

Identification of a Dual Inhibitor of Secreted Phospholipase A₂ (GIIA sPLA₂) and SARS-CoV-2 Main Protease

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Table S1. Structural details of structures 1-7 concerning the configuration of 2-carbon and the dihedral angle $C\beta_{C145}-S_{C145}-C2-O2$ related to the covalent bond formatted between Cys145 and 2-oxoamide moiety. Molecular Dynamics RMSDs for the protein and the ligand along with MM-GBSA Binding Energy.

Structure	Configuration of 2-carbon	MM-GBSA Binding Energy (kcal/mol)	Average Protein RMSD \pm StDev	Average Ligand RMSD \pm StDev	Major Interactions	Dihedral Angle $C\beta-S-C2-O2$
1	<i>R</i>	-58.69	2.84 \pm 0.48	3.36 \pm 0.95	S144	-73.68
2	<i>R</i>	-56.37	1.77 \pm 0.24	4.63 \pm 0.70	E166	-175.218
3	<i>S</i>	-54.51	1.73 \pm 0.28	7.36 \pm 3.73	-	86.525
4	<i>S</i>	-47.03	2.29 \pm 0.35	5.53 \pm 0.89	T26, T24	-45.037
5	<i>R</i>	-46.41	2.00 \pm 0.32	3.97 \pm 1.10	H41, E166	-50.033
6	<i>S</i>	-45.17	1.95 \pm 0.31	5.05 \pm 0.95	H41, N142, H164	134.808
7	<i>S</i>	-43.26	1.79 \pm 0.26	3.22 \pm 0.79	T26, H41, N142	-55.195

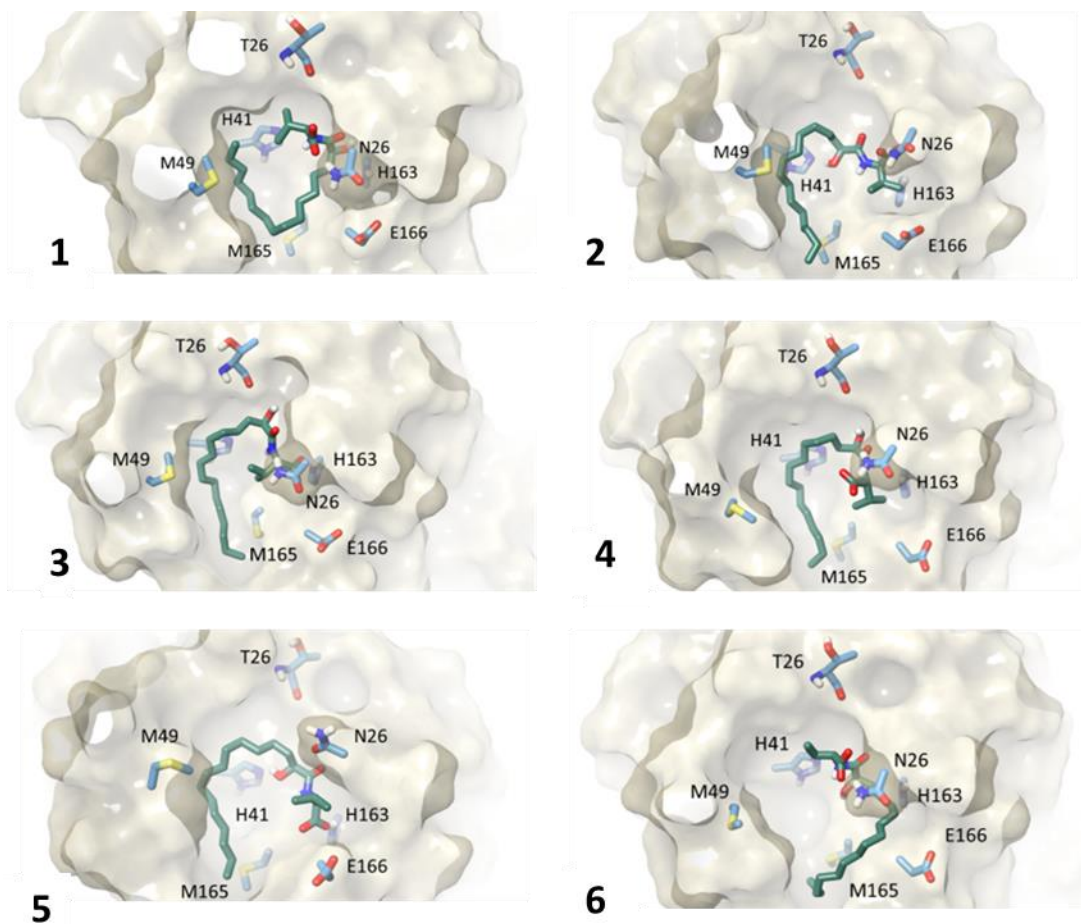


Figure S1. Structural details for the M^{pro} -GK241 interactions in structures 1-6 resulted by covalent docking calculations.

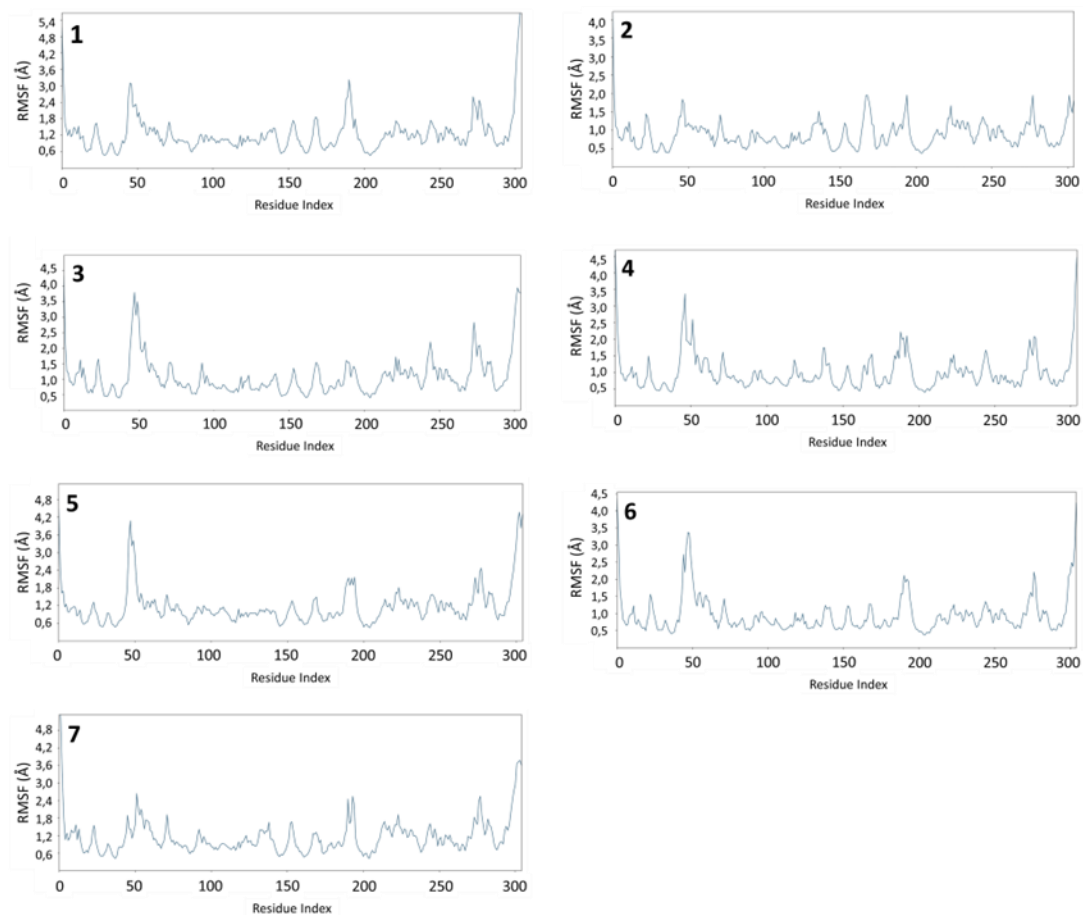


Figure S2. Protein residues Root Mean Square Fluctuation during the 50 nsec MD simulation showing local changes among the protein chain.

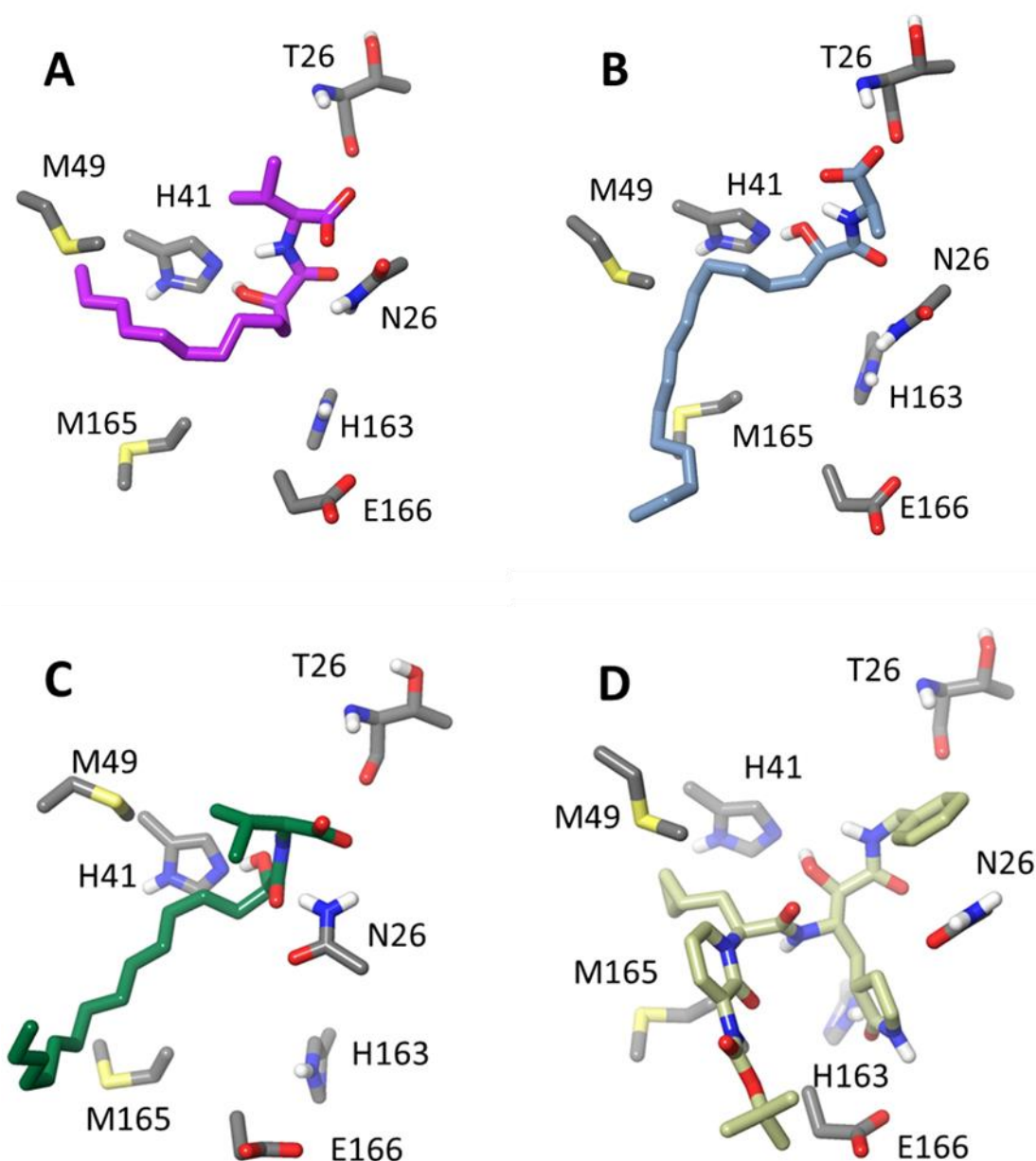
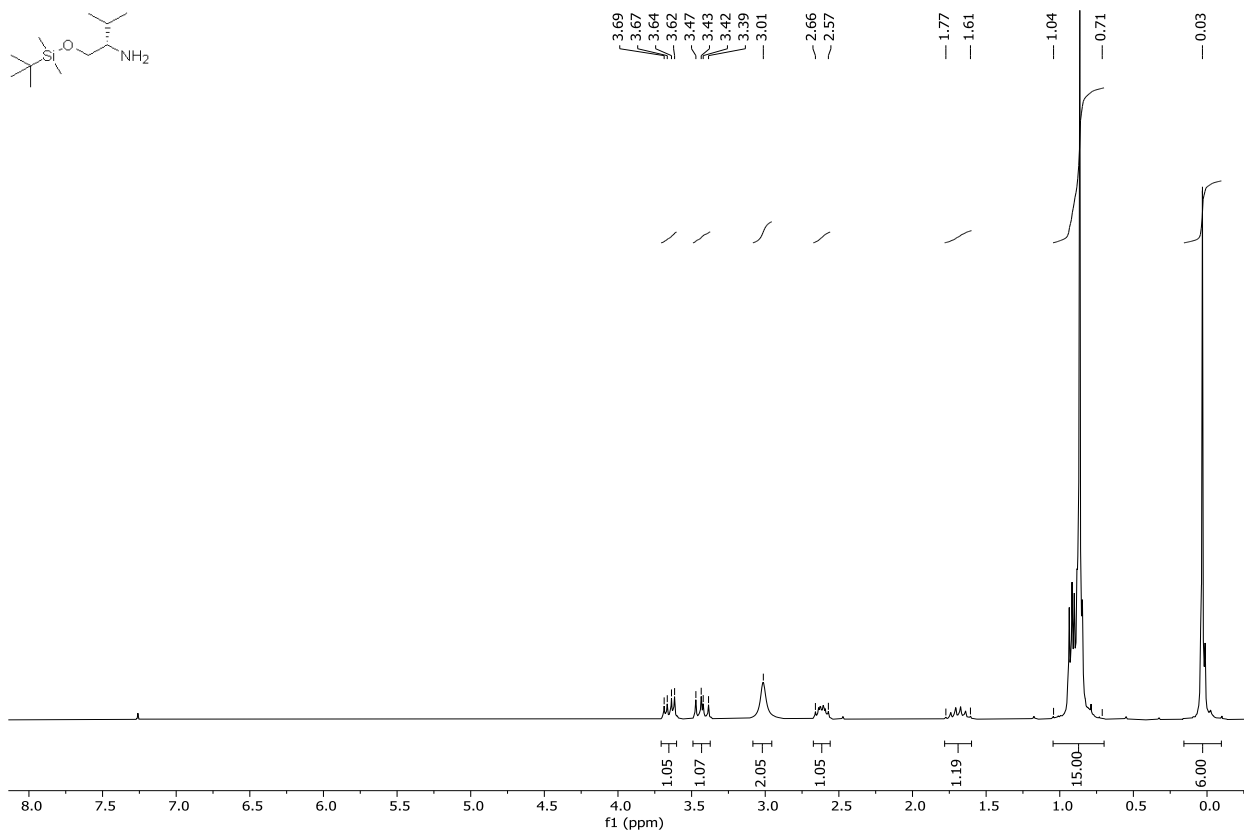
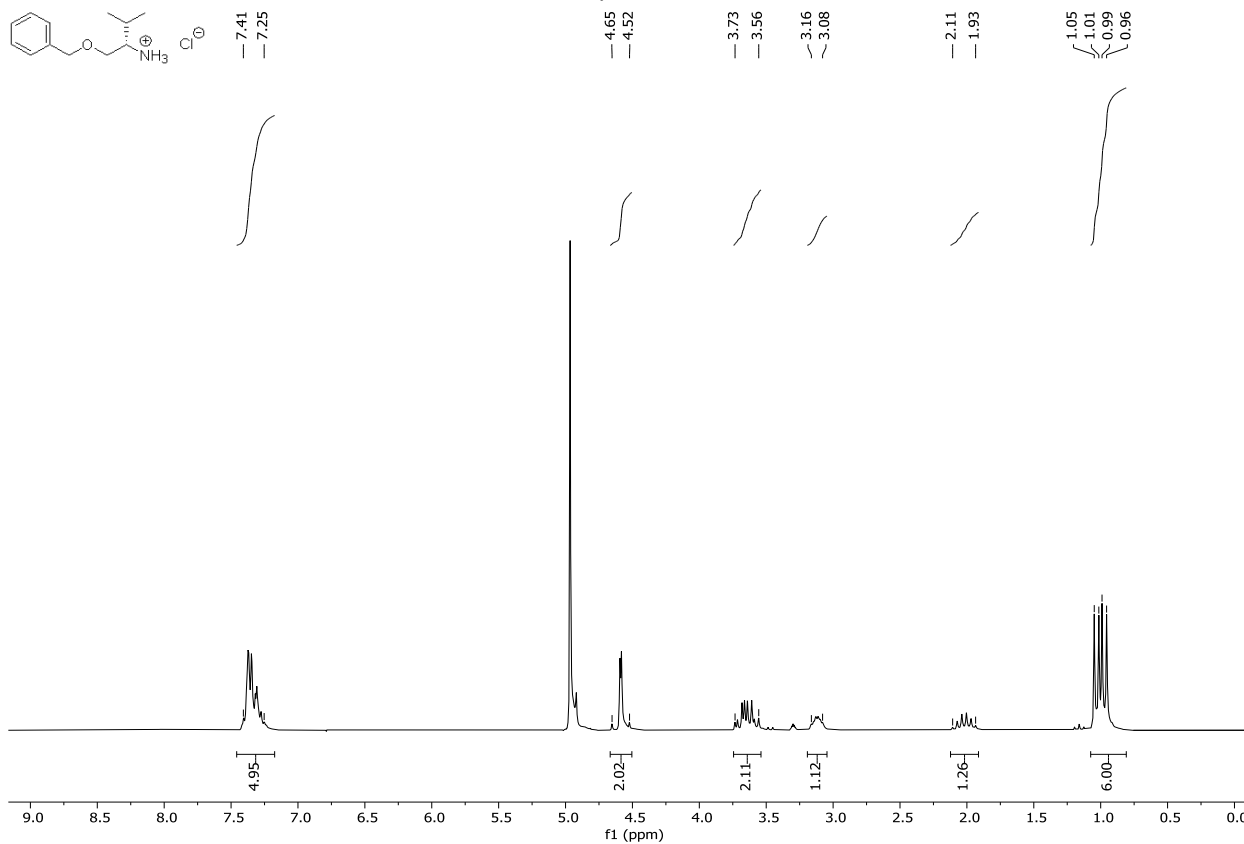


