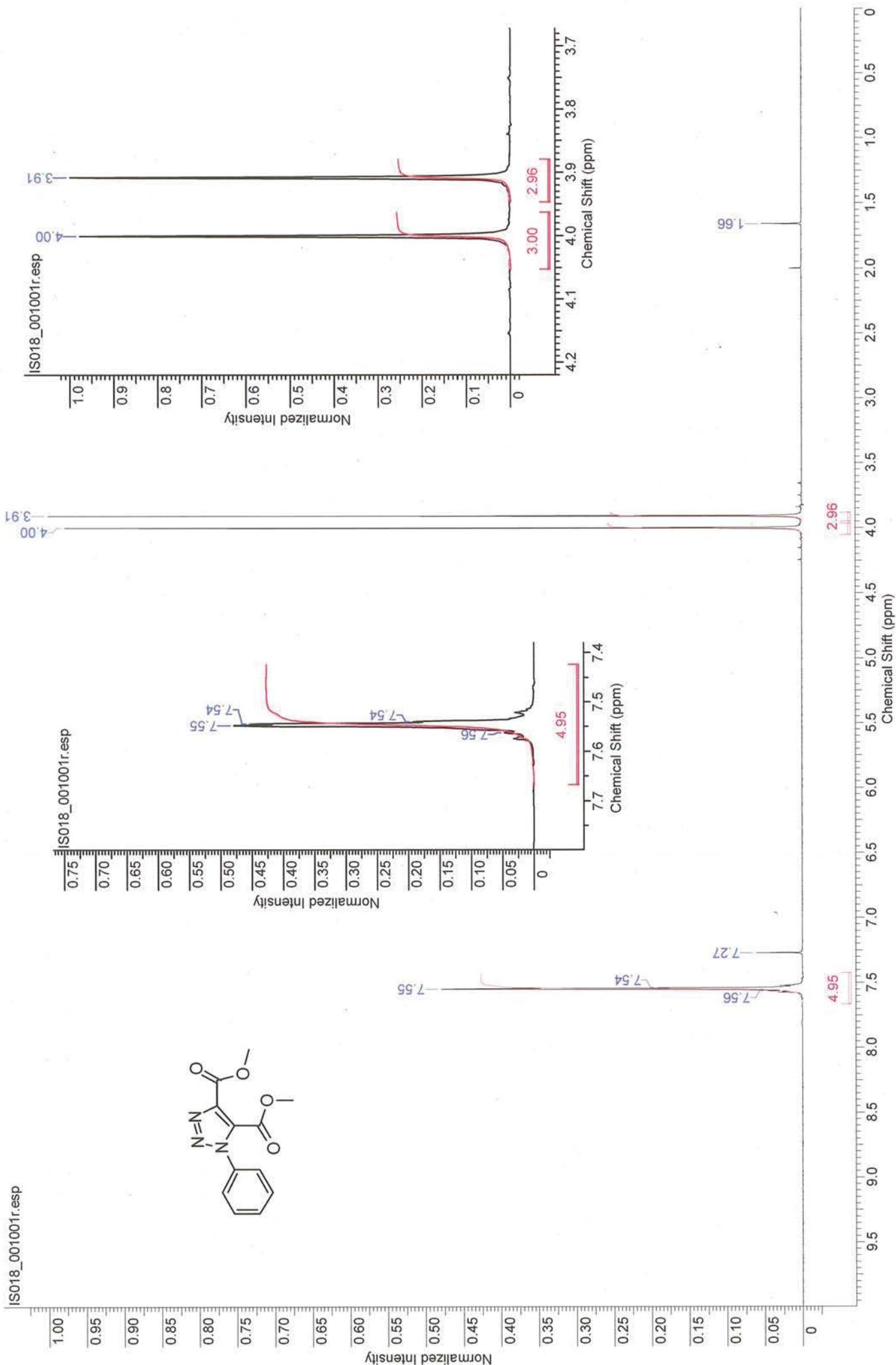
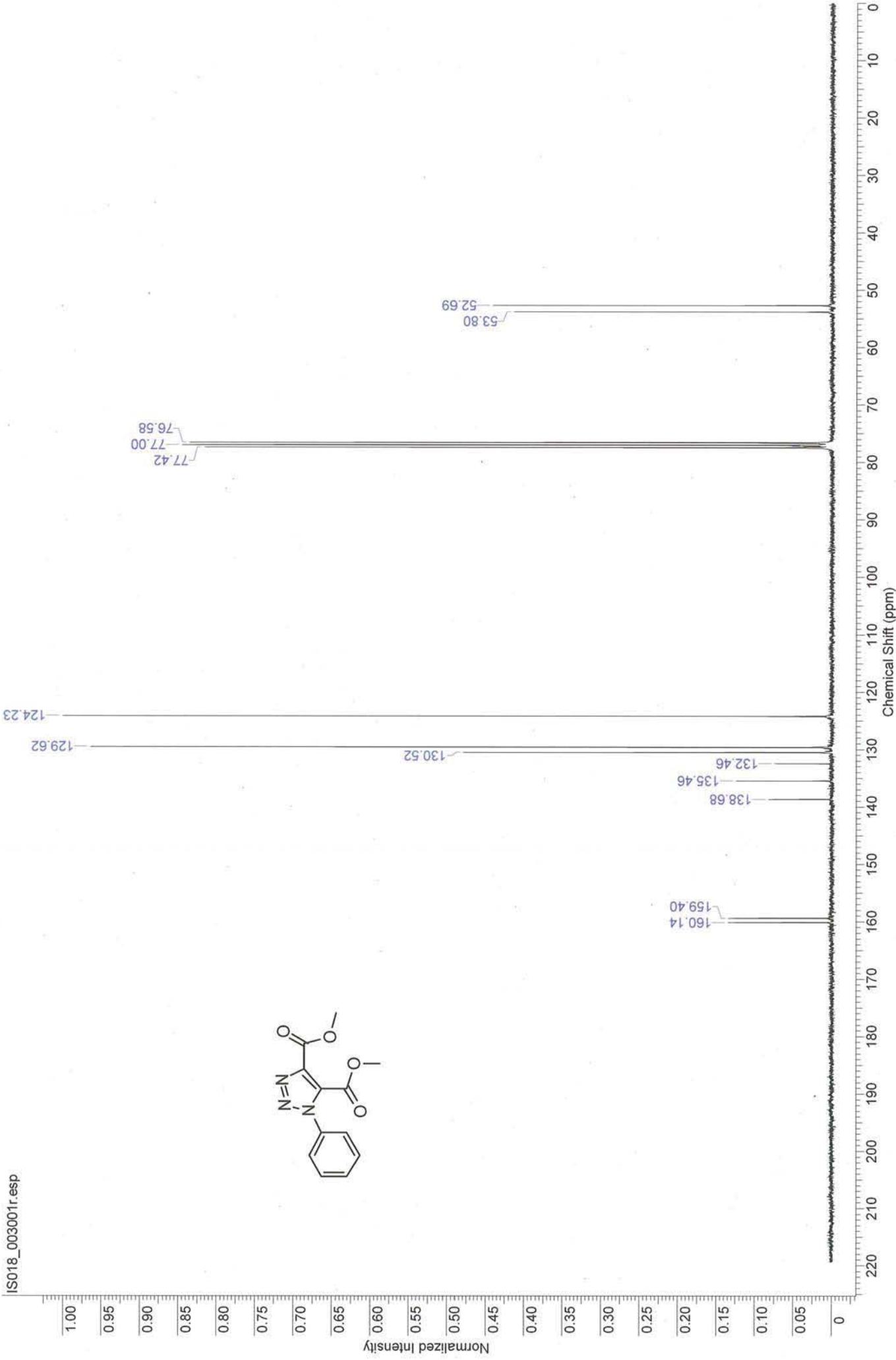


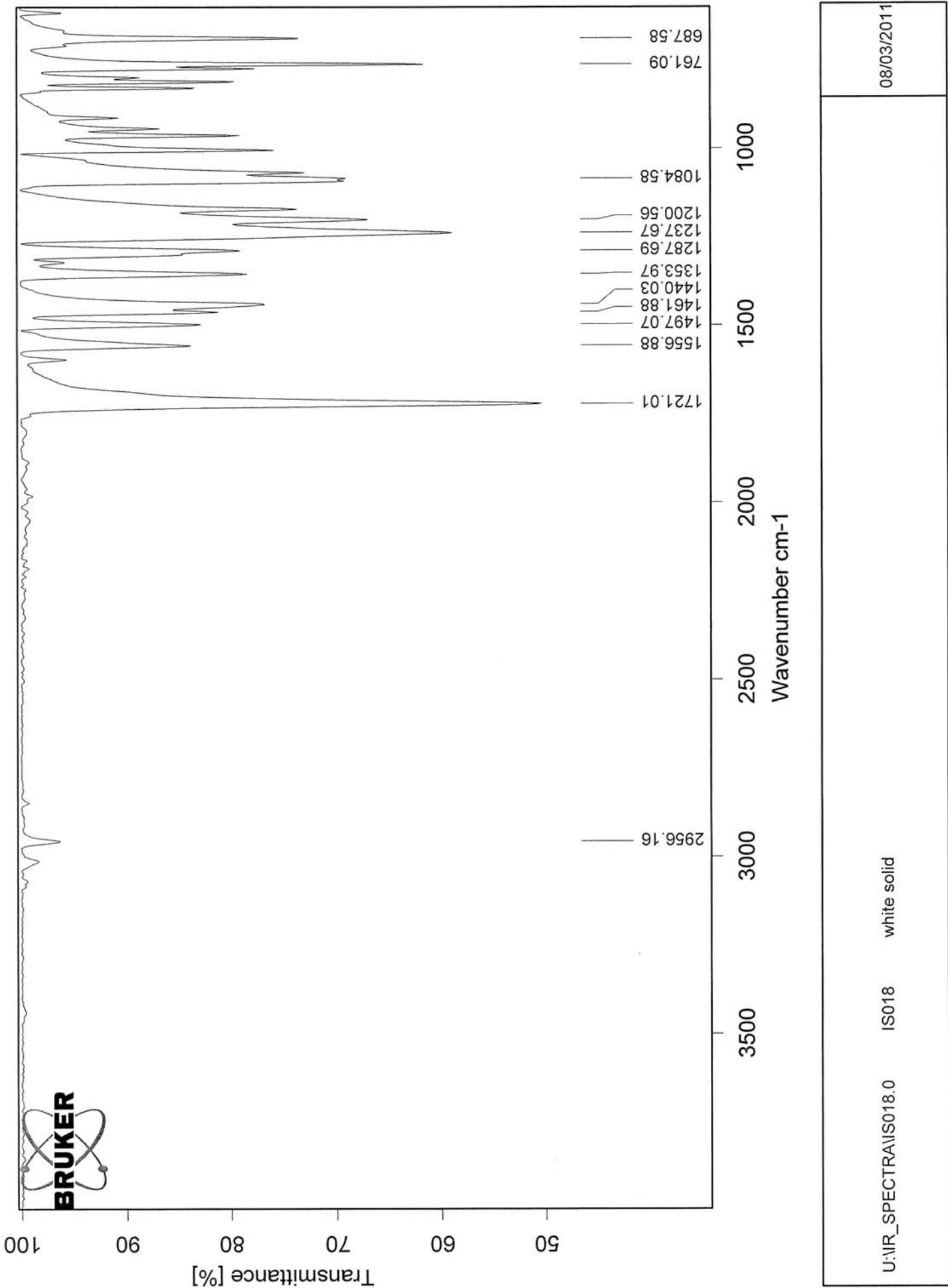
Table of Contents

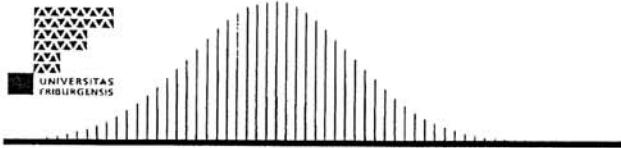
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 1a	S3
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 1b	S7
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 1c	S11
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 1c'	S15
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 1d	S19
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 1d'	S23
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 1e	S27
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 1e'	S31
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 1f	S35
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 1f'	S39
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 1g	S43
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 1g'	S47
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 1h	S51
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 1h'	S55
^1H NMR and IR spectra of 1i	S59
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 1j	S61
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 1k	S65
^1H NMR spectrum of 1l	S69
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 1m	S70
^1H NMR spectrum of 1n	S74
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 2a	S75
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 2b	S79
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 2c	S83
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 2d	S87
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 2e	S91
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 2f	S95
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 2g	S99
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 2h	S103
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 2i	S107
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 2j	S111
^1H NMR, ^{13}C NMR, IR and ESI-HRMS spectra of 2k	S115

¹ H NMR, ¹³ C NMR and IR spectra of 2l	S119
¹ H NMR, ¹³ C NMR, IR and ESI-HRMS spectra of 2m	S122
¹ H NMR spectrum of 2n	S126
¹ H NMR, ¹³ C NMR, IR and ESI-HRMS spectra of 4a	S127
¹ H NMR, ¹³ C NMR, IR and ESI-HRMS spectra of 4a'	S131
¹ H NMR, ¹³ C NMR, IR and ESI-HRMS spectra of 4b	S135
¹ H NMR, ¹³ C NMR, IR and ESI-HRMS spectra of 4b'	S139
¹ H NMR, ¹³ C NMR, IR and MALDI-TOF-HRMS spectra of 4c	S143
¹ H NMR, ¹³ C NMR, IR and MALDI-TOF-HRMS spectra of 4d	S147
¹ H NMR, ¹³ C NMR, IR and ESI-HRMS spectra of 4e	S151
¹ H NMR, ¹³ C NMR, IR and MALDI-TOF-HRMS spectra of 4f	S155
¹ H NMR, ¹³ C NMR, IR and ESI-HRMS spectra of 4g	S159
¹ H NMR, ¹³ C NMR, IR and ESI-HRMS spectra of 4h	S163
¹ H NMR, ¹³ C NMR, IR and EI-MS spectra of 4i	S167
¹ H NMR, ¹³ C NMR, IR and EI-MS spectra of 4j	S171
¹ H NMR, ¹³ C NMR, IR and EI-MS spectra of 4k	S175
¹ H NMR, ¹³ C NMR, IR and ESI-HRMS spectra of 5b	S179
¹ H NMR, ¹³ C NMR, IR and ESI-HRMS spectra of 6	S183
¹ H NMR, ¹³ C NMR, IR and ESI-HRMS spectra of 7	S187
¹ H NMR, ¹³ C NMR, IR and ESI-HRMS spectra of 9a	S191
¹ H NMR, ¹³ C NMR, IR and ESI-HRMS spectra of 9b	S195
¹ H NMR, ¹³ C NMR, IR and ESI-HRMS spectra of 3a	S199
¹ H NMR, ¹³ C NMR, IR and ESI-HRMS spectra of 3b	S203









BRUKER FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

ESI-MS: is-018

XMASS Mass Analysis for /Data/UNI_FR/ALIM4569_ESI/2/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

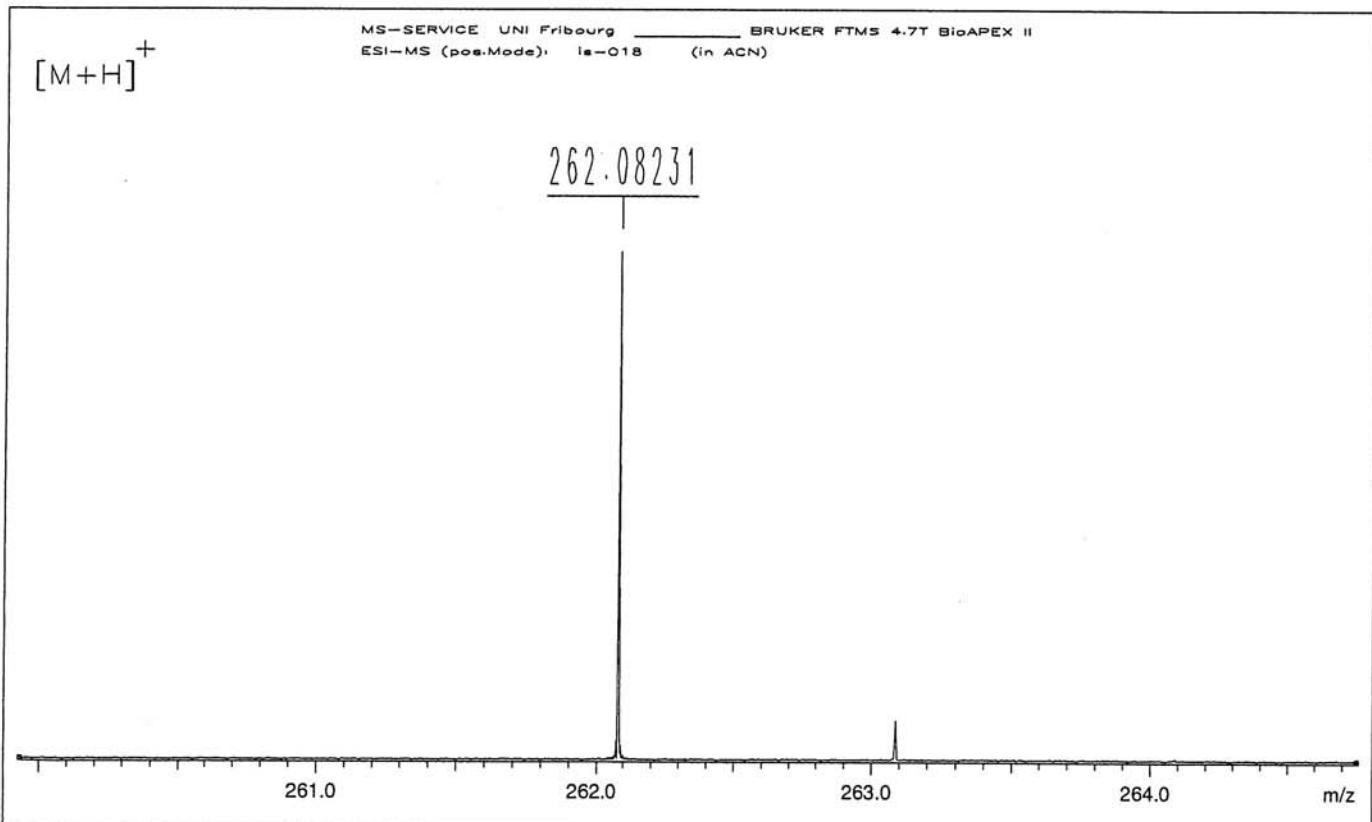
Ion mass = 262.0823110

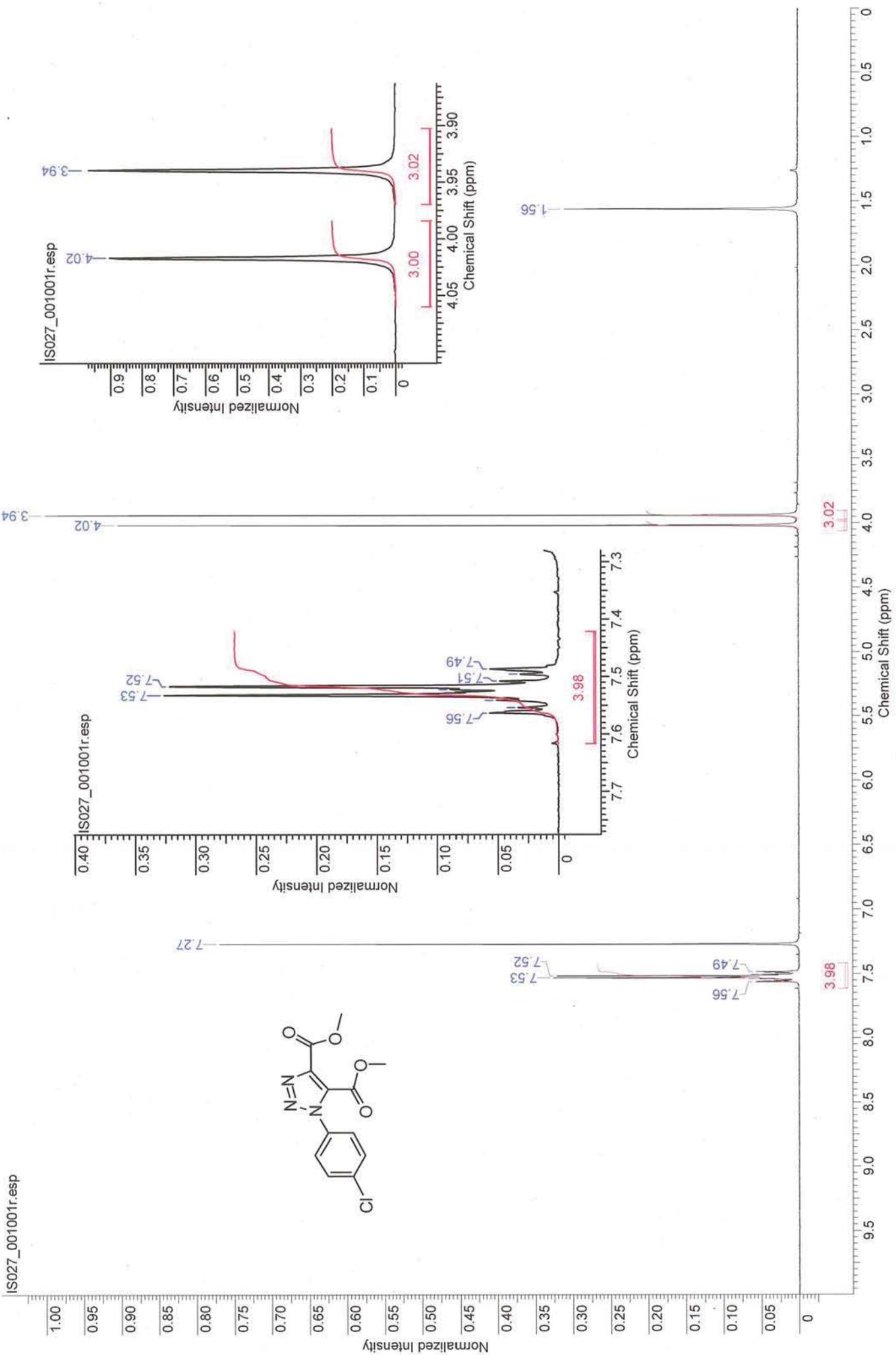
Charge = +1

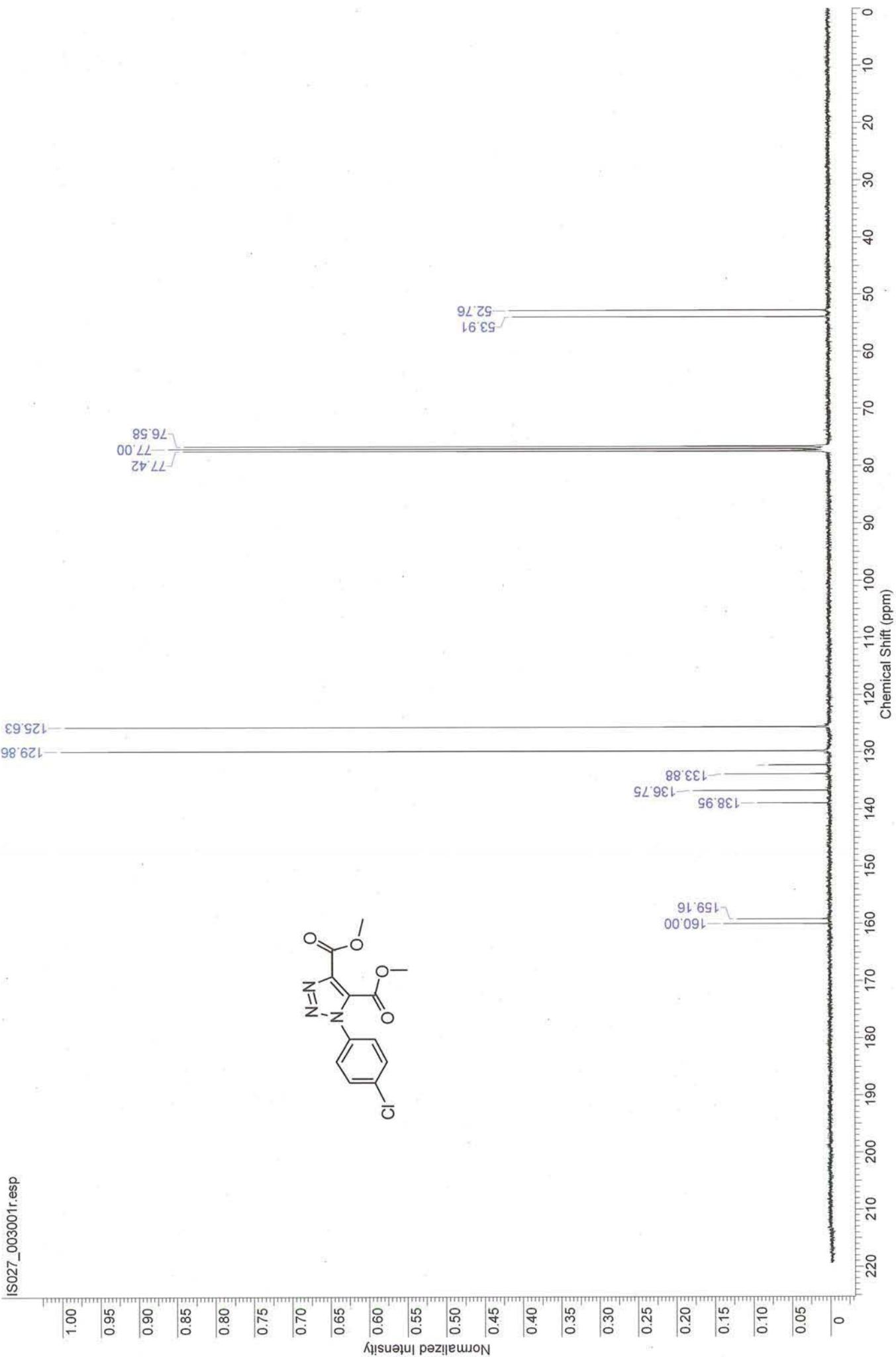
#	C	H	N	O	Na	mass	DBE	error
---	---	---	---	---	----	------	-----	-------

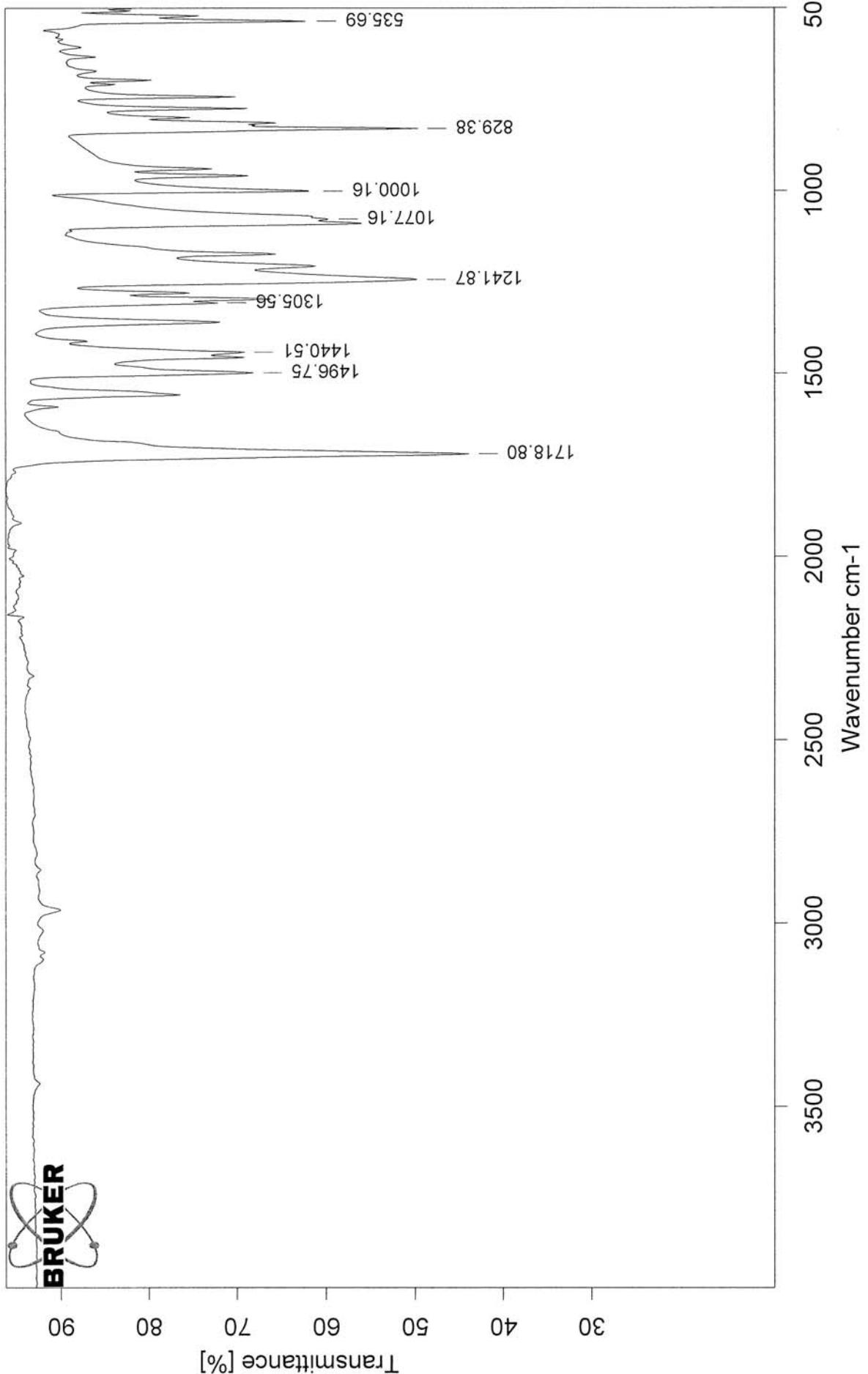
*** Mass Analysis for mass 262.0823110

1	12	12	3	4	0	262.0822323	8.5	7.869e-05
2	13	11	4	1	1	262.0825071	10.0	1.961e-04
3	10	10	6	3	0	262.0808896	9.0	1.421e-03
4	15	13	1	2	1	262.0838498	9.5	1.539e-03
5	10	13	3	4	1	262.0798270	5.5	2.484e-03
6	15	10	4	1	0	262.0849124	13.0	2.601e-03
7	9	14	2	7	0	262.0795522	4.0	2.759e-03
8	8	8	9	2	0	262.0795470	9.5	2.764e-03
9	8	11	6	3	1	262.0784843	6.0	3.827e-03
10	17	12	1	2	0	262.0862551	12.5	3.944e-03



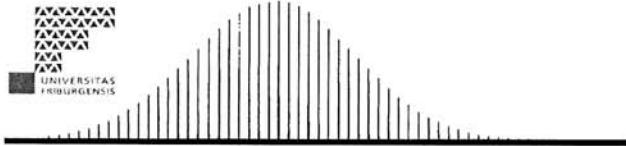






C:\PROGRAM FILES\OPUS\DATA\Isaki\is-027.2 is-027 is-027

26/01/2010



FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

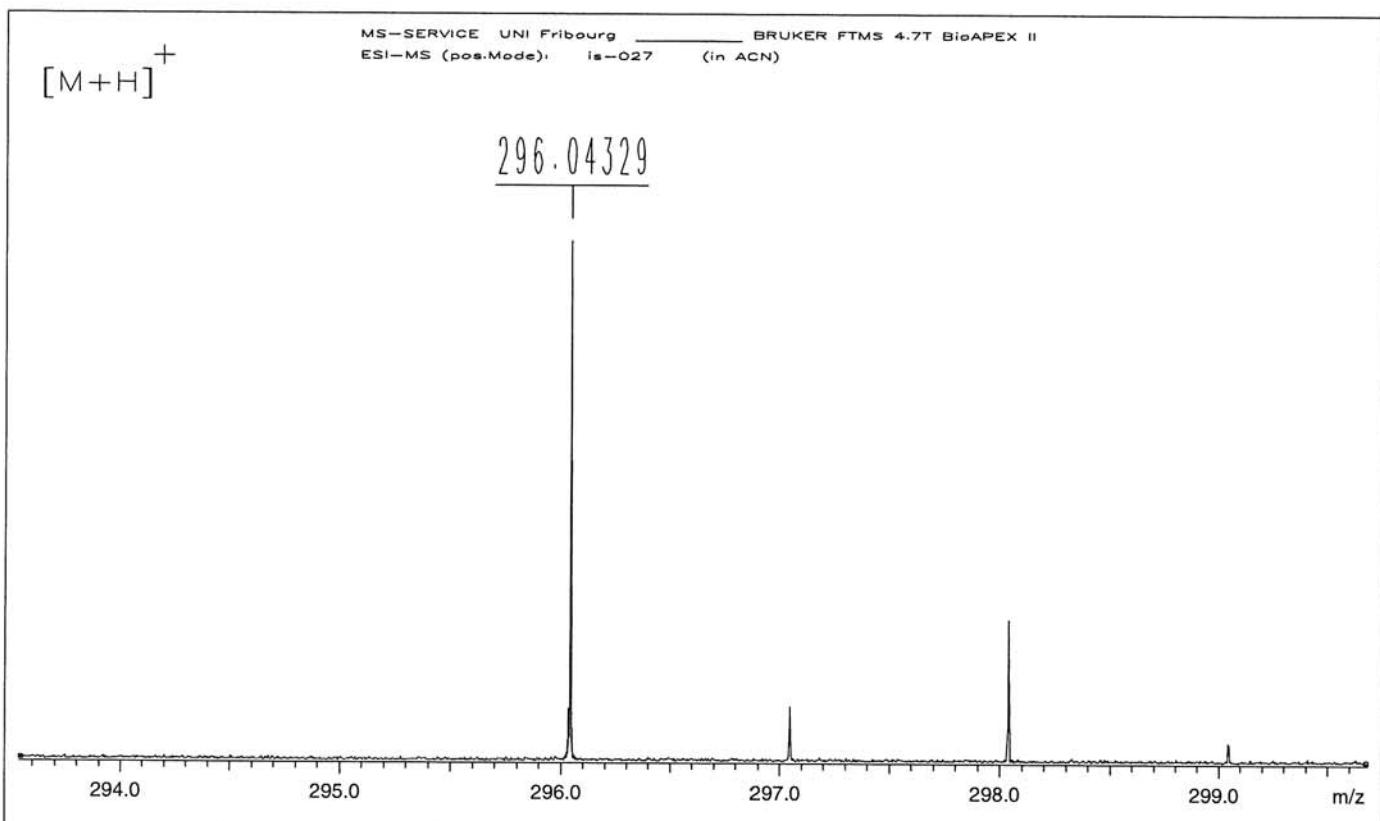
ESI-MS: is-027

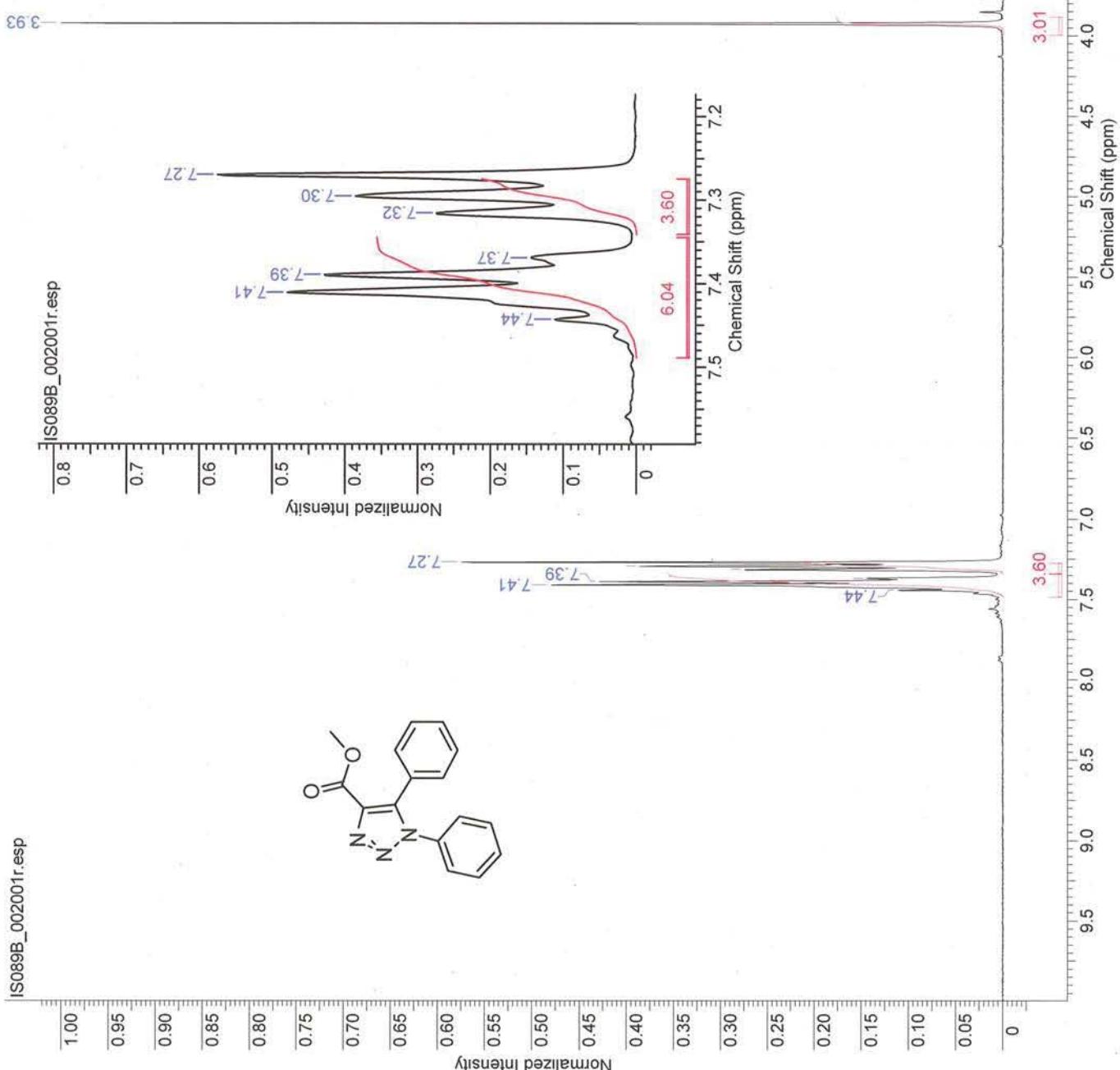
XMASS Mass Analysis for /Data/UNI_FR/ALIM4674_ESI/2/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

Ion mass = 296.0432860

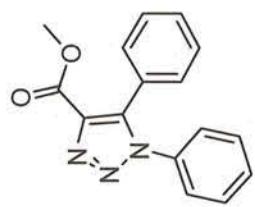
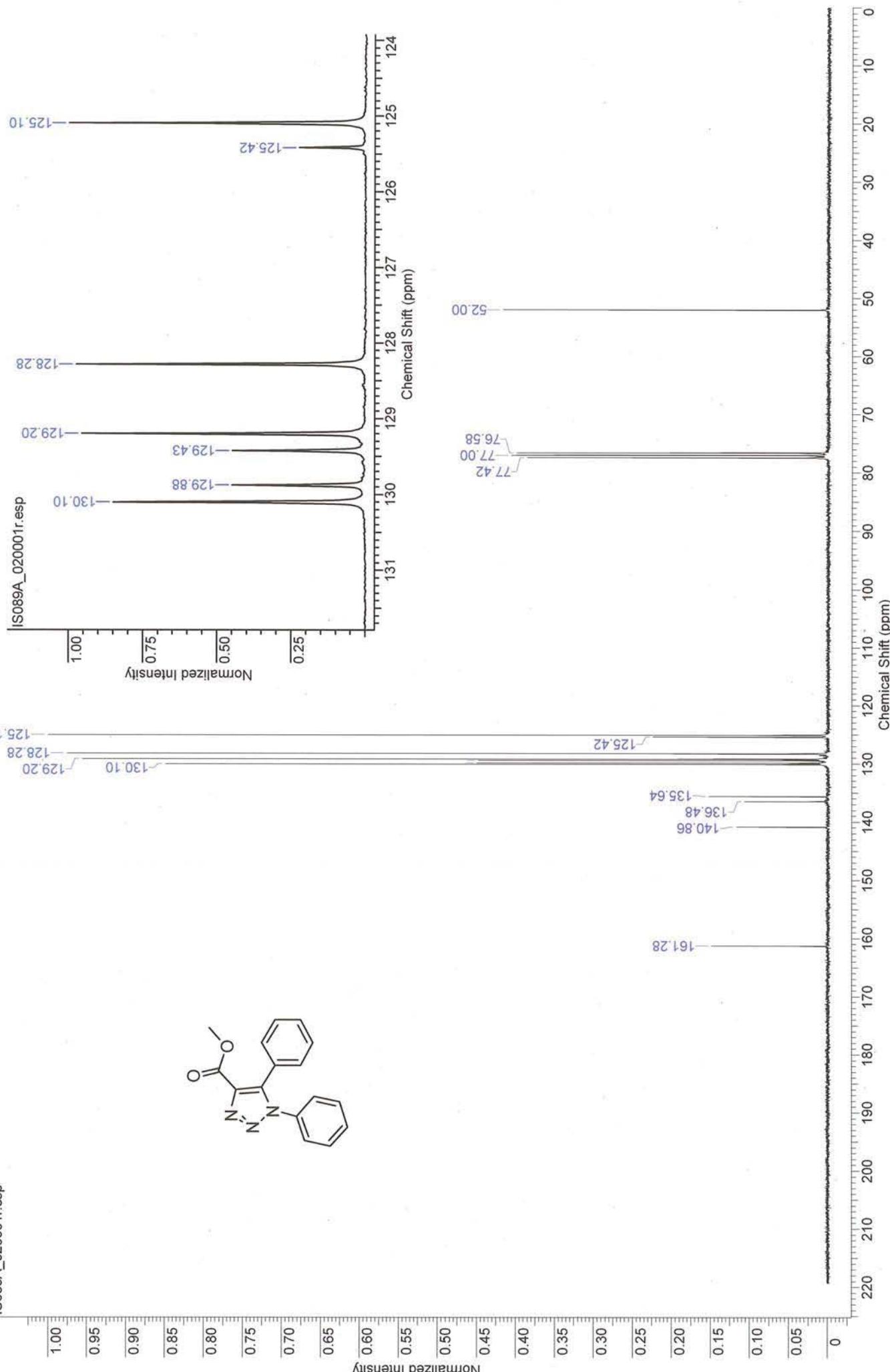
Charge = +1

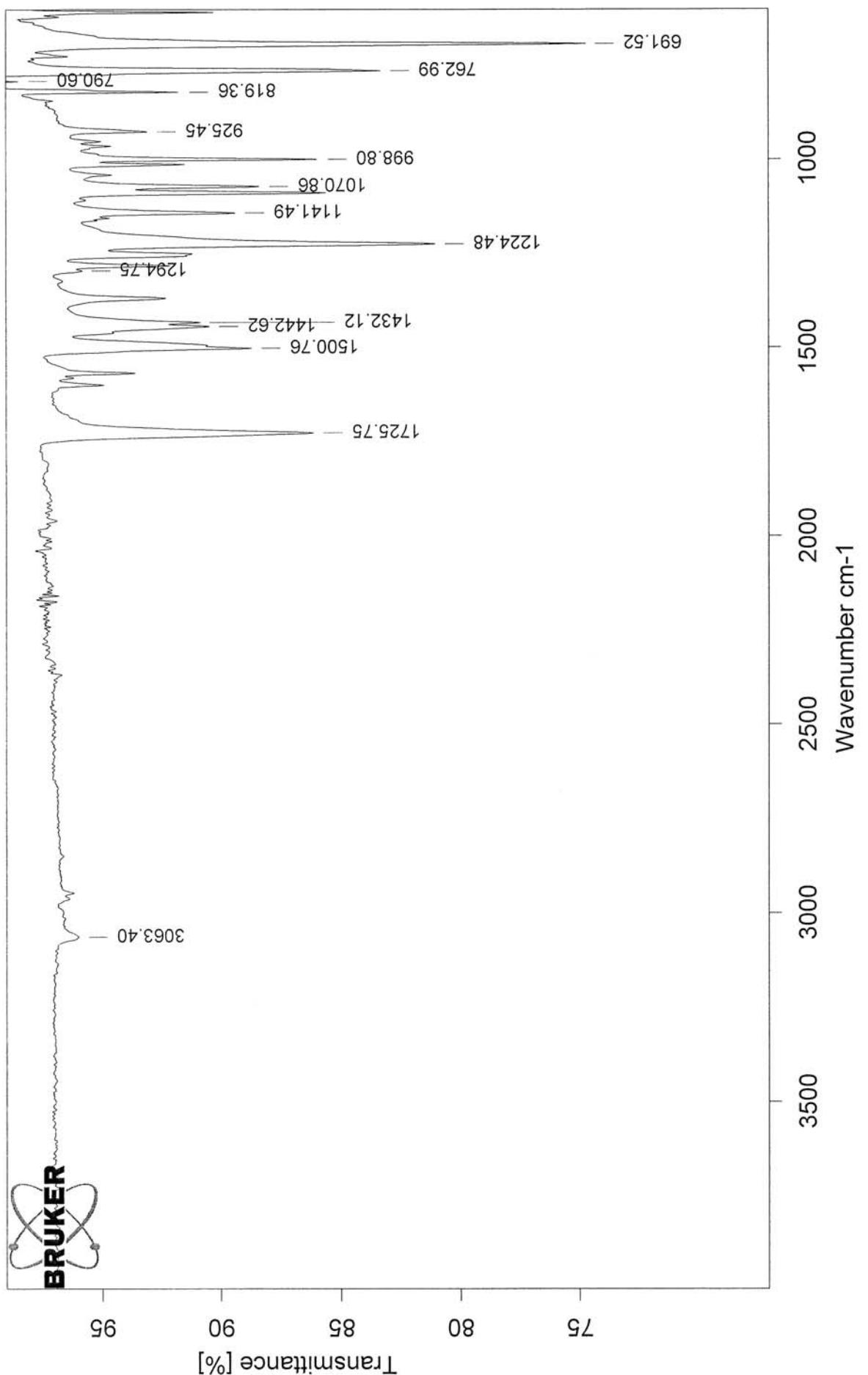
#	C	H	35Cl	N	O	mass	DBE	error
*** Mass Analysis for mass 296.0432860								
1	12	11	1	3	4	296.0432600	8.5	2.601e-05
2	9	14	2	4	3	296.0437472	4.0	4.612e-04
3	11	16	2	1	4	296.0450898	3.5	1.804e-03
4	6	16	2	3	6	296.0410671	-0.5	2.219e-03
5	15	9	1	4	1	296.0459401	13.0	2.654e-03
6	9	13	1	2	7	296.0405799	4.0	2.706e-03
7	17	11	1	1	2	296.0472827	12.5	3.997e-03
8	7	11	1	5	6	296.0392372	4.5	4.049e-03
9	14	14	2	2	1	296.0477699	8.0	4.484e-03
10	6	15	1	1	10	296.0378998	-0.5	5.386e-03





