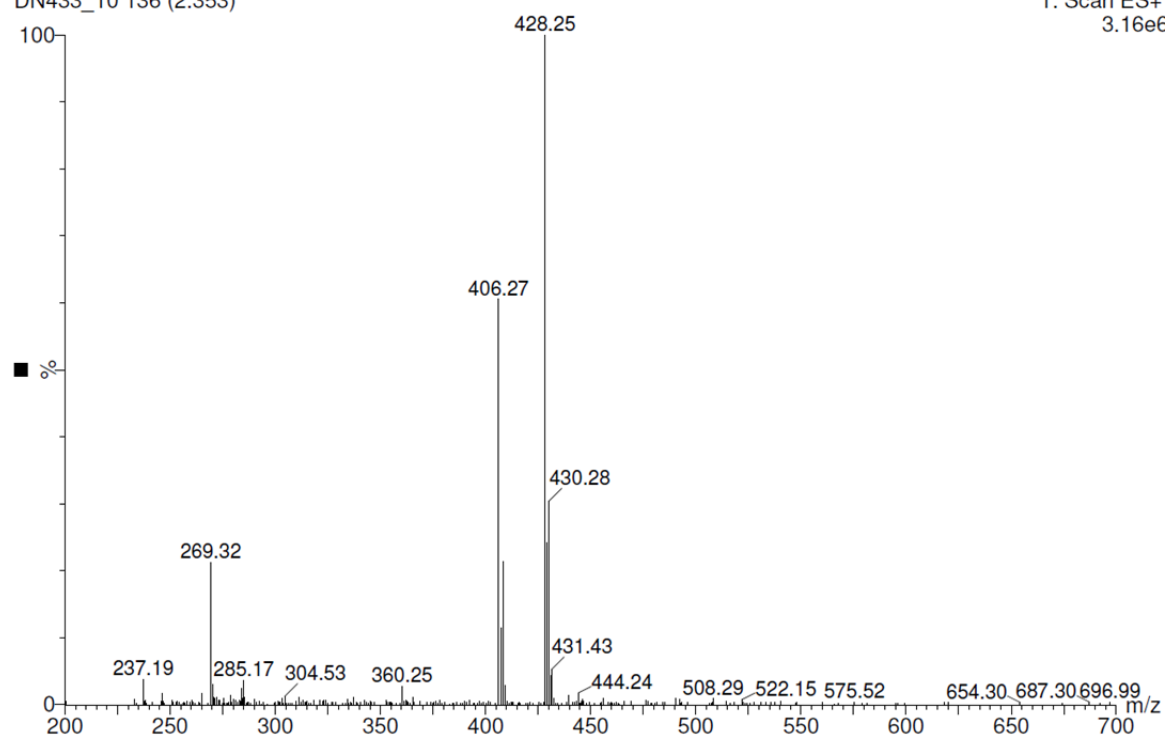


Chromatogram and MS spectra of **14f**, DN433

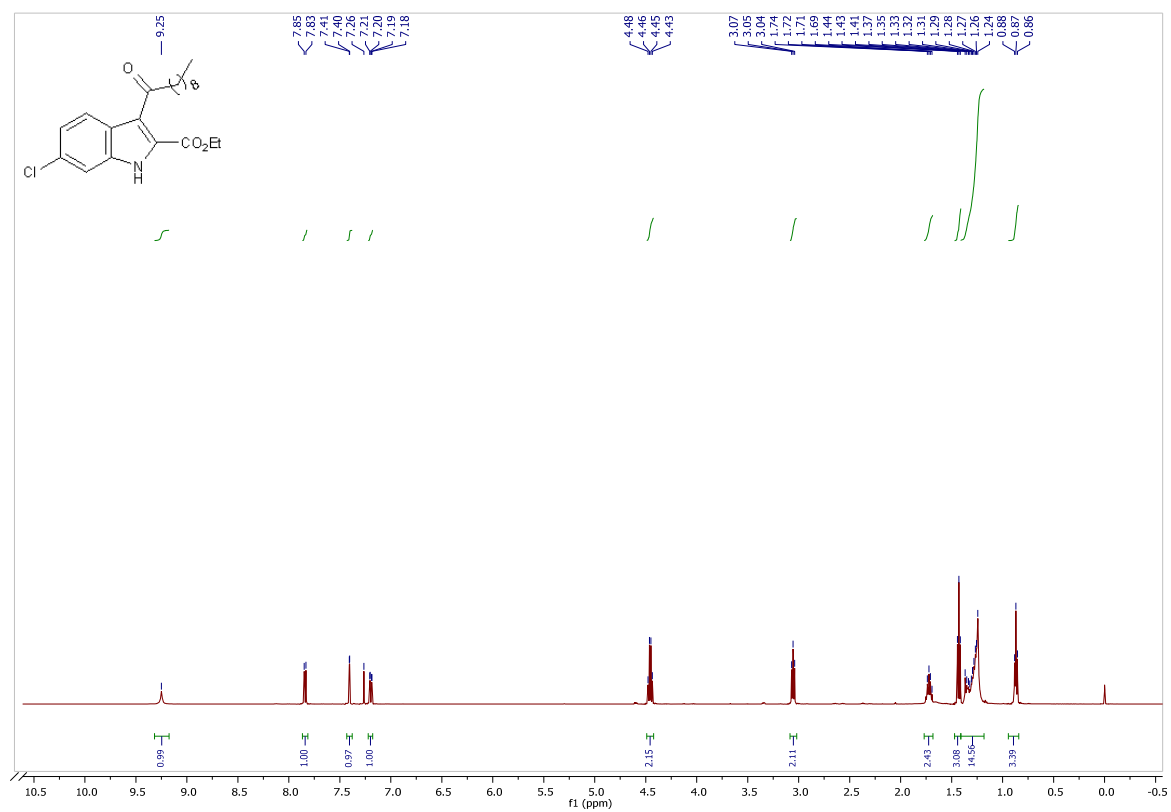
DN433\_10\_Col4\_sol1\_5-30%\_6min  
DN433\_10



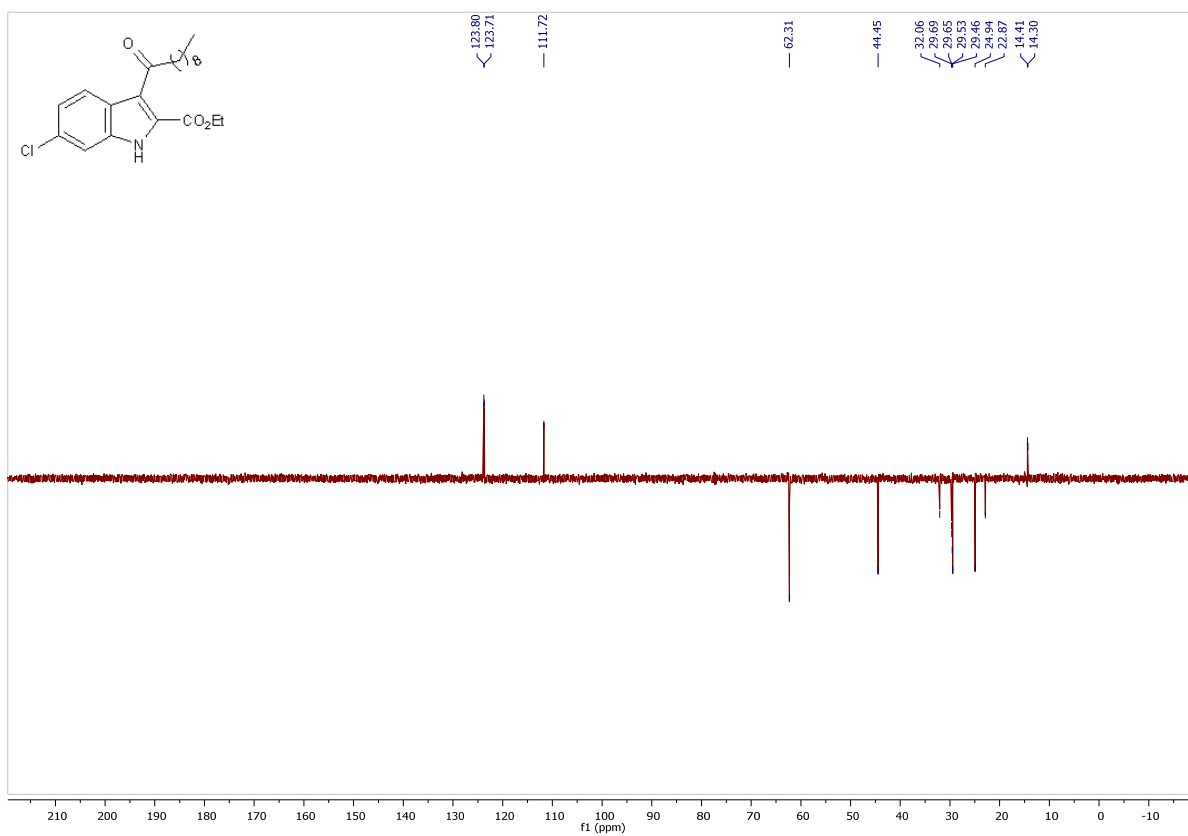
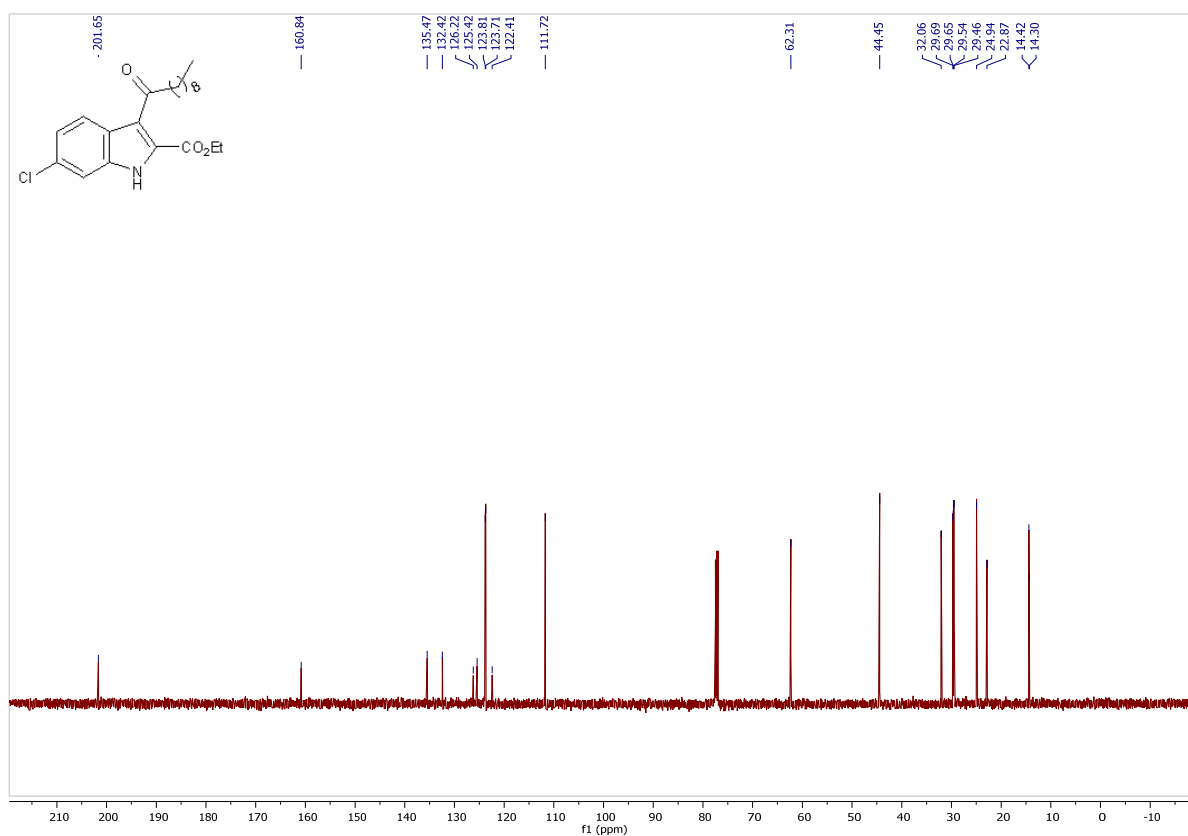
DN433\_10\_Col4\_sol1\_5-30%\_6min  
DN433\_10 136 (2.353)



$^1\text{H}$  NMR spectrum of **14g**, DN432



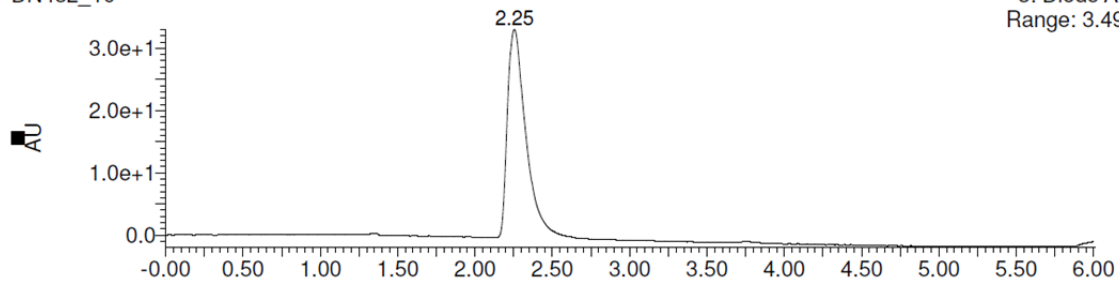
<sup>13</sup>C NMR and DEPT-135 spectra of **14g**, DN432