Figure S3. Structures of **8d** (A) in magenta and **8a** (B) in blue bound to M^{pro} in comparison with GK241 (C) in green and crystal structure PDB 6Y2F (D) in chaki. The side chain in derivative **8d** adopts a curved conformation exploring part of S1 and S2 protease cavities. Derivative **8a** aliphatic chain adopts a very similar structural arrangement to GK241, however the alanine moiety appears to interact weakly with N142 and T26.

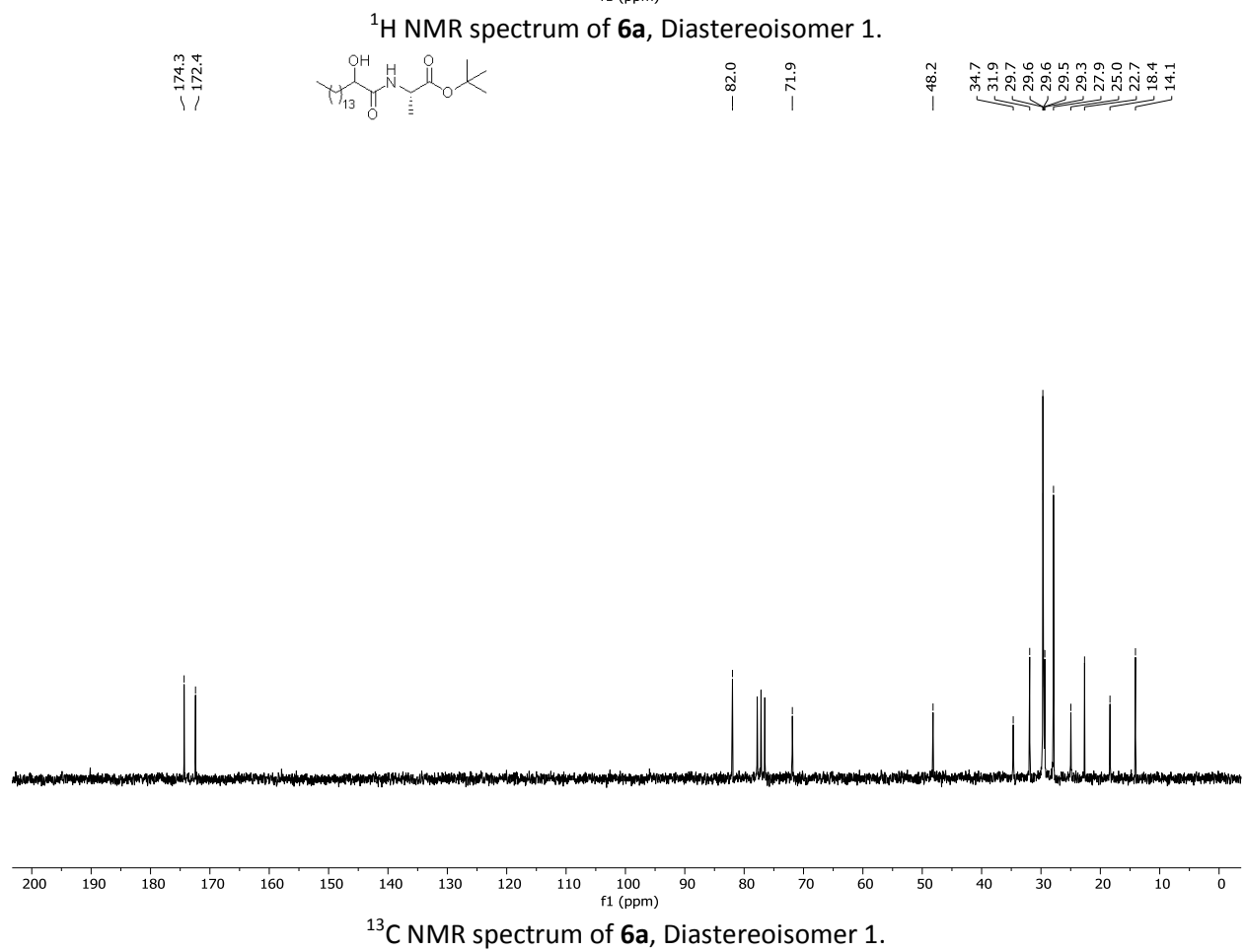
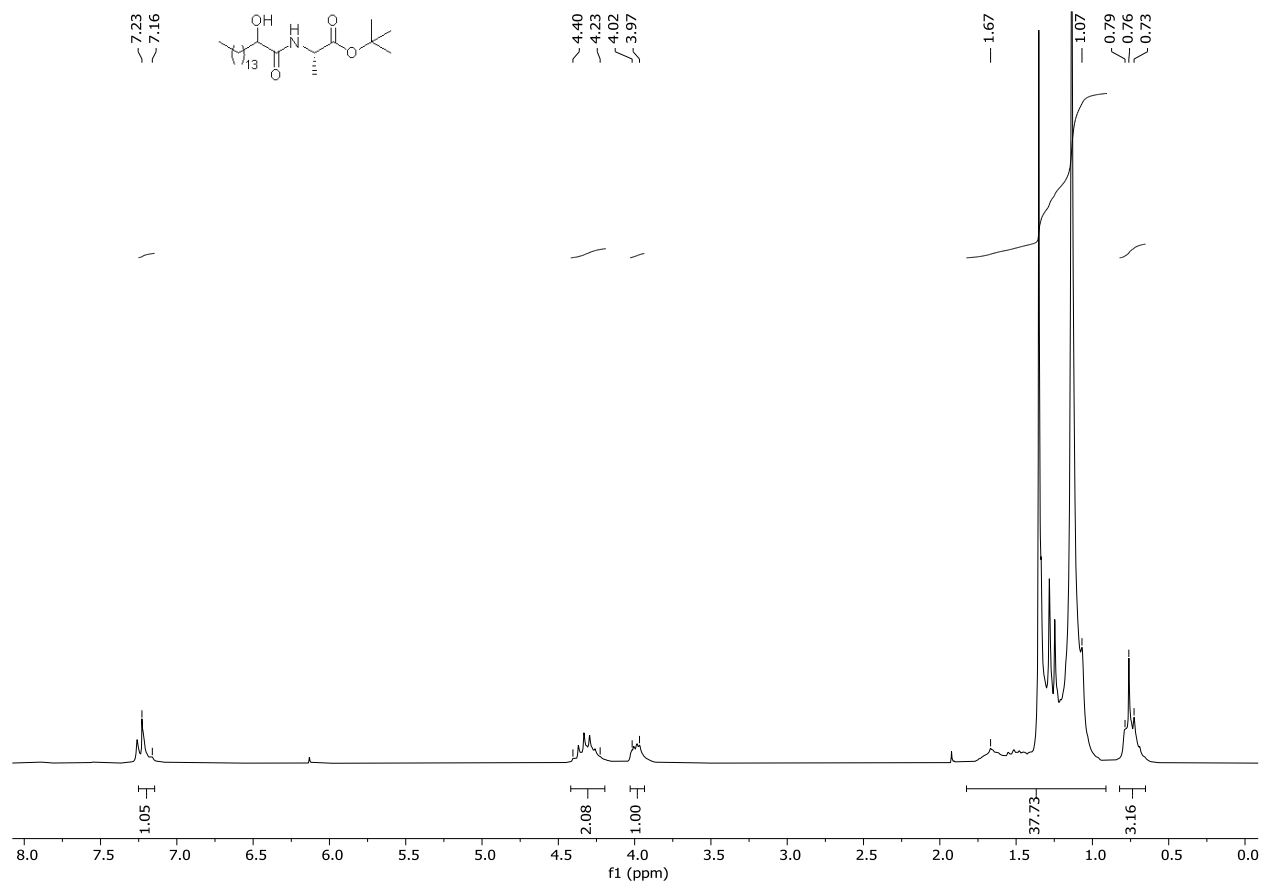
NMR Spectra

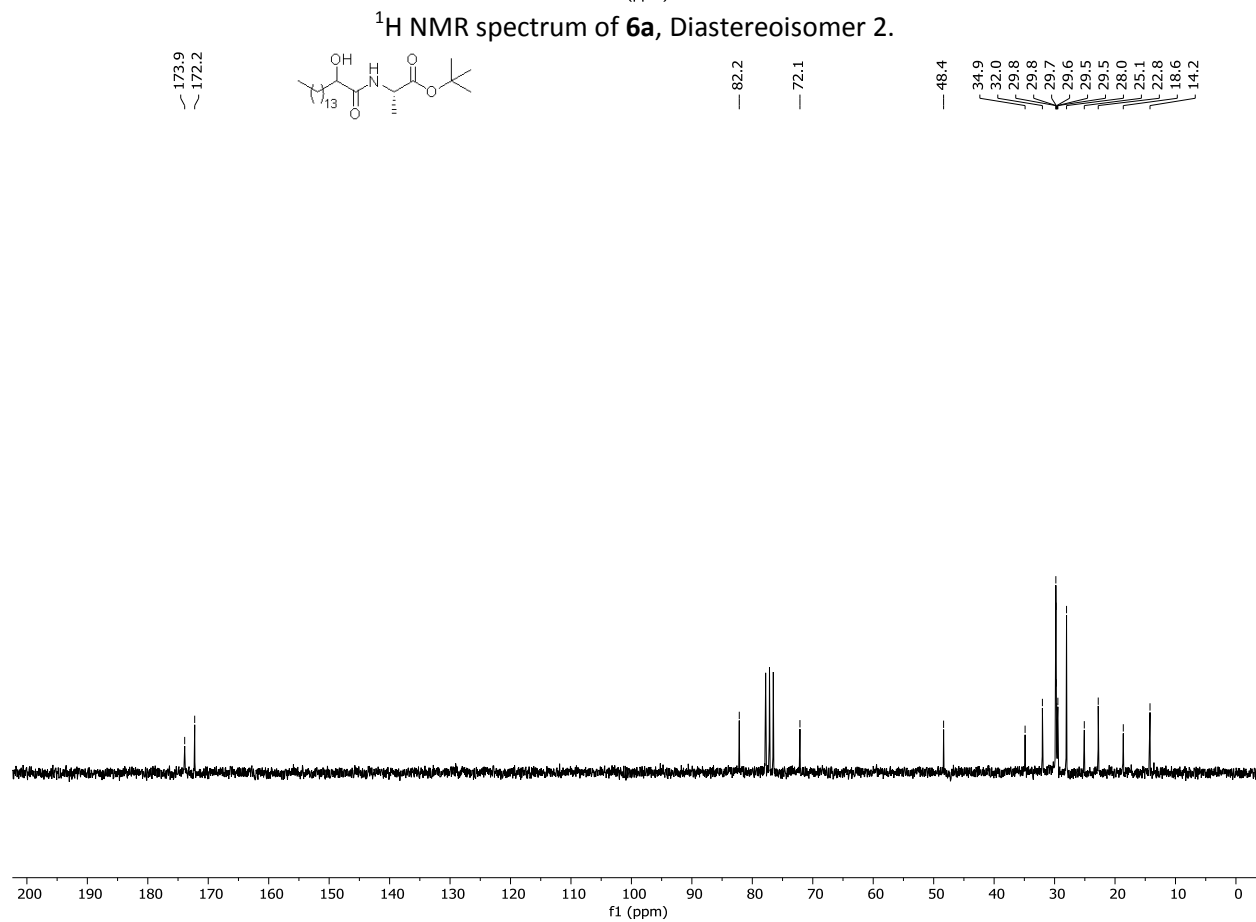
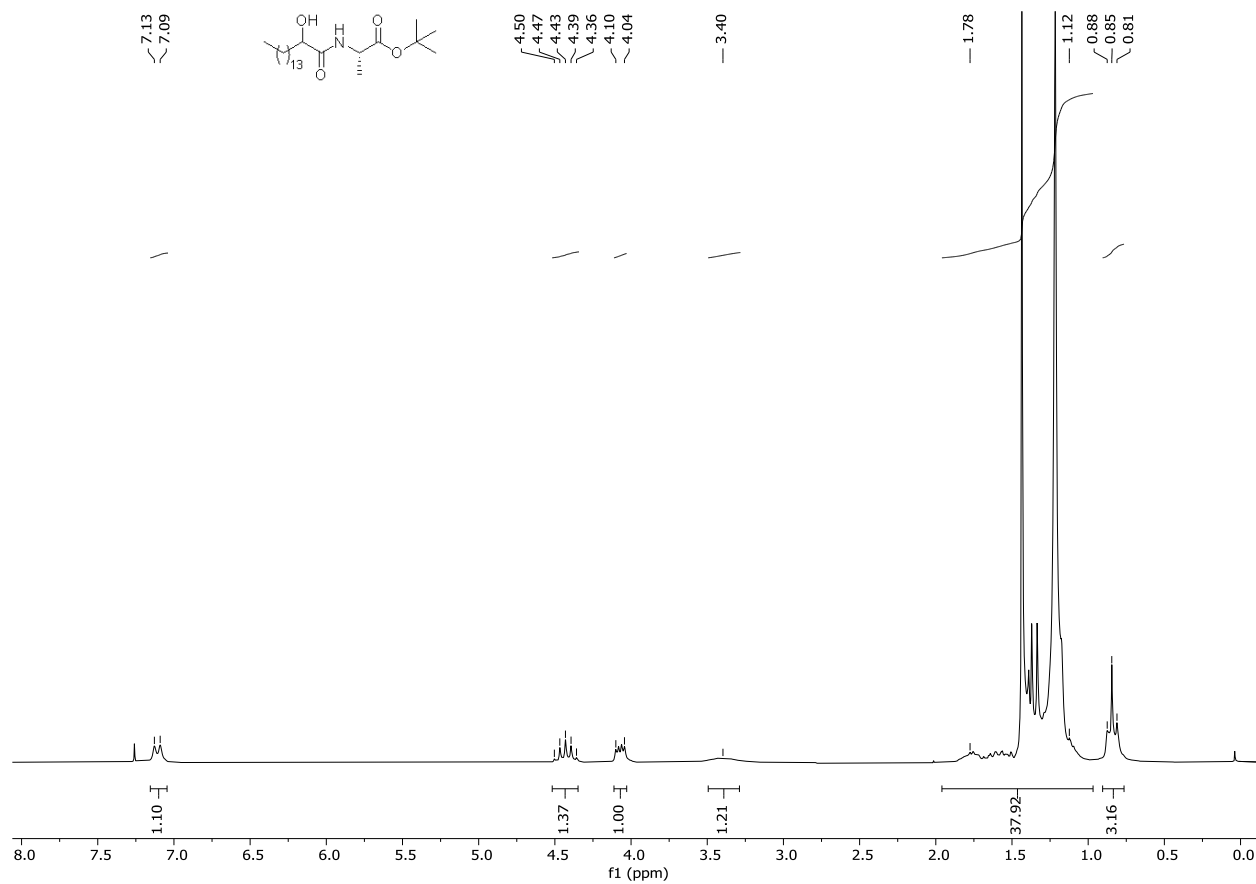