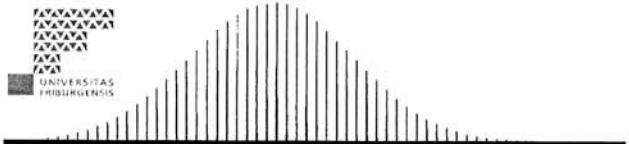
IS089A_020001r.esp





01/07/2010

U:\IR_spectra\IS089.0 IS089 white solid



FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

ESI-MS: IS089

XMASS Mass Analysis for /Data/UNI_FR/ALIM4950_ESI/5/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

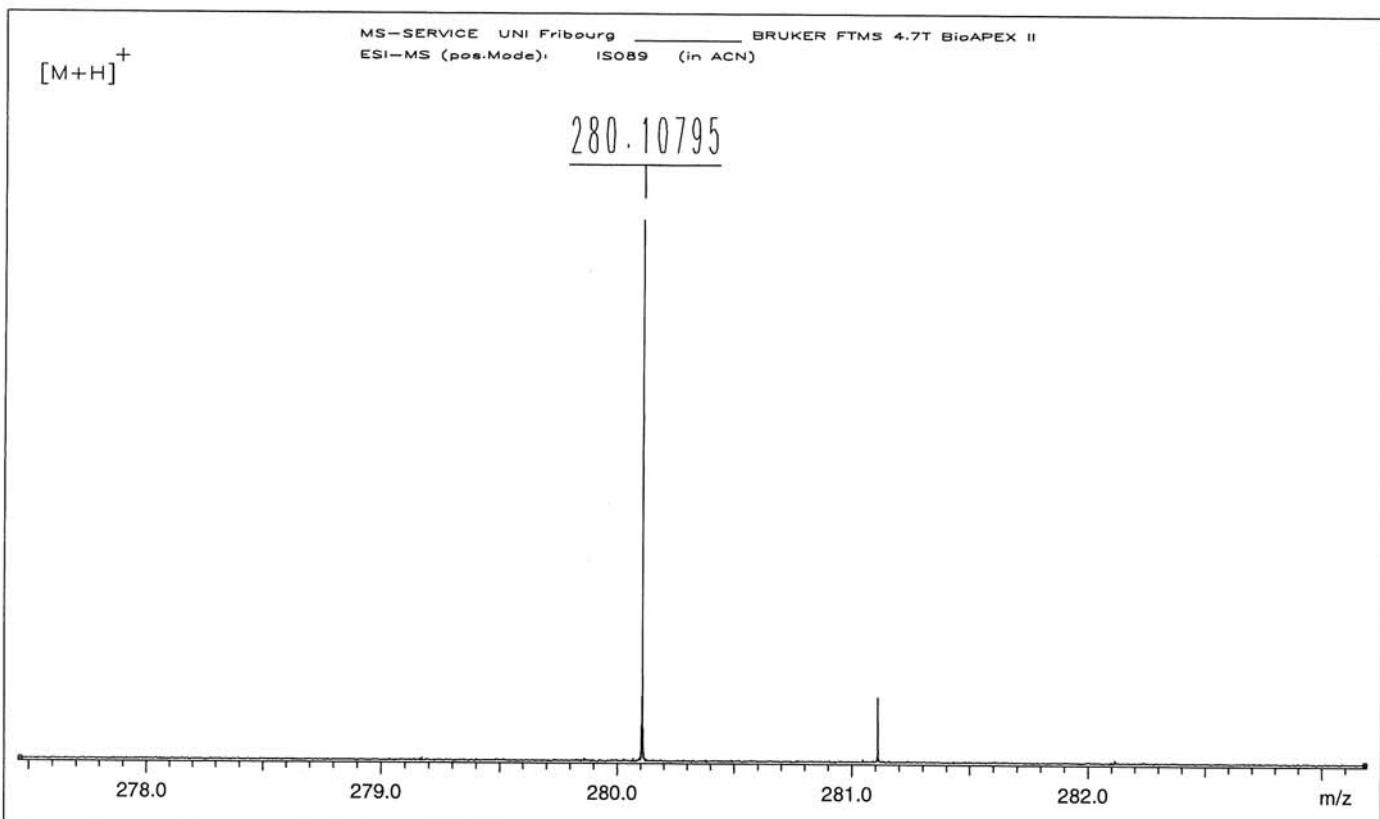
Ion mass = 280.1079540

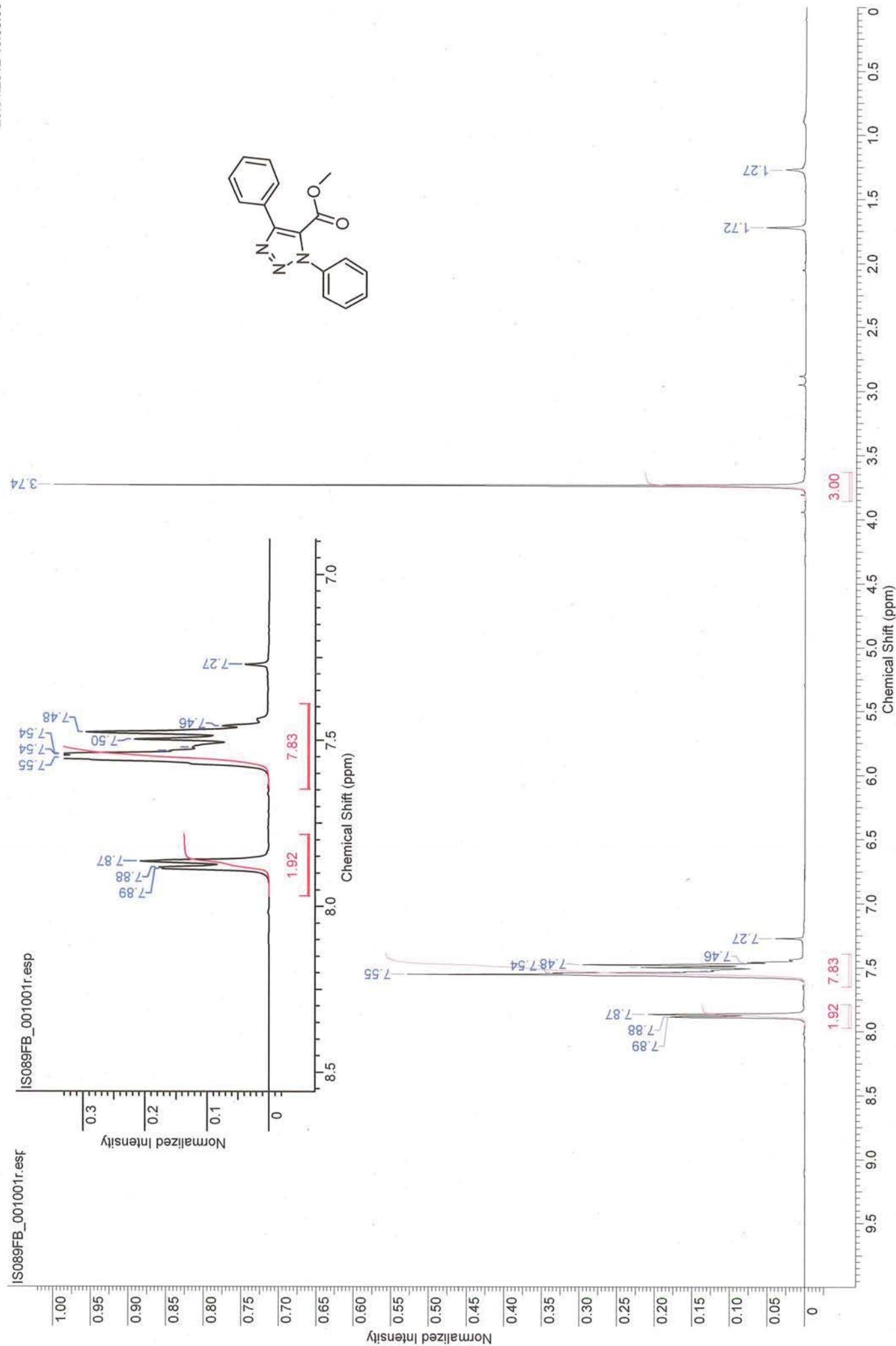
Charge = +1

#	C	H	N	O	mass	DBE	error
---	---	---	---	---	------	-----	-------

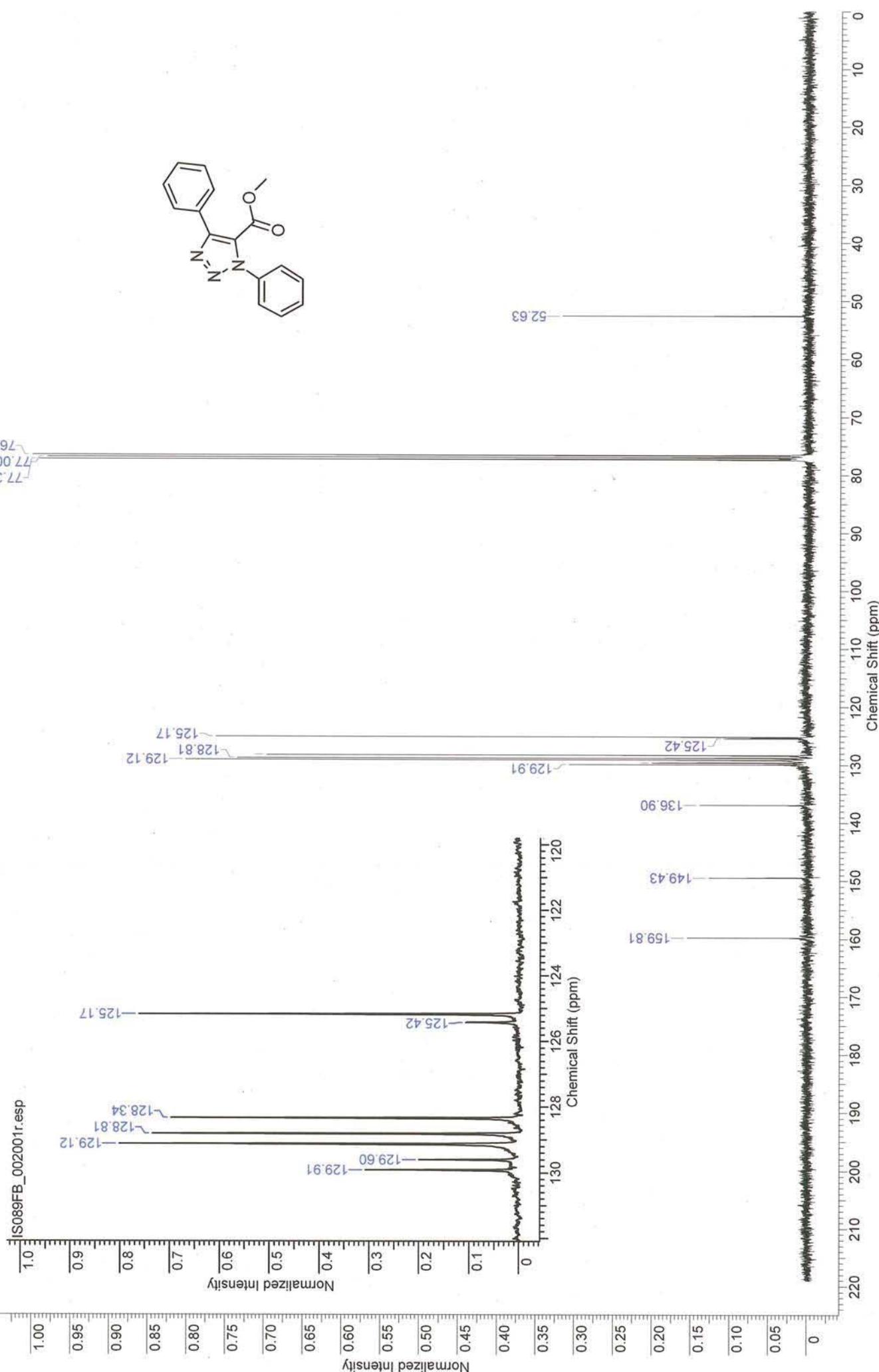
*** Mass Analysis for mass 280.1079540

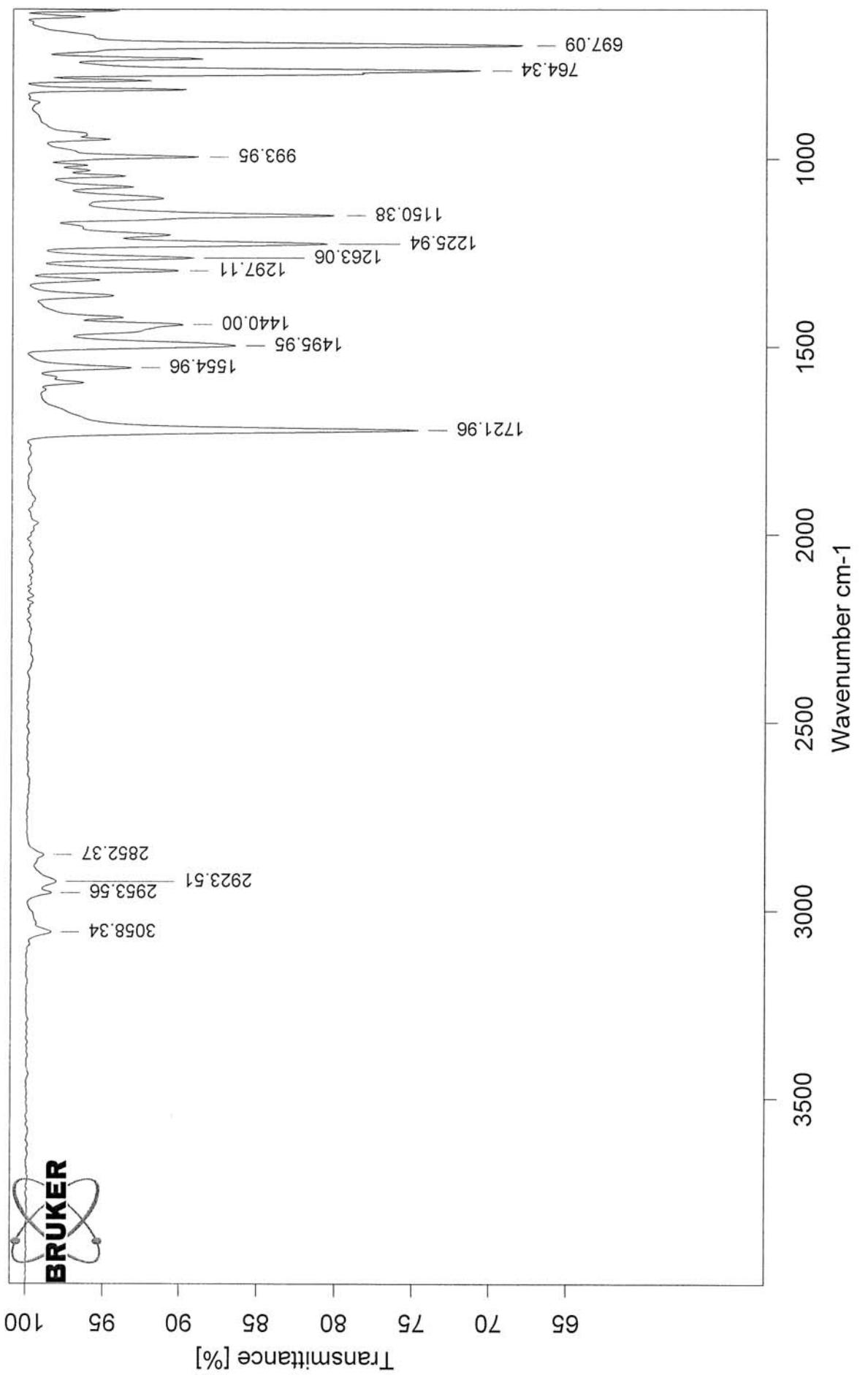
1	16	14	3	2	280.1080531	11.5	9.913e-05
2	14	12	6	1	280.1067105	12.0	1.244e-03
3	13	16	2	5	280.1053731	7.0	2.581e-03
4	5	14	9	5	280.1112410	3.5	3.287e-03
5	6	20	2	10	280.1112463	-2.0	3.292e-03
6	11	14	5	4	280.1040304	7.5	3.924e-03
7	7	16	6	6	280.1125837	3.0	4.630e-03
8	10	18	1	8	280.1026930	2.5	5.261e-03
9	9	12	8	3	280.1026877	8.0	5.266e-03
10	8	12	10	2	280.1139211	8.0	5.967e-03





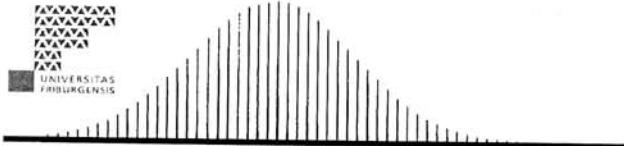
IS089FB_002001r.esp





09/07/2010

U:\IR_spectra\IS089.2 IS089 white yellow solid



FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

ESI-MS: IS089B

XMASS Mass Analysis for /Data/UNI_FR/ALIM5004_ESI/2/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

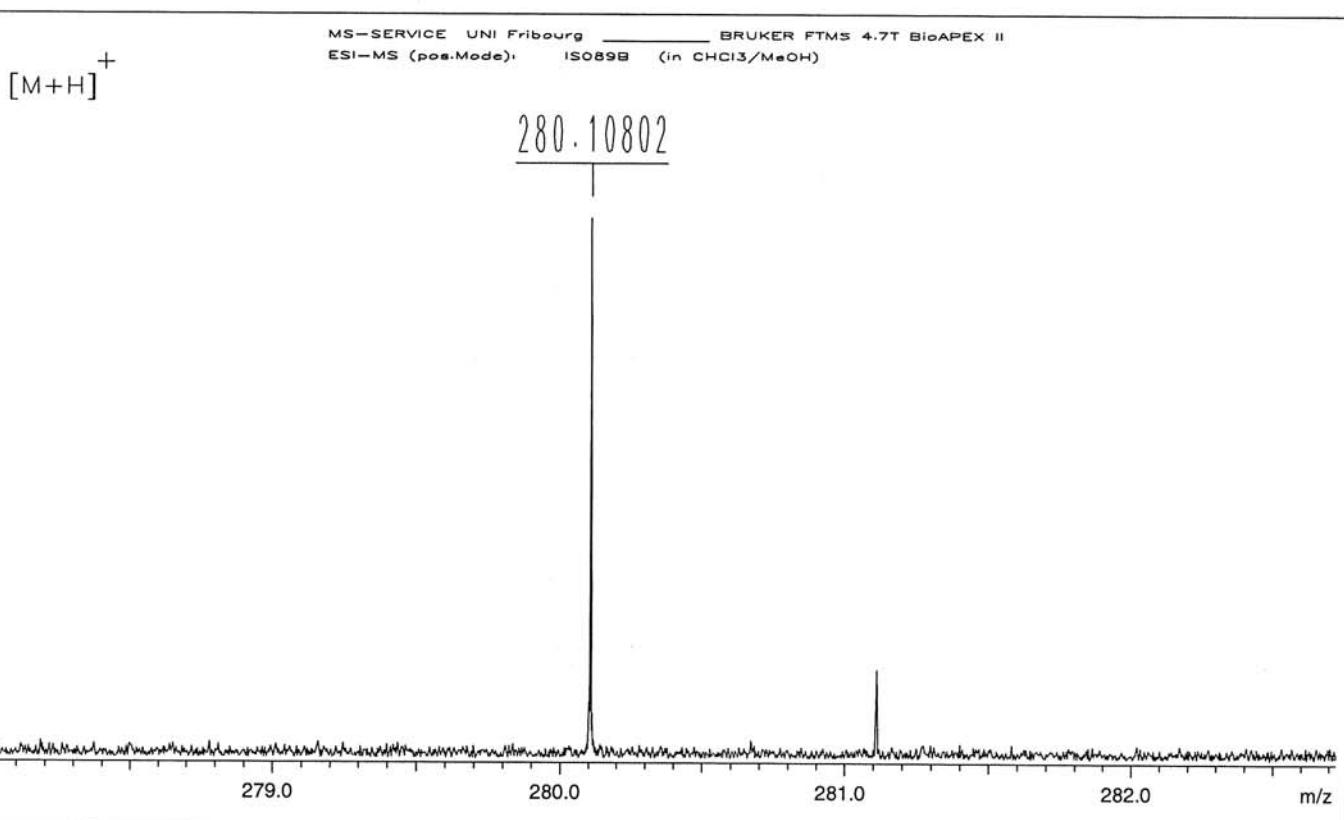
Ion mass = 280.1080220

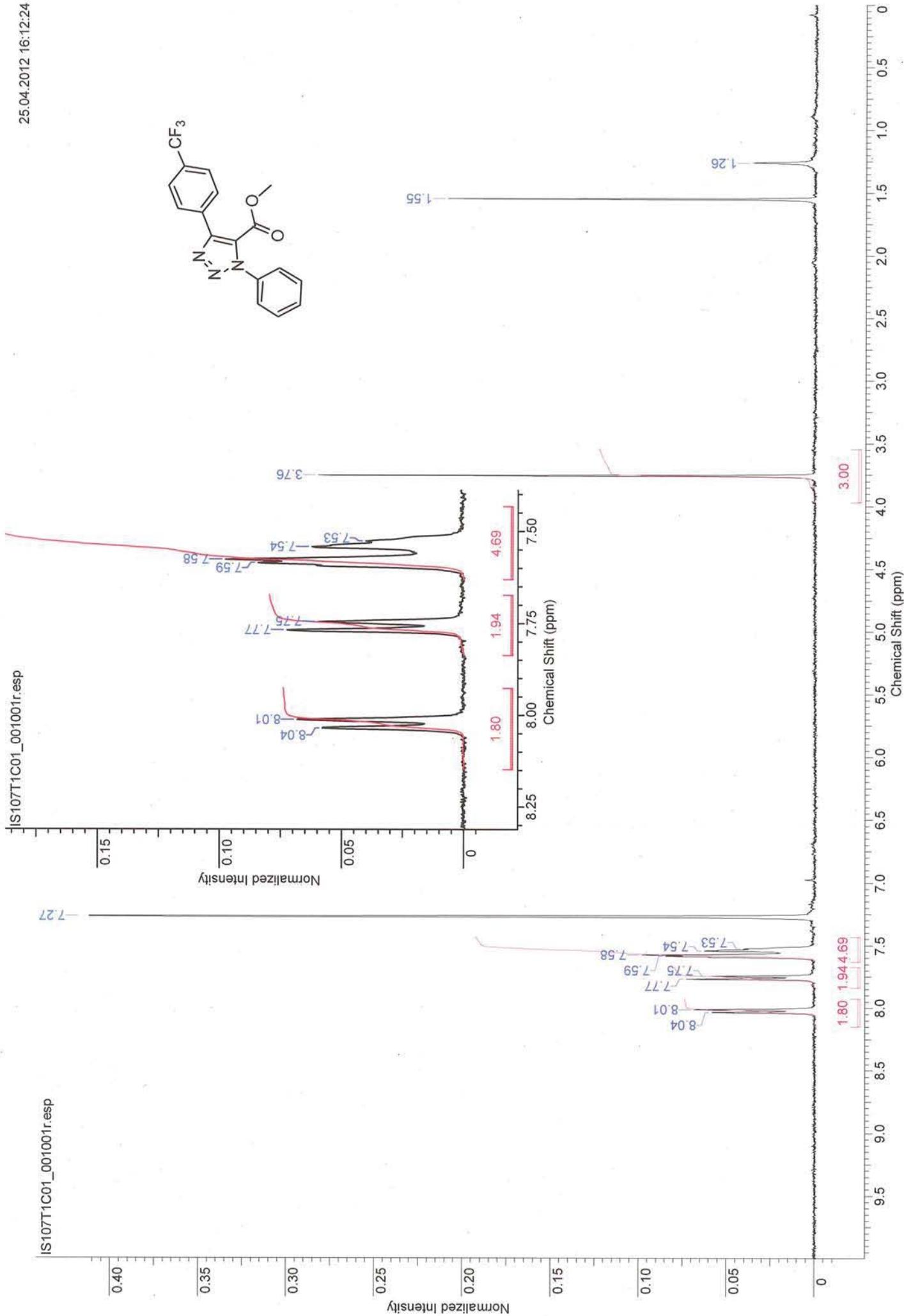
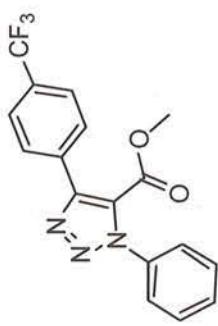
Charge = +1

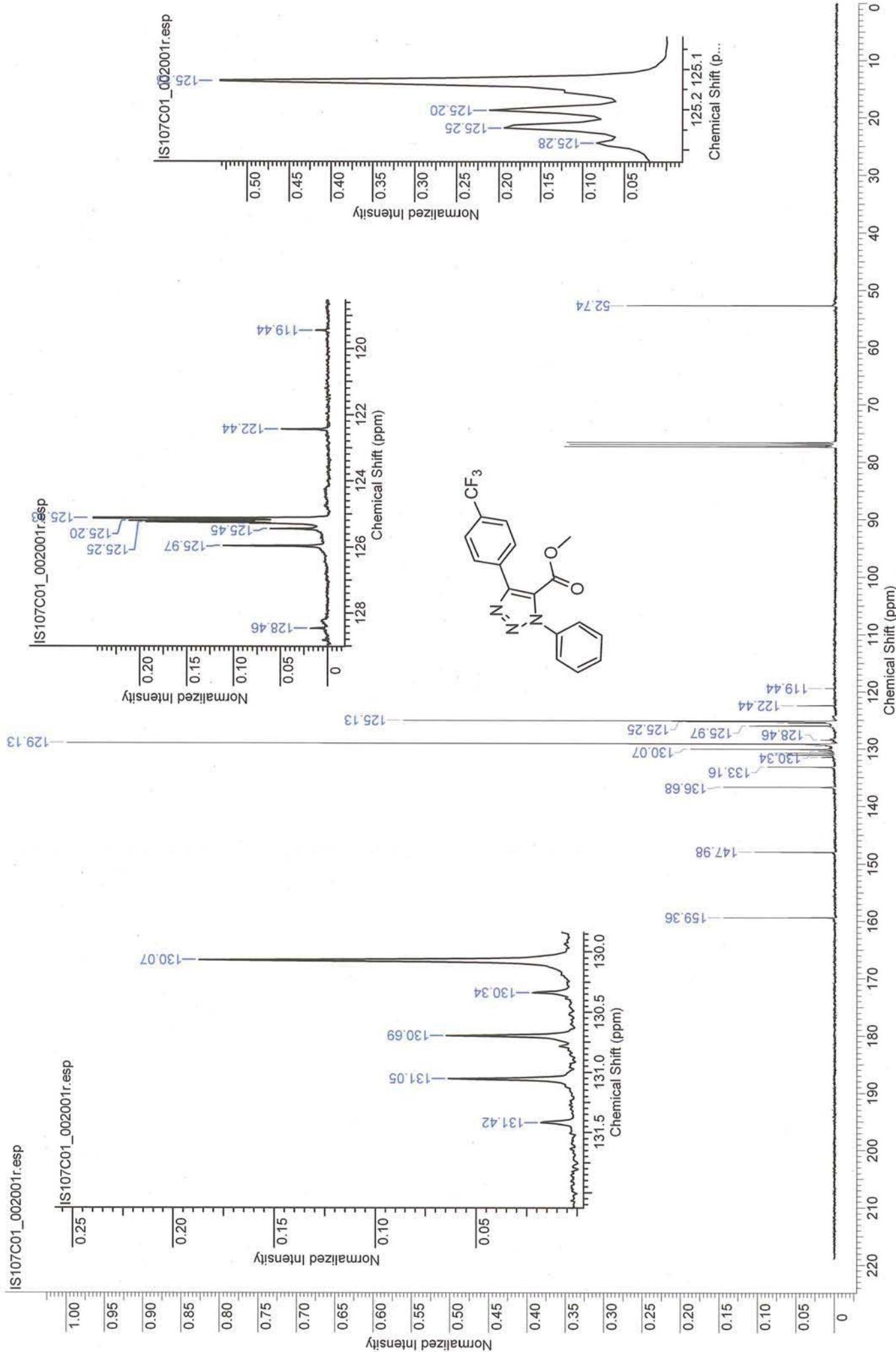
#	C	H	N	O	mass	DBE	error
---	---	---	---	---	------	-----	-------

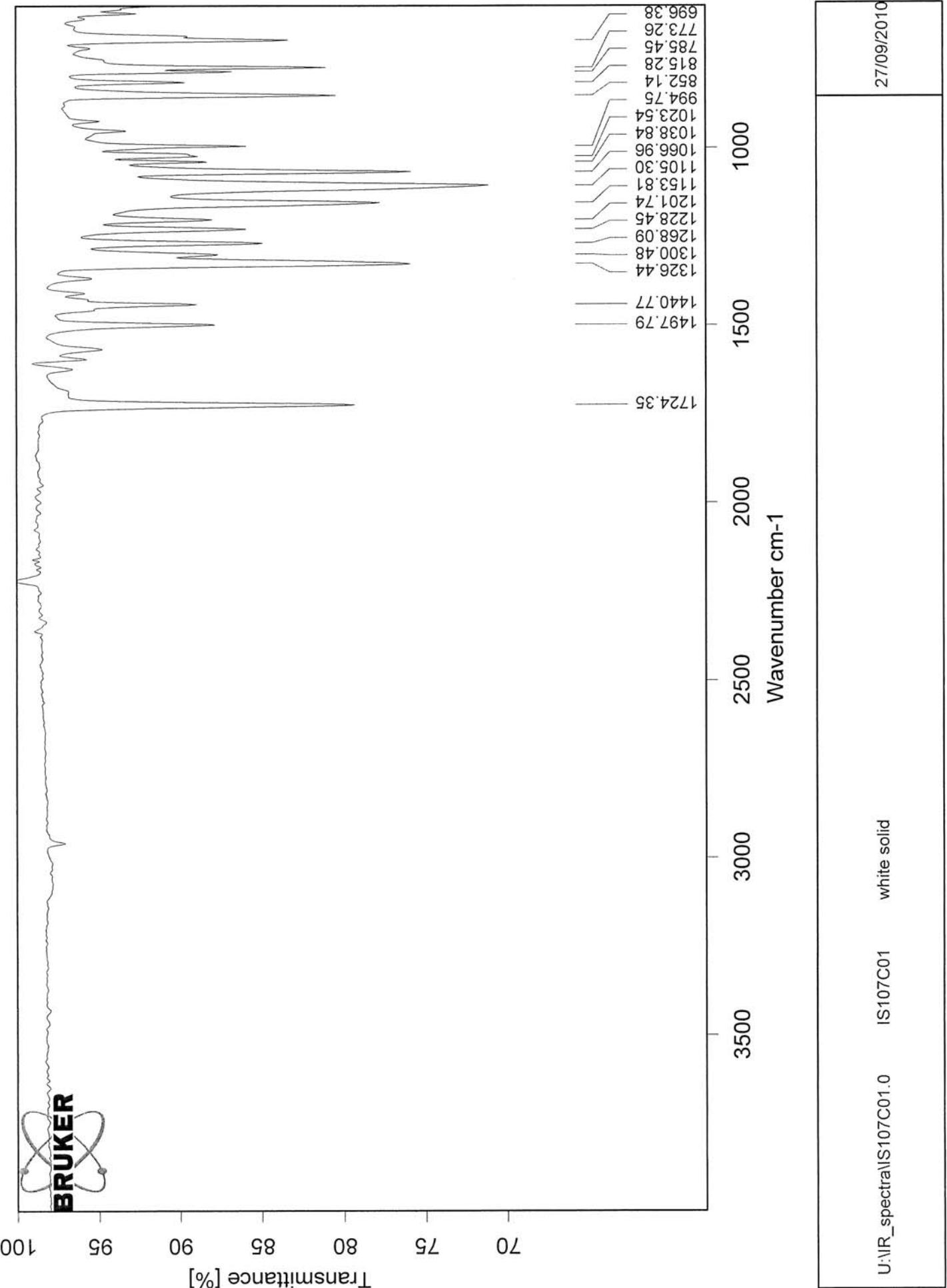
*** Mass Analysis for mass 280.1080220

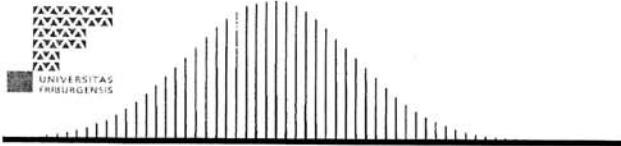
1	16	14	3	2	280.1080531	11.5	3.113e-05
2	14	12	6	1	280.1067105	12.0	1.312e-03
3	13	16	2	5	280.1053731	7.0	2.649e-03
4	5	14	9	5	280.1112410	3.5	3.219e-03
5	6	20	2	10	280.1112463	-2.0	3.224e-03
6	11	14	5	4	280.1040304	7.5	3.992e-03
7	7	16	6	6	280.1125837	3.0	4.562e-03
8	10	18	1	8	280.1026930	2.5	5.329e-03
9	9	12	8	3	280.1026877	8.0	5.334e-03
10	8	12	10	2	280.1139211	8.0	5.899e-03











FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

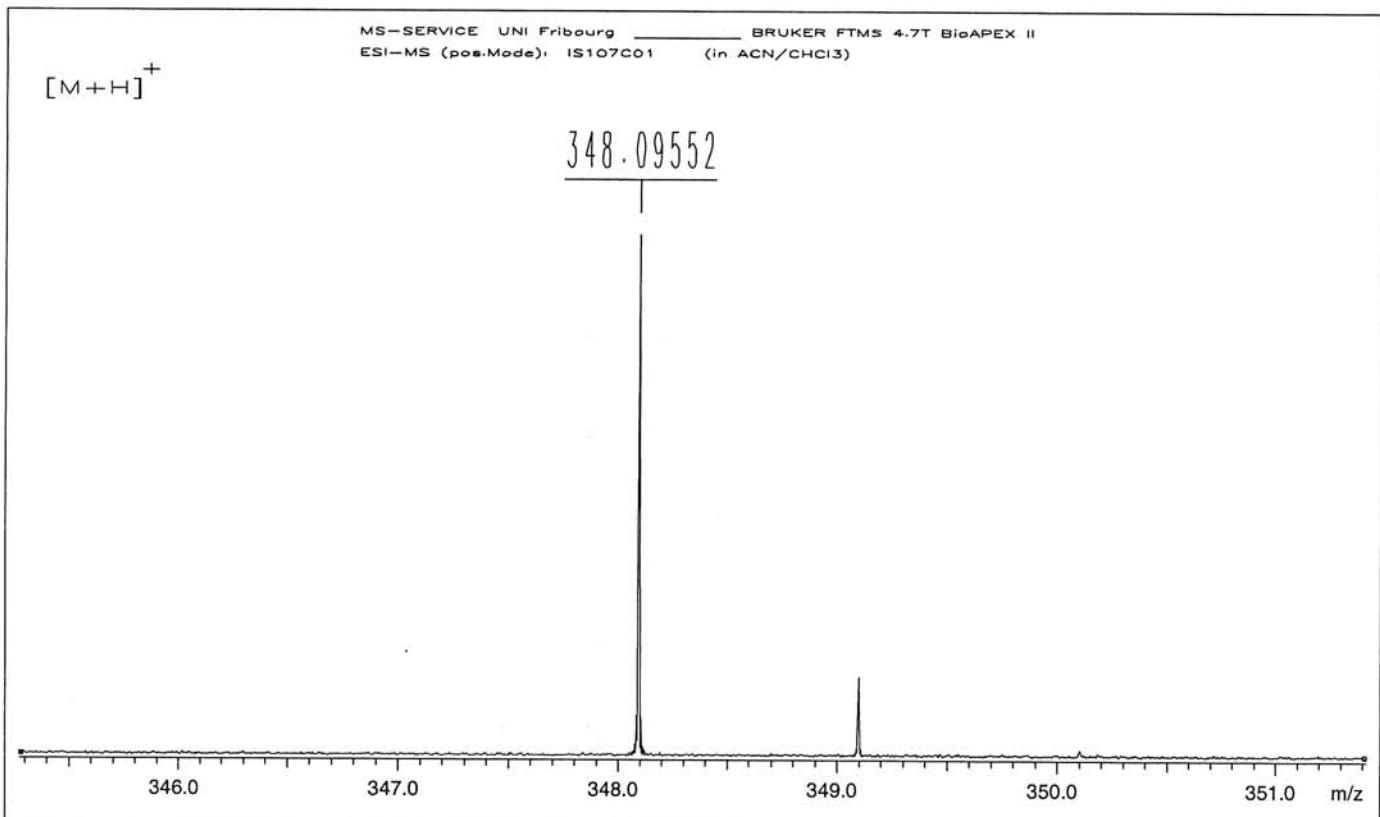
ESI-MS: IS107C01

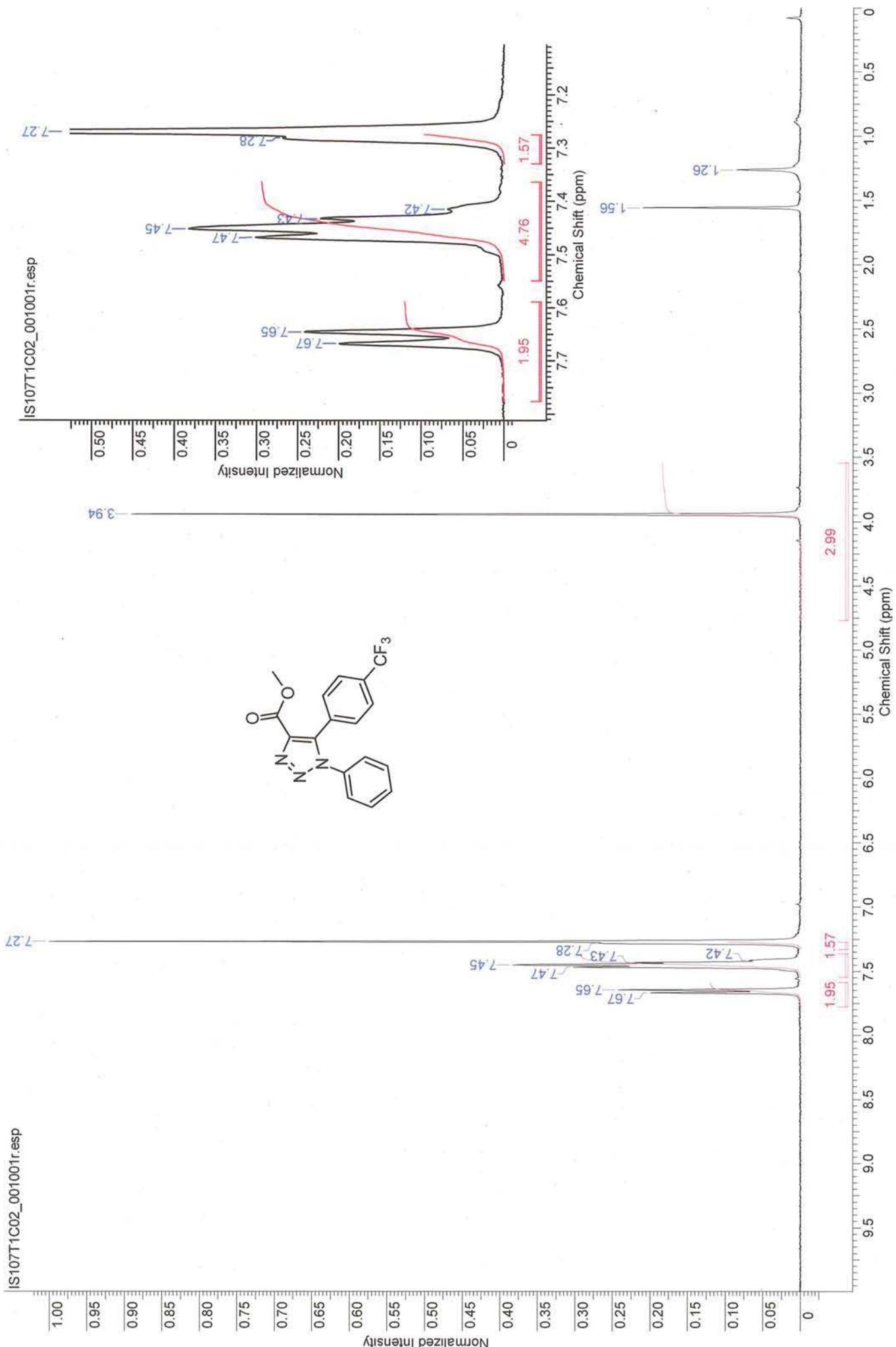
XMASS Mass Analysis for /Data/UNI_FR/ALIM5062_ESI/2/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

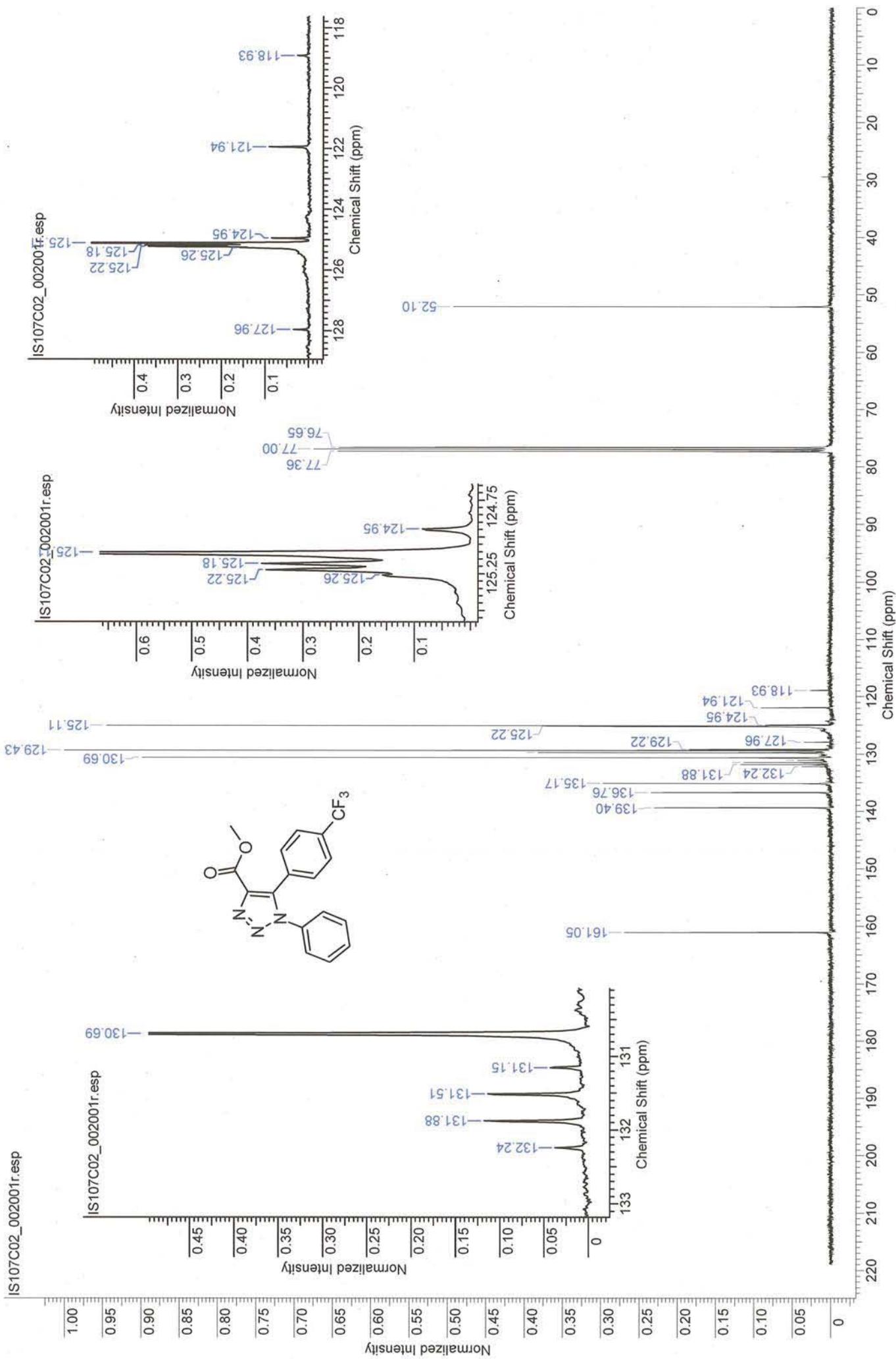
Ion mass = 348.0955240

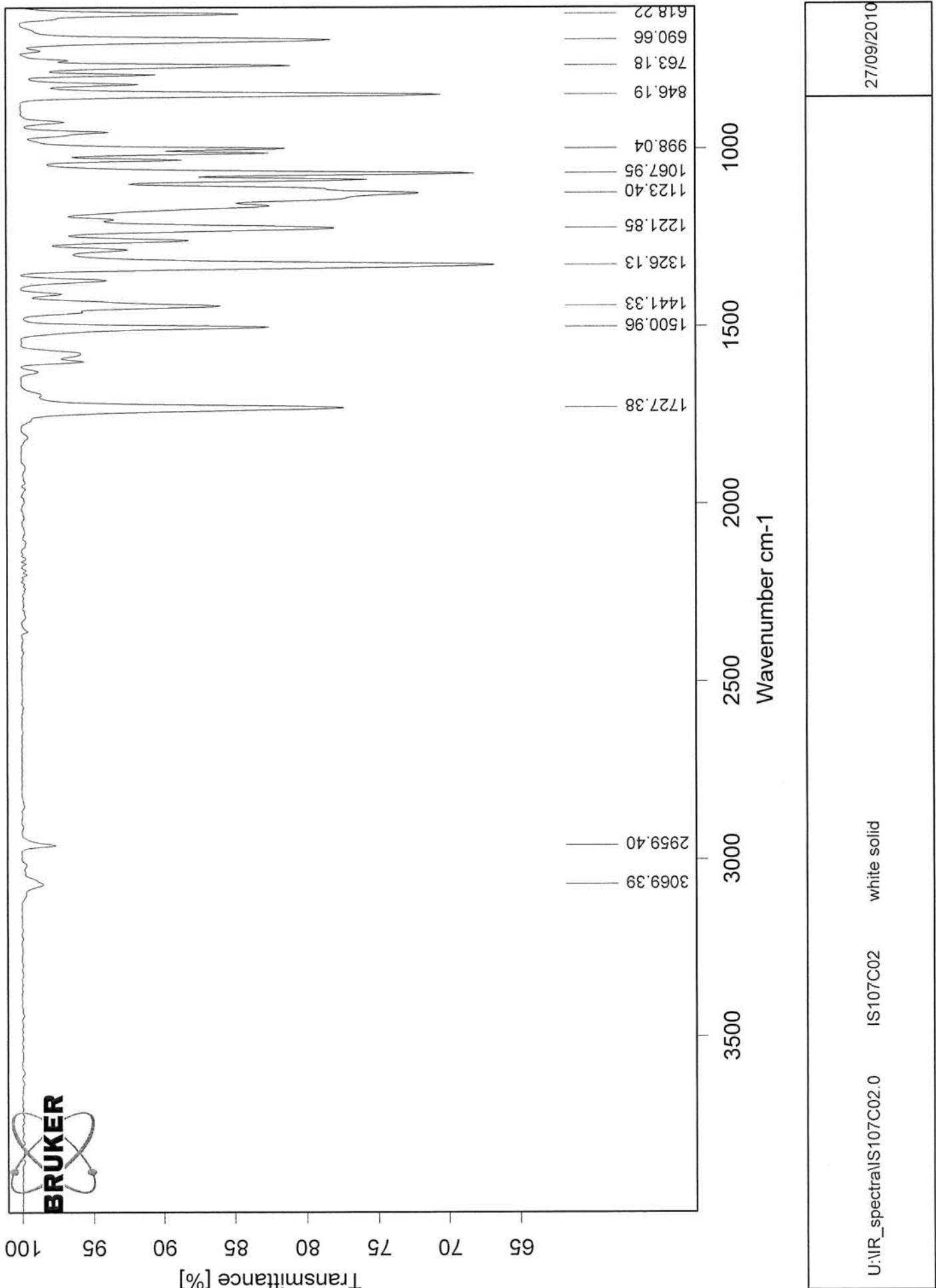
Charge = +1

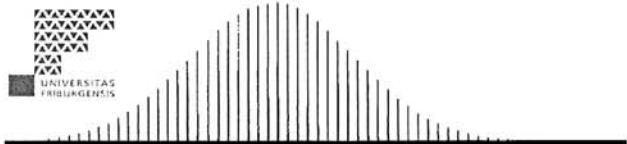
#	C	H	F	N	O	mass	DBE	error
*** Mass Analysis for mass 348.0955240								
1	17	13	3	3	2	348.0954377	11.5	8.630e-05
2	14	14	4	3	3	348.0965806	7.5	1.057e-03
3	20	12	2	3	1	348.0942948	15.5	1.229e-03
4	11	15	5	3	4	348.0977234	3.5	2.199e-03
5	14	15	3	2	5	348.0927576	7.0	2.766e-03
6	16	15	1	3	5	348.0990252	10.5	3.501e-03
7	17	14	2	2	4	348.0916148	11.0	3.909e-03
8	12	13	3	5	4	348.0914150	7.5	4.109e-03
9	14	13	5	4	1	348.1004035	8.0	4.879e-03
10	20	13	1	2	3	348.0904719	15.0	5.052e-03











FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

ESI-MS: IS107C02

XMASS Mass Analysis for /Data/UNI_FR/ALIM5063_ESI/2/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

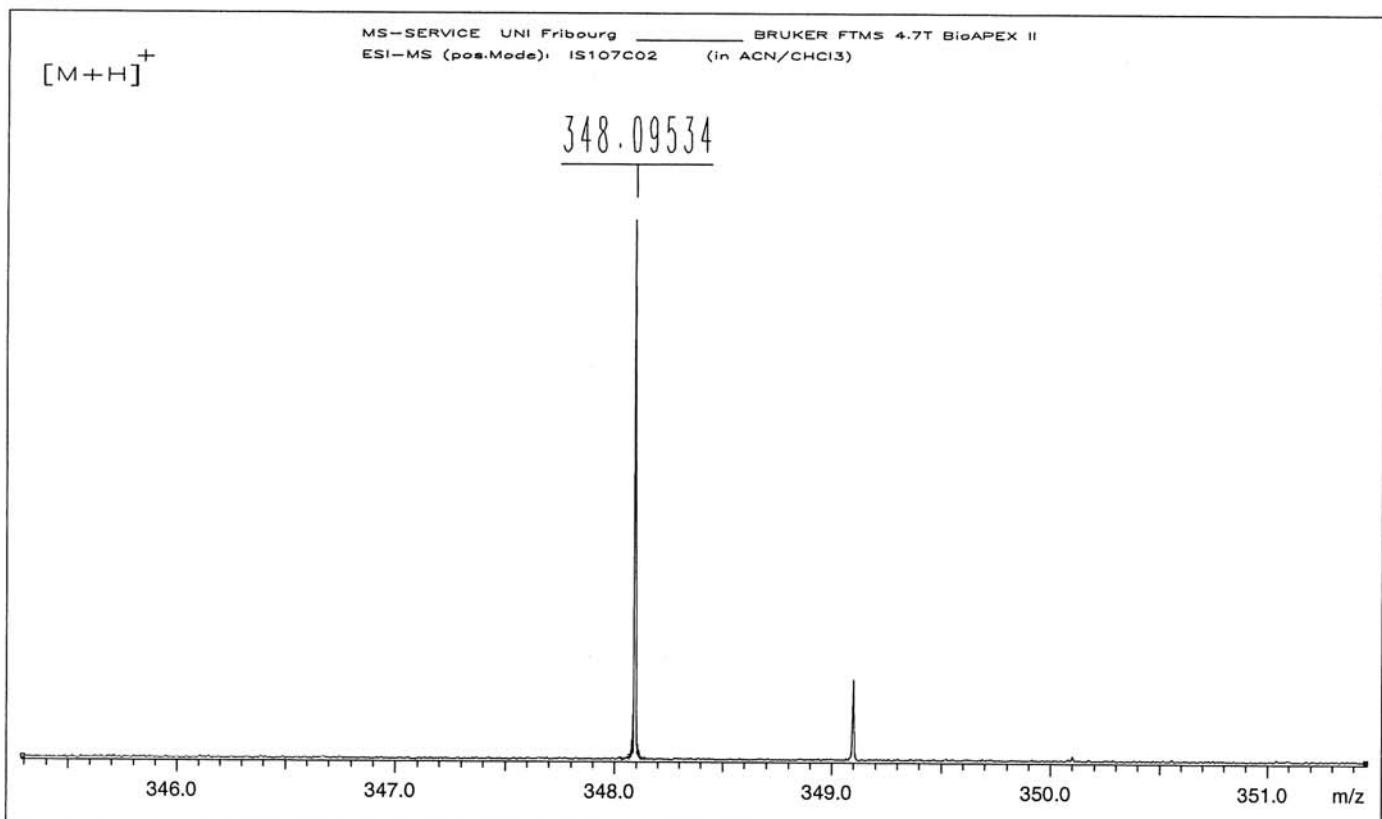
Ion mass = 348.0953440

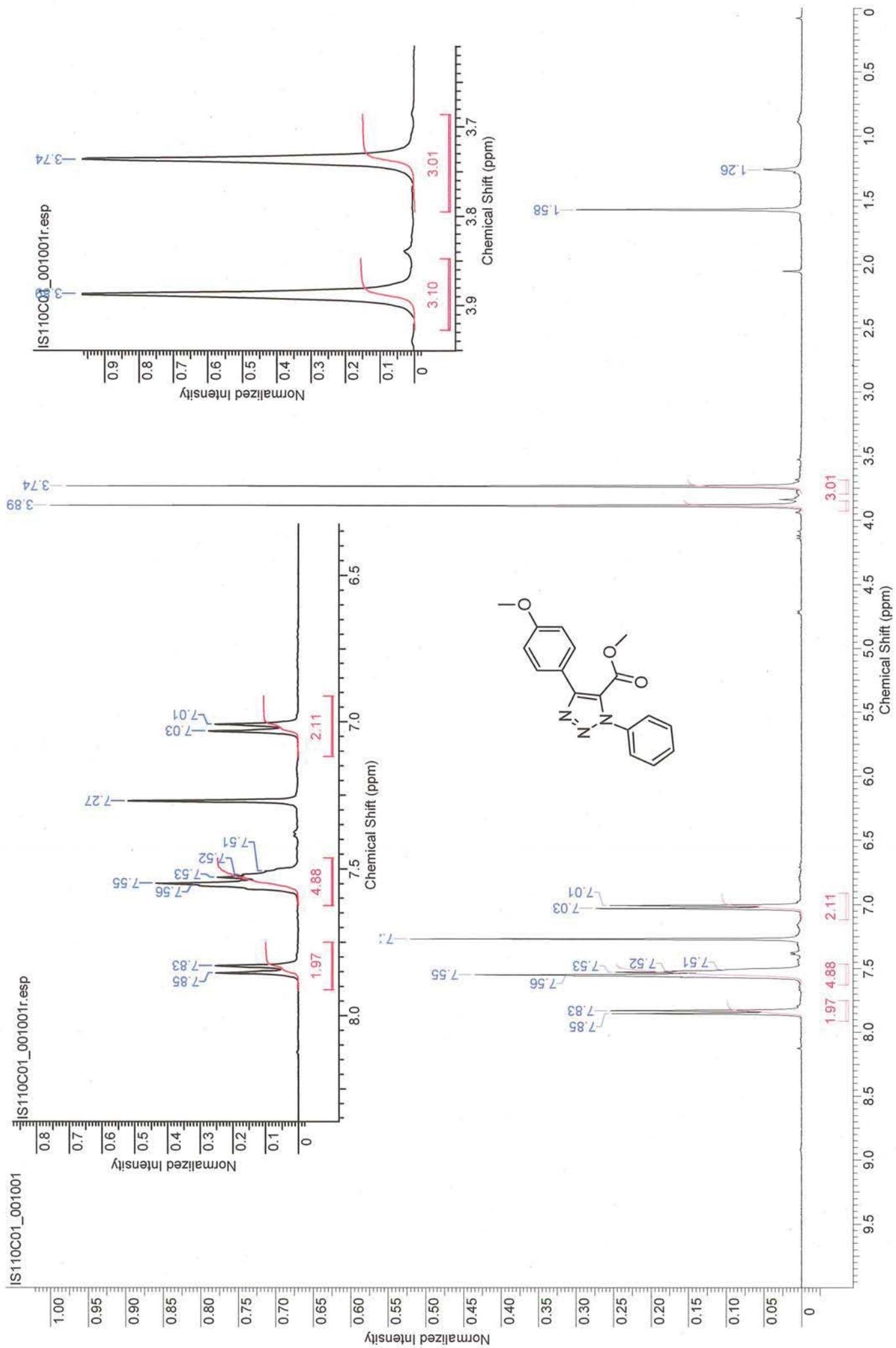
Charge = +1

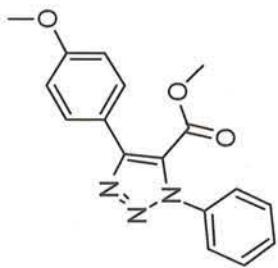
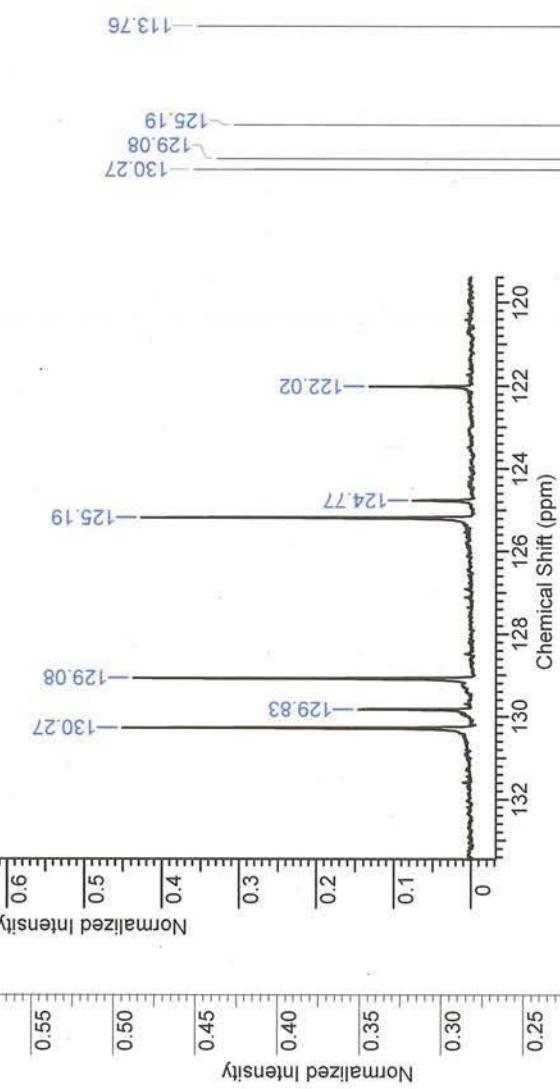
#	C	H	F	N	O	mass	DBE	error
---	---	---	---	---	---	------	-----	-------

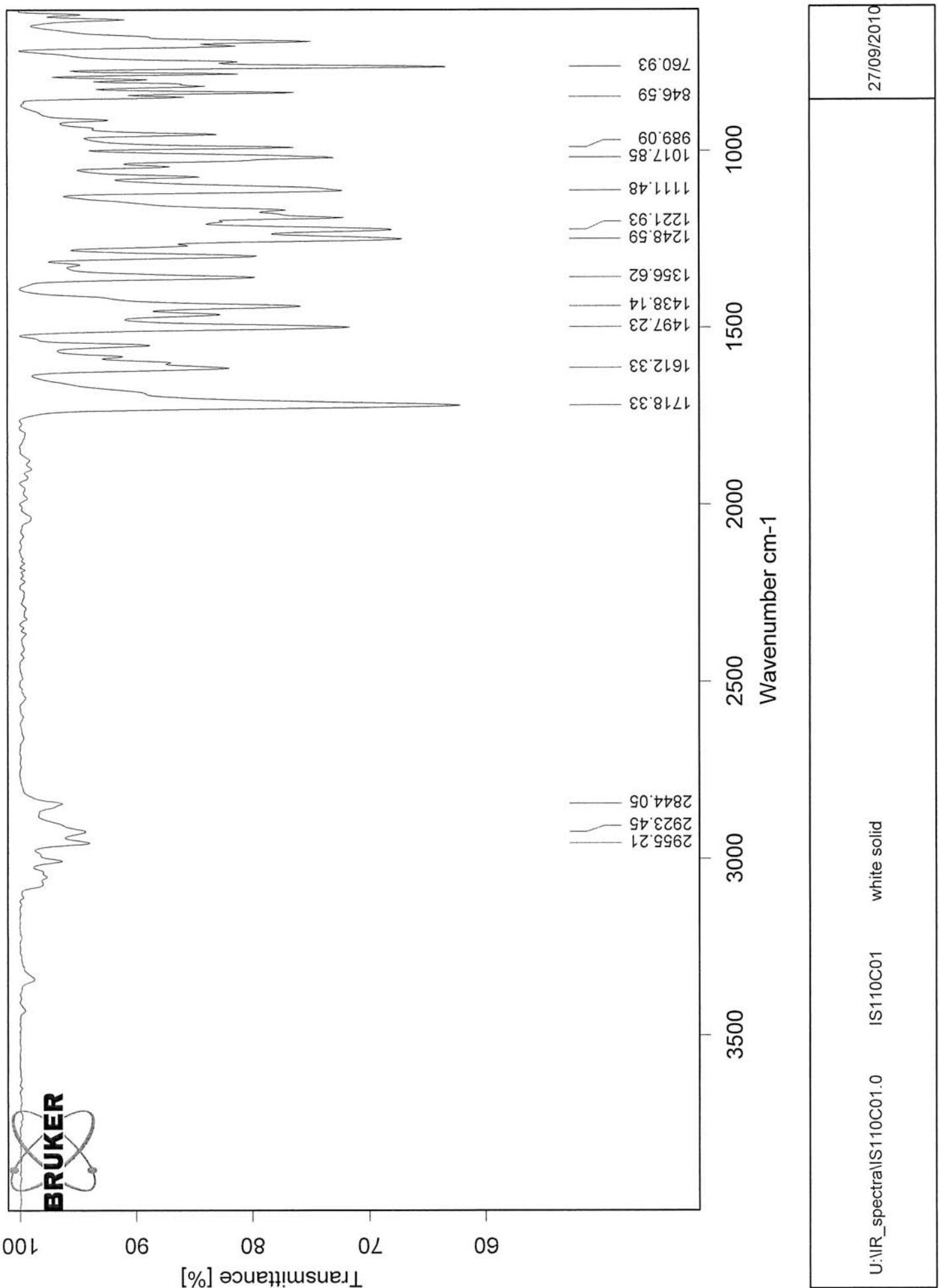
*** Mass Analysis for mass 348.0953440

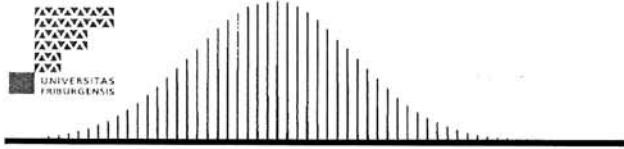
1	17	13	3	3	2	348.0954377	11.5	9.370e-05
2	20	12	2	3	1	348.0942948	15.5	1.049e-03
3	14	14	4	3	3	348.0965806	7.5	1.237e-03
4	11	15	5	3	4	348.0977234	3.5	2.379e-03
5	14	15	3	2	5	348.0927576	7.0	2.586e-03
6	16	15	1	3	5	348.0990252	10.5	3.681e-03
7	17	14	2	2	4	348.0916148	11.0	3.729e-03
8	12	13	3	5	4	348.0914150	7.5	3.929e-03
9	20	13	1	2	3	348.0904719	15.0	4.872e-03
10	14	13	5	4	1	348.1004035	8.0	5.059e-03





77.35
77.00
76.65IS110C01_002001r.esp
IS110C01_002001r.esp





FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

ESI-MS: IS0C01

XMASS Mass Analysis for /Data/UNI_FR/ALIM50778_ESI/5/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

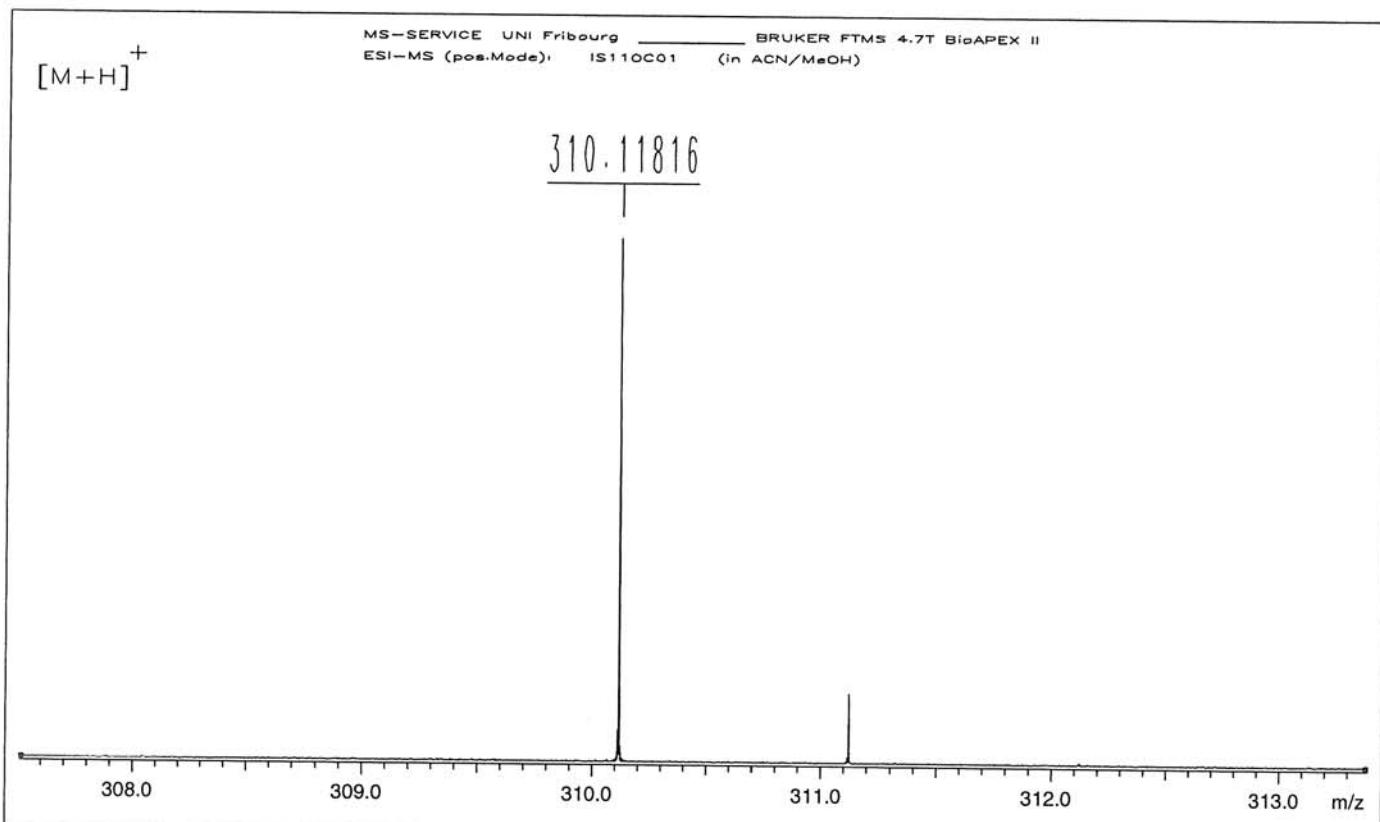
Ion mass = 310.1181560

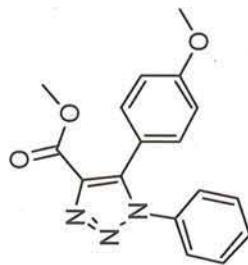
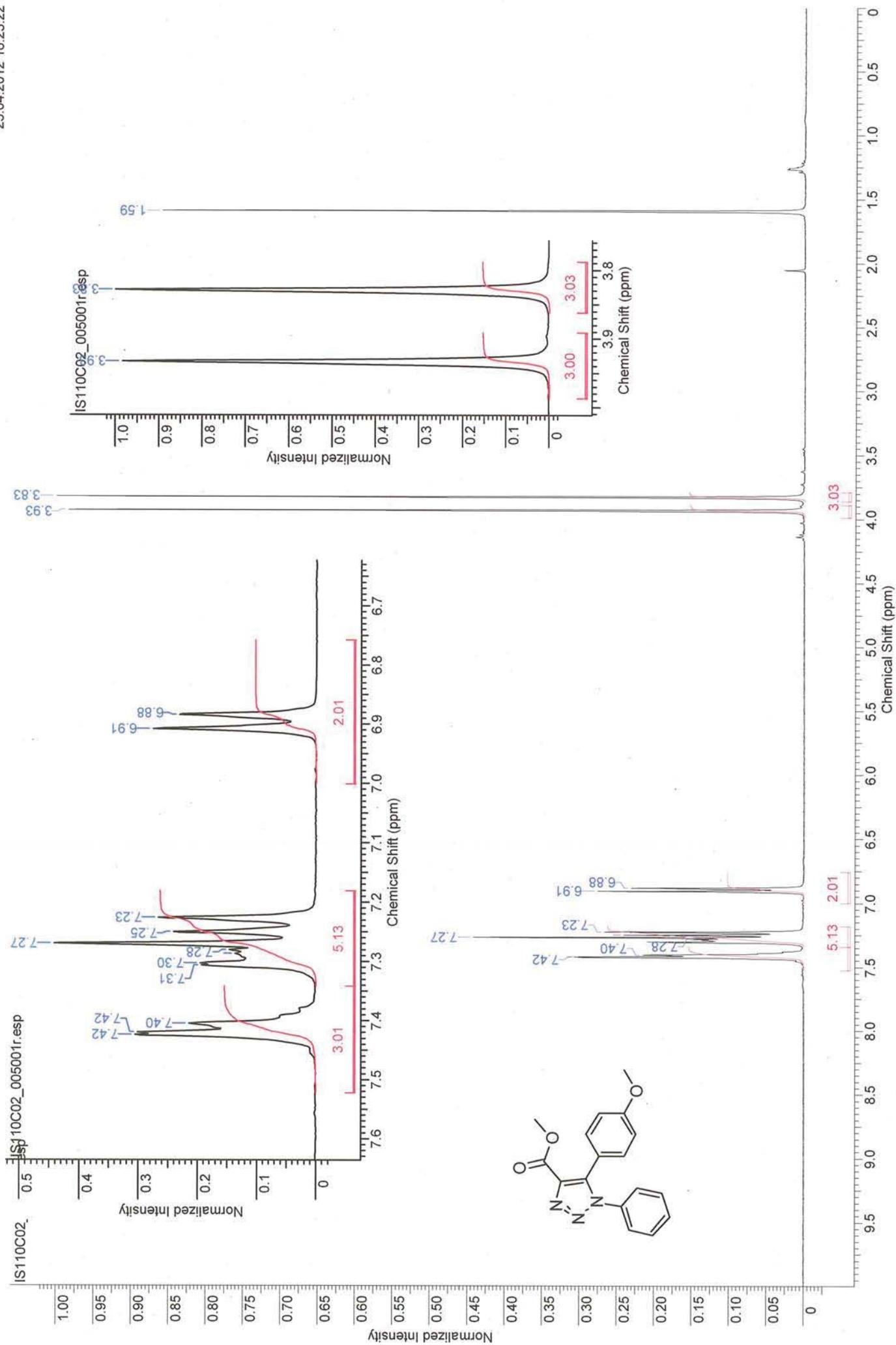
Charge = +1

#	C	H	N	O	mass	DBE	error
---	---	---	---	---	------	-----	-------

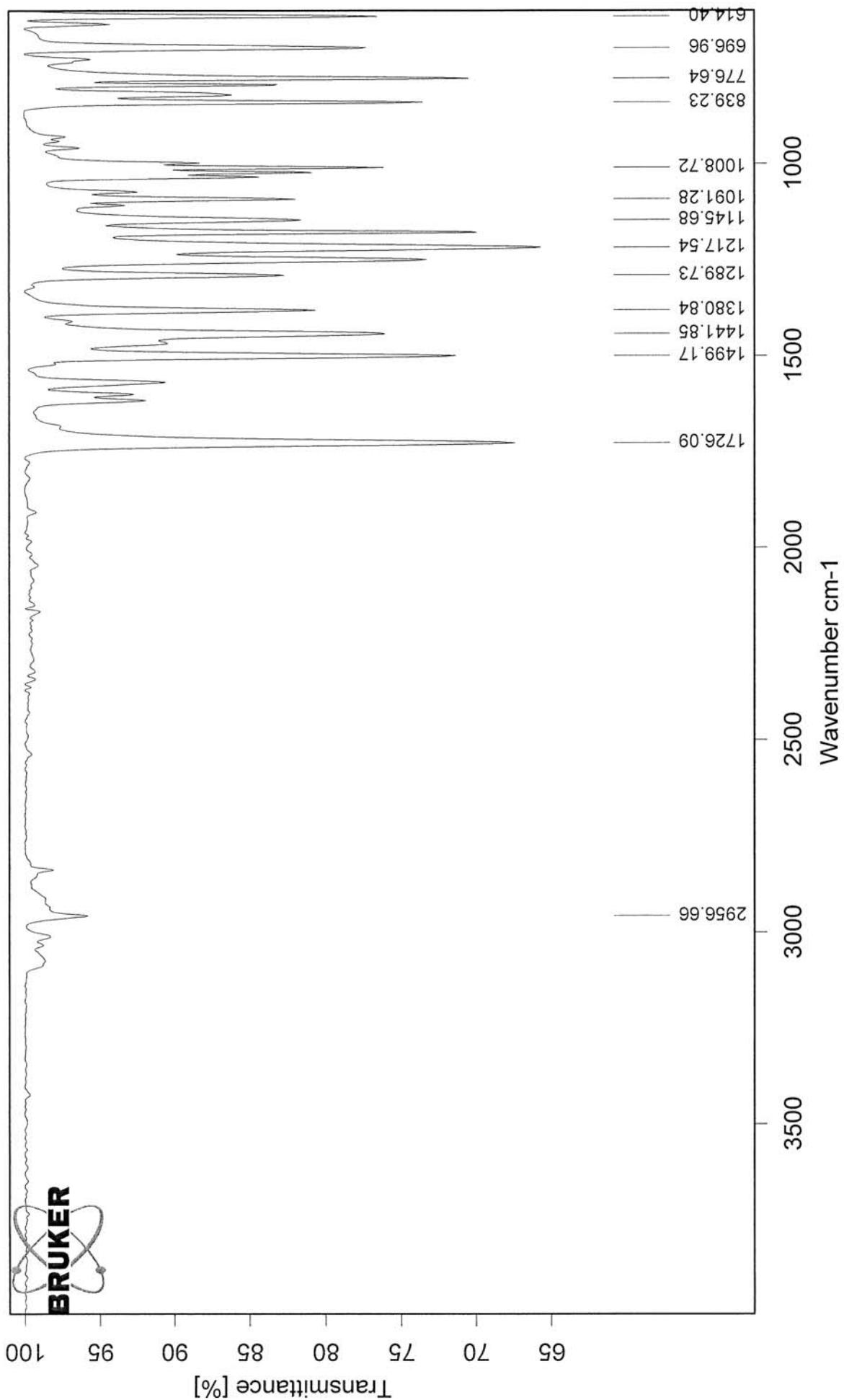
*** Mass Analysis for mass 310.1181560

1	17	16	3	3	310.1186178	11.5	4.618e-04
2	15	14	6	2	310.1172751	12.0	8.809e-04
3	12	16	5	5	310.1145951	7.5	3.561e-03
4	22	16	1	1	310.1226406	15.5	4.485e-03
5	11	16	7	4	310.1258285	7.5	7.672e-03
6	21	14	2	1	310.1100645	16.0	8.091e-03
7	13	18	4	5	310.1271711	7.0	9.015e-03
8	18	16	1	4	310.1073844	11.5	1.077e-02
9	16	16	5	2	310.1298512	11.5	1.170e-02
10	16	14	4	3	310.1060418	12.0	1.211e-02

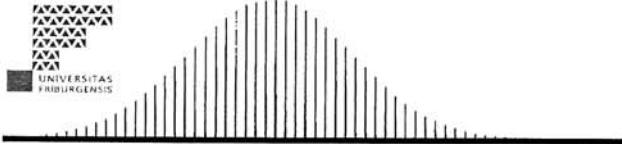








U:\IR_spectra\IS110C02.0	IS110C02	white solid	27/09/2010
--------------------------	----------	-------------	------------



FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

ESI-MS: ISOC02

XMASS Mass Analysis for /Data/UNI_FR/ALIM5077_ESI/5/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

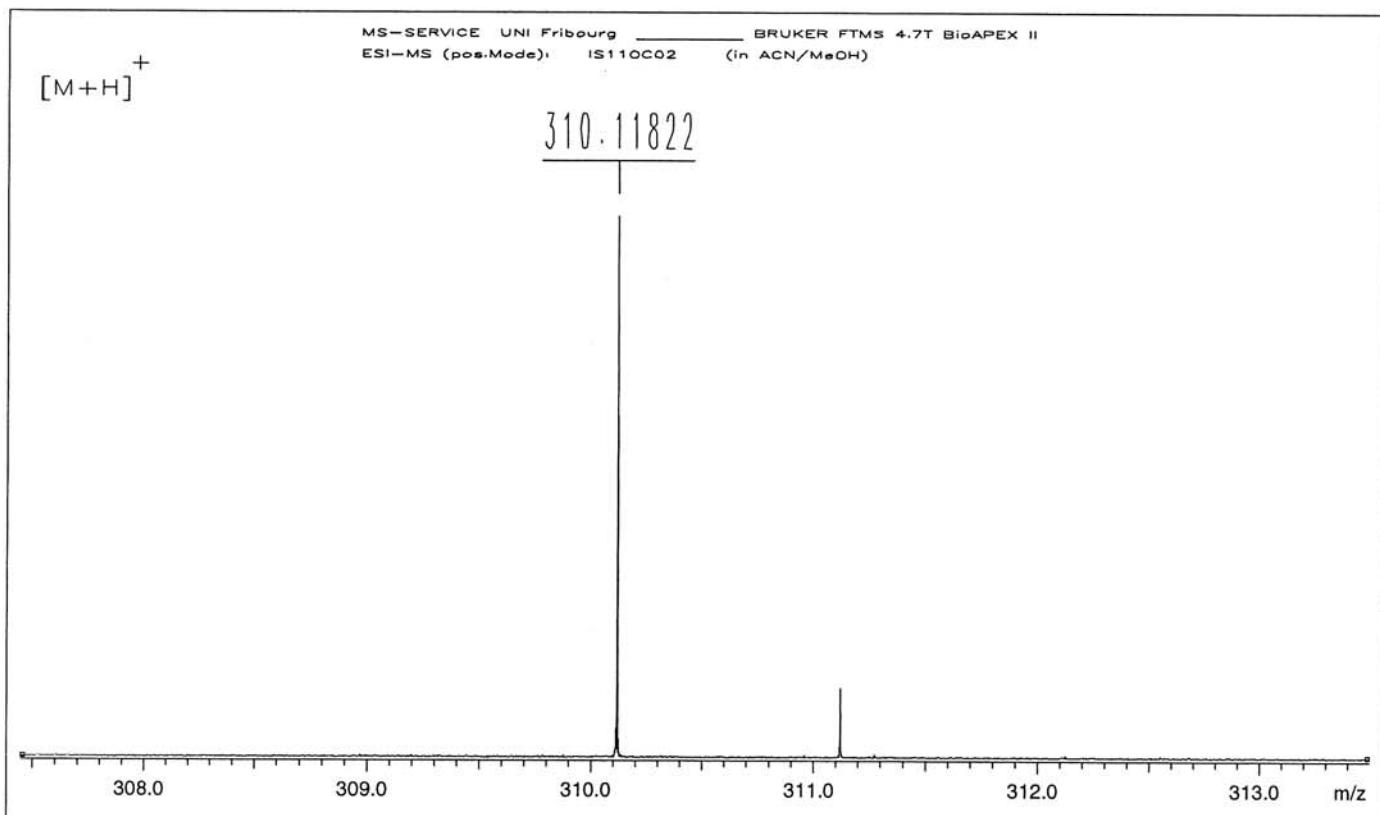
Ion mass = 310.1182180

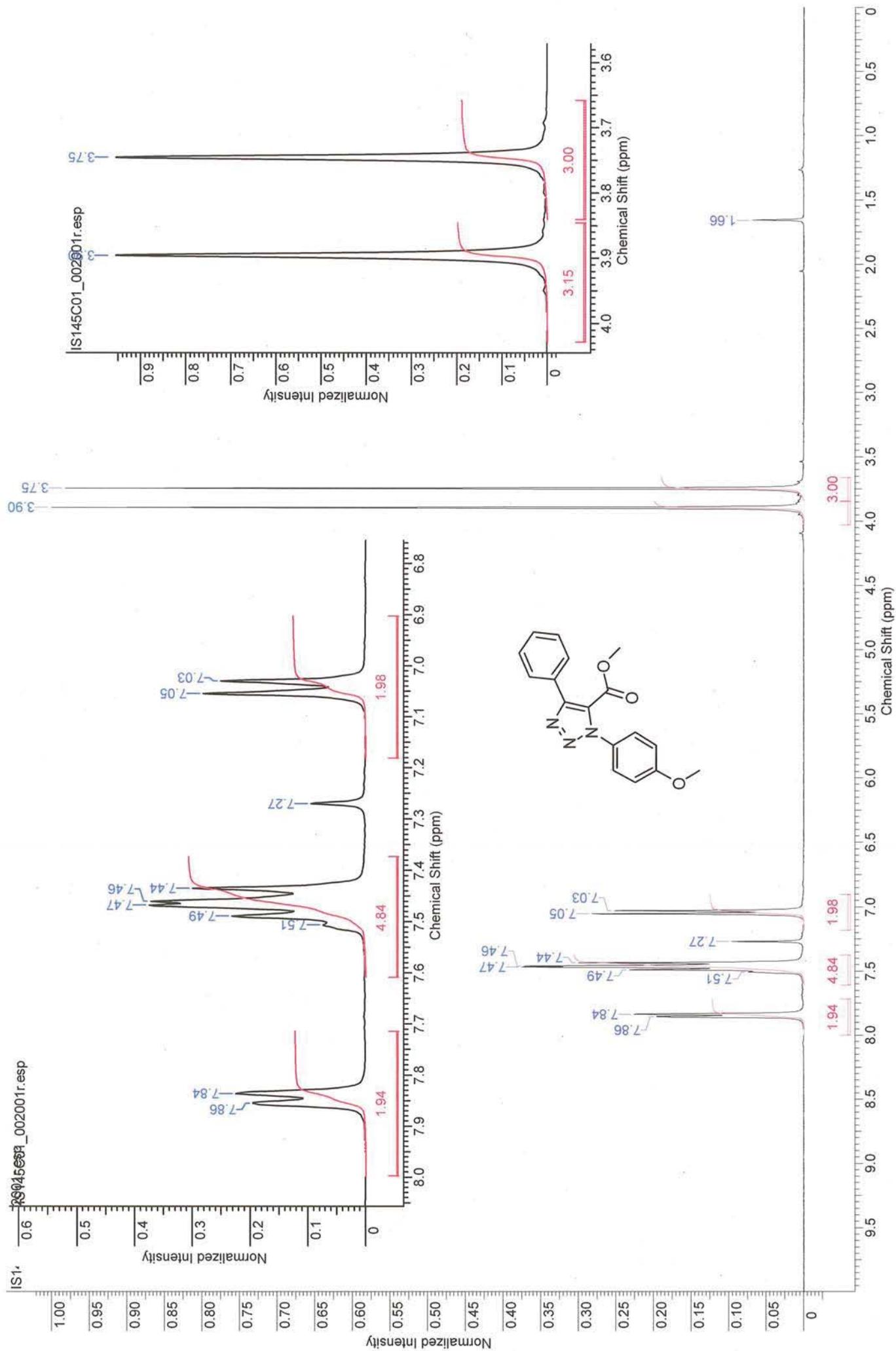
Charge = +1

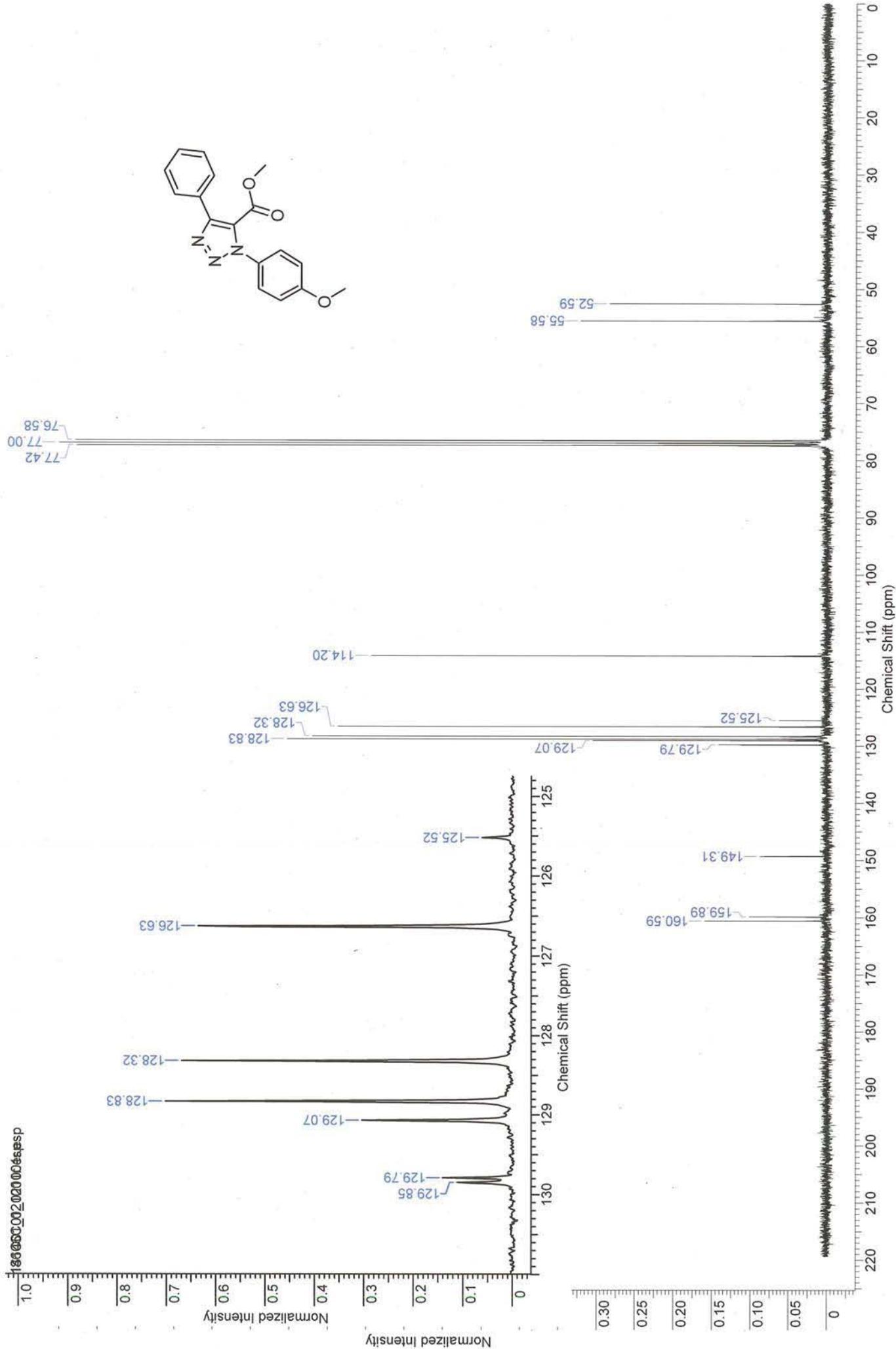
#	C	H	N	O	mass	DBE	error
---	---	---	---	---	------	-----	-------

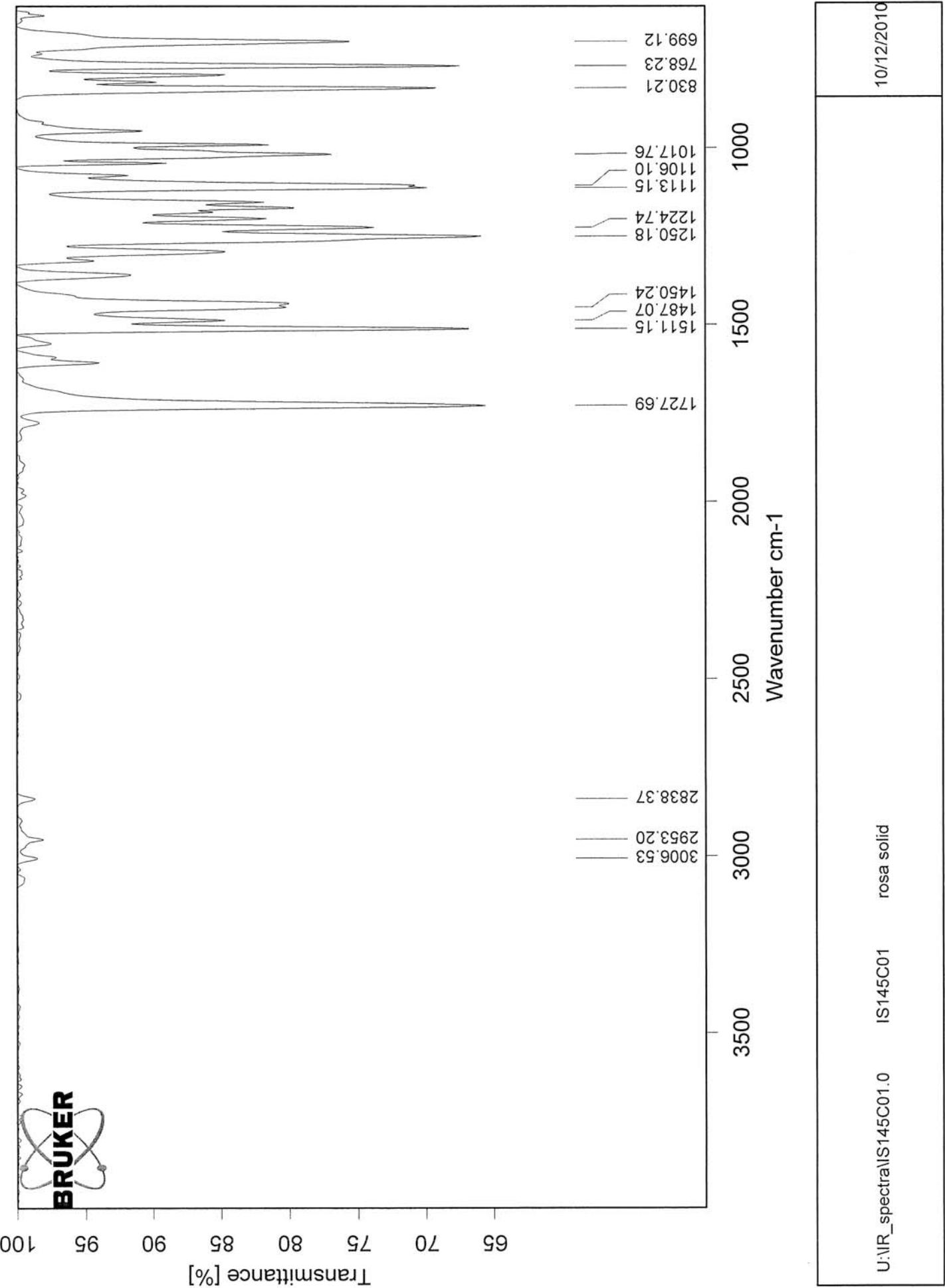
*** Mass Analysis for mass 310.1182180

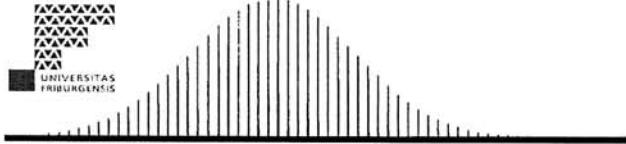
1	17	16	3	3	310.1186178	11.5	3.998e-04
2	15	14	6	2	310.1172751	12.0	9.429e-04
3	12	16	5	5	310.1145951	7.5	3.623e-03
4	22	16	1	1	310.1226406	15.5	4.423e-03
5	11	16	7	4	310.1258285	7.5	7.610e-03
6	21	14	2	1	310.1100645	16.0	8.153e-03
7	13	18	4	5	310.1271711	7.0	8.953e-03
8	18	16	1	4	310.1073844	11.5	1.083e-02
9	16	16	5	2	310.1298512	11.5	1.163e-02
10	16	14	4	3	310.1060418	12.0	1.218e-02











