

Supplementary materials for:

Synthesis of SARS-CoV-2 M^{pro} Inhibitors bearing a Cinnamic Ester Warhead with *In Vitro* Activity against Human Coronaviruses

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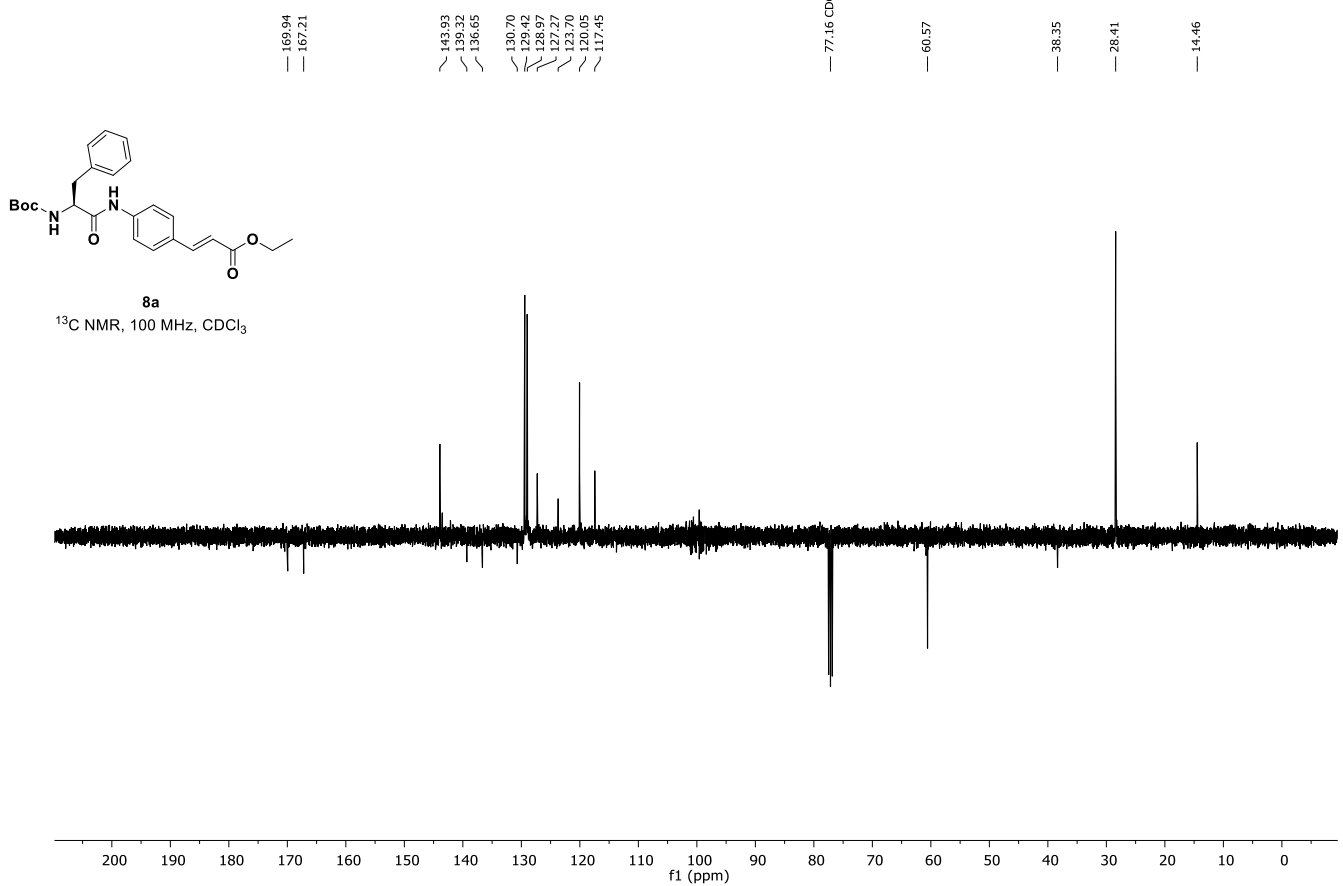
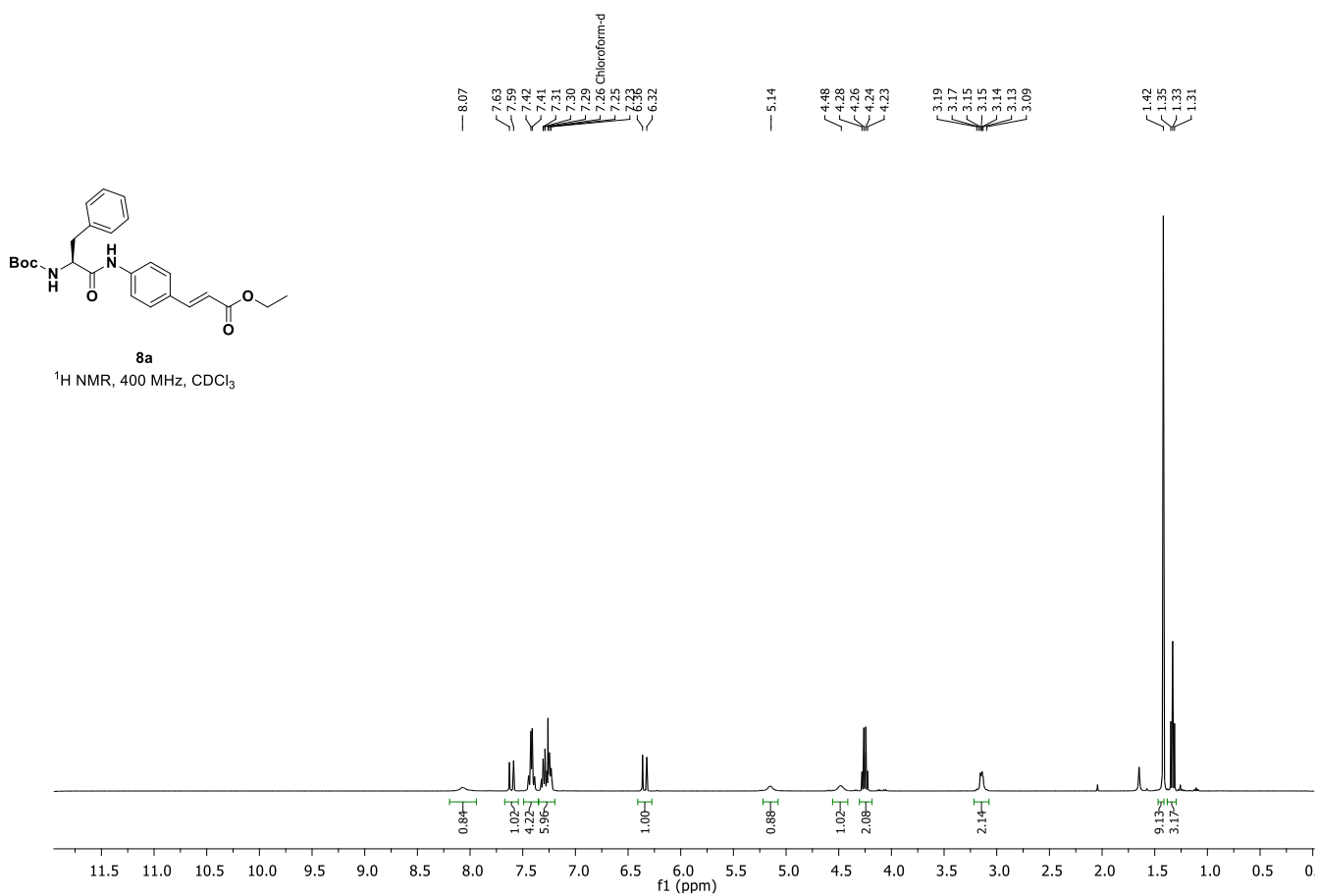
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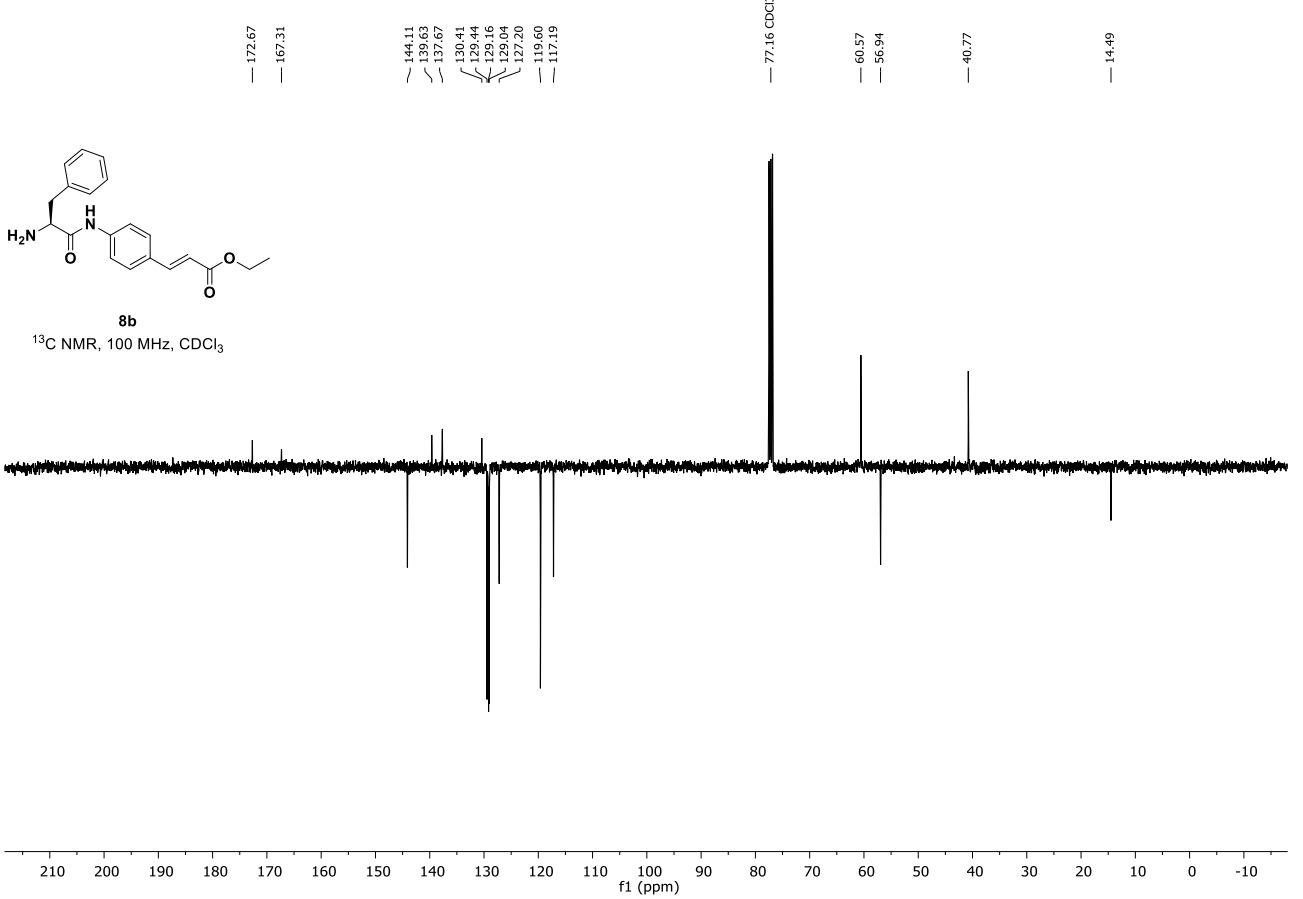
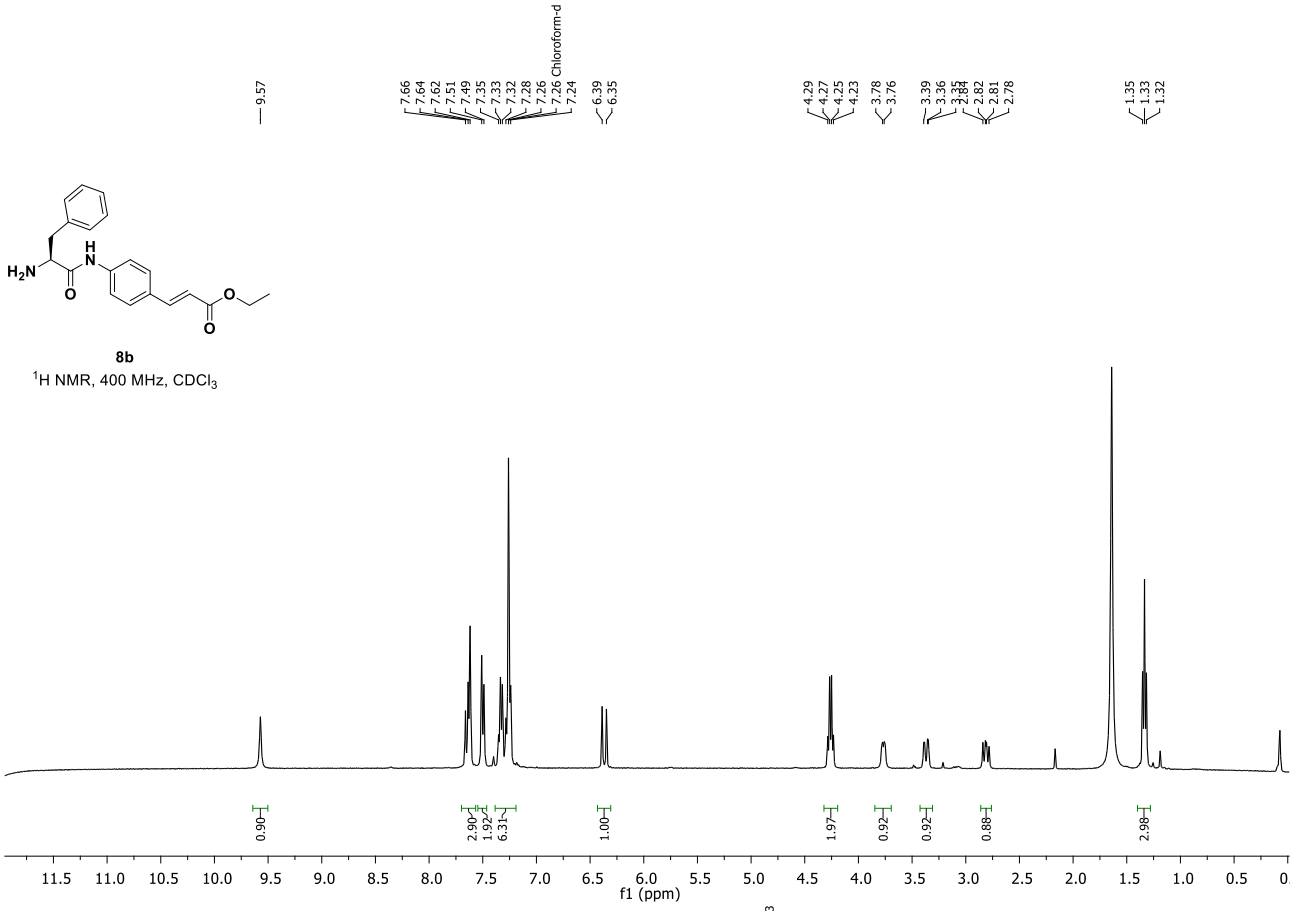
