

Supporting Information for

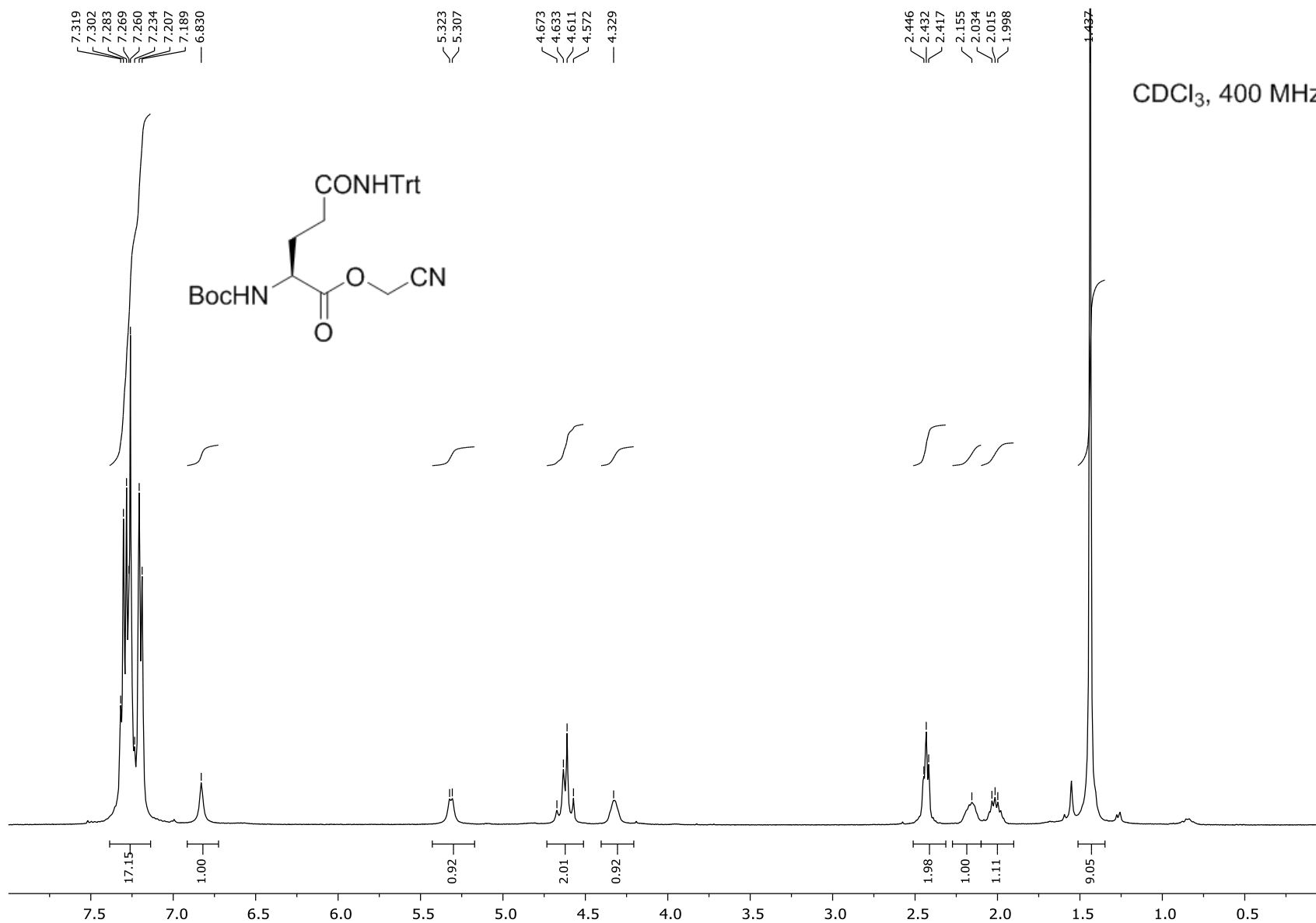
Total Synthesis of the Fellutamides, Lipopeptide Proteasome Inhibitors. Sustainable Peptide Bond Formation

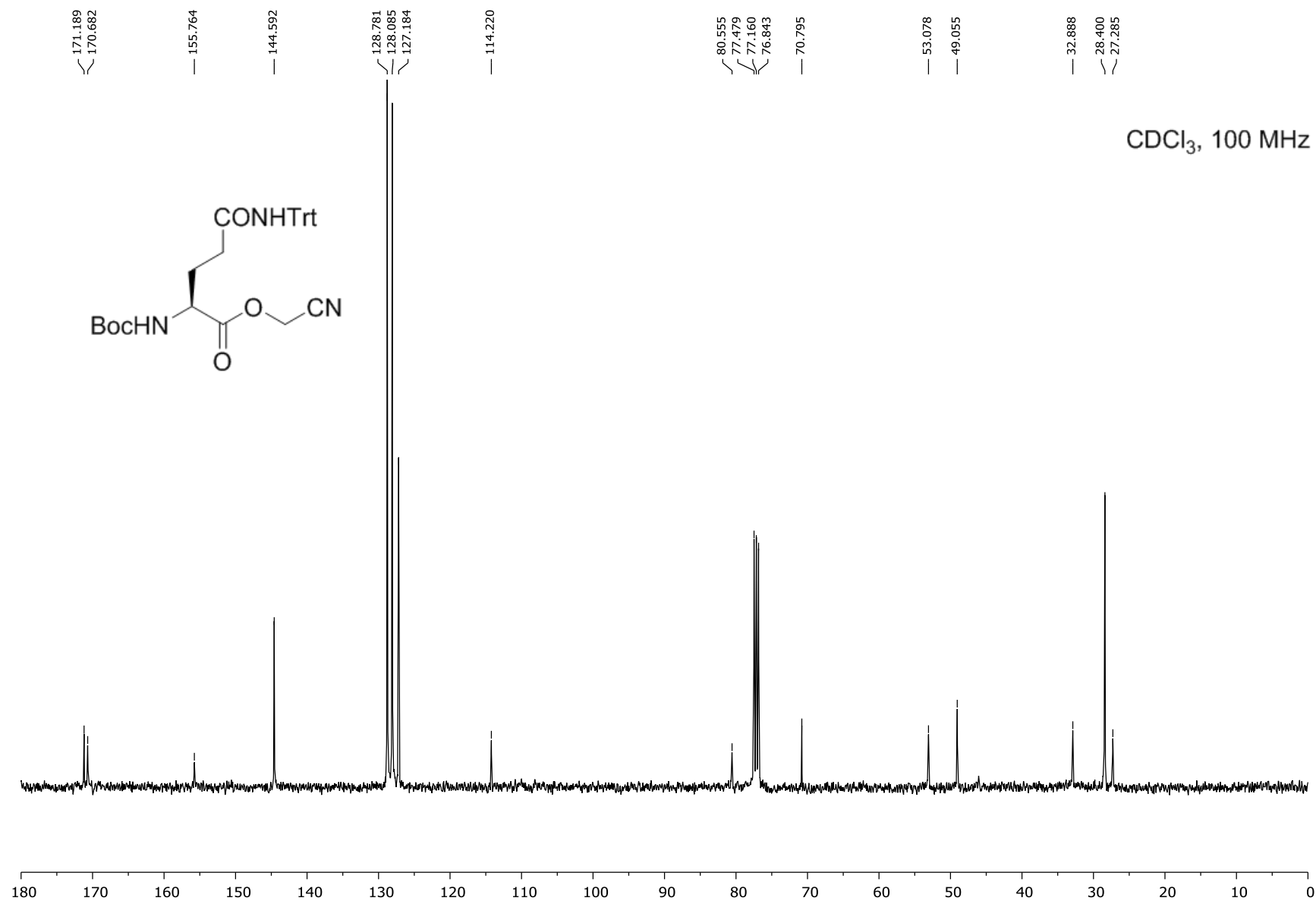
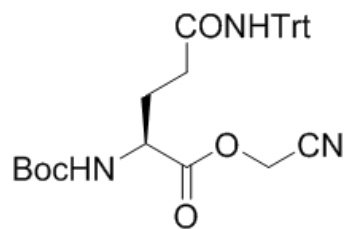
Michael C. Pirrung,* Fa Zhang, Sudhakar Ambadi, and Y. Gangadhara Rao

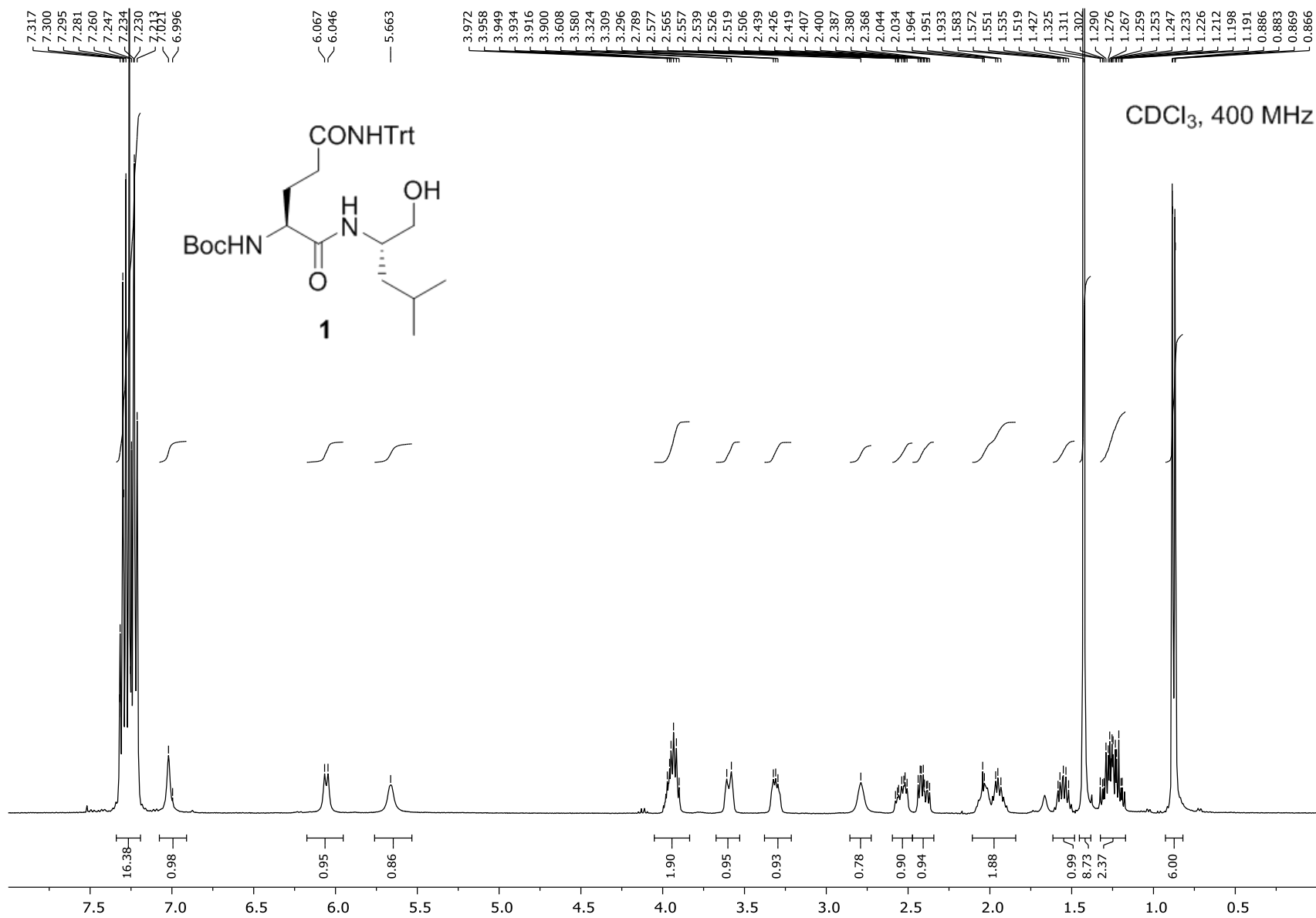
Department of Chemistry, University of California, Riverside, CA 92521, USA. Fax: 1 951 827 2749; *E-mail: michael.pirrung@ucr.edu

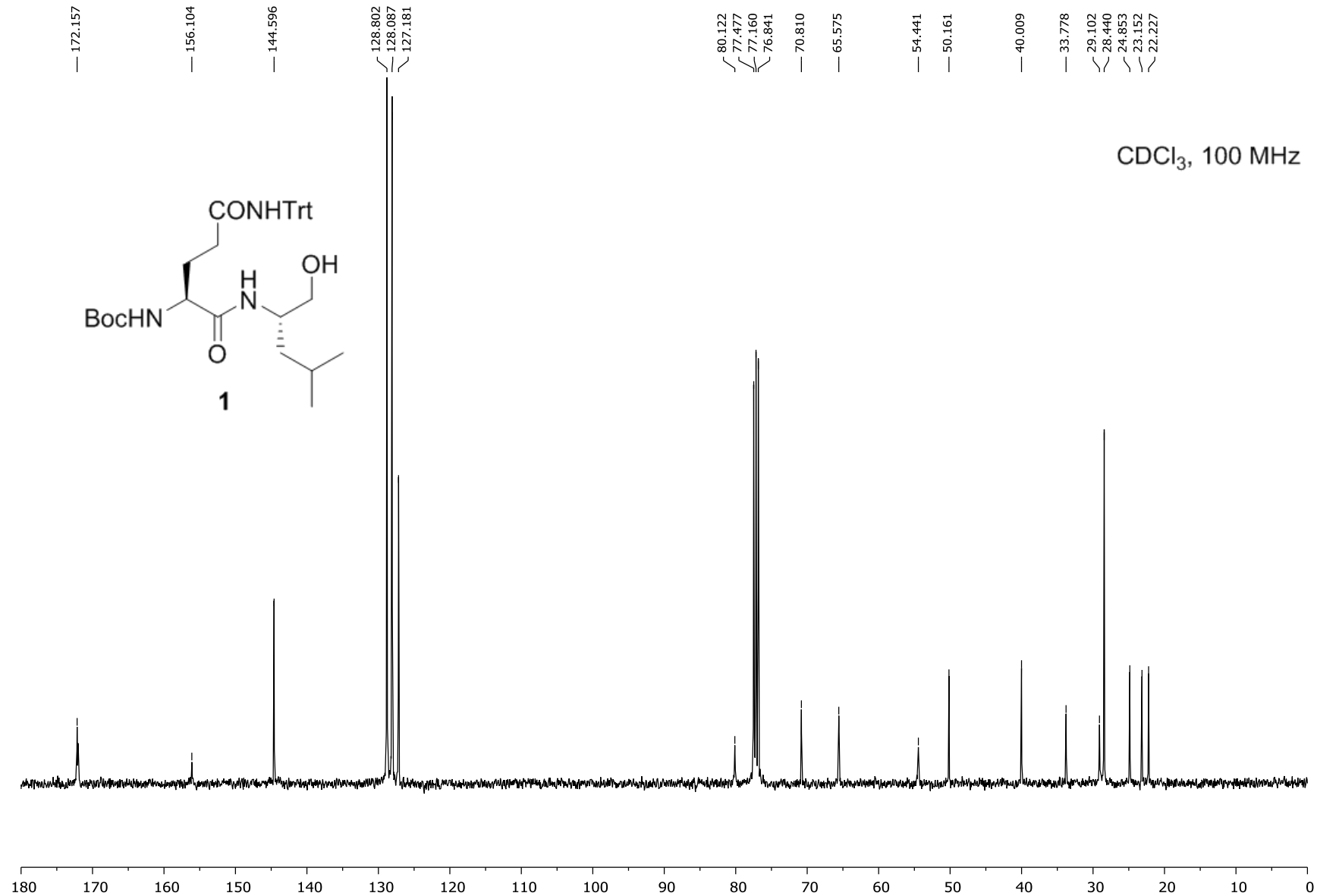
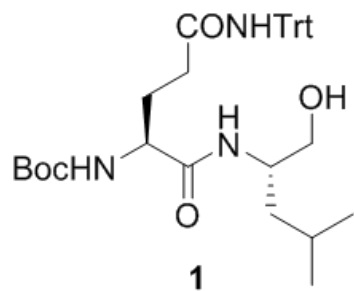
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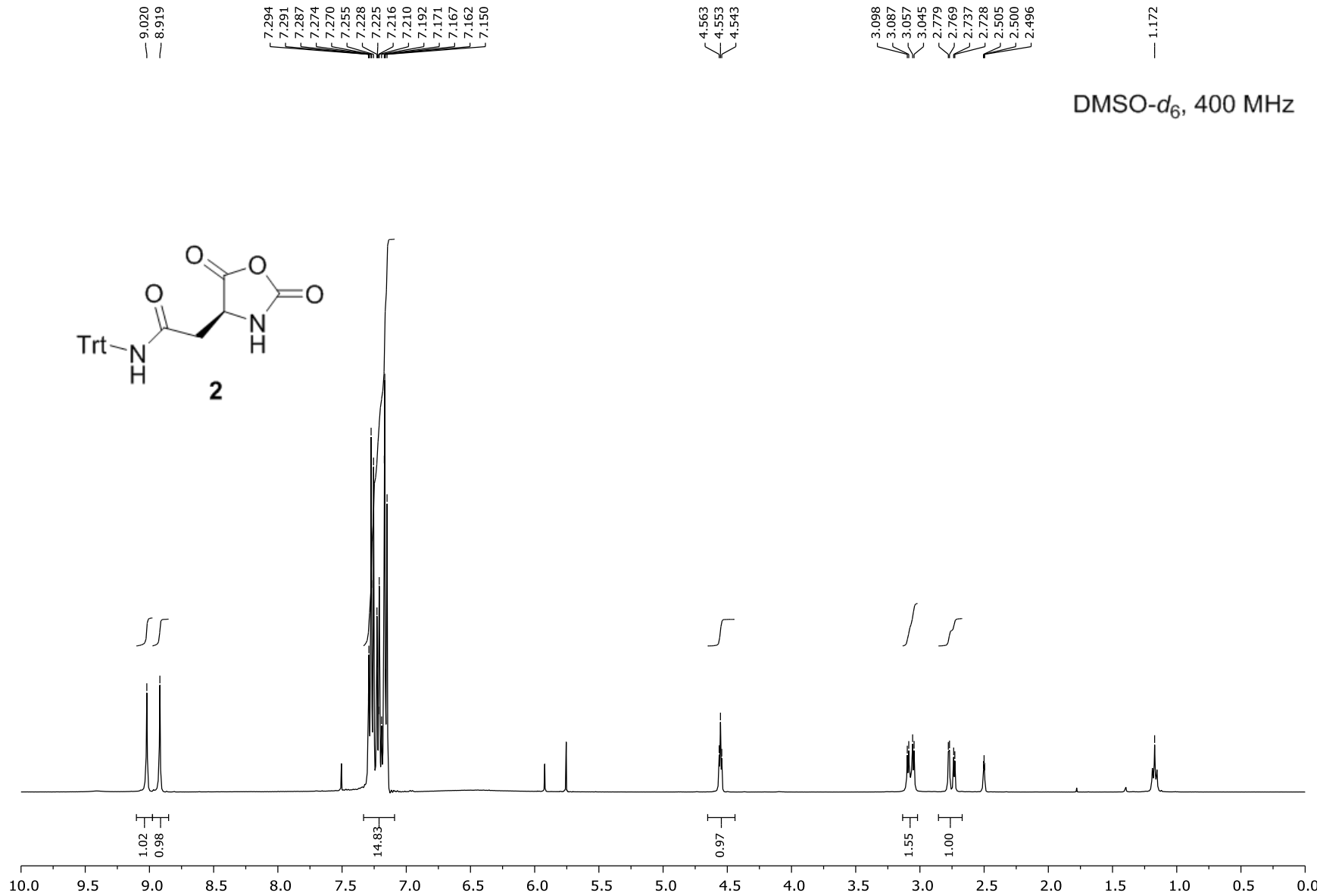
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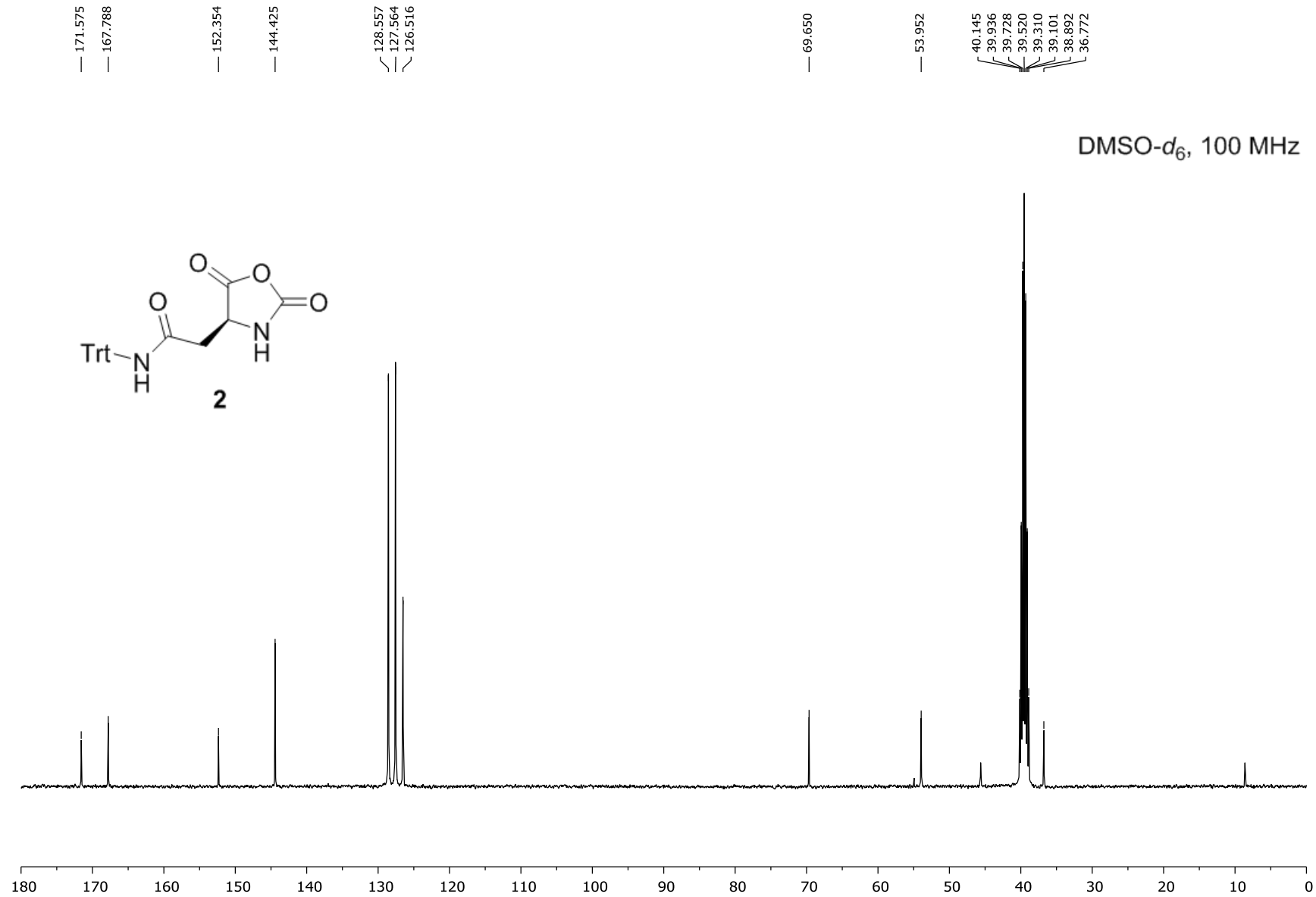