BRUKER FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

ESI-MS: IS145C01

XMASS Mass Analysis for /Data/UNI_FR/ALIM5267_ESI/2/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

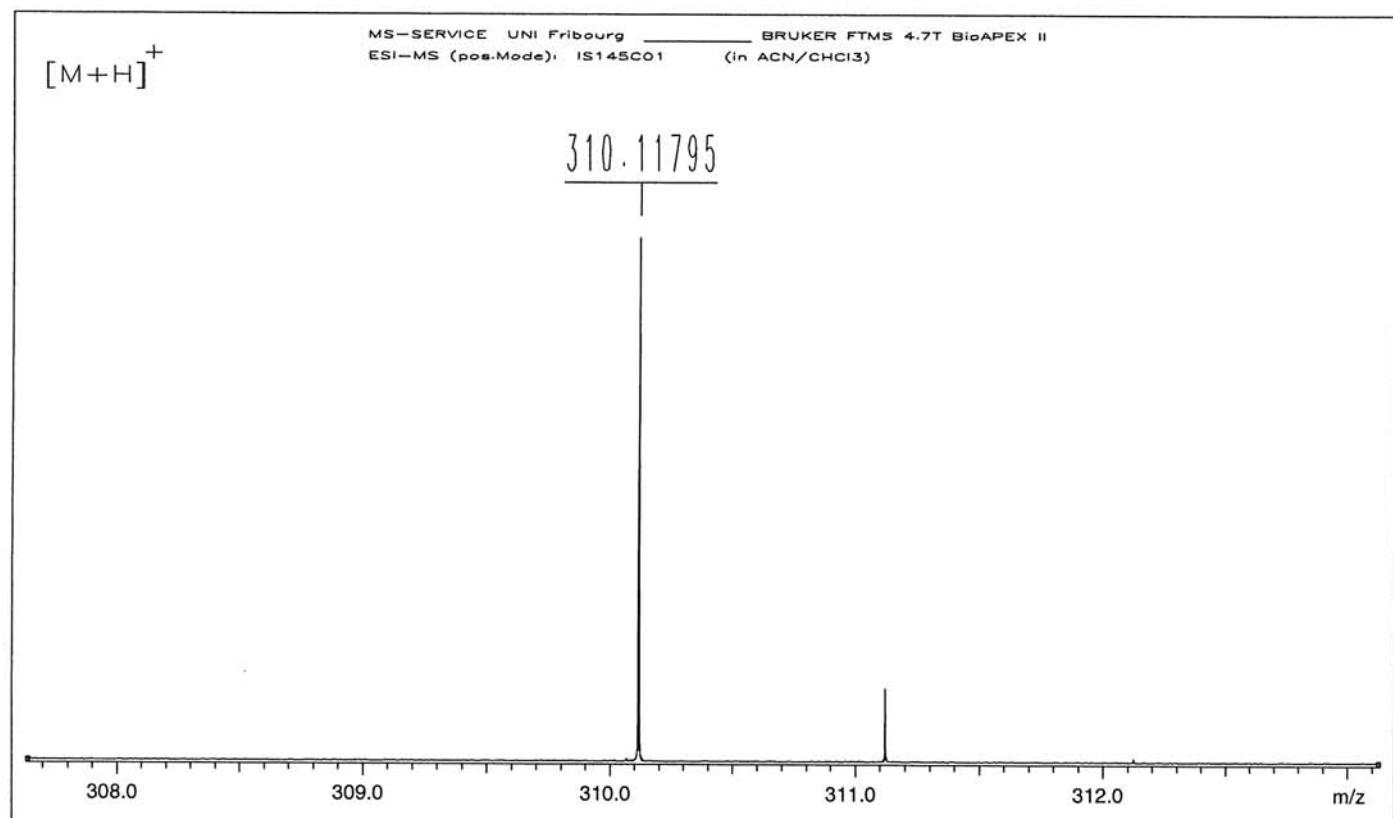
Ion mass = 310.1179500

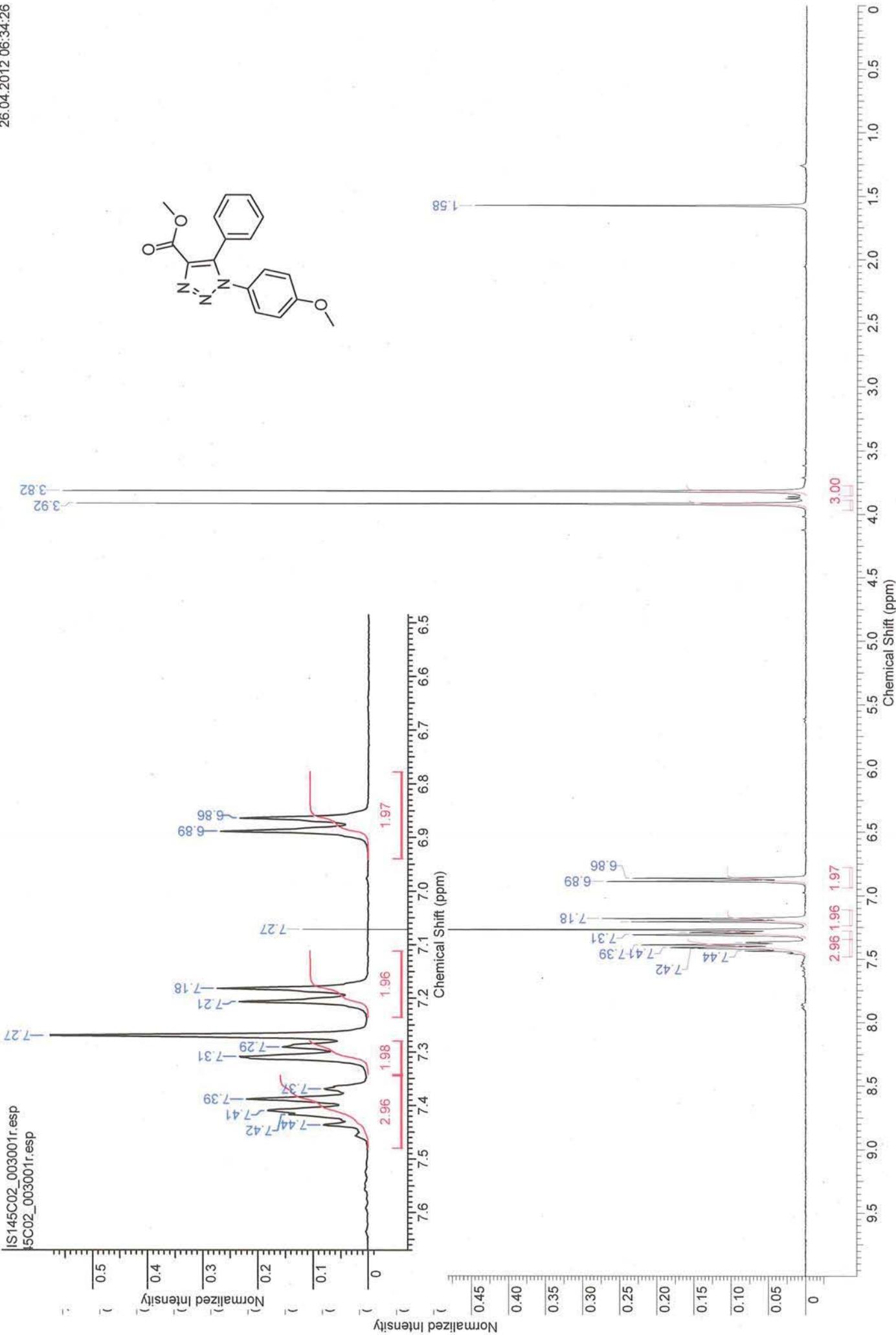
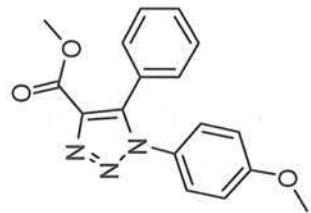
Charge = +1

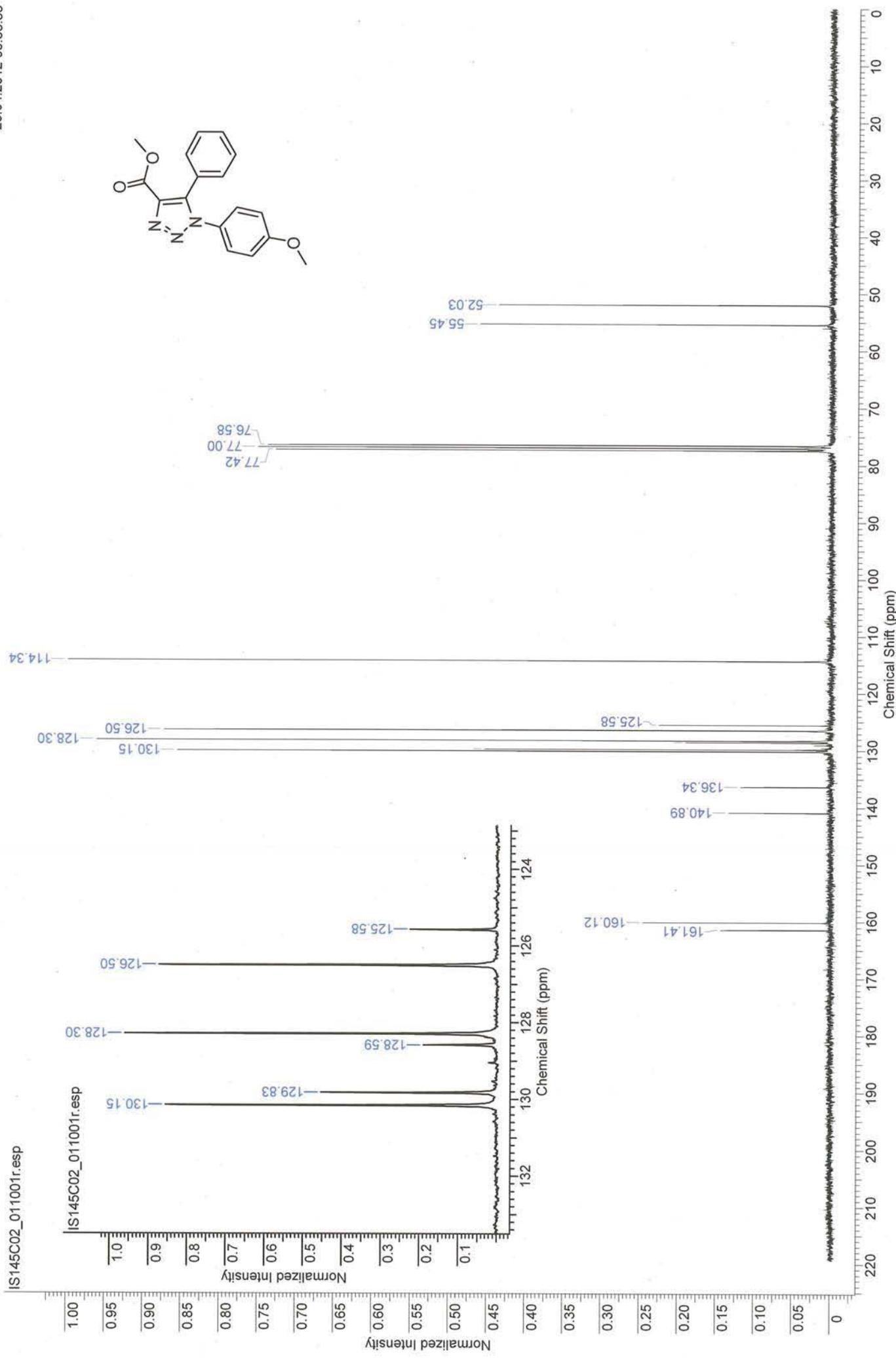
#	C	H	N	O	Na	mass	DBE	error
---	---	---	---	---	----	------	-----	-------

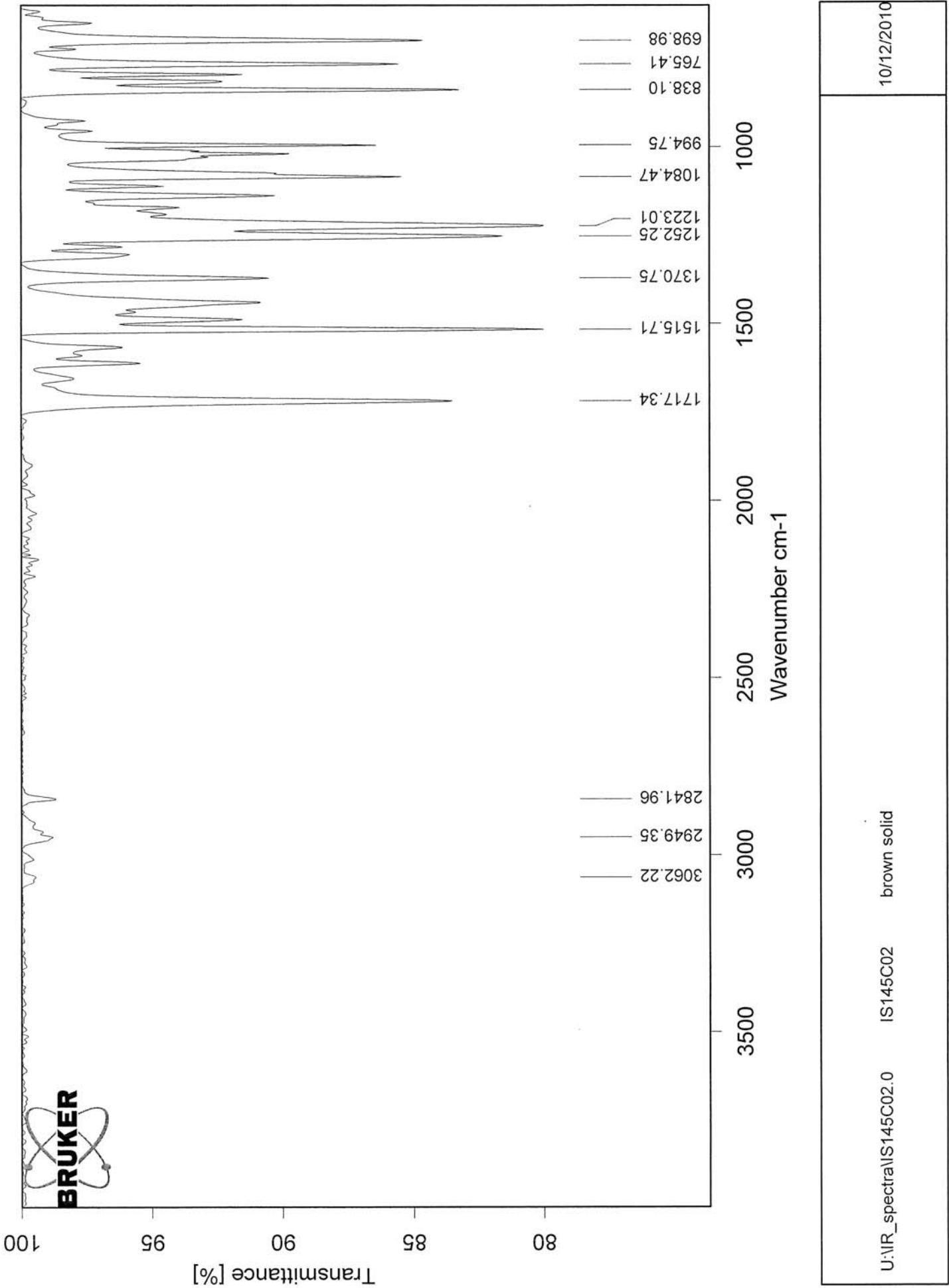
*** Mass Analysis for mass 310.1179500

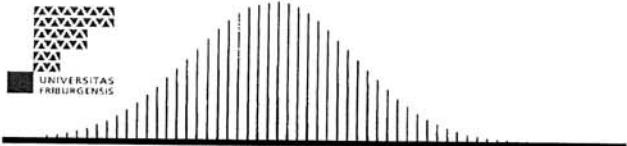
1	17	16	3	3	0	310.1186178	11.5	6.678e-04
2	15	17	3	3	1	310.1162125	8.5	1.737e-03
3	14	18	2	6	0	310.1159377	7.0	2.012e-03
4	20	17	1	1	1	310.1202353	12.5	2.285e-03
5	8	21	3	8	1	310.1220858	-0.5	4.136e-03
6	12	19	2	6	1	310.1135324	4.0	4.418e-03
7	22	16	1	1	0	310.1226406	15.5	4.691e-03
8	11	20	1	9	0	310.1132577	2.5	4.692e-03
9	9	18	4	8	0	310.1119150	3.0	6.035e-03
10	10	20	3	8	0	310.1244911	2.5	6.541e-03











FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

ESI-MS: IS145C02

XMASS Mass Analysis for /Data/UNI_FR/ALIM5265_ESI/2/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

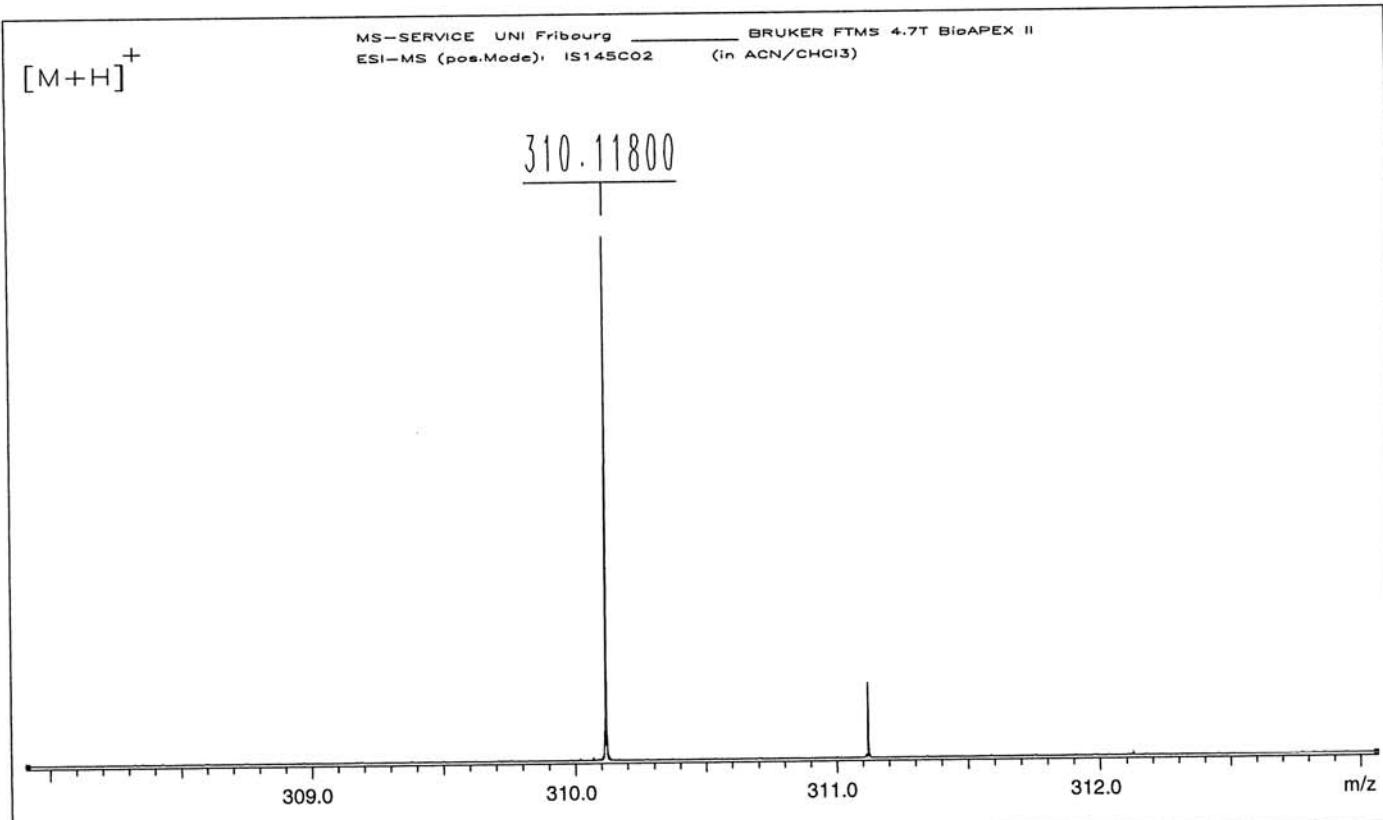
Ion mass = 310.1179980

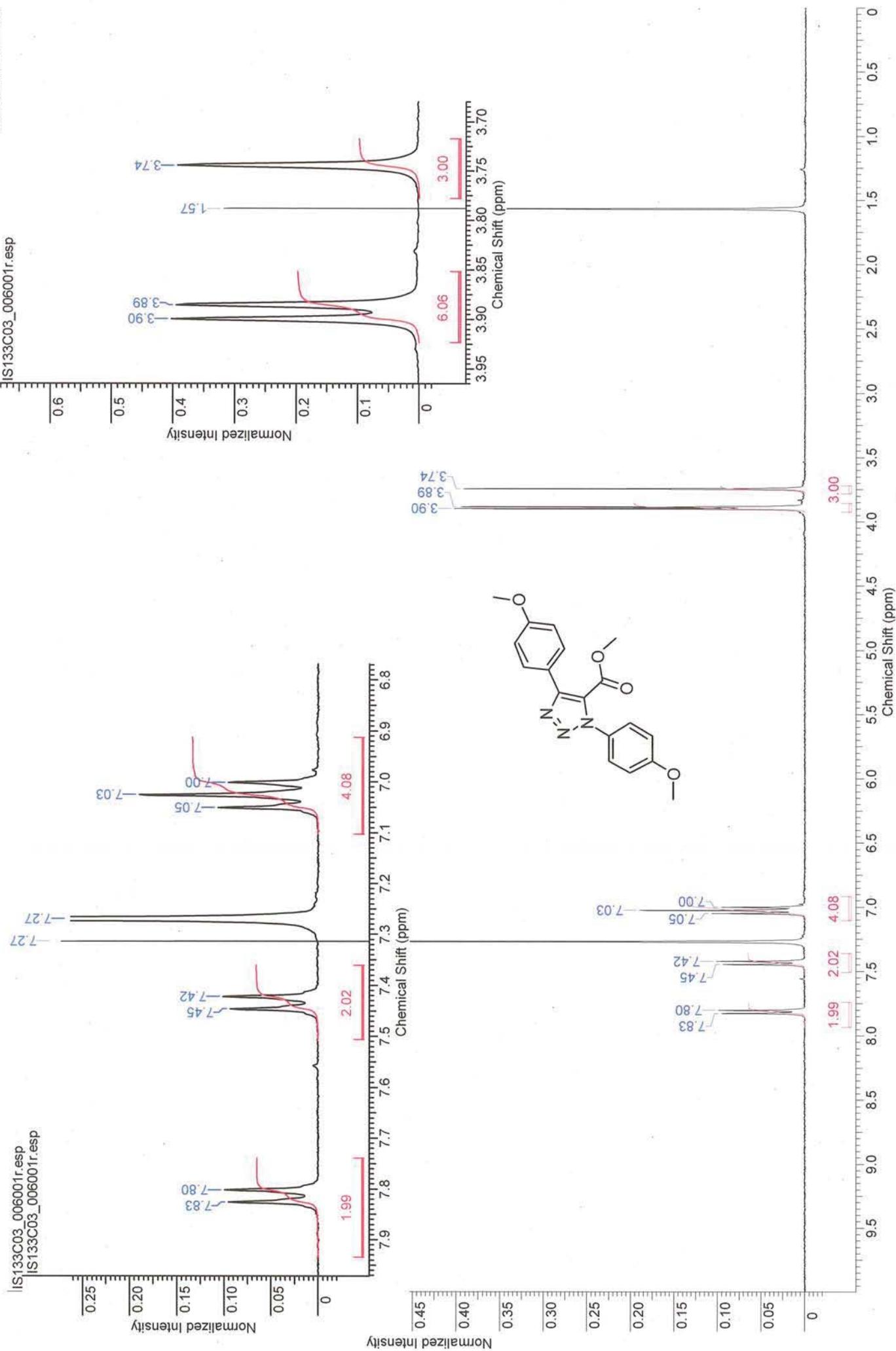
Charge = +1

#	C	H	N	O	Na	mass	DBE	error
---	---	---	---	---	----	------	-----	-------

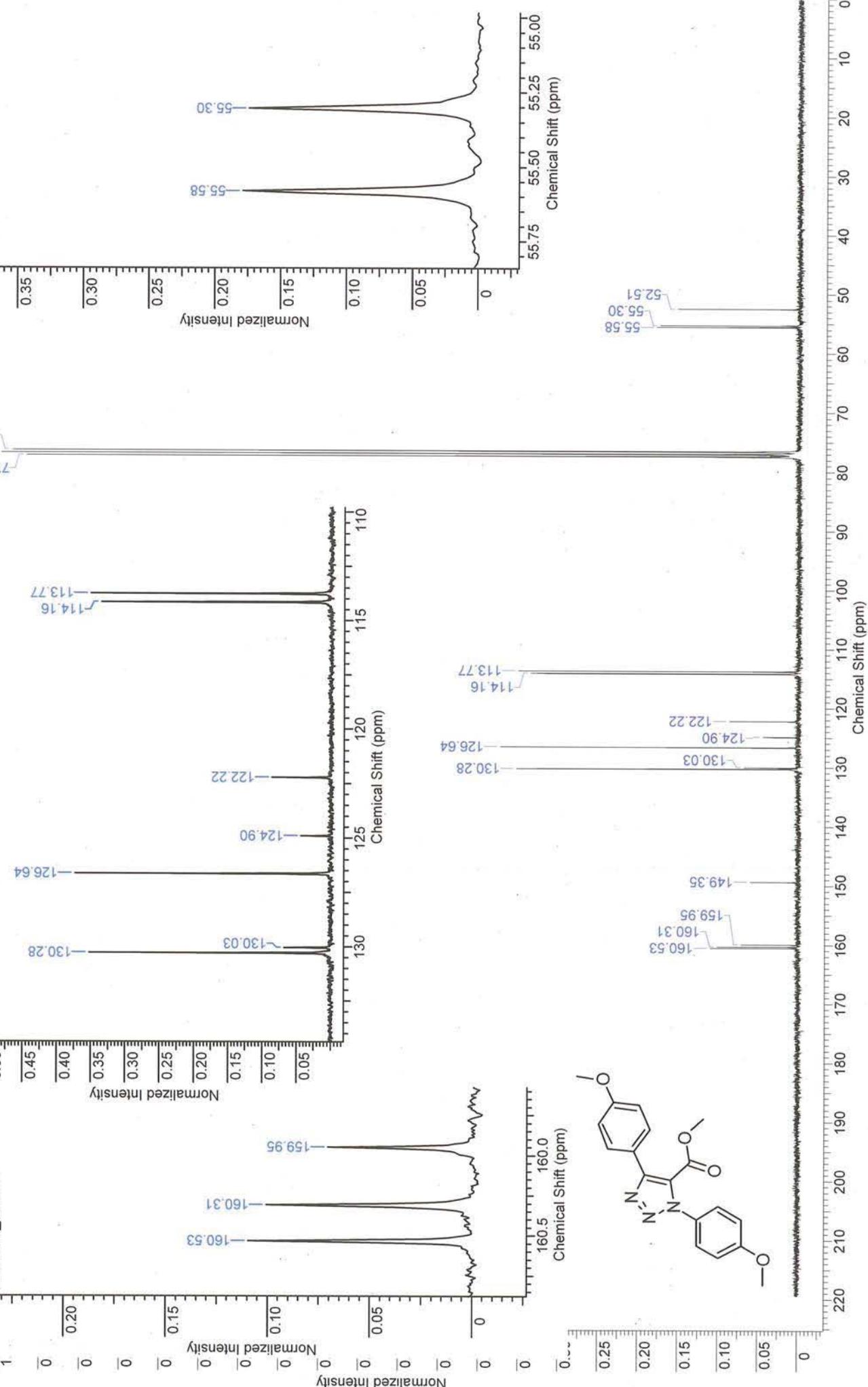
*** Mass Analysis for mass 310.1179980

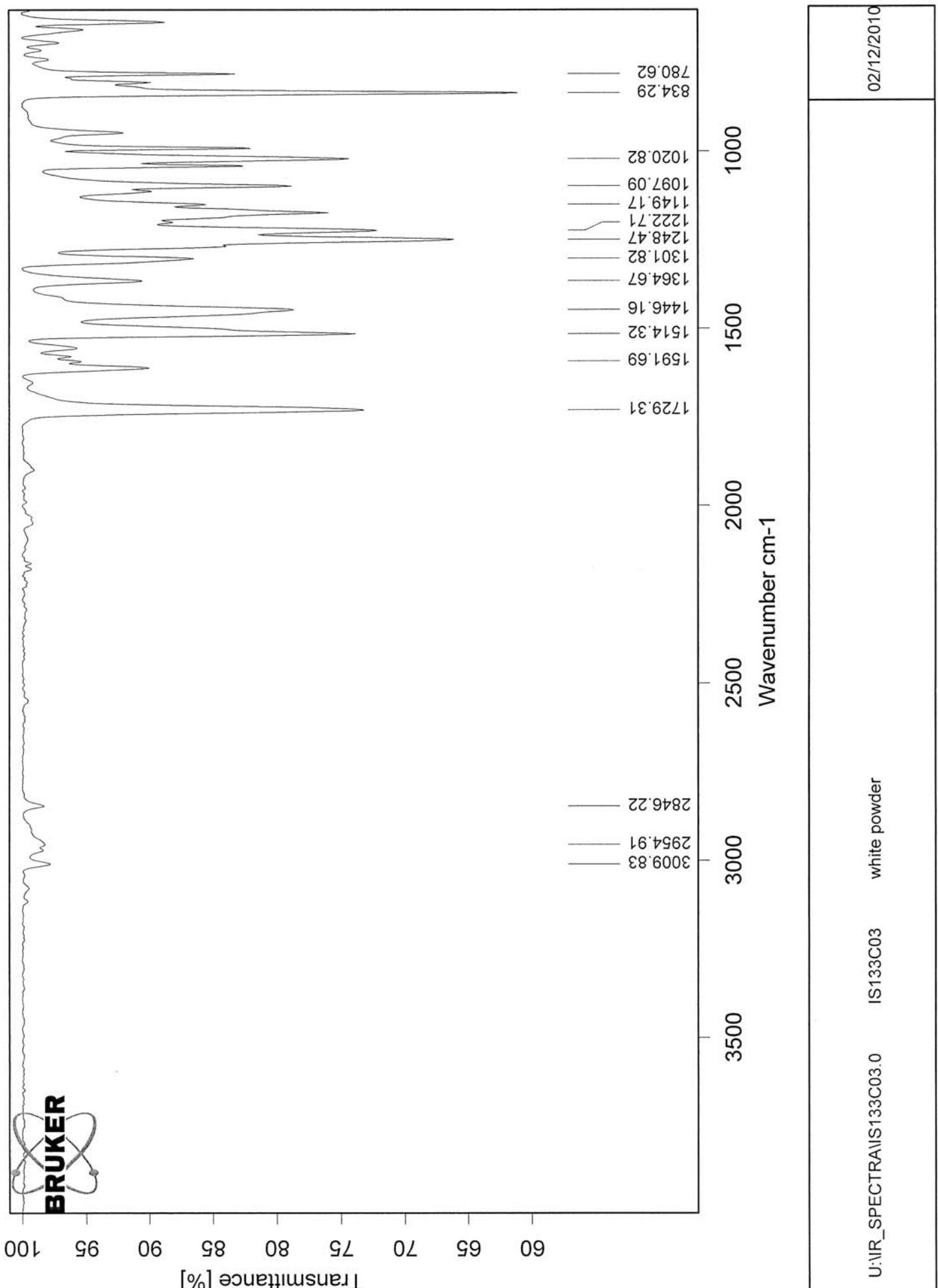
1	17	16	3	3	0	310.1186178	11.5	6.198e-04
2	15	14	6	2	0	310.1172751	12.0	7.229e-04
3	15	17	3	3	1	310.1162125	8.5	1.785e-03
4	14	18	2	6	0	310.1159377	7.0	2.060e-03
5	13	12	9	1	0	310.1159325	12.5	2.066e-03
6	20	17	1	1	1	310.1202353	12.5	2.237e-03
7	5	20	5	10	0	310.1204683	-1.5	2.470e-03
8	6	19	6	7	1	310.1207431	0.0	2.745e-03
9	13	15	6	2	1	310.1148698	9.0	3.128e-03
10	12	16	5	5	0	310.1145951	7.5	3.403e-03

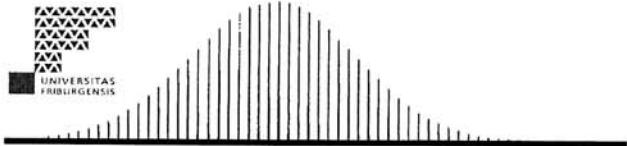




IS133C03_0200001r







FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

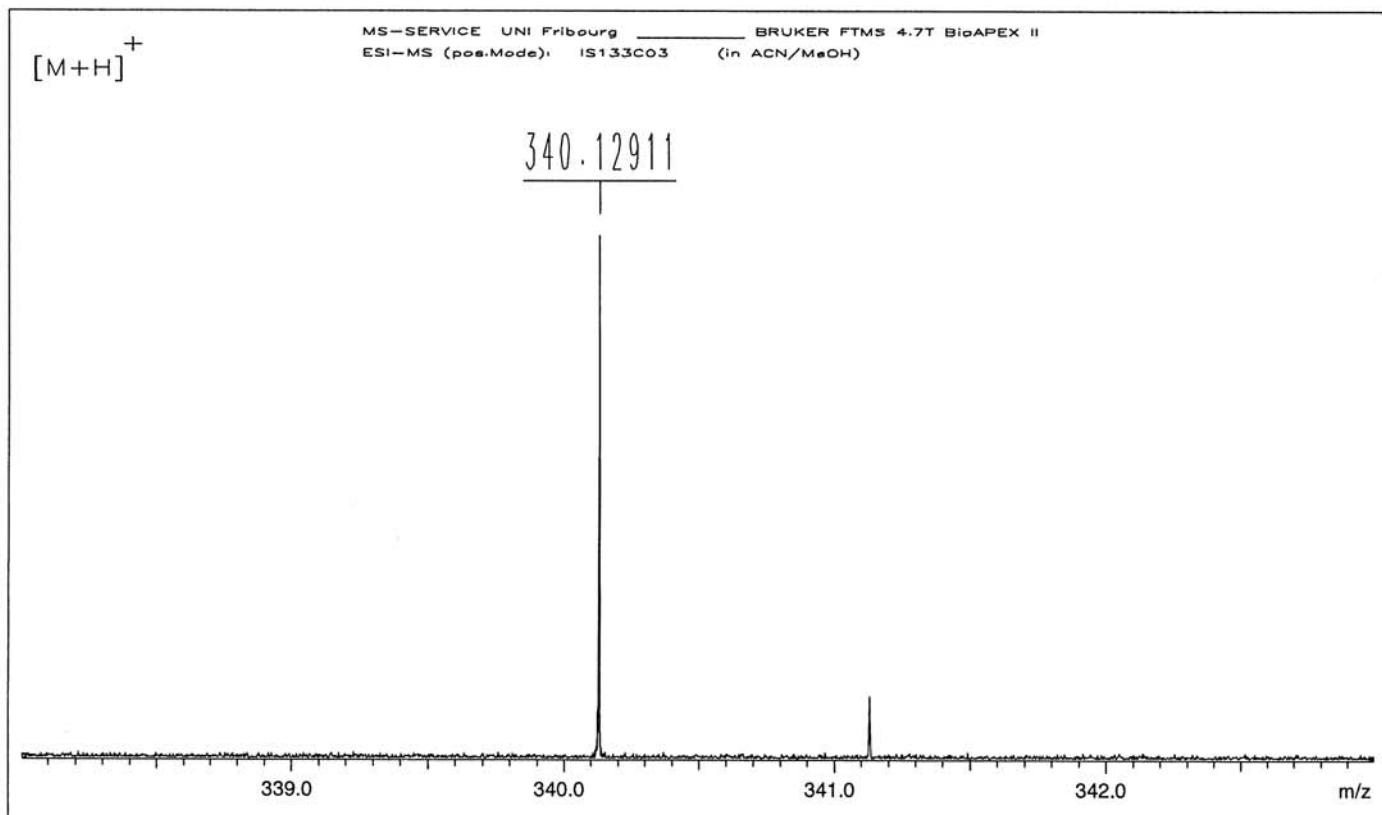
ESI-MS: IS133C03

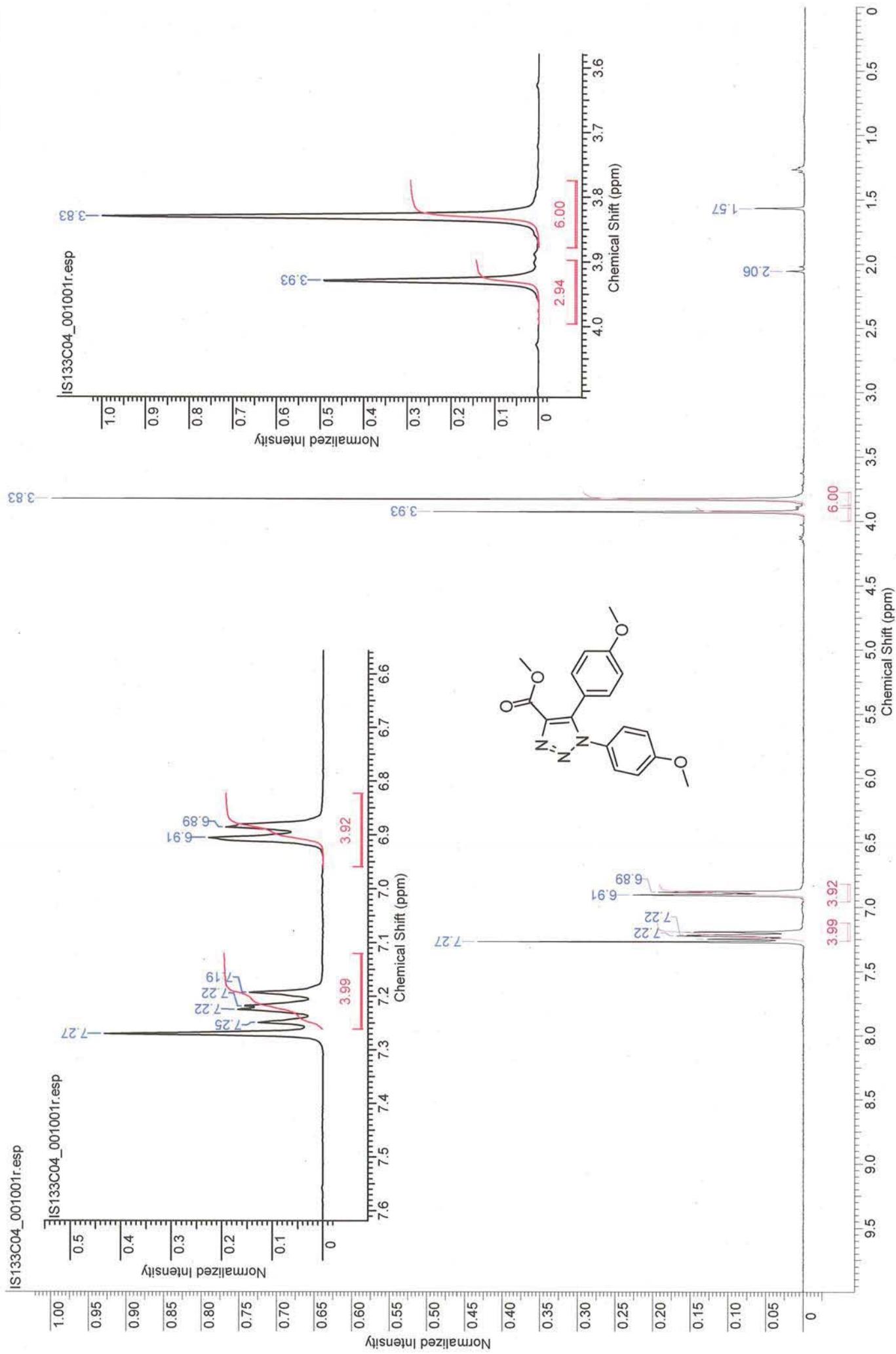
XMASS Mass Analysis for /Data/UNI_FR/ALIM5246_ESI/5/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

Ion mass = 340.1291080

Charge = +1

#	C	H	N	O	mass	DBE	error
*** Mass Analysis for mass 340.1291080							
1	18	18	3	4	340.1291825	11.5	7.450e-05
2	16	16	6	3	340.1278398	12.0	1.268e-03
3	15	20	2	7	340.1265024	7.0	2.606e-03
4	14	14	9	2	340.1264972	12.5	2.611e-03
5	21	16	4	1	340.1318626	16.0	2.755e-03
6	7	18	9	7	340.1323704	3.5	3.262e-03
7	13	18	5	6	340.1251598	7.5	3.948e-03
8	23	18	1	2	340.1332052	15.5	4.097e-03
9	9	20	6	8	340.1337131	3.0	4.605e-03
10	12	22	1	10	340.1238224	2.5	5.286e-03

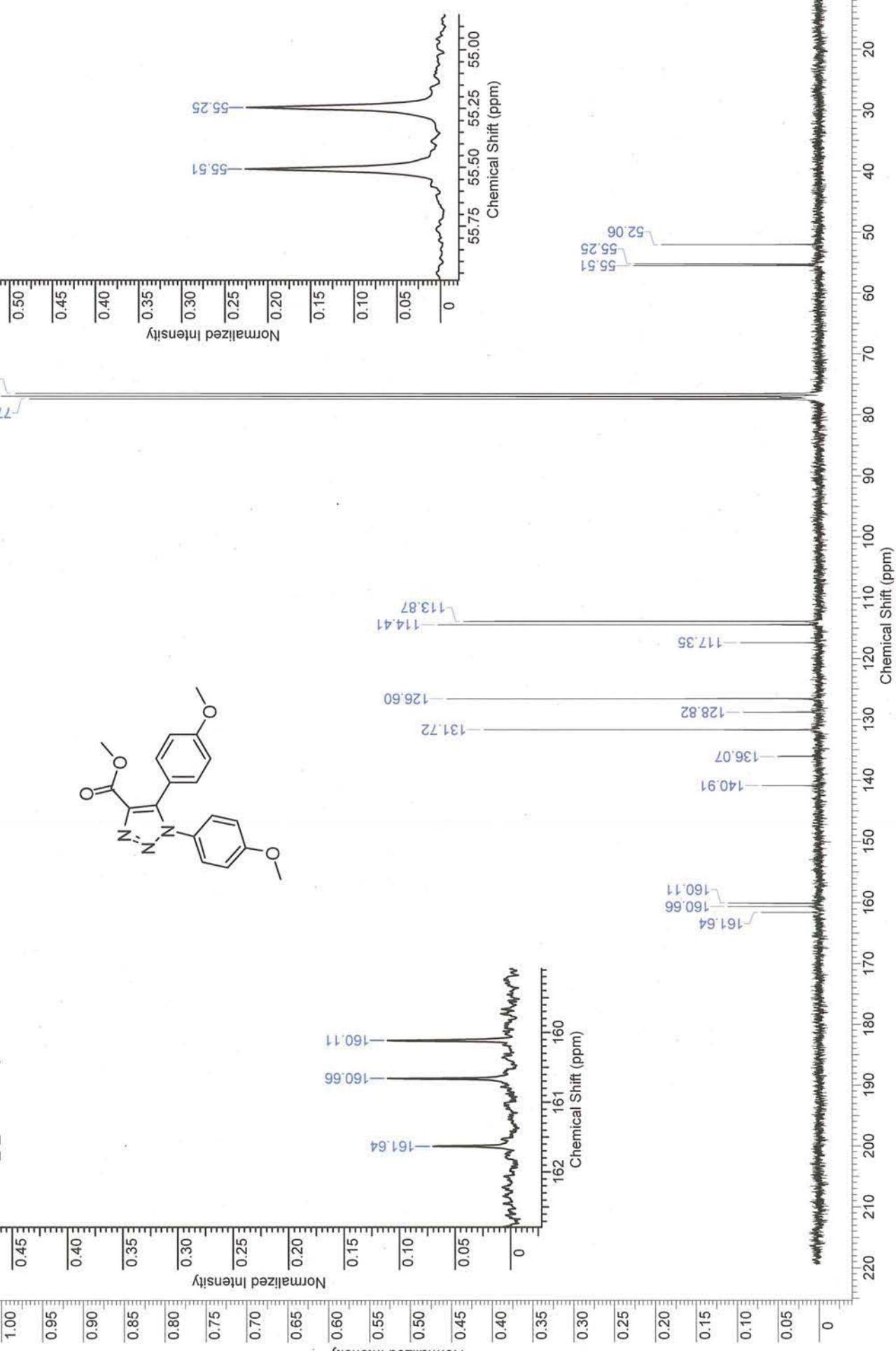


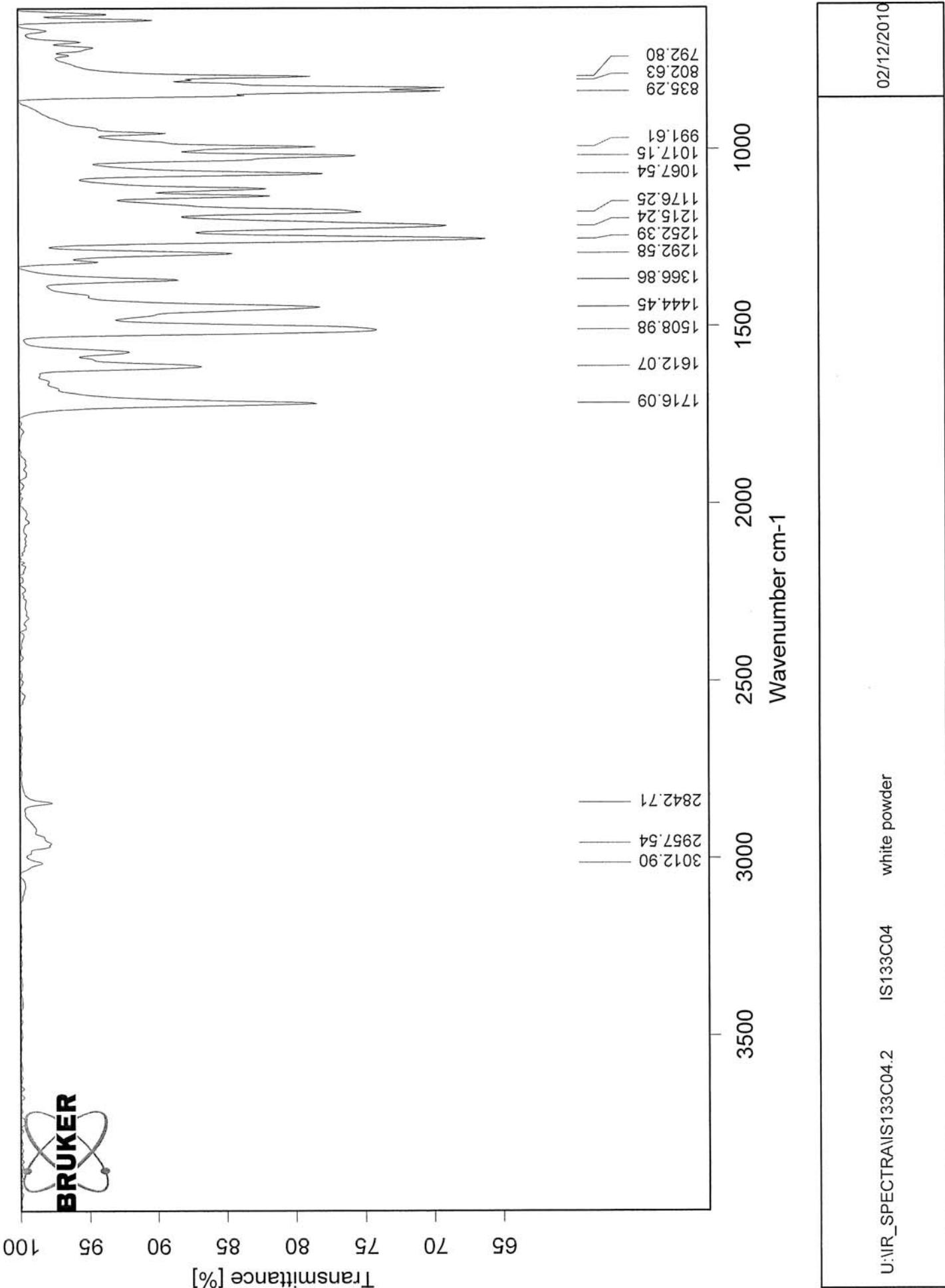


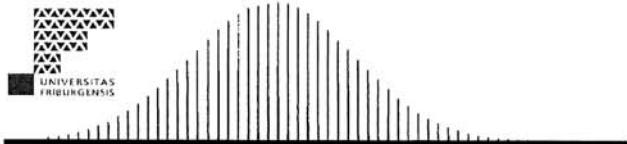
IS133C04_2_021001r.esp

77.42
77.00
76.58

IS133C04_2_021001r.esp







FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

ESI-MS: IS133C04

XMASS Mass Analysis for /Data/UNI_FR/ALIM5244_ESI/5/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