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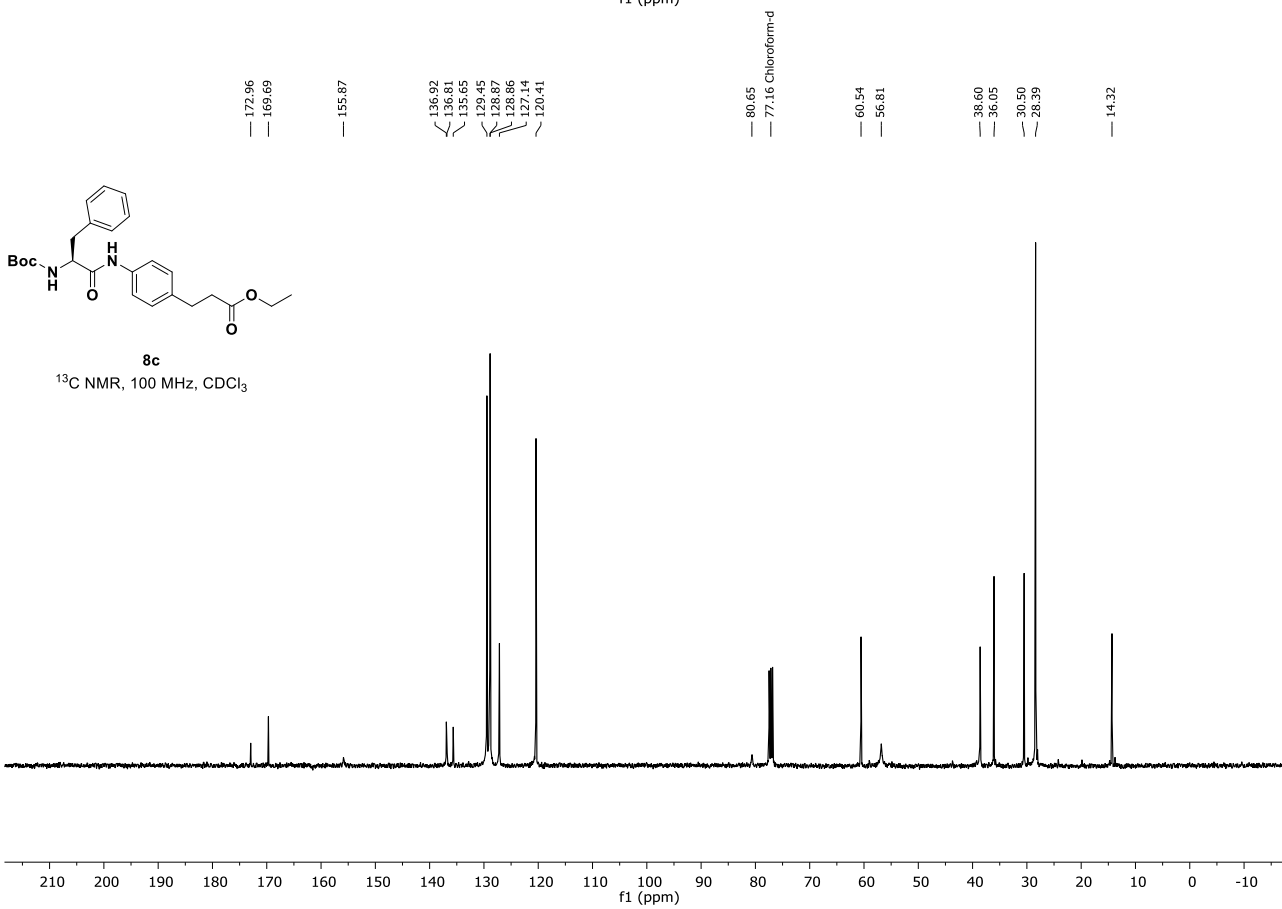
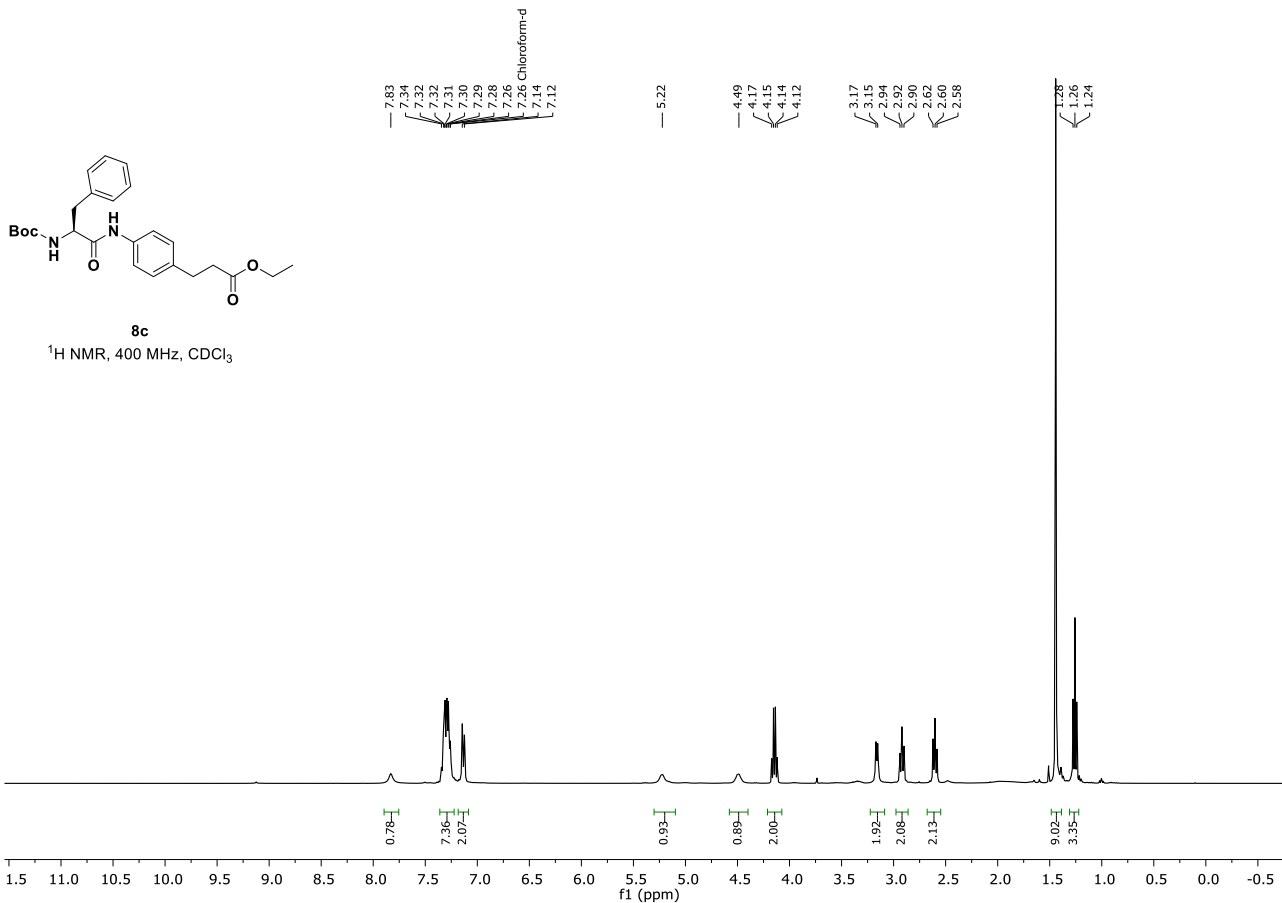
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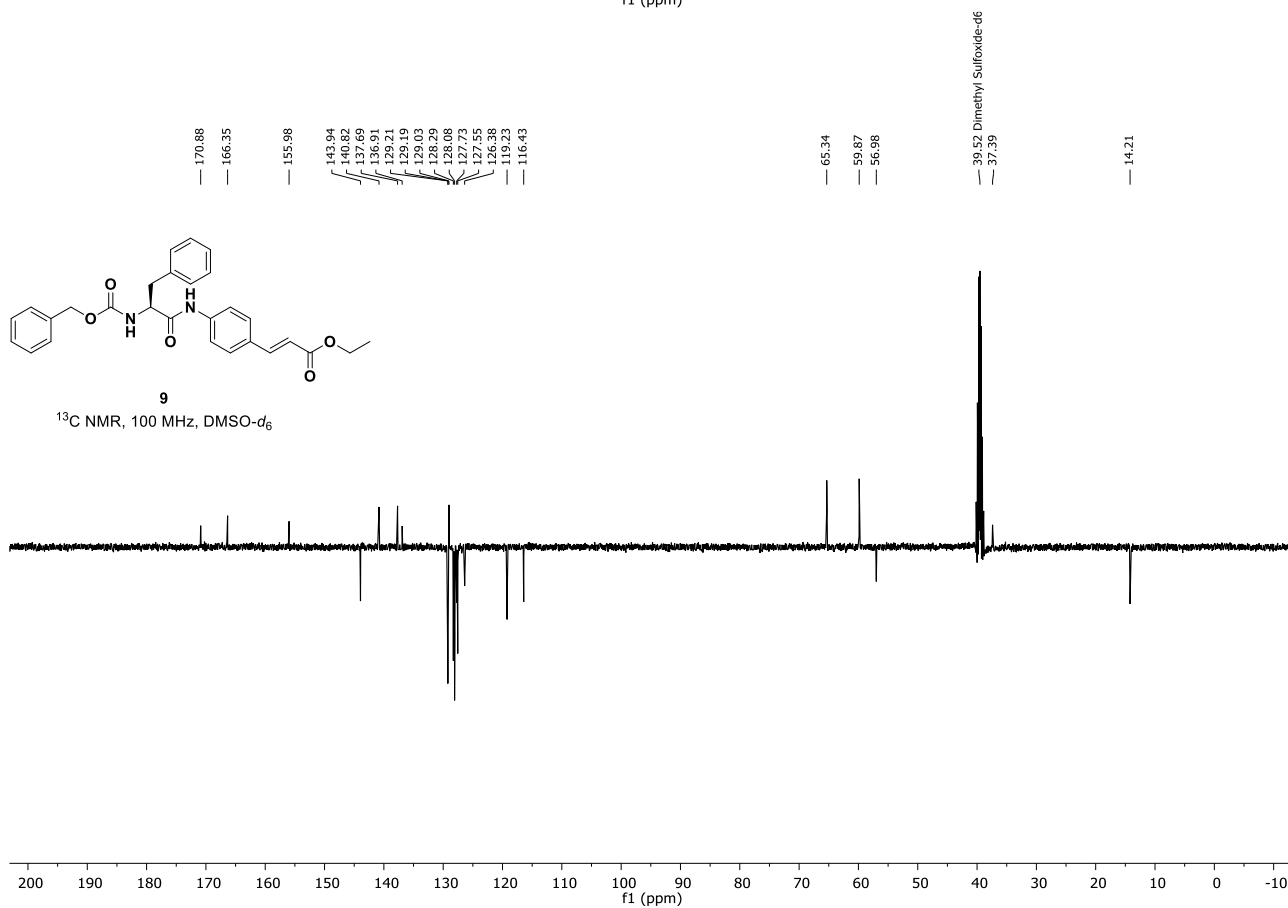
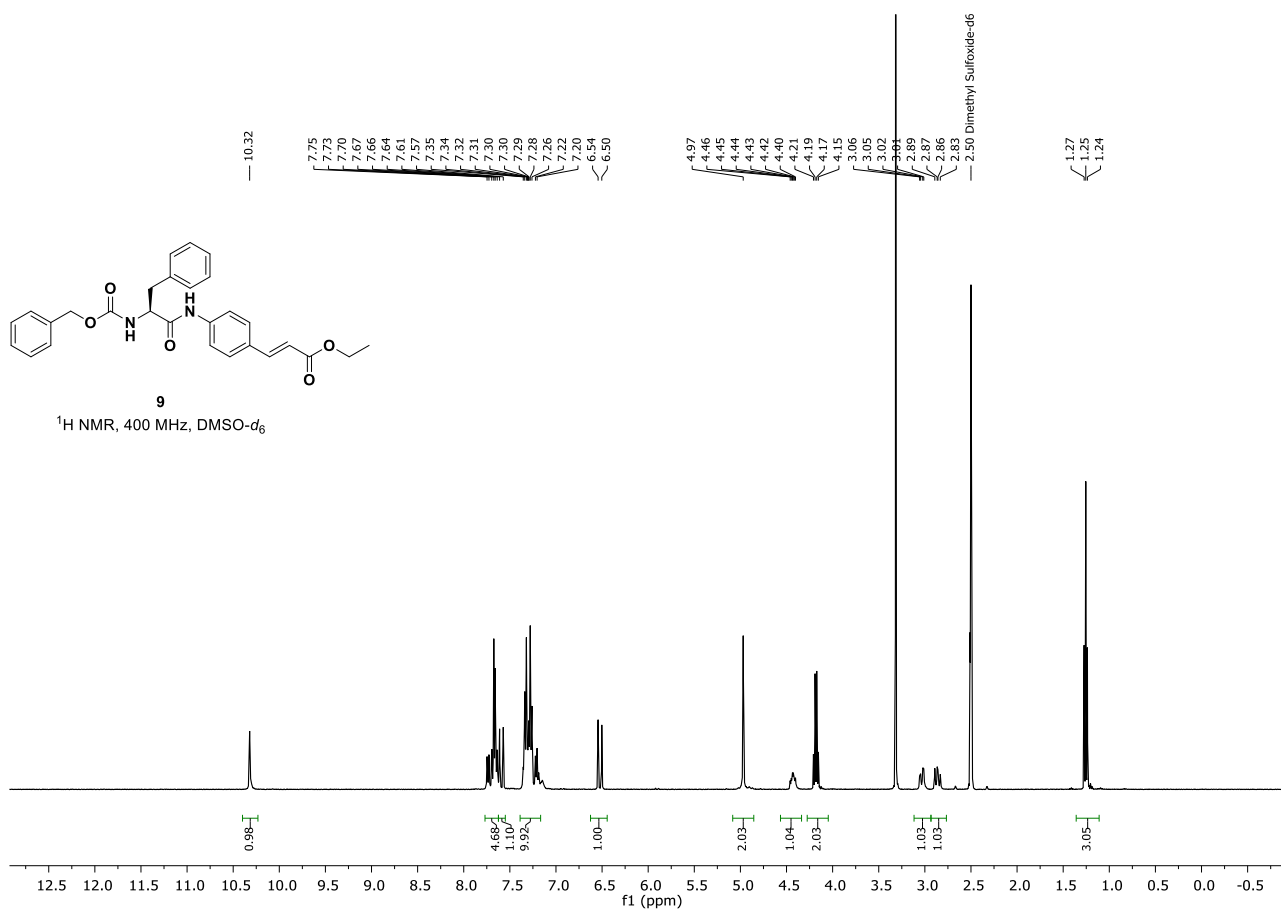
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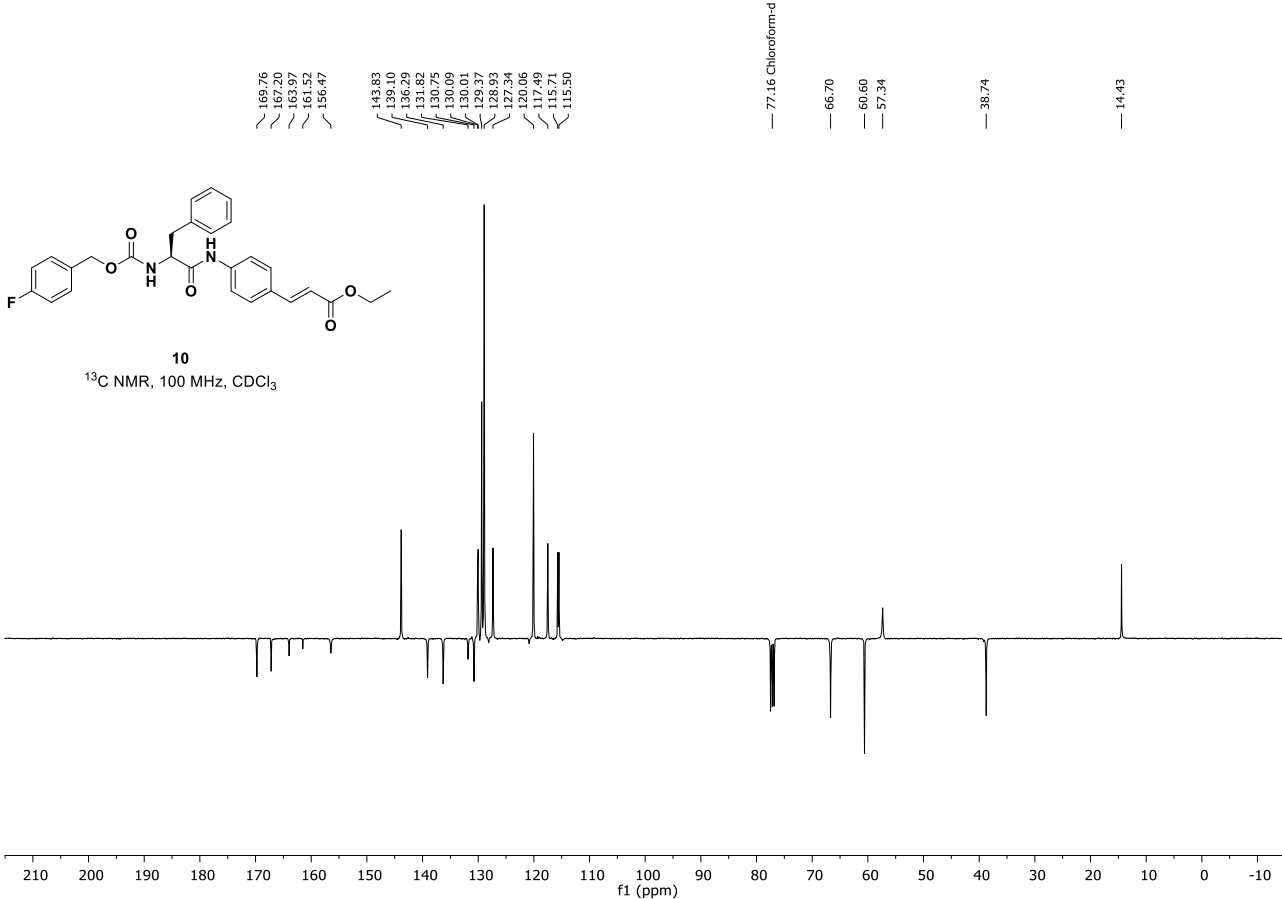
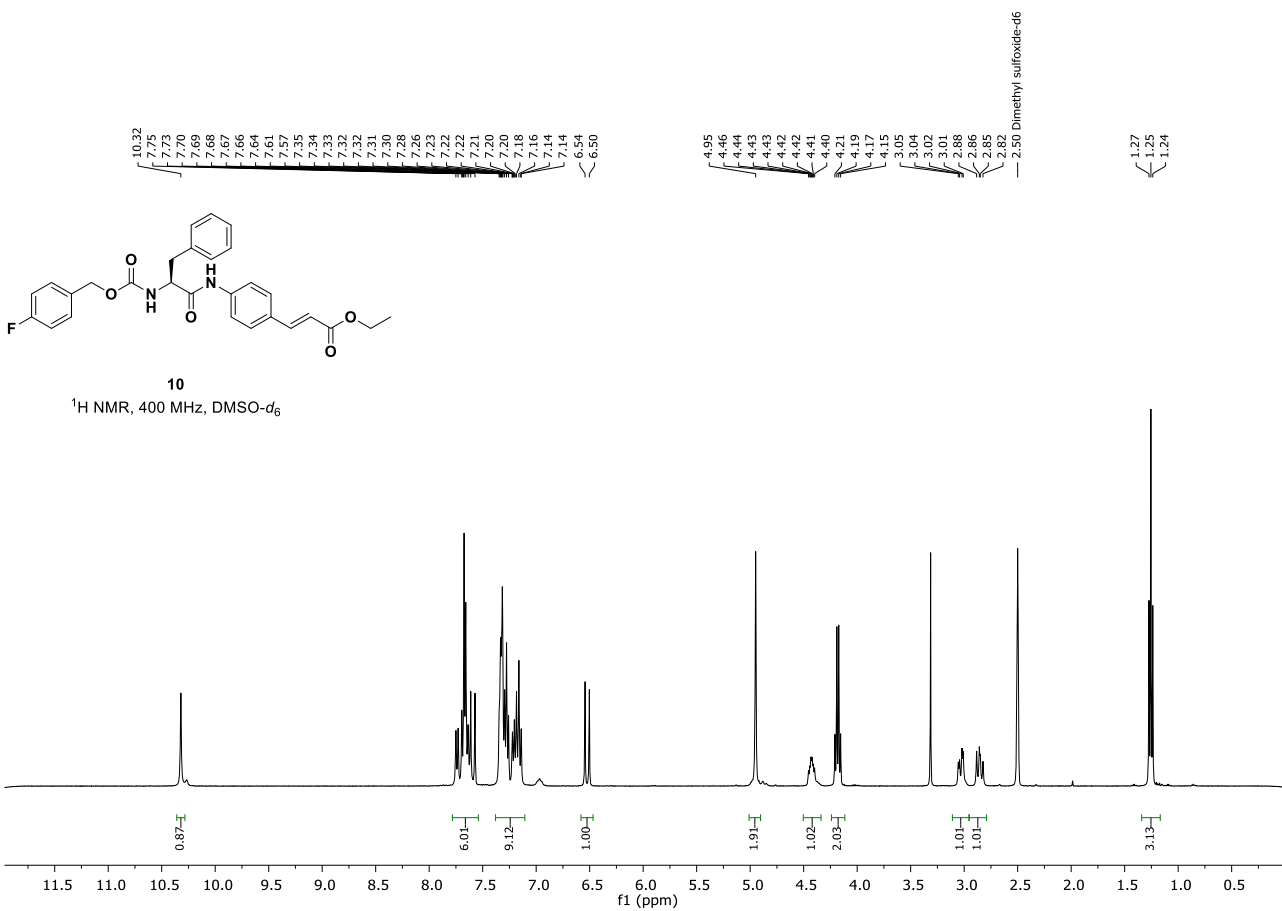
1. ¹H- and ¹³C- Spectra of the Compounds