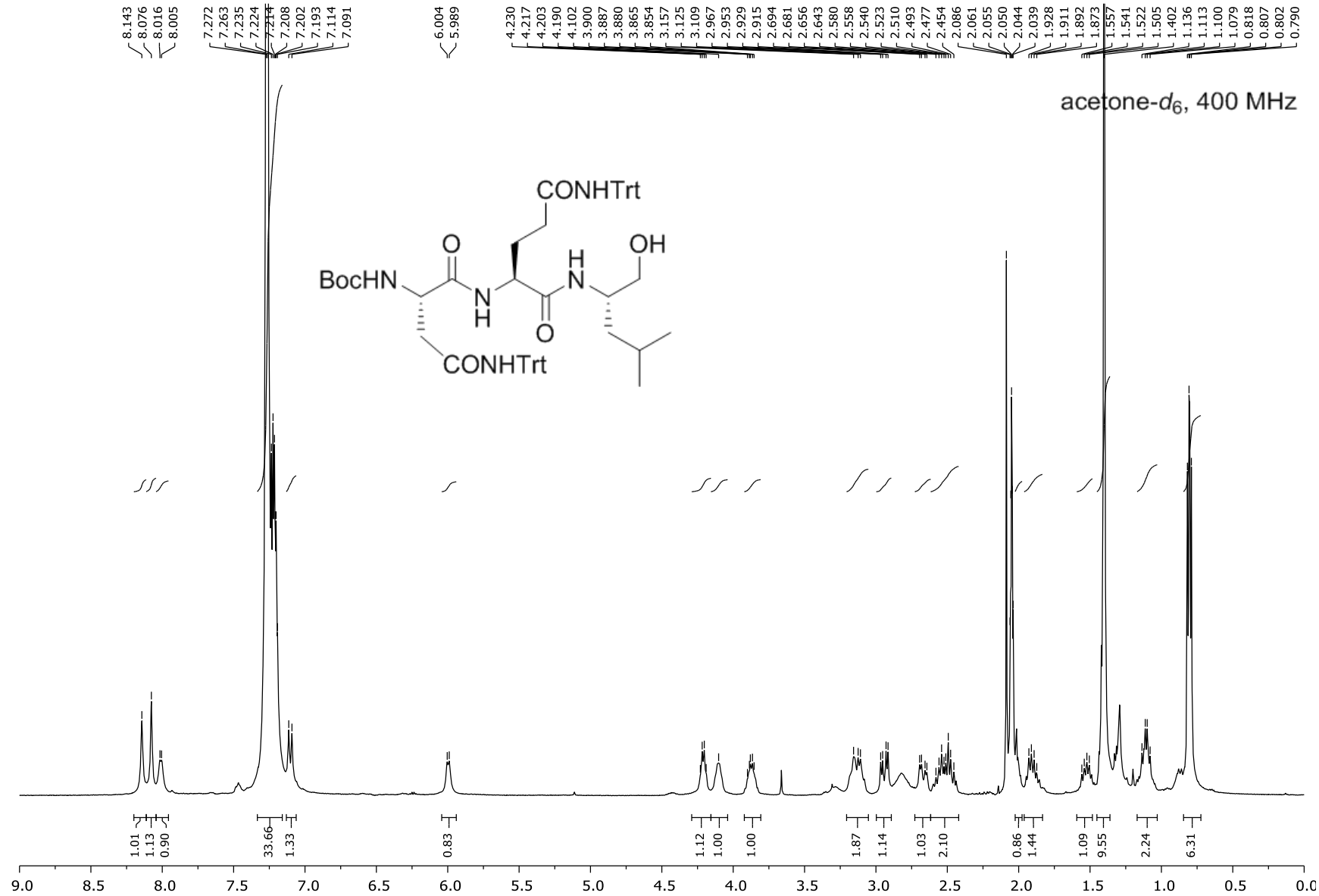


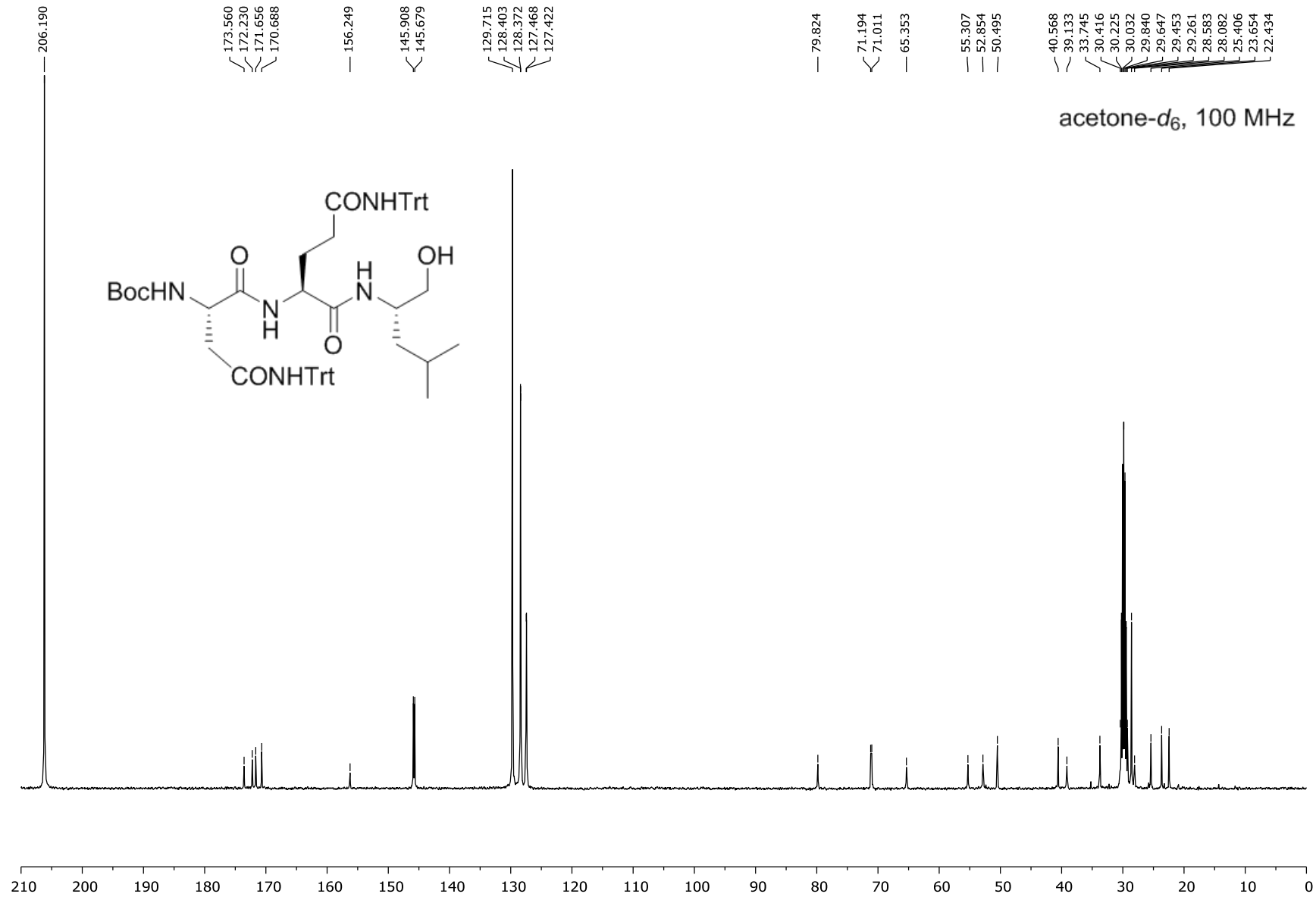


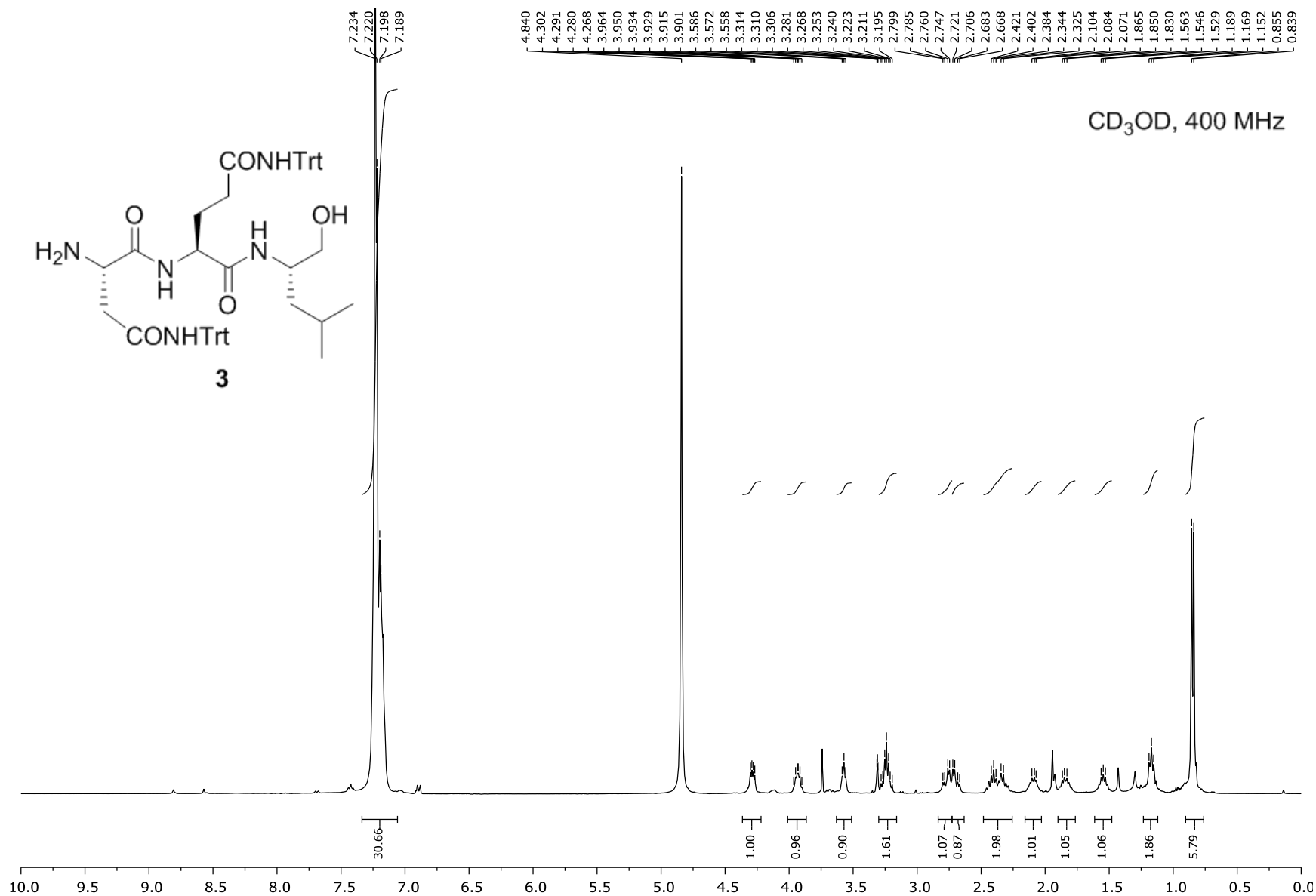
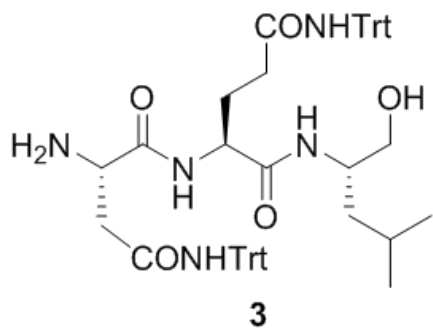


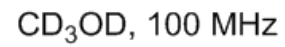
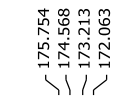