Chromatogram and MS spectra of **14g**, DN432

DN432\_10\_Col4\_sol1\_5-30%\_6min

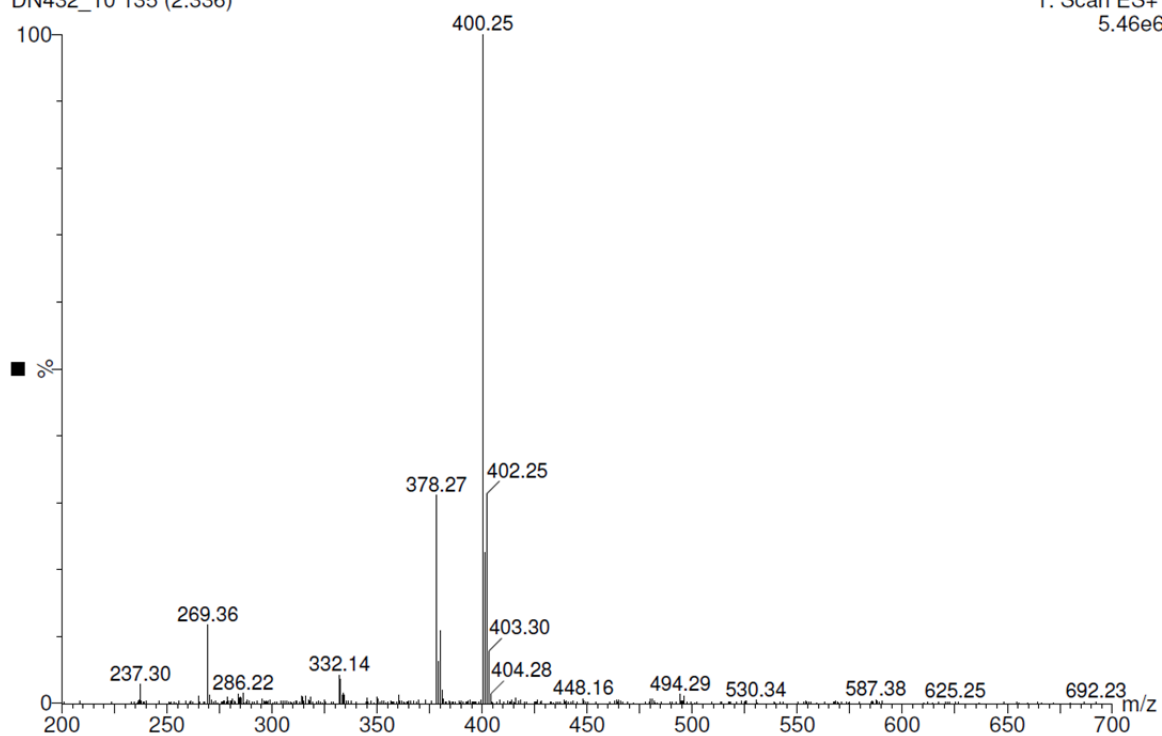
DN432\_10



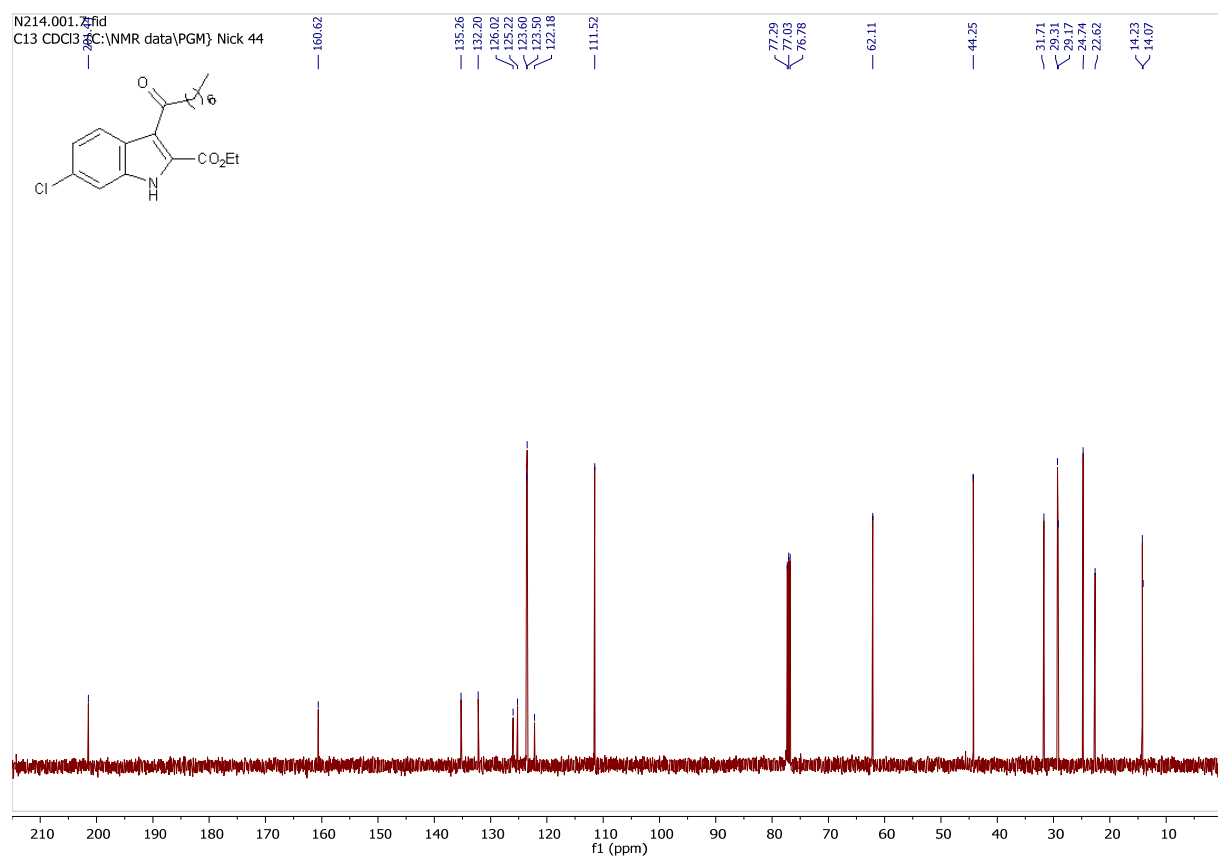
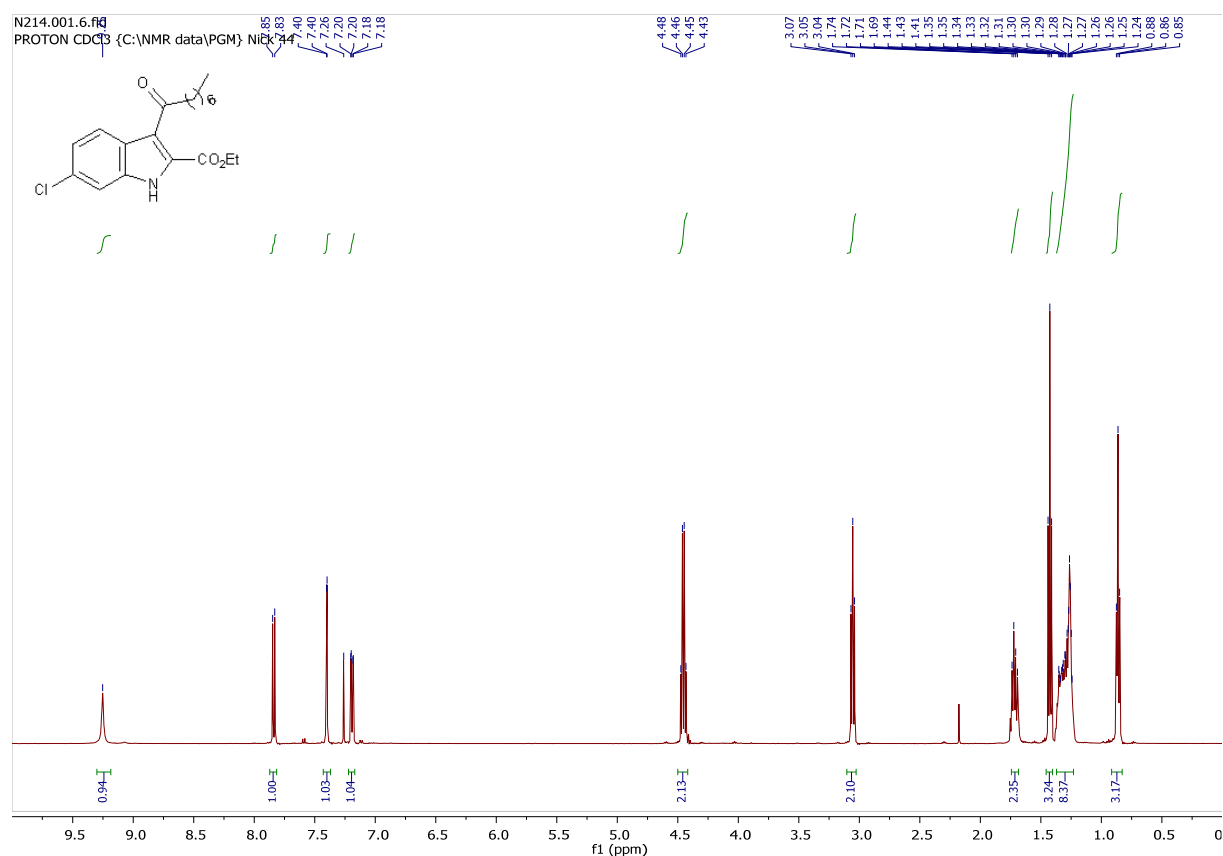
DN432\_10\_Col4\_sol1\_5-30%\_6min

DN432\_10 135 (2.336)

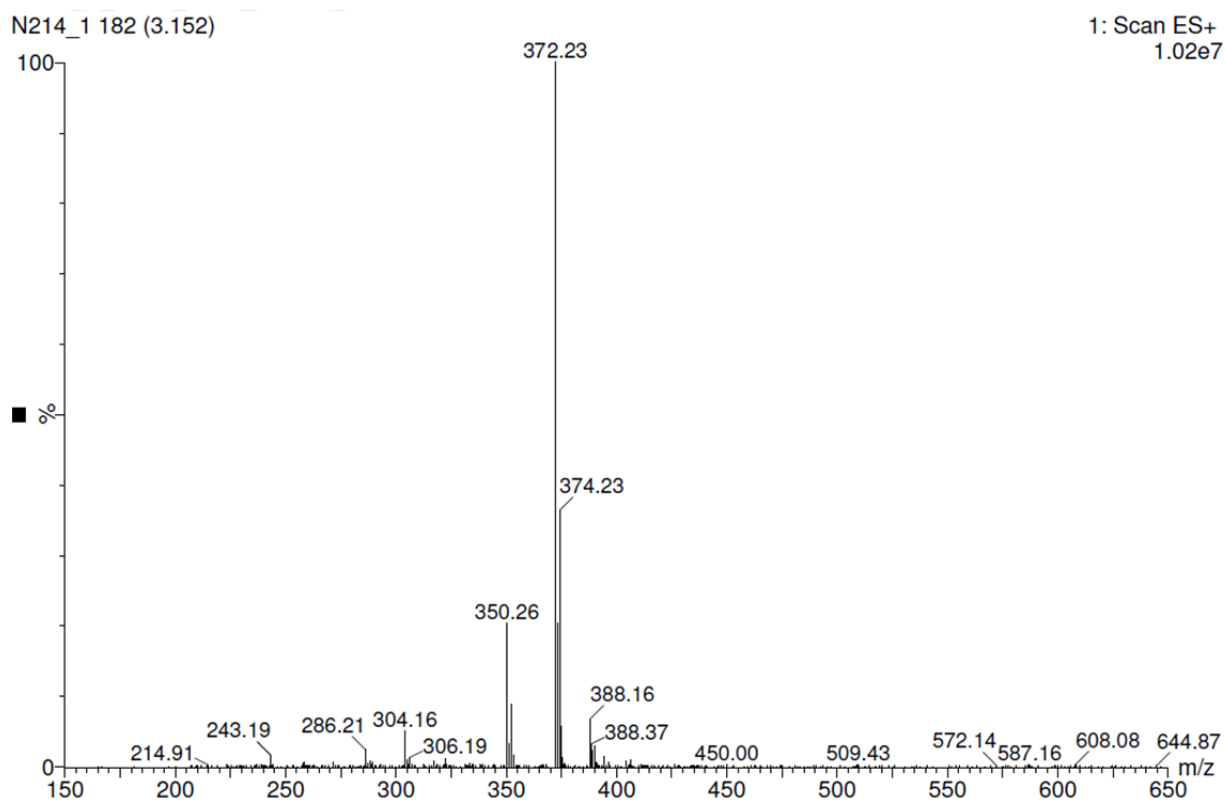
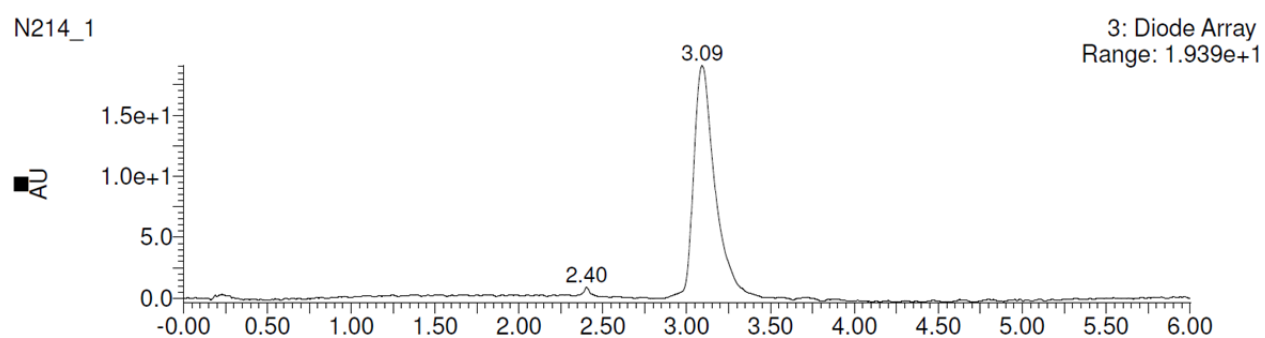
1: Scan ES+  
5.46e6



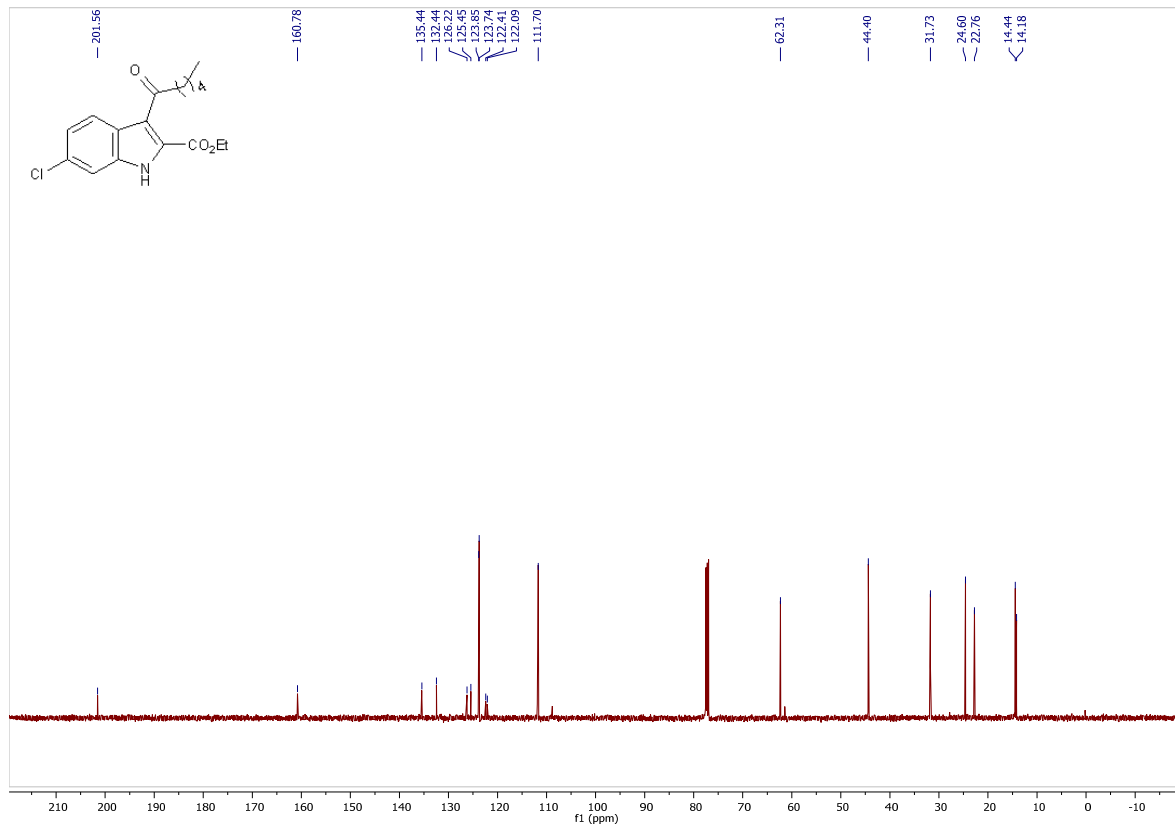
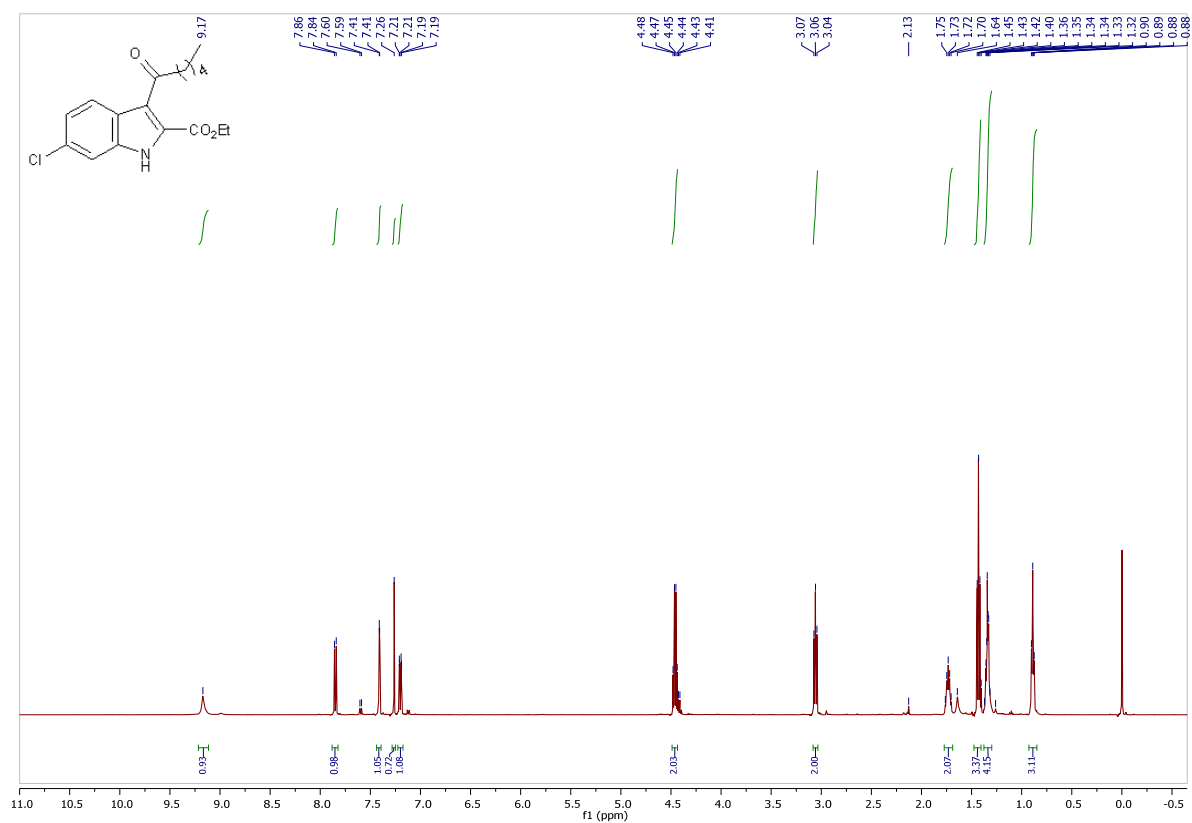
$^1\text{H}$  NMR and  $^{13}\text{C}$  NMR spectra of **14h**, N214