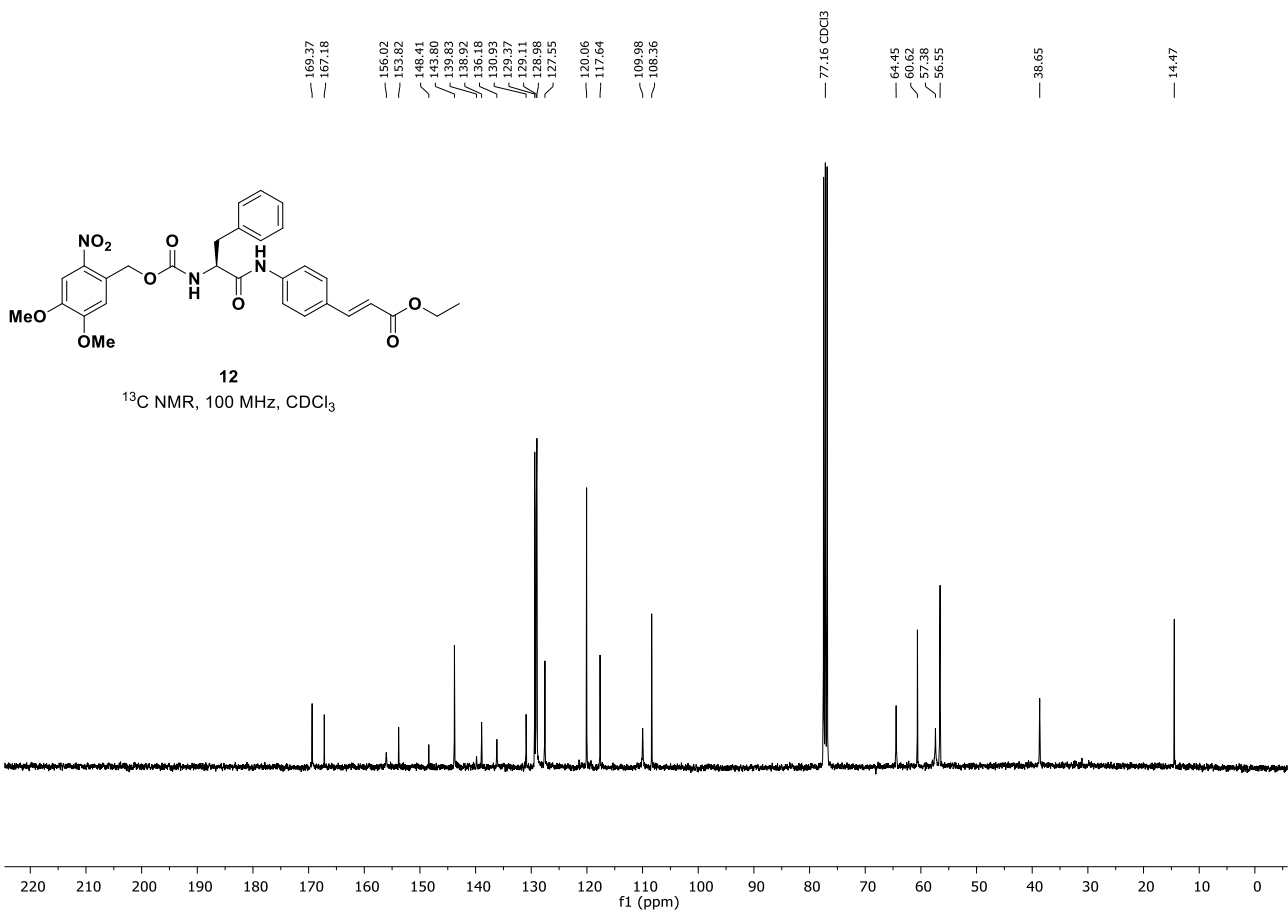
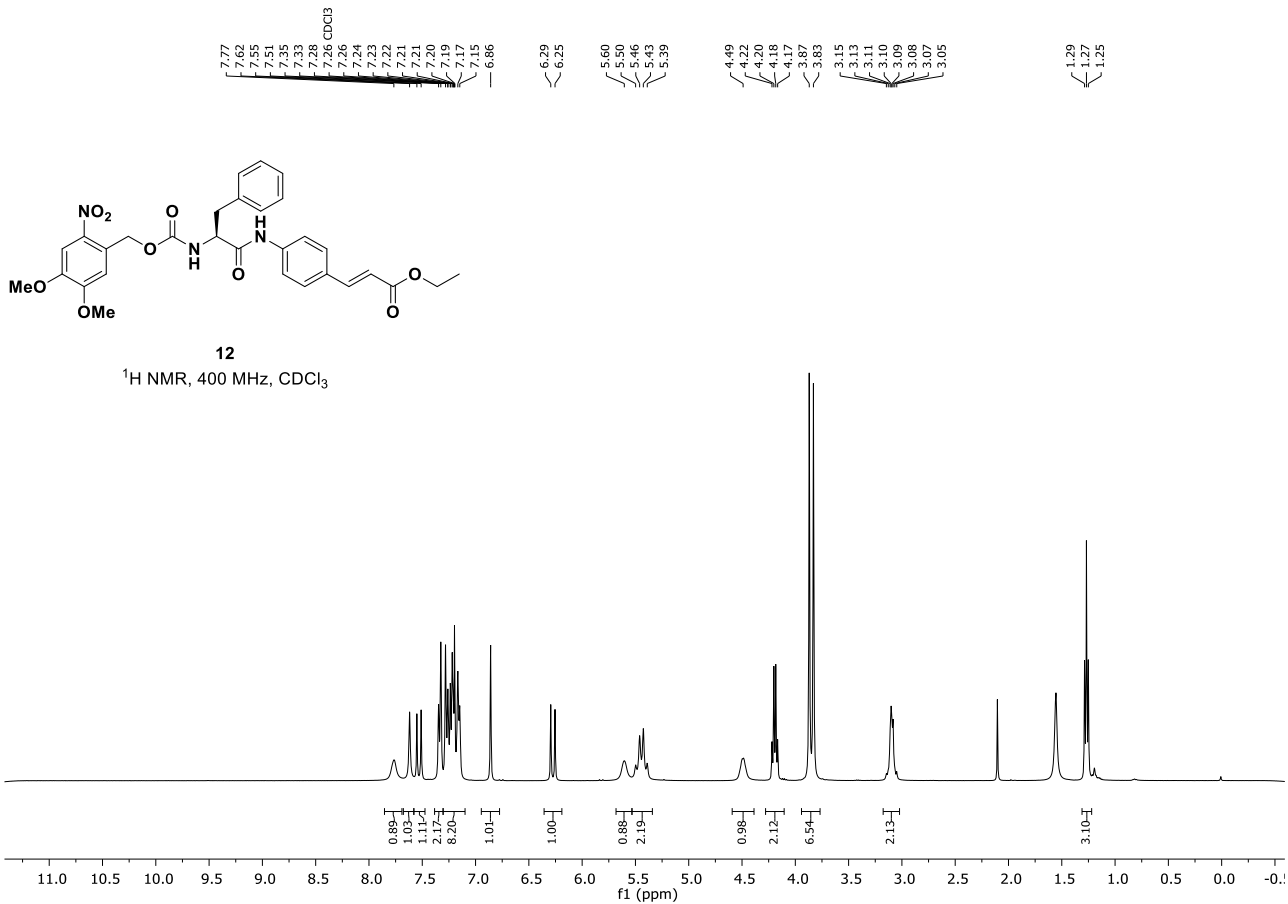


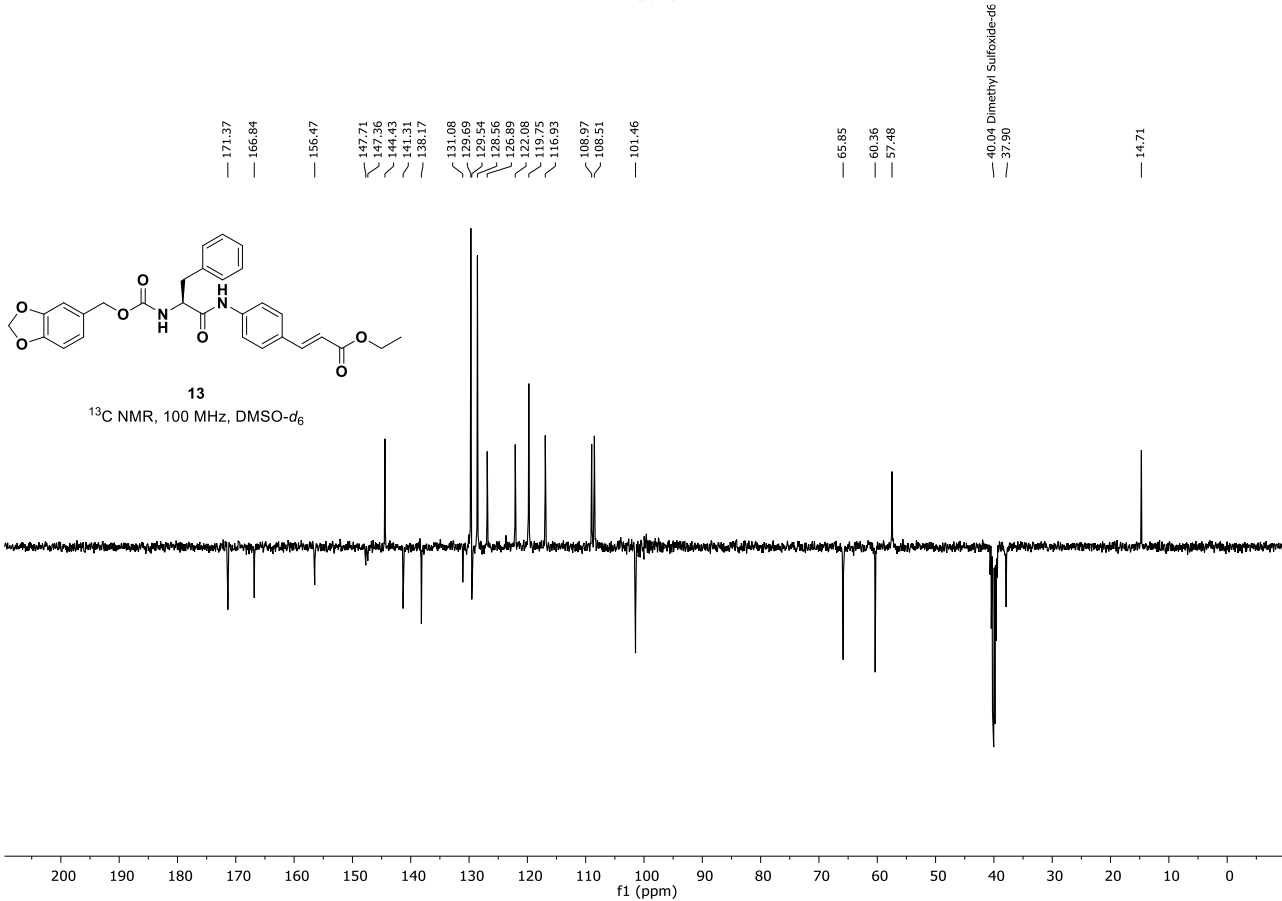
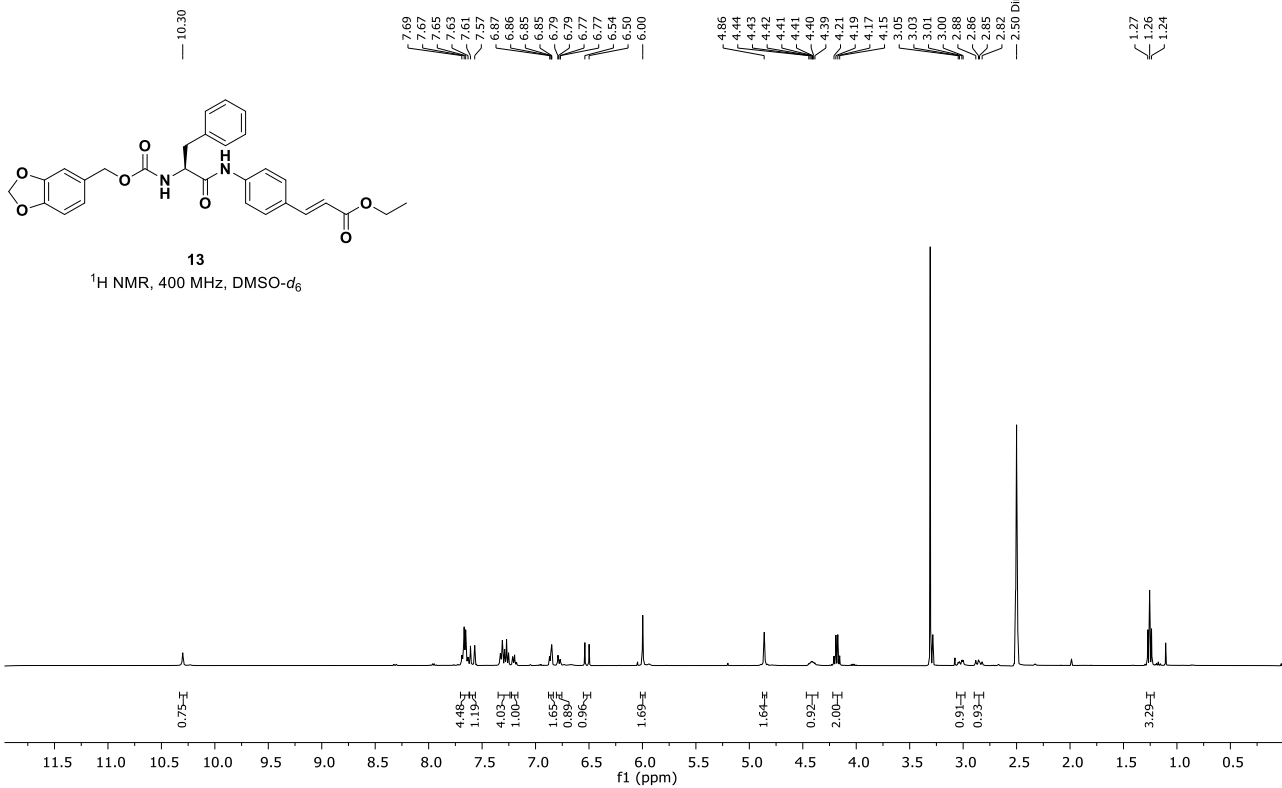