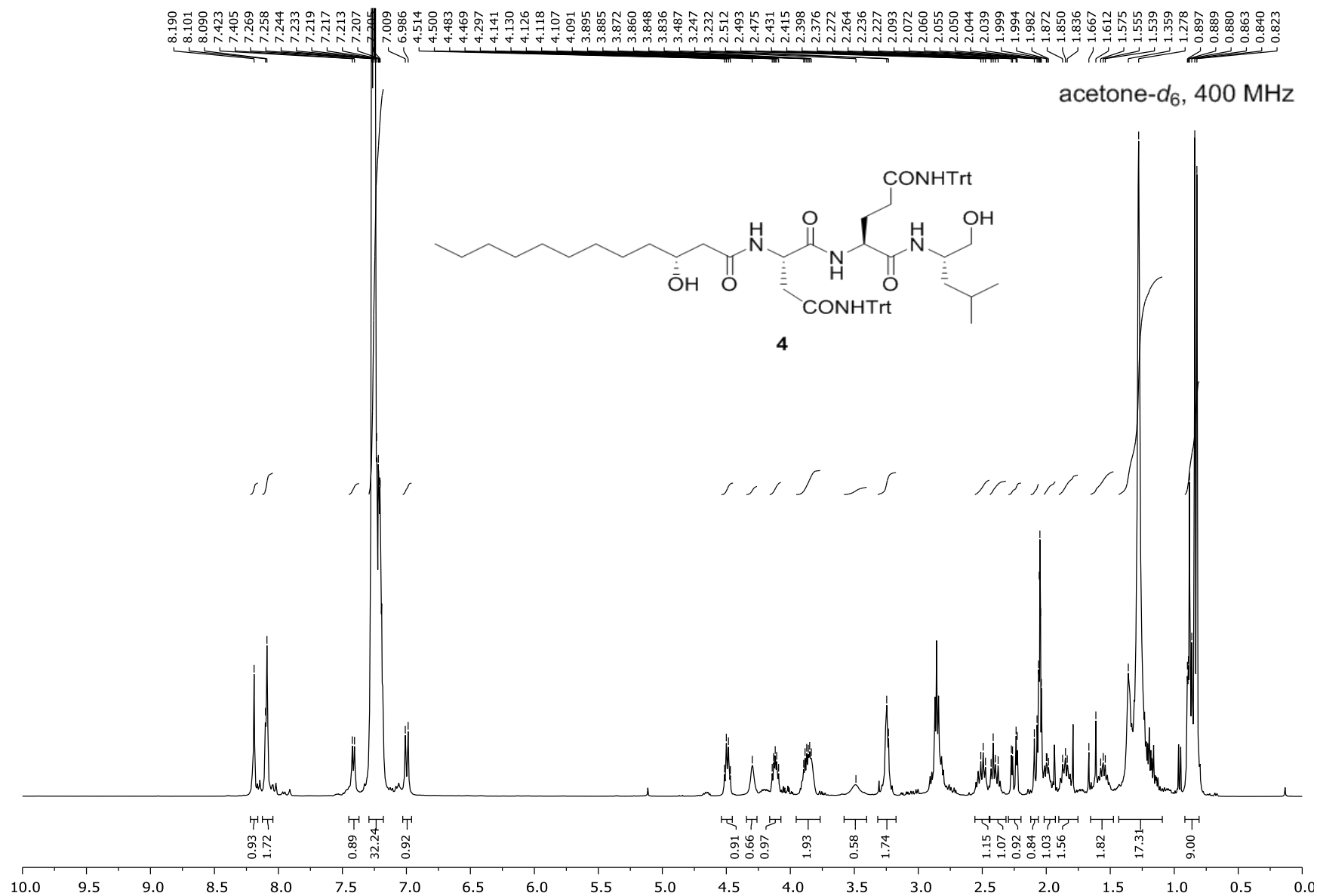


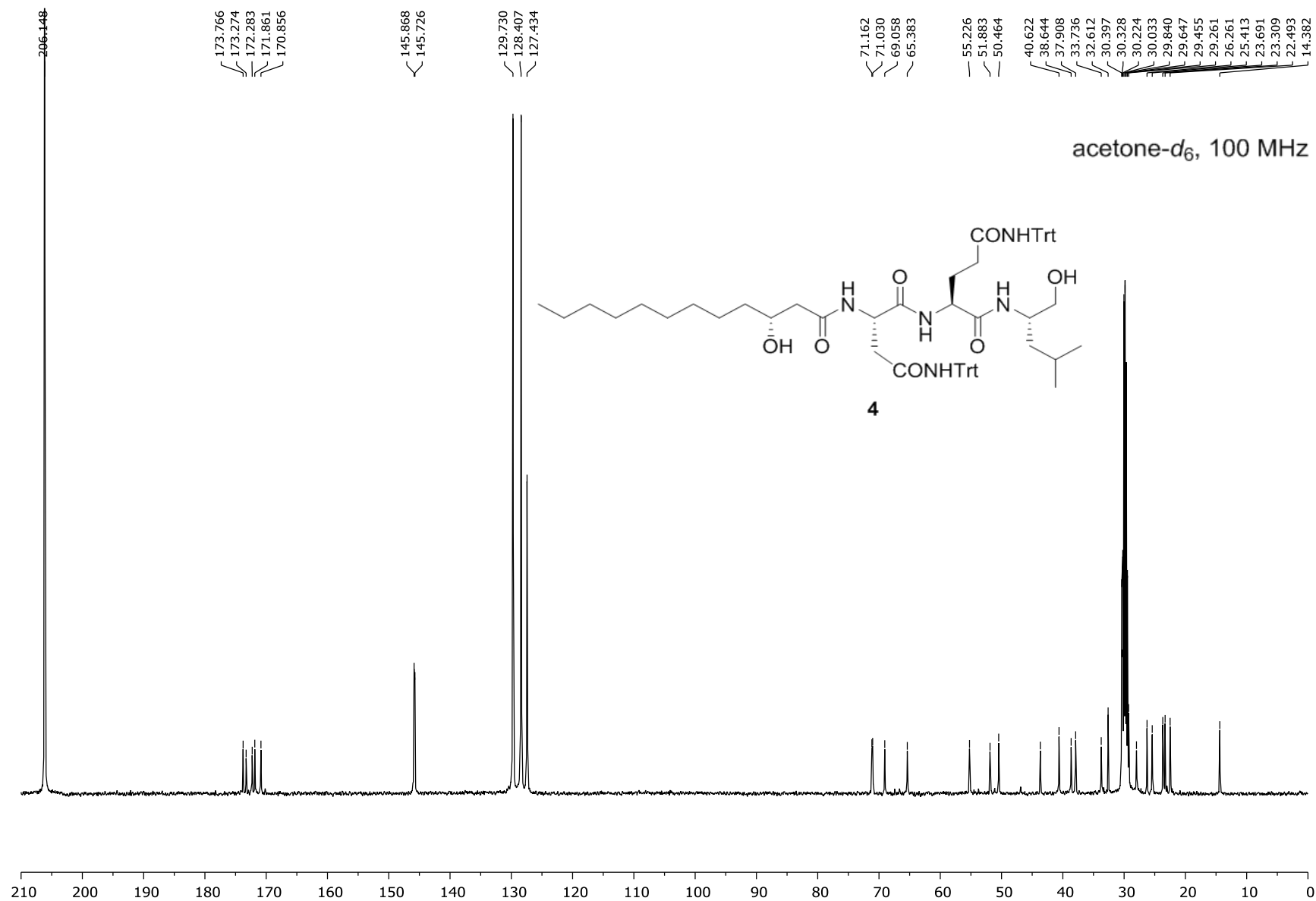


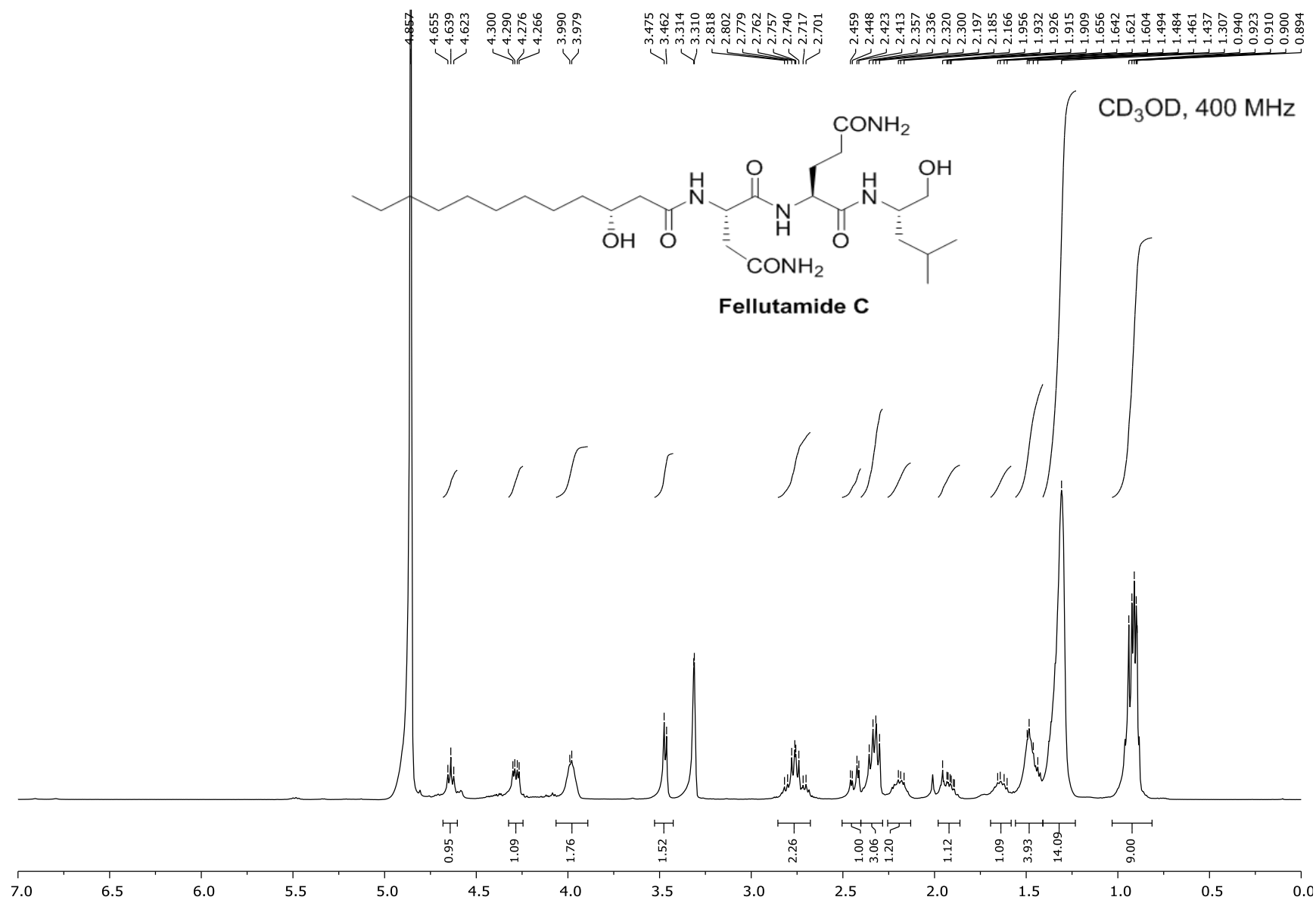


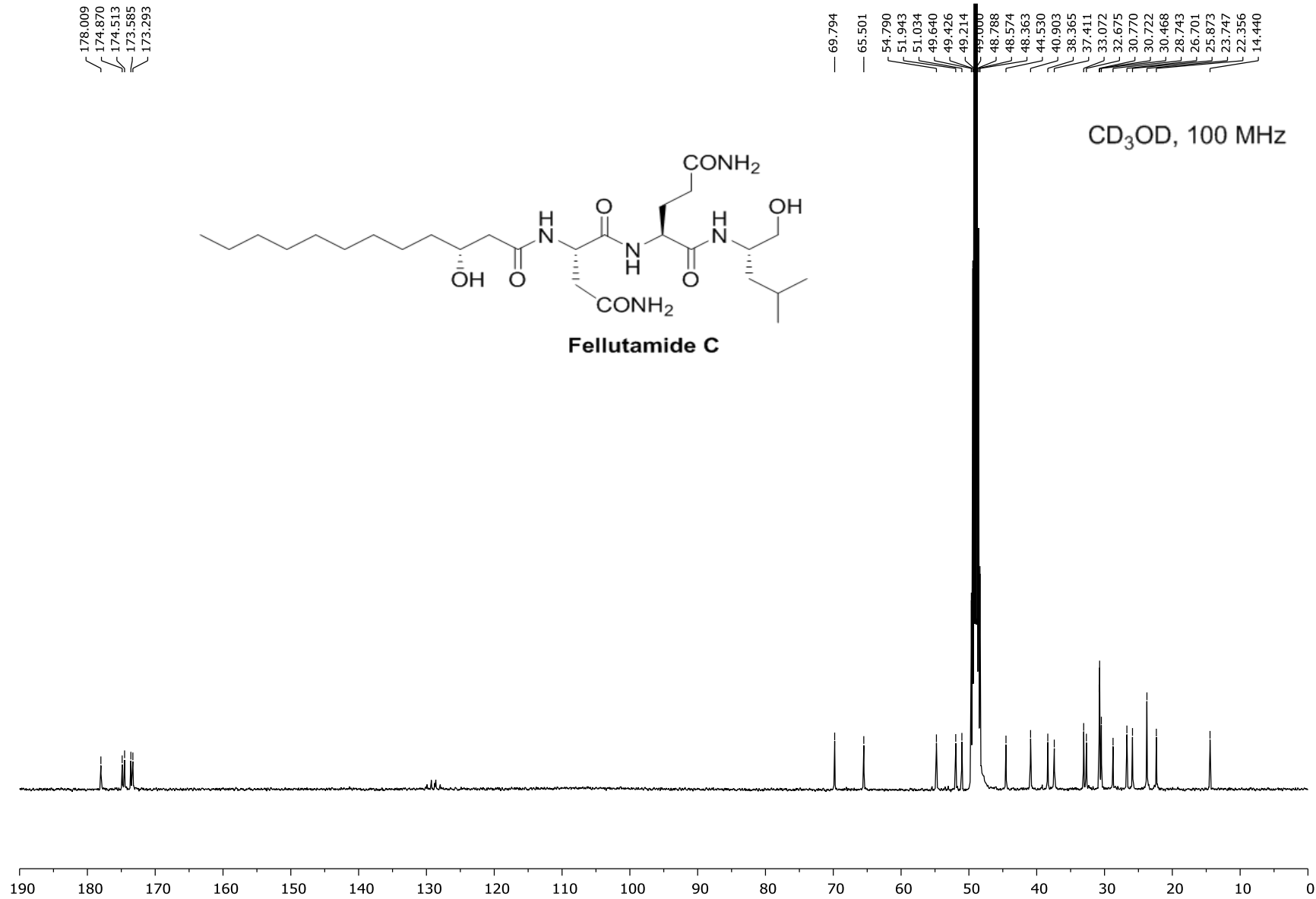


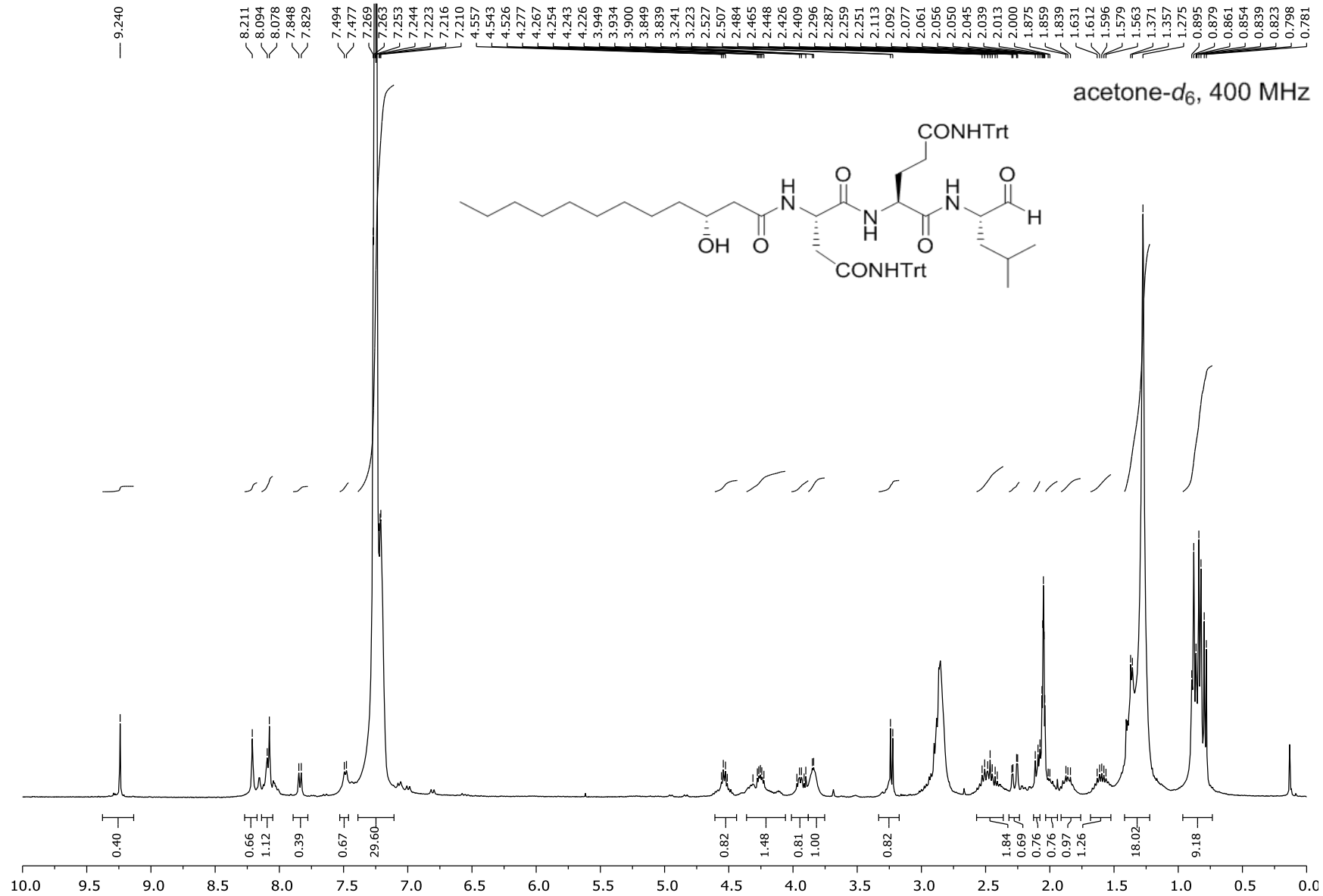
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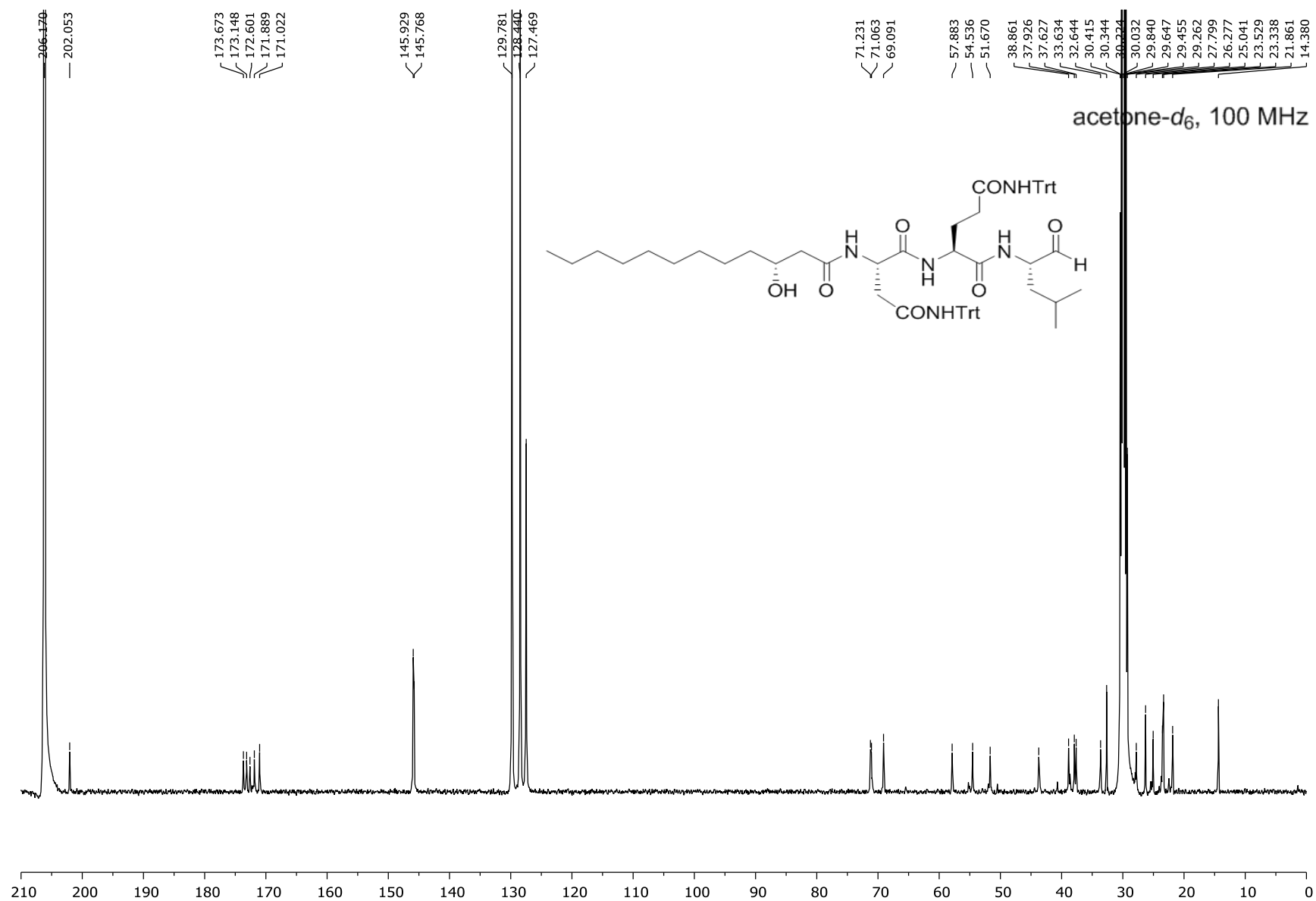