Chromatogram and MS spectra of **14h**, N214



$^1\text{H}$  NMR and  $^{13}\text{C}$  NMR spectra of **14i**, DN441

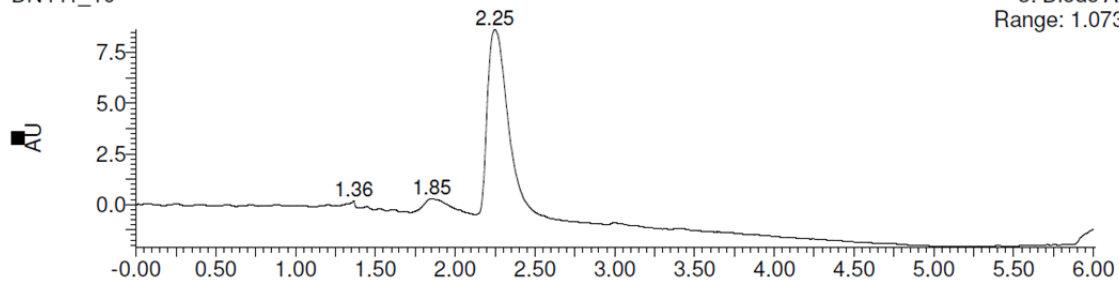


Chromatogram and MS spectra of **14i**, DN441

DN441\_10\_Col4\_sol1\_5-30%\_6min

DN441\_10

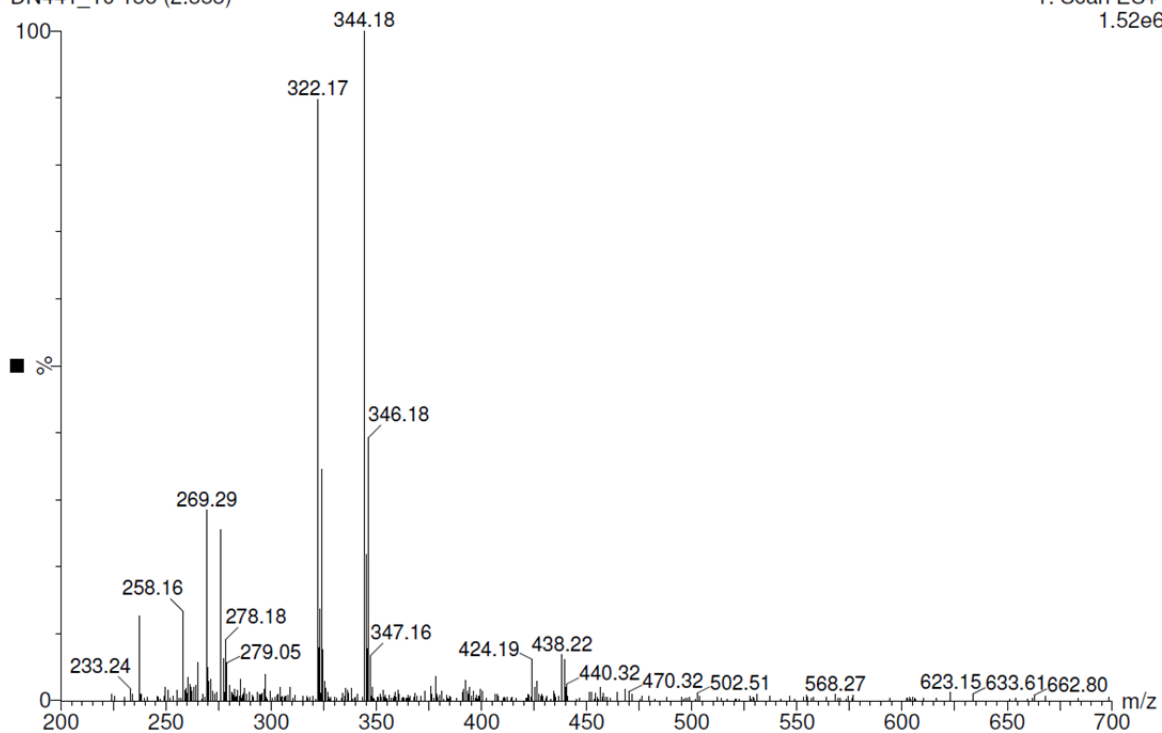
3: Diode Array  
Range: 1.073e+1



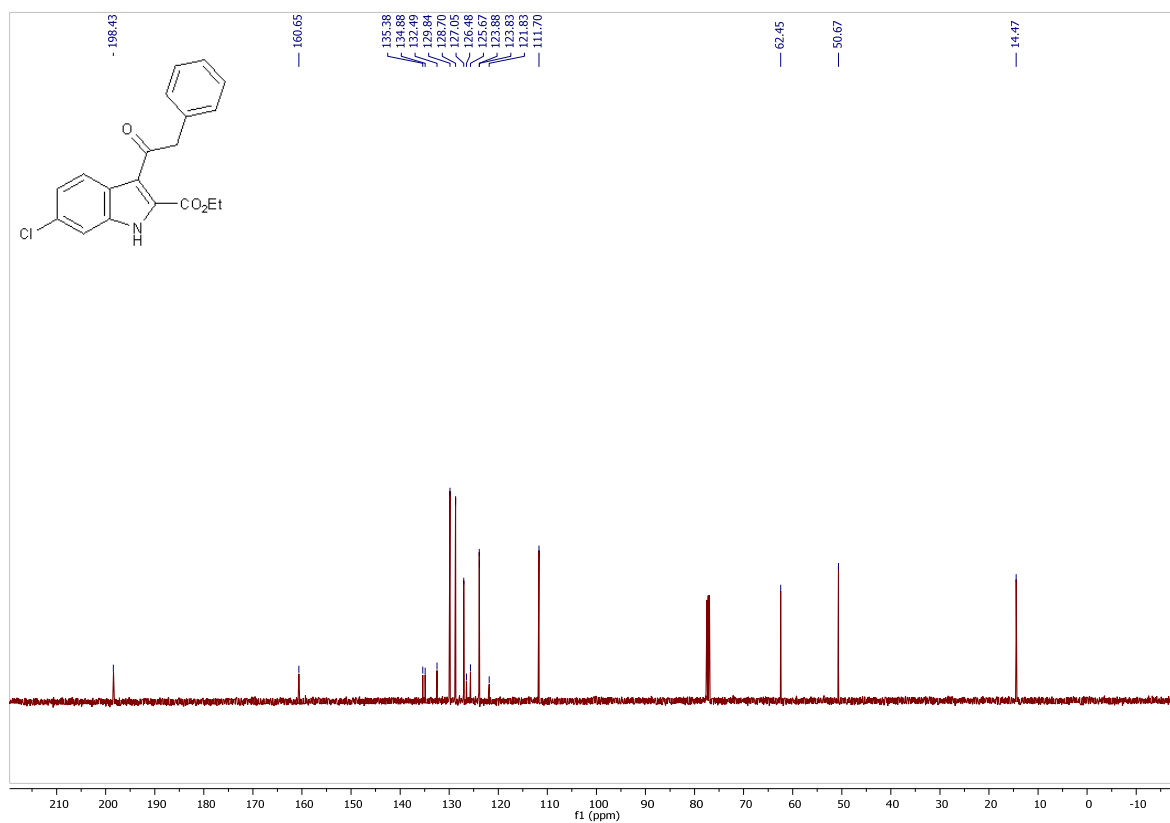
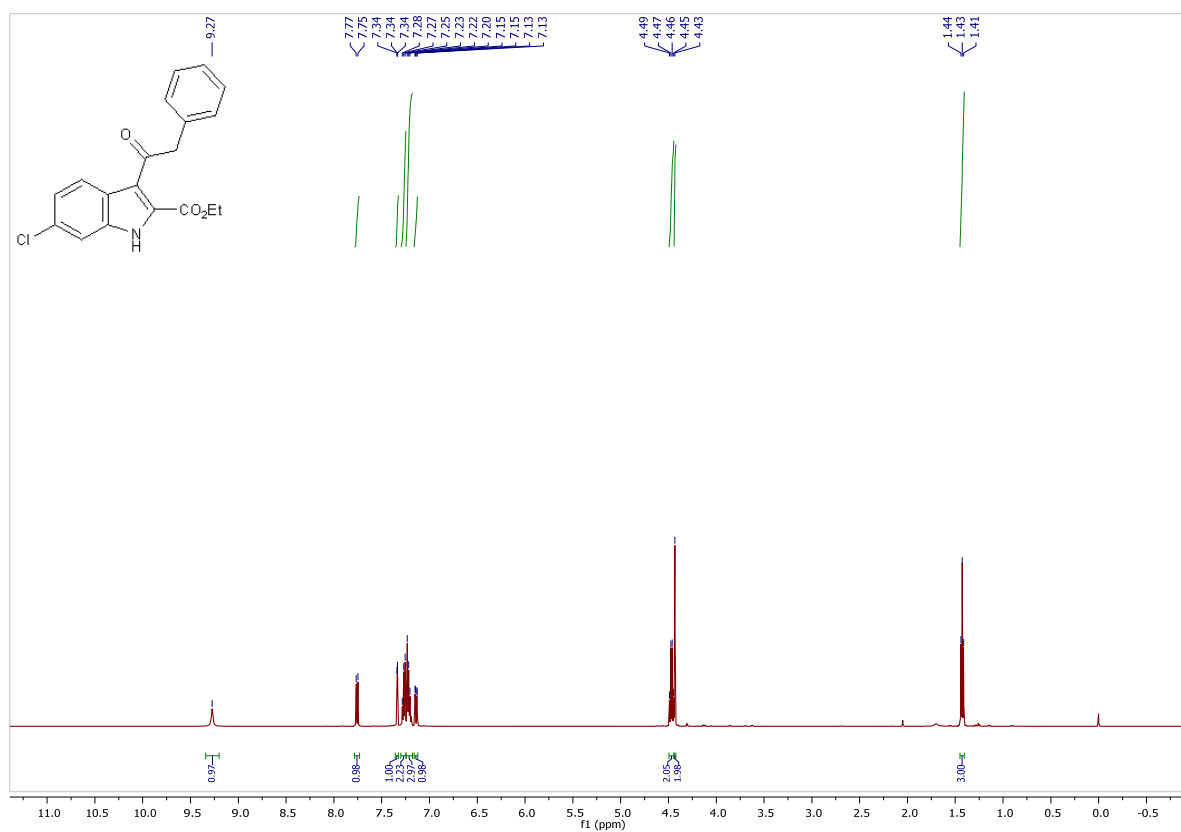
DN441\_10\_Col4\_sol1\_5-30%\_6min

DN441\_10 136 (2.353)

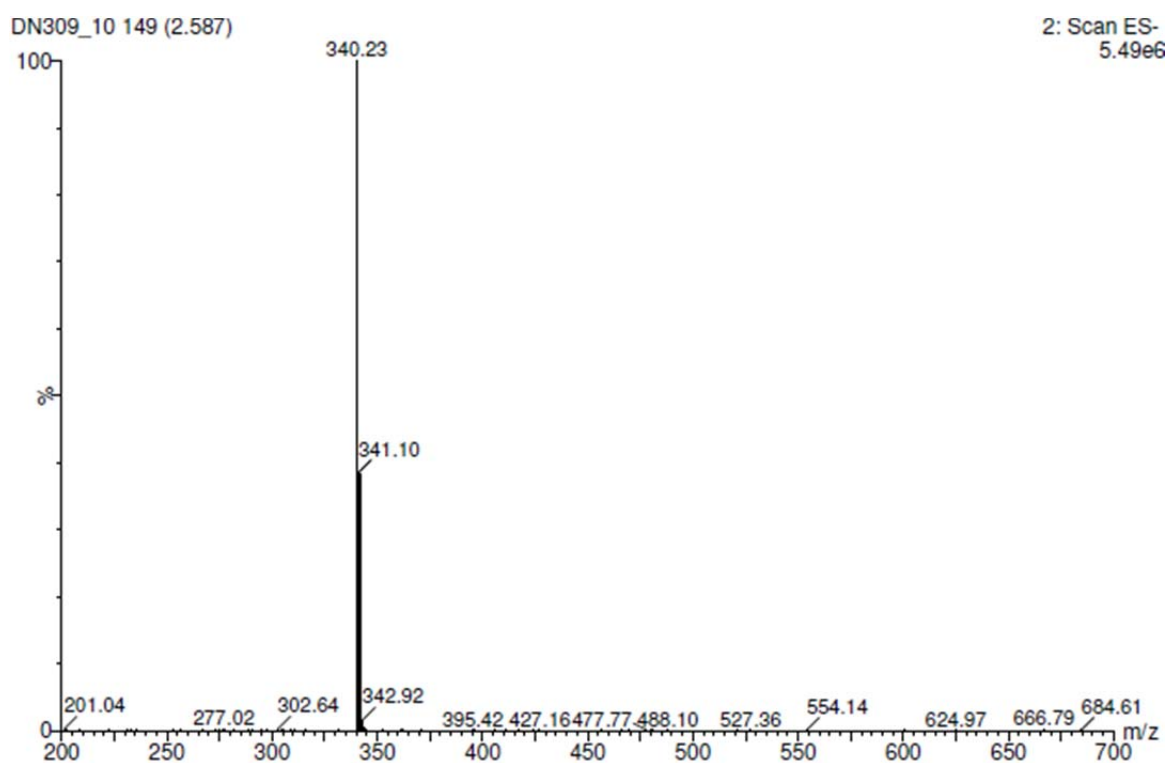
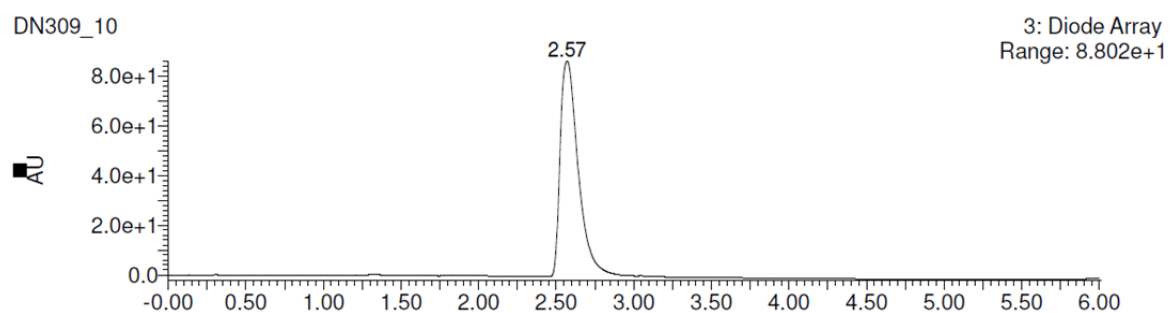
1: Scan ES+  
1.52e6