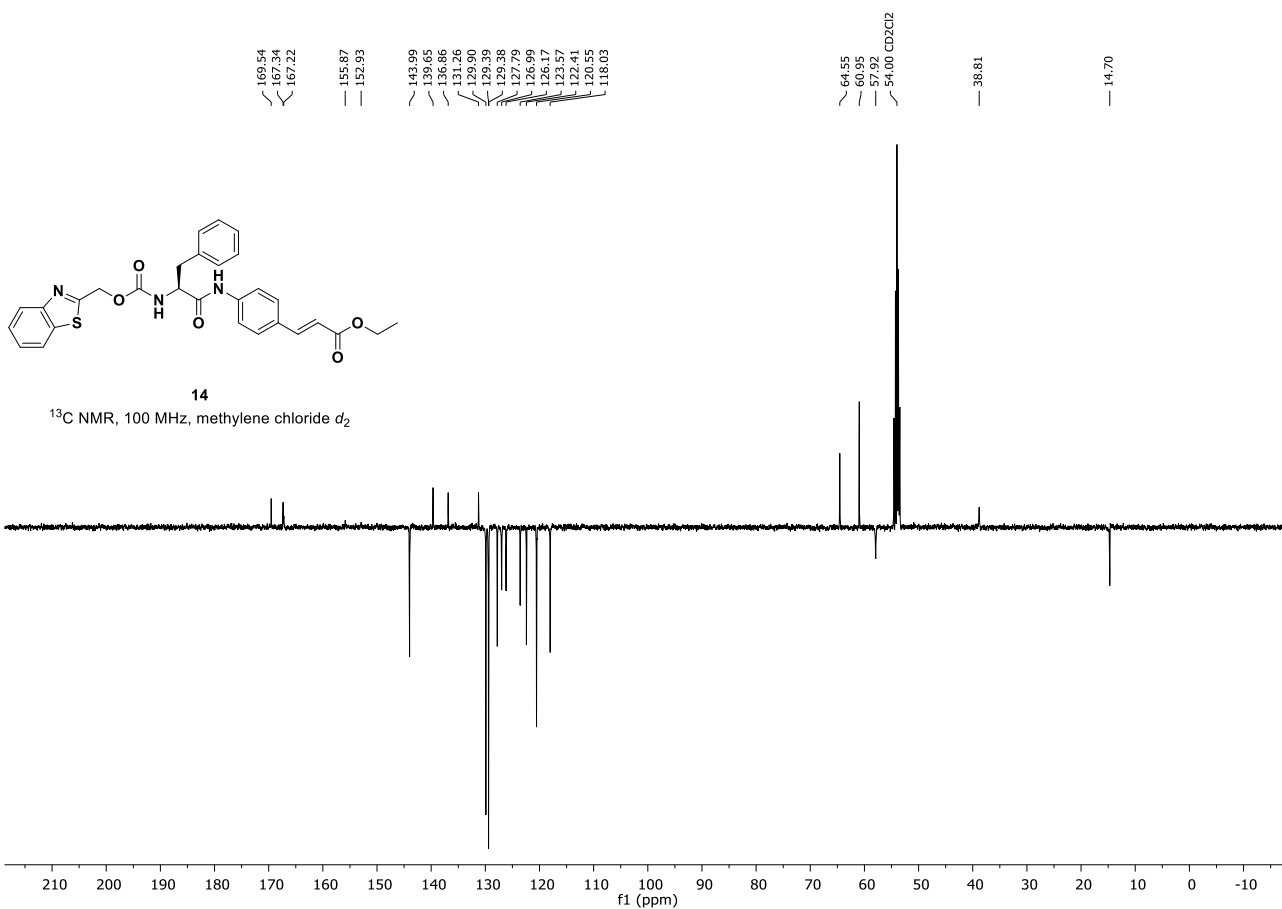
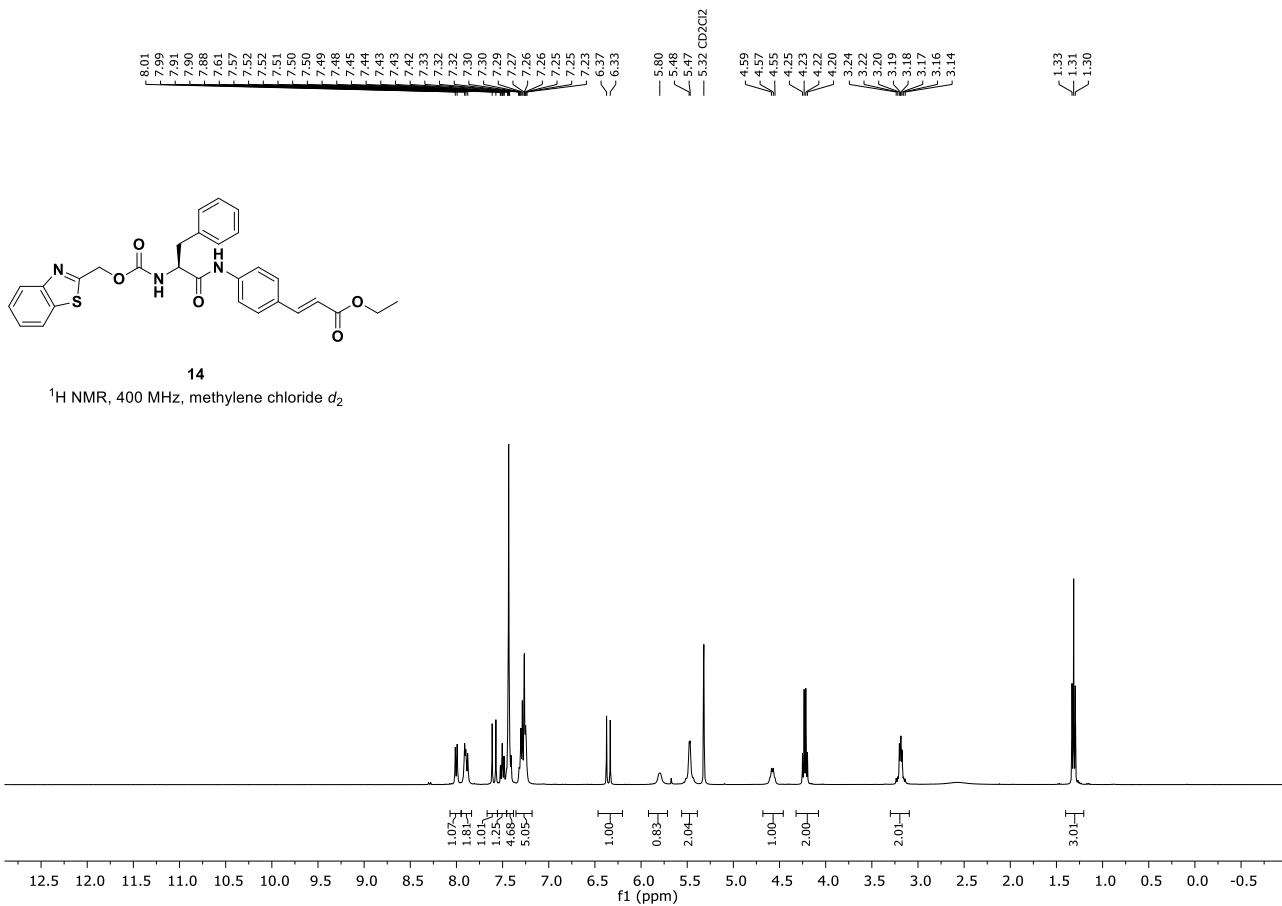


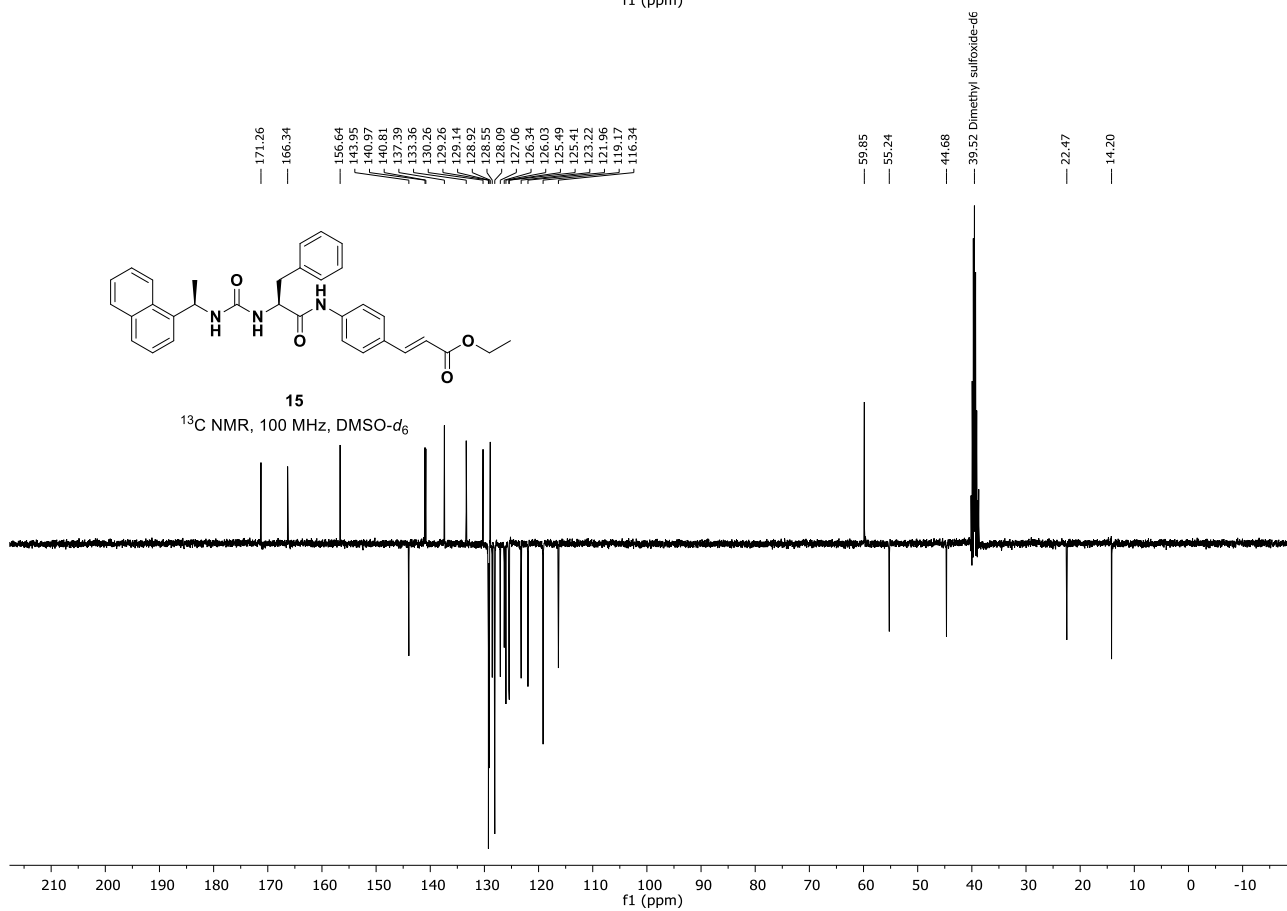
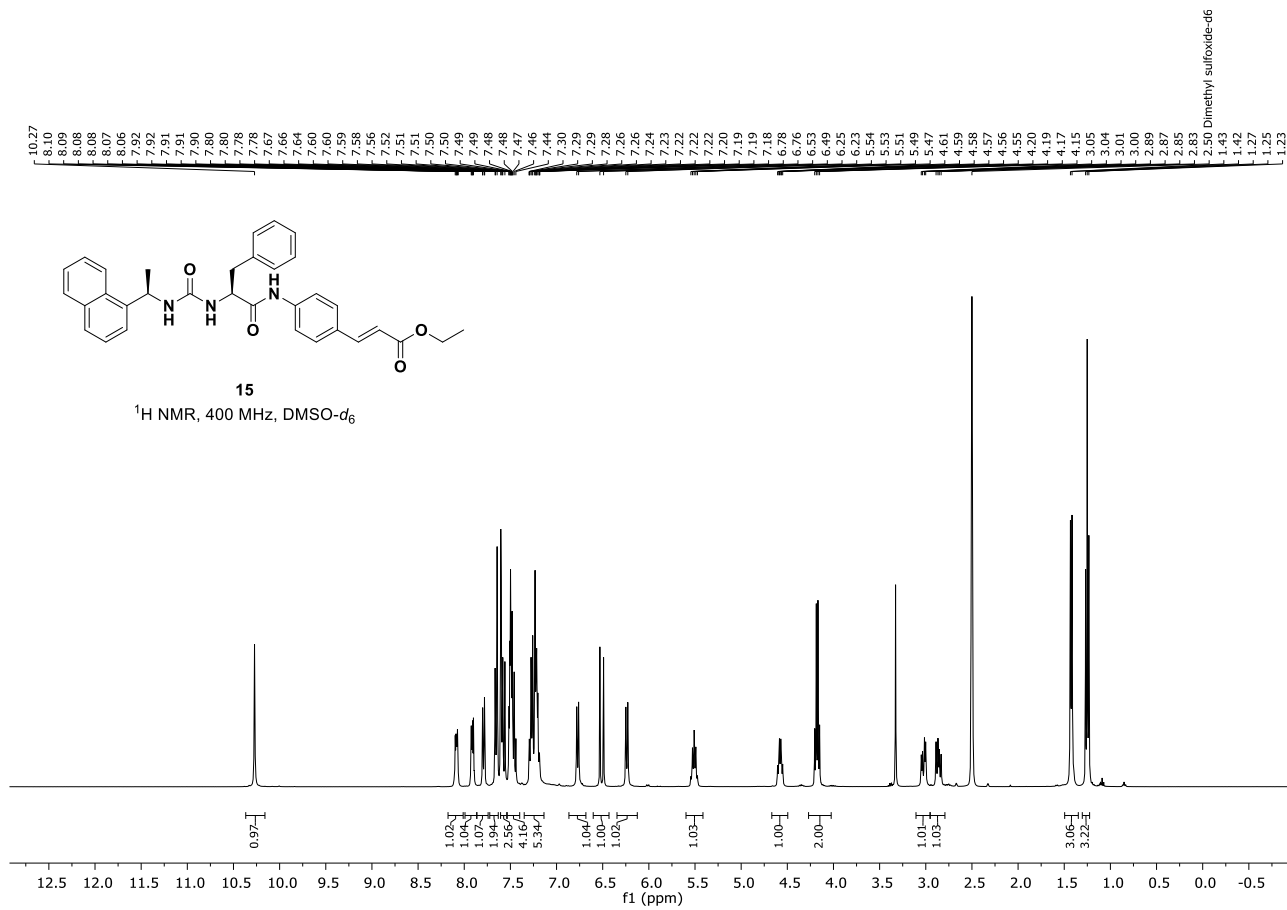


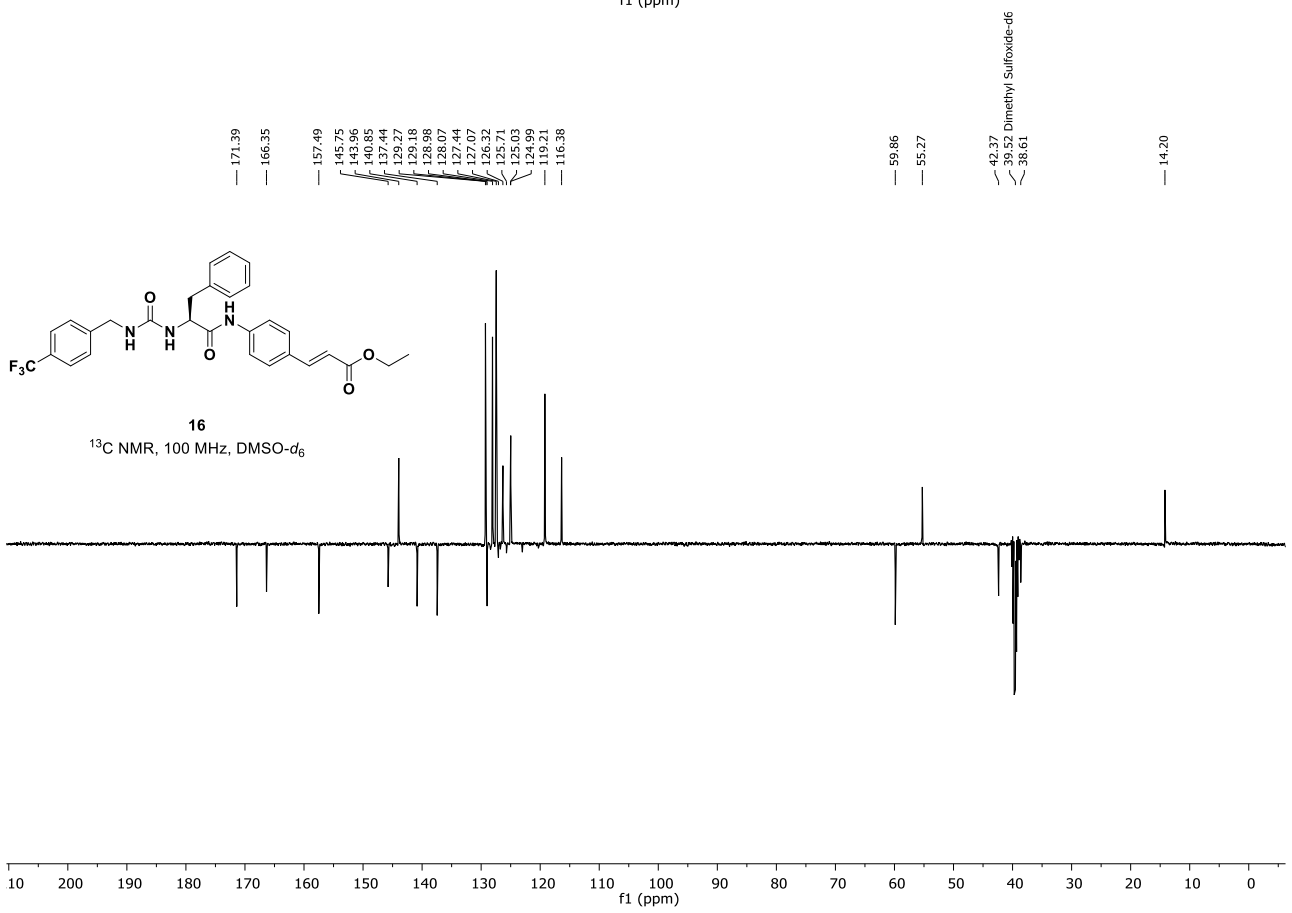
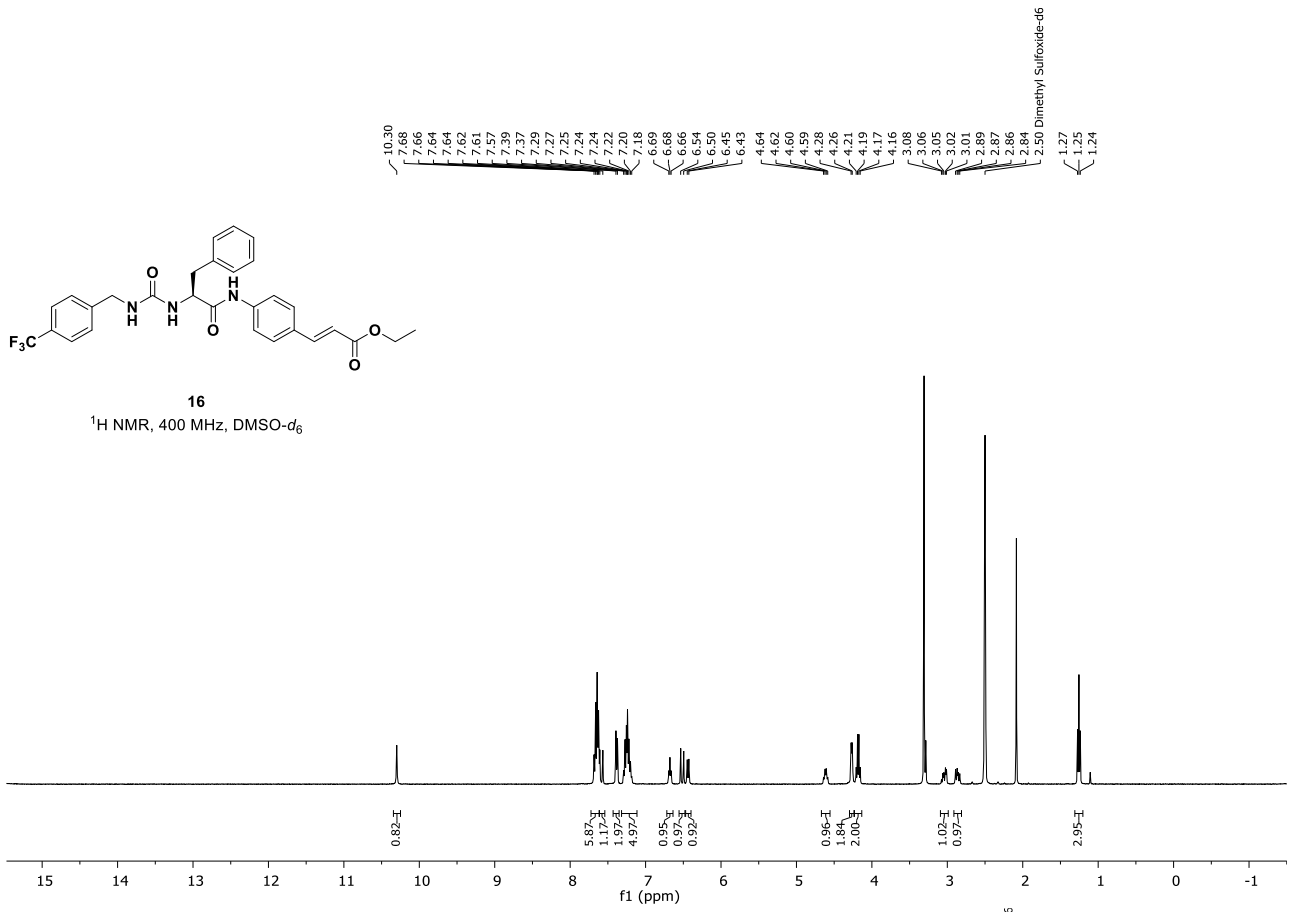