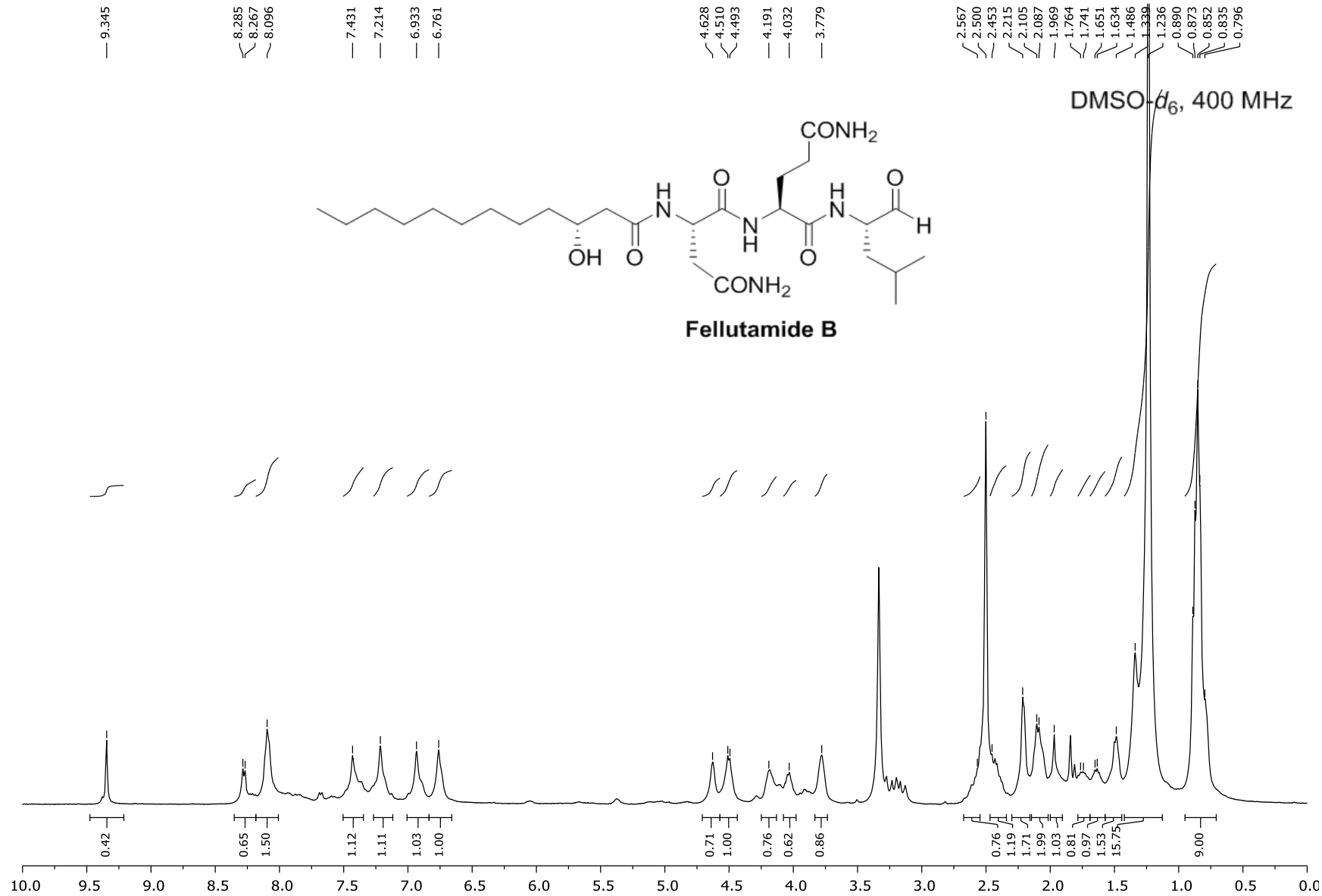


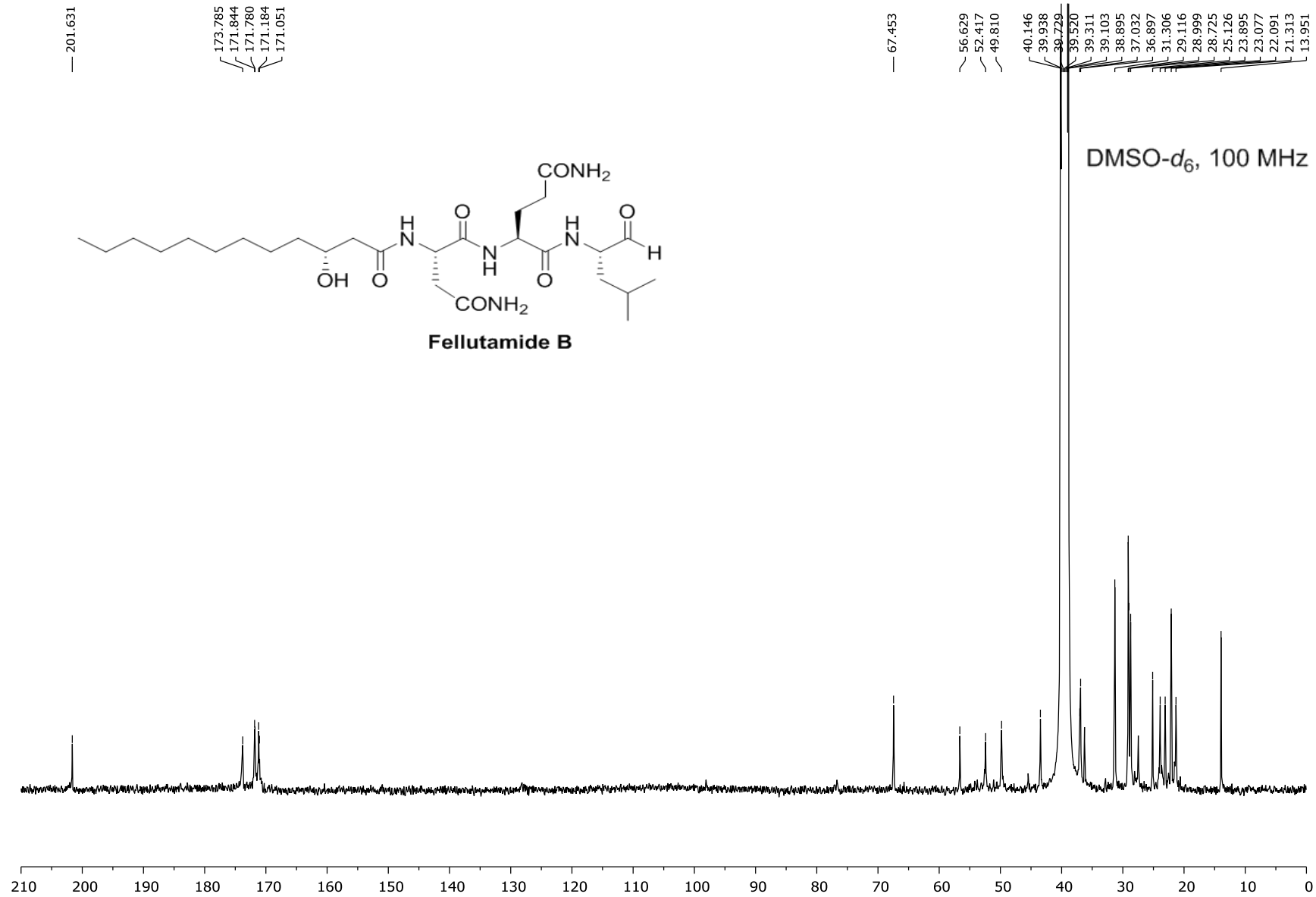


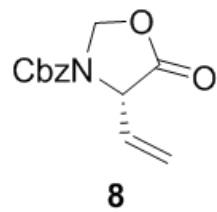




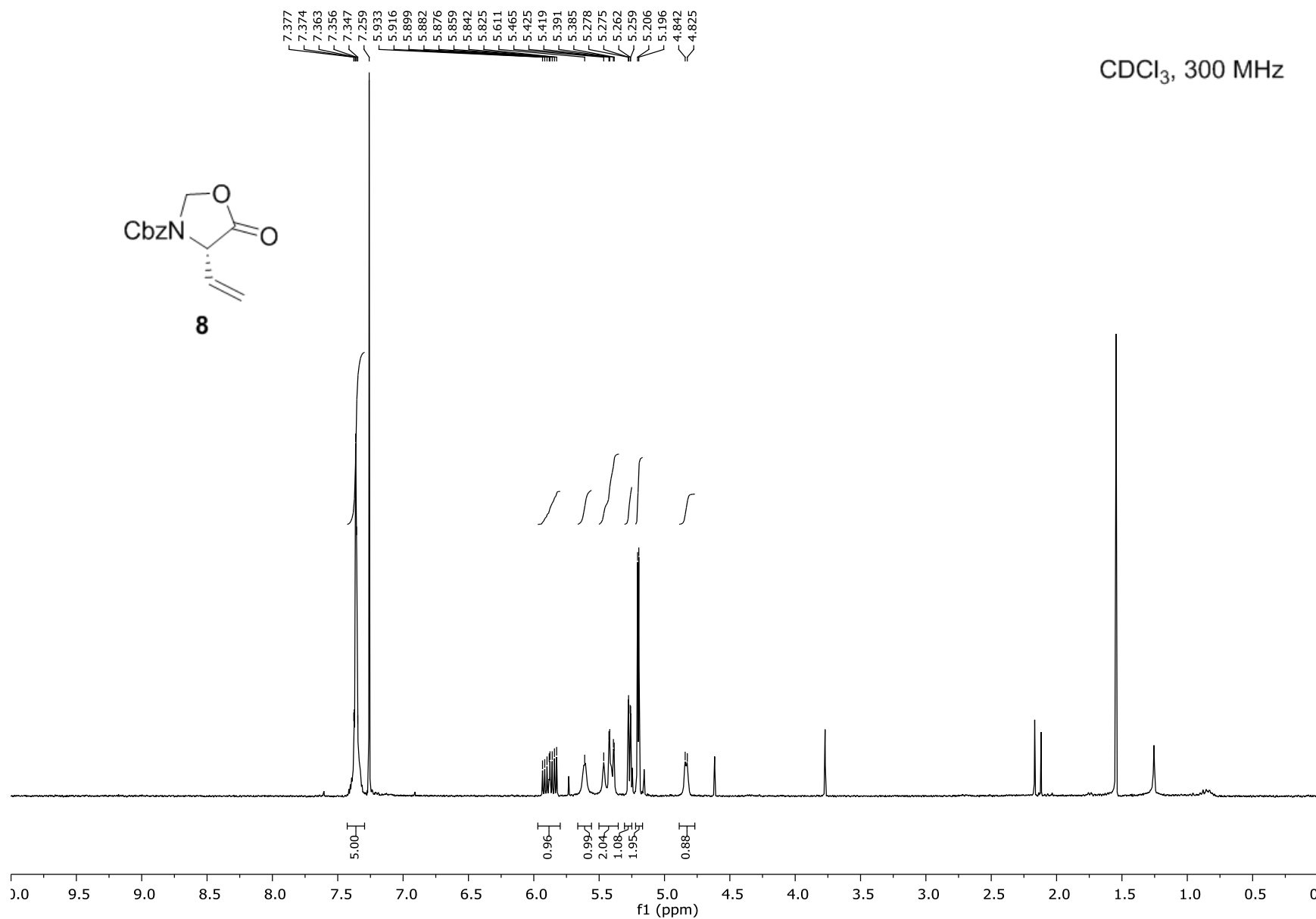


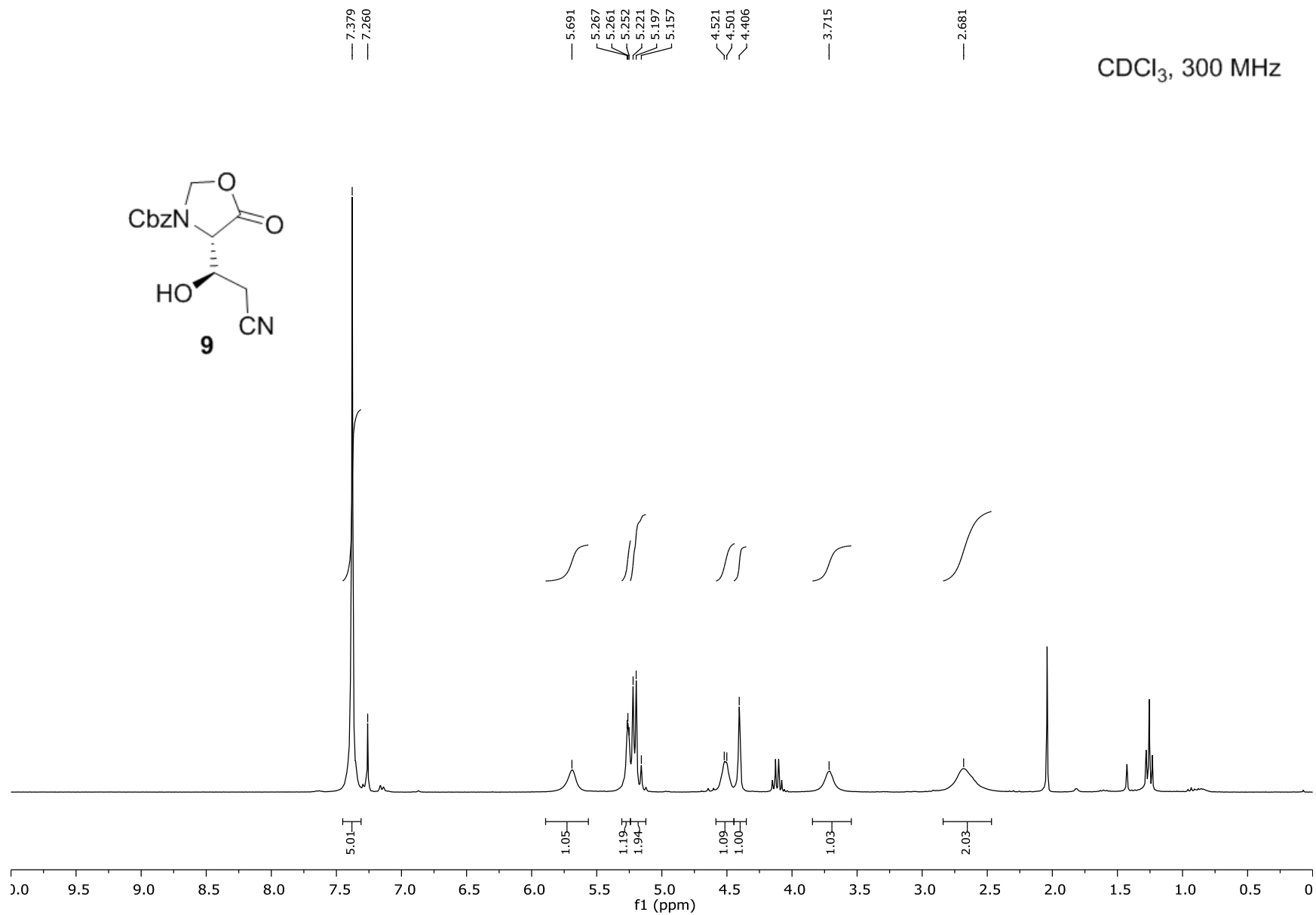
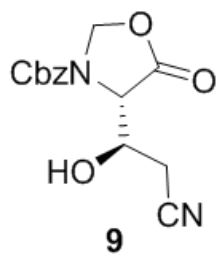