$^1\text{H}$  NMR and  $^{13}\text{C}$  NMR spectra of **14J**, DN309



Chromatogram and MS spectra of **14J**, DN309

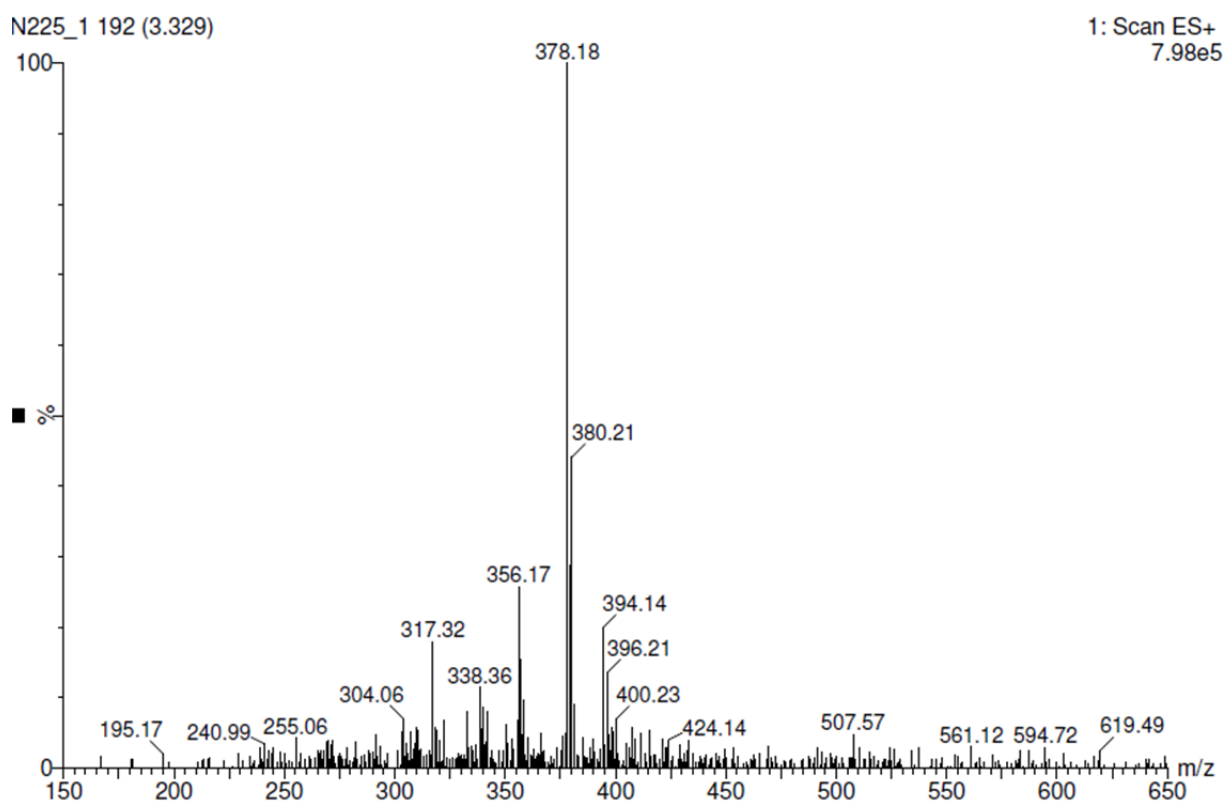
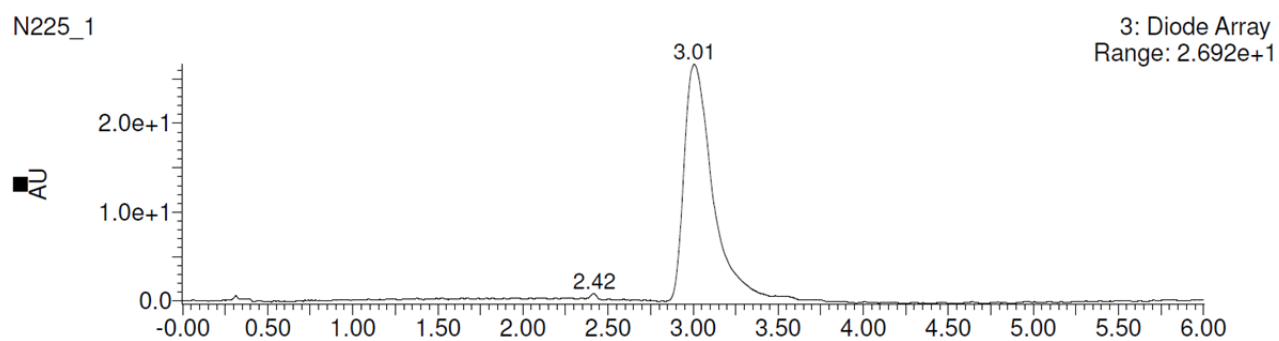


<sup>1</sup>H NMR spectrum (400 MHz, CDCl<sub>3</sub>) of the compound. The chemical structure is shown in the top left corner. The spectrum displays peaks corresponding to the protons in the molecule, with integration values and chemical shifts (ppm) indicated.

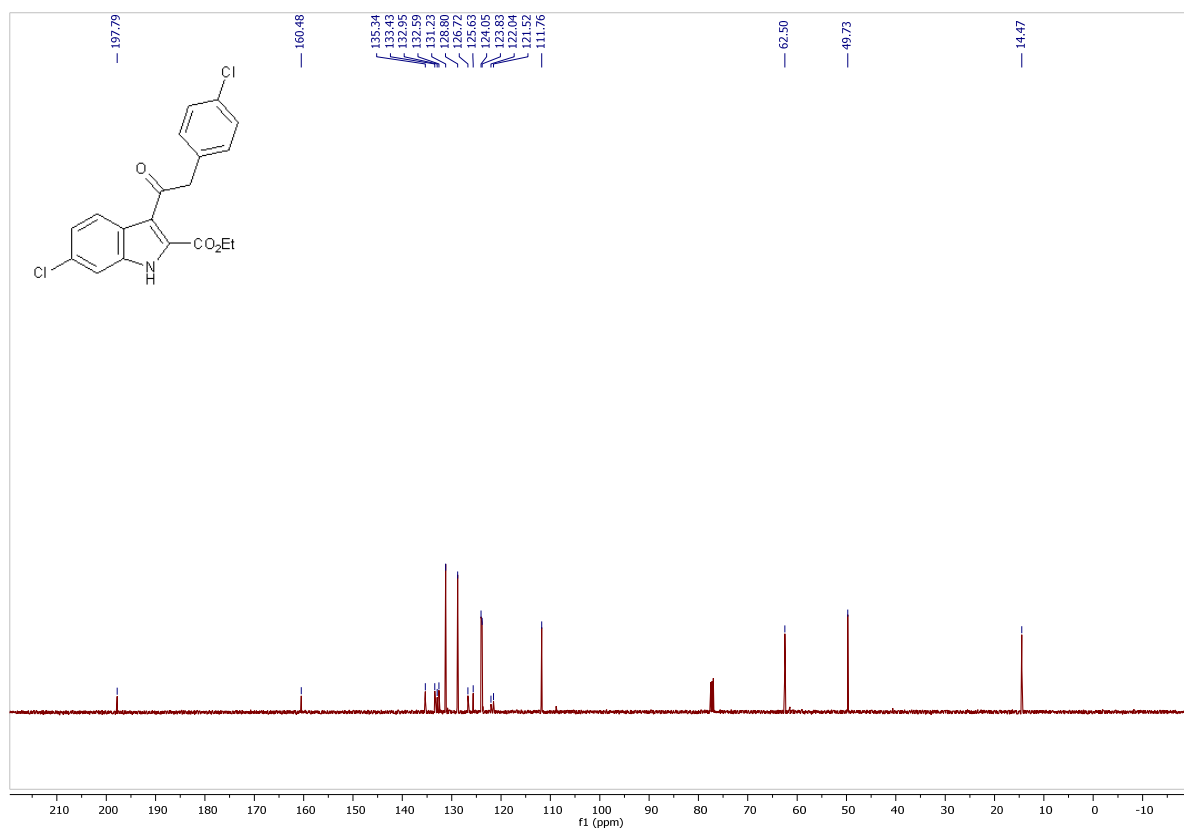
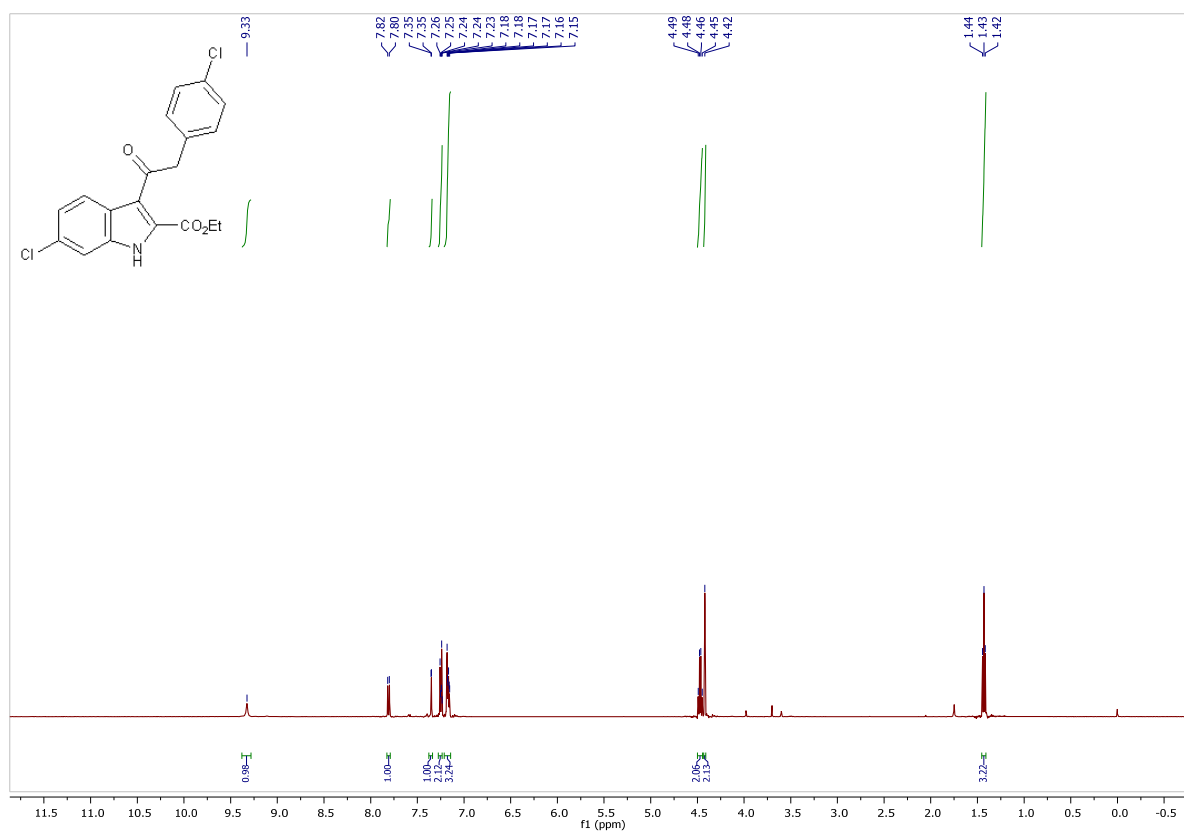
| Chemical Shift (ppm) | Integration |
|----------------------|-------------|
| ~9.1                 | 1.00        |
| ~7.4                 | 1.00        |
| ~7.3                 | 0.94        |
| ~7.2                 | 5.41        |
| ~4.4                 | 2.19        |
| ~3.6                 | 1.90        |
| ~2.3                 | 3.00        |
| ~1.4                 | 3.27        |