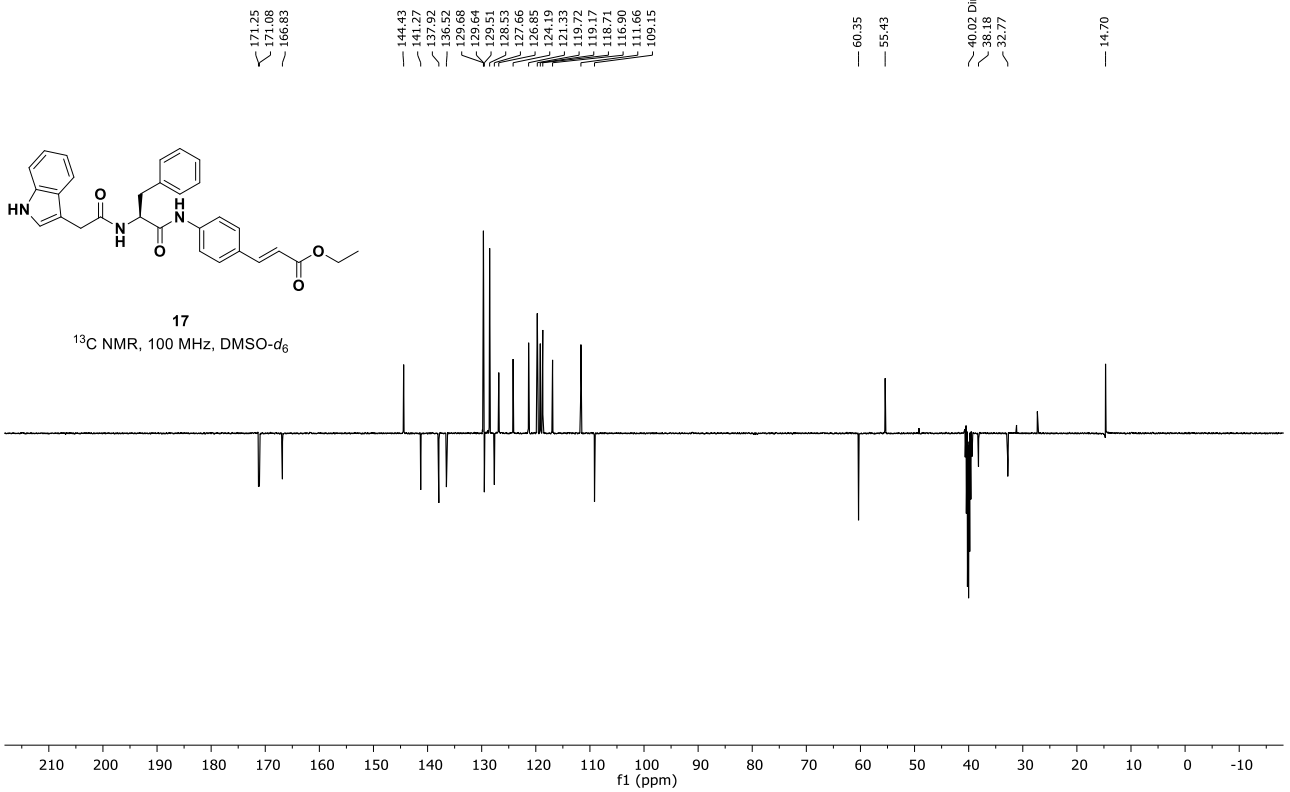
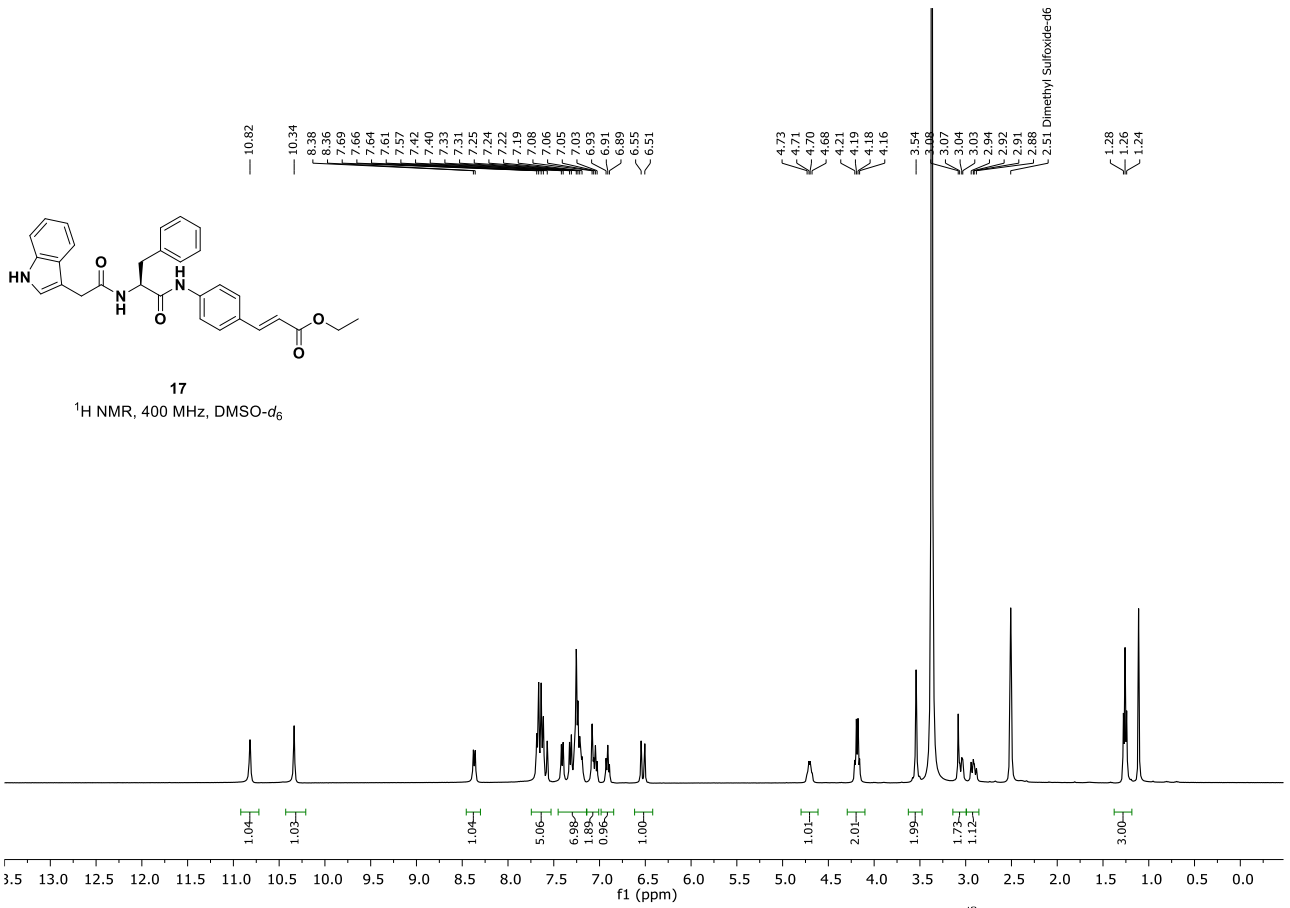


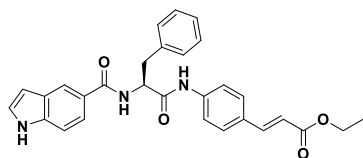






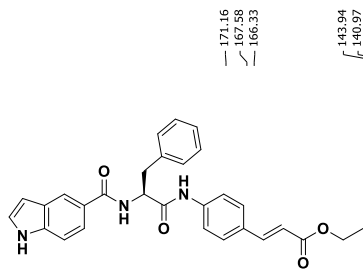
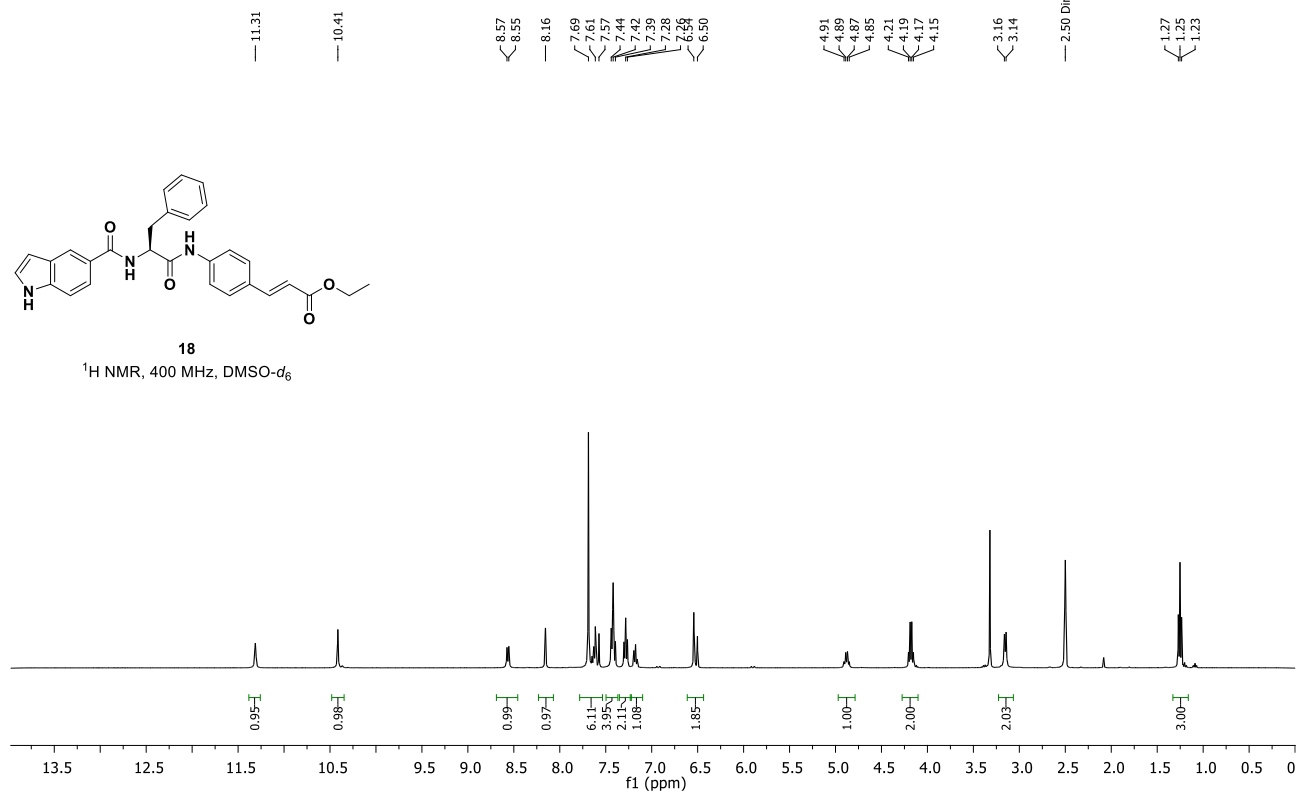