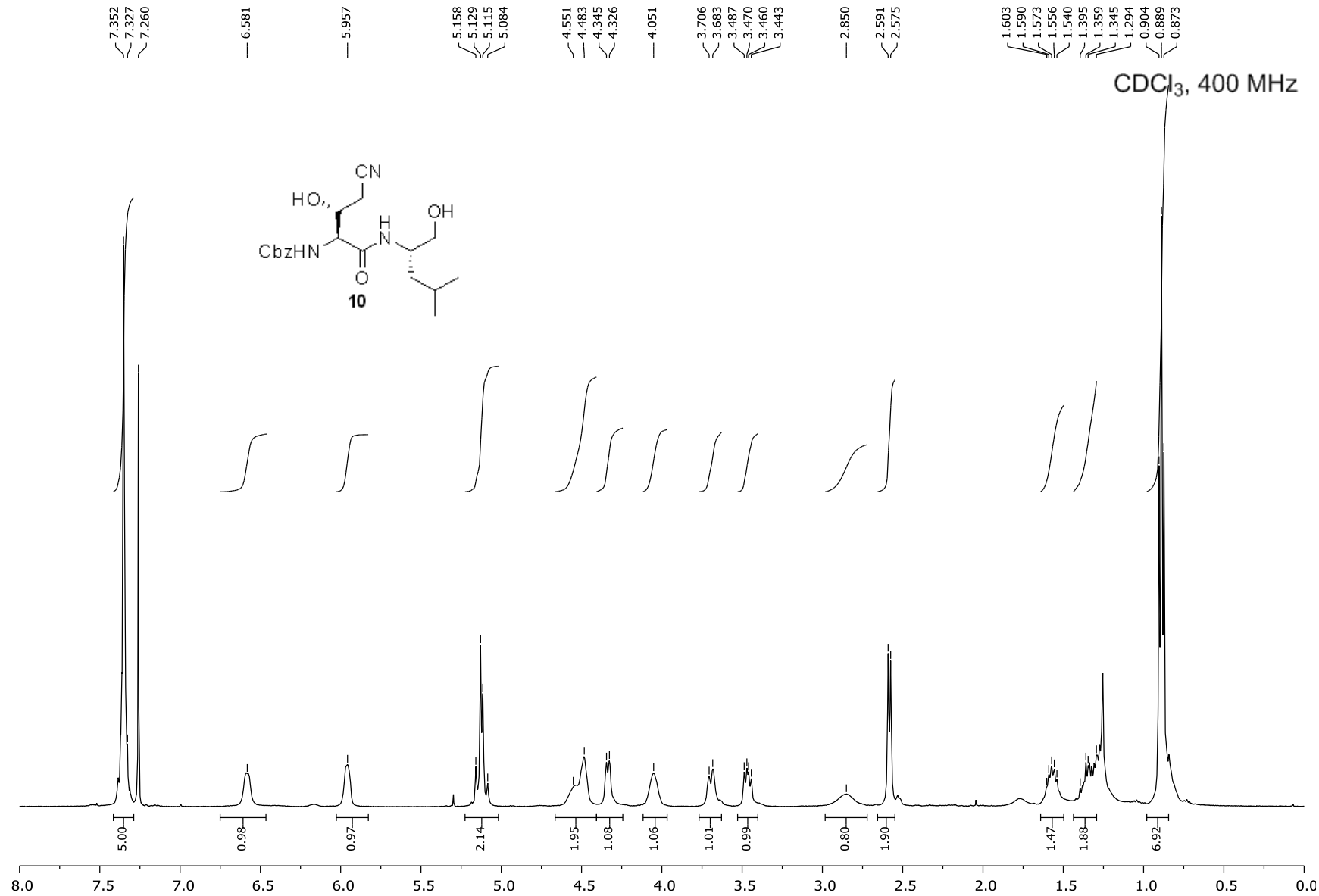


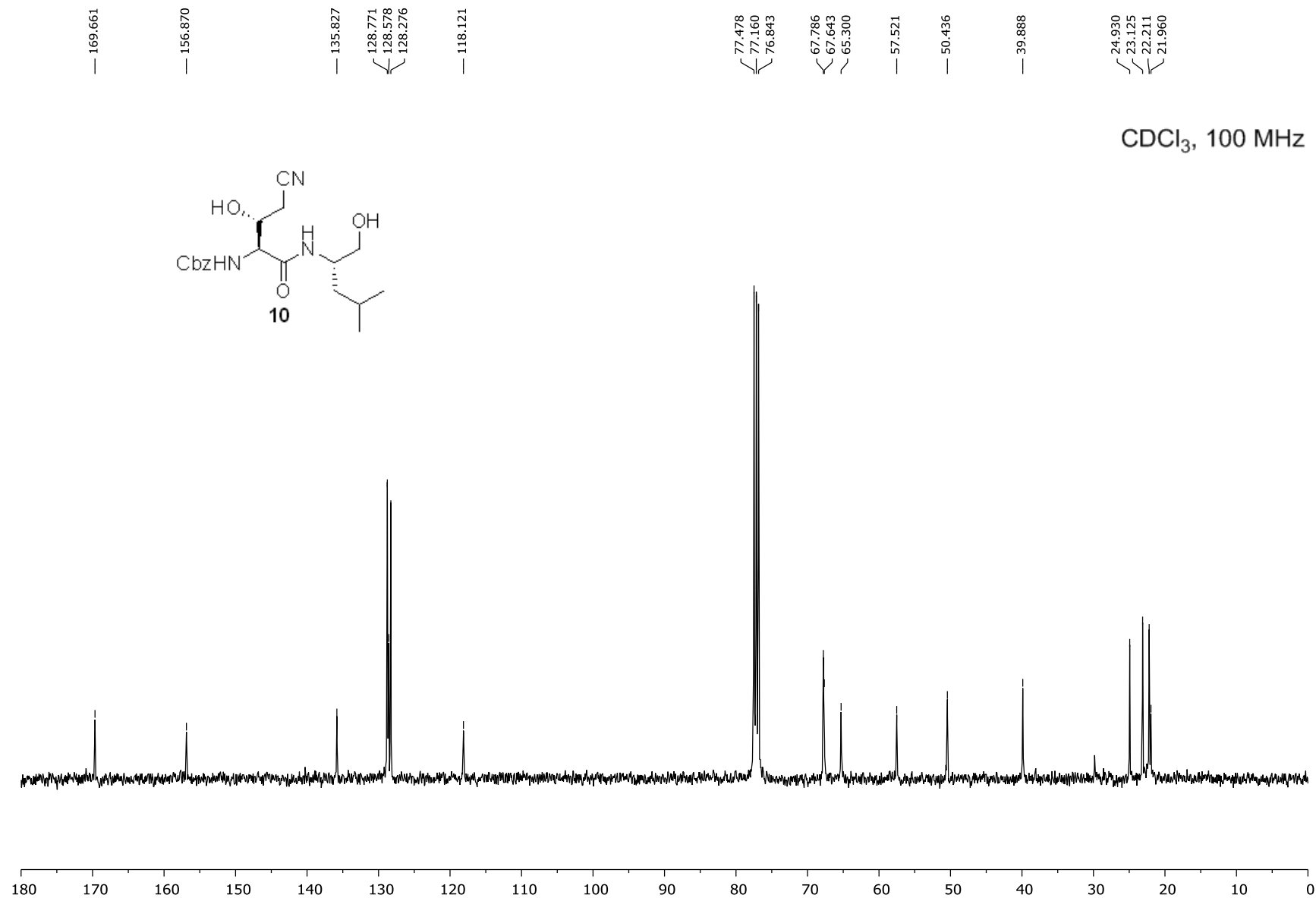
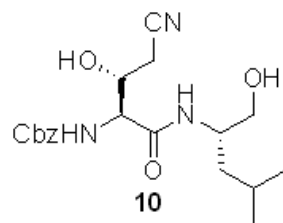


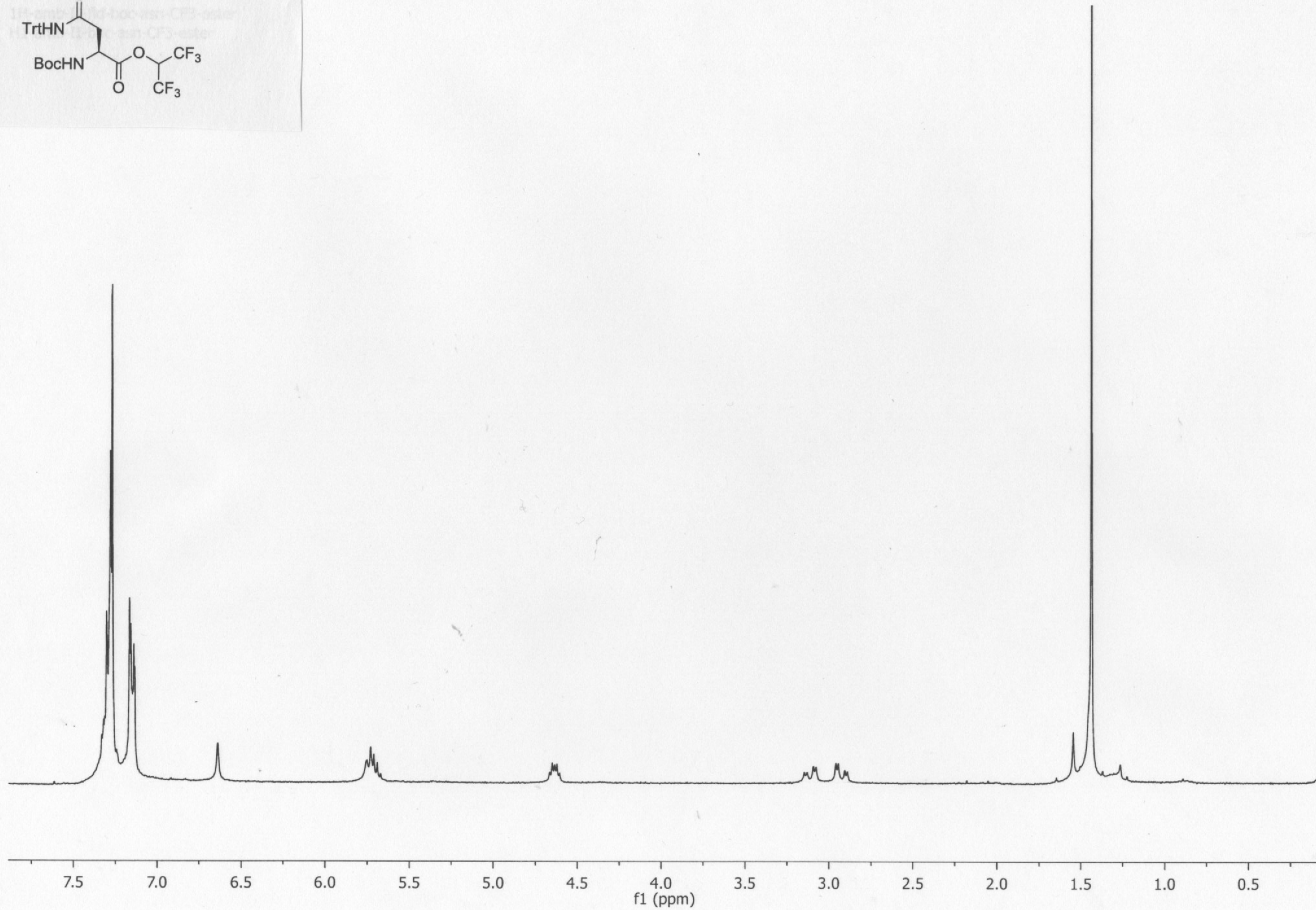
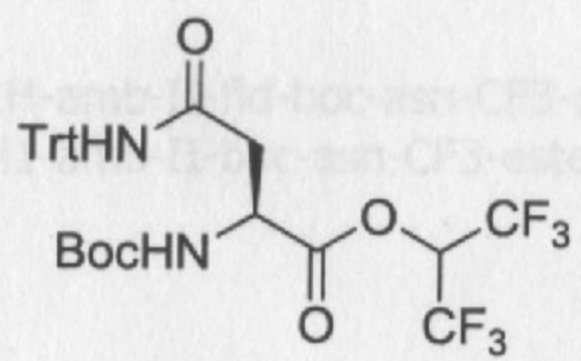
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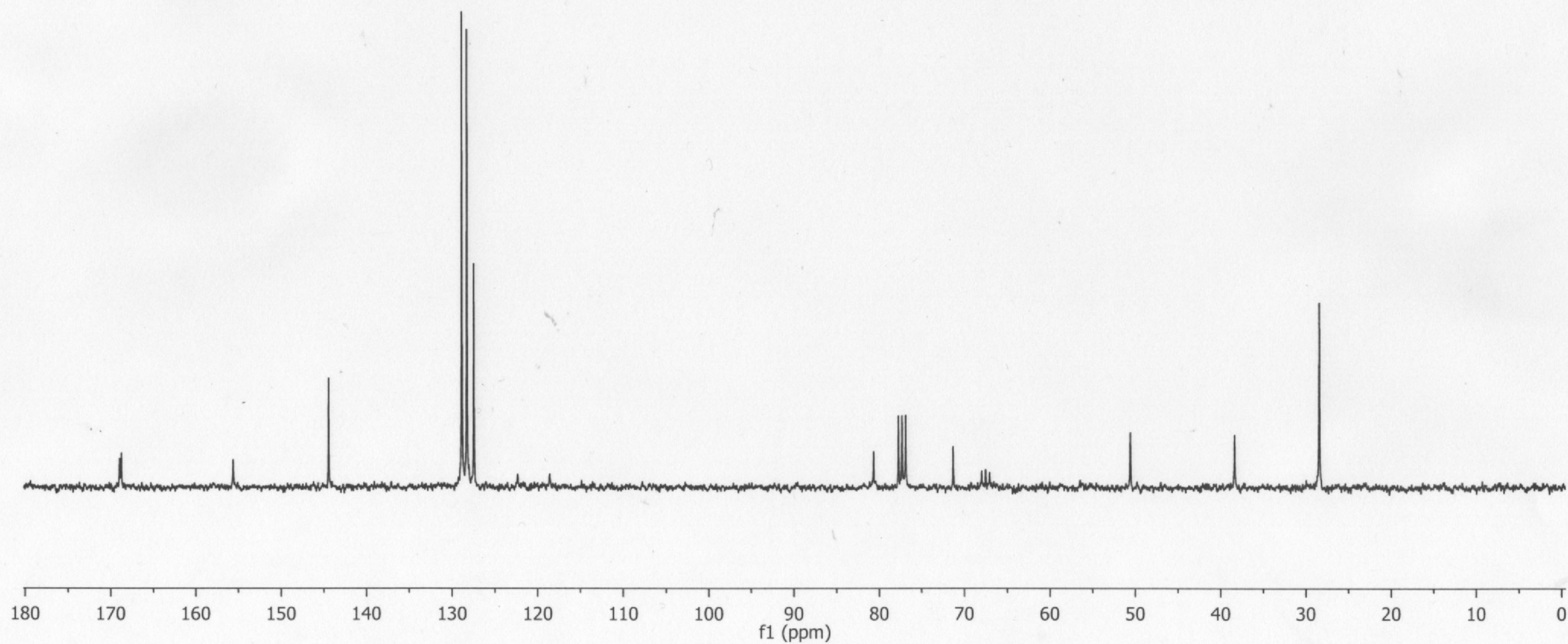
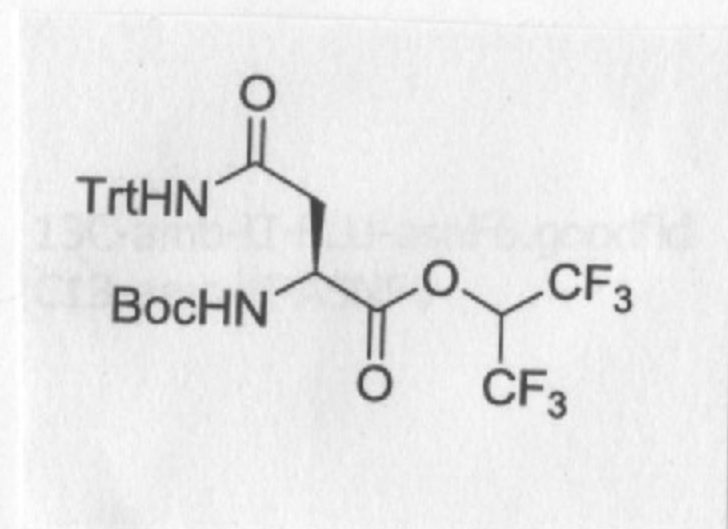