Chromatogram and MS spectra of **14k**, N225



$^1\text{H}$  NMR and  $^{13}\text{C}$  NMR spectra of **14I**, DN522

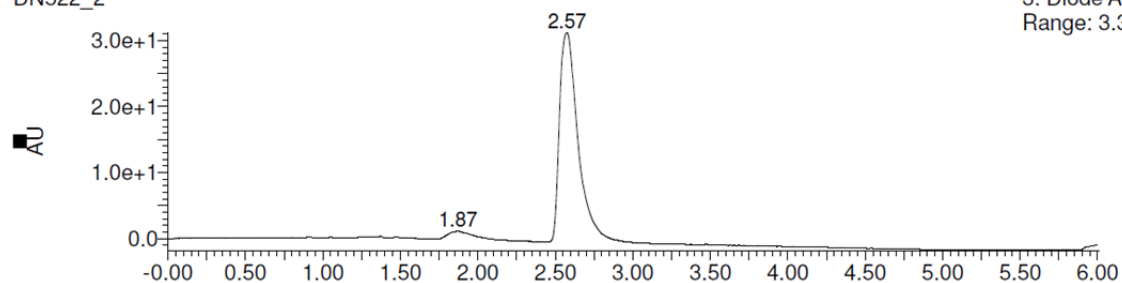


Chromatogram and MS spectra of **14I**, DN522

DN522\_2\_Col4\_sol1\_5-30%\_6min

DN522\_2

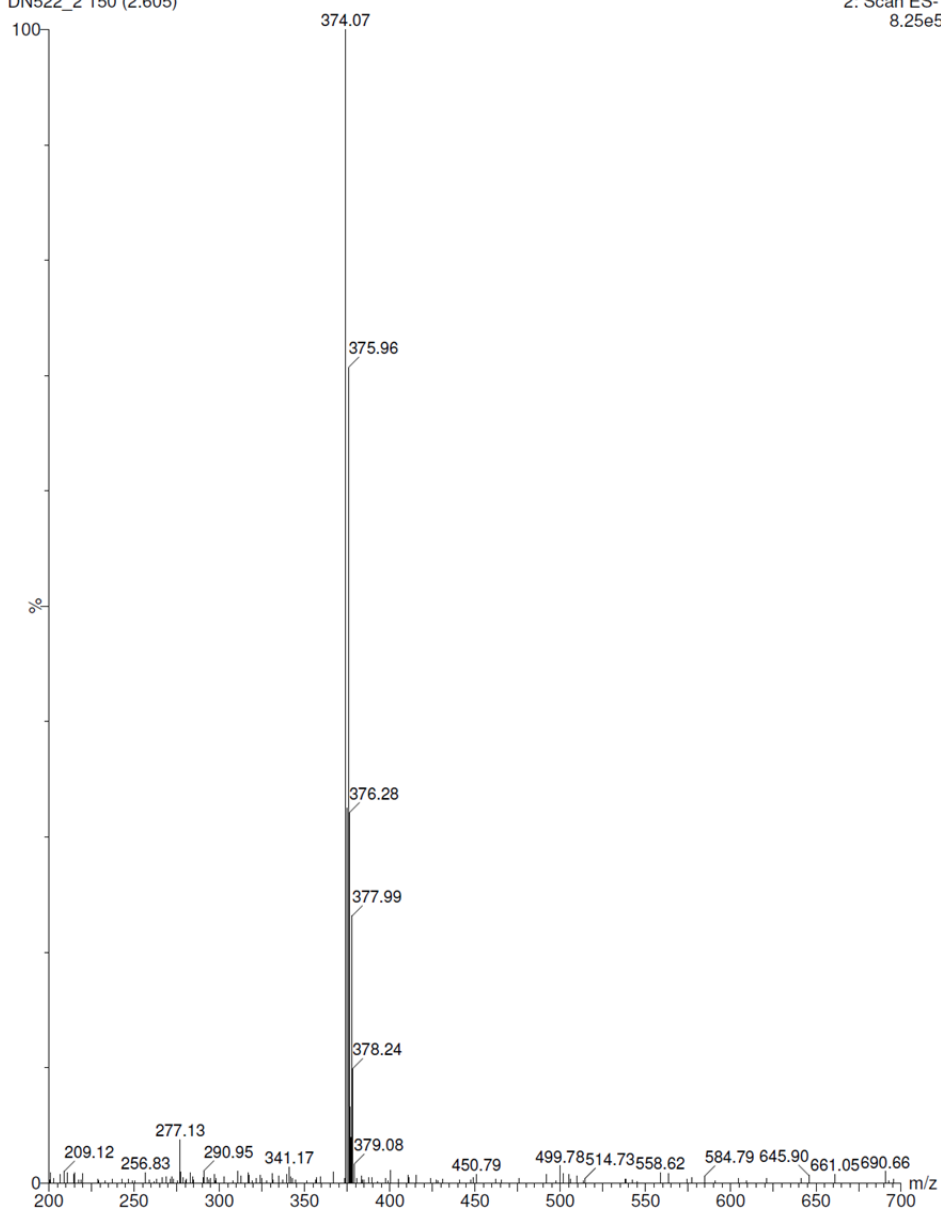
3: Diode Array  
Range: 3.3e+1



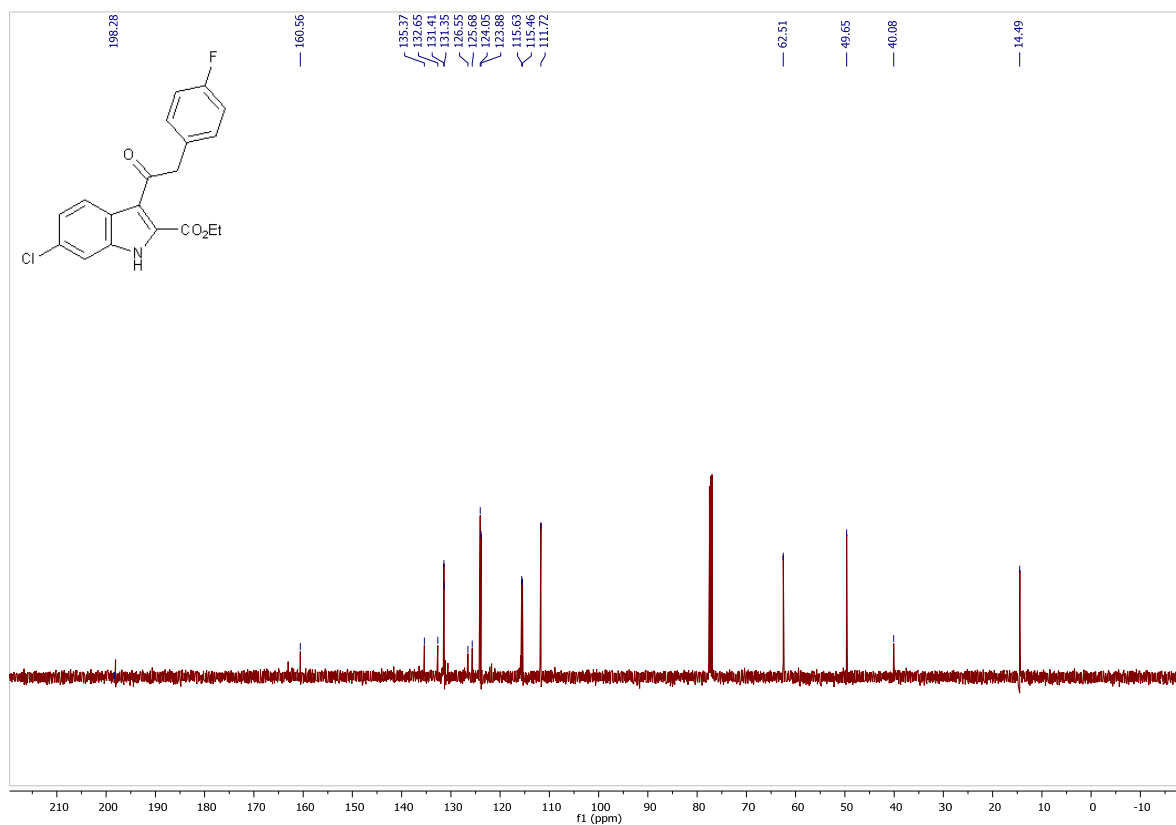
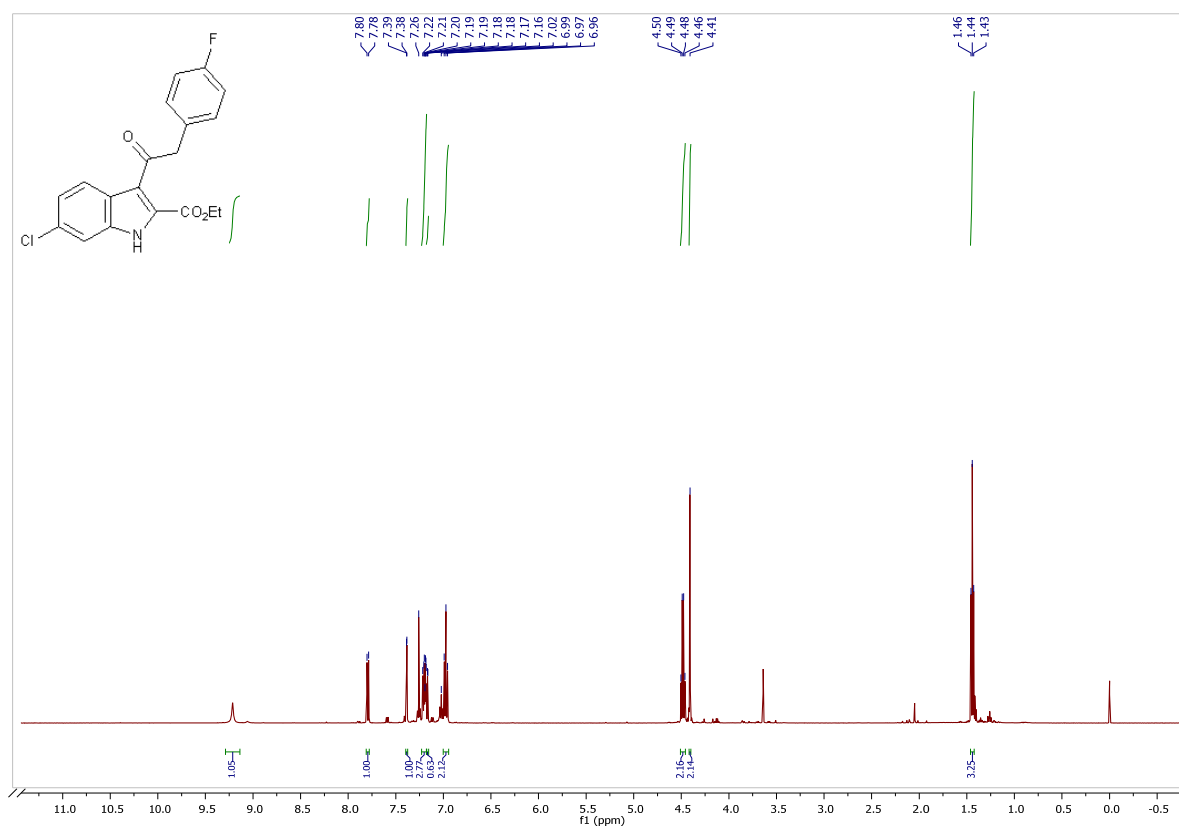
DN522\_2\_Col4\_sol1\_5-30%\_6min

DN522\_2 150 (2.605)