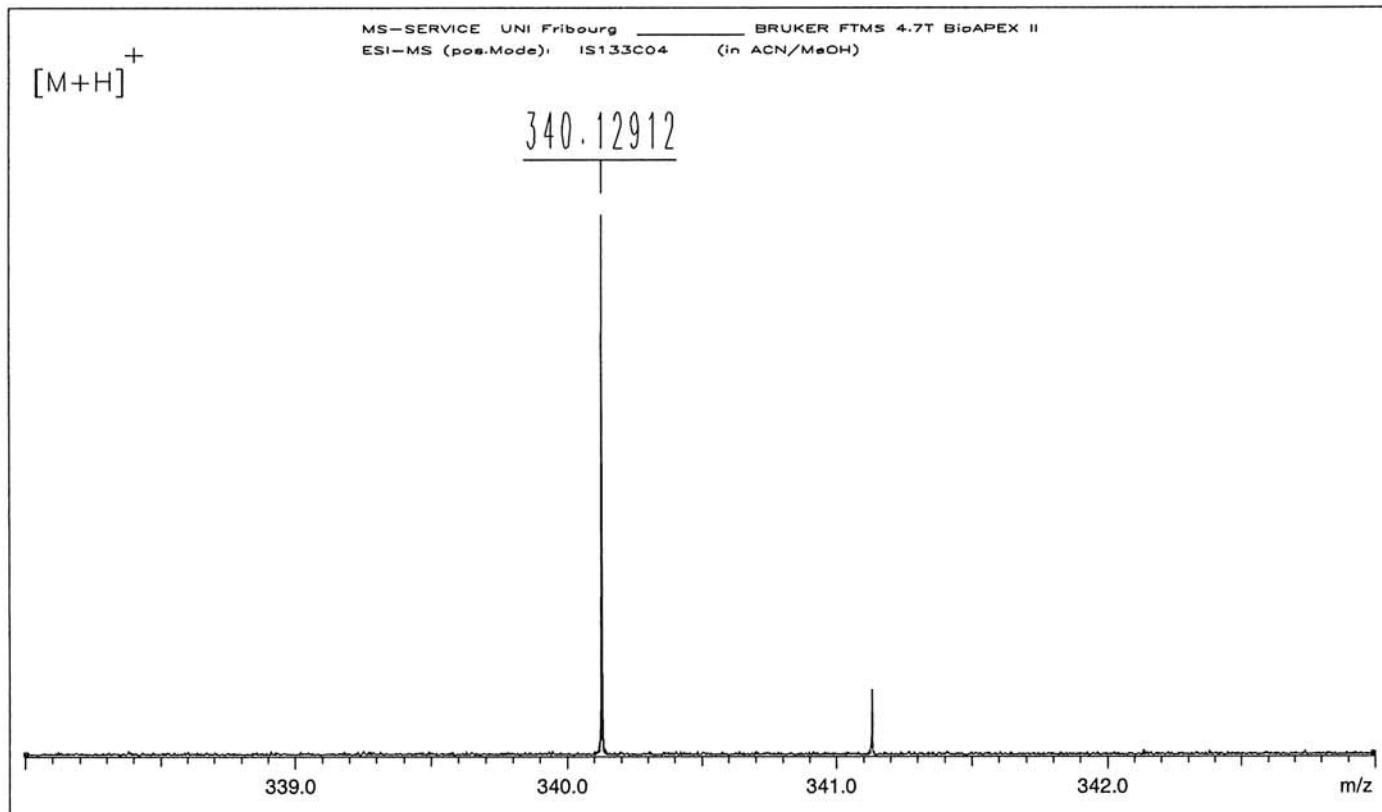
Ion mass = 340.1291160

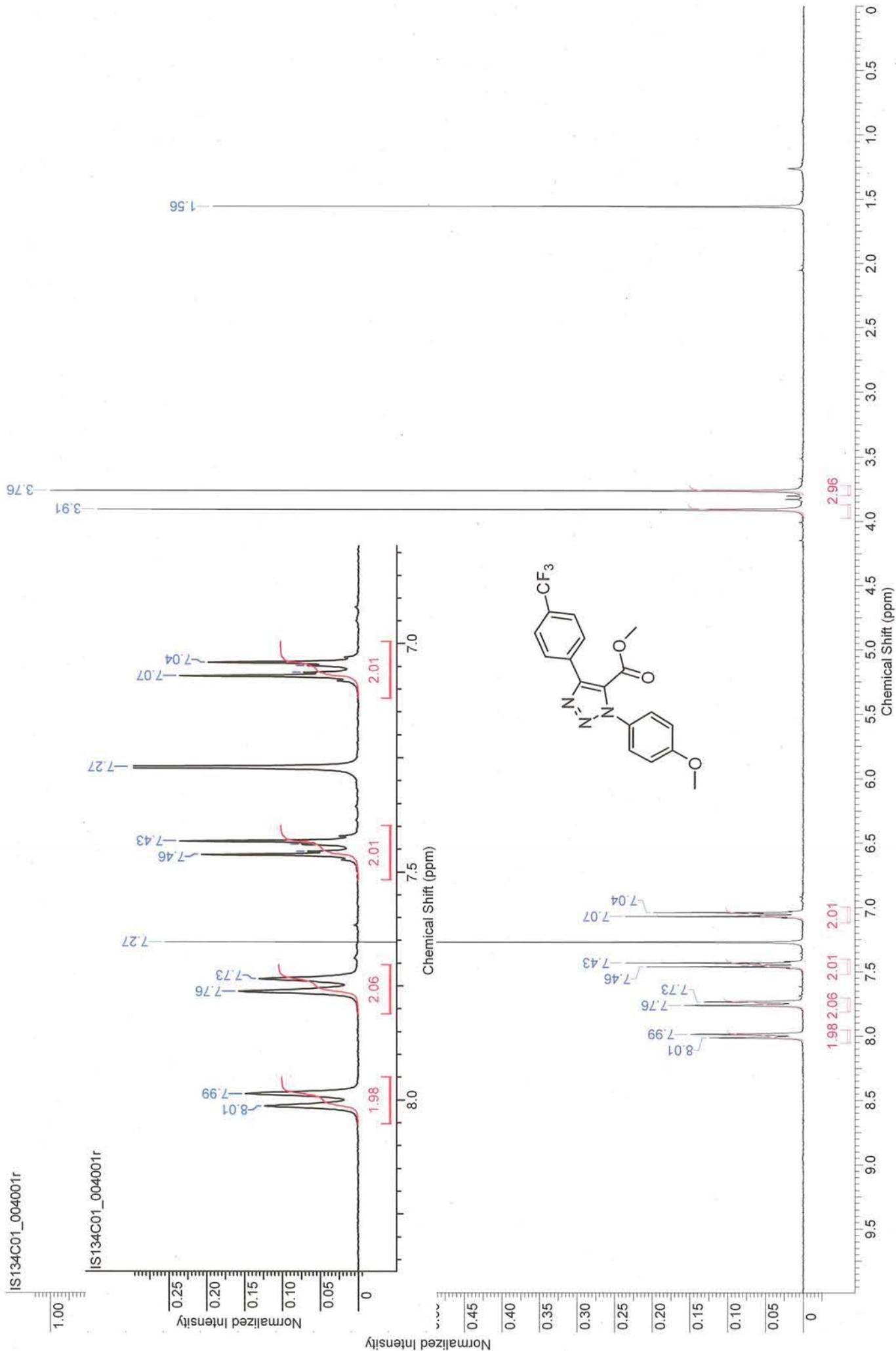
Charge = +1

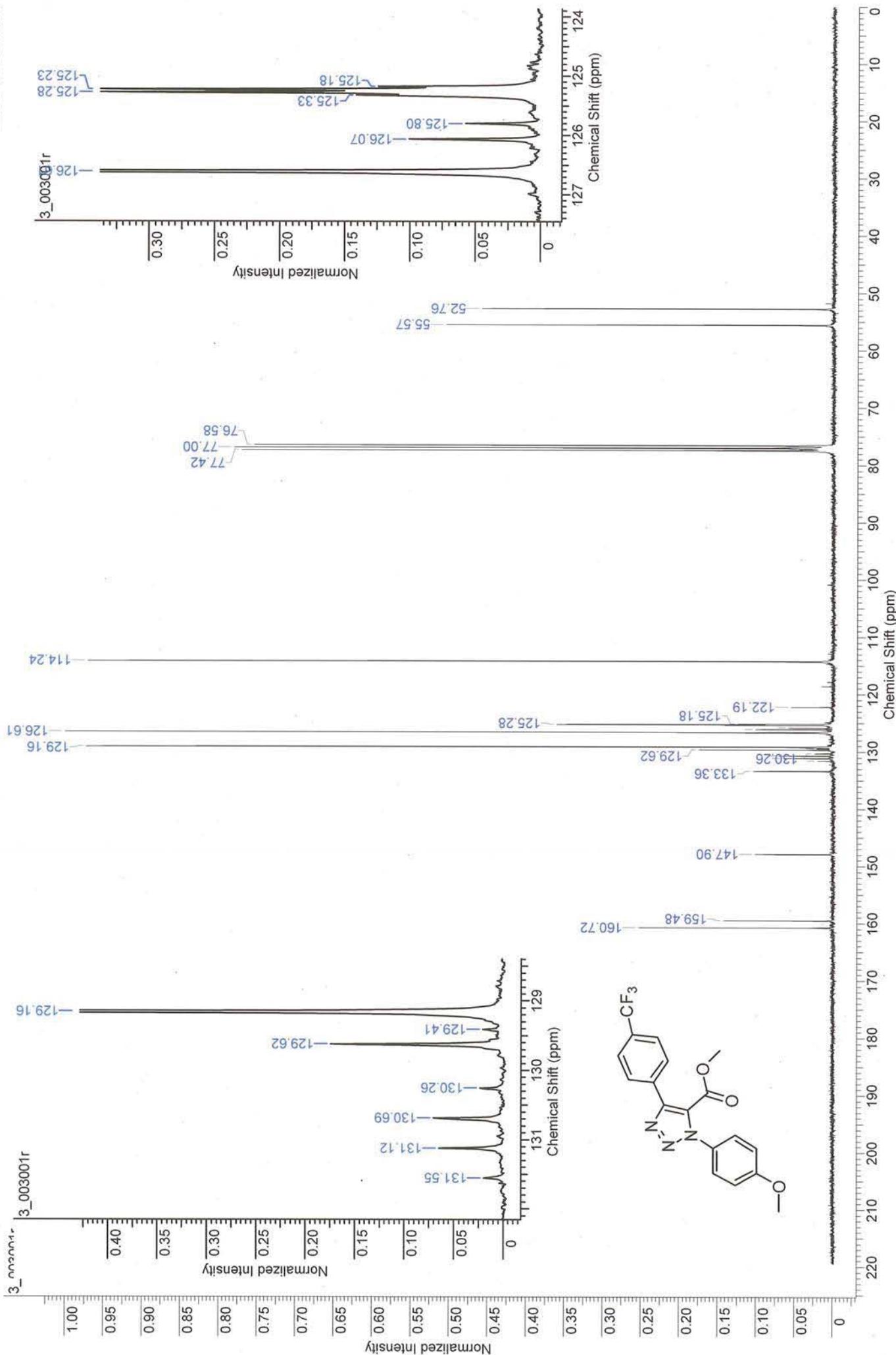
#	C	H	N	O	mass	DBE	error
---	---	---	---	---	------	-----	-------

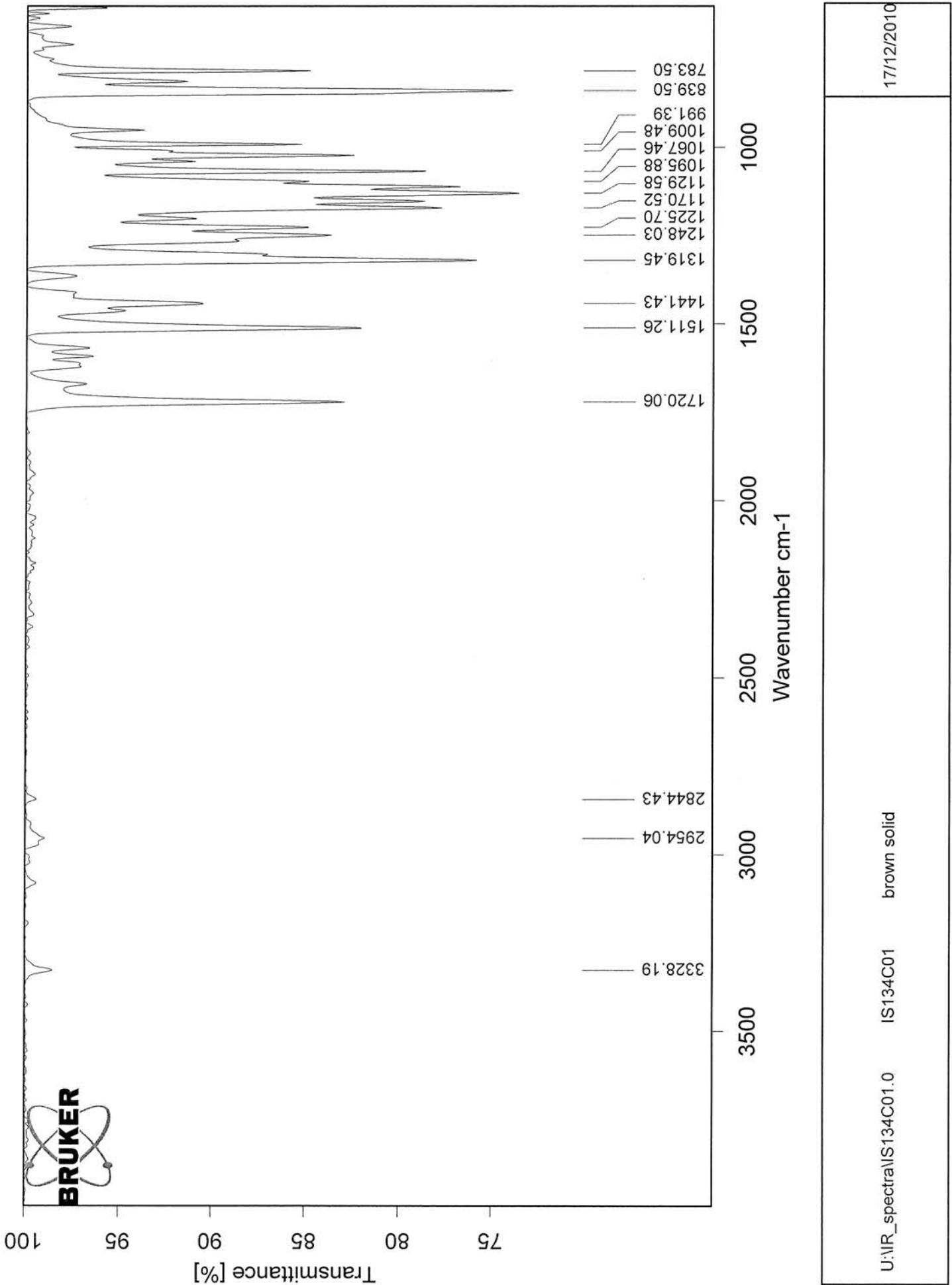
*** Mass Analysis for mass 340.1291160

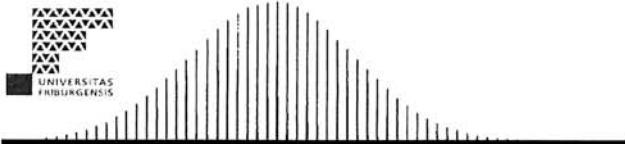
1	18	18	3	4	340.1291825	11.5	6.650e-05
2	16	16	6	3	340.1278398	12.0	1.276e-03
3	15	20	2	7	340.1265024	7.0	2.614e-03
4	14	14	9	2	340.1264972	12.5	2.619e-03
5	21	16	4	1	340.1318626	16.0	2.747e-03
6	7	18	9	7	340.1323704	3.5	3.254e-03
7	13	18	5	6	340.1251598	7.5	3.956e-03
8	23	18	1	2	340.1332052	15.5	4.089e-03
9	9	20	6	8	340.1337131	3.0	4.597e-03
10	12	22	1	10	340.1238224	2.5	5.294e-03











FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

ESI-MS: IS134C01

XMASS Mass Analysis for /Data/UNI_FR/ALIM5264_ESI/2/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

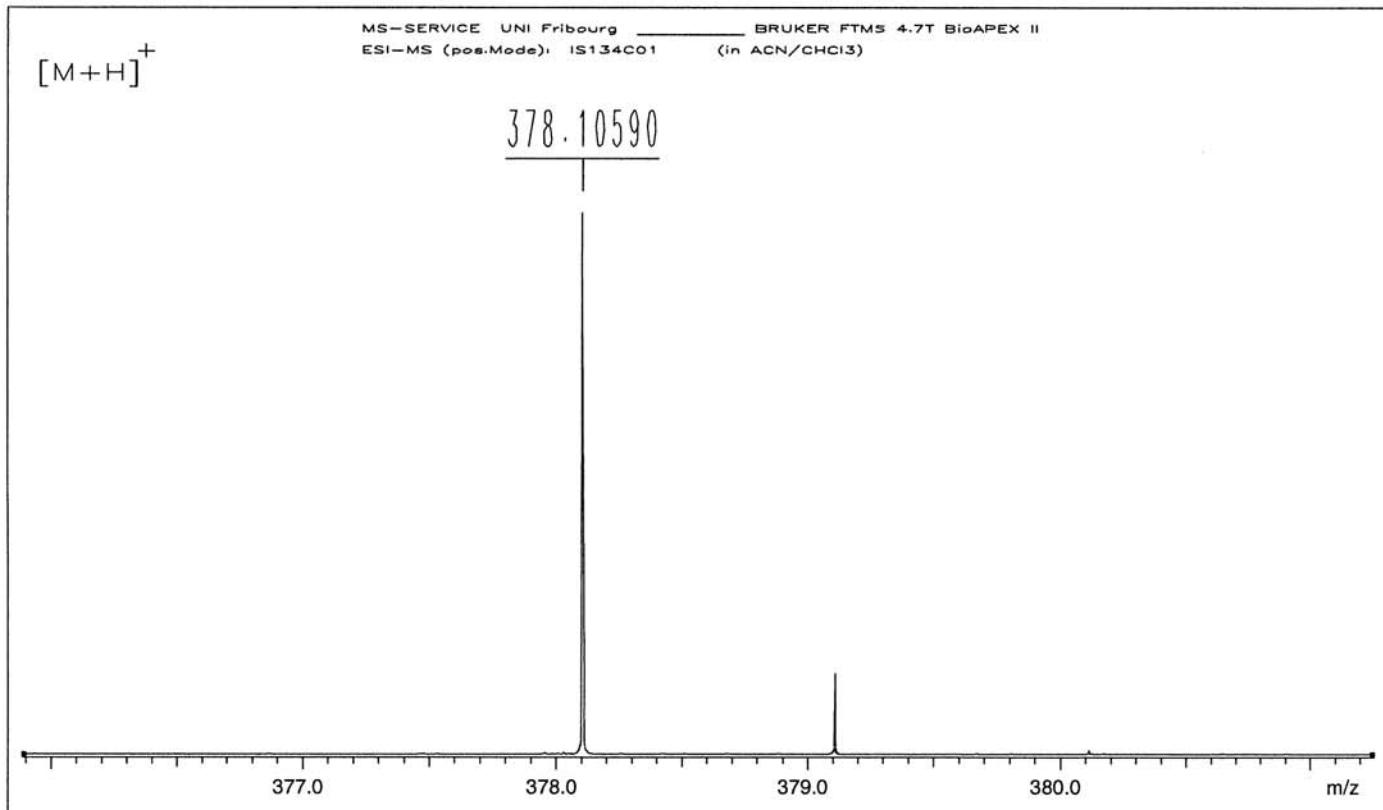
Ion mass = 378.1058990

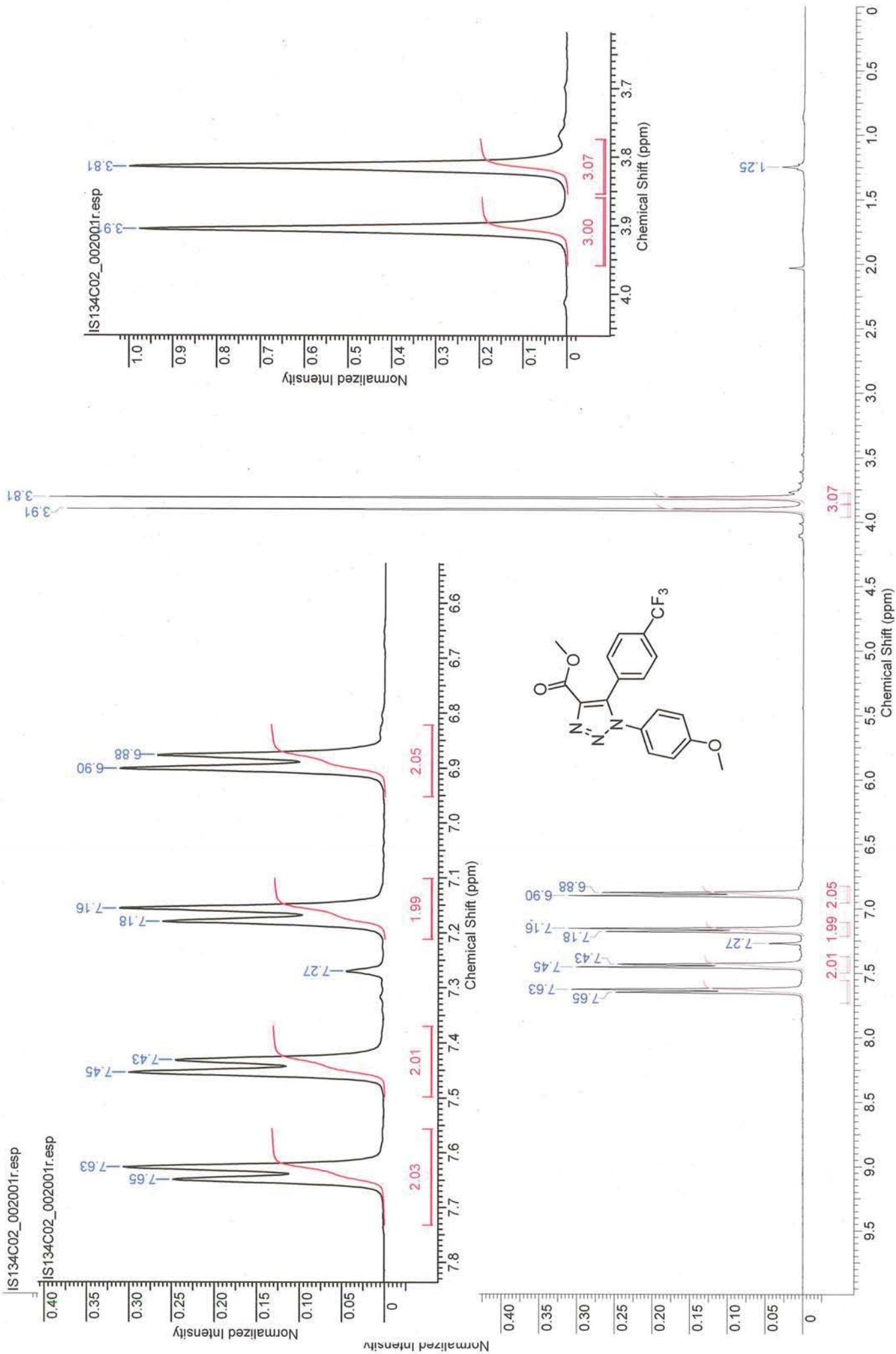
Charge = +1

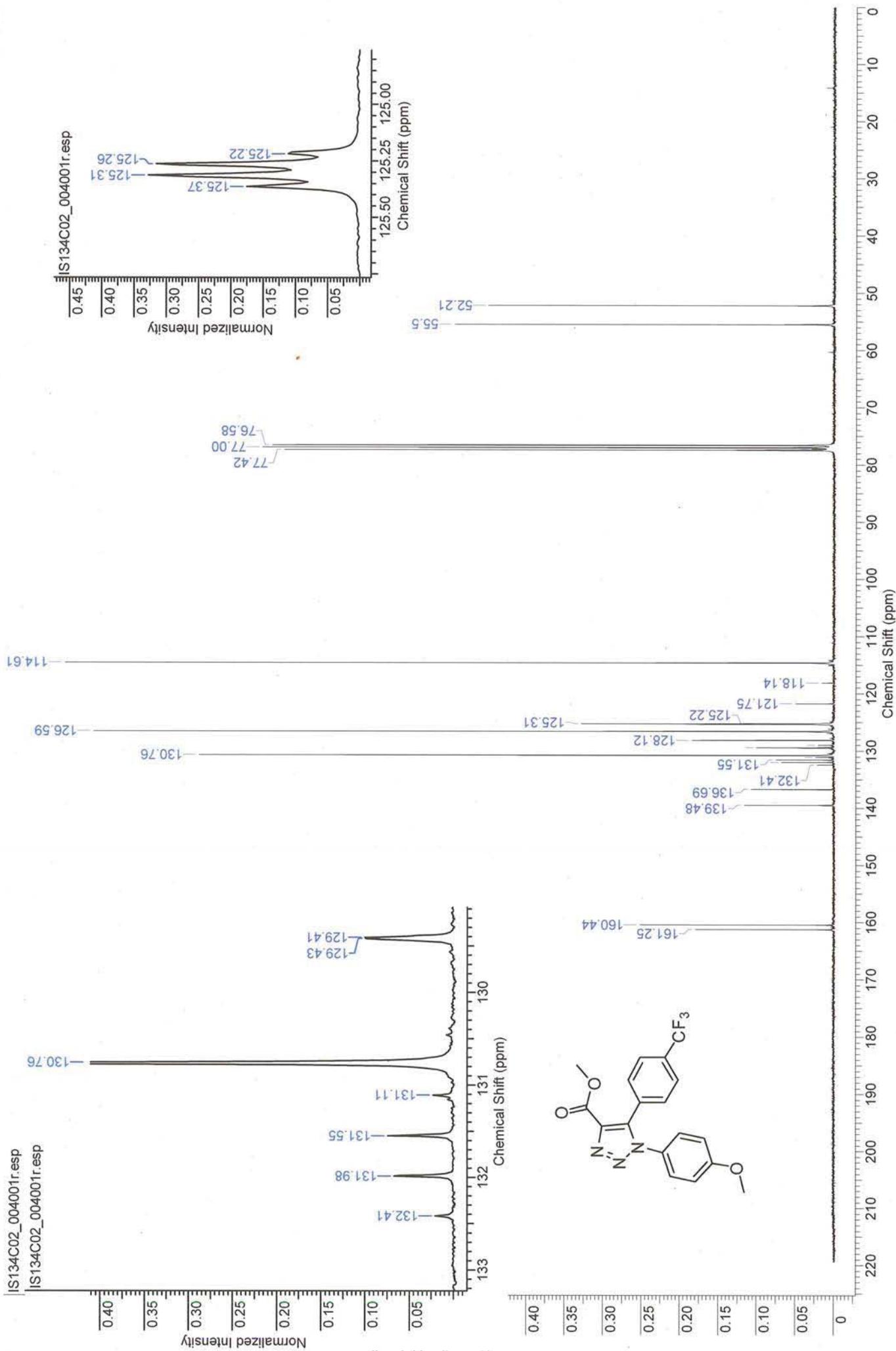
#	C	H	N	F	O	Na	mass	DBE	error
---	---	---	---	---	---	----	------	-----	-------

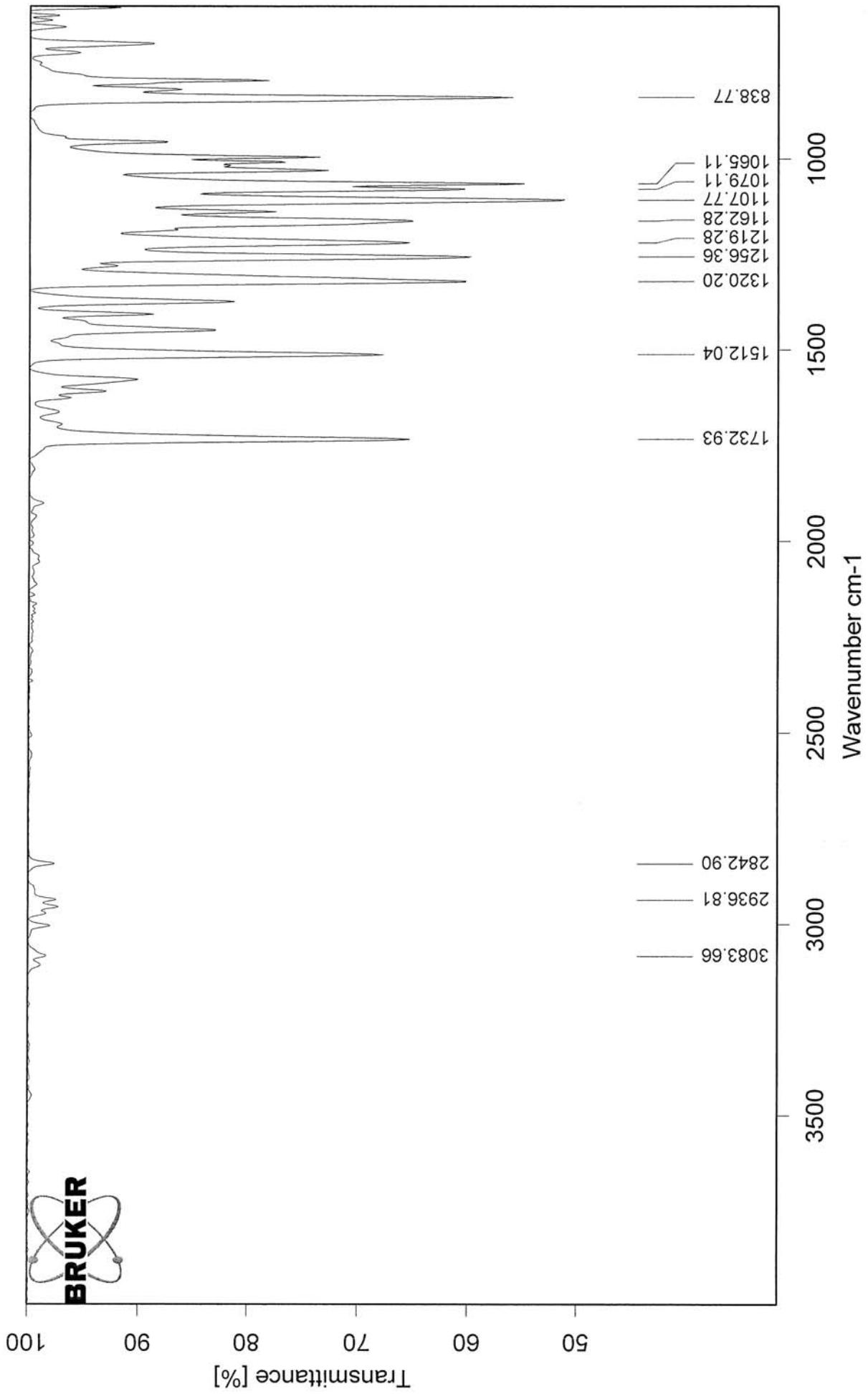
*** Mass Analysis for mass 378.1058990

1	18	15	3	3	3	0	378.1060024	11.5	1.034e-04
2	21	14	3	2	2	0	378.1048595	15.5	1.039e-03
3	13	17	3	4	4	1	378.1047399	4.5	1.159e-03
4	15	16	3	4	4	0	378.1071452	7.5	1.246e-03
5	16	15	4	4	1	1	378.1074200	9.0	1.521e-03
6	21	16	1	3	1	1	378.1076198	12.5	1.721e-03
7	24	13	3	1	1	0	378.1037167	19.5	2.182e-03
8	16	16	3	3	3	1	378.1035971	8.5	2.302e-03
9	16	14	3	6	1	0	378.1035577	8.5	2.341e-03
10	13	16	4	5	2	1	378.1085629	5.0	2.664e-03

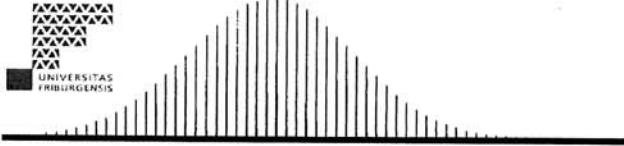








U:\IR_spectra\IS134C02.0	IS134C02	brown solid	17/12/2010
--------------------------	----------	-------------	------------



FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

ESI-MS: IS134C02

XMASS Mass Analysis for /Data/UNI_FR/ALIM5263_ESI/5/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

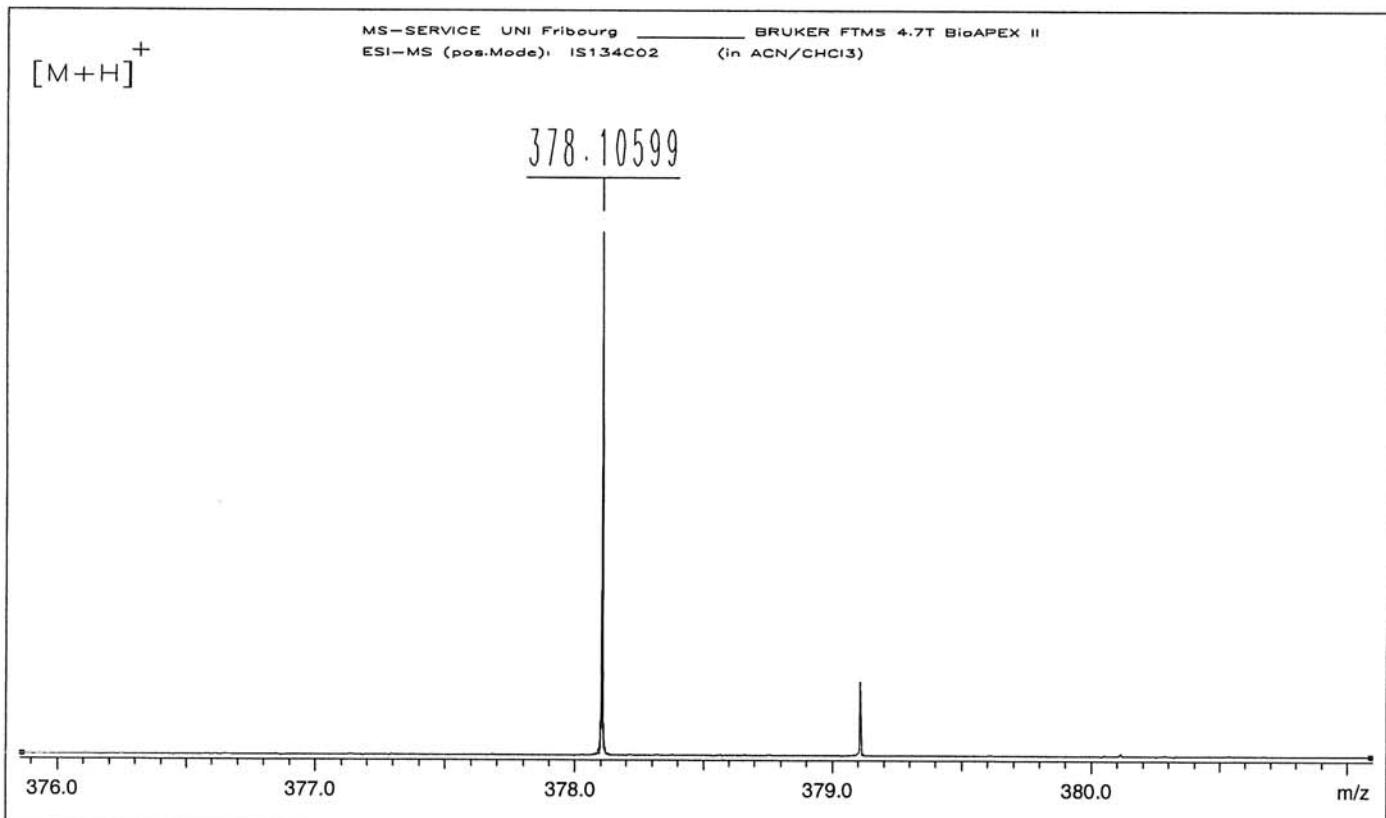
Ion mass = 378.1059890

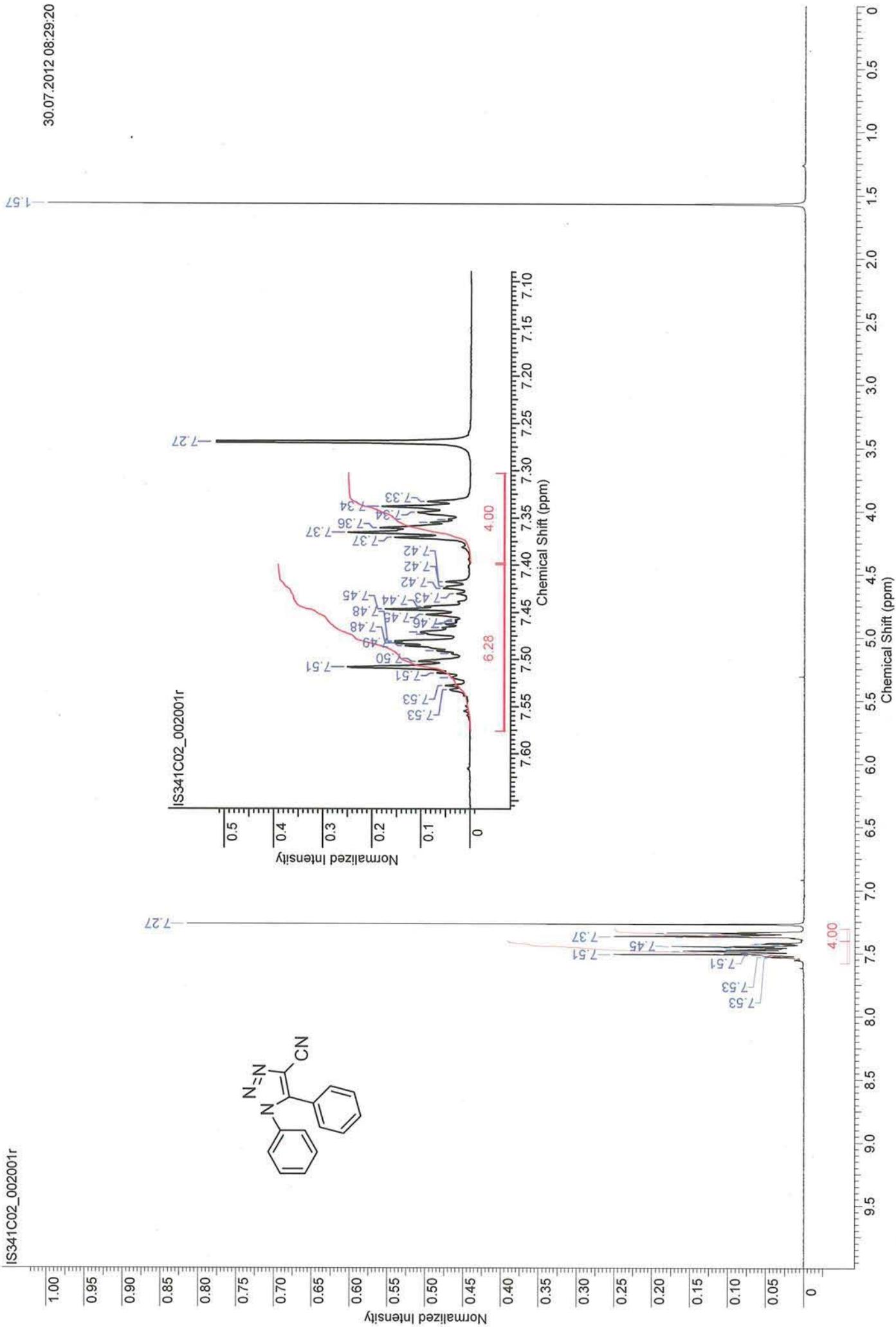
Charge = +1

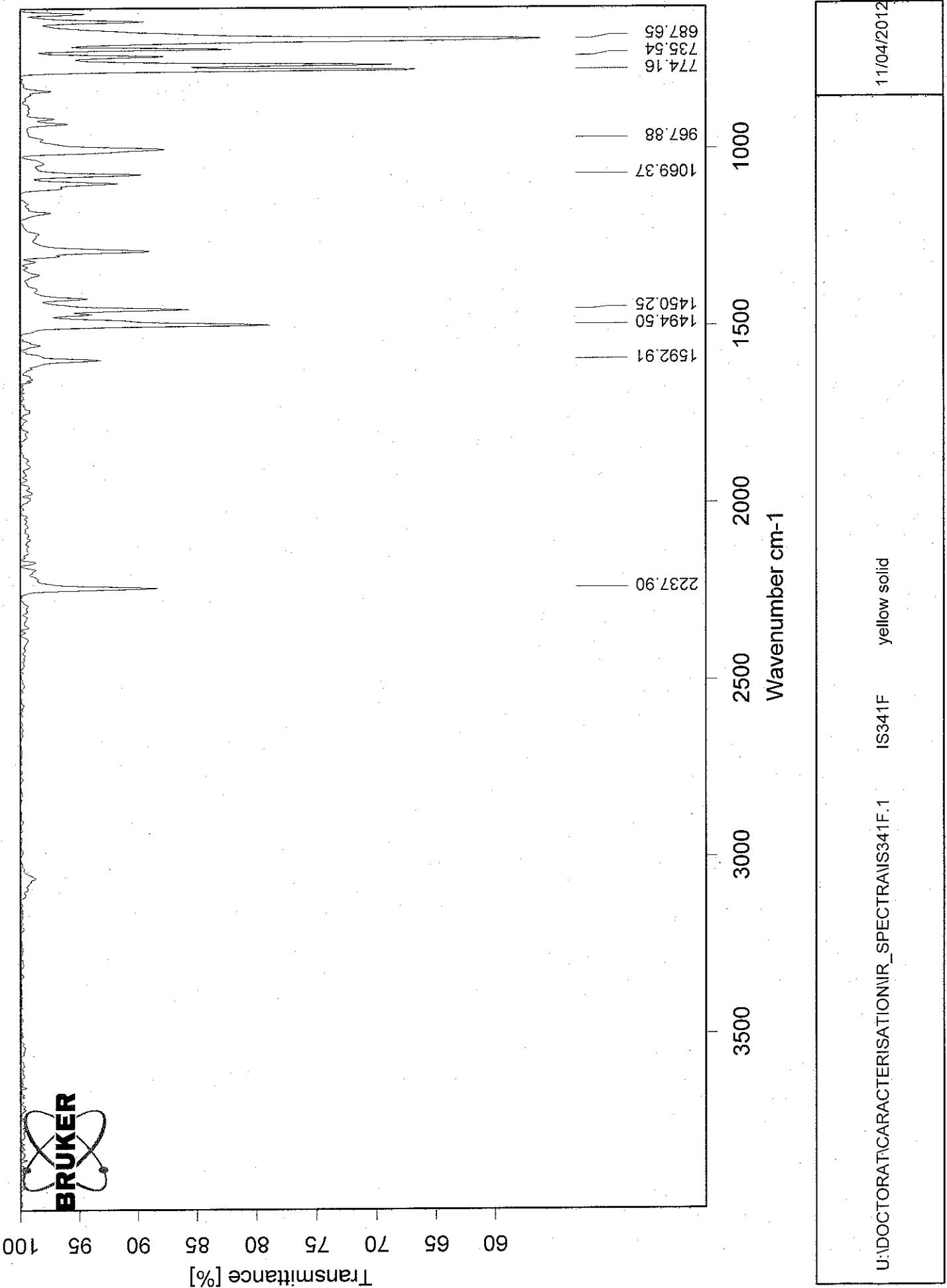
#	C	H	N	F	O	Na	mass	DBE	error
---	---	---	---	---	---	----	------	-----	-------