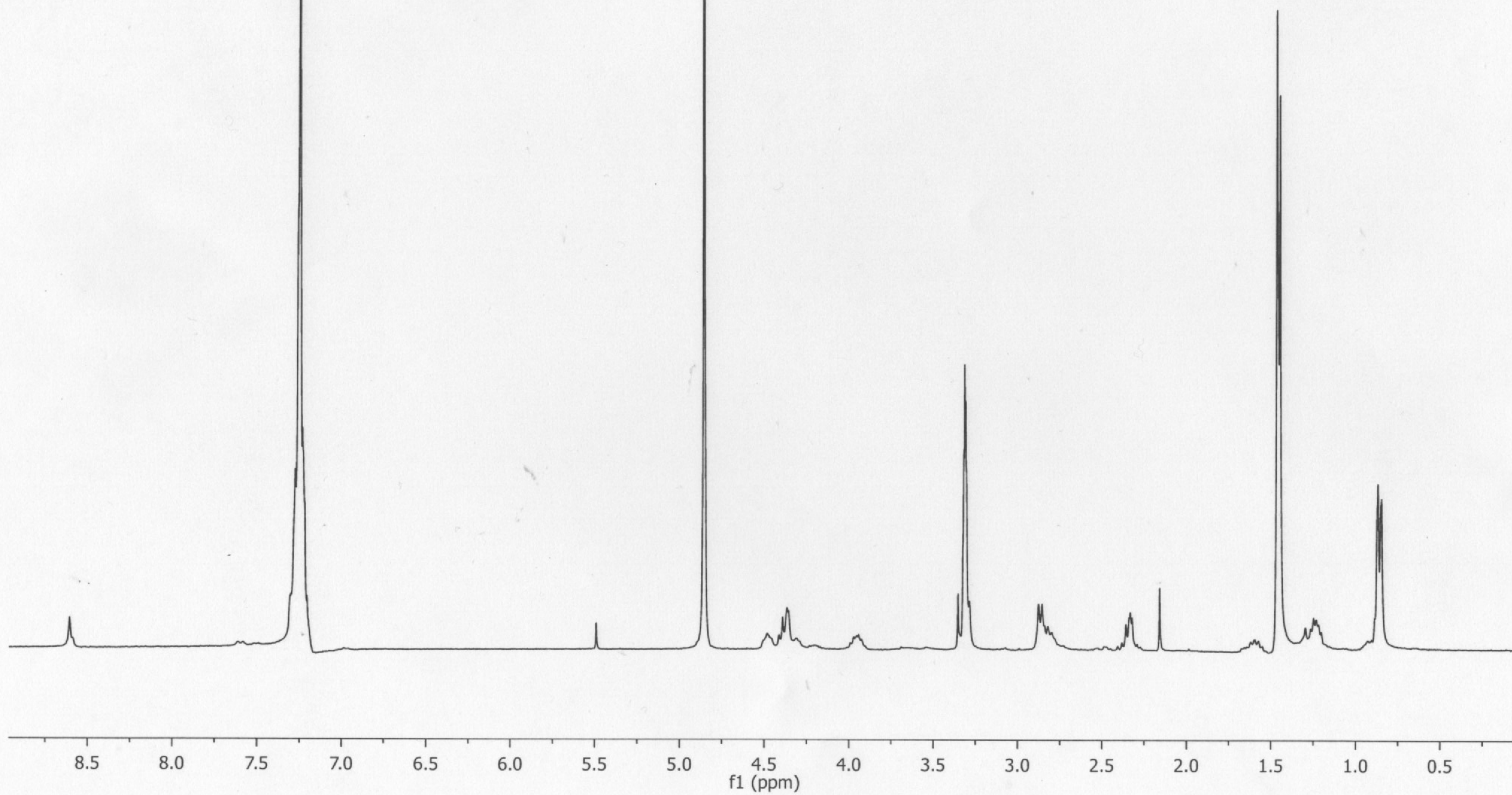
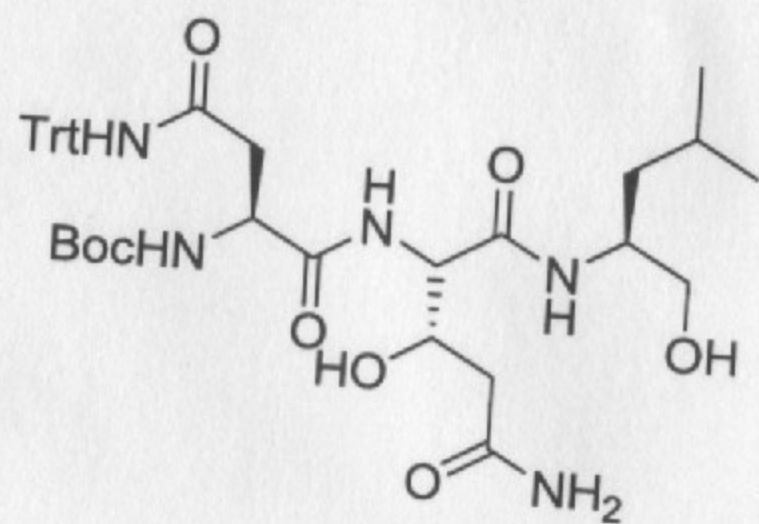




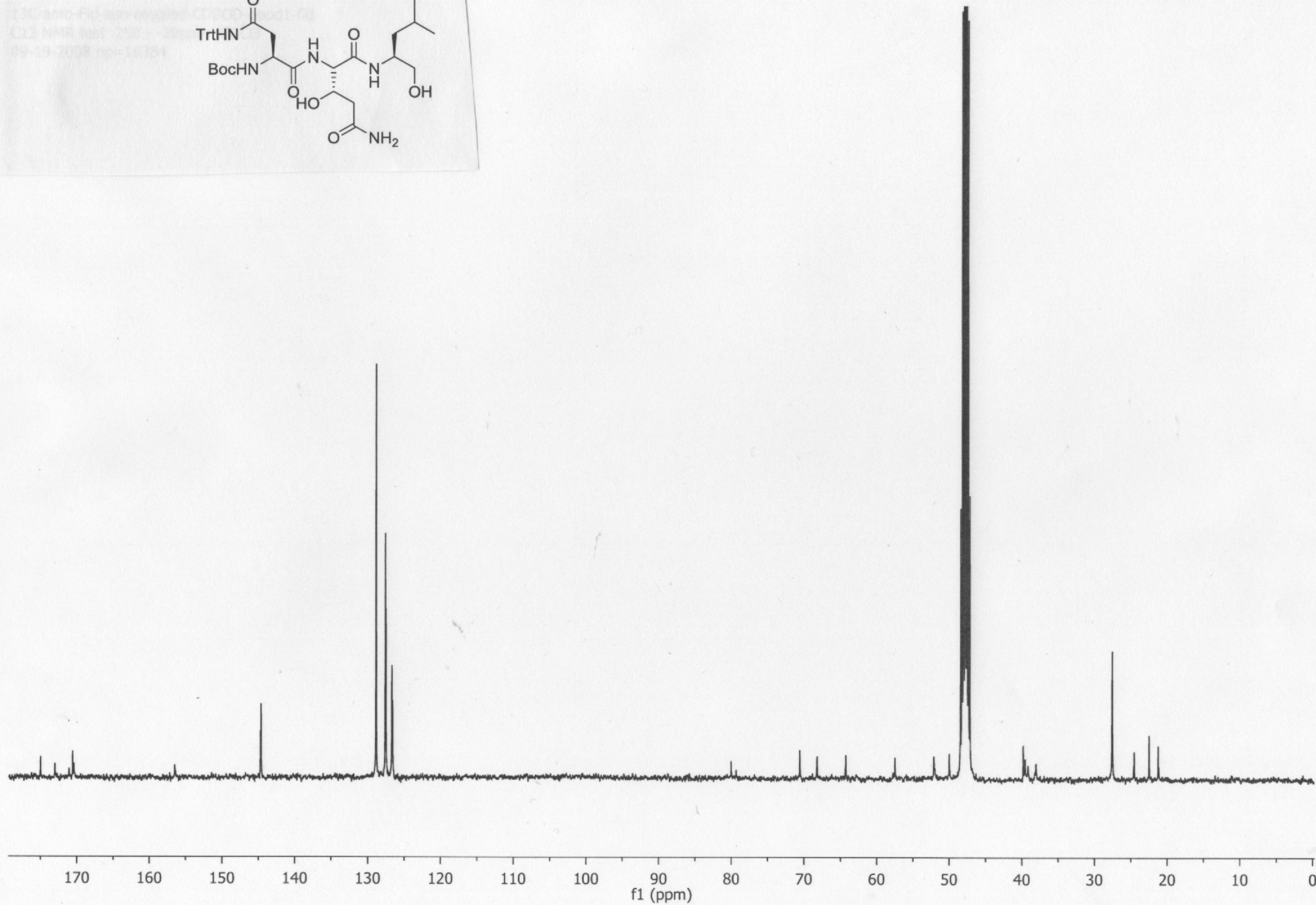
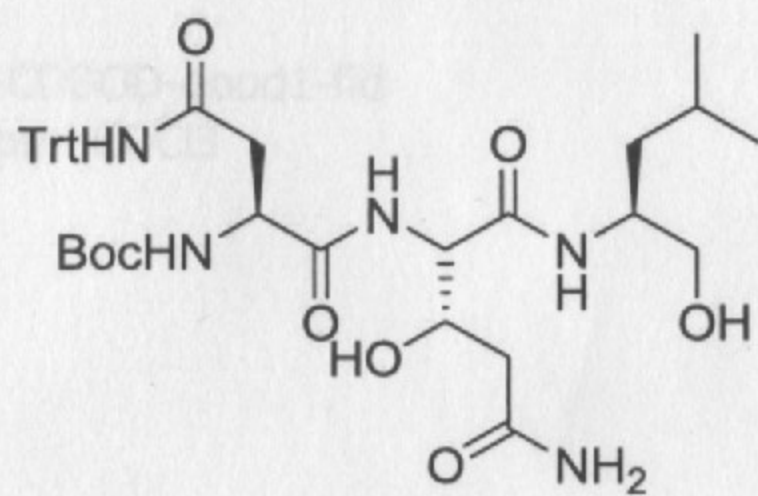


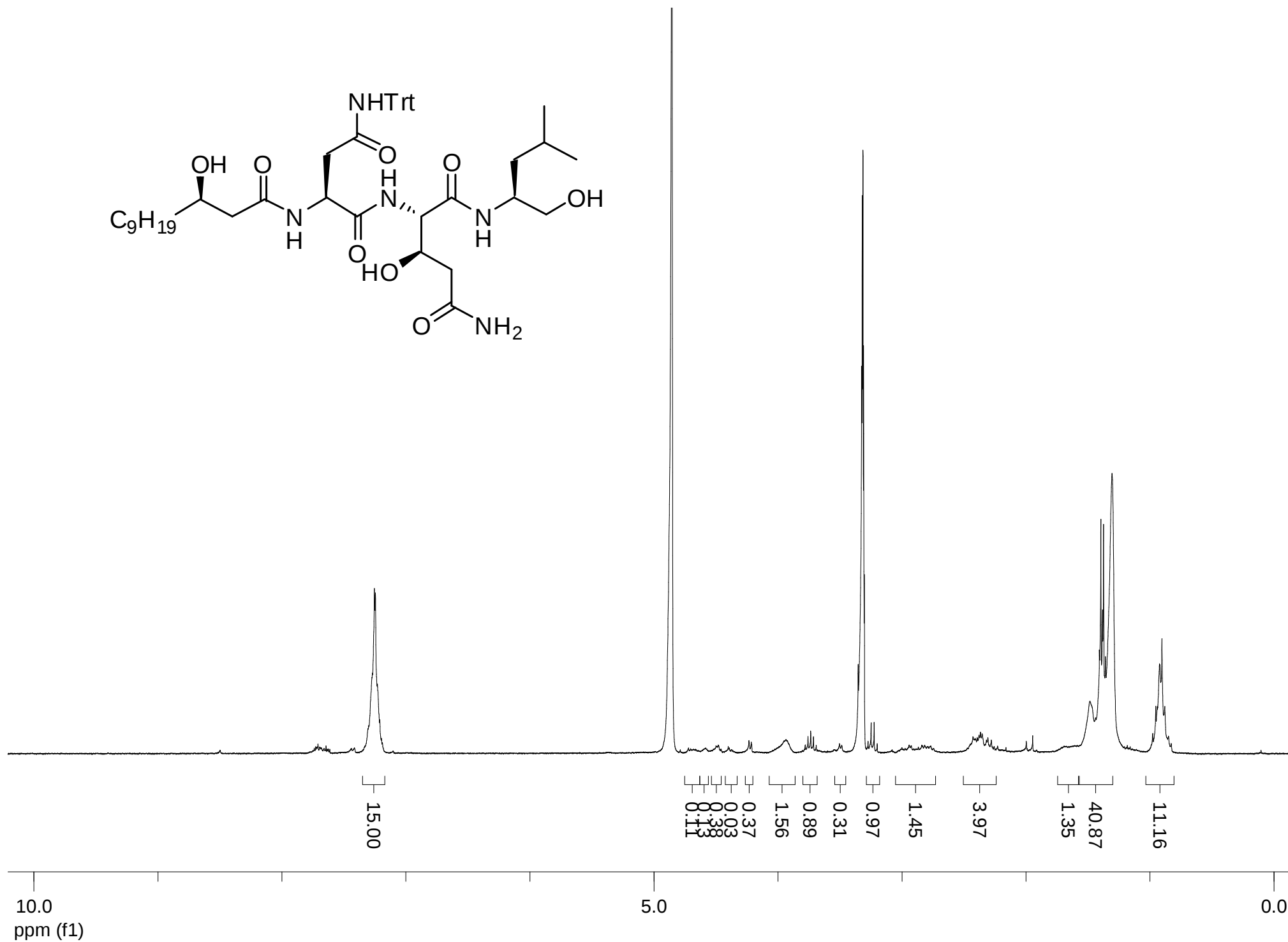
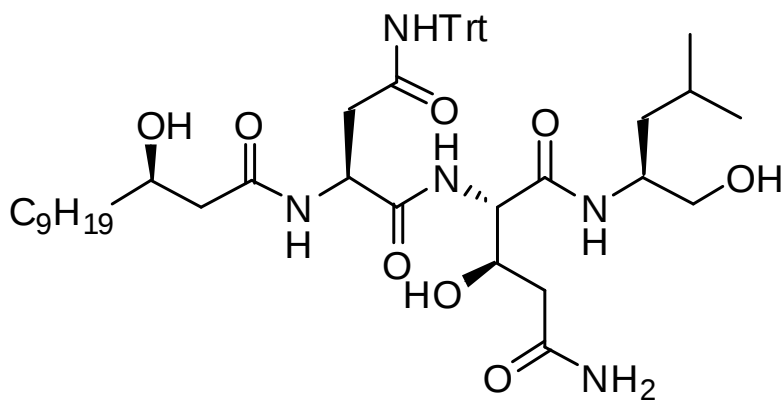


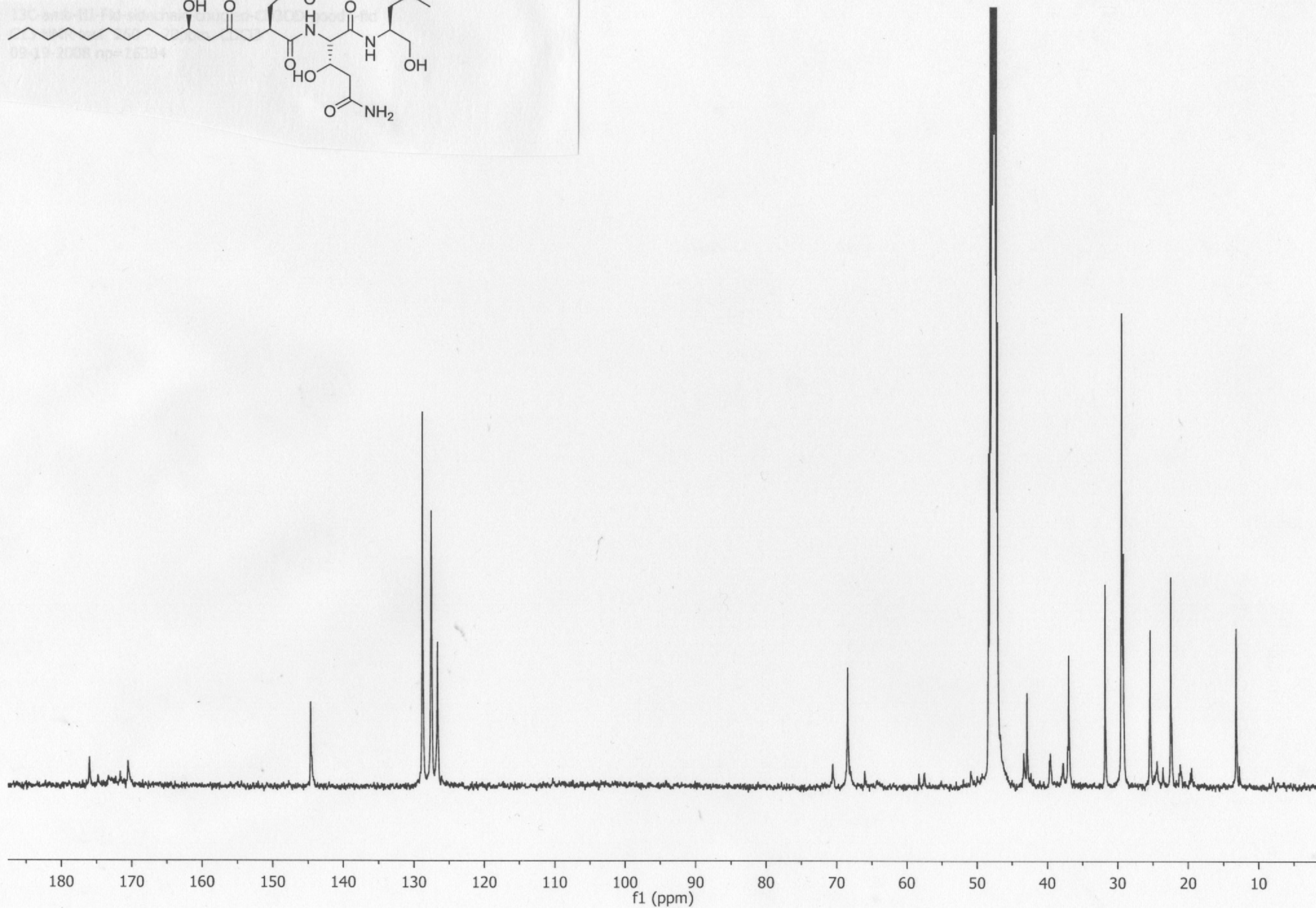
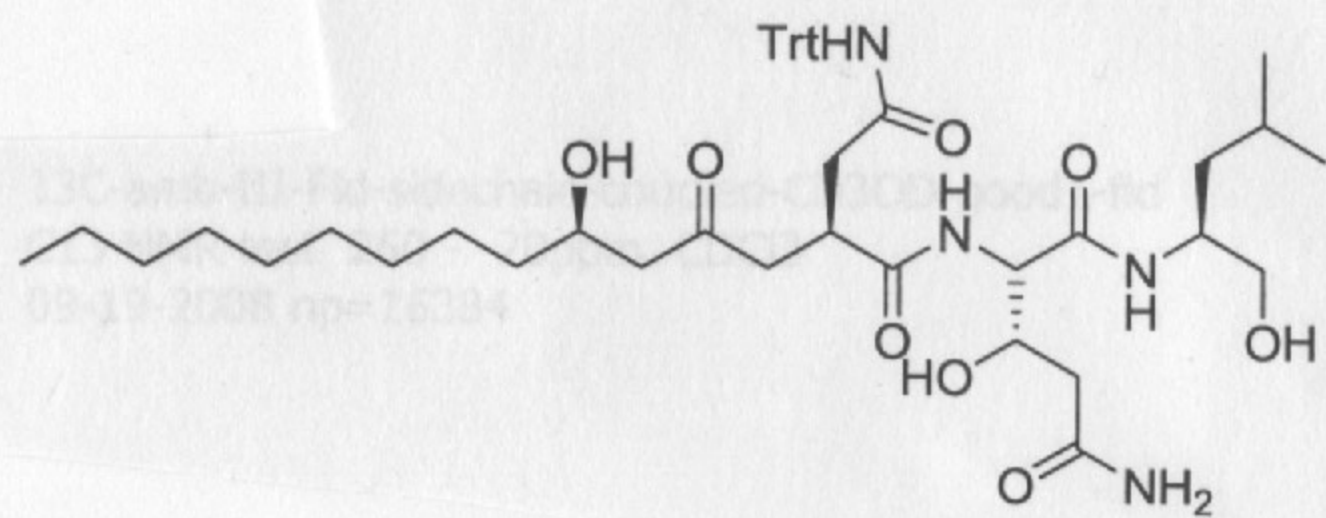
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H1-amb-III-FLD-asn-coupling-good