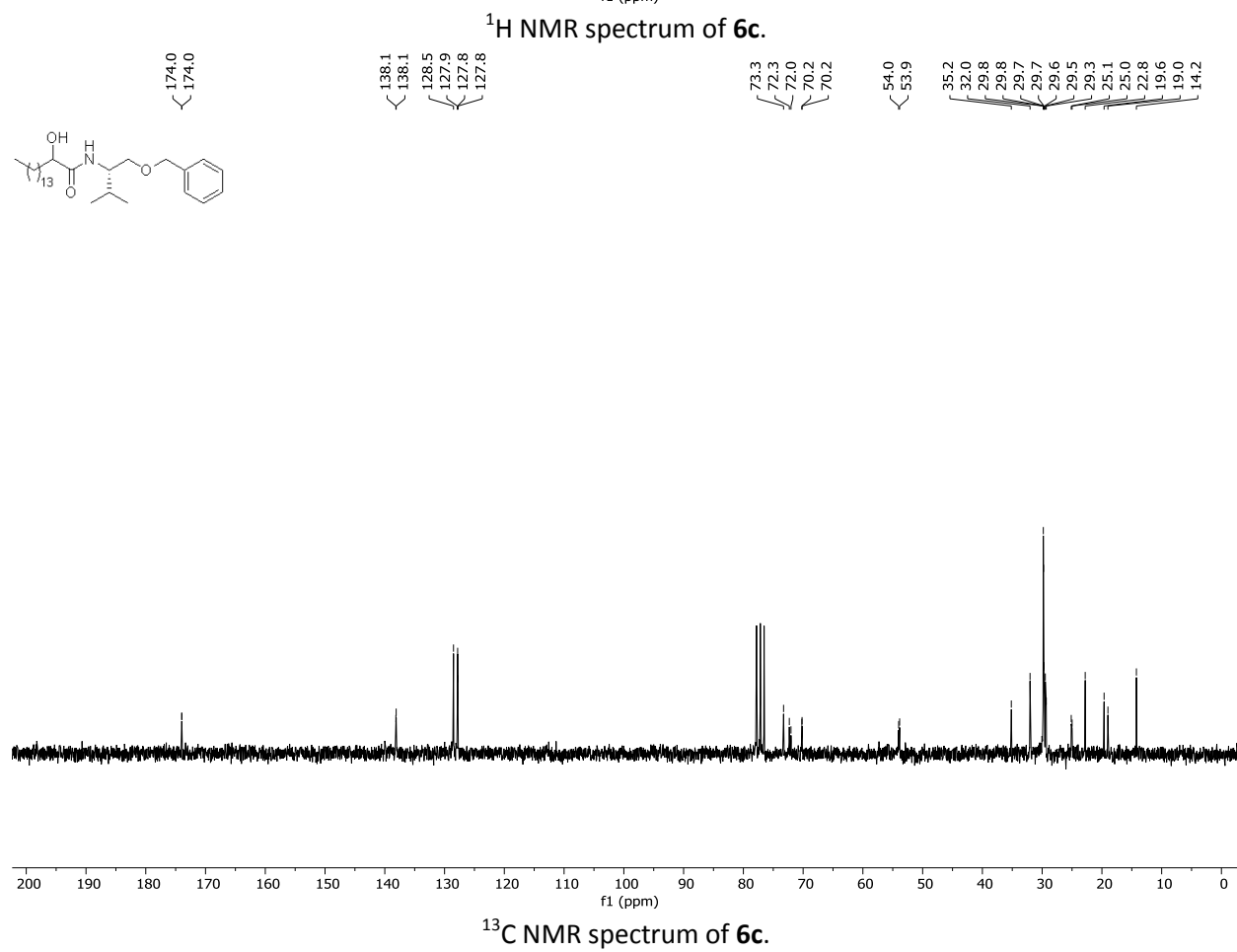
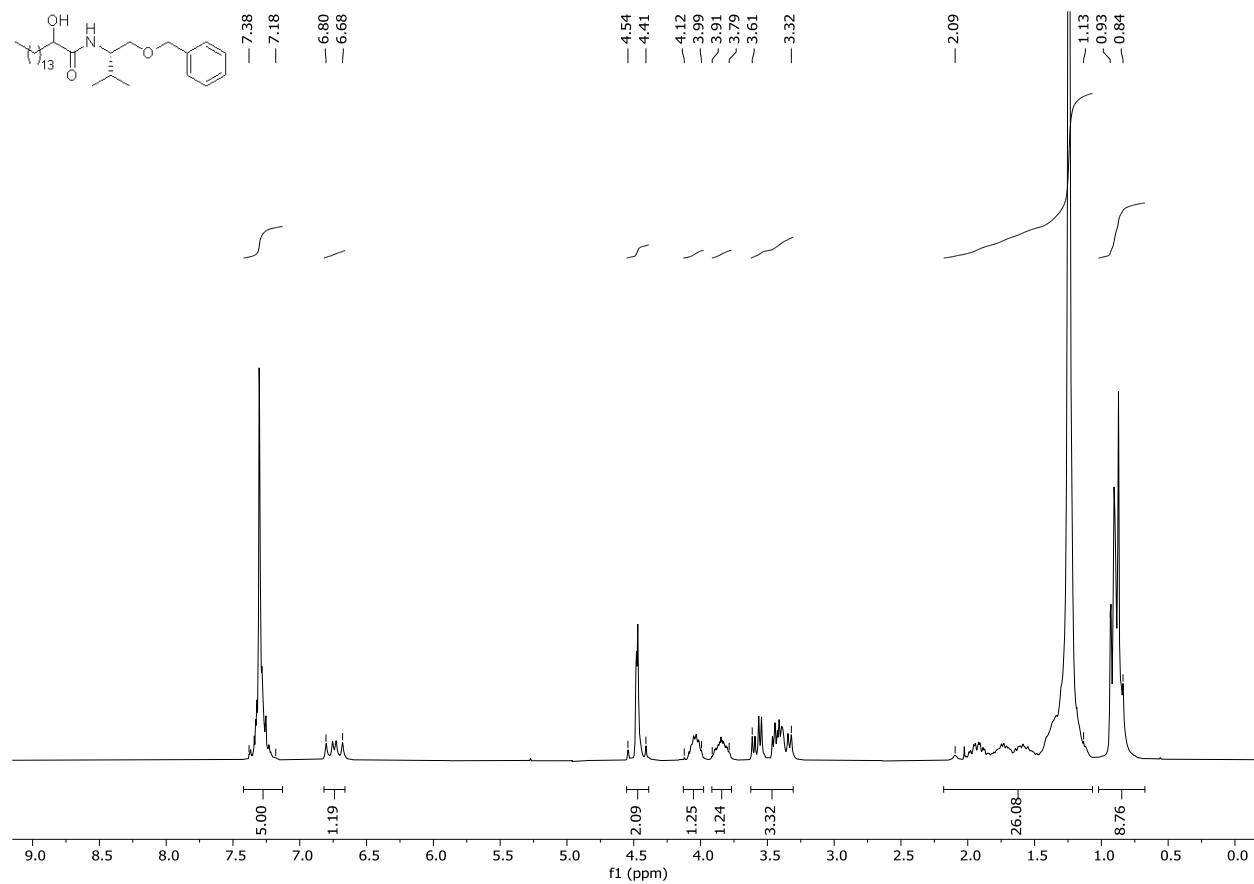
¹H NMR spectrum of **5b**.

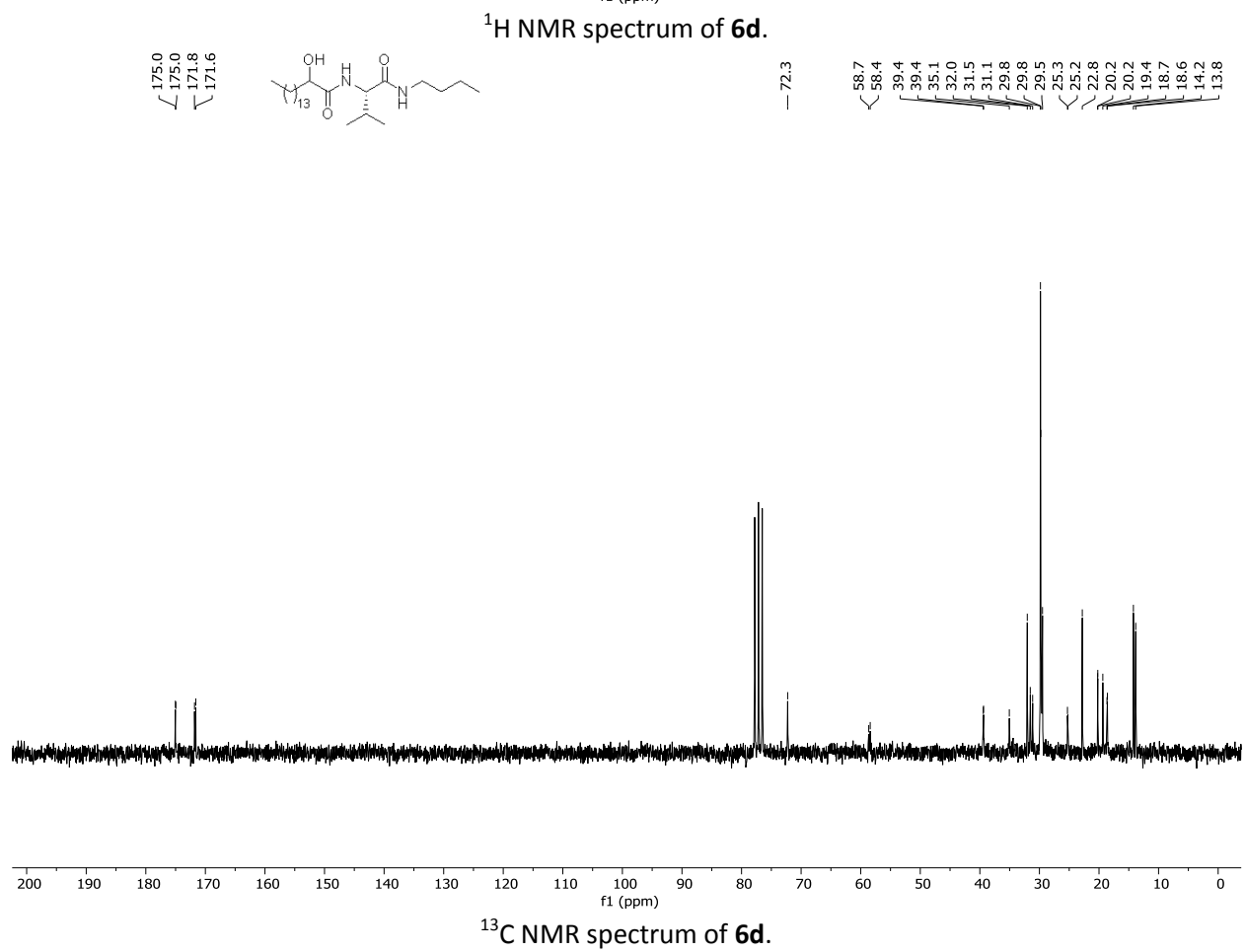
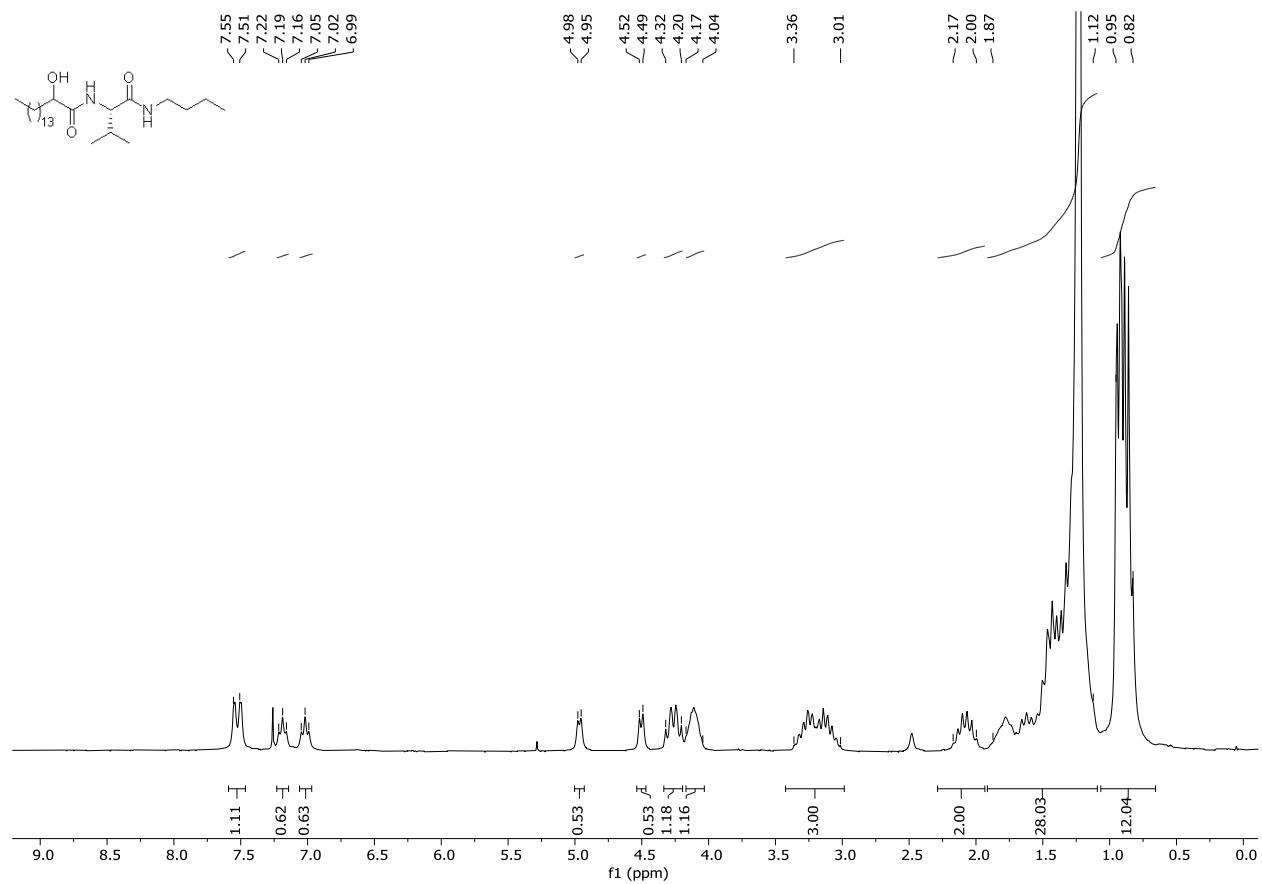