2: Scan ES-  
8.25e5



$^1\text{H}$  NMR and  $^{13}\text{C}$  NMR spectra of **14m**, DN312

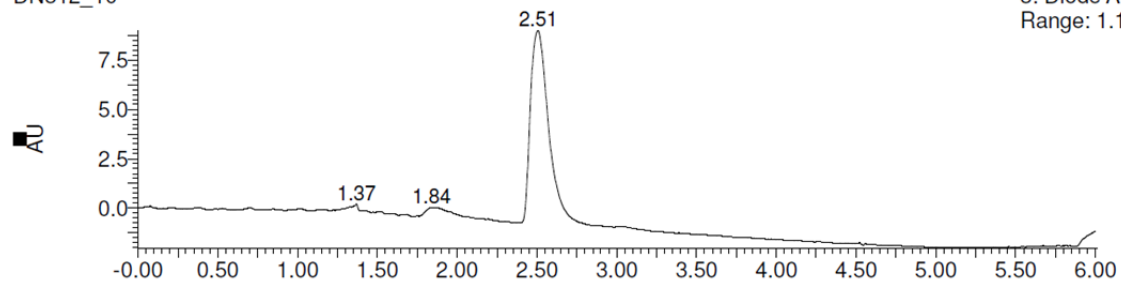


Chromatogram and MS spectra of **14m**, DN312

DN312\_10\_Col4\_sol1\_5-30%\_6min

DN312\_10

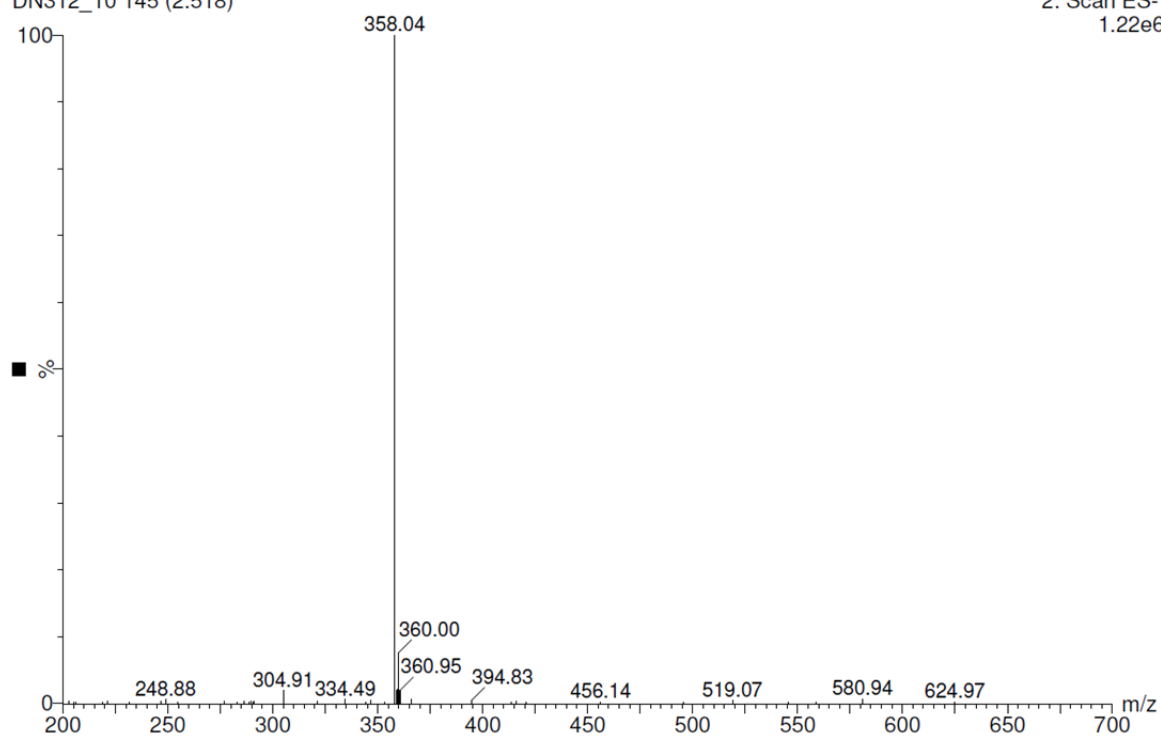
3: Diode Array  
Range: 1.1e+1



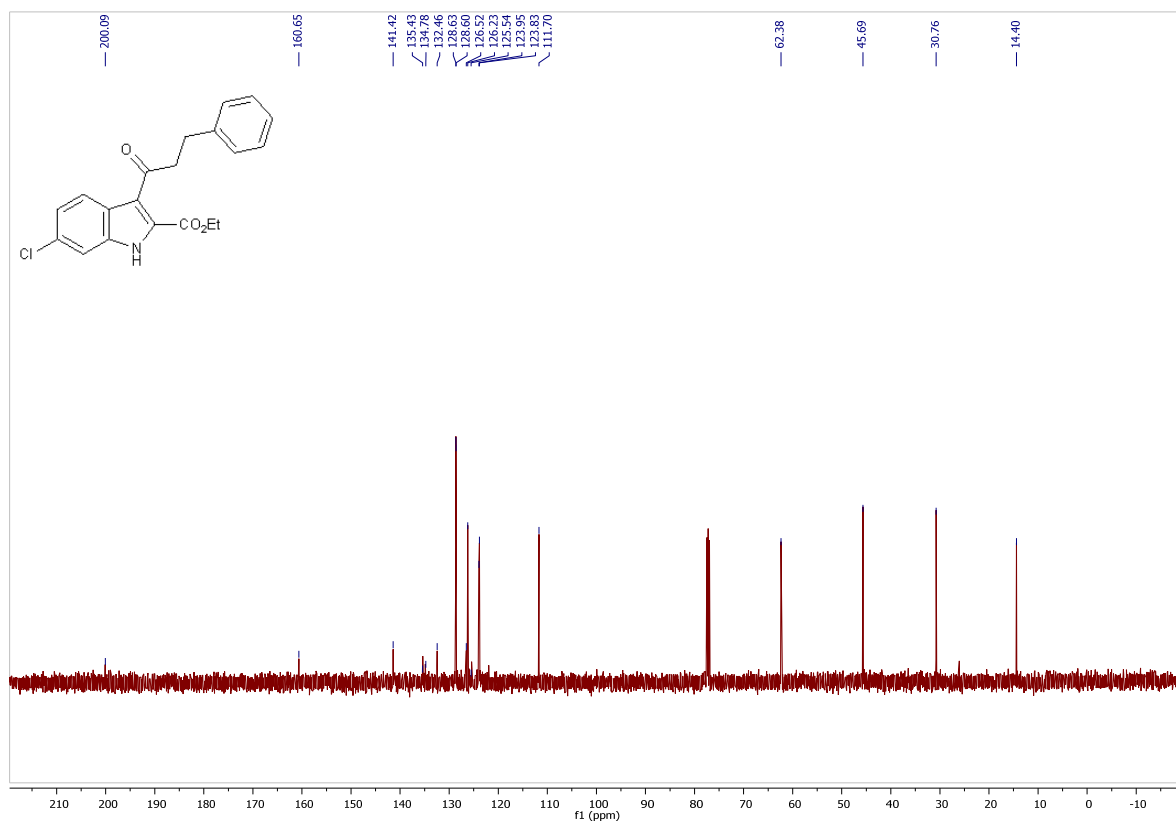
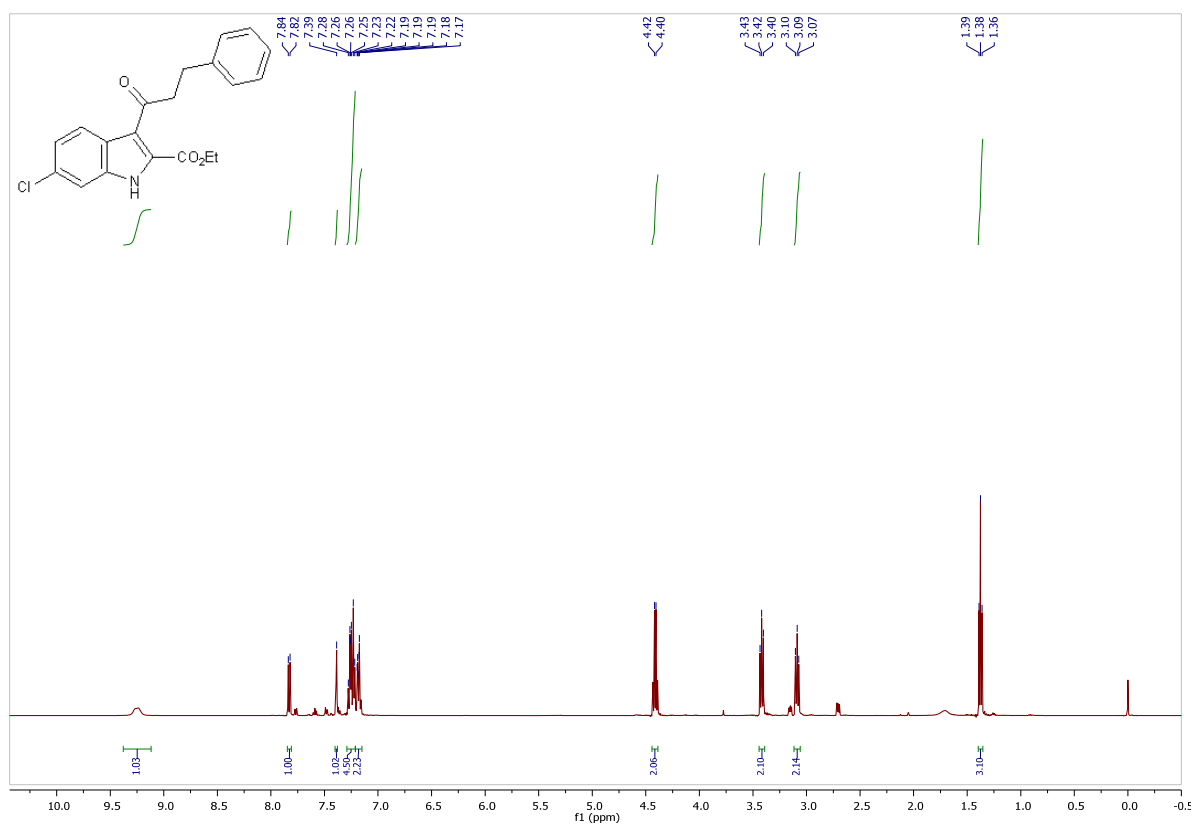
DN312\_10\_Col4\_sol1\_5-30%\_6min

DN312\_10 145 (2.518)

2: Scan ES-  
1.22e6



$^1\text{H}$  NMR and  $^{13}\text{C}$  NMR spectra of **14n**, DN427

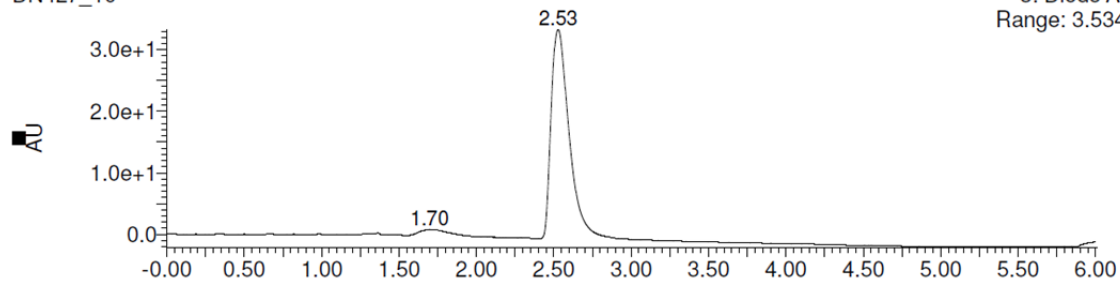


Chromatogram and MS spectra of **14n**, DN427

DN427\_10\_Col4\_sol1\_5-30%\_6min

DN427\_10

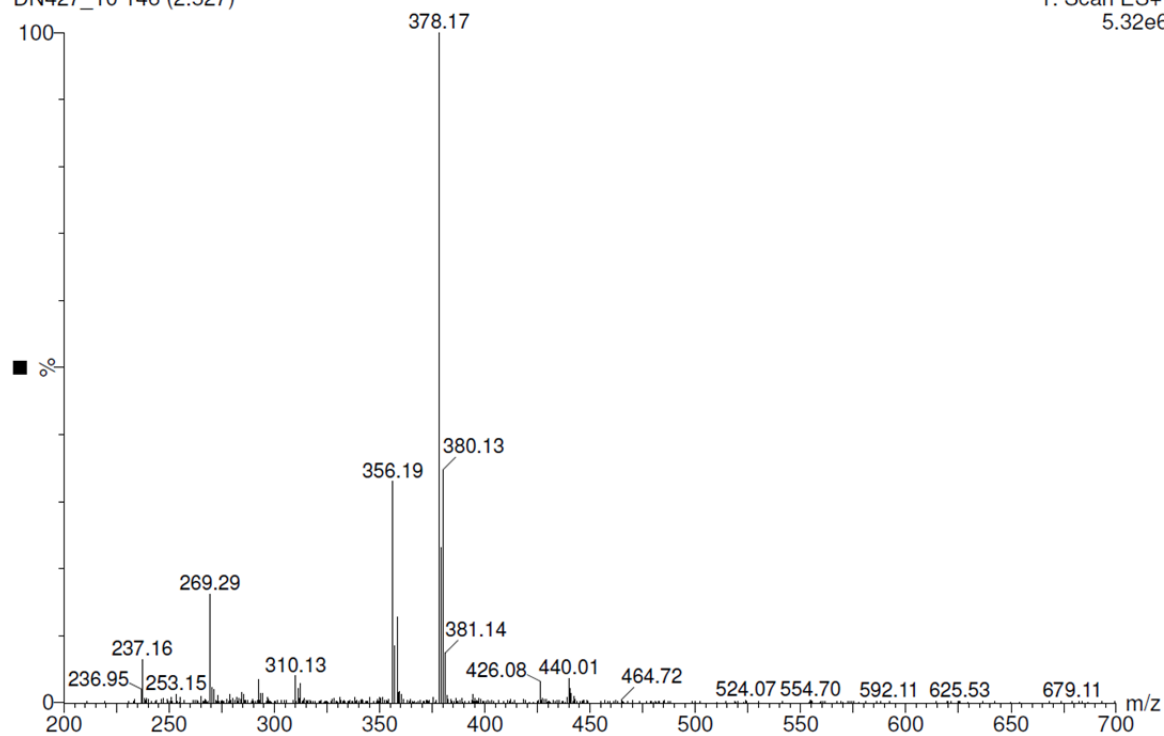
3: Diode Array  
Range: 3.534e+1



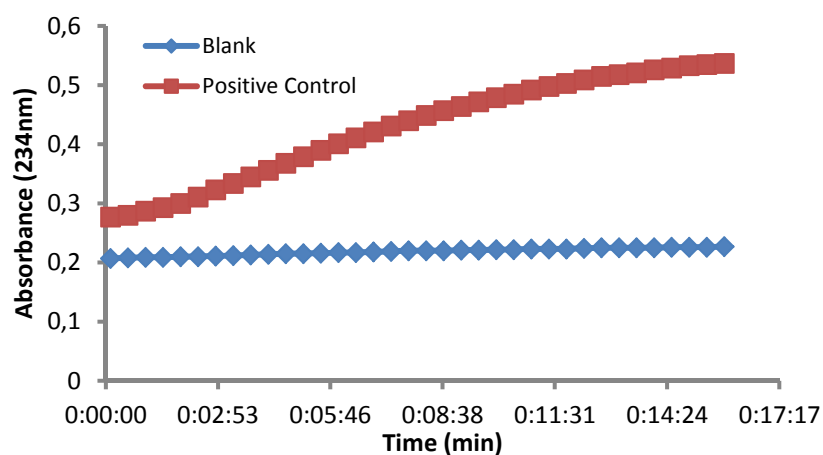
DN427\_10\_Col4\_sol1\_5-30%\_6min

DN427\_10 146 (2.527)

1: Scan ES+  
5.32e6



## Enzyme inhibition studies



**Figure S1:** Conversion of linoleic acid in present (Positive Control) or absent of the enzyme (Blank) following with UV-absorbance assay at 234 nm over time.

## Focused library

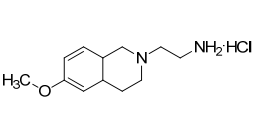
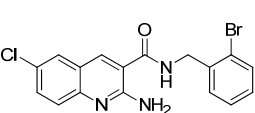
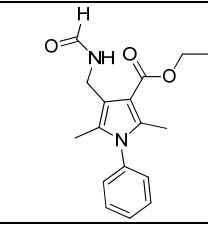
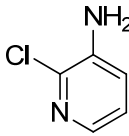
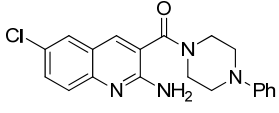
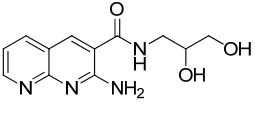
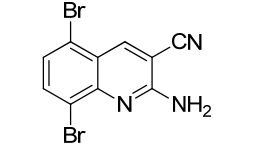
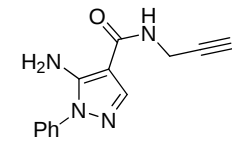
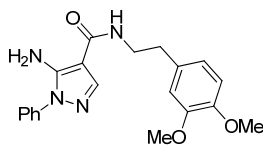
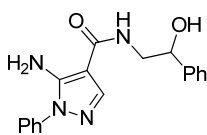
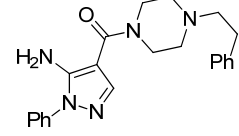
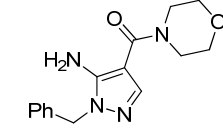
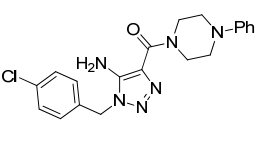
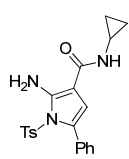
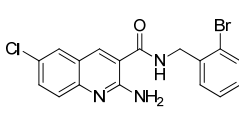
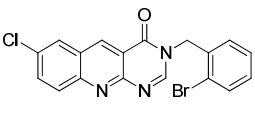
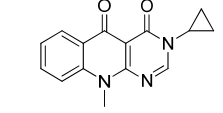
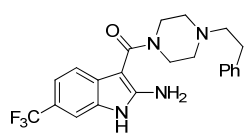
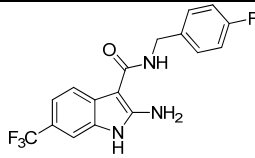
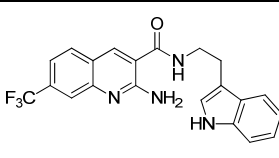
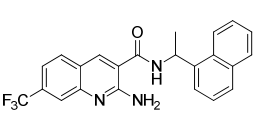
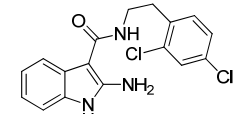
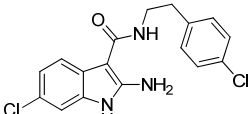
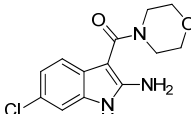
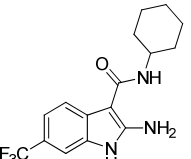
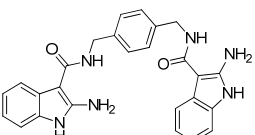
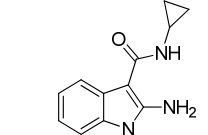
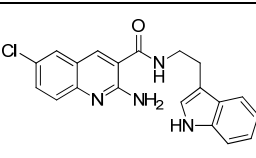
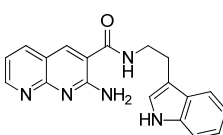
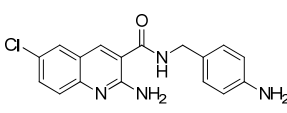
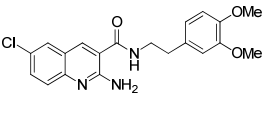
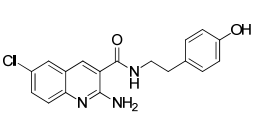
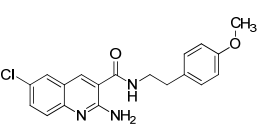
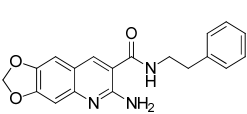
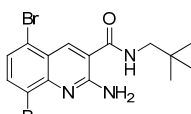
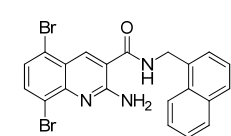
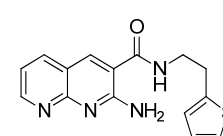
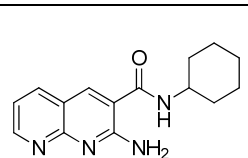
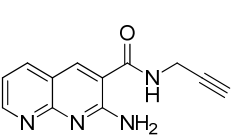
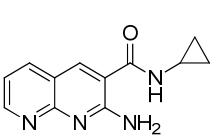
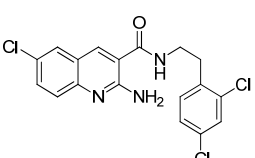
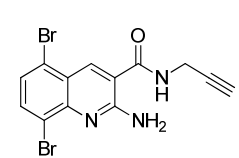
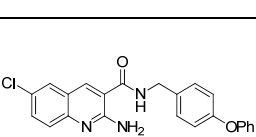
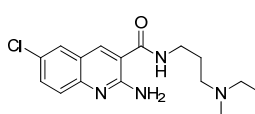
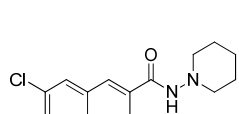
Library of heterocycle nitrogen containing compounds (200). The name of every compound corresponds with the position of the compound in the next table.