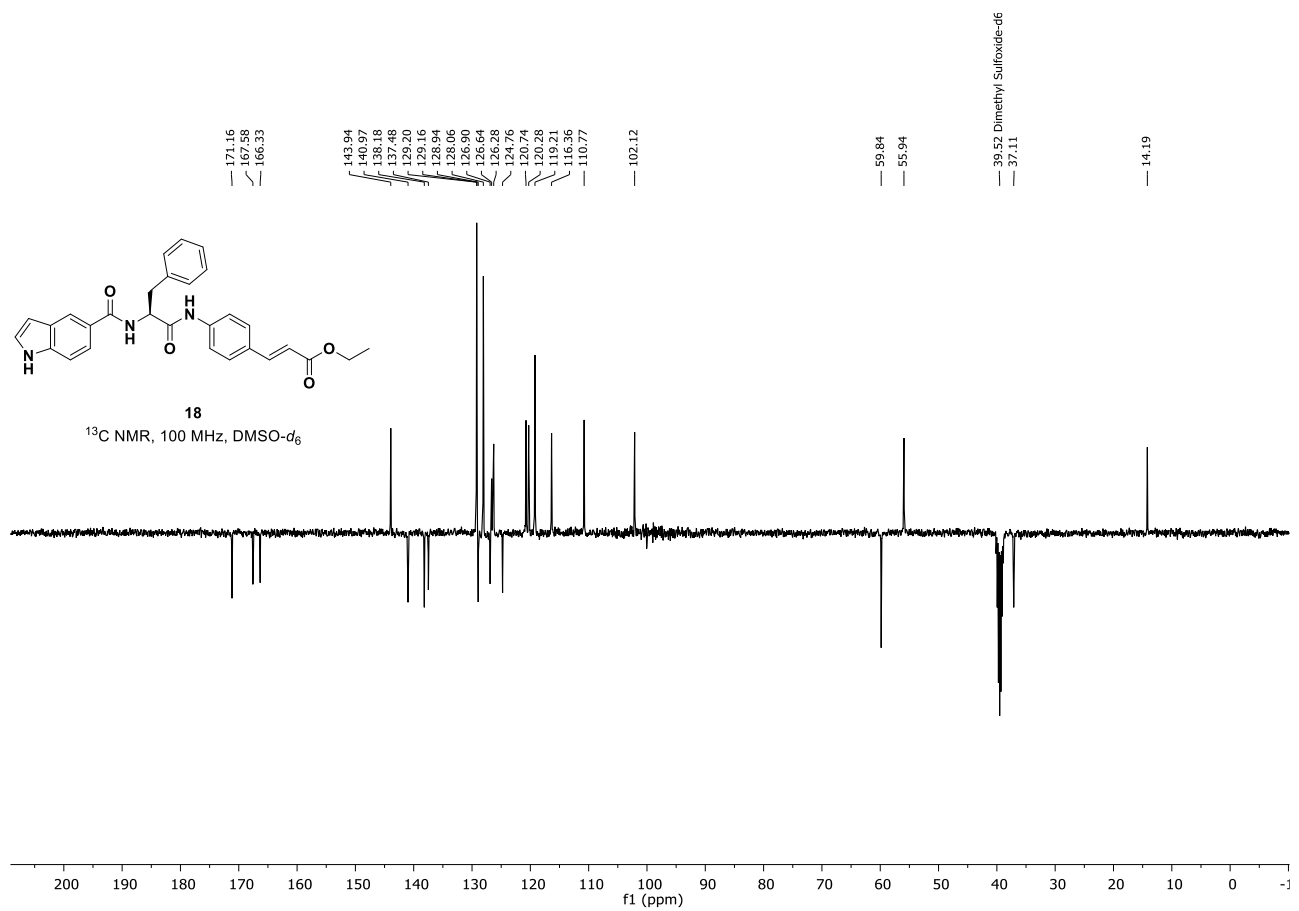
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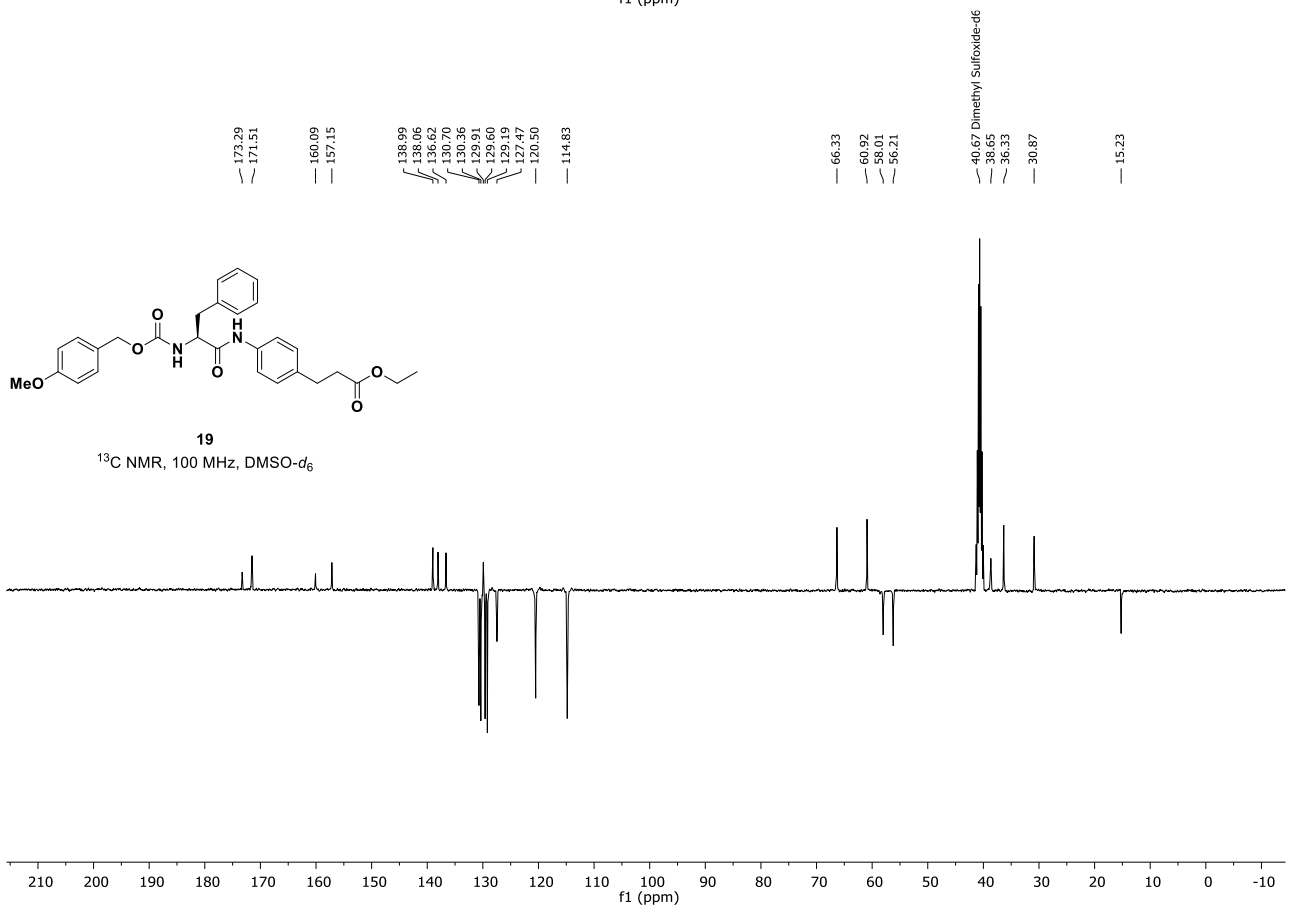
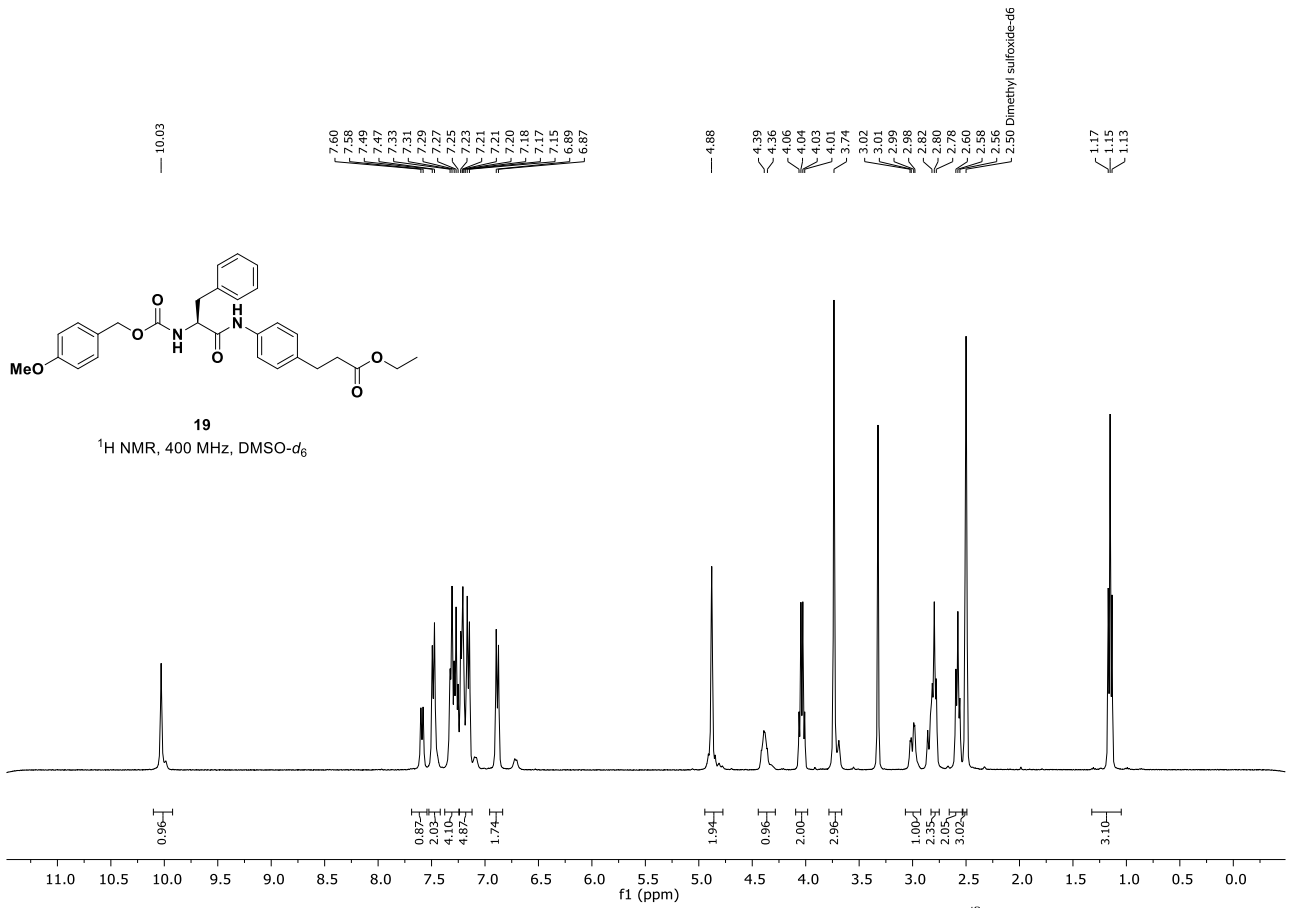
¹H NMR, 400 MHz, DMSO-*d*₆



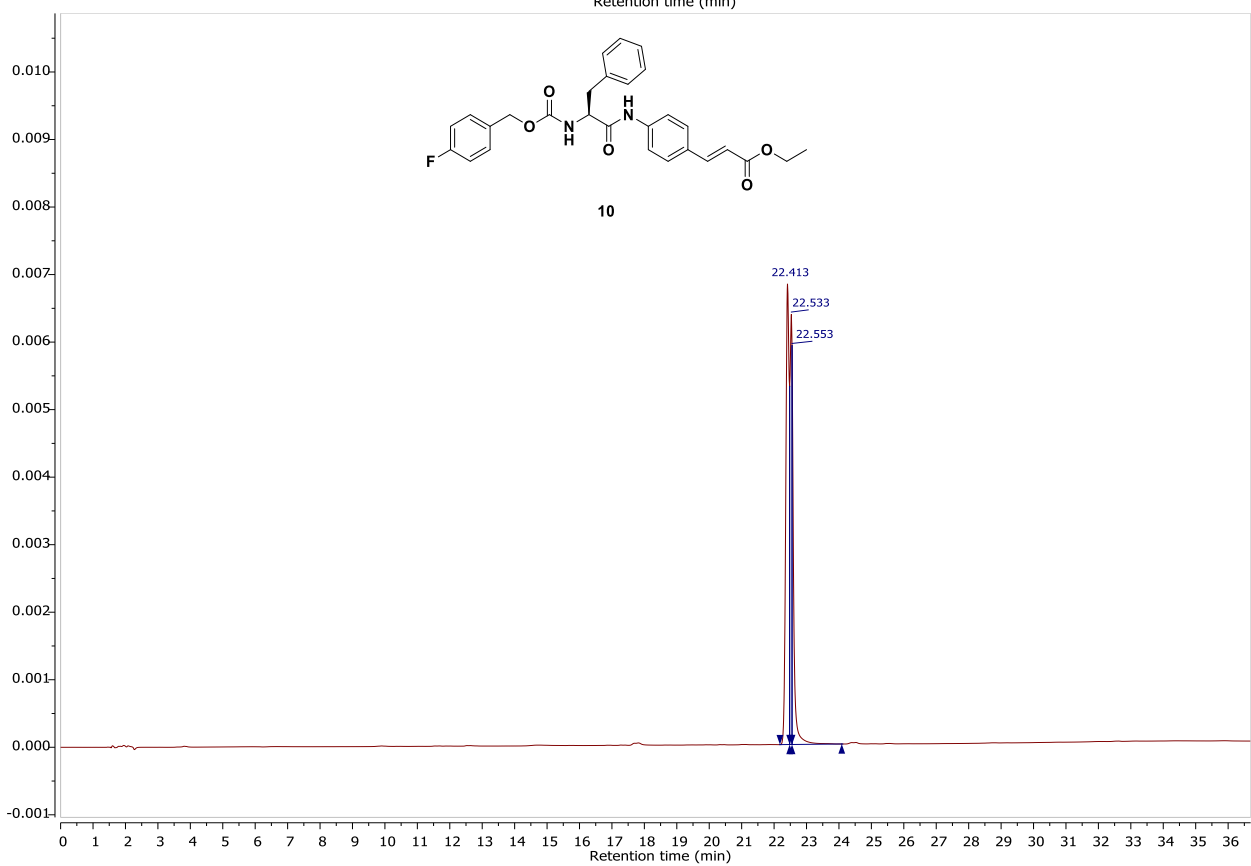
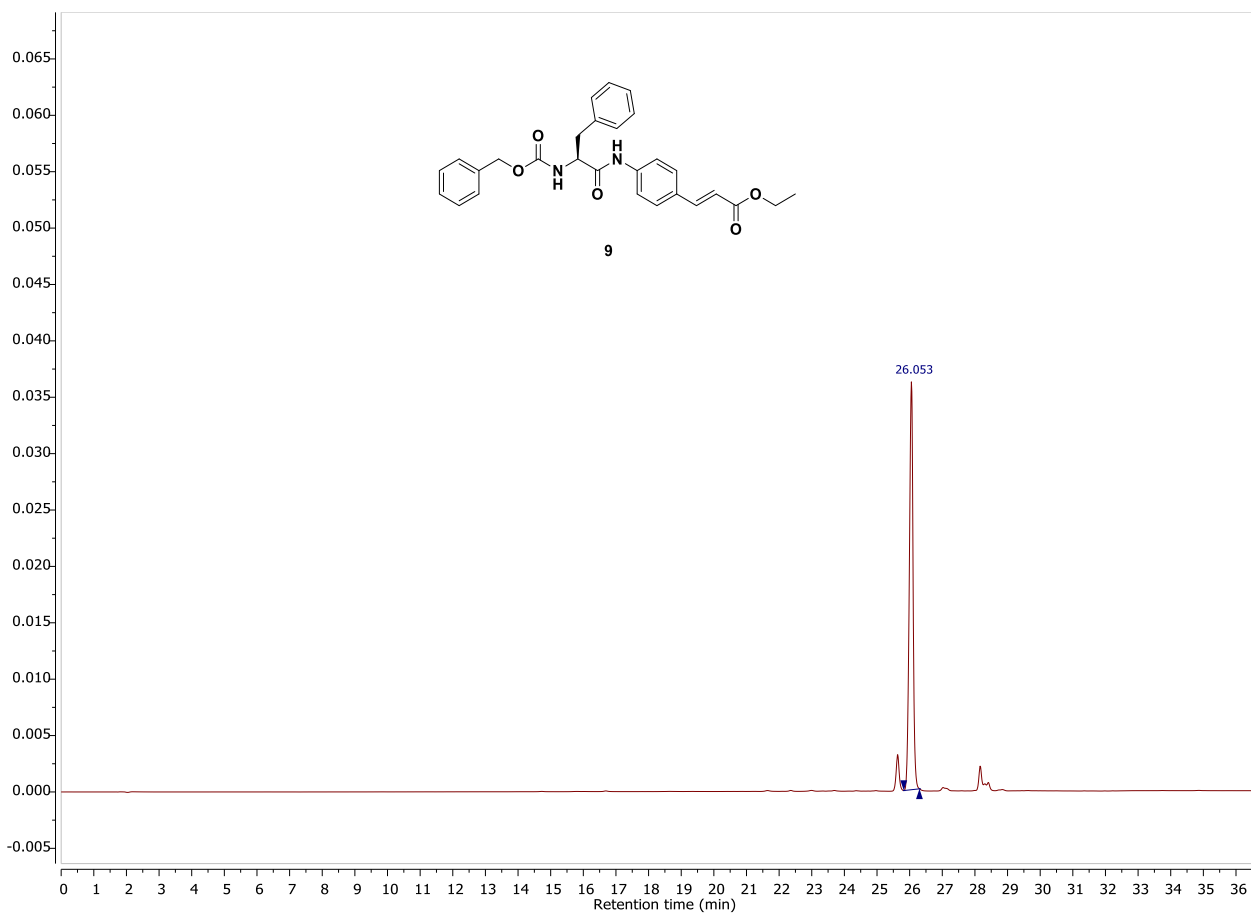
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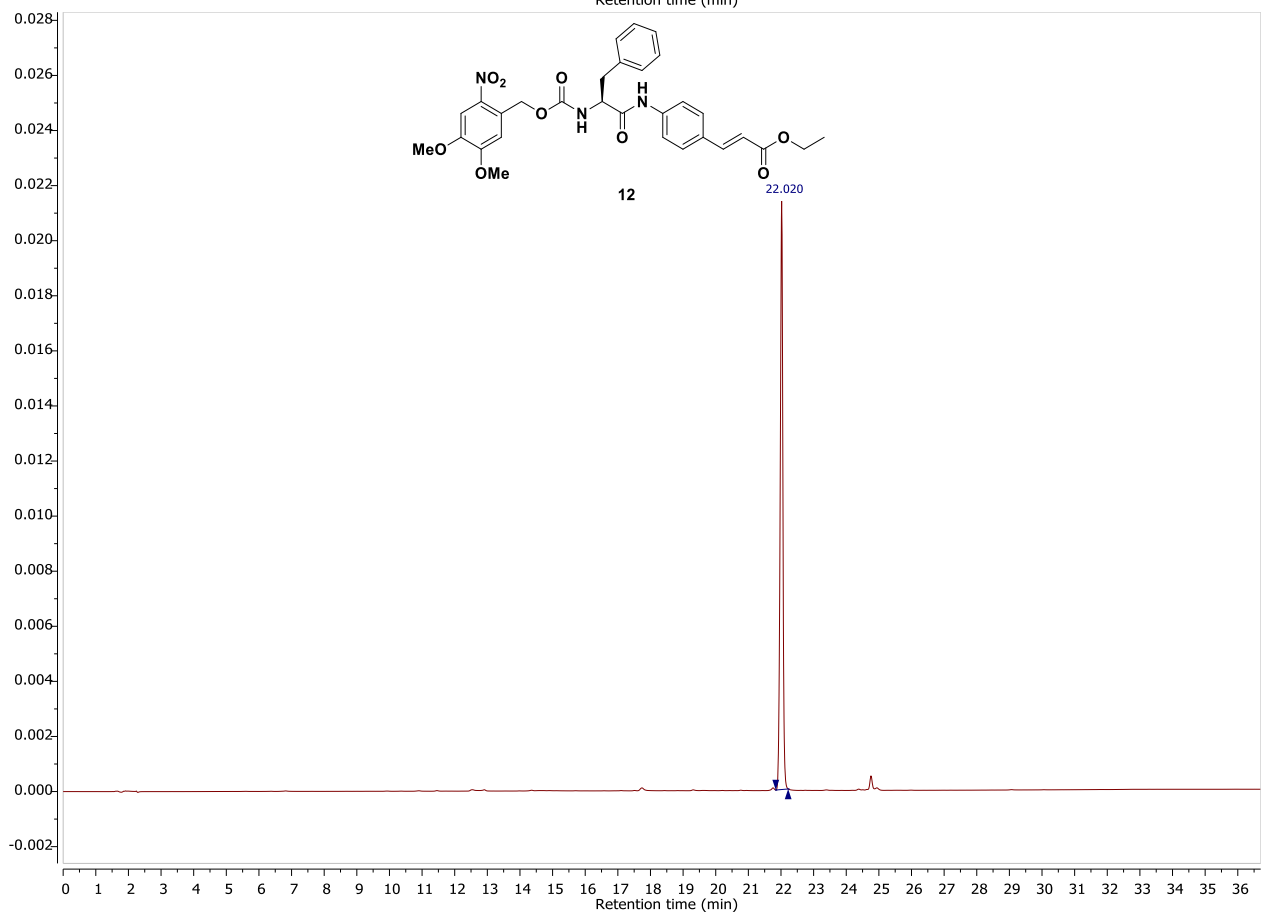
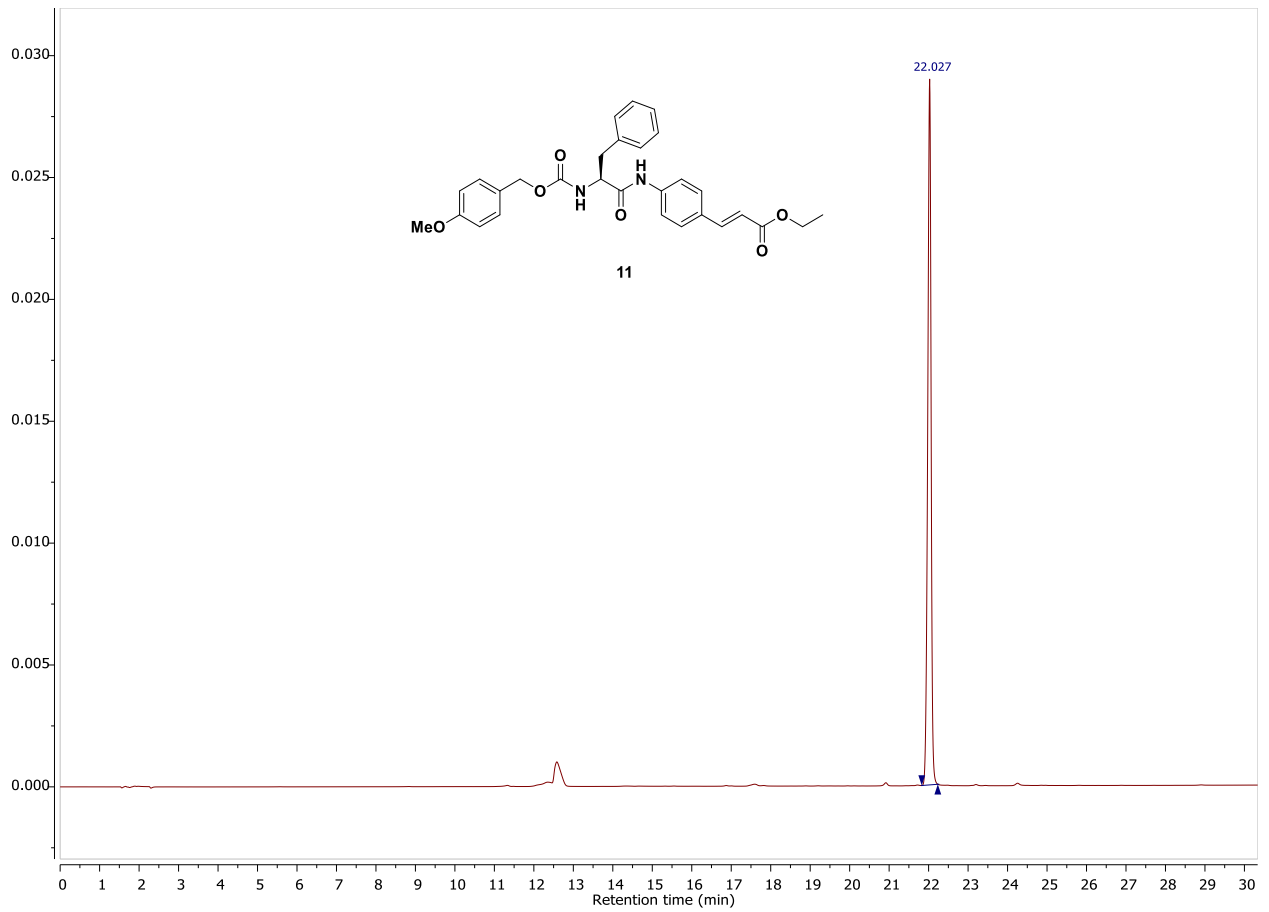
¹³C NMR, 100 MHz, DMSO-*d*₆