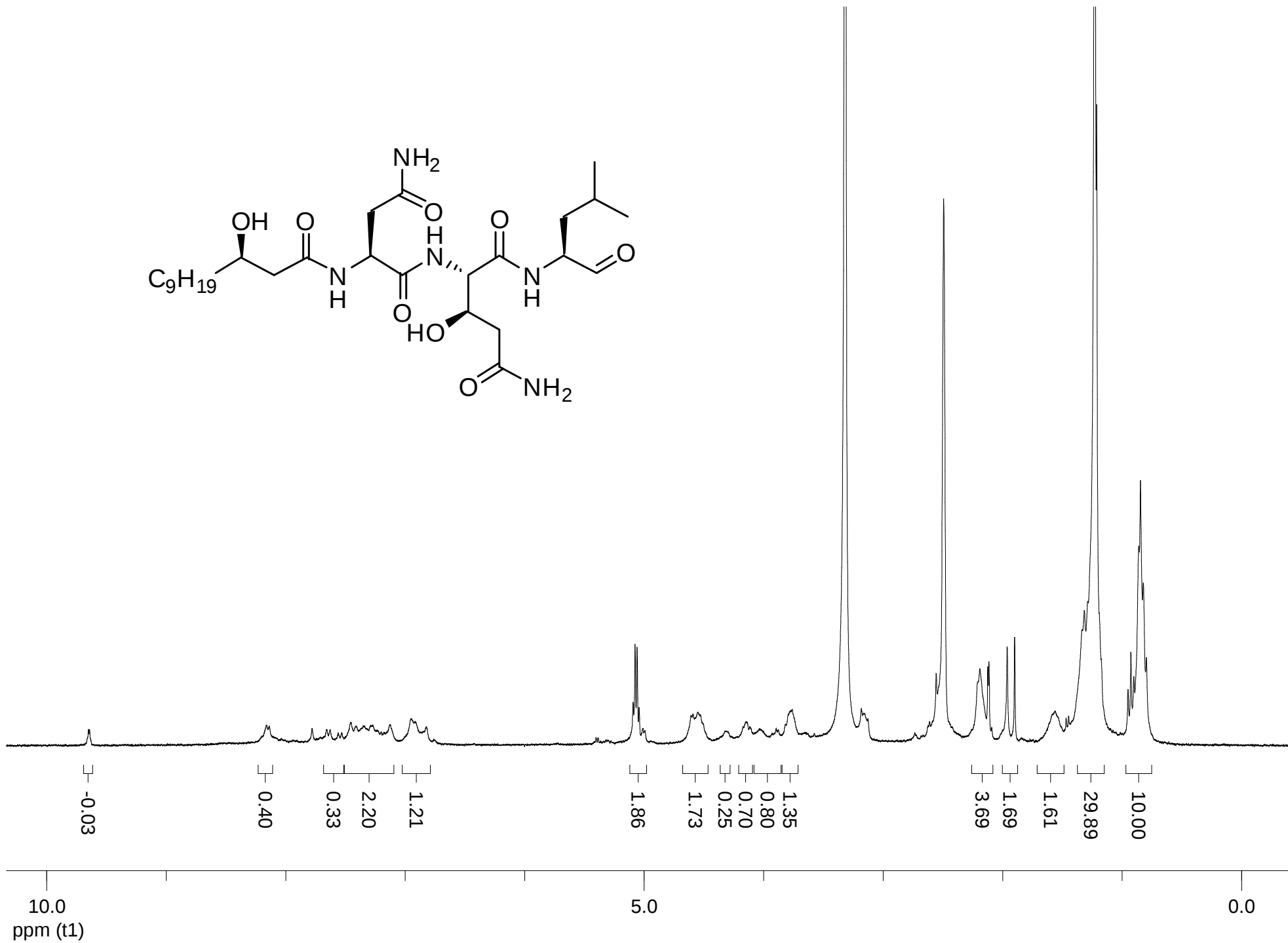
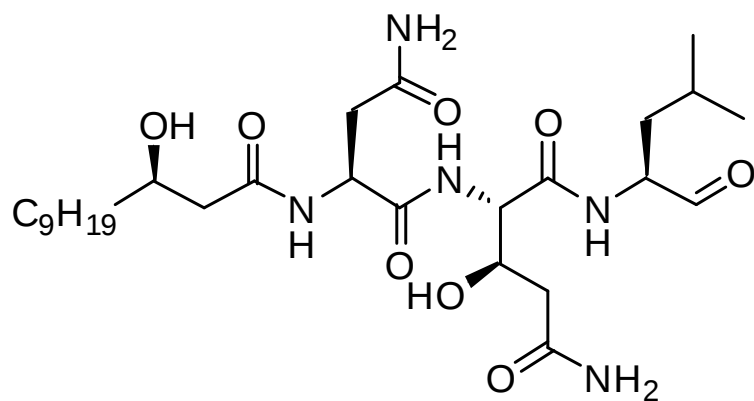


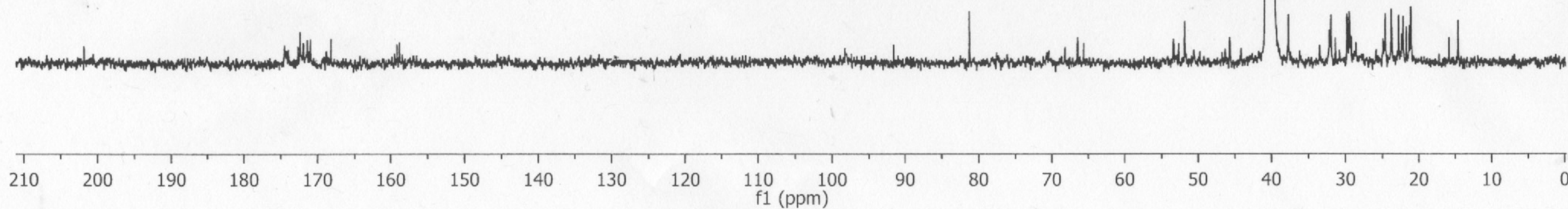
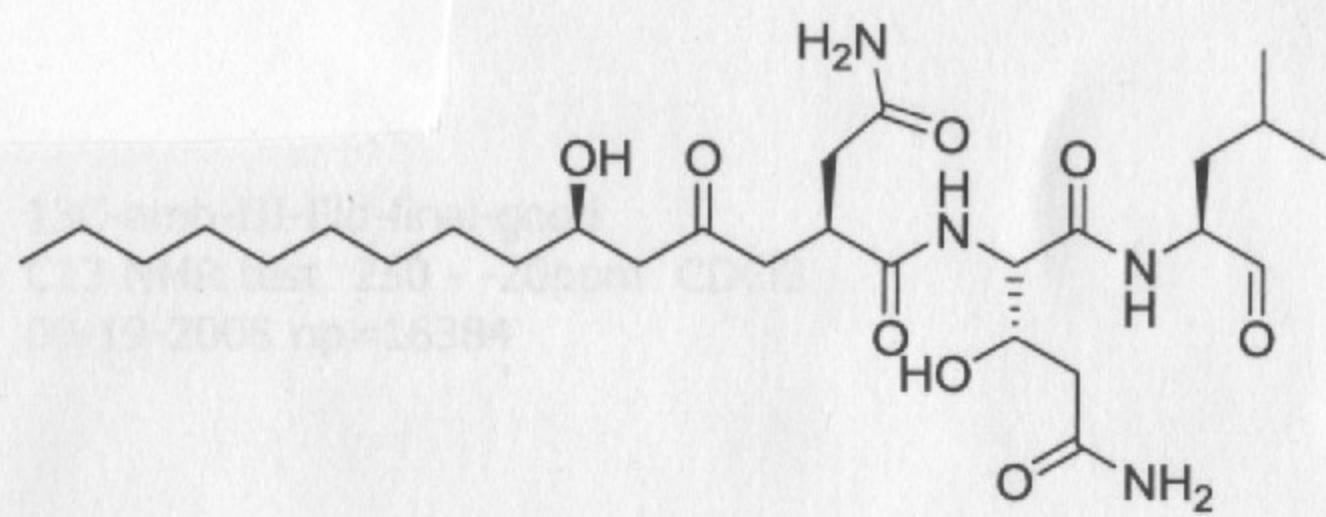
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09-19-2008 mp=163.4







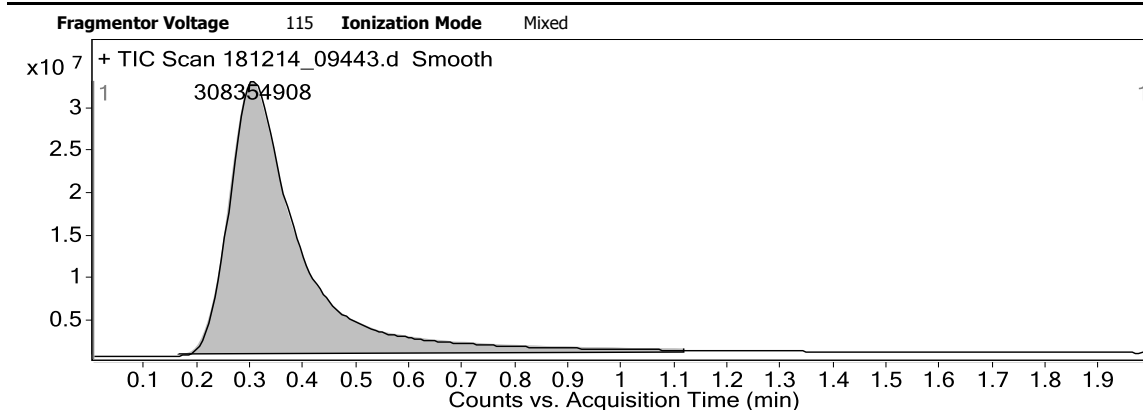
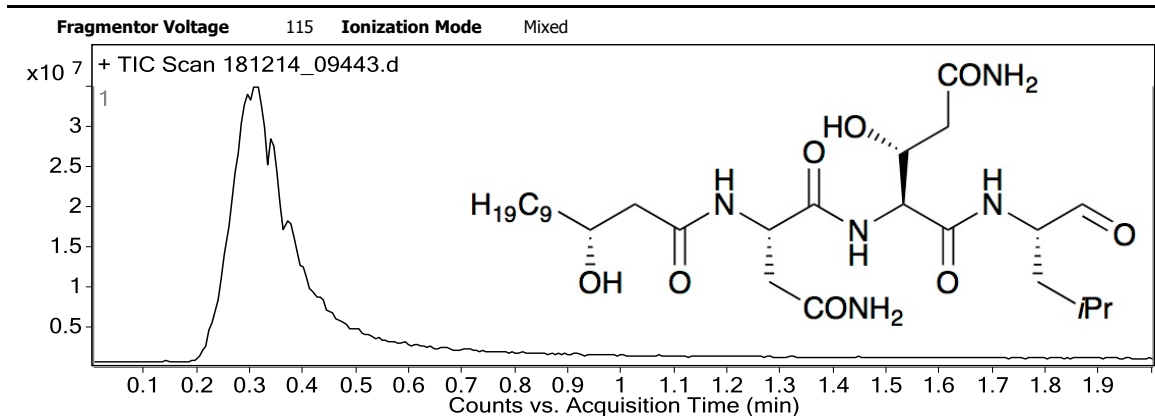




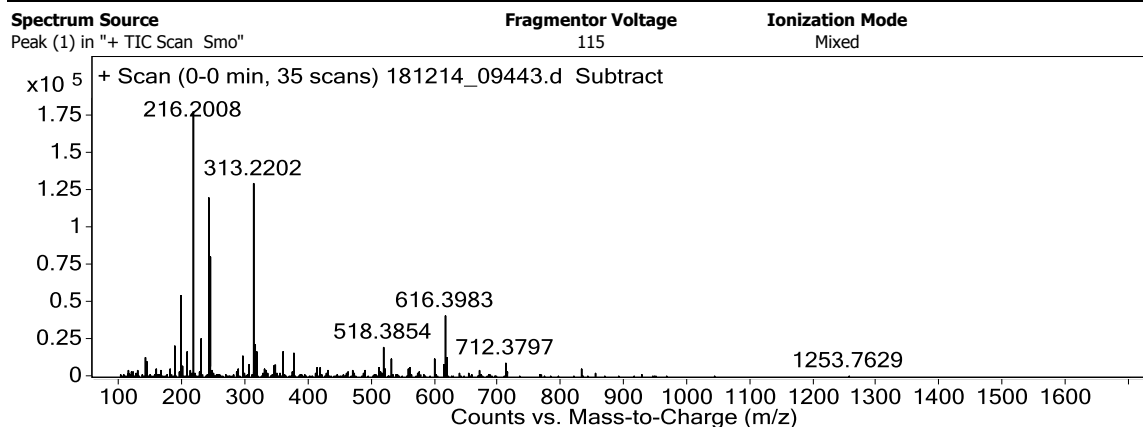
Qualitative Analysis Report

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Instrument Name	UCR Chem TOF	User Name	Pirrings Account for NIH

Acq Method	MM_MIXED_FI_POS.M	IRM Calibration Status	Success
DA Method	DA_QUAL_ALL PURPOSE.M	Comment	FN Easy-Access Method: 'FI MM Mixed positive ion' C27H49N5O8



User Spectra



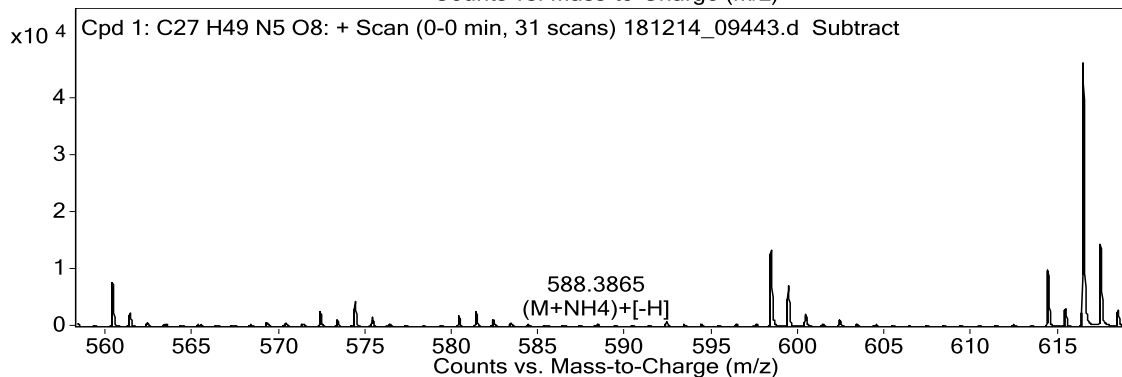
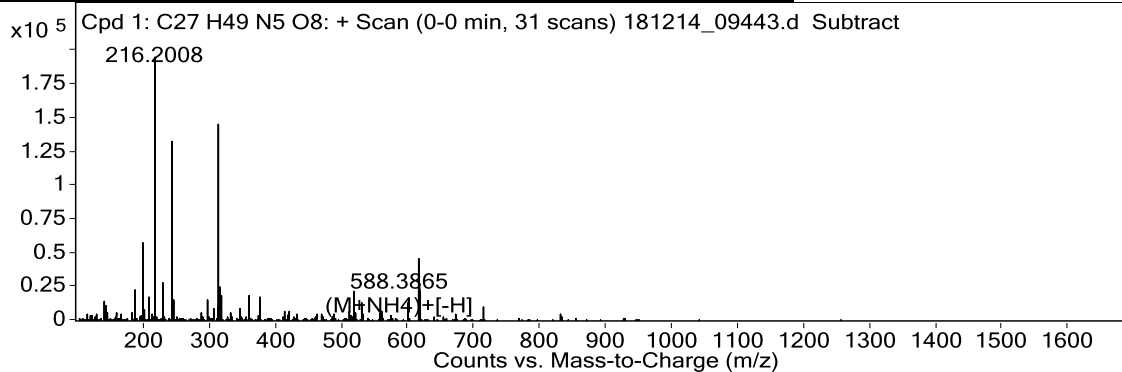
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Formula Confirmation Report