**Table S1. Screening Compound**

|   | A | B | C | D | E |
|---|---|---|---|---|---|
| 1 |   |   |   |   |   |
| 2 |   |   |   |   |   |
| 3 |   |   |   |   |   |
| 4 |   |   |   |   |   |

|    |  |  |  |  |  |
|----|--|--|--|--|--|
| 5  |  |  |  |  |  |
| 6  |  |  |  |  |  |
| 7  |  |  |  |  |  |
| 8  |  |  |  |  |  |
| 9  |  |  |  |  |  |
| 10 |  |  |  |  |  |
| 11 |  |  |  |  |  |
| 12 |  |  |  |  |  |
| 13 |  |  |  |  |  |

|    |  |  |  |  |  |
|----|--|--|--|--|--|
| 14 |  |  |  |  |  |
| 15 |  |  |  |  |  |
| 16 |  |  |  |  |  |
| 17 |  |  |  |  |  |
| 18 |  |  |  |  |  |
| 19 |  |  |  |  |  |
| 20 |  |  |  |  |  |
| 21 |  |  |  |  |  |
| 22 |  |  |  |  |  |

|    |                                                                                     |                                                                                     |                                                                                     |                                                                                       |                                                                                       |
|----|-------------------------------------------------------------------------------------|-------------------------------------------------------------------------------------|-------------------------------------------------------------------------------------|---------------------------------------------------------------------------------------|---------------------------------------------------------------------------------------|
| 23 |    |    |    |    |    |
| 24 |    |    |    |     |    |
| 25 |    |    |    |    |    |
| 26 |    |    |    |     |    |
| 27 |  |  |  |  |   |
| 28 |  |  |  |  |  |
| 29 |  |  |  |  |  |
| 30 |  |  |  |  |  |
| 31 |  |  |  |   |  |

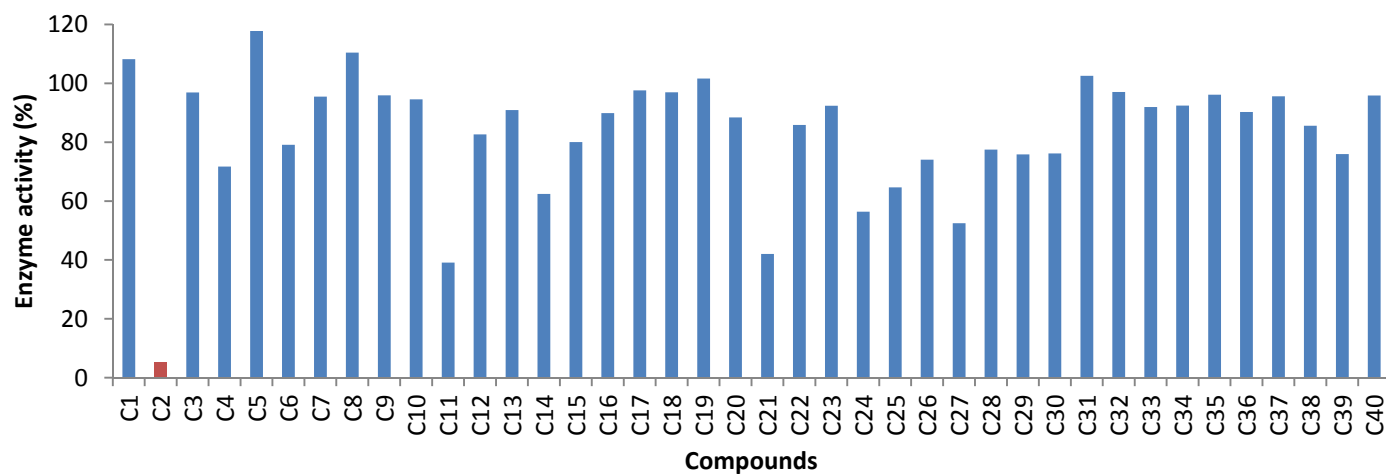
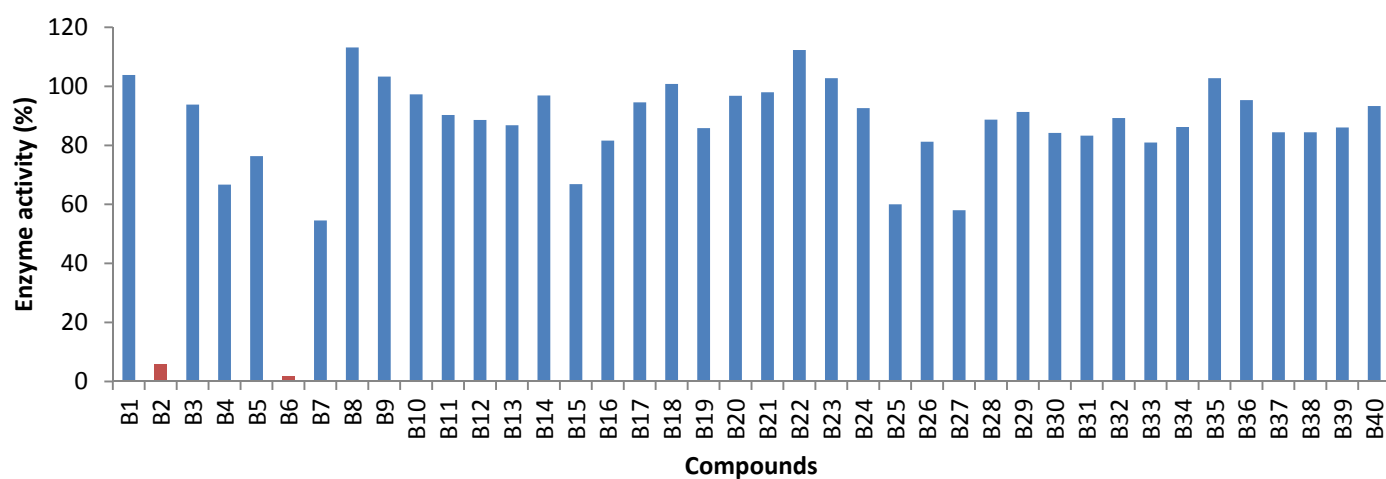
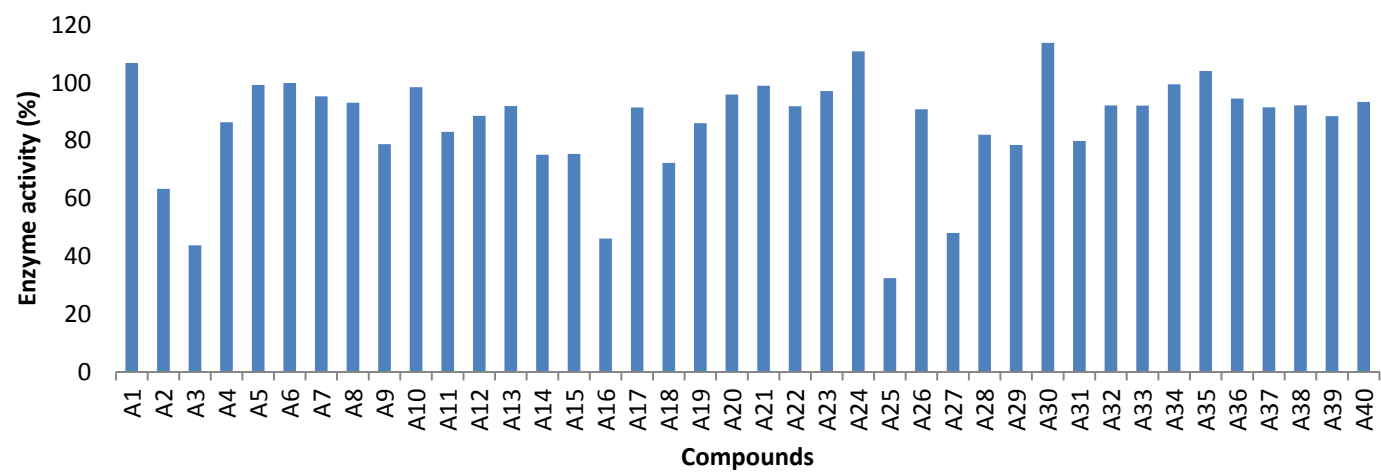
|    |  |  |  |  |  |
|----|--|--|--|--|--|
| 32 |  |  |  |  |  |
| 33 |  |  |  |  |  |
| 34 |  |  |  |  |  |
| 35 |  |  |  |  |  |
| 36 |  |  |  |  |  |
| 37 |  |  |  |  |  |
| 38 |  |  |  |  |  |
| 39 |  |  |  |  |  |
| 40 |  |  |  |  |  |

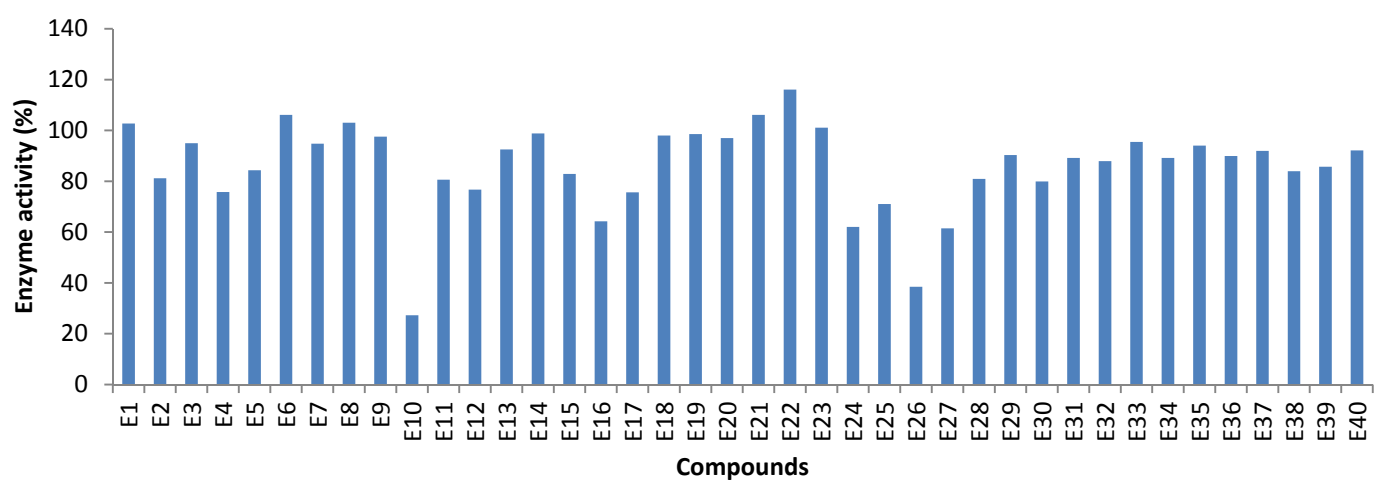
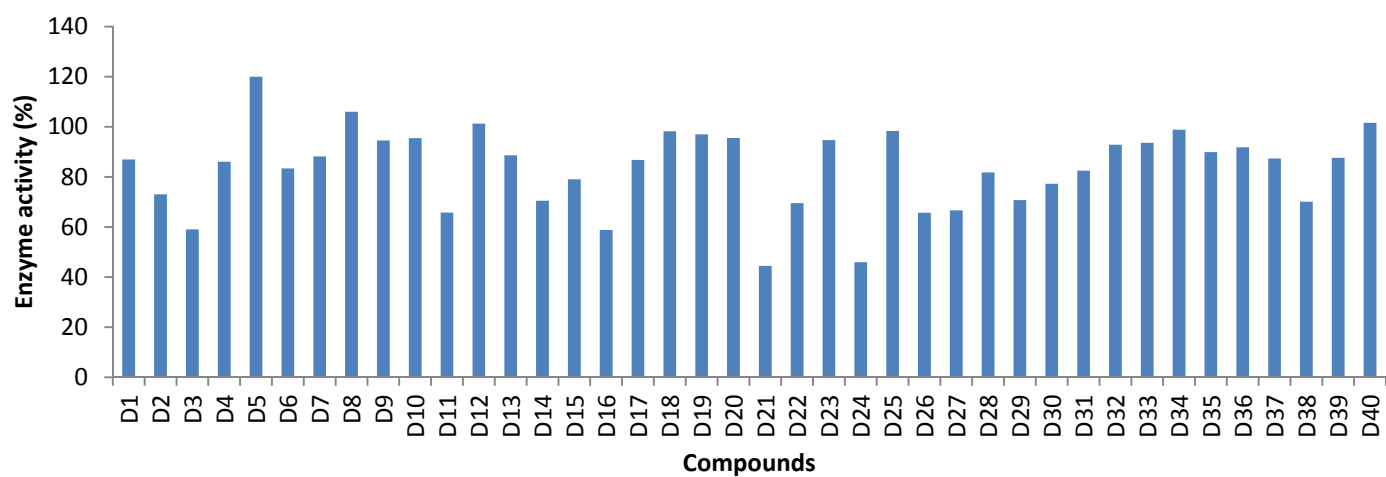
## Screening Results

**Table S2. Residual enzyme activity (%) that was observed for the screening of a compounds collection for inhibition of h-15-LOX-1 in presence of 50  $\mu$ M of the respective compounds.**

|    | A     | B     | C     | D     | E     |
|----|-------|-------|-------|-------|-------|
| 1  | 106,8 | 103,8 | 108,2 | 86,96 | 102,7 |
| 2  | 63,33 | 6,060 | 5,285 | 73,01 | 81,14 |
| 3  | 43,78 | 93,78 | 96,89 | 59,06 | 94,94 |
| 4  | 86,36 | 66,66 | 71,71 | 86,04 | 75,75 |
| 5  | 99,27 | 76,36 | 117,7 | 120,0 | 84,33 |
| 6  | 99,95 | 2,020 | 79,09 | 83,38 | 106,1 |
| 7  | 95,34 | 54,54 | 95,45 | 88,18 | 94,79 |
| 8  | 93,15 | 113,1 | 110,4 | 106,0 | 103,0 |
| 9  | 78,75 | 103,2 | 95,89 | 94,54 | 97,57 |
| 10 | 98,48 | 97,27 | 94,54 | 95,45 | 27,27 |
| 11 | 83,03 | 90,30 | 39,09 | 65,75 | 80,60 |
| 12 | 88,60 | 88,60 | 82,64 | 101,2 | 76,68 |
| 13 | 91,96 | 86,78 | 90,93 | 88,60 | 92,48 |
| 14 | 75,12 | 96,89 | 62,43 | 70,46 | 98,78 |
| 15 | 75,38 | 66,83 | 80,05 | 79,01 | 82,90 |
| 16 | 46,11 | 81,60 | 89,89 | 58,80 | 64,24 |
| 17 | 91,45 | 94,55 | 97,57 | 86,78 | 75,64 |
| 18 | 72,27 | 100,8 | 96,96 | 98,18 | 97,97 |
| 19 | 86,06 | 85,85 | 101,6 | 96,96 | 98,58 |
| 20 | 95,95 | 96,76 | 88,41 | 95,54 | 96,96 |
| 21 | 98,98 | 97,97 | 42,02 | 44,44 | 106,1 |
| 22 | 91,91 | 112,2 | 85,85 | 69,49 | 116,0 |
| 23 | 97,16 | 102,7 | 92,35 | 94,70 | 101,0 |
| 24 | 110,9 | 92,61 | 56,37 | 45,95 | 62,02 |
| 25 | 32,44 | 60,04 | 64,62 | 98,35 | 71,02 |
| 26 | 90,84 | 81,25 | 74,06 | 65,65 | 38,45 |
| 27 | 48,04 | 57,99 | 52,46 | 66,64 | 61,47 |
| 28 | 82,04 | 88,71 | 77,46 | 81,76 | 80,93 |
| 29 | 78,52 | 91,32 | 75,84 | 70,75 | 90,33 |
| 30 | 113,8 | 84,21 | 76,19 | 77,26 | 79,91 |
| 31 | 79,87 | 83,26 | 102,5 | 82,47 | 89,19 |
| 32 | 92,19 | 89,23 | 97,04 | 92,86 | 87,92 |
| 33 | 92,11 | 80,95 | 91,95 | 93,58 | 95,46 |
| 34 | 99,50 | 86,19 | 92,44 | 98,85 | 89,16 |
| 35 | 104,1 | 102,7 | 96,14 | 89,88 | 94,03 |
| 36 | 94,56 | 95,31 | 90,26 | 91,80 | 89,92 |
| 37 | 91,54 | 84,41 | 95,61 | 87,32 | 91,95 |
| 38 | 92,22 | 84,42 | 85,58 | 70,09 | 83,96 |
| 39 | 88,49 | 86,07 | 75,97 | 87,62 | 85,70 |
| 40 | 93,39 | 93,31 | 95,83 | 101,5 | 92,16 |