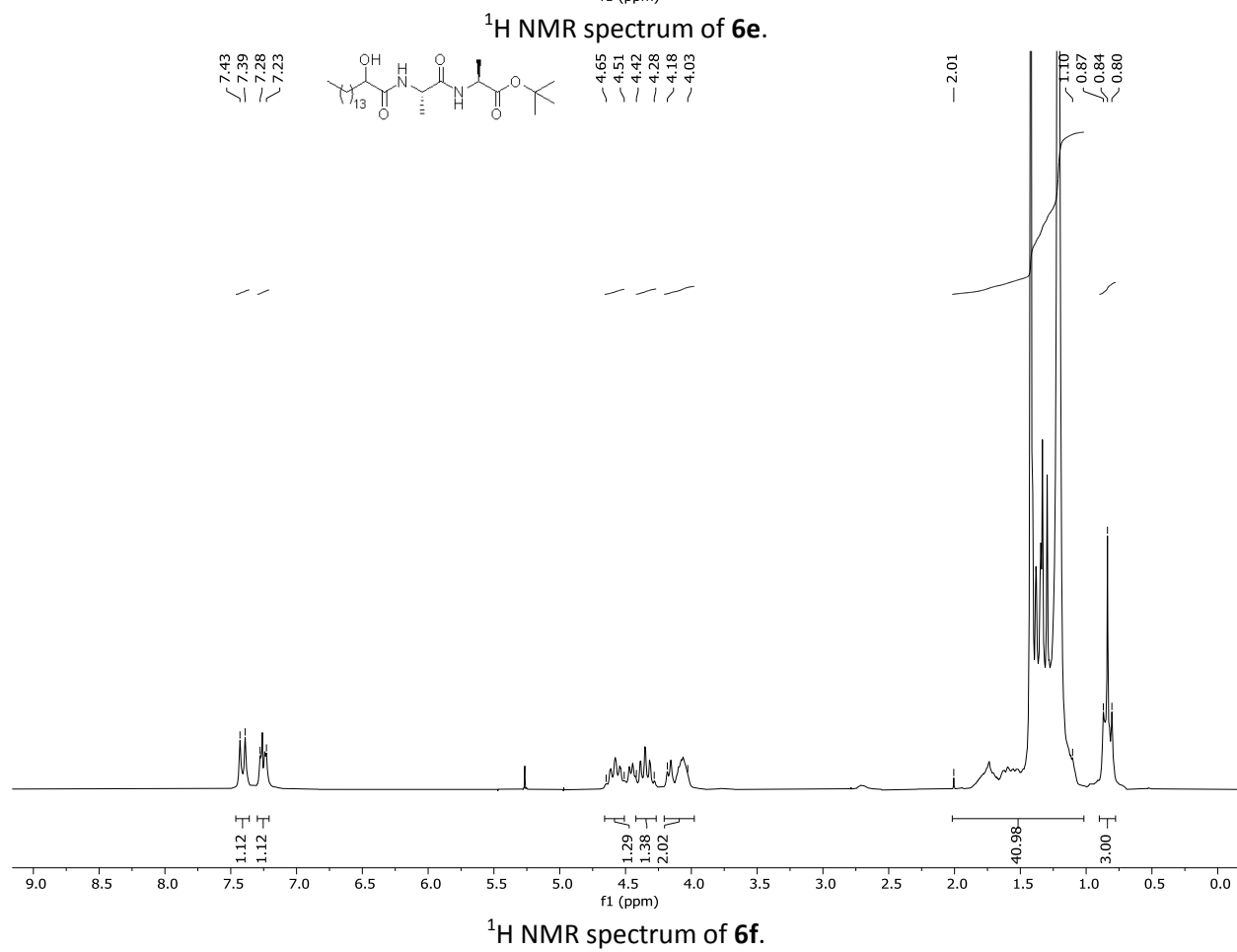
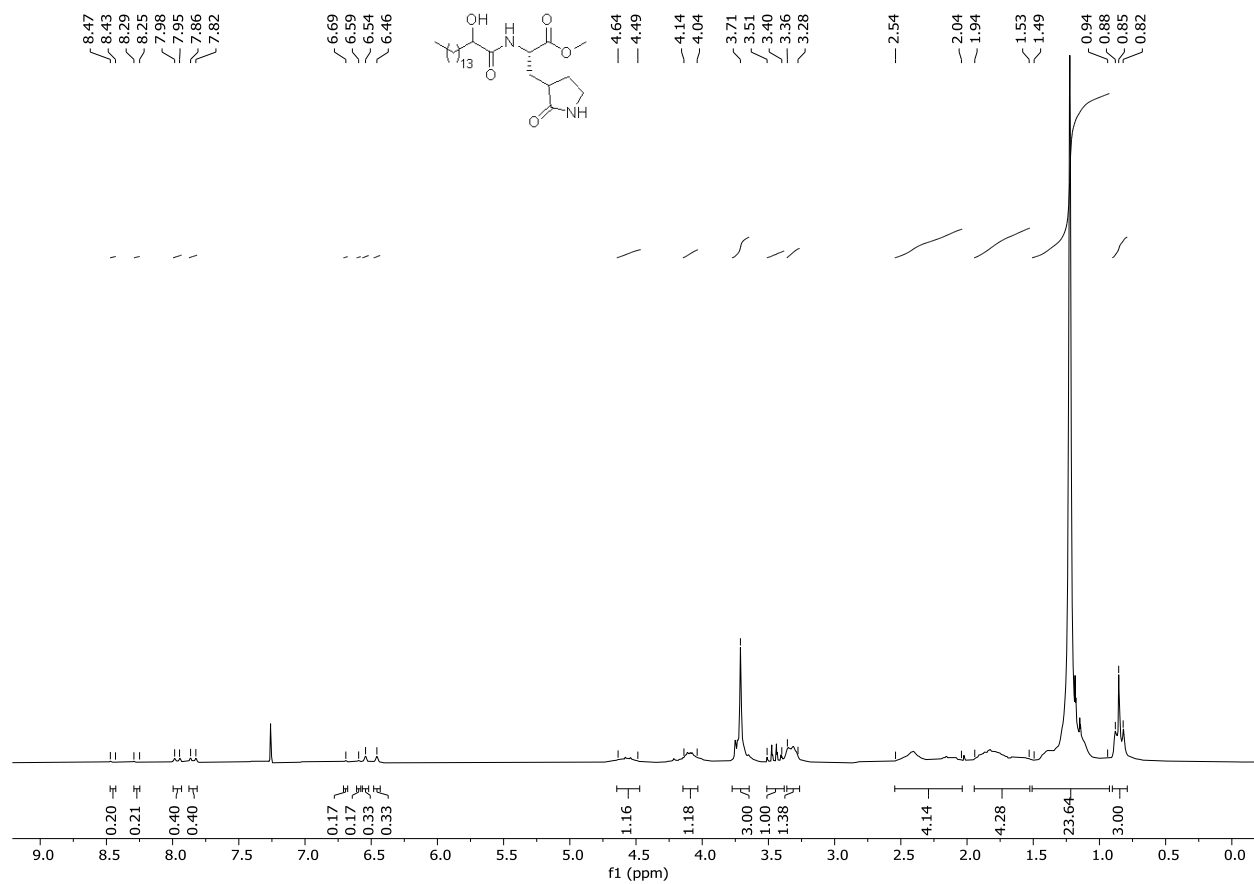
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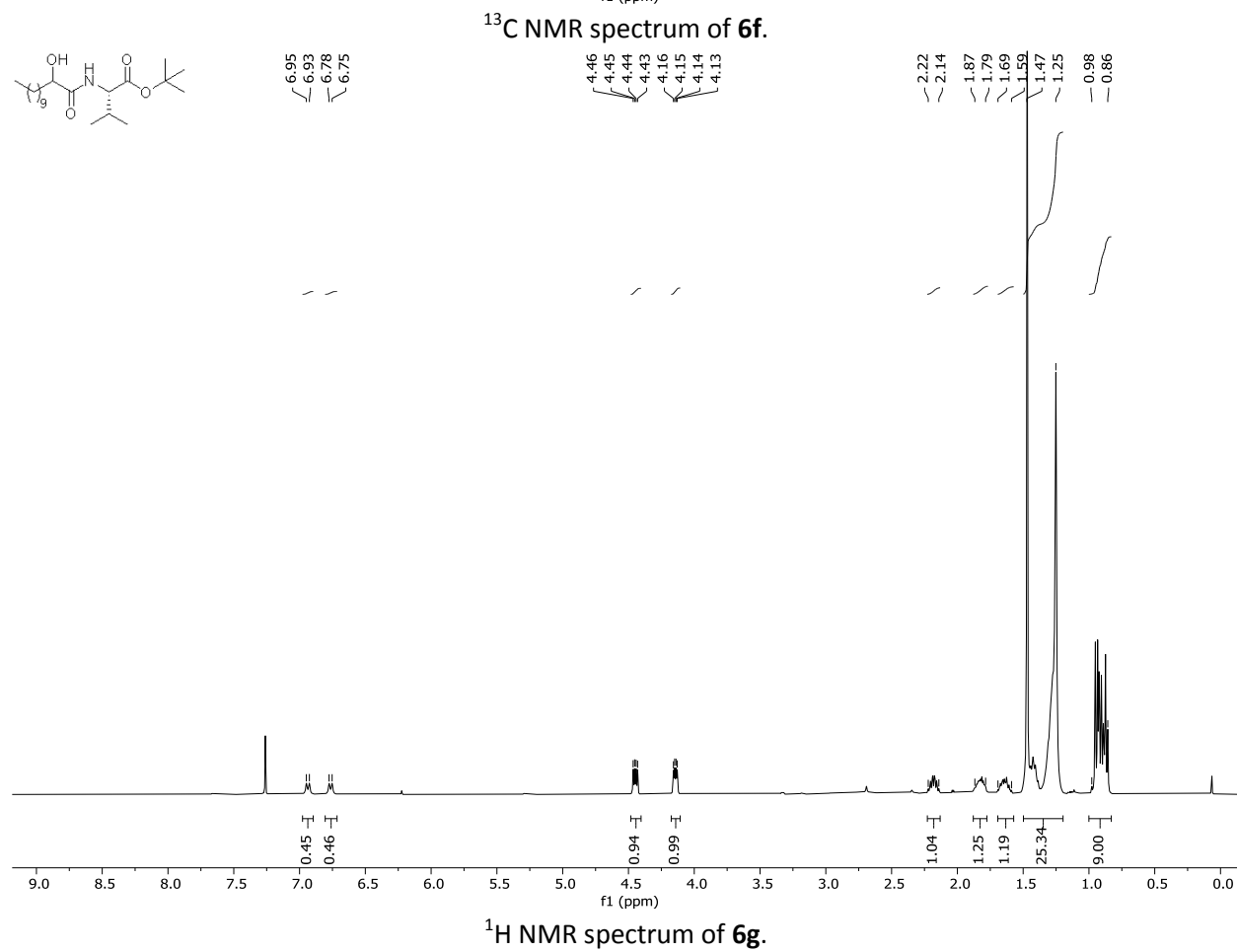
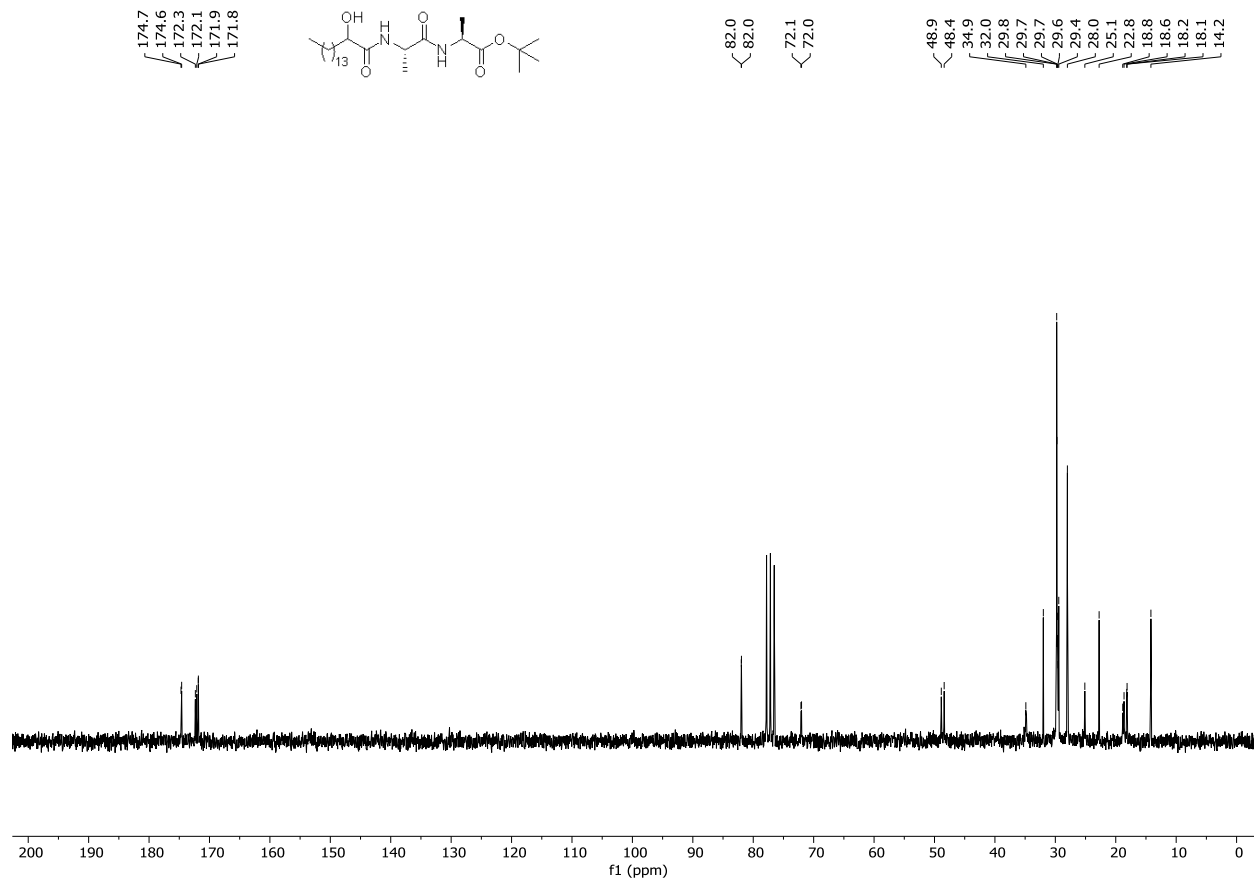