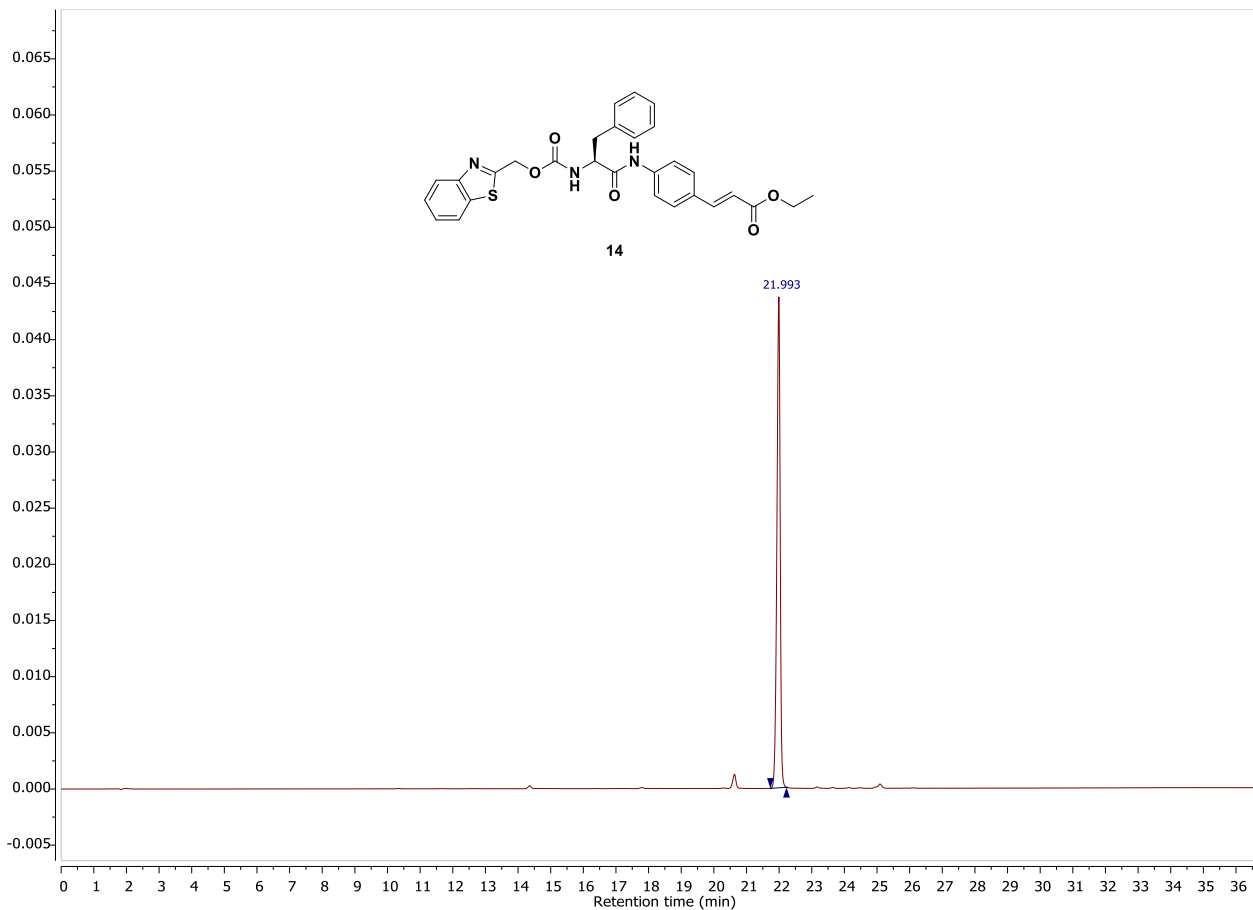
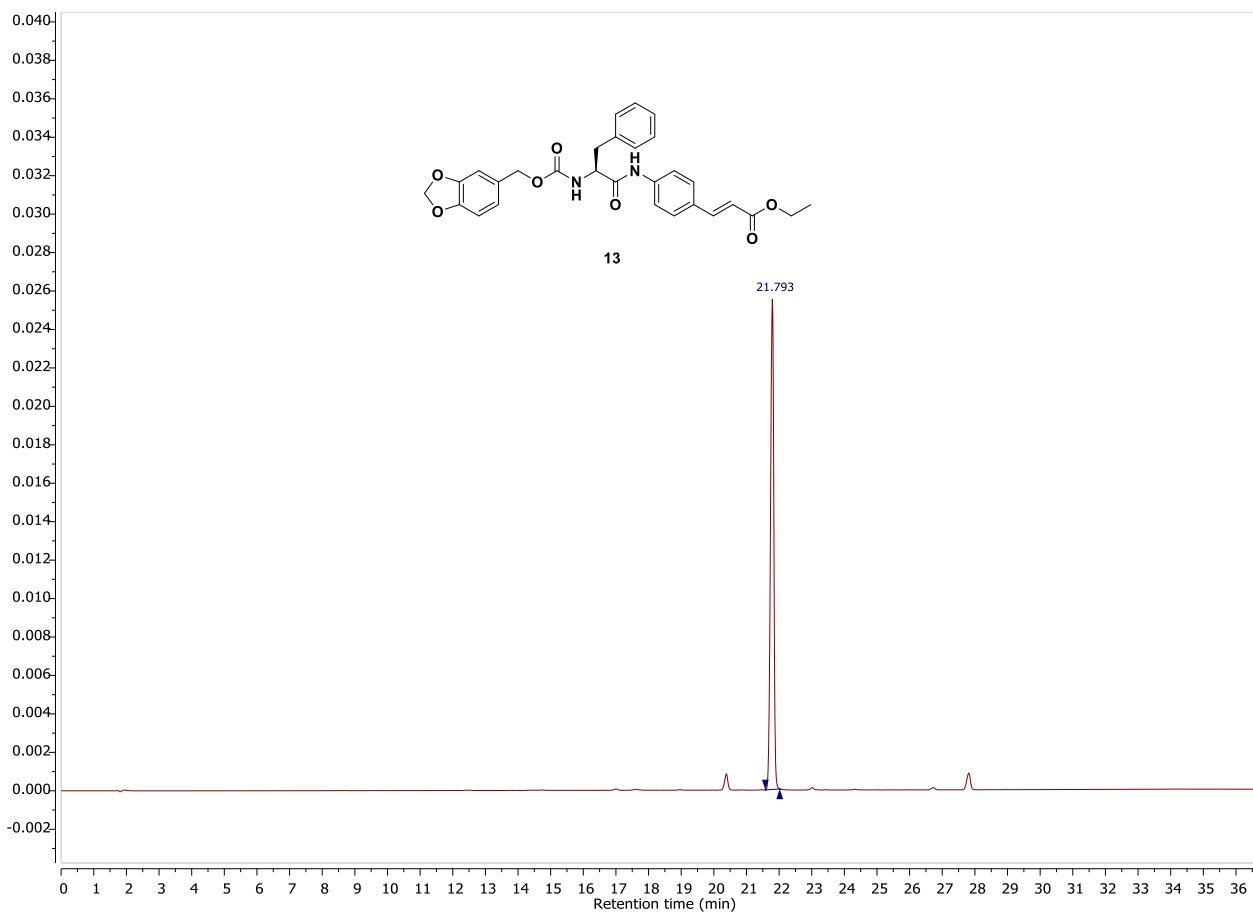


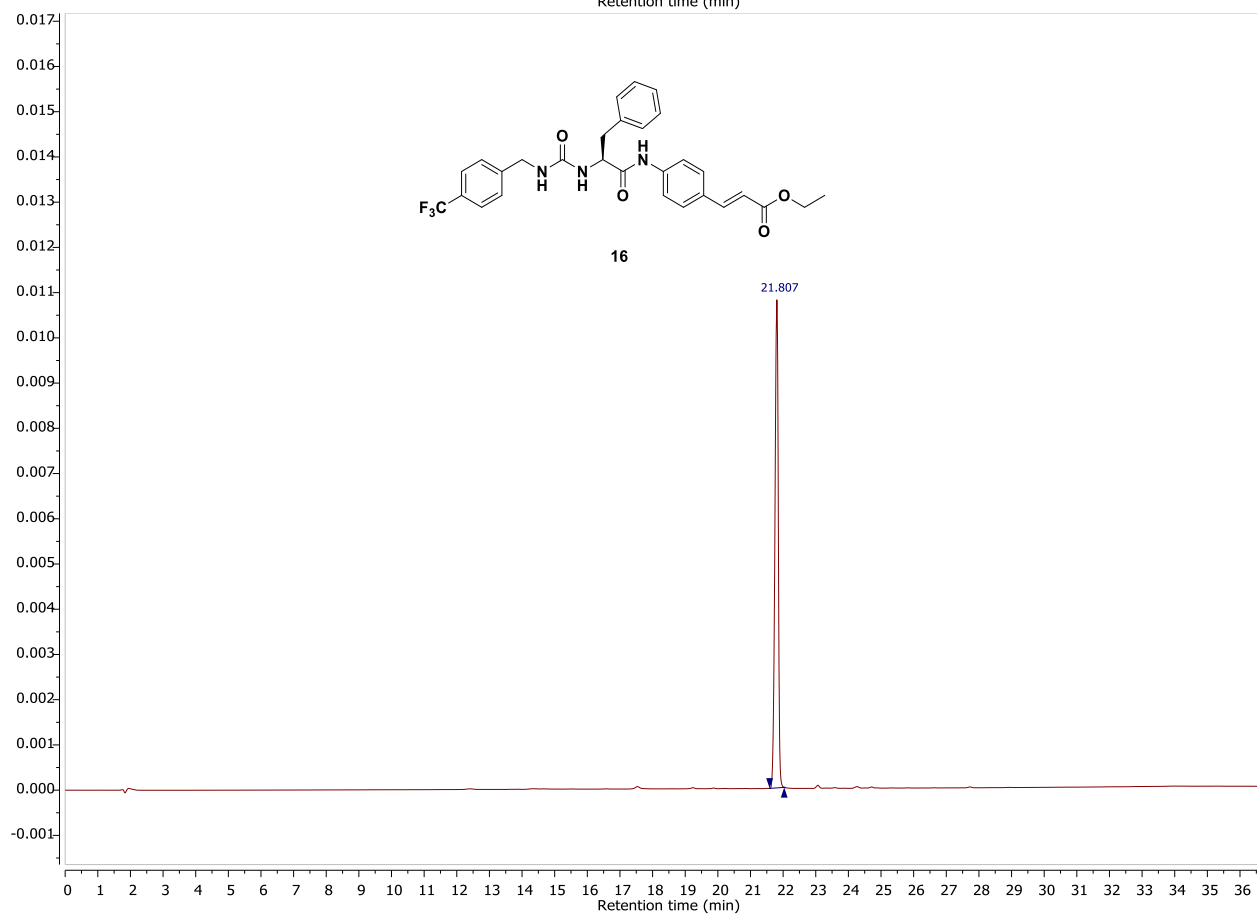
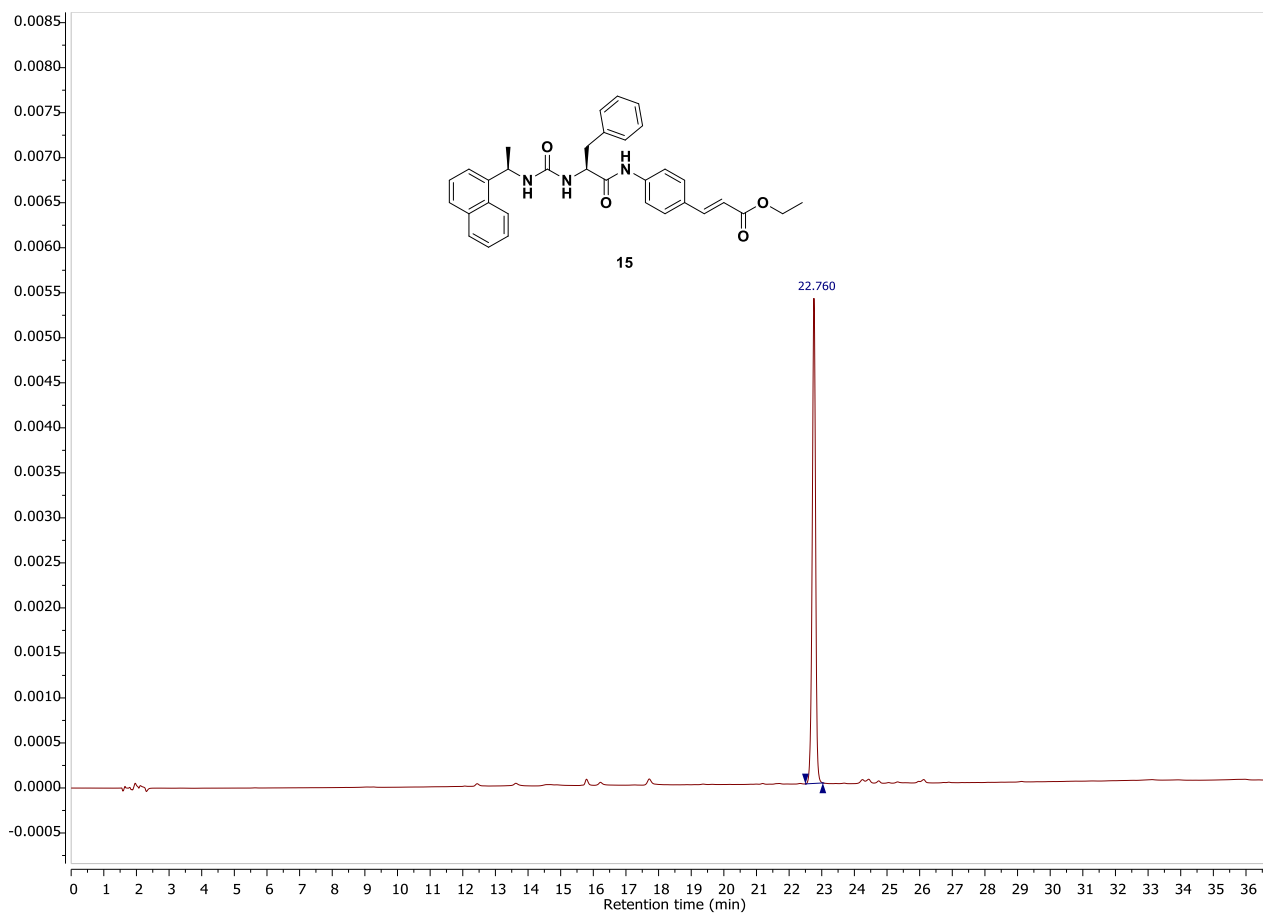