Sample Name	fid-final	Position	P1-E-04	Instrument Name	UCR Chem TOF
User Name	Pirrings Account for	Inj Vol	0.5	IRM Calibration Status	Success
Data Filename	181214_09443.d	ACQ Method	MM_MIXED_FI_POS.M	Comment	FN Easy-Access Method: 'FI MM Mixed posi

Acquired Time 12/18/2014 1:20:43 PM

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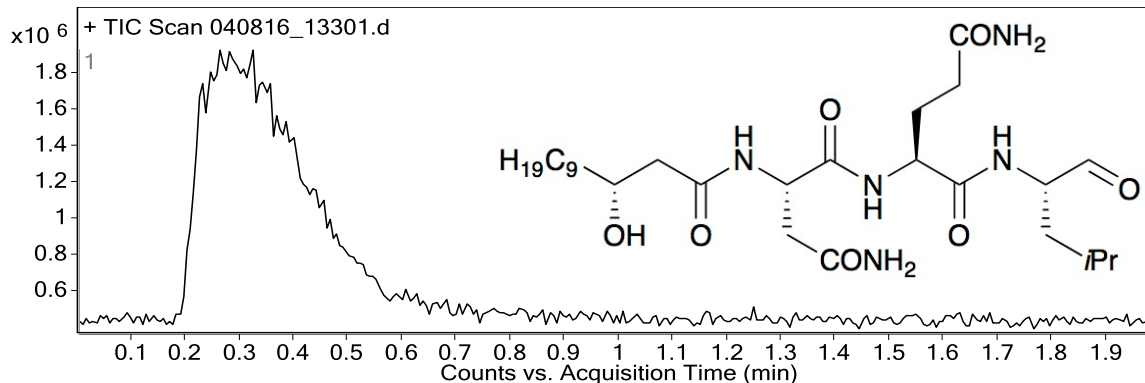


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216.2008				195587		
216.3638				6768		
217.2018				24670		
588.3865	588.3841	4.14	1	390	C27 H52 N6 O8	(M+NH4)+[-H]
589.3945	589.3919	4.31	1	302	C27 H53 N6 O8	(M+NH4)+

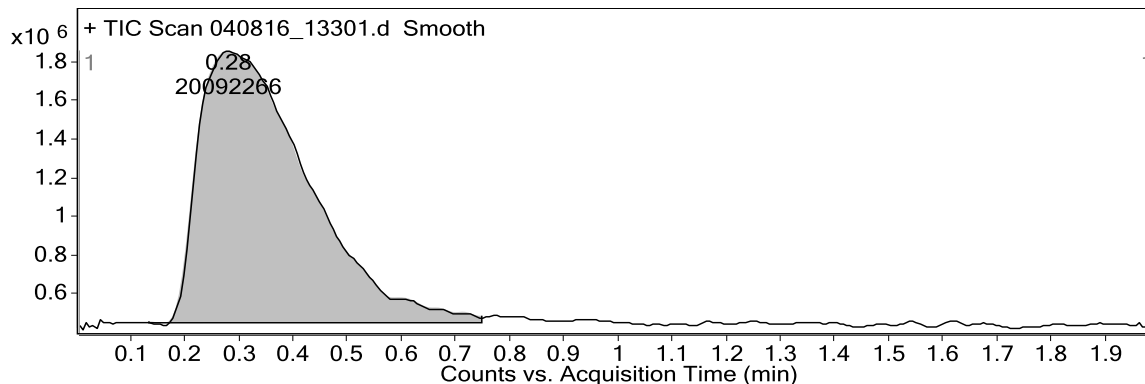
Qualitative Analysis Report

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Instrument Name	UCR Chem TOF	User Name	Pirrungs NSF Account
Acq Method	MM_MIXED_FI_POS.M	IRM Calibration Status	Success
DA Method	DA_QUAL_ALL PURPOSE.M	Comment	FELLUTAMIDE B Easy-Access Method: 'FI MM Mixed positive ion' C27H49N5O7

Fragmentor Voltage 115 **Ionization Mode** Mixed

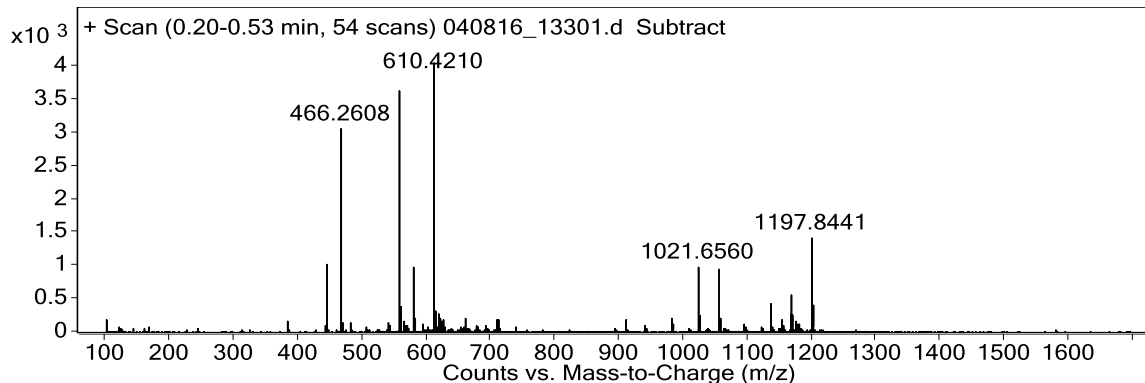


Fragmentor Voltage 115 **Ionization Mode** Mixed



User Spectra

Spectrum Source Peak (1) in "+ TIC Scan Smo"
Fragmentor Voltage 115 **Ionization Mode** Mixed



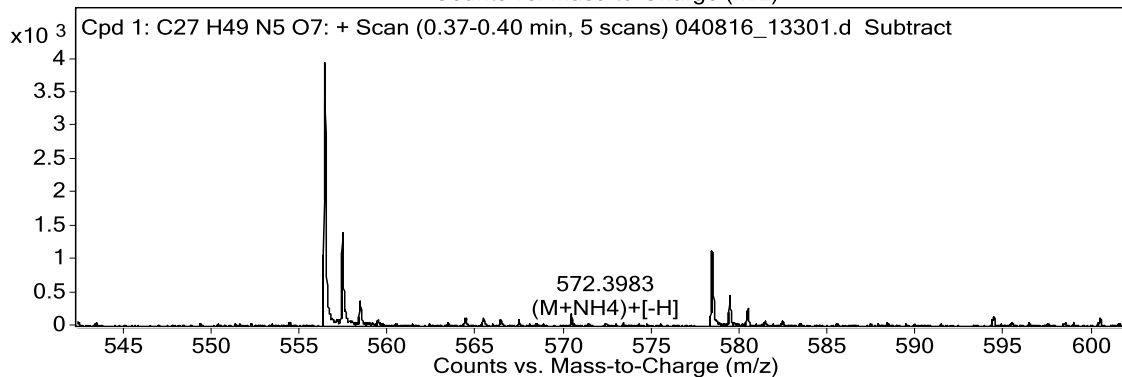
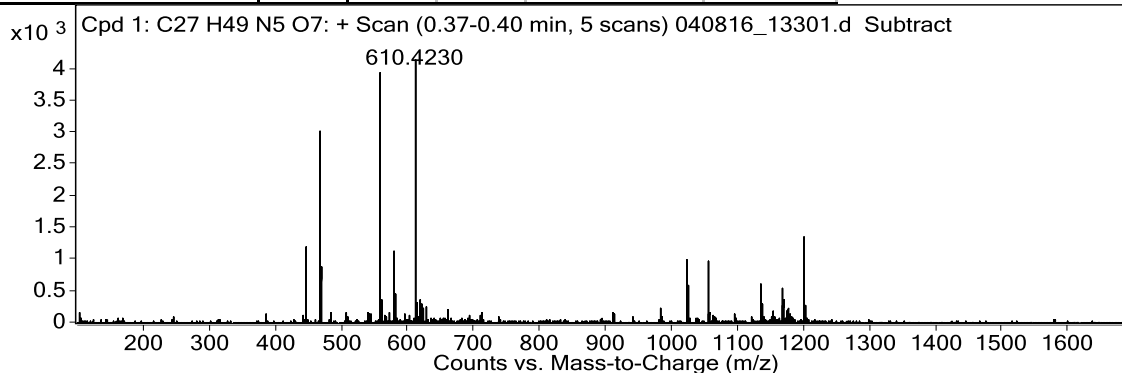
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Formula Confirmation Report

Sample Name	FZ-2-173	Position	P1-A-07	Instrument Name	UCR Chem TOF
User Name	Pirrungs NSF Account	Inj Vol	0.5	IRM Calibration Status	Success
Data Filename	040816_13301.d	ACQ Method	MM_MIXED_FI_POS.M	Comment	FELLUTAMIDE B Easy- Access Method: 'FI MM

Acquired Time 8/4/2016 2:47:03 PM

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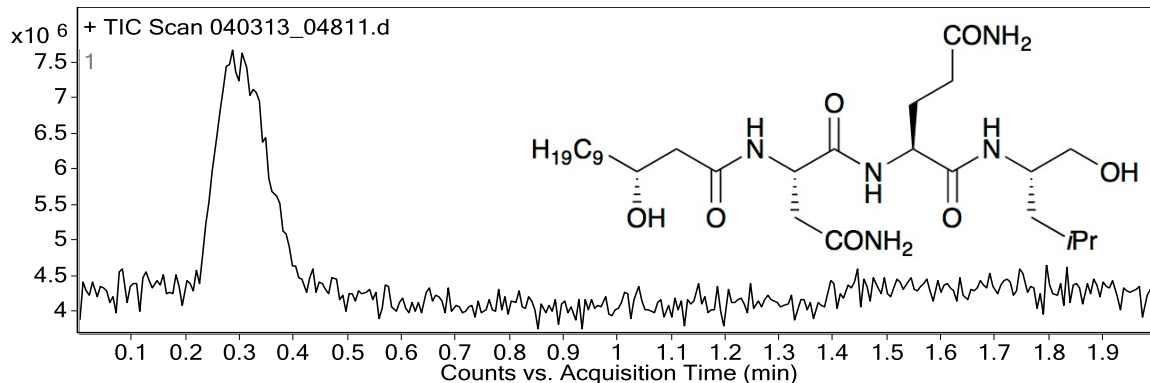


m/z	Calc m/z	Diff(ppm)	z	Abund	Formula	Ion
572.3983	572.3892	15.93	1	52	C27 H52 N6 O7	(M+NH4)+[-H]
610.423				4142		
610.8688				183		
610.9545				198		
611.4226				1338		
612.4376				346		

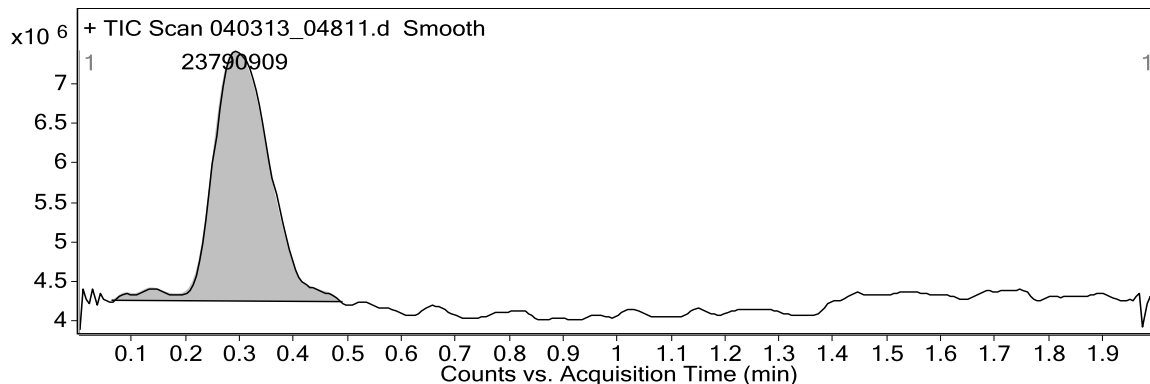
Qualitative Analysis Report

Data Filename	040313_04811.d	Sample Name	FZ-2-175
Sample Type	Unknown	Position	P1-A-01
Instrument Name	UCR Chem TOF	User Name	Pirrungs DOD Acct.
Acq Method	MM_MIXED_FI_POS.M	IRM Calibration Status	Success
DA Method	DA_QUAL_ALL PURPOSE.M	Comment	Fellutamide C Easy-Access Method: 'FI MM Mixed positive ion' C27H51N5O7

Fragmentor Voltage 115 **Ionization Mode** Mixed

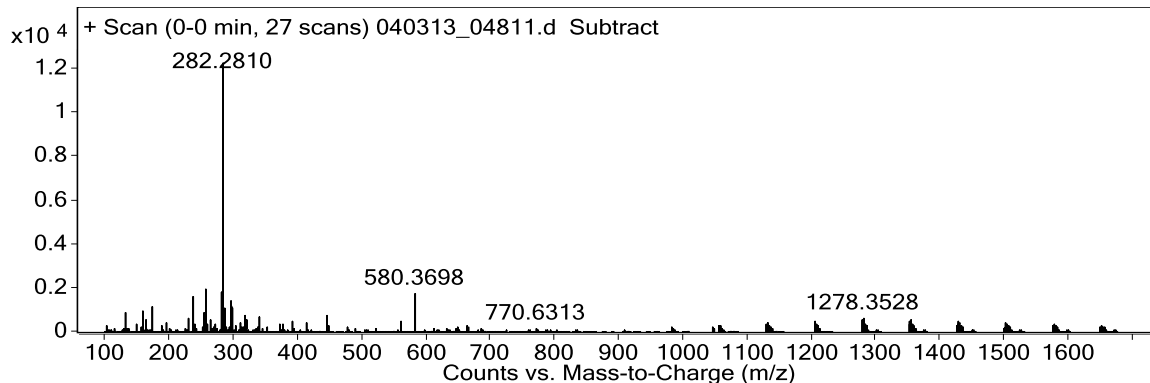


Fragmentor Voltage 115 **Ionization Mode** Mixed



User Spectra

Spectrum Source Peak (1) in "+ TIC Scan Smo"
Fragmentor Voltage 115 **Ionization Mode** Mixed



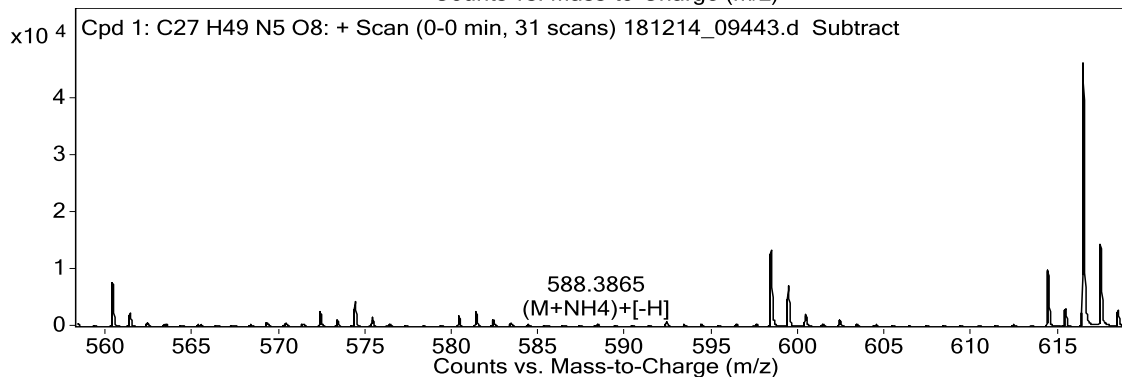
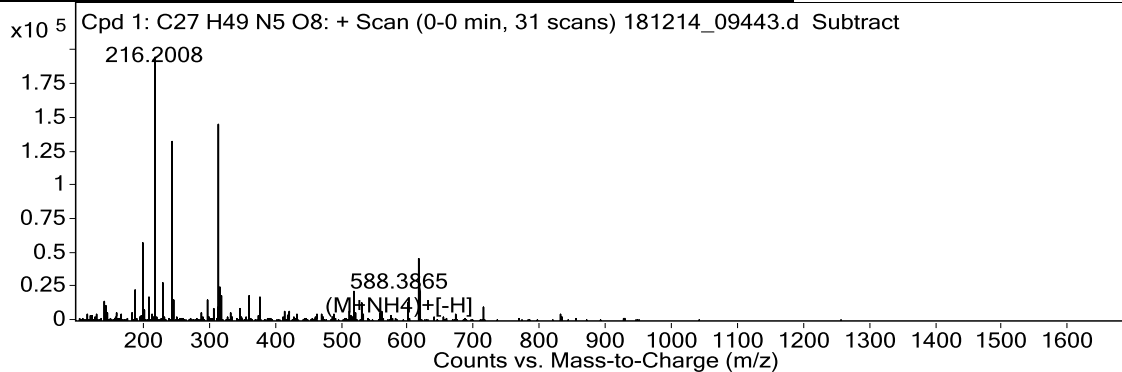
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Formula Confirmation Report

Sample Name	fid-final	Position	P1-E-04	Instrument Name	UCR Chem TOF
User Name	Pirrings Account for	Inj Vol	0.5	IRM Calibration Status	Success
Data Filename	181214_09443.d	ACQ Method	MM_MIXED_FI_POS.M	Comment	FN Easy-Access Method: 'FI MM Mixed posi

Acquired Time 12/18/2014 1:20:43 PM

Compound Label	RT	Mass	Abund	Formula	Tgt Mass
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m/z	Calc m/z	Diff(ppm)	z	Abund	Formula	Ion
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216.2008				195587		
216.3638				6768		
217.2018				24670		
588.3865	588.3841	4.14	1	390	C27 H52 N6 O8	(M+NH4)+[-H]
589.3945	589.3919	4.31	1	302	C27 H53 N6 O8	(M+NH4)+