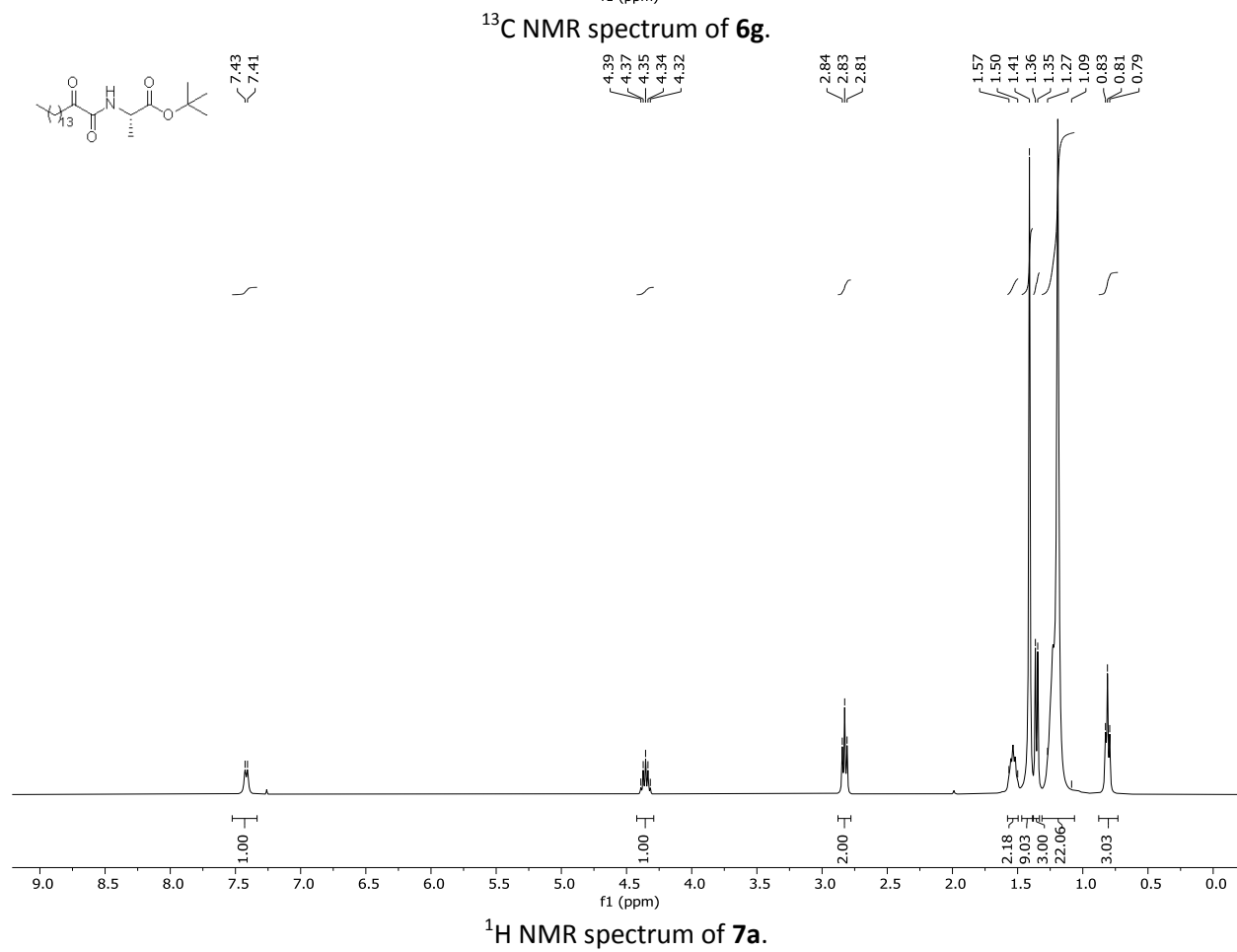
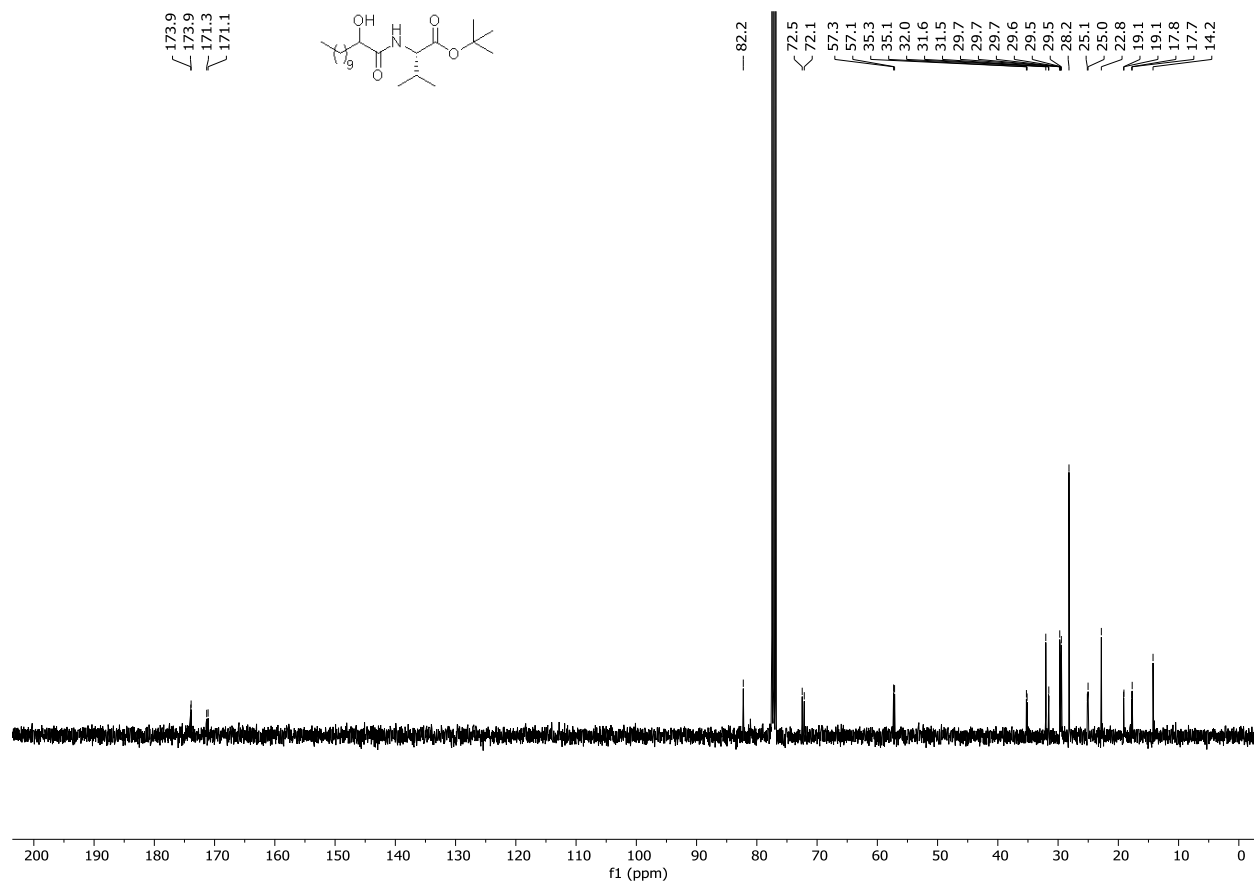


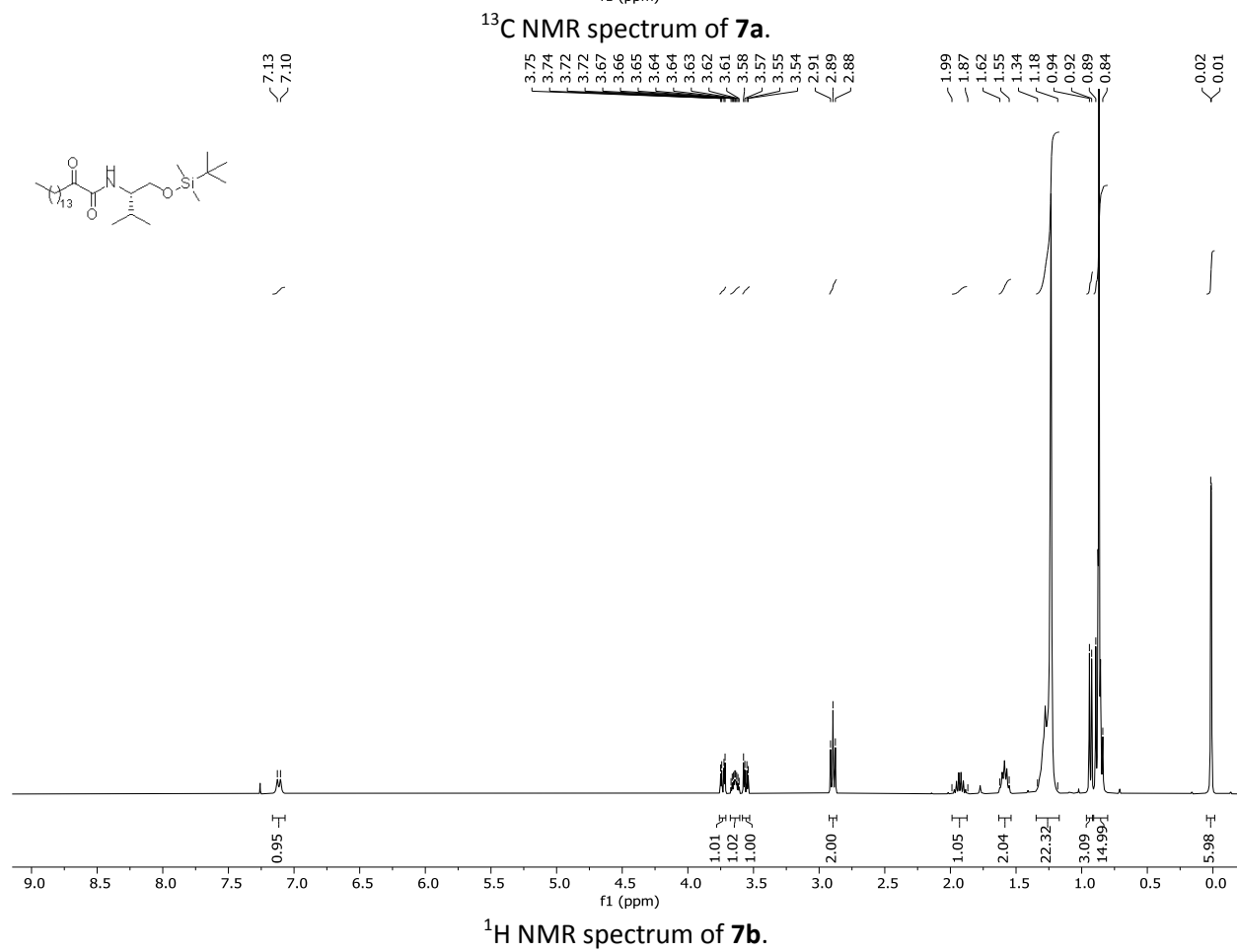
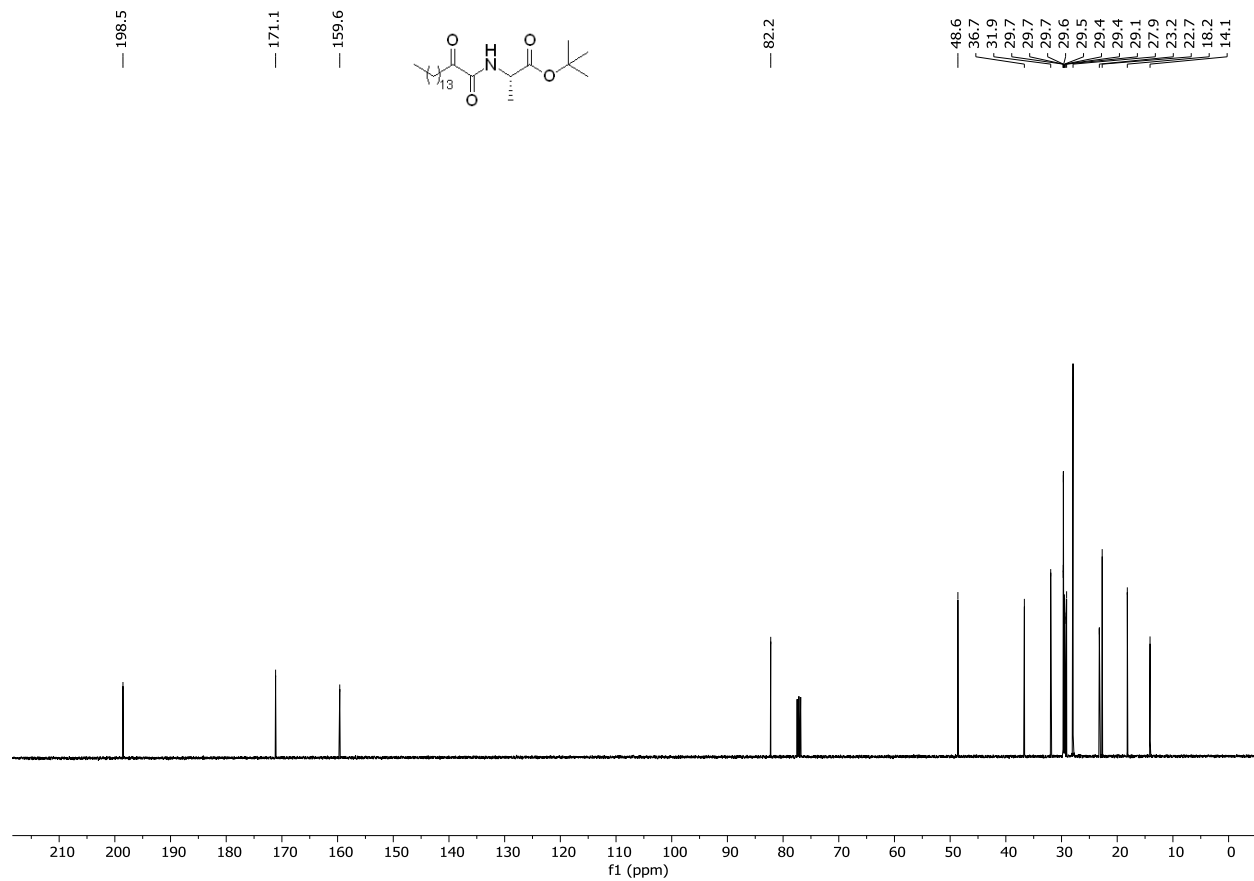