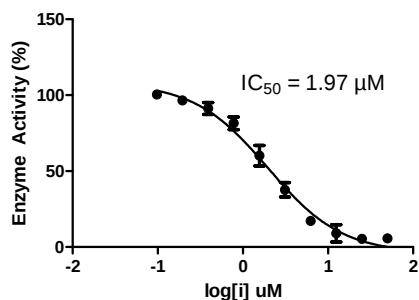
## Screening Results (column graphs)



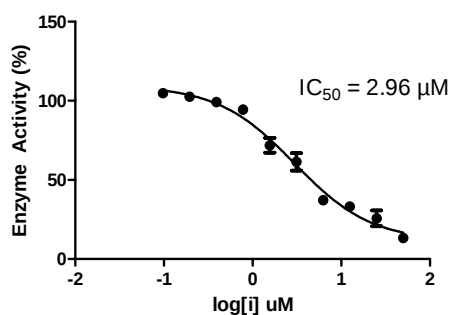


## IC<sub>50</sub> graphs

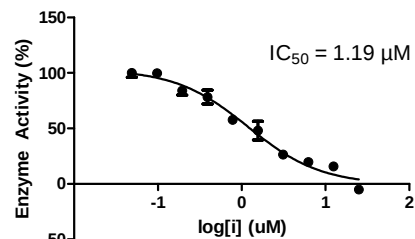
Compound 1



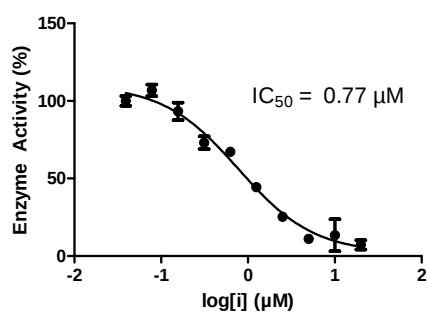
Compound 2



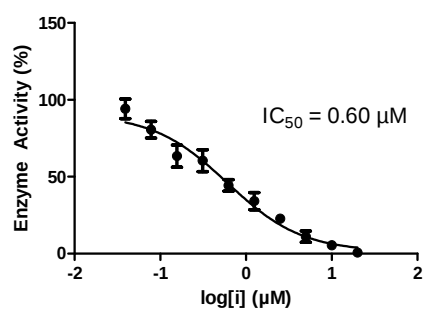
Compound 14l



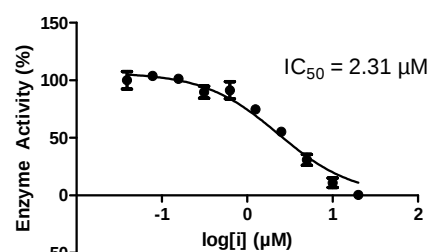
Compound 14m



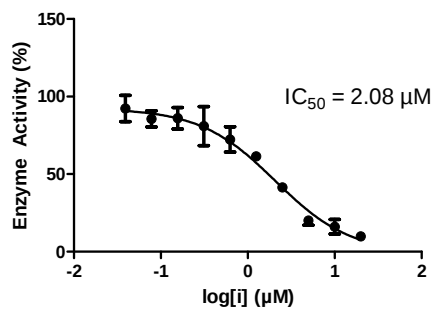
Compound 14j



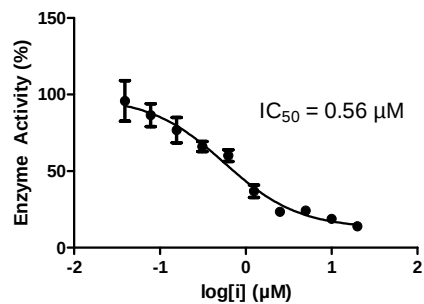
Compound 14a



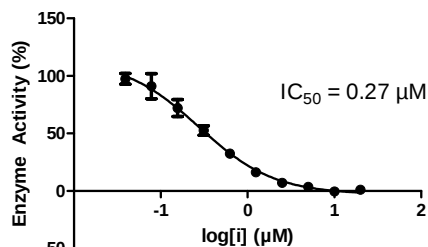
Compound 14k



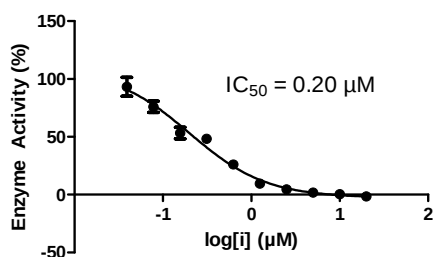
Compound 14n



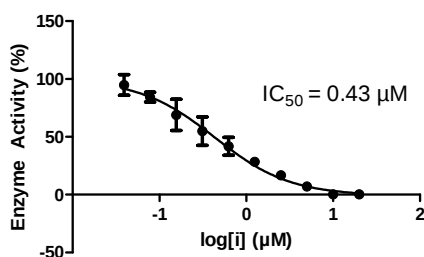
Compound 14h



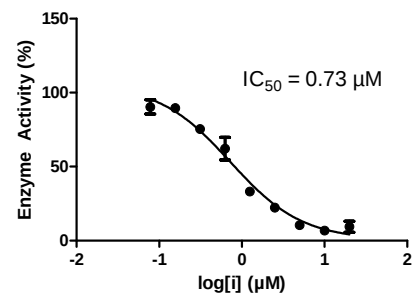
Compound 14g



Compound 14f



Compound 14c



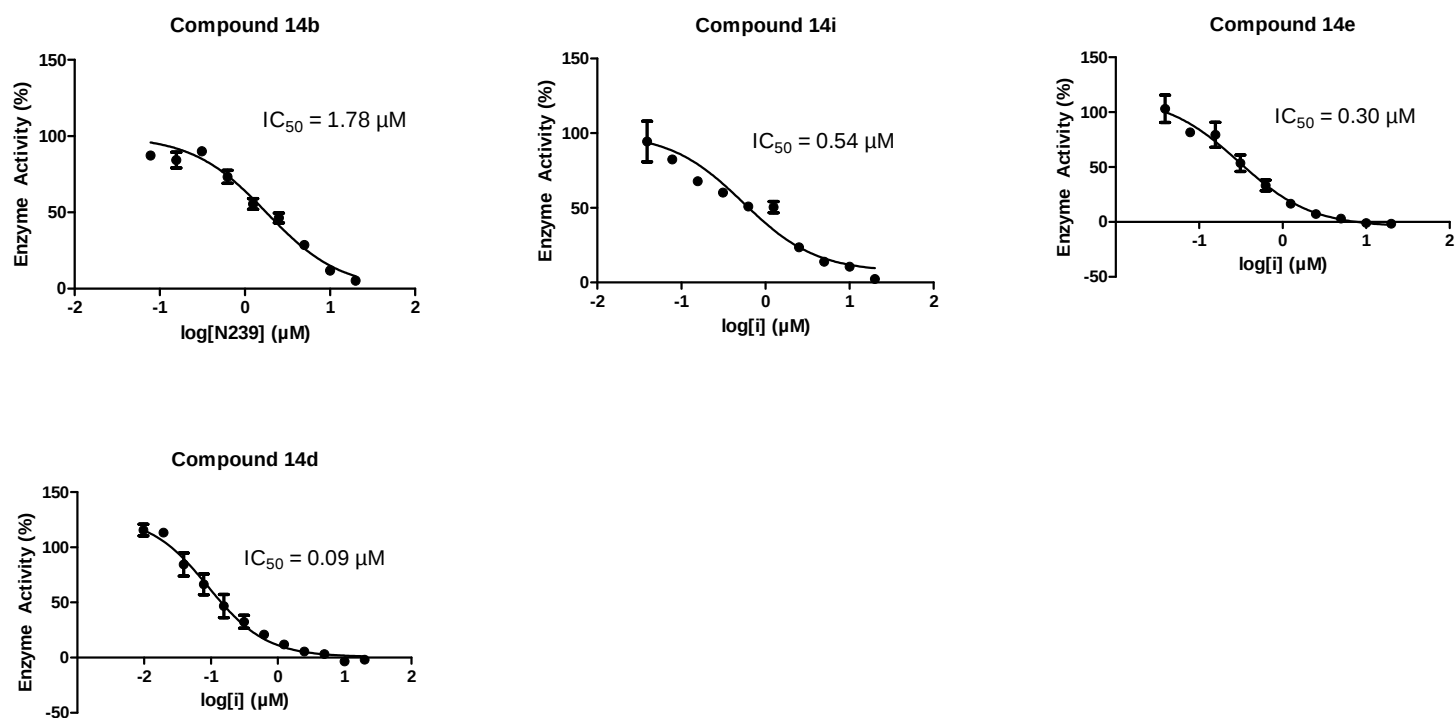


Figure S2.  $\text{IC}_{50}$  graphs

## Enzyme Kinetics

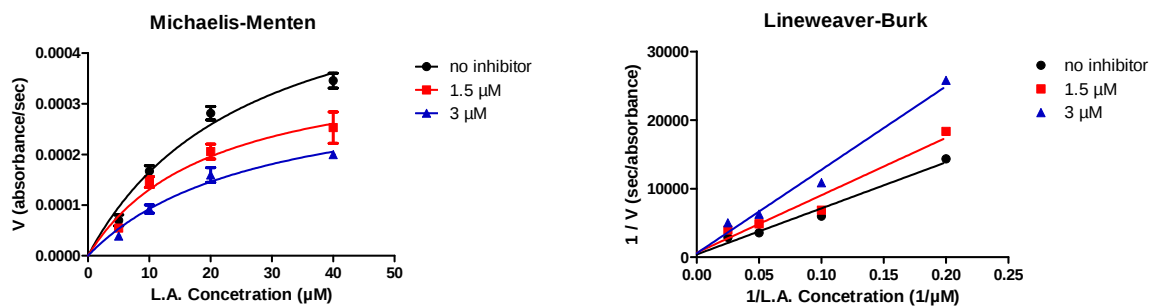


Figure S3. Steady-State kinetic characterization of 15-LOX-1 in the presence of different concentrations of compound 14l: A) Michaelis-Menten representation and B) Lineweaver-Burk representation.

Table S3. Enzyme kinetic parameters for inhibition of 15-LOX-1 by compound 14l.

| 14l ( $\mu\text{M}$ ) | $K_m^{\text{app}}$ ( $\mu\text{M}$ ) | $V_{\text{max}}^{\text{app}}$ (absorbance/s) |
|-----------------------|--------------------------------------|----------------------------------------------|
| 0                     | $16.9 \pm 9.8$                       | $4.6 \times 10^{-4} \pm 1.1 \times 10^{-4}$  |
| 1.5                   | $20.0 \pm 15$                        | $3.9 \times 10^{-4} \pm 1.4 \times 10^{-4}$  |
| 3                     | $27.5 \pm 17$                        | $3.4 \times 10^{-4} \pm 1.1 \times 10^{-4}$  |

## Molecular Modeling

Since competitive inhibition was observed, the inhibitors were docked in the active site of the enzyme. The molecular modeling studies were performed in MOE software (2012.10) and highest scoring docking poses were chosen. The experiments were performed with rescoring model 1 London dG (refinement: forcefield) and rescoring 2: GBVI/WSA dG, followed by minimization energy (forcefield: MMFF94X; eps = r, cutoff {8,10})

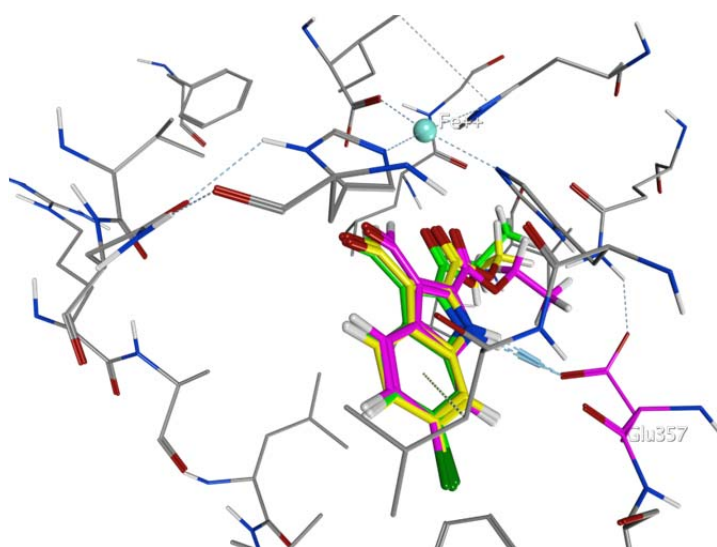


Figure S4. Three of the top five highest scoring poses of the compound 2. The docking score of the pink structure pose is -5.51, from the yellow structure pose is -5.47 and from the green structure pose -5.36.

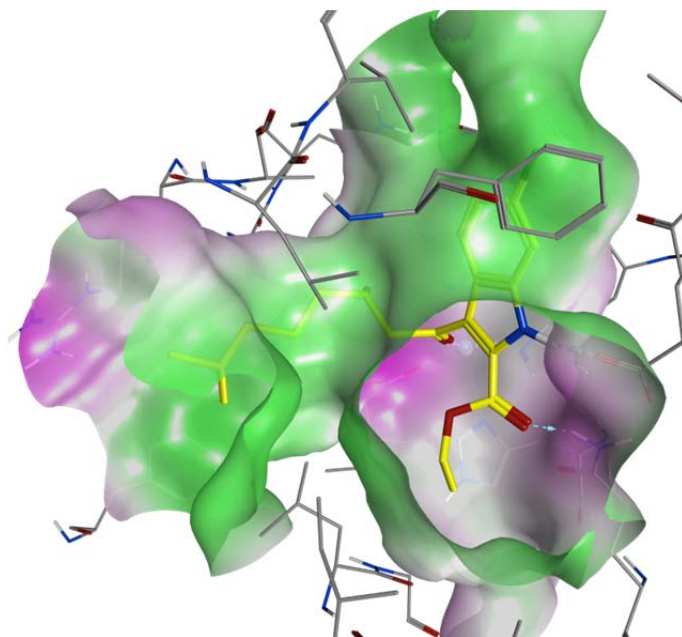


Figure S5. The surface of the active site with the compound 14d in the highest docking score pose. With green is the lipophilic and with red the lipophobic area in the active site of 15-LOX.