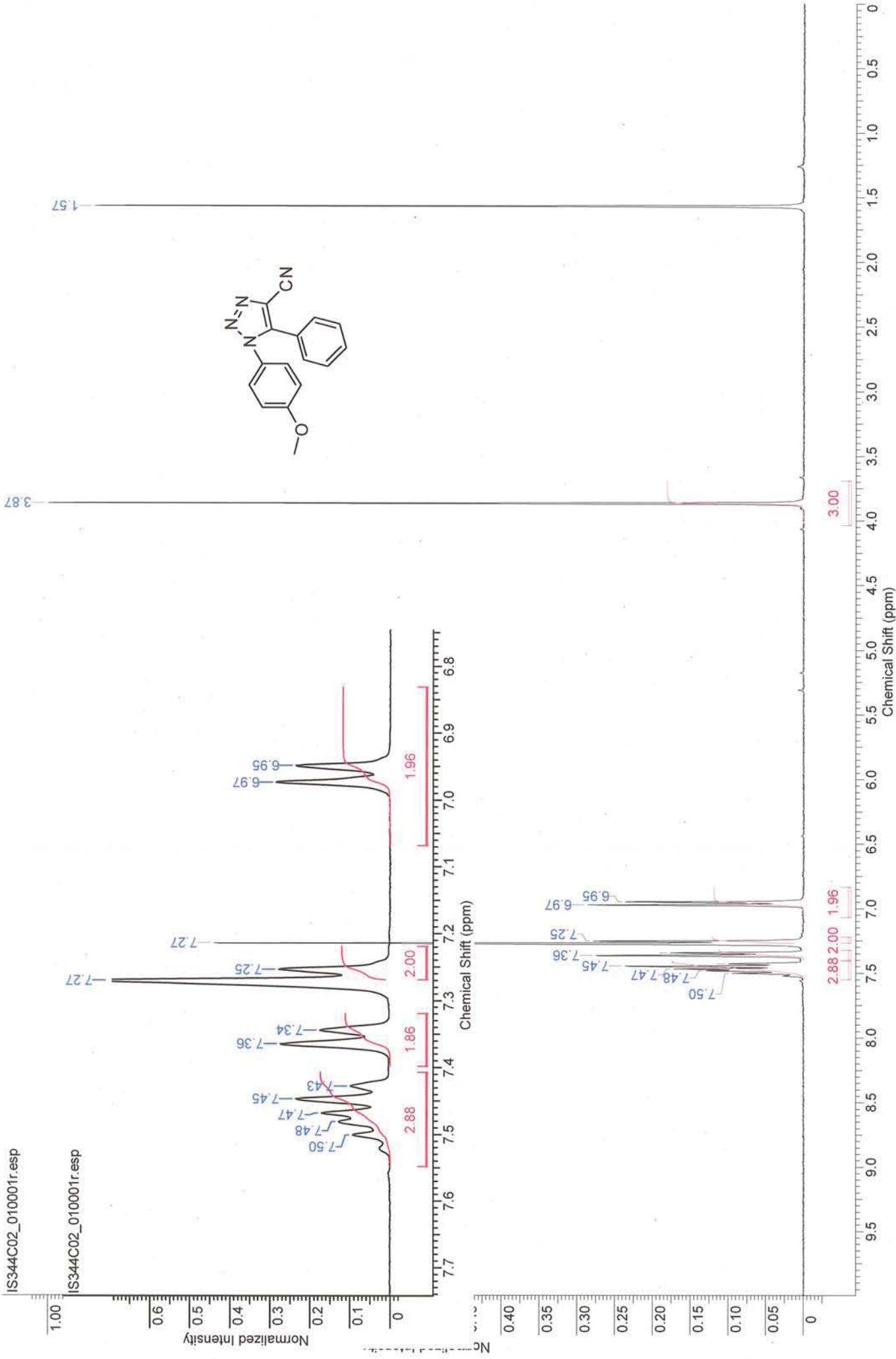
*** Mass Analysis for mass 378.1059890

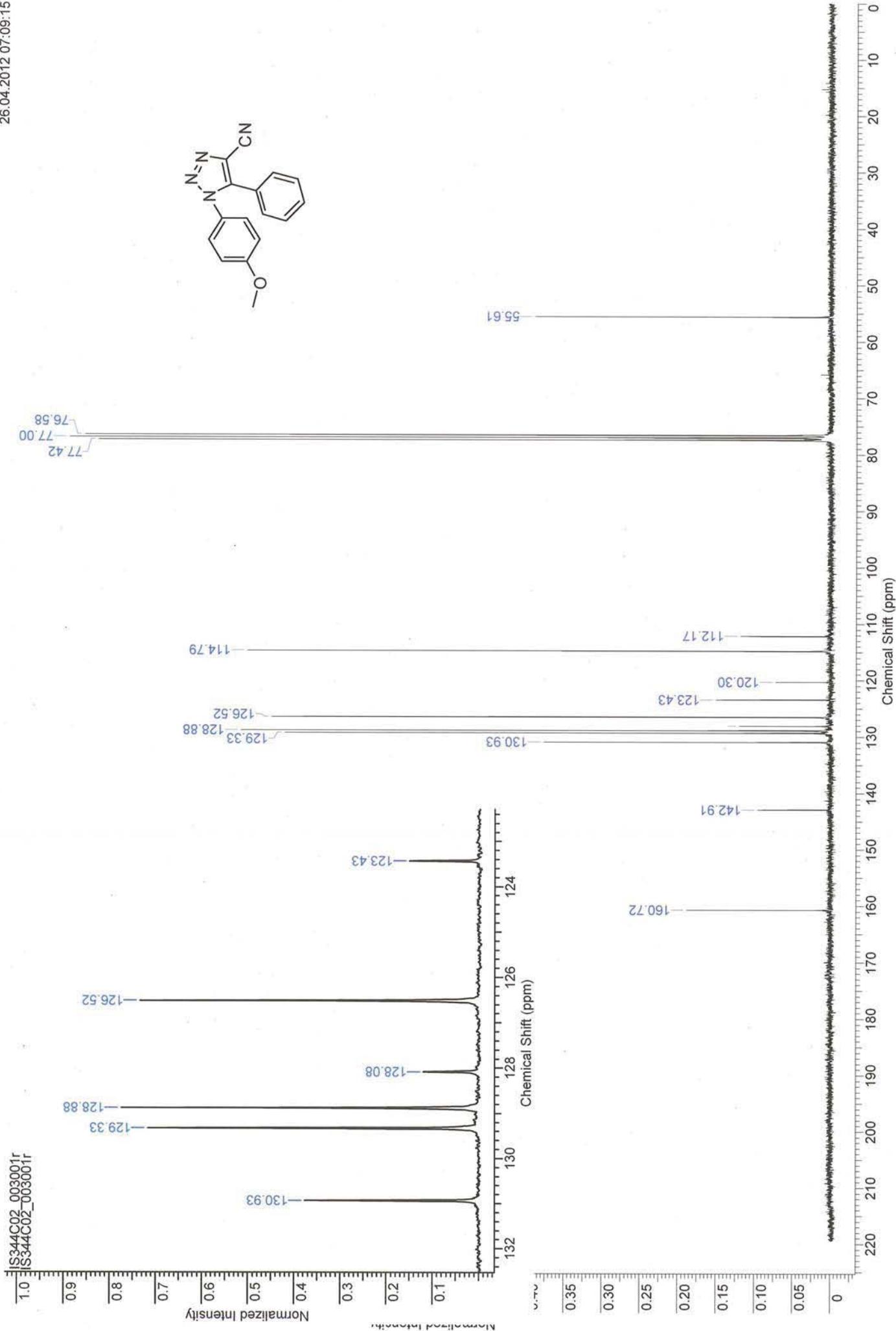
1	18	15	3	3	3	0	378.1060024	11.5	1.338e-05
2	21	14	3	2	2	0	378.1048595	15.5	1.129e-03
3	15	16	3	4	4	0	378.1071452	7.5	1.156e-03
4	13	17	3	4	4	1	378.1047399	4.5	1.249e-03
5	16	15	4	4	1	1	378.1074200	9.0	1.431e-03
6	21	16	1	3	1	1	378.1076198	12.5	1.631e-03
7	24	13	3	1	1	0	378.1037167	19.5	2.272e-03
8	16	16	3	3	3	1	378.1035971	8.5	2.392e-03
9	16	14	3	6	1	0	378.1035577	8.5	2.431e-03
10	13	16	4	5	2	1	378.1085629	5.0	2.574e-03



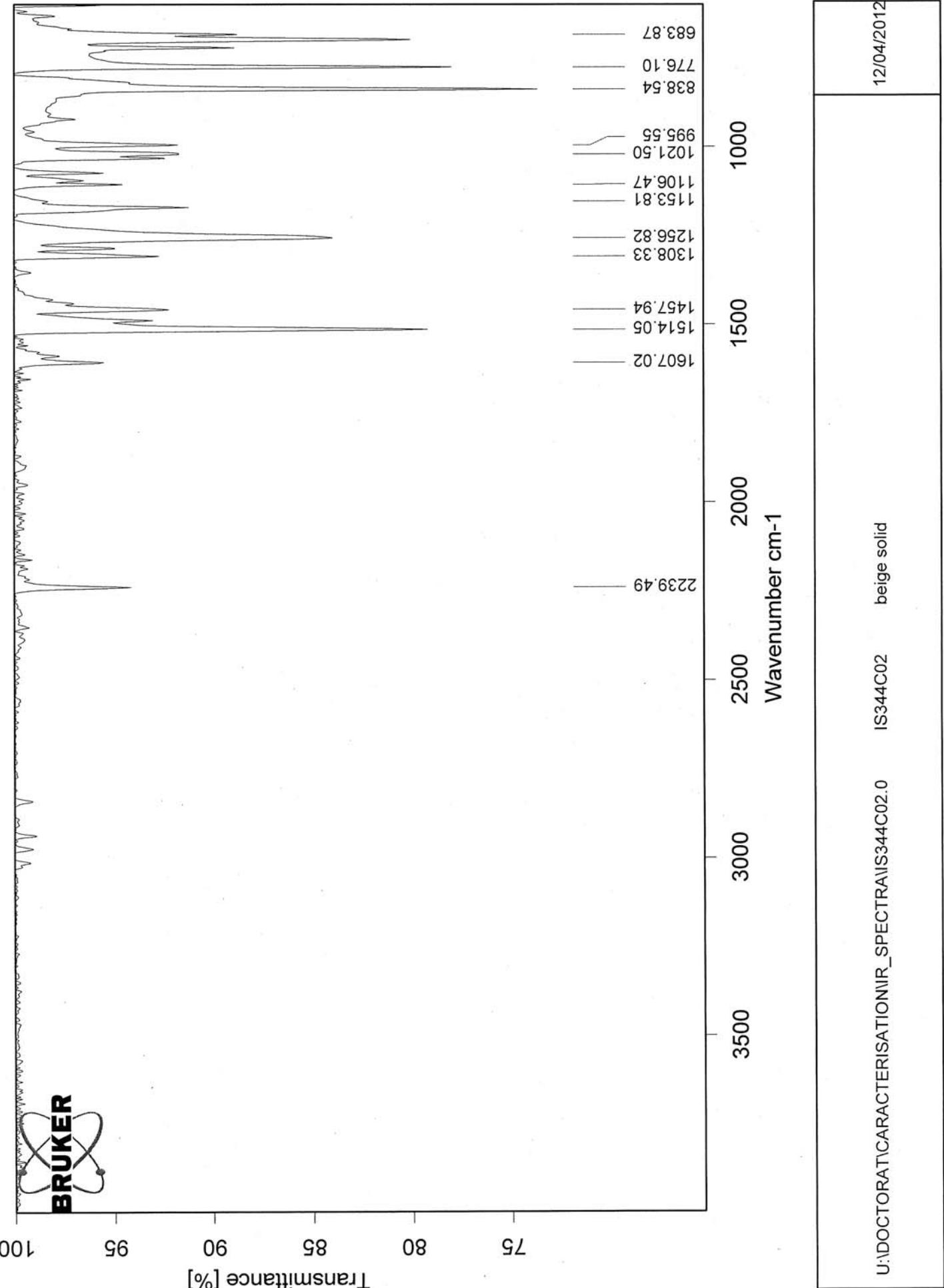


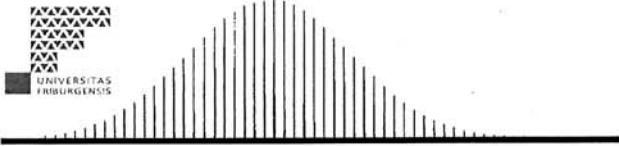






S344C02_003001
S344C02_003001





FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

ESI-MS: IS344CO2

XMASS Mass Analysis for /Data/UNI_FR/ALIM6523_ESI/2/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

Ion mass = 277.1086490

Charge = +1

#	C	H	N	O	mass	DBE	error
---	---	---	---	---	------	-----	-------

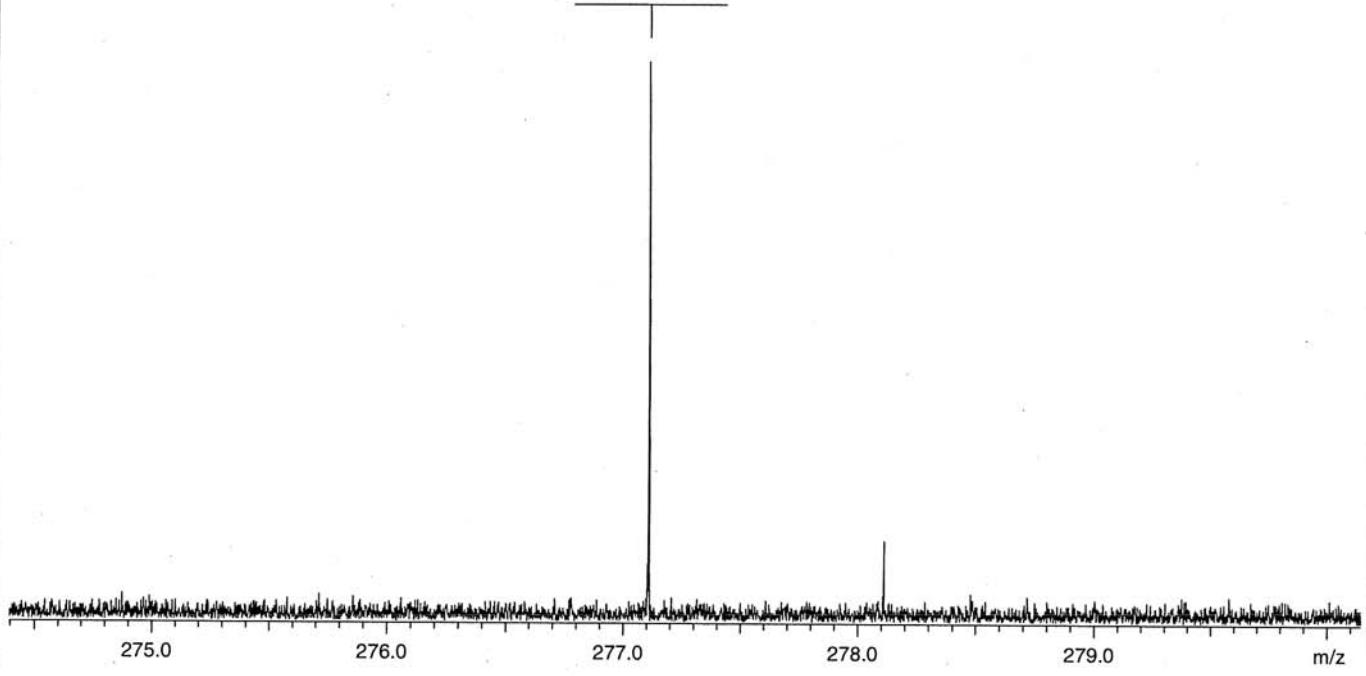
*** Mass Analysis for mass 277.1086490

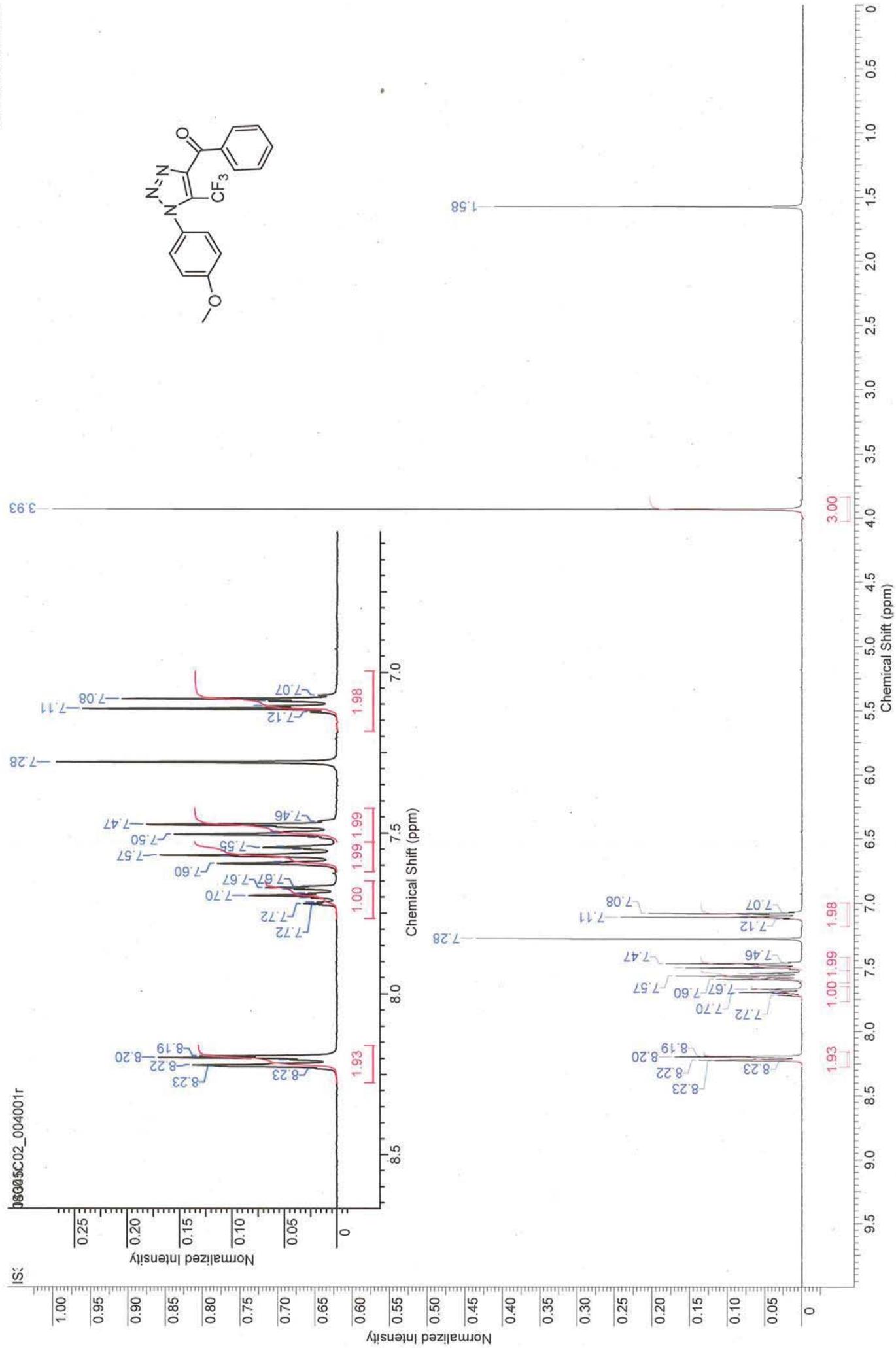
1	16	13	4	1	277.1083875	12.5	2.615e-04
2	18	15	1	2	277.1097302	12.0	1.081e-03
3	5	13	10	4	277.1115754	4.5	2.926e-03
4	6	19	3	9	277.1115806	-1.0	2.932e-03
5	13	15	3	4	277.1057074	8.0	2.942e-03
6	7	15	7	5	277.1129180	4.0	4.269e-03
7	11	13	6	3	277.1043647	8.5	4.284e-03
8	9	17	4	6	277.1142607	3.5	5.612e-03
9	10	17	2	7	277.1030273	3.5	5.622e-03
10	9	11	9	2	277.1030221	9.0	5.627e-03

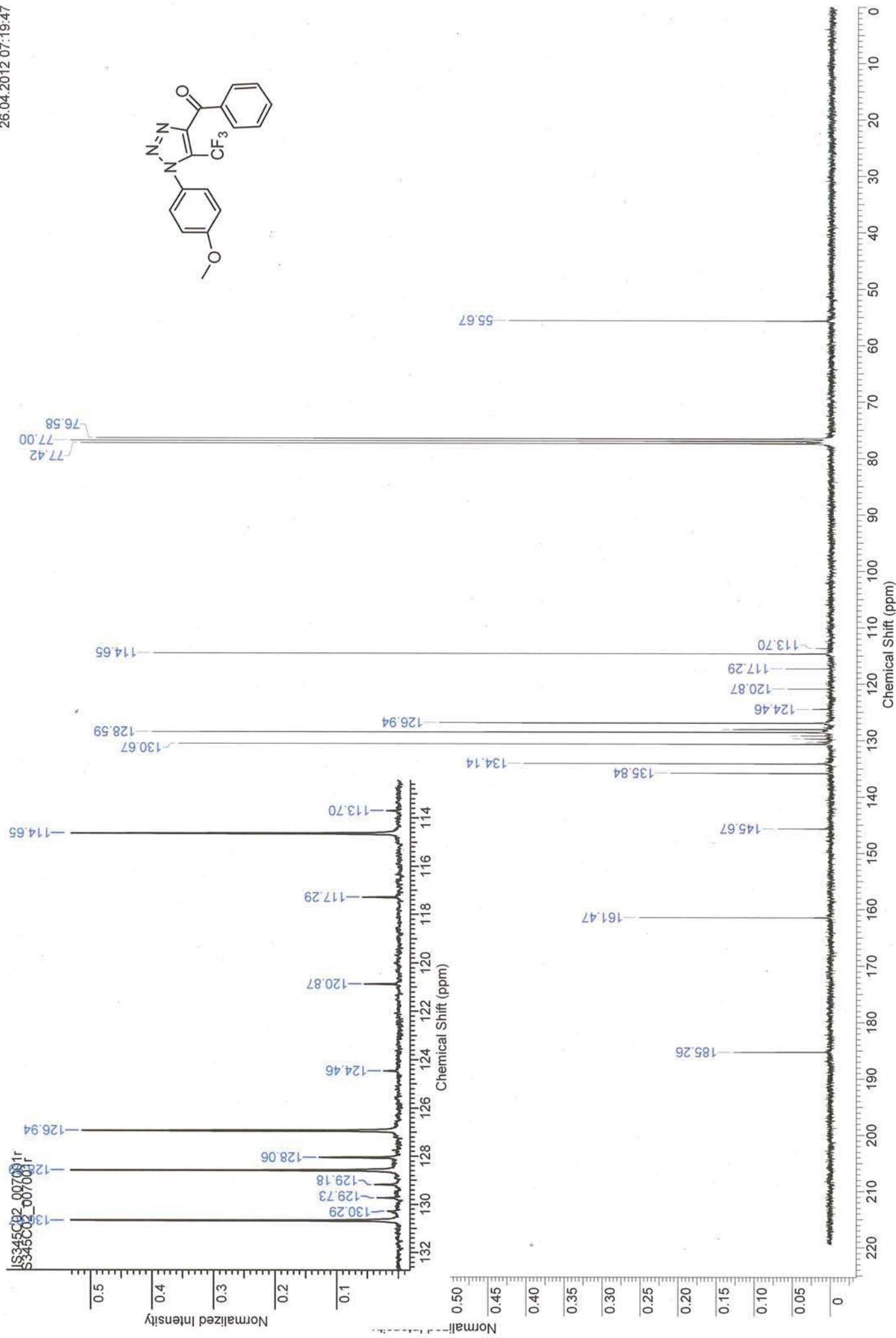
MS-SERVICE UNI Fribourg BRUKER FTMS 4.7T BioAPEX II
ESI-MS (pos.Mode): IS344CO2 (in ACN)

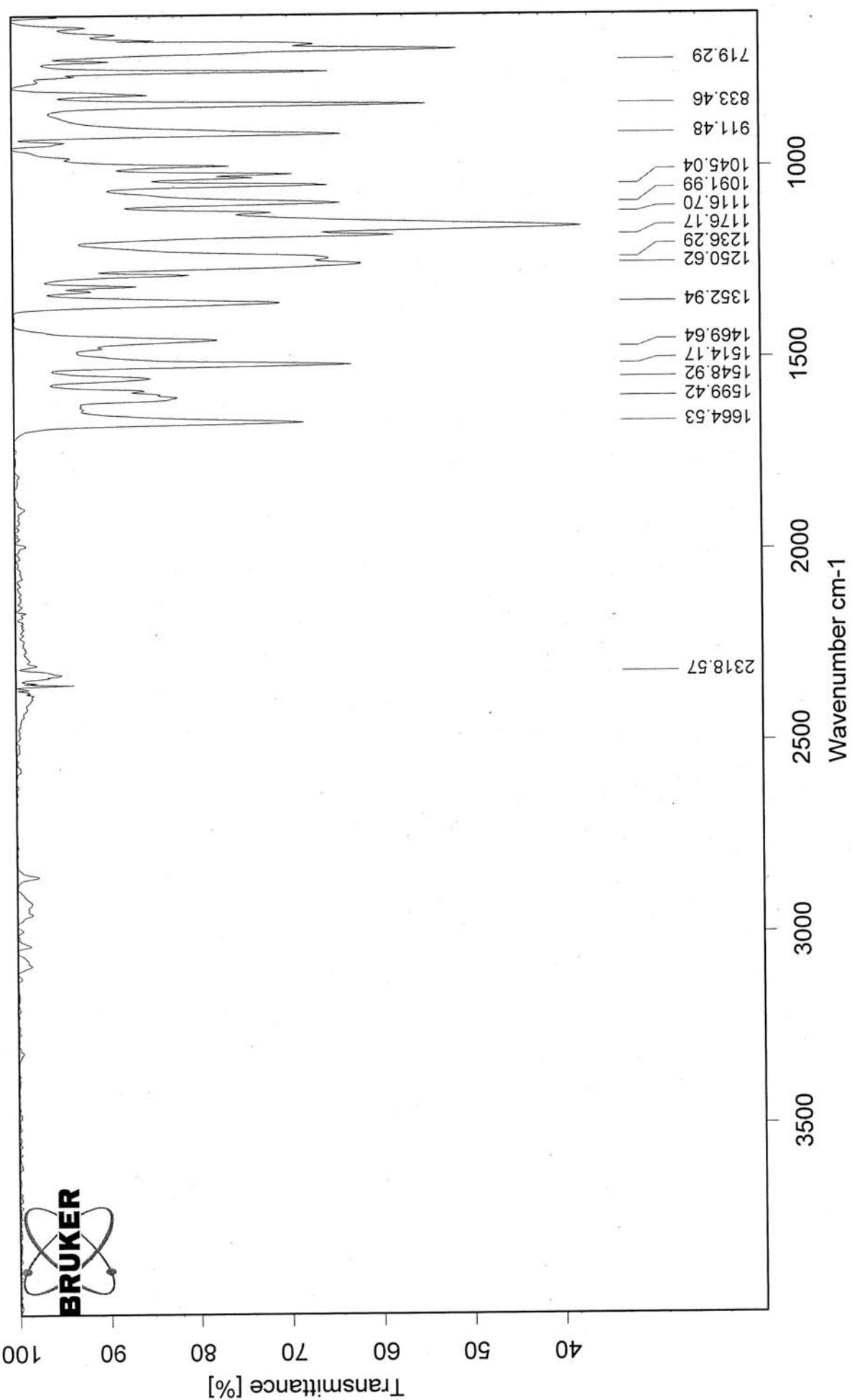
[M+H]⁺

277.10865

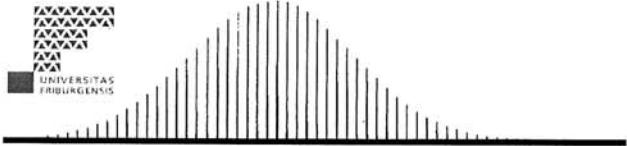








U:\DOCTORAT\CHARACTERISATION\IR_SPECTRA\IS345C02.0	IS345C02	yellow solid	11/04/2012
--	----------	--------------	------------



FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

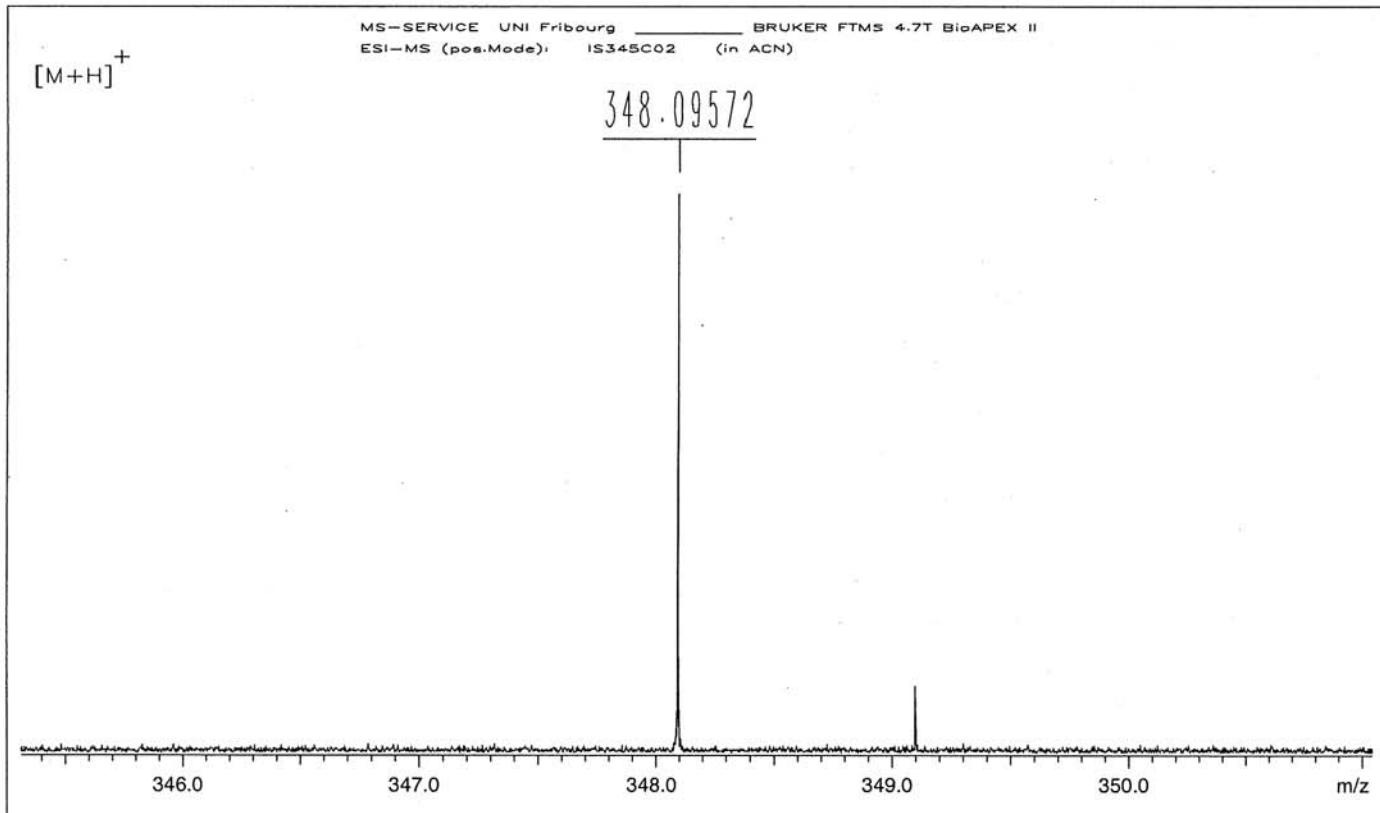
ESI-MS: IS345CO2

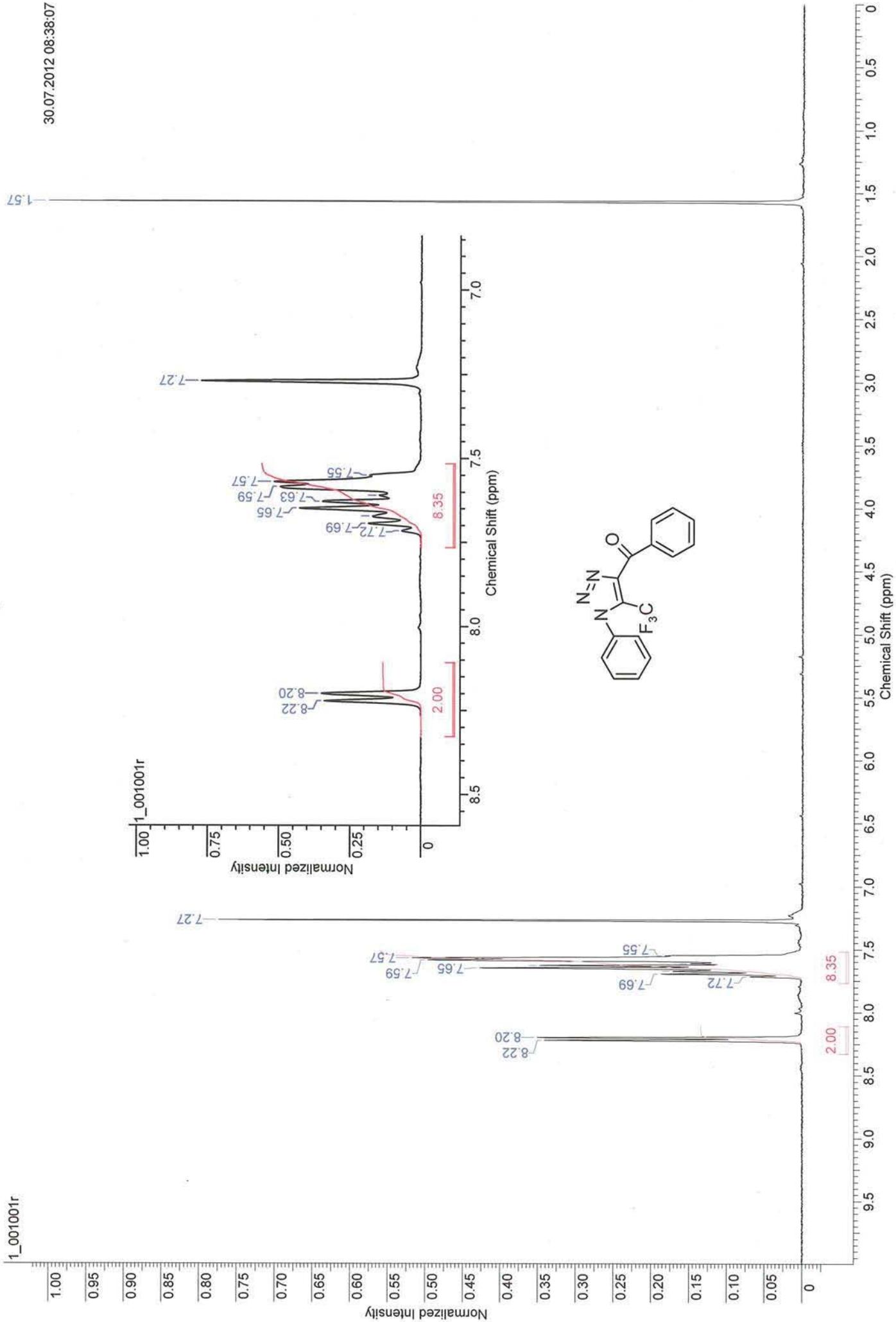
XMASS Mass Analysis for /Data/UNI_FR/ALIM6524_ESI/2/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

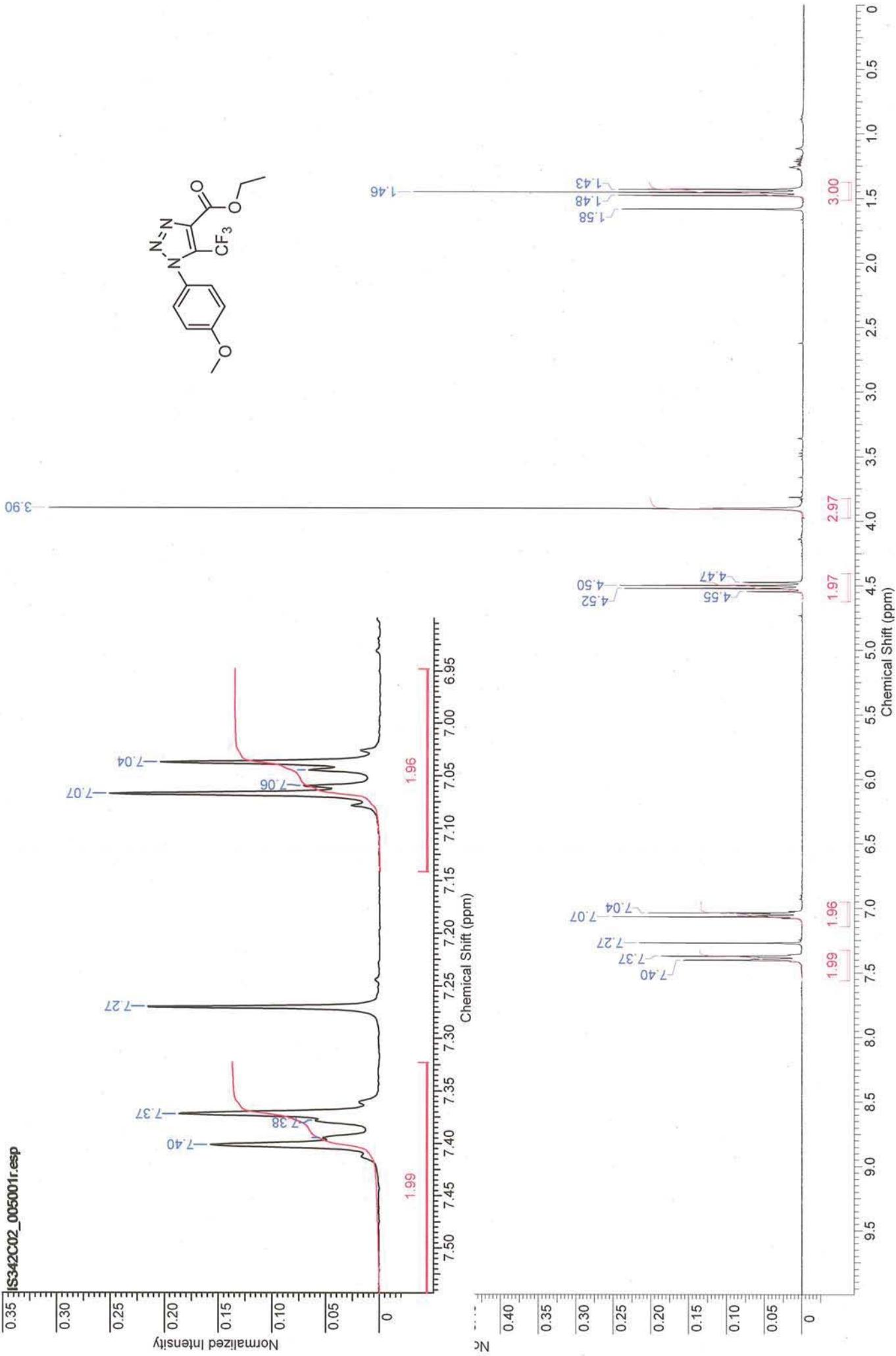
Ion mass = 348.0957160

Charge = +1

#	C	H	F	N	O	mass	DBE	error
*** Mass Analysis for mass 348.0957160								
1	17	13	3	3	2	348.0954377	11.5	2.783e-04
2	8	16	2	5	8	348.0961453	2.5	4.293e-04
3	12	12	4	6	2	348.0952379	8.0	4.781e-04
4	12	11	1	9	3	348.0963399	11.5	6.239e-04
5	13	17	1	2	8	348.0963452	6.0	6.292e-04
6	9	13	5	6	3	348.0963807	4.0	6.647e-04
7	8	17	5	2	7	348.0950433	-1.0	6.727e-04
8	7	11	5	9	2	348.0950381	4.5	6.779e-04
9	11	15	1	5	7	348.0950025	6.5	7.135e-04
10	14	14	4	3	3	348.0965806	7.5	8.646e-04