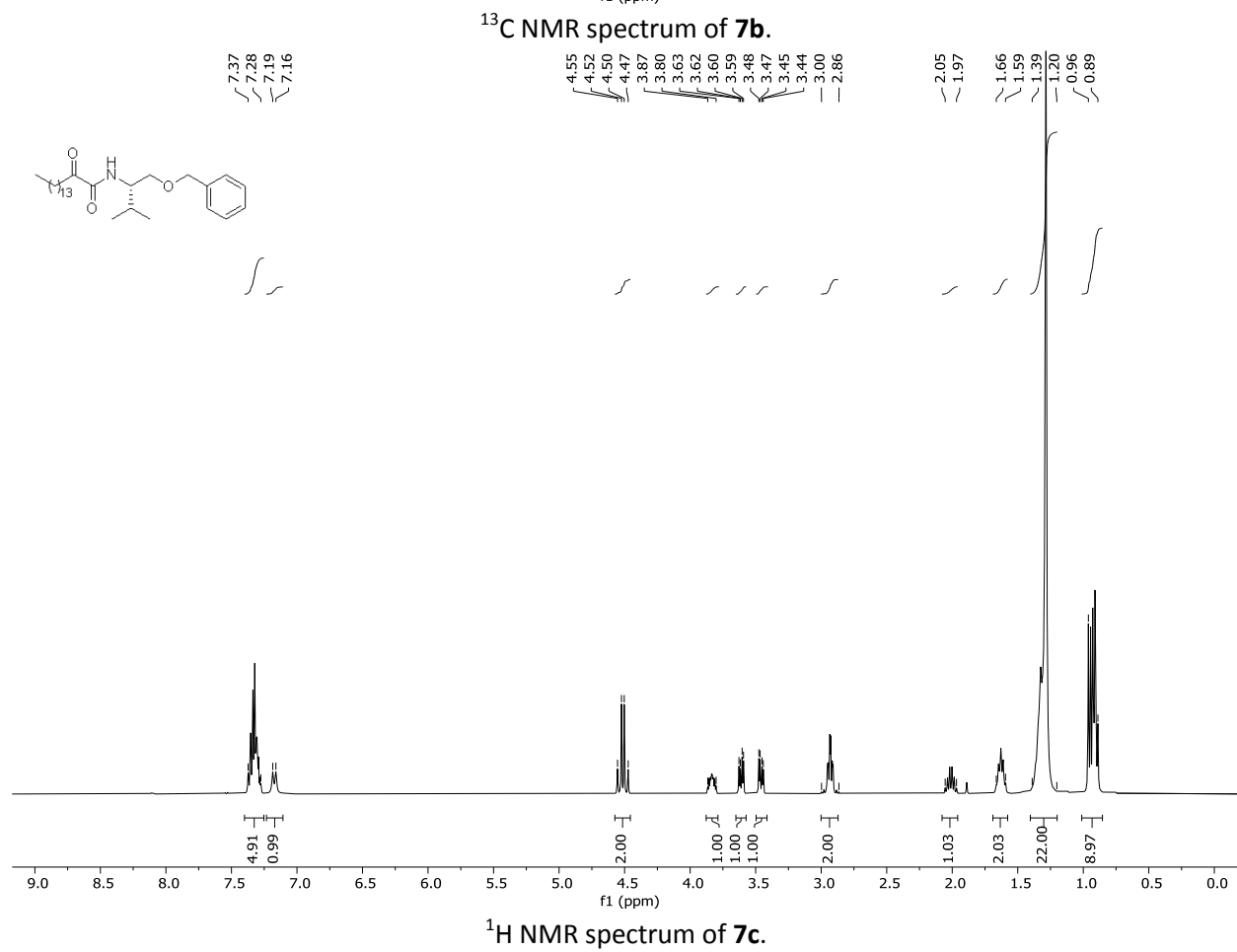
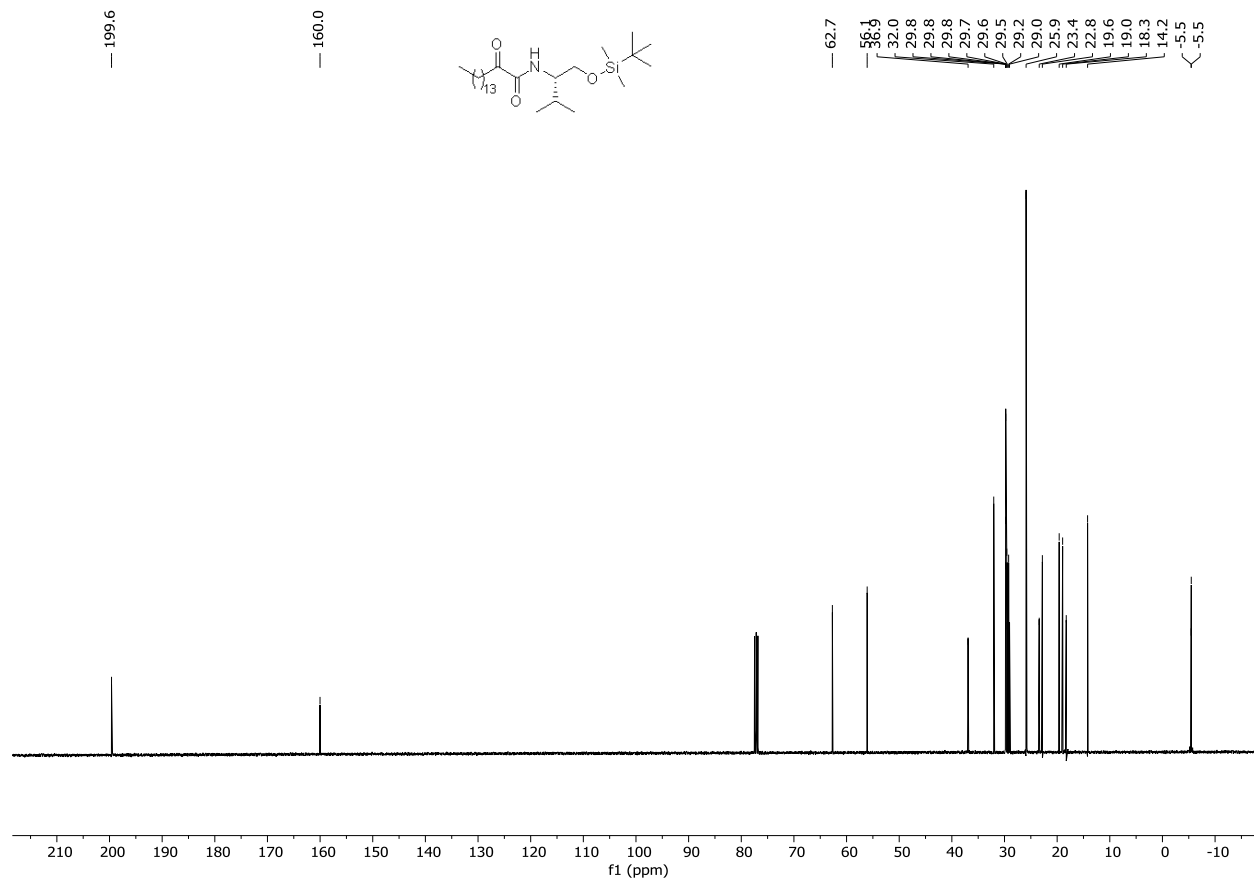


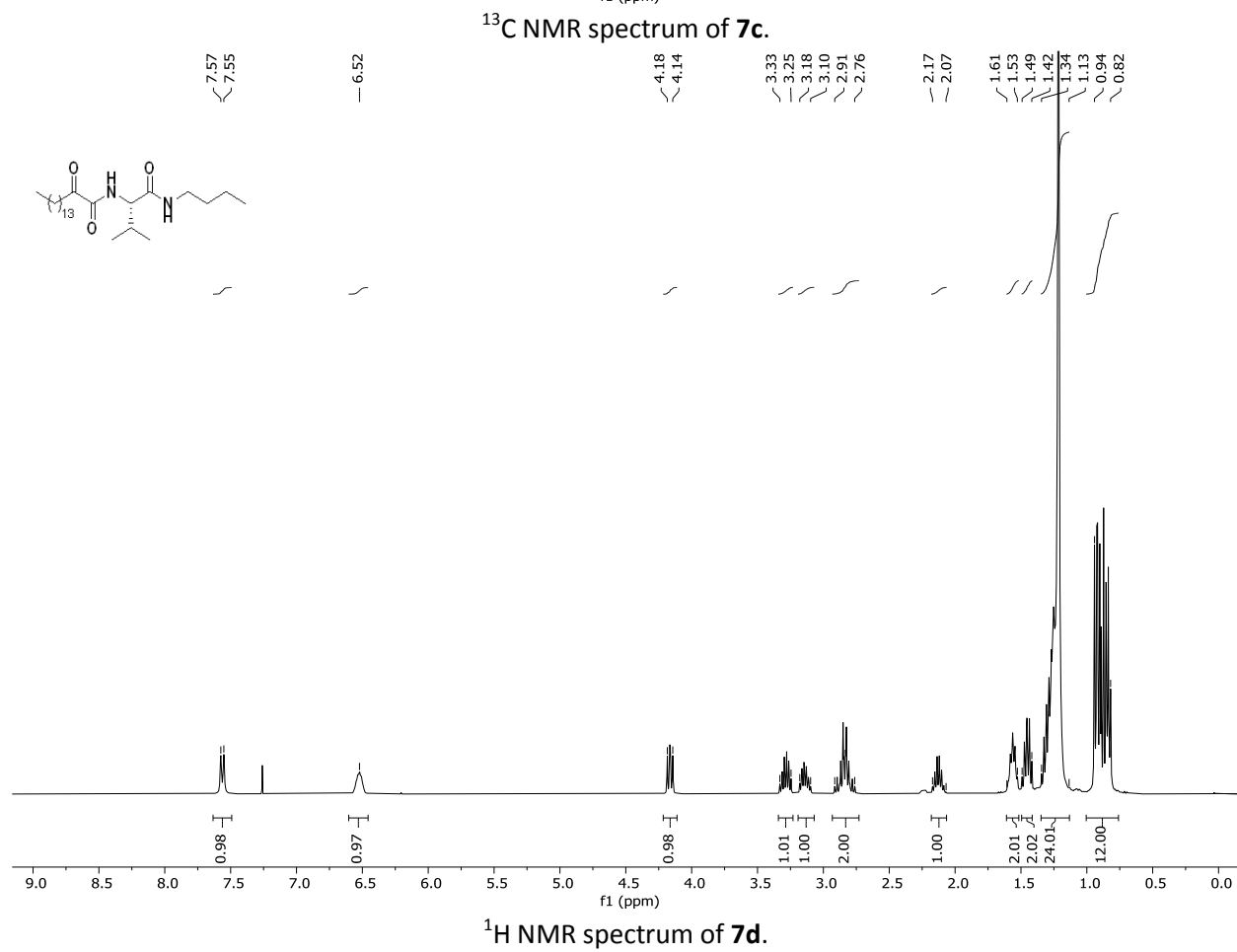
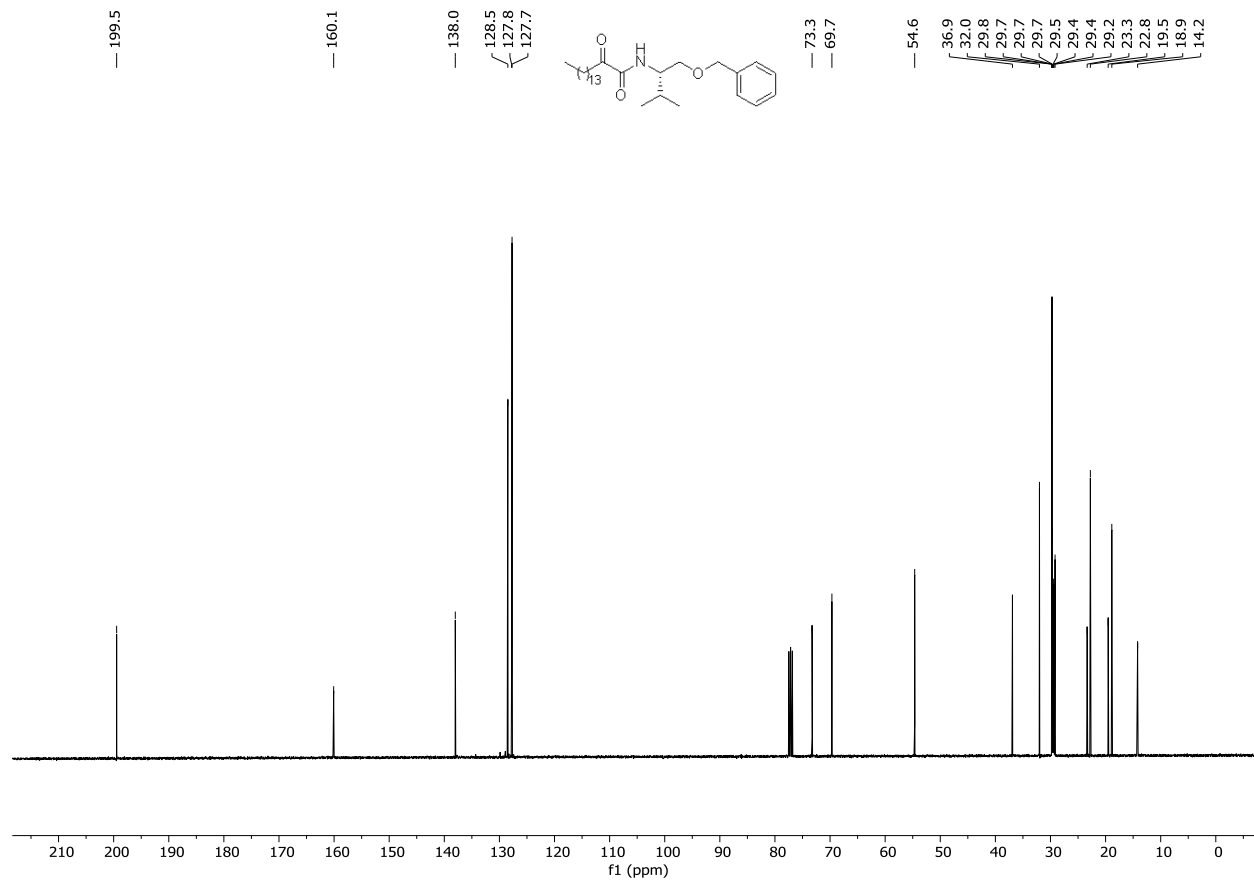