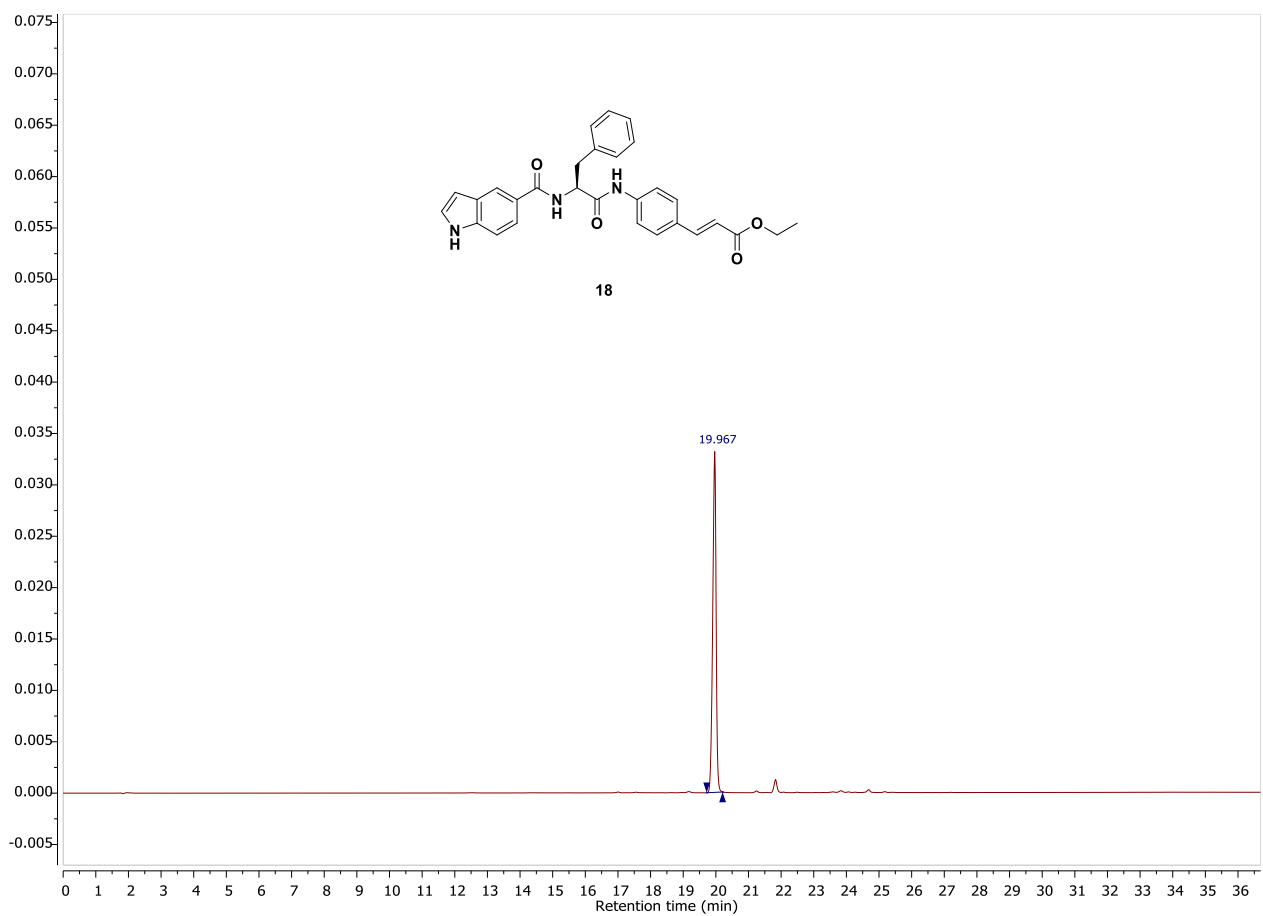
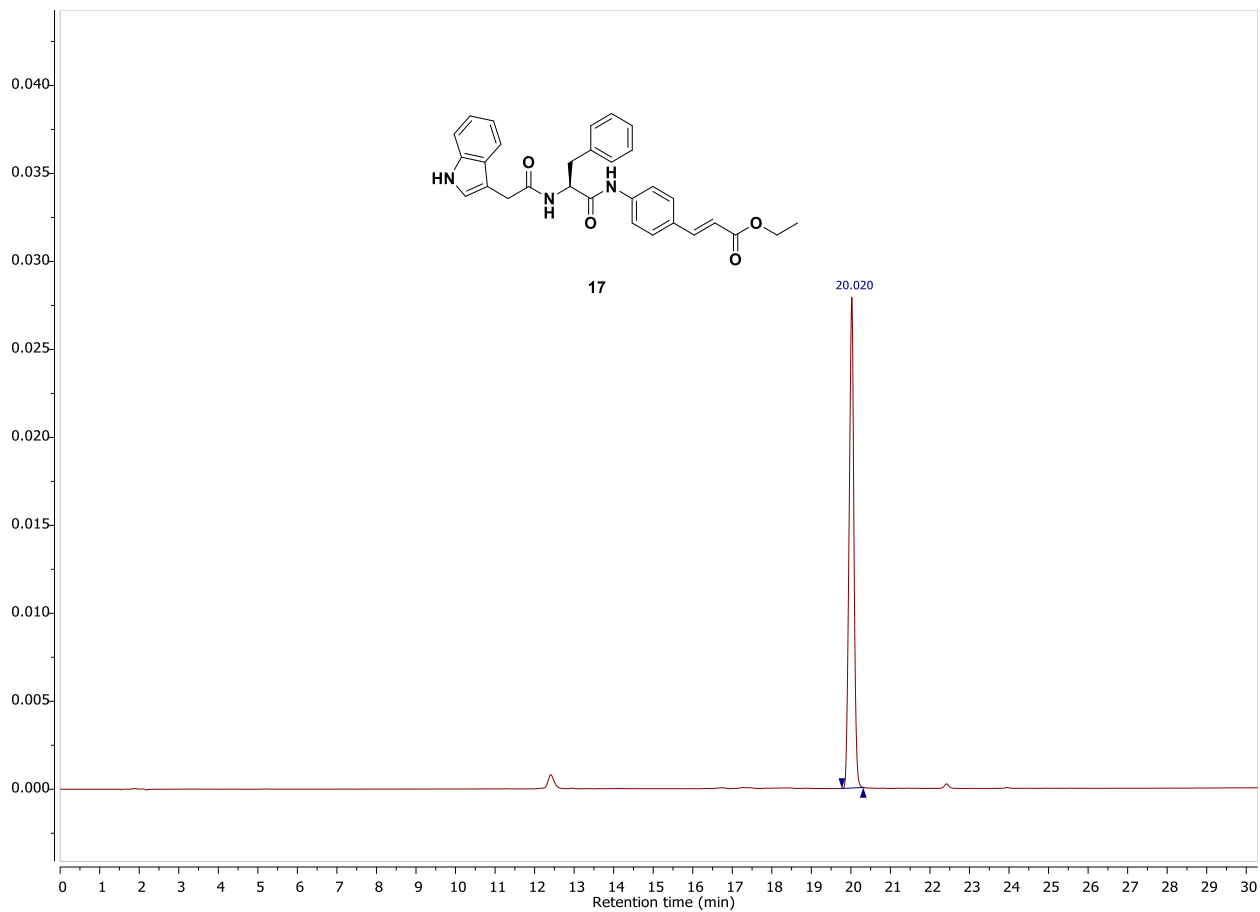
2. HPLC traces of the final compounds

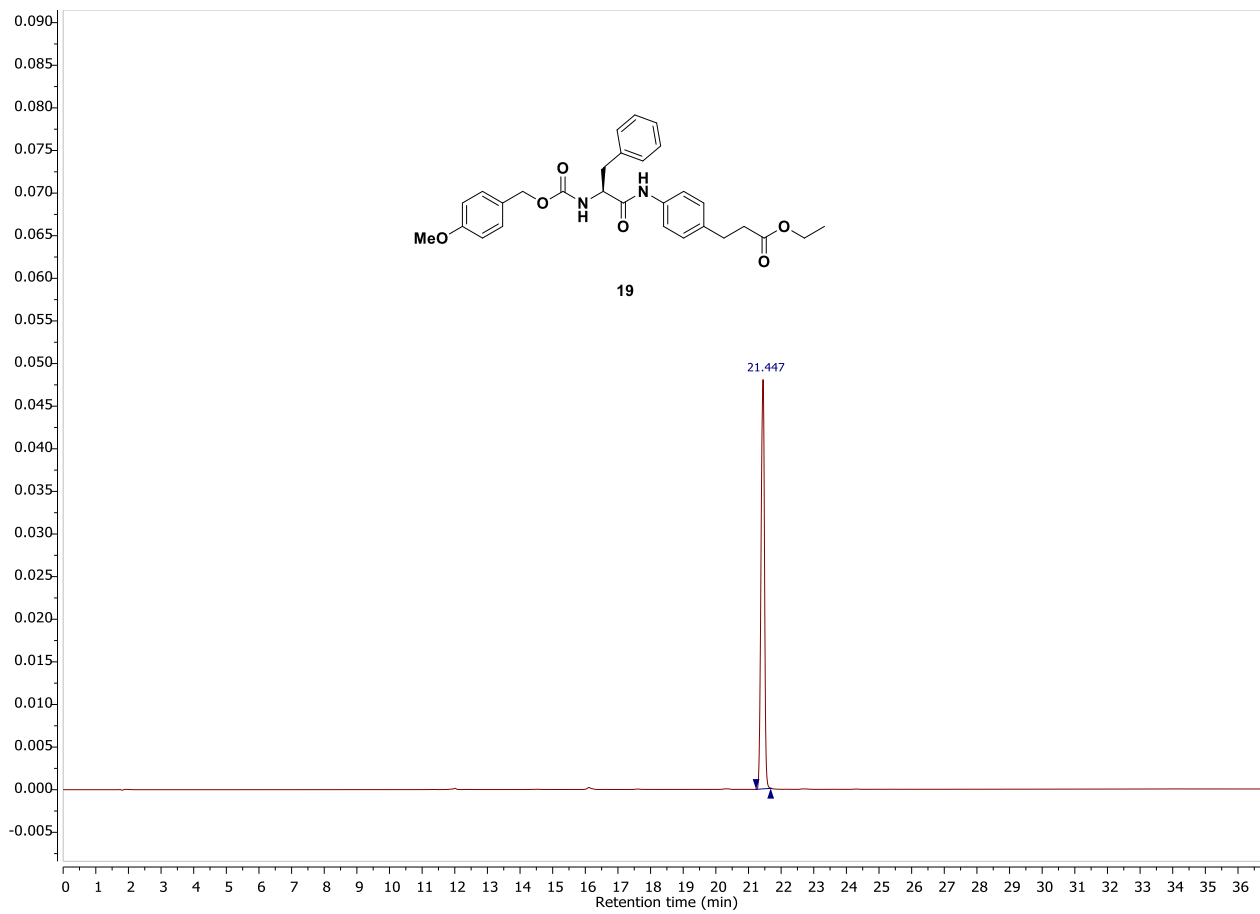












3. Figure S1

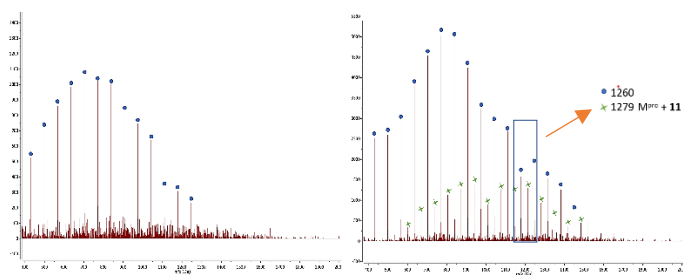


Figure S1. (A) Representative ESI-MS spectrum of a solution containing 0.4 μ M SARS-CoV-2 M^{pro} in water/MeOH (1:1) with 0.1% HCOOH. The spectrum was acquired in positive ion mode. The blue dots correspond to the unmodified protein. (B) Representative ESI-MS spectrum of a mixture containing 0.4 mM SARS-CoV-2 M^{pro} after incubation with **11** in water/MeOH (1:1) with 0.1% HCOOH. The spectrum was acquired in positive ion mode. The blue dots correspond to the unmodified protein, and the green stars correspond to the modified protein.

4. Figure S2

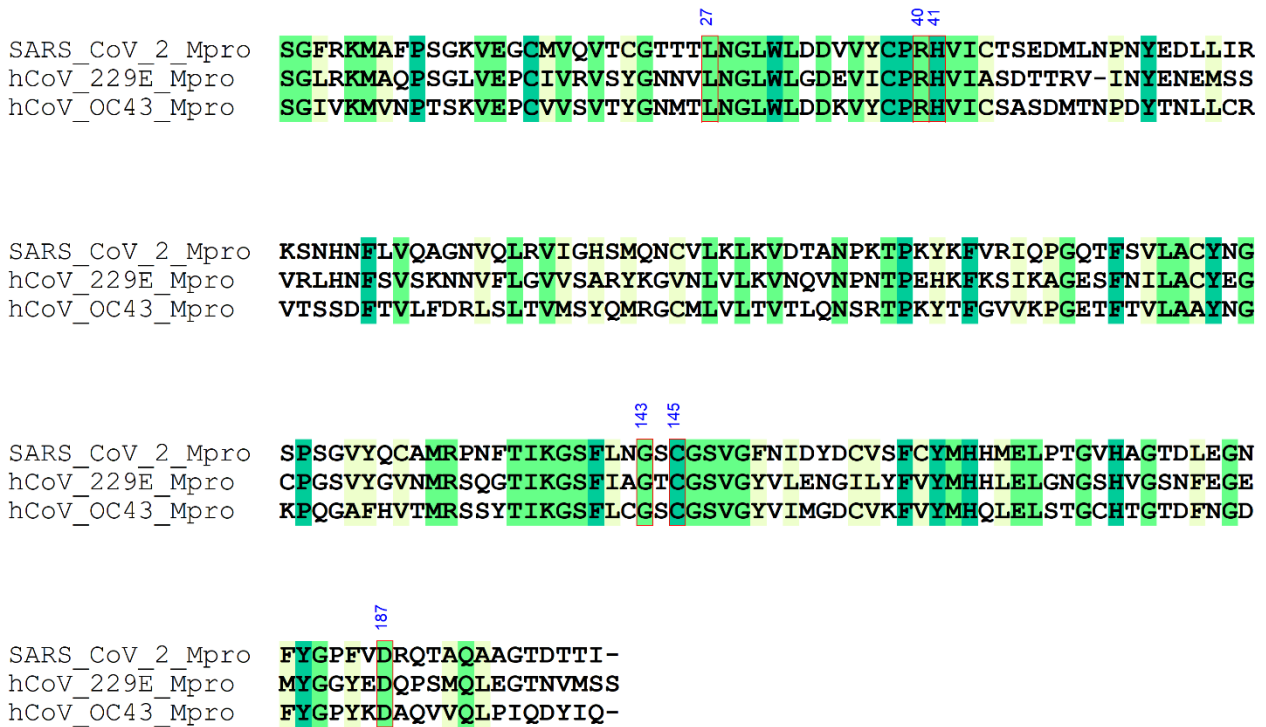


Figure S2. Sequence alignment of SARS-CoV-2 M^{pro}, hCoV-229E M^{pro}, and hCoV-OC43 M^{pro}. Conserved residues are labeled in dark green, the same residues are indicated in lighter green, and residues with similar properties are yellow. The key residues are highlighted by a red square and the number of sequence positions in navy blue.

Table S1. Antiviral activity of cinnamic esters against hCoV-229E and hCoV-OC43. Notes: EC₅₀ was measured for hCoV-229E by evaluating the residual MRC-5 cell viability as a surrogate of viral CPE, or by focus forming reduction assay (FFRA) against hCoV-OC43 in HCT-8 cells.

Compound	hCoV-229E EC₅₀ (μM)	hCoV-OC43 EC₅₀ (μM)
11	> 50	> 50
12	5.27 ± 0.25	> 50
13	> 50	> 50
15	> 50	> 50
17	> 50	9.14 ± 0.70
18	> 50	10.1 ± 0.17
19	> 50	> 50