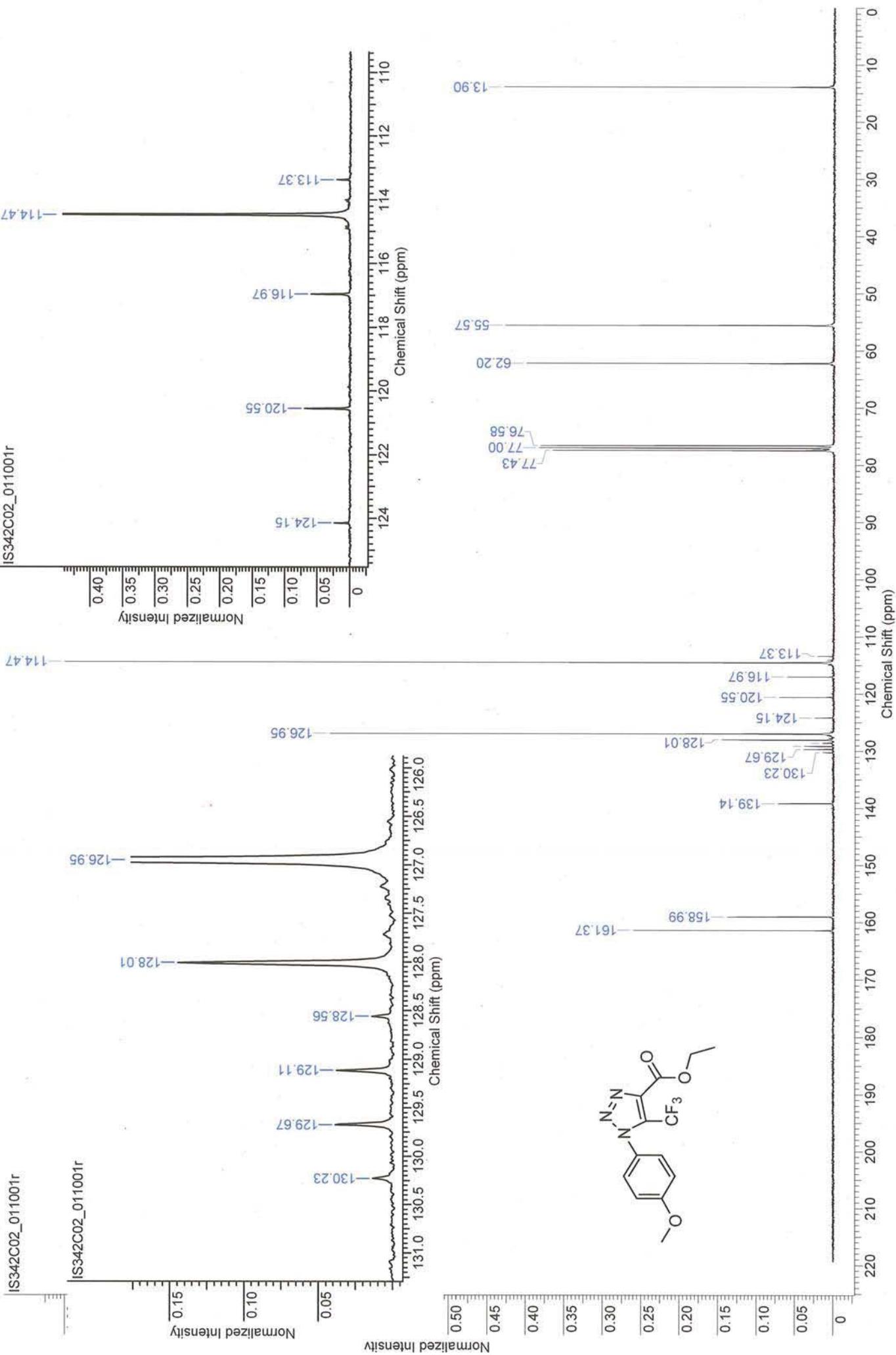


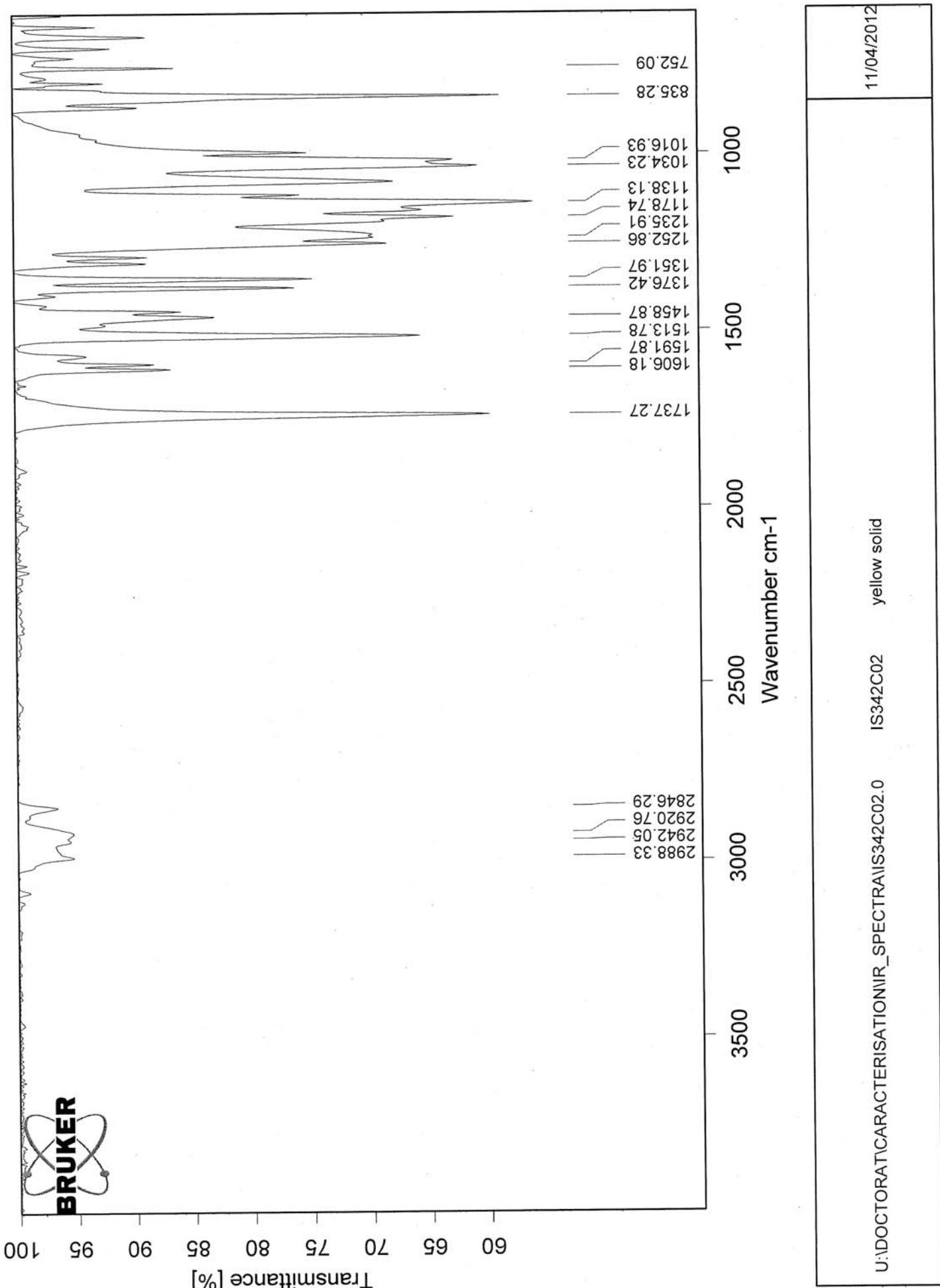


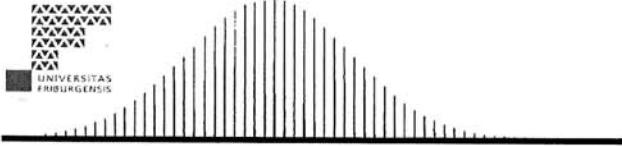


IS342C02_011001r

IS342C02_011001r







BRUKER FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

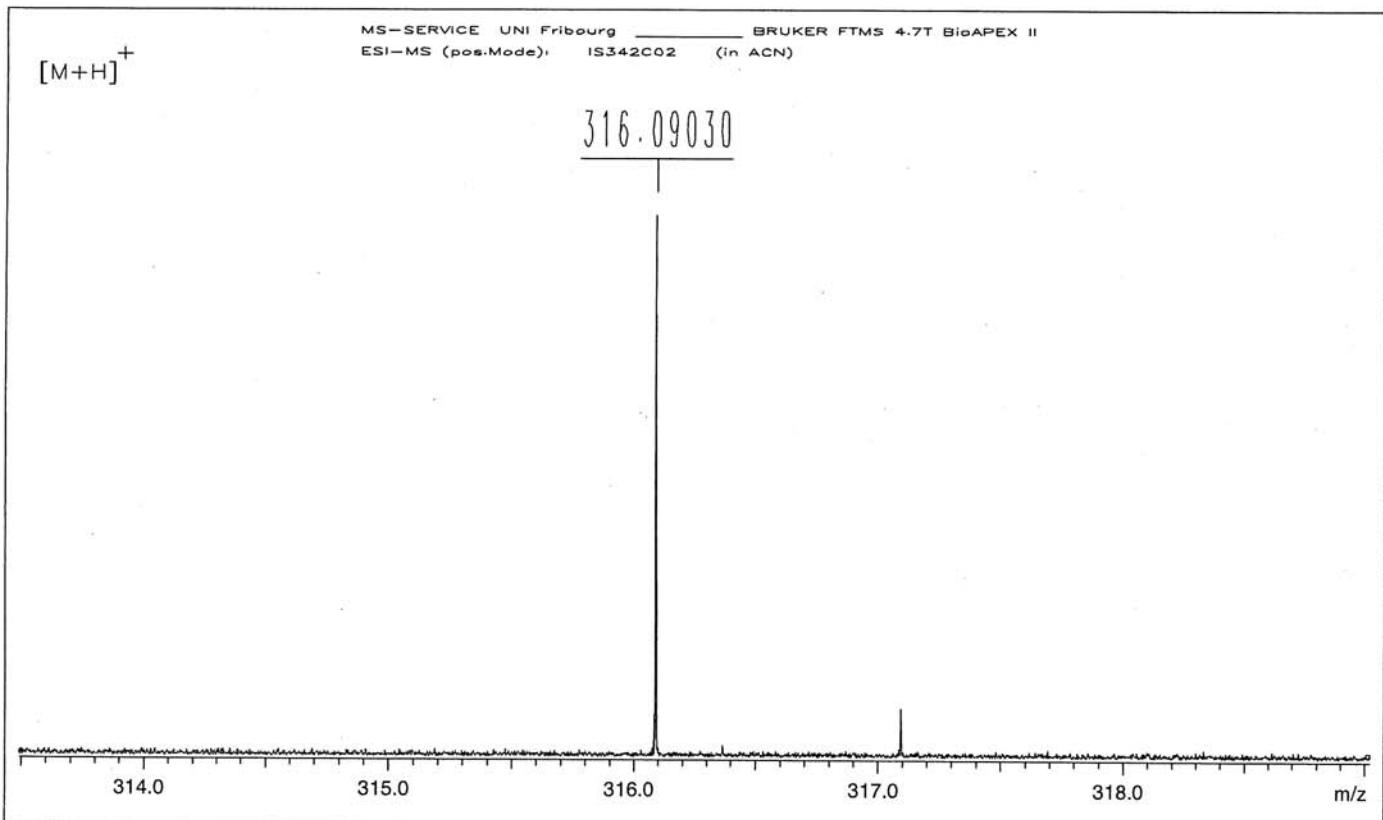
ESI-MS: IS342CO2

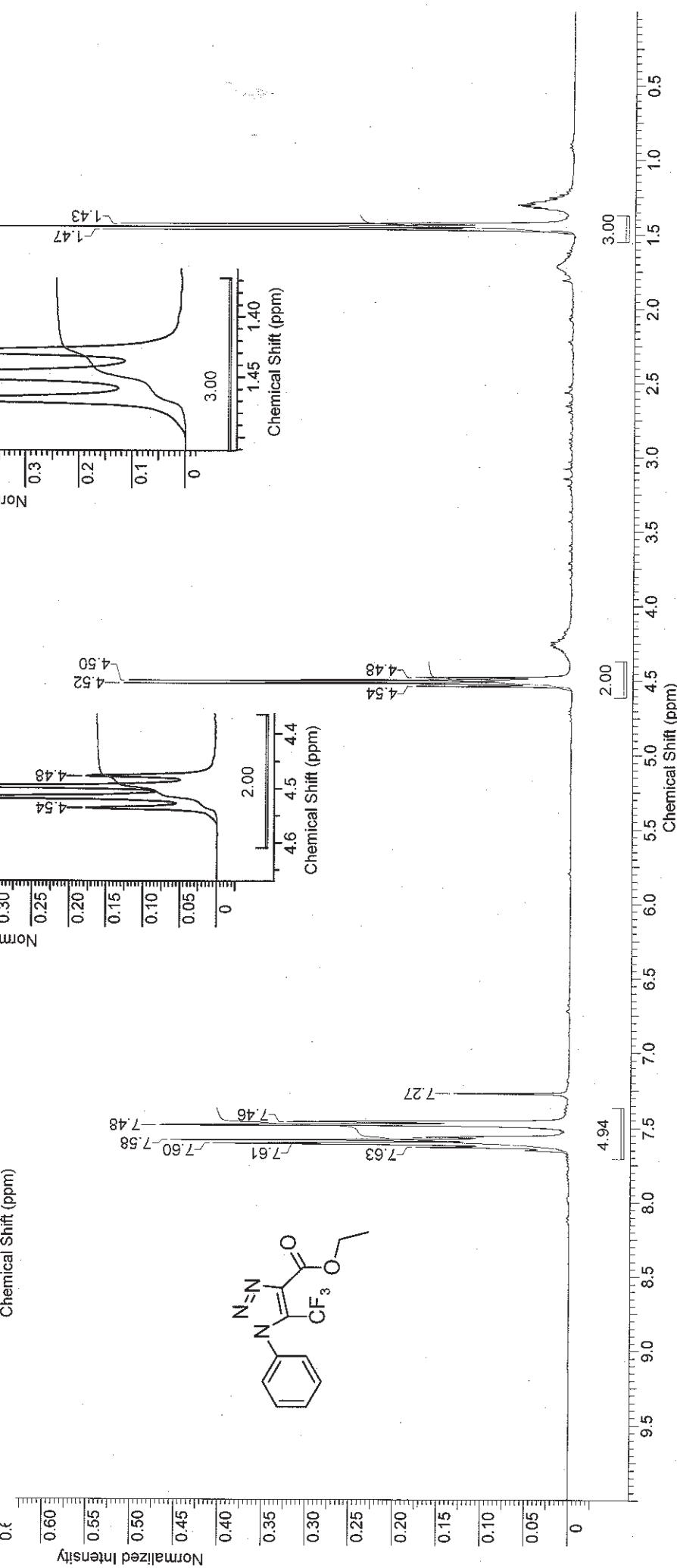
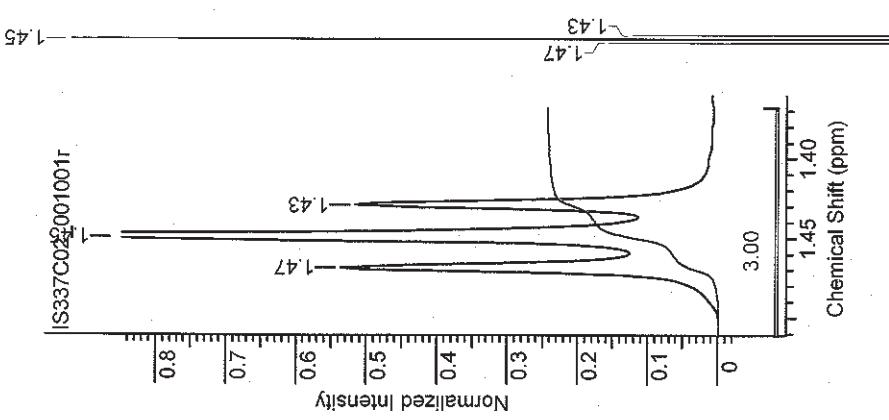
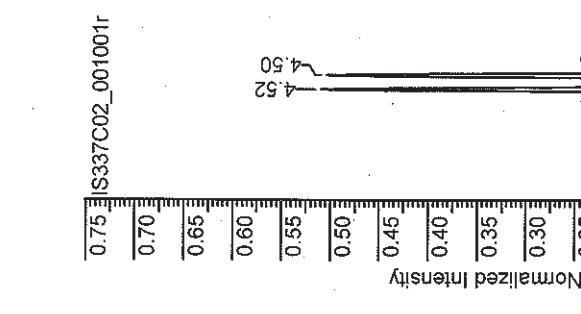
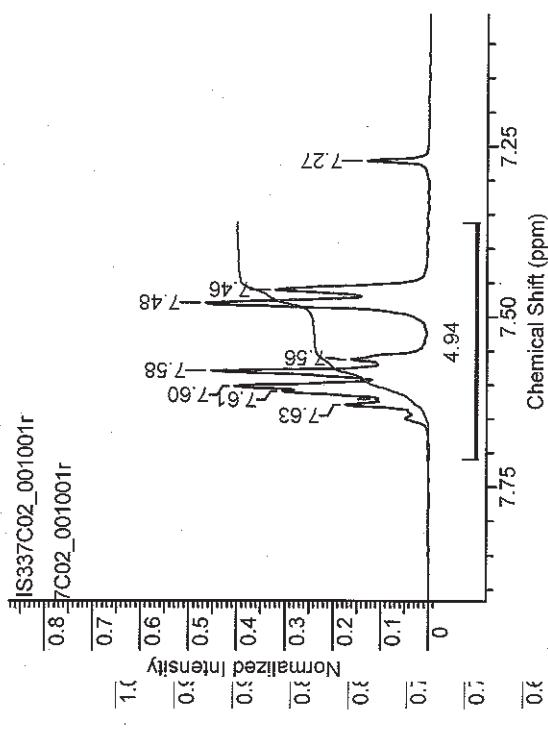
XMASS Mass Analysis for /Data/UNI_FR/ALIM6522_ESI/2/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

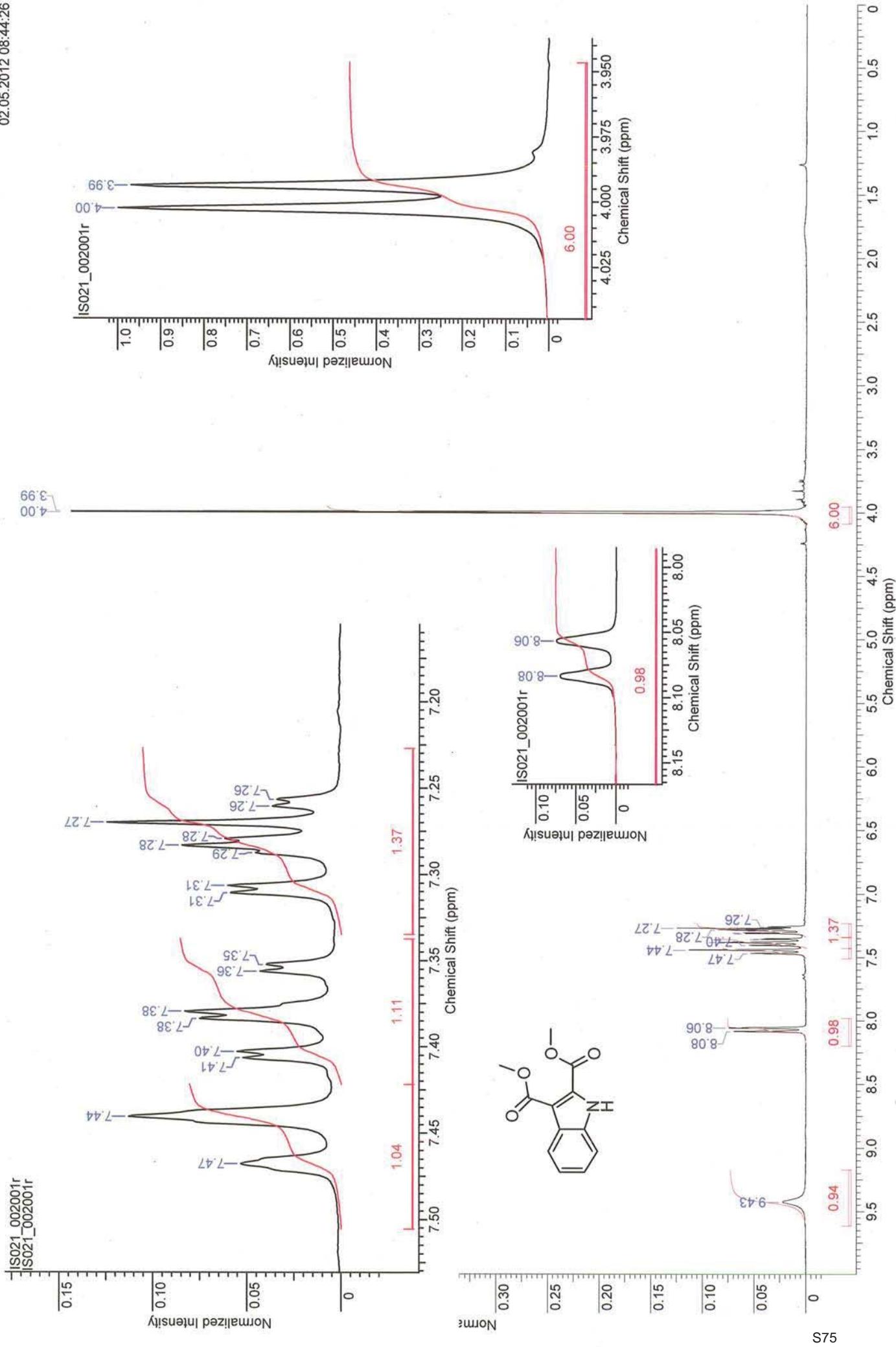
Ion mass = 316.0902960

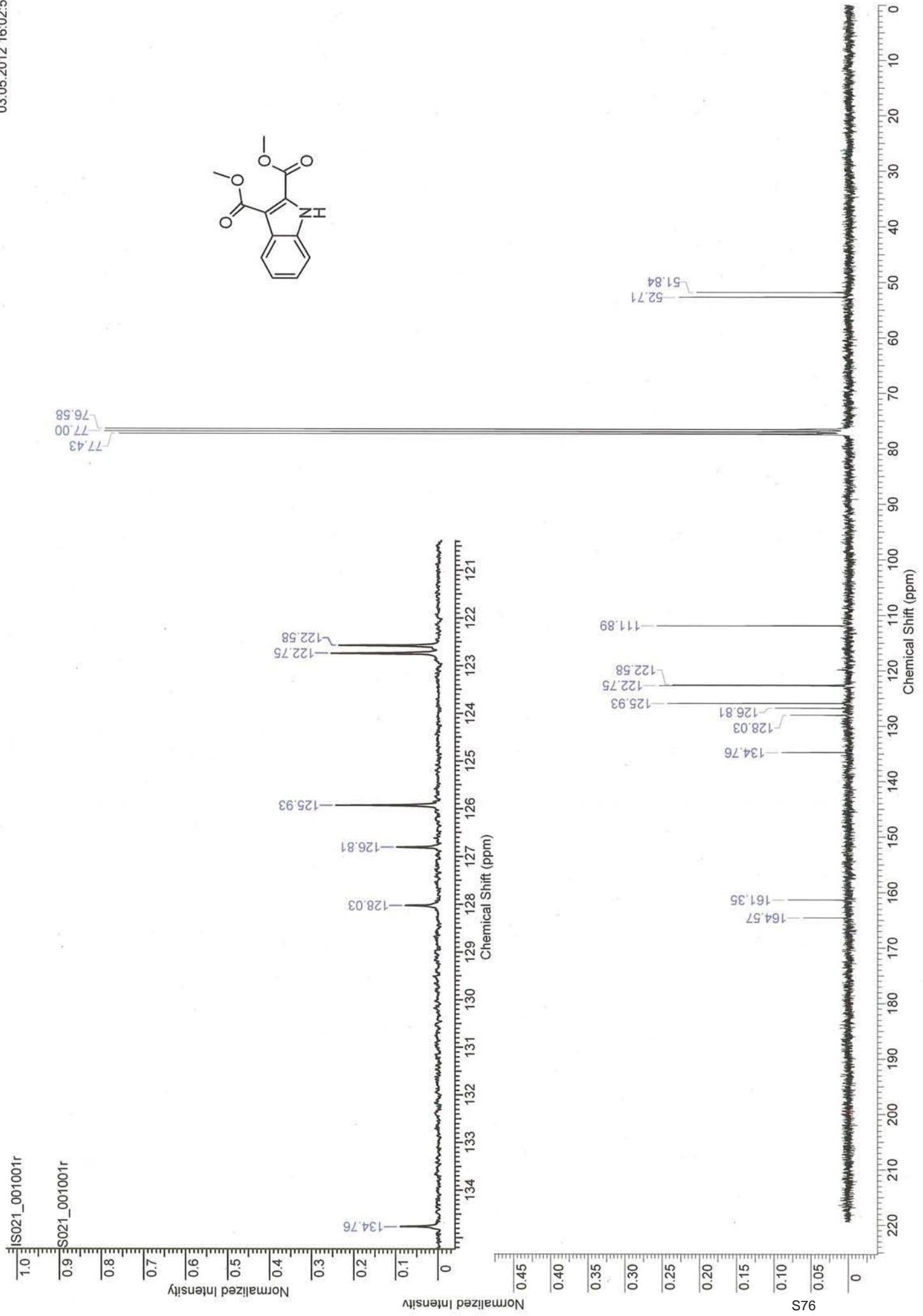
Charge = +1

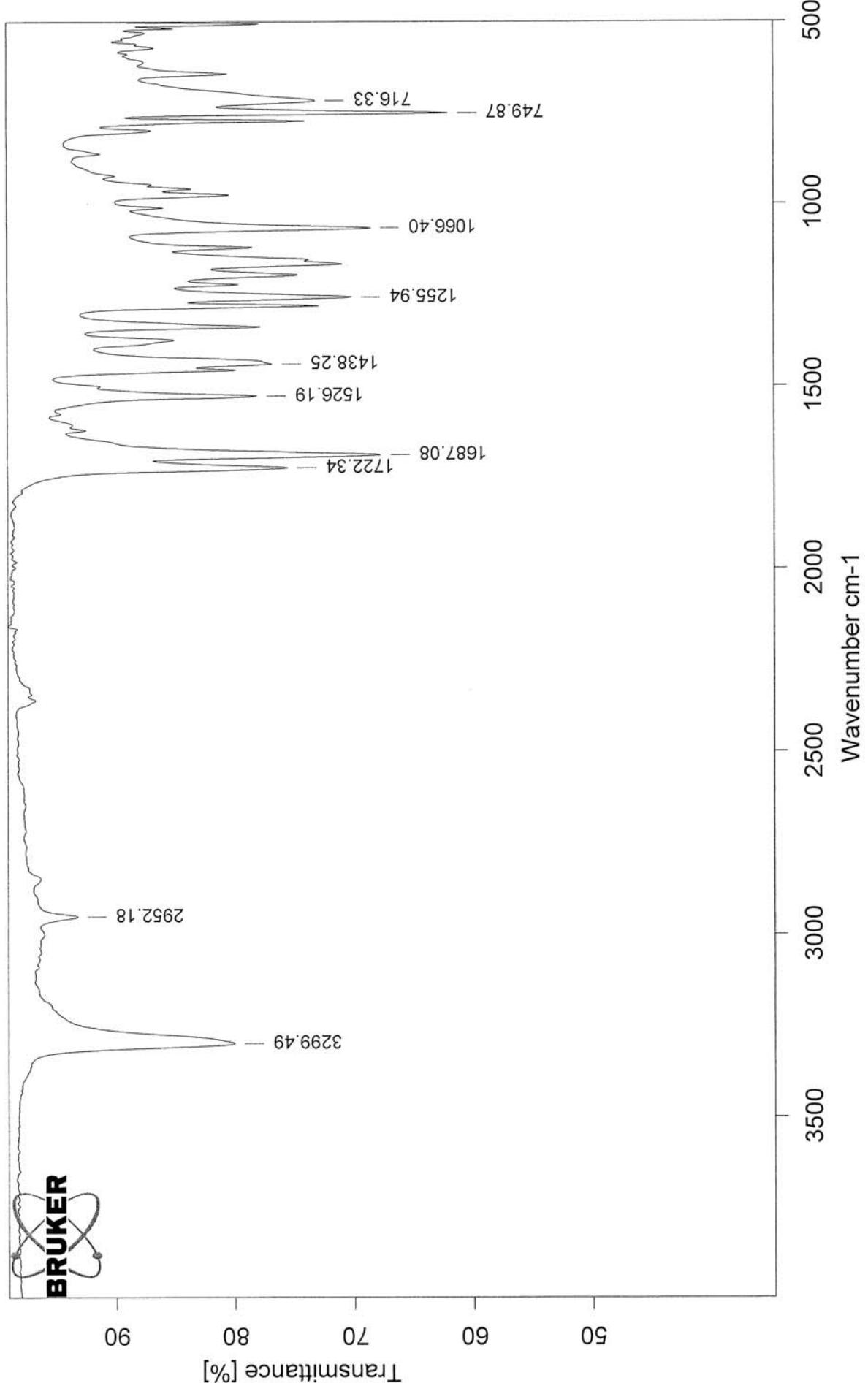
#	C	H	F	N	O	mass	DBE	error
*** Mass Analysis for mass 316.0902960								
1	13	13	3	3	3	316.0903523	7.5	5.632e-05
2	8	12	4	6	3	316.0901525	4.0	1.435e-04
3	7	15	1	5	8	316.0899171	2.5	3.789e-04
4	8	11	1	9	4	316.0912545	7.5	9.585e-04
5	9	17	1	2	9	316.0912598	2.0	9.638e-04
6	5	13	5	6	4	316.0912954	0.0	9.994e-04
7	16	12	2	3	2	316.0892095	11.5	1.087e-03
8	10	14	4	3	4	316.0914952	3.5	1.199e-03
9	11	11	3	6	2	316.0890096	8.0	1.286e-03
10	7	16	4	2	7	316.0888151	-1.0	1.481e-03





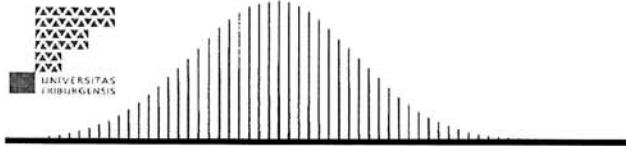






C:\PROGRAM FILES\OPUS\DATA\lsakis-021.0 is-021 is-021

26/01/2010



BRUKER FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

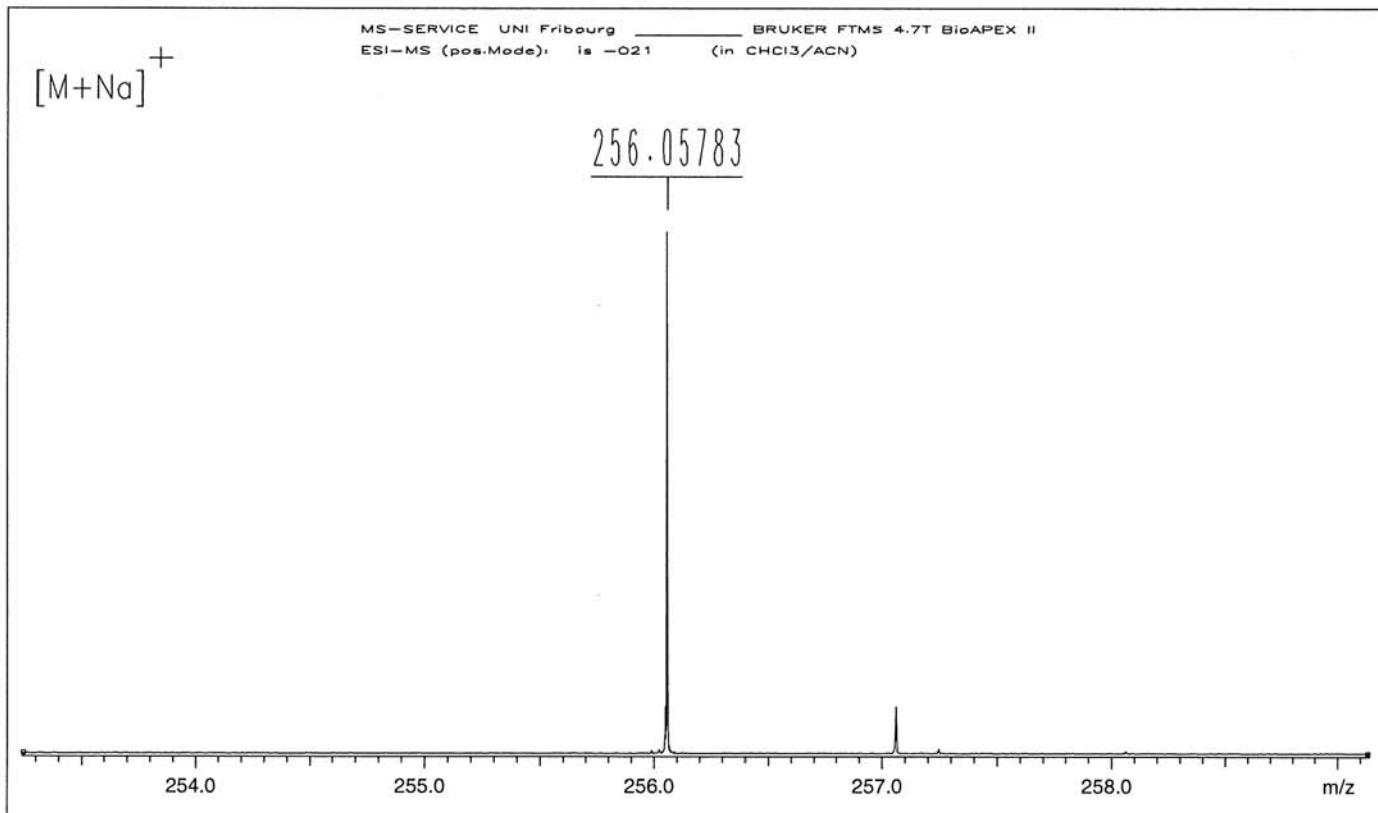
ESI-MS: is-021

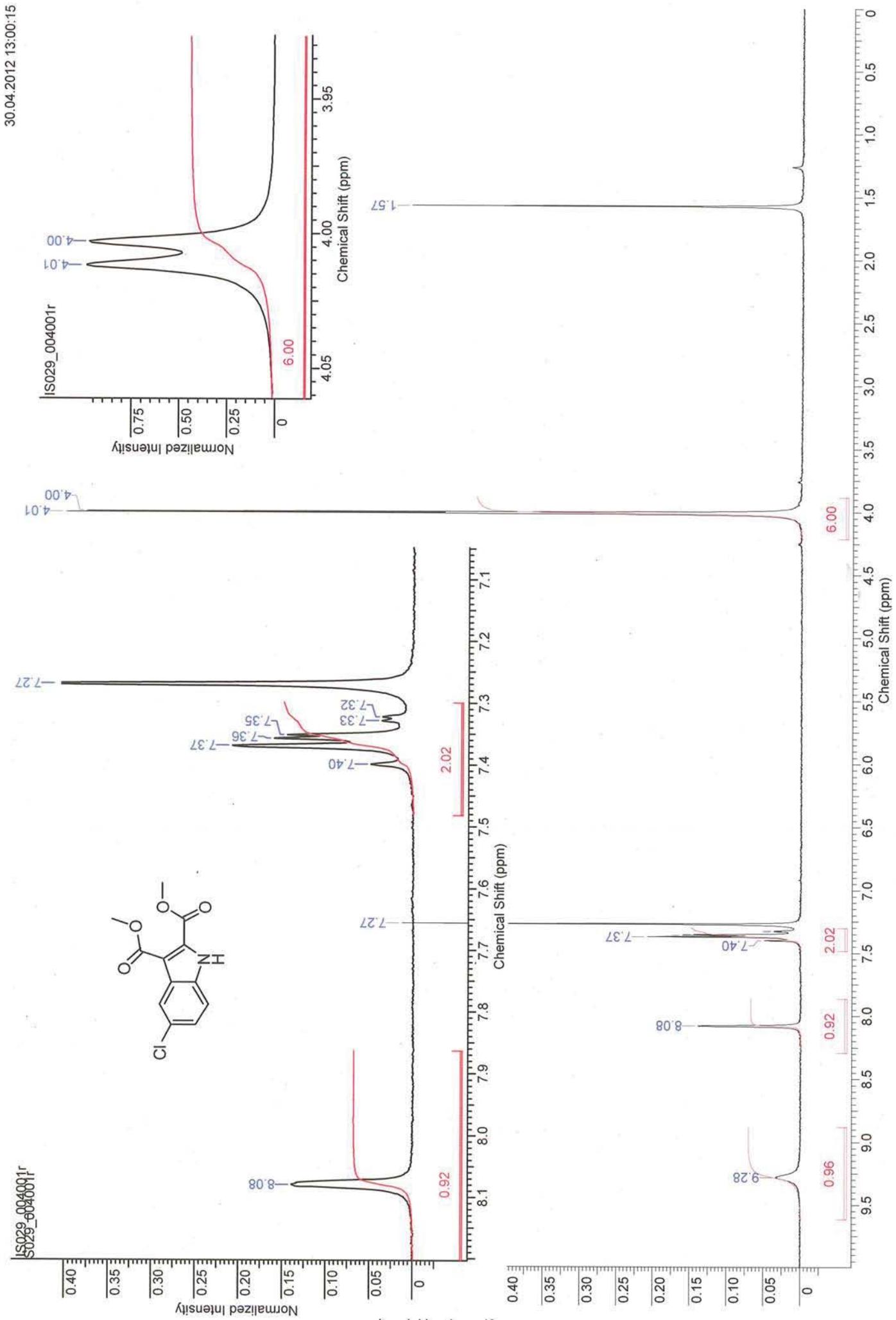
XMASS Mass Analysis for /Data/UNI_FR/ALIM4616_ESI/2/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

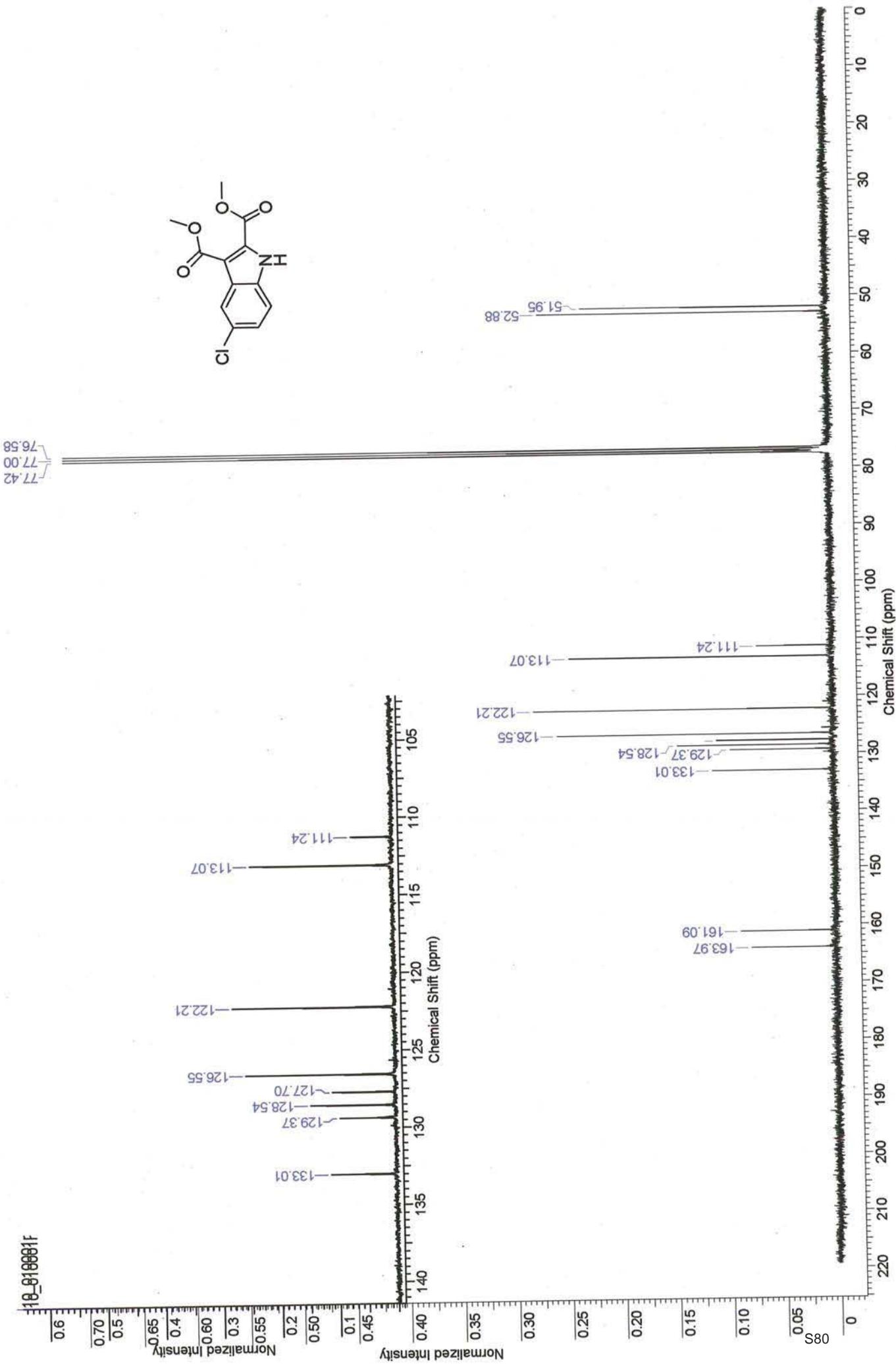
Ion mass = 256.0578260

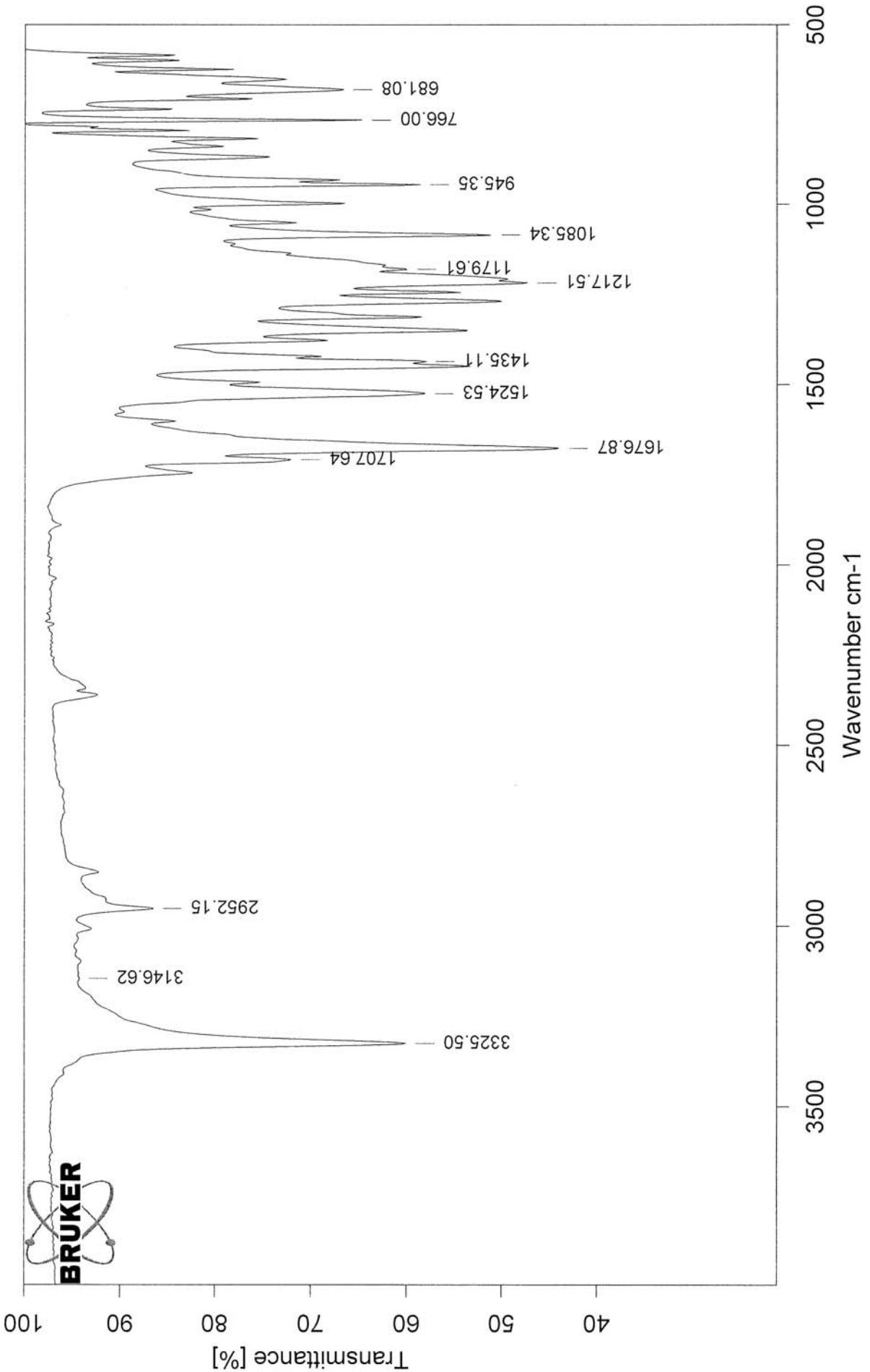
Charge = +1

#	C	H	N	O	Na	mass	DBE	error
*** Mass Analysis for mass 256.0578260								
1	12	11	1	4	1	256.0580289	7.5	2.029e-04
2	10	9	4	3	1	256.0566863	8.0	1.140e-03
3	12	8	4	3	0	256.0590916	11.0	1.266e-03
4	9	10	3	6	0	256.0564115	6.5	1.415e-03
5	14	10	1	4	0	256.0604342	10.5	2.608e-03
6	15	9	2	1	1	256.0607090	12.0	2.883e-03
7	7	11	3	6	1	256.0540062	3.5	3.820e-03
8	6	12	2	9	0	256.0537314	2.0	4.095e-03
9	17	8	2	1	0	256.0631143	15.0	5.288e-03
10	5	15	1	9	1	256.0639022	-1.5	6.076e-03



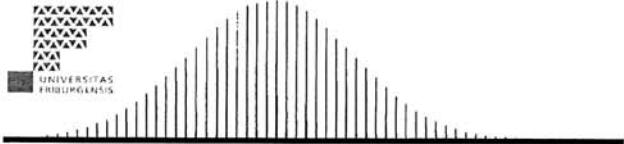






C:\PROGRAM FILES\OPUS\DATA\isak\is-029.0 is-029

26/01/2010



FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

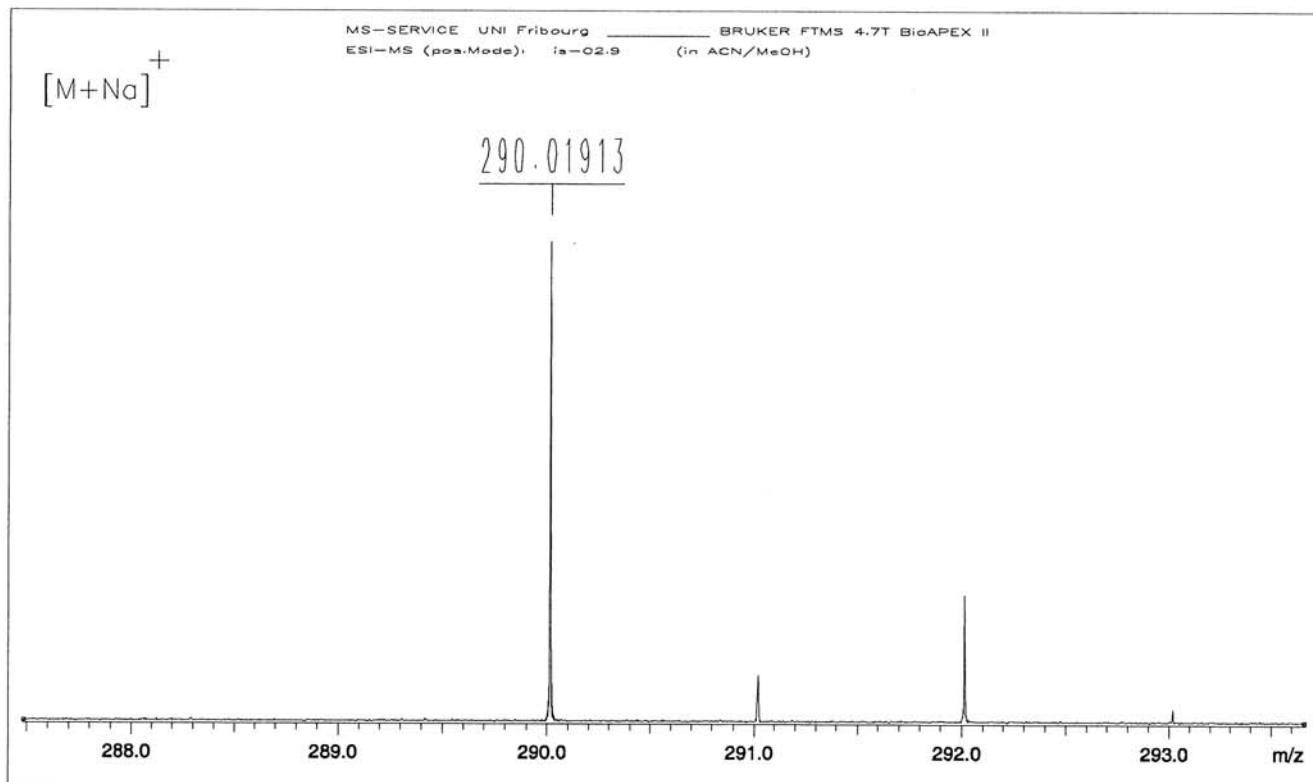
ESI-MS: is-029

XMASS Mass Analysis for /Data/UNI_FR/ALIM4617_ESI/2/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