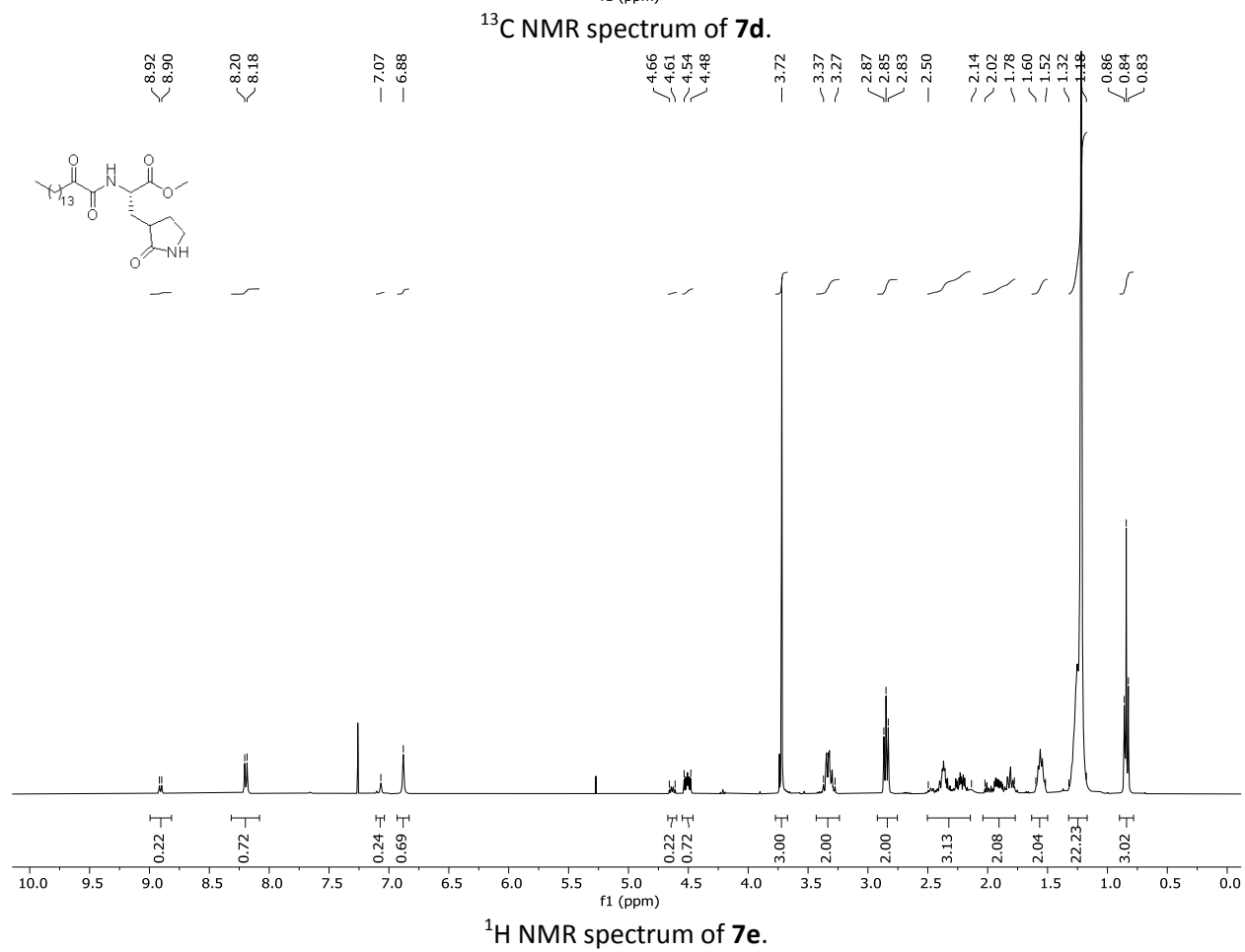
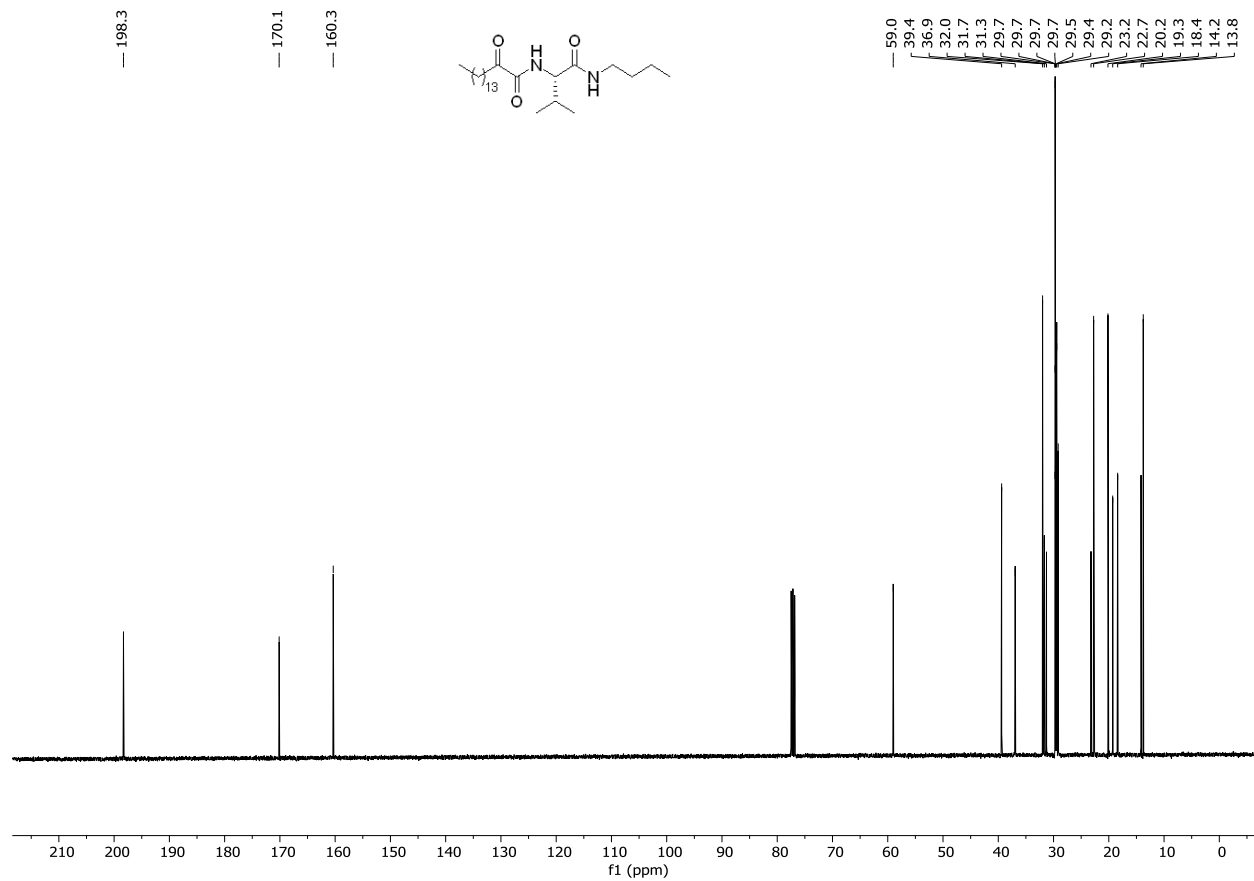


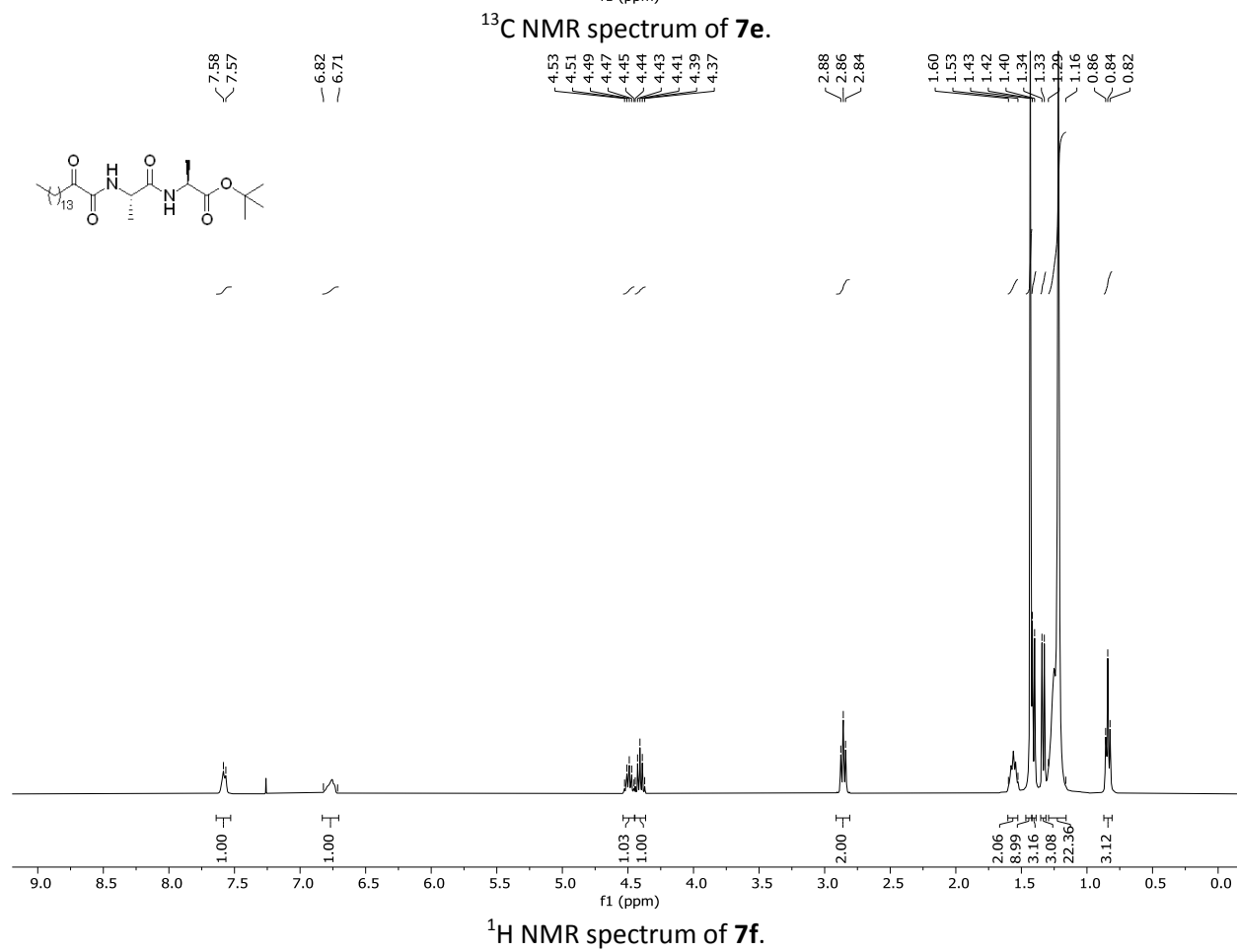
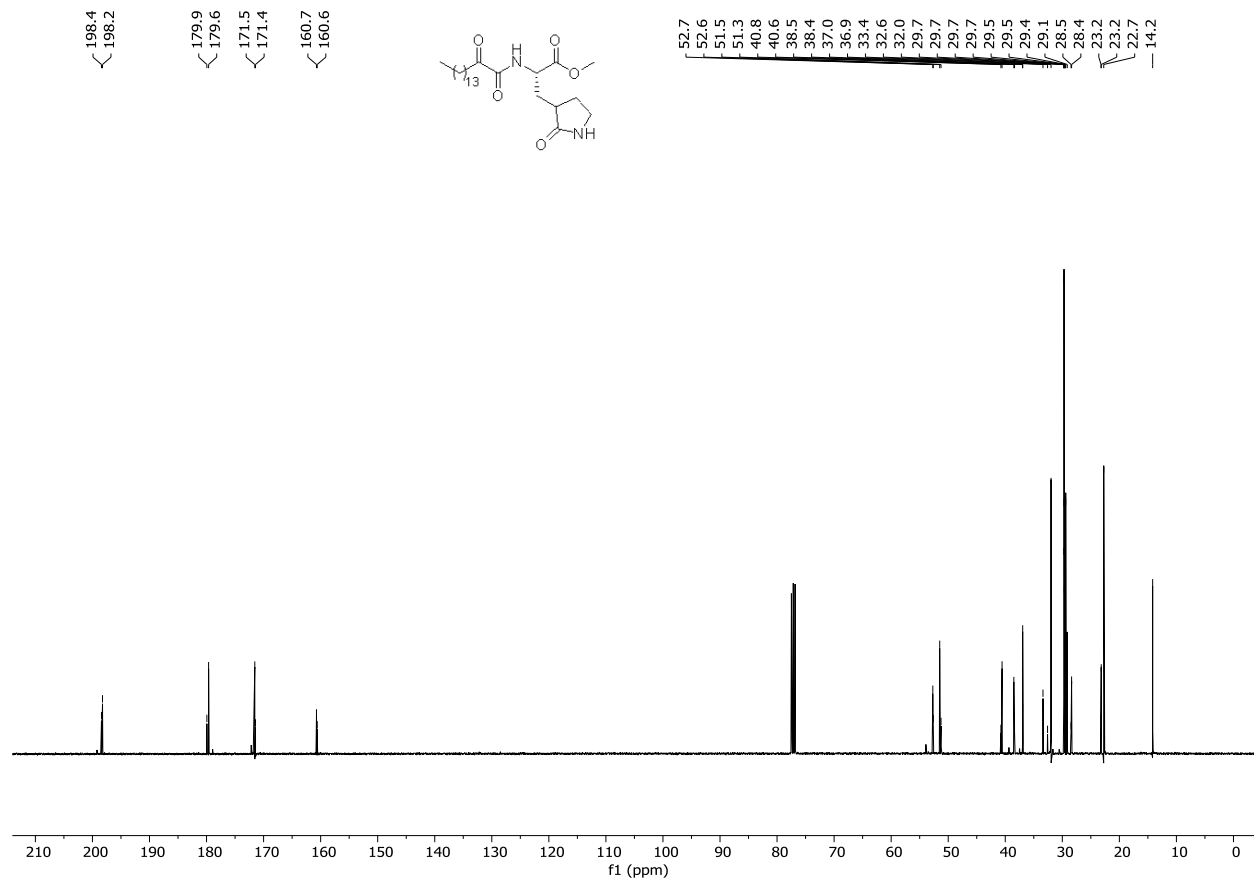


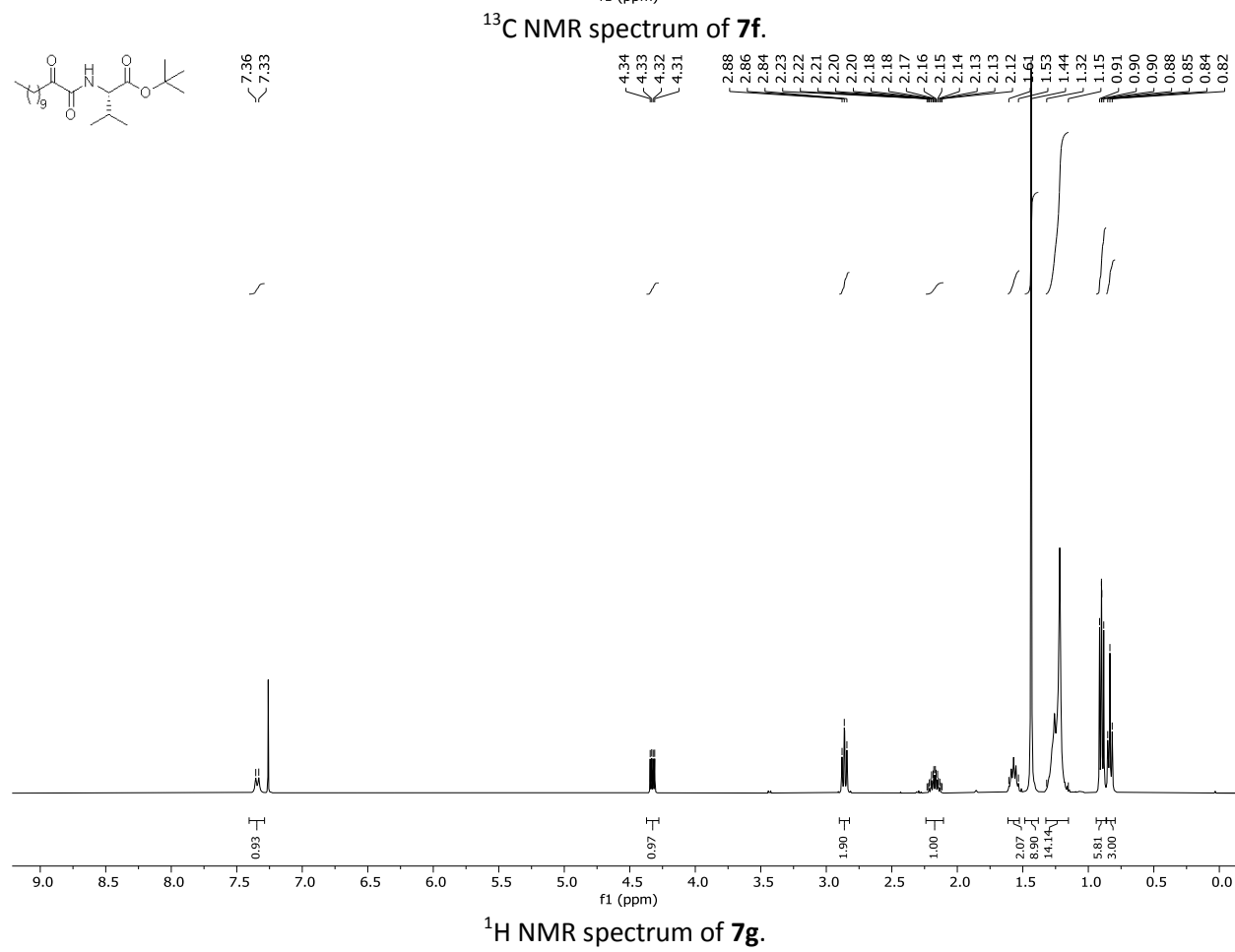
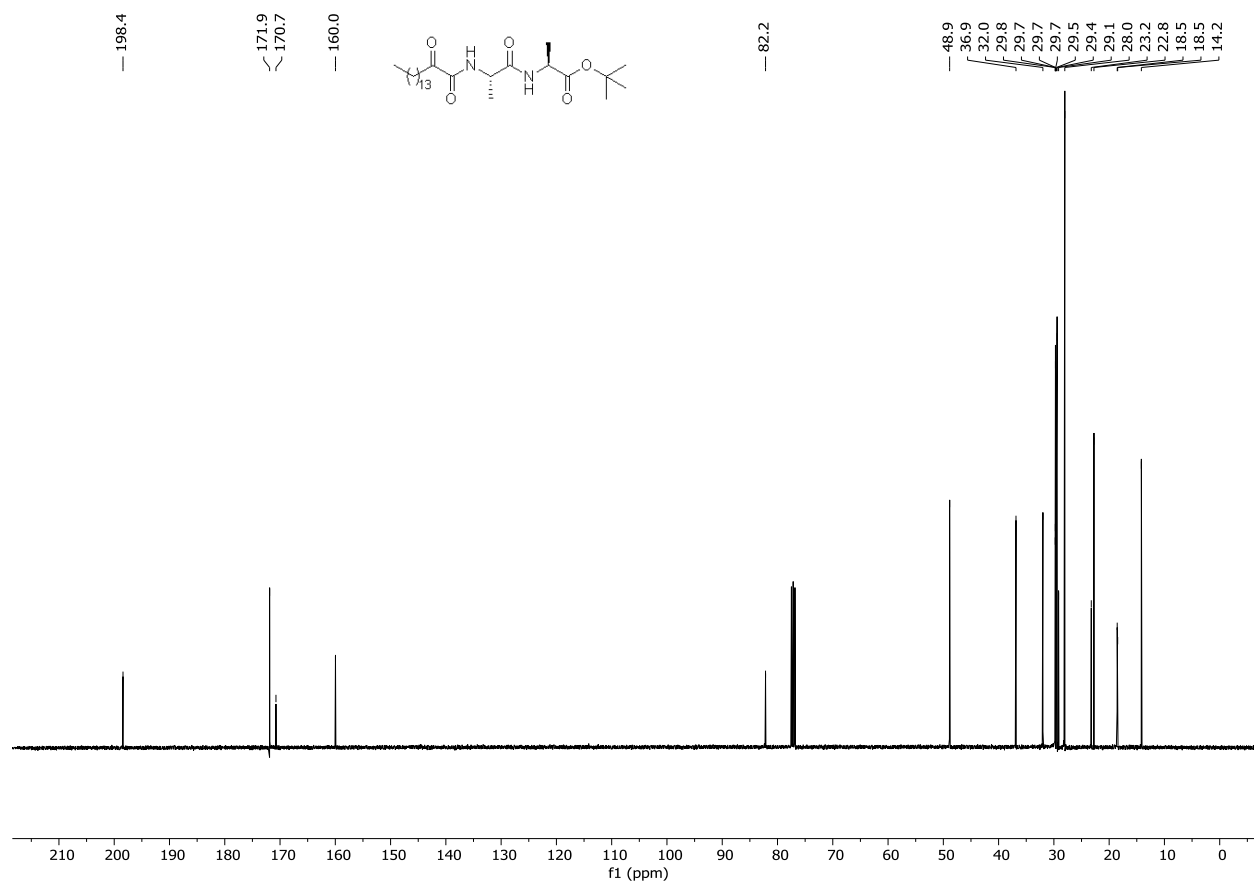


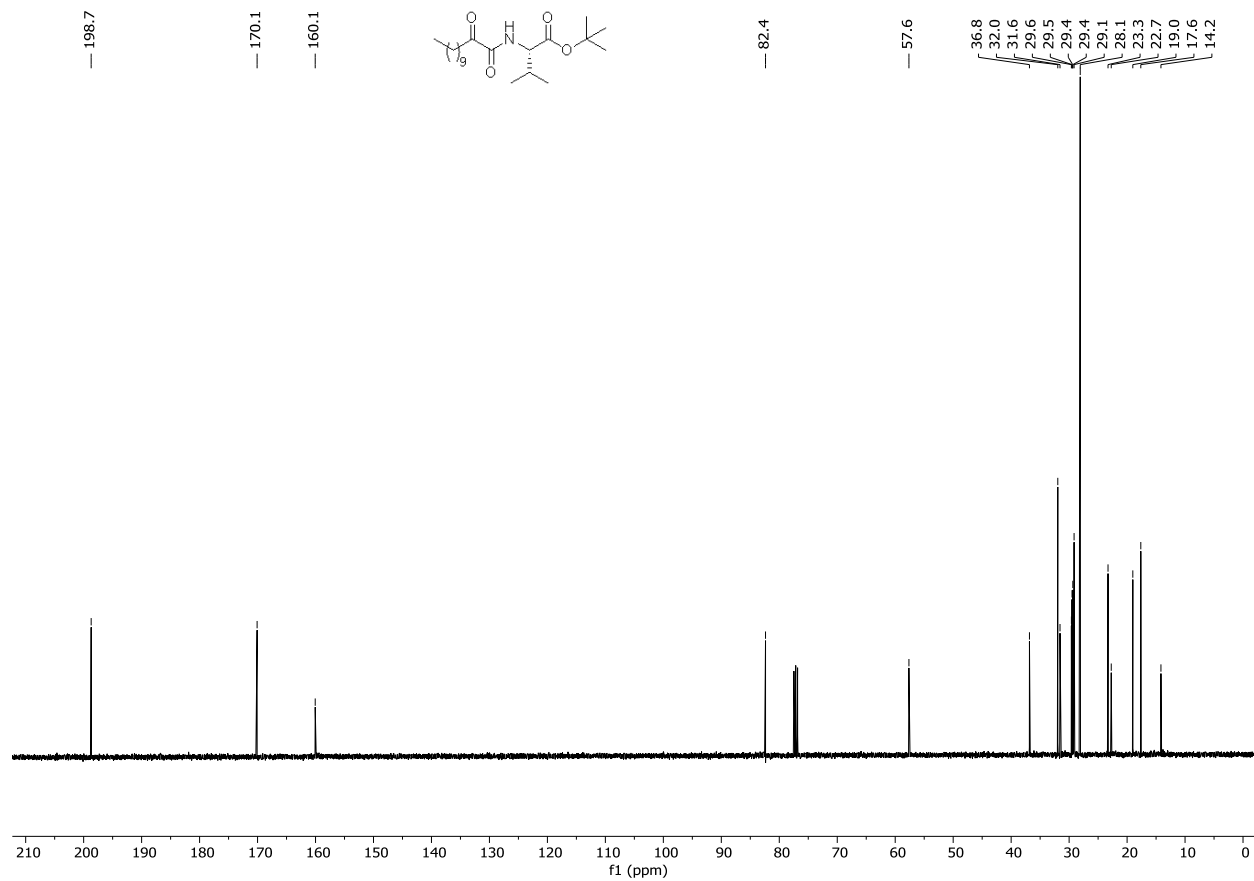




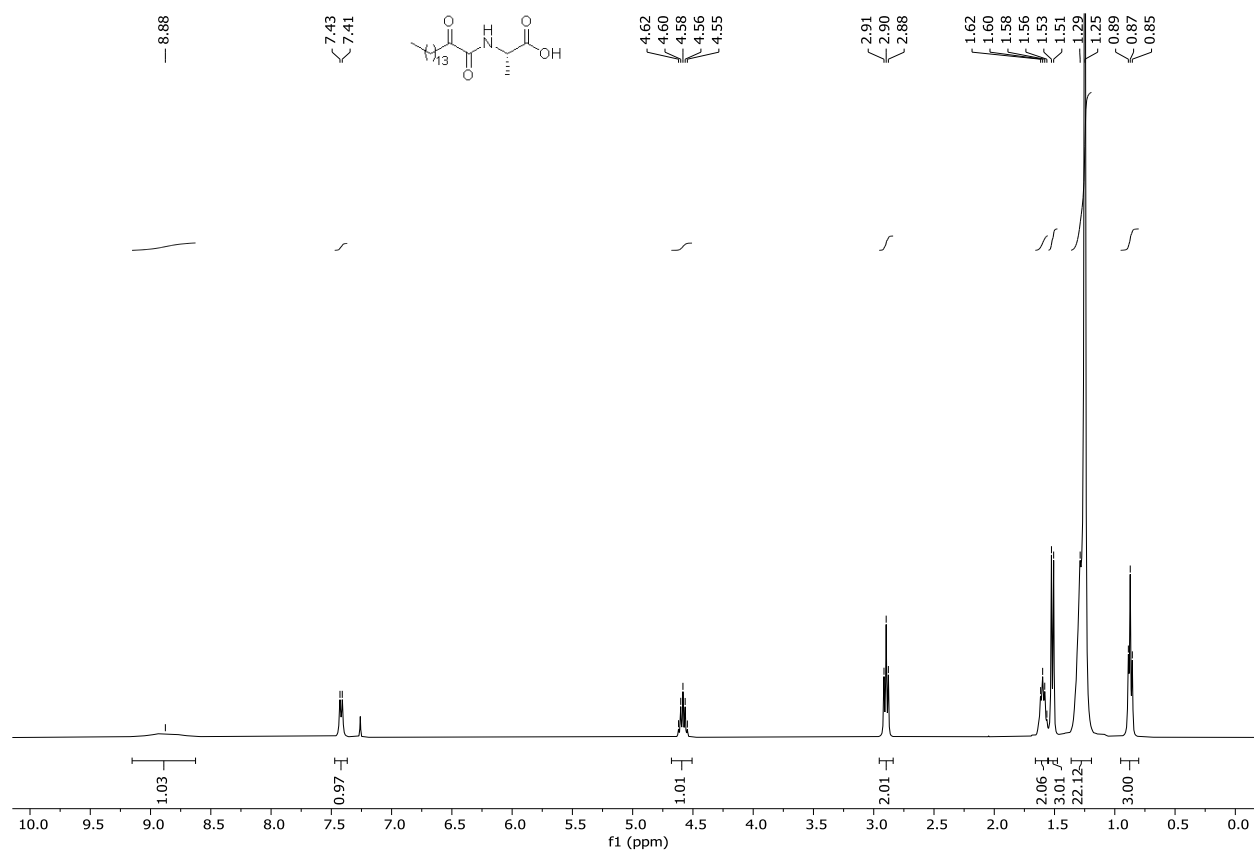








¹³C NMR spectrum of 7g.



¹H NMR spectrum of 8a.