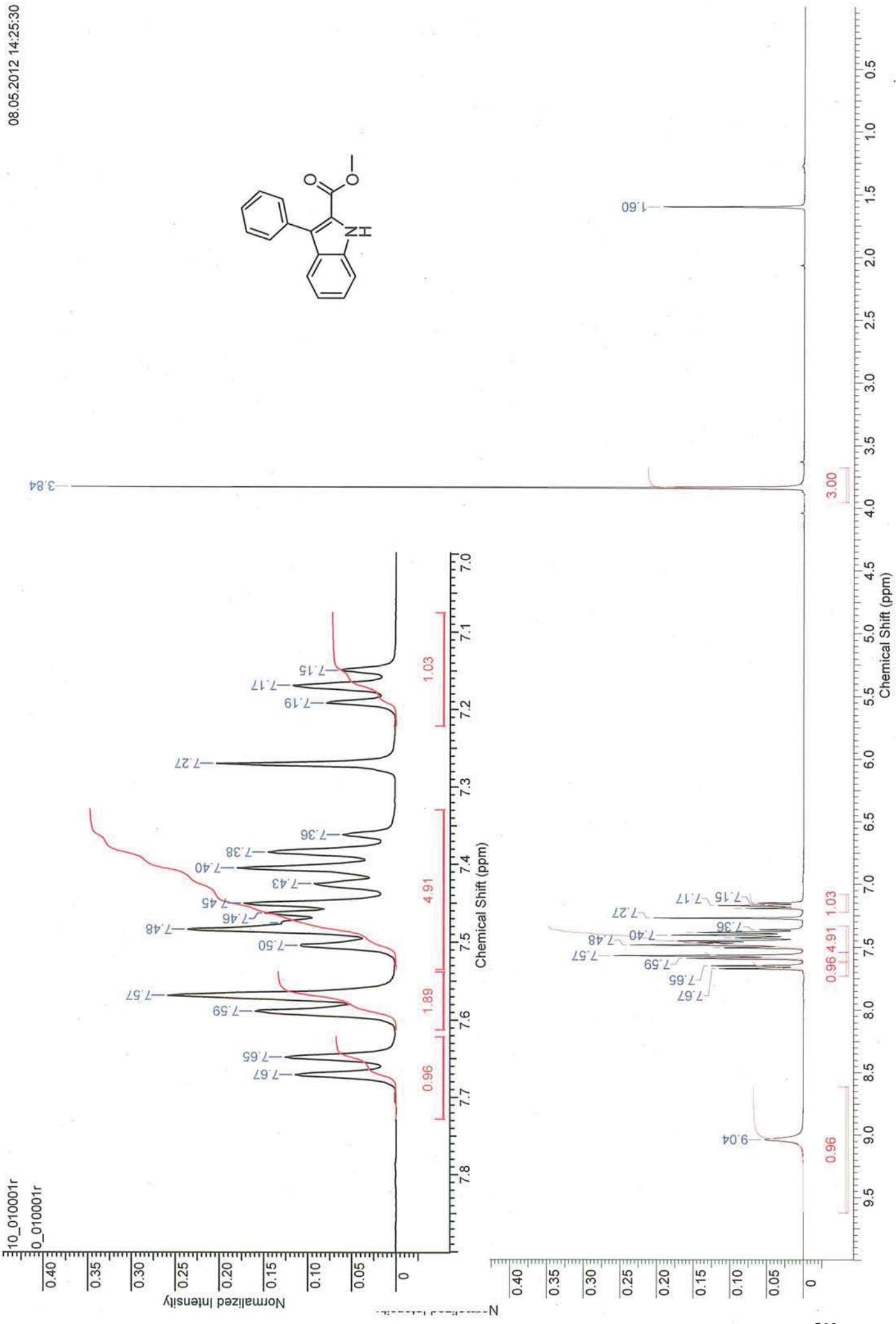
Ion mass = 290.0191340

Charge = +1

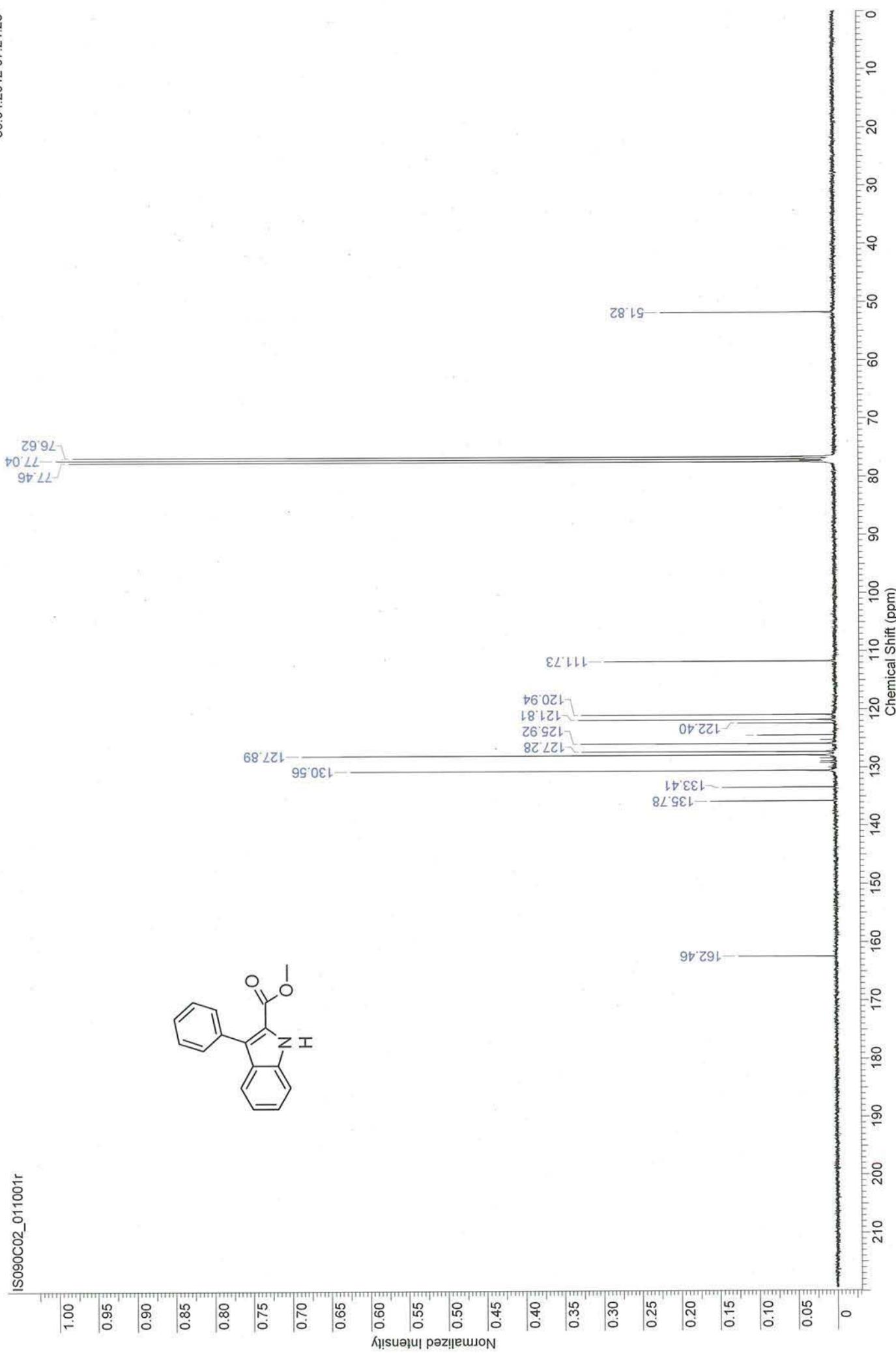
#	C	H	35Cl	N	O	Na	mass	DBE	error
*** Mass Analysis for mass 290.0191340									
1	12	10	1	1	4	1	290.0190566	7.5	7.739e-05
2	10	5	1	7	2	0	290.0187766	11.5	3.574e-04
3	12	7	1	4	3	0	290.0201192	11.0	9.852e-04
4	10	8	1	4	3	1	290.0177139	8.0	1.420e-03
5	14	9	1	1	4	0	290.0214619	10.5	2.328e-03
6	15	8	1	2	1	1	290.0217367	12.0	2.603e-03
7	11	12	2	2	3	0	290.0219491	6.0	2.815e-03
8	17	7	1	2	1	0	290.0241420	15.0	5.008e-03
9	15	10	2	1	1	0	290.0133958	10.5	5.738e-03
10	16	5	1	3	1	0	290.0115659	15.5	7.568e-03

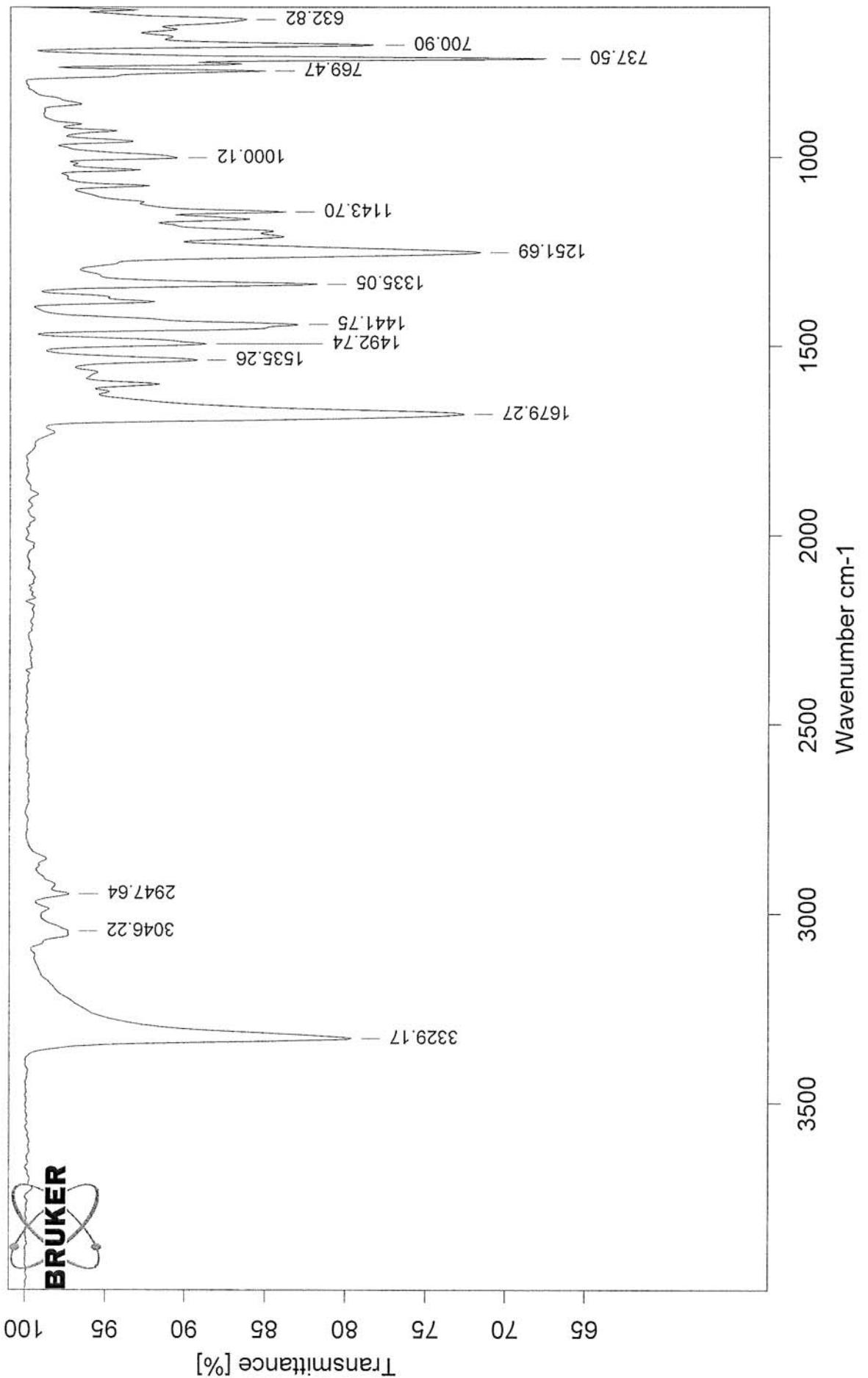


/Data/UNI_FR/ALIM4617_ESI/2/pdata/1 FTMS USER Mon Jan 4 10:40:37 2010



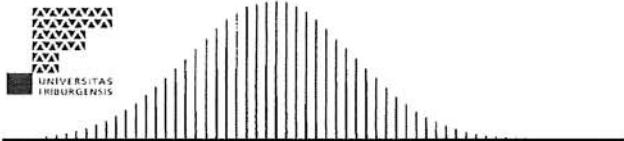
IS090C02_011001r





U:\IR_spectra\IS090.1 IS090 white yellow solid

09/07/2010



FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

ESI-MS: IS090

XMASS Mass Analysis for /Data/UNI_FR/ALIM4973_ESI/1/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

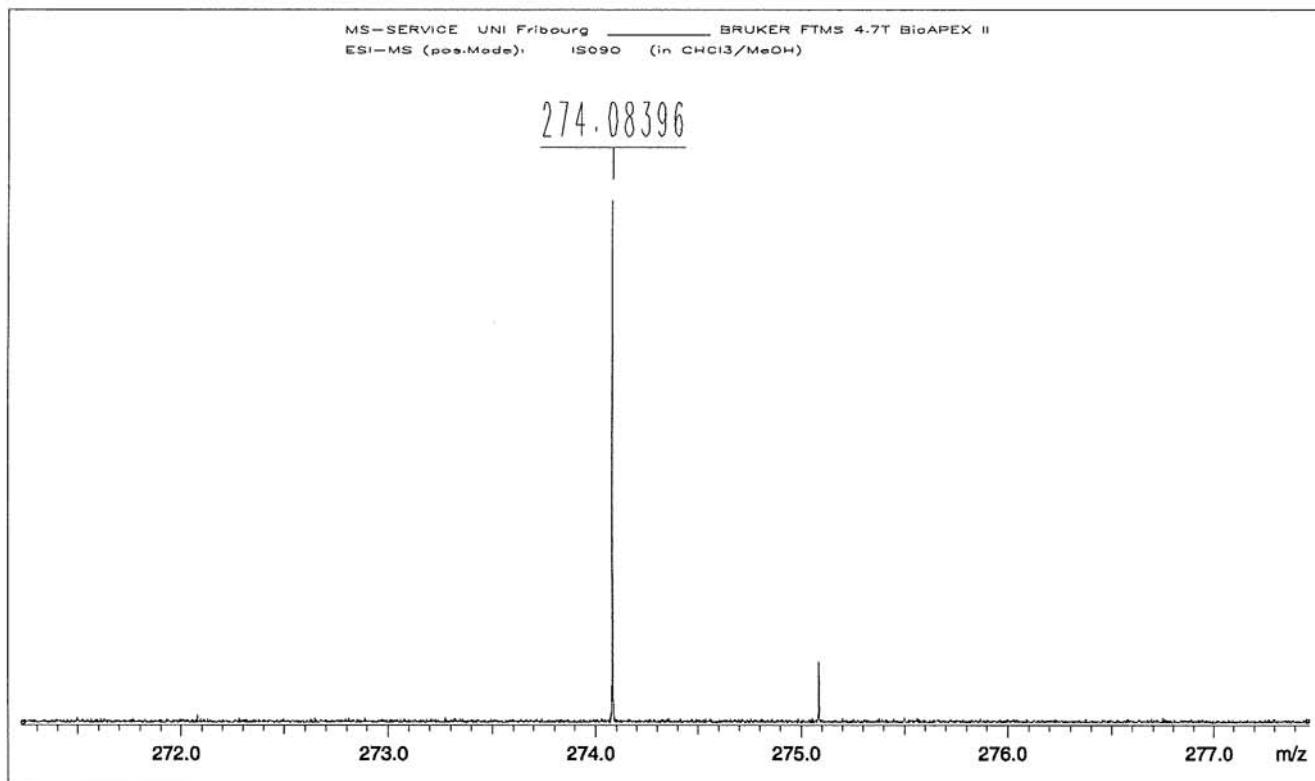
Ion mass = 274.0839610

Charge = +1

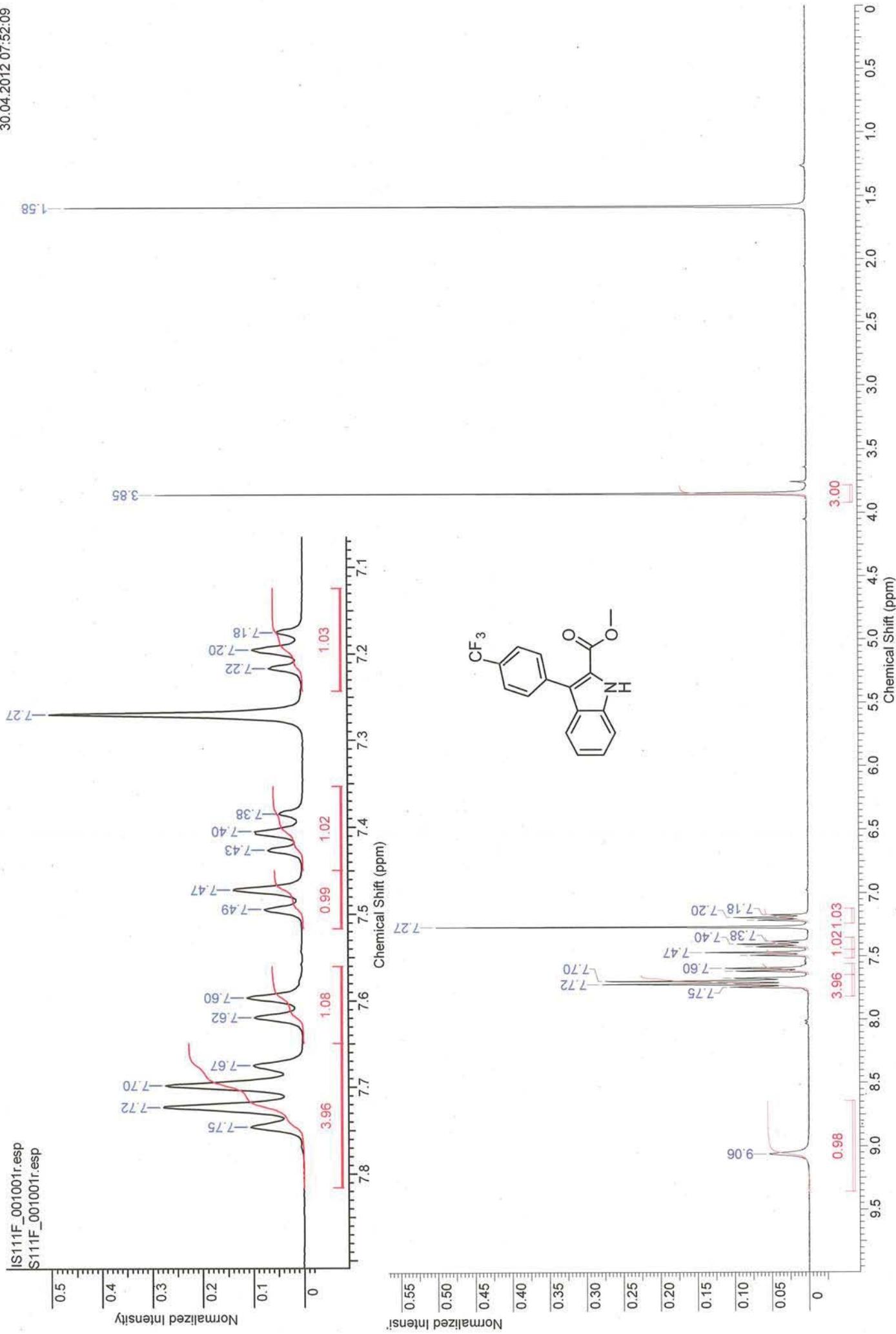
#	C	H	N	O	Na	mass	DBE	error
---	---	---	---	---	----	------	-----	-------

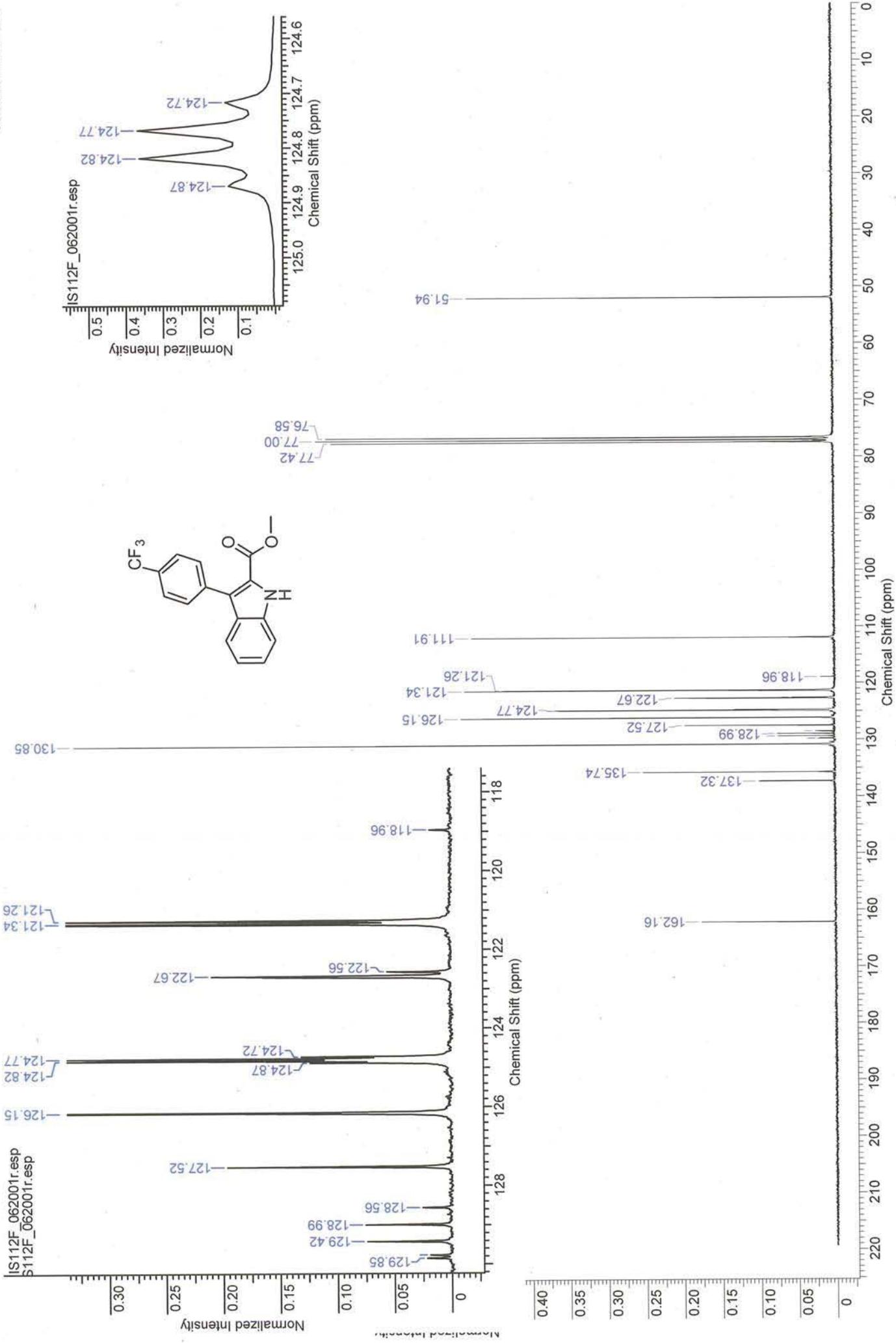
*** Mass Analysis for mass 274.0839610

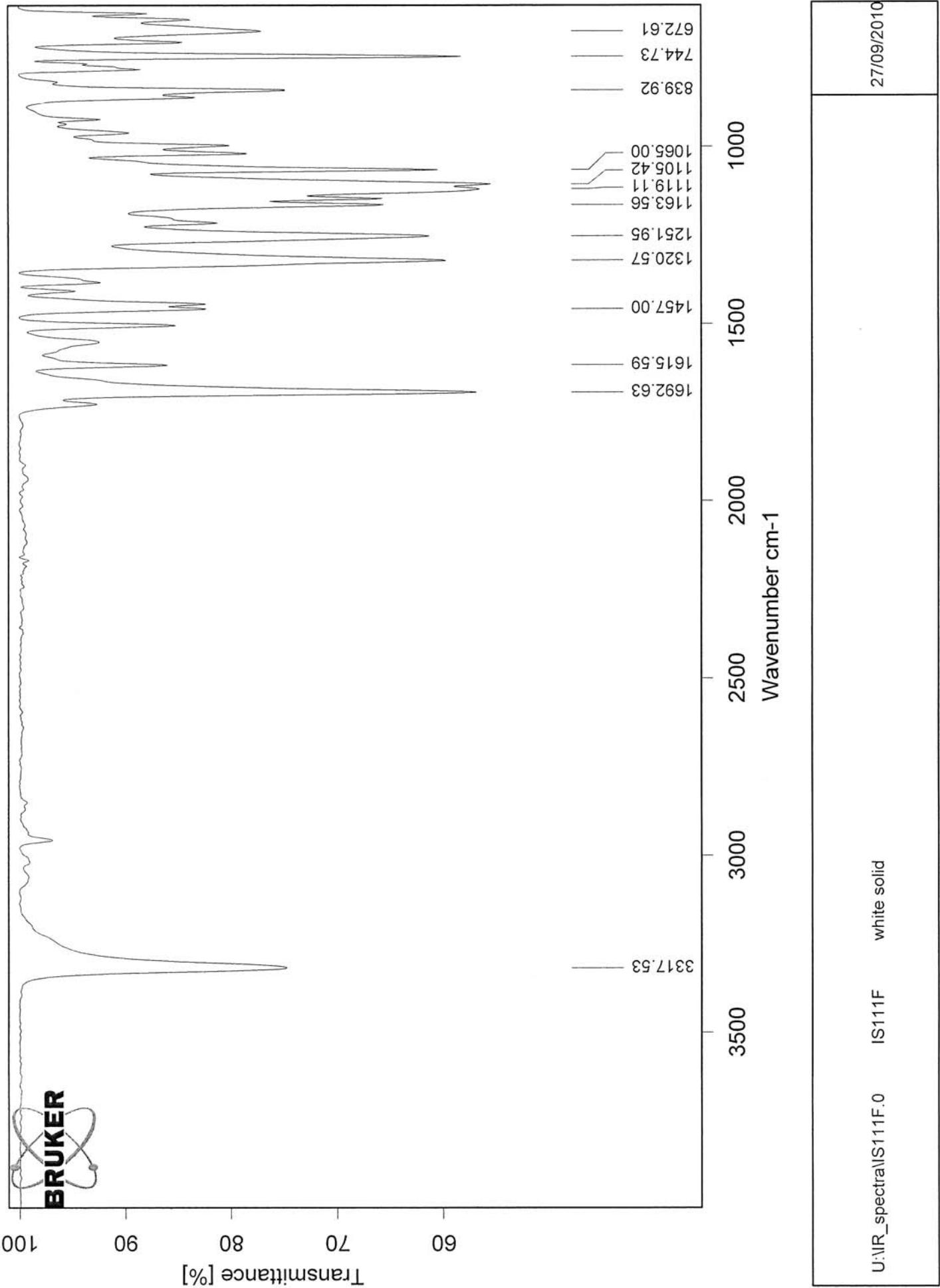
1	16	13	1	2	1	274.0838498	10.5	1.112e-04
2	16	10	4	1	0	274.0849124	14.0	9.514e-04
3	14	11	4	1	1	274.0825071	11.0	1.454e-03
4	13	12	3	4	0	274.0822323	9.5	1.729e-03
5	18	12	1	2	0	274.0862551	13.5	2.294e-03
6	11	10	6	3	0	274.0808896	10.0	3.071e-03
7	5	13	7	5	1	274.0870377	2.5	3.077e-03
8	11	13	3	4	1	274.0798270	6.5	4.134e-03
9	5	10	10	4	0	274.0881003	6.0	4.139e-03
10	6	16	3	9	0	274.0881055	0.5	4.145e-03

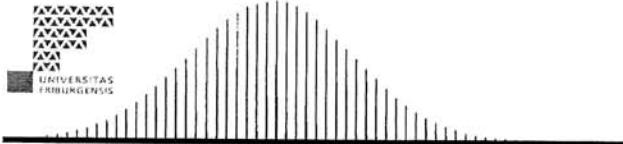


/Data/UNI_FR/ALIM4973_ESI/1/pdata/1 FTMS USER Thu Jul 8 11:57:50 2010









FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

ESI-MS: IS111F

XMASS Mass Analysis for /Data/UNI_FR/ALIM5098_ESI/3/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

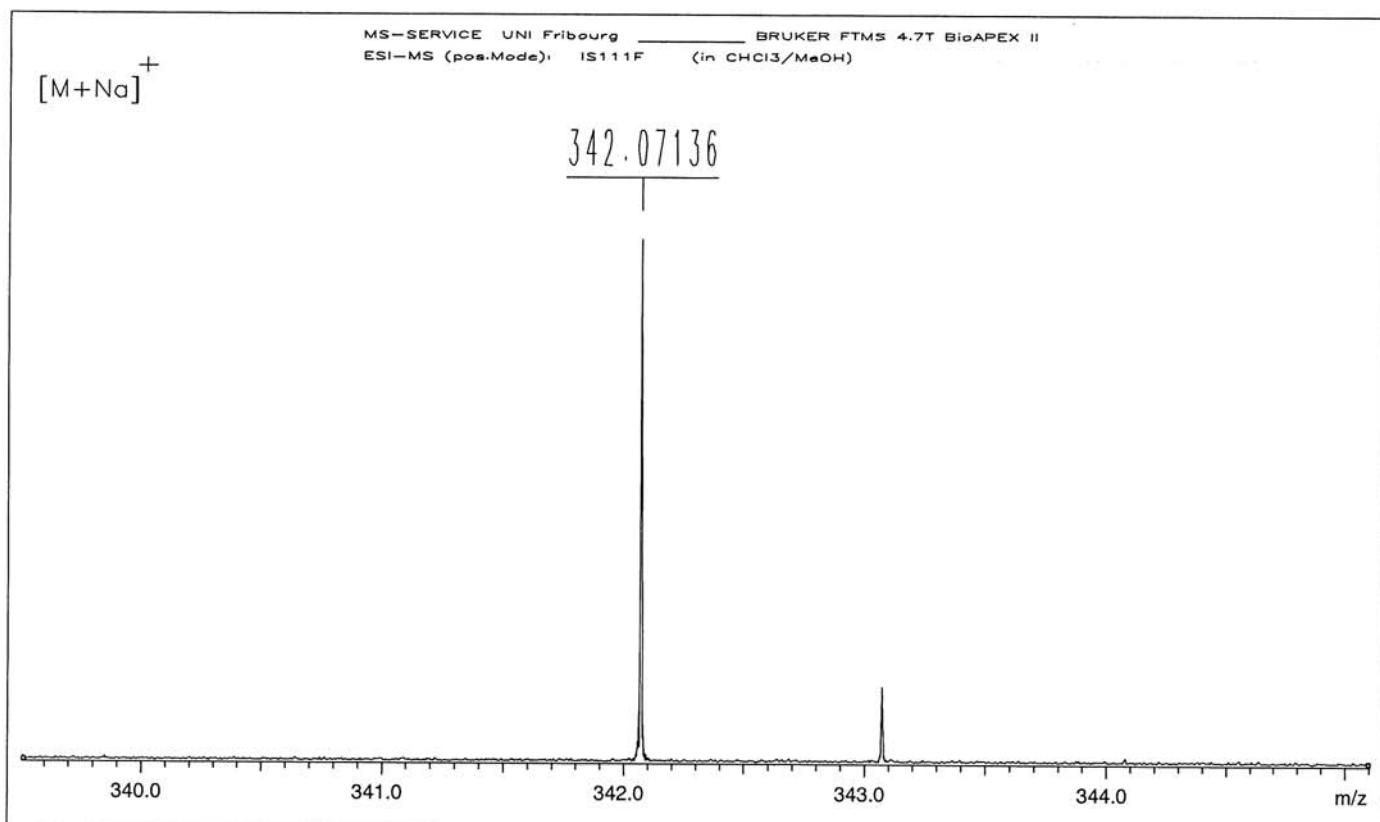
Ion mass = 342.0713610

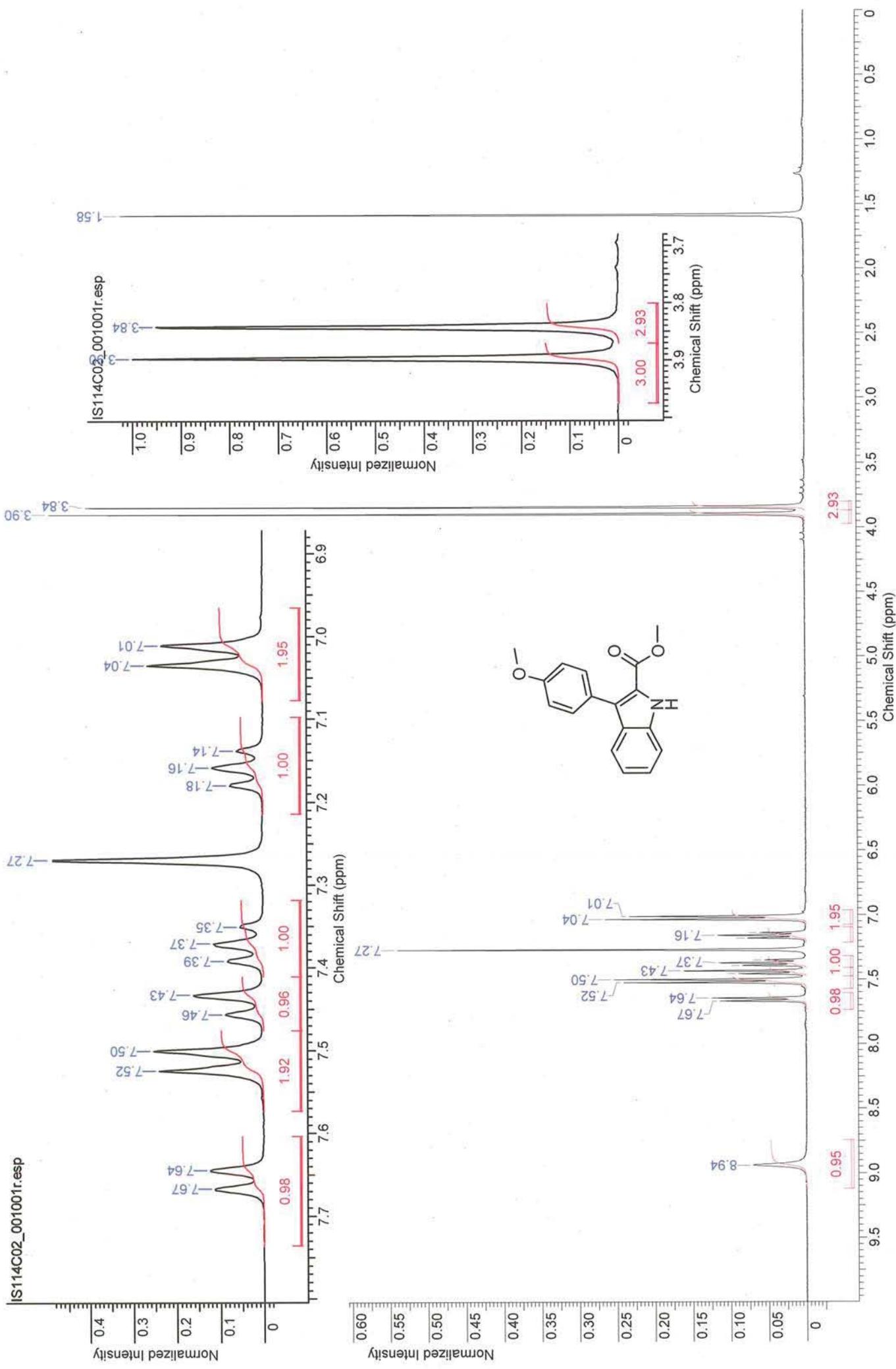
Charge = +1

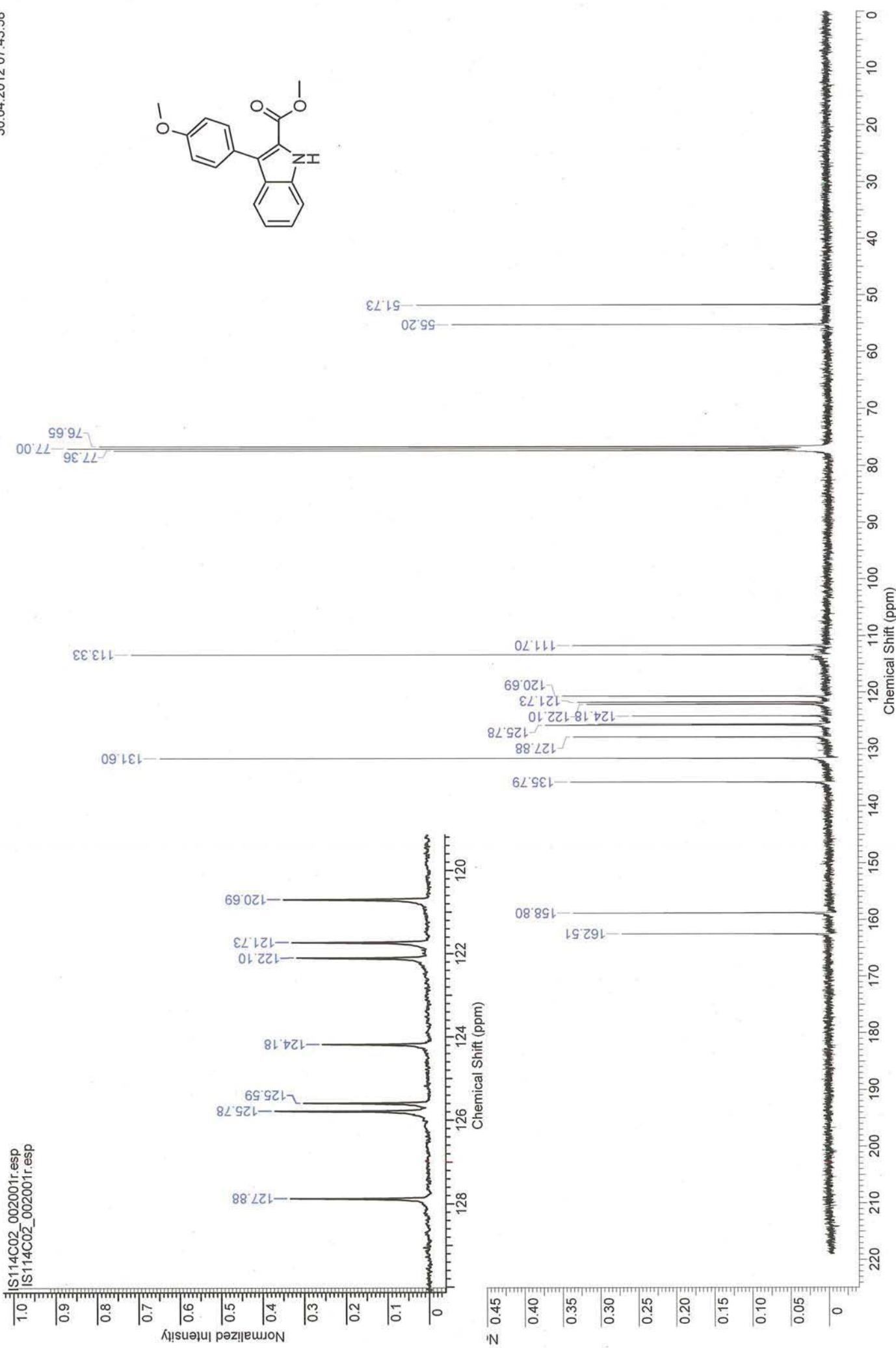
#	C	H	F	N	O	Na	mass	DBE	error
---	---	---	---	---	---	----	------	-----	-------

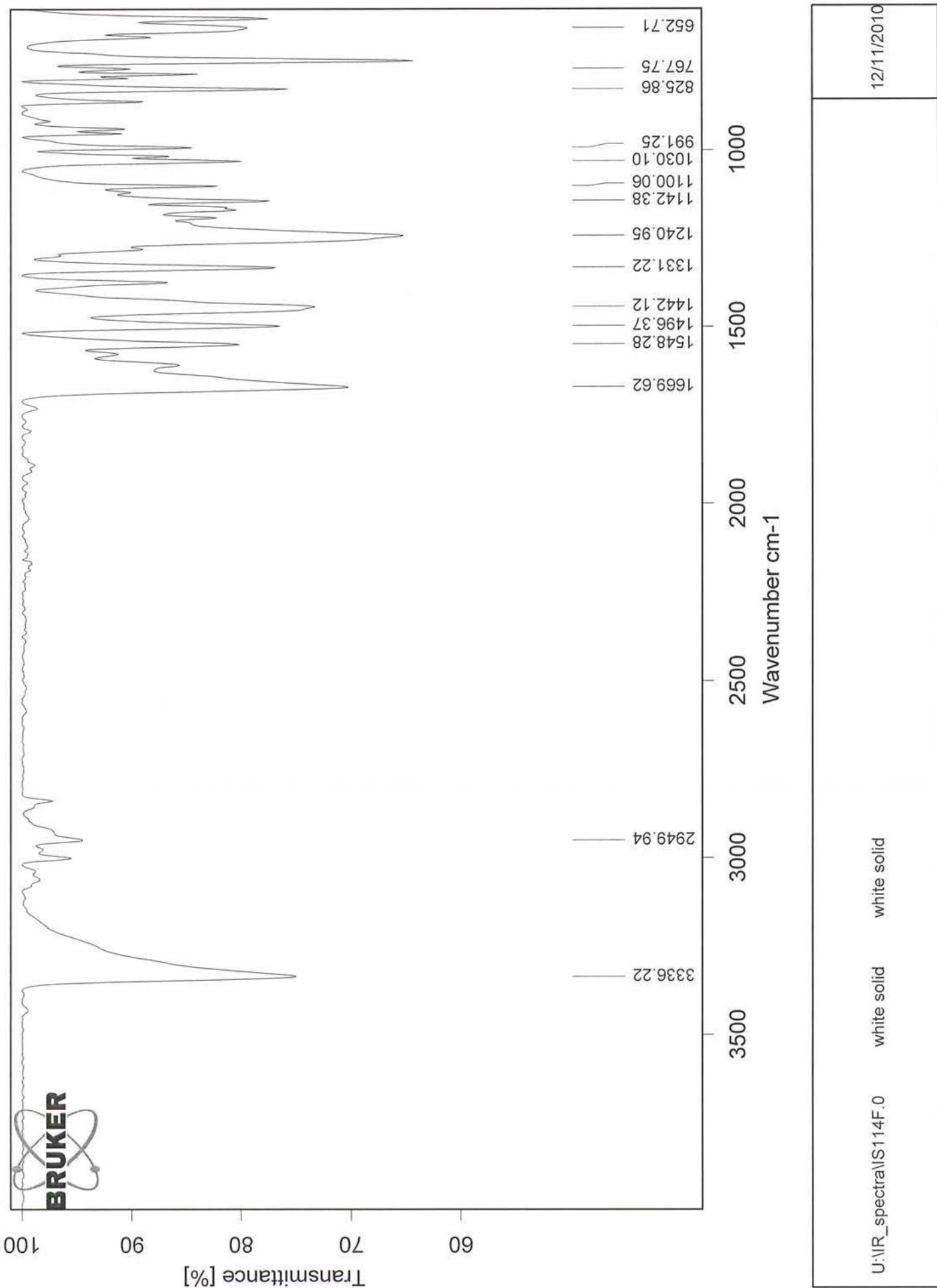
*** Mass Analysis for mass 342.0713610

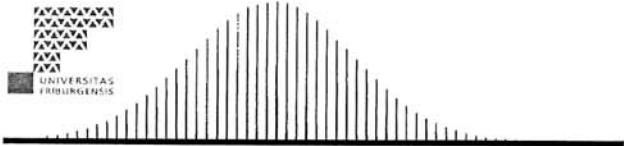
1	17	12	3	1	2	1	342.0712343	10.5	1.267e-04
2	12	11	4	4	2	1	342.0710345	7.0	3.265e-04
3	11	12	4	3	5	0	342.0707597	5.5	6.013e-04
4	9	12	5	4	3	1	342.0721774	3.0	8.164e-04
5	17	9	3	4	1	0	342.0722970	14.0	9.360e-04
6	14	13	4	1	3	1	342.0723772	6.5	1.016e-03
7	22	10	2	1	1	0	342.0724968	17.5	1.136e-03
8	20	11	2	1	1	1	342.0700915	14.5	1.270e-03
9	15	10	3	4	1	1	342.0698917	11.0	1.469e-03
10	14	11	3	3	4	0	342.0696169	9.5	1.744e-03











FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

ESI-MS: IS114F

XMASS Mass Analysis for /Data/UNI_FR/ALIM5099_ESI/3/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

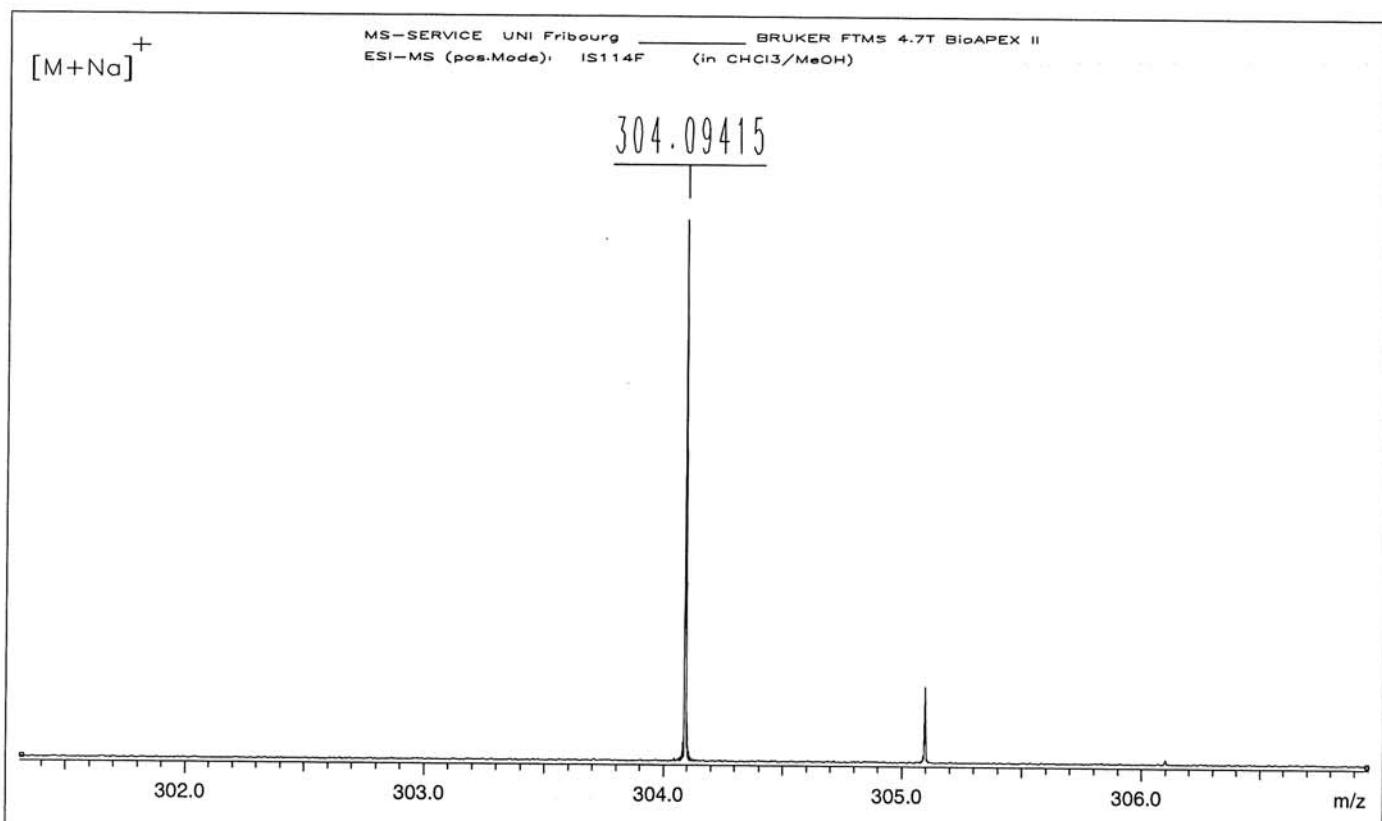
Ion mass = 304.0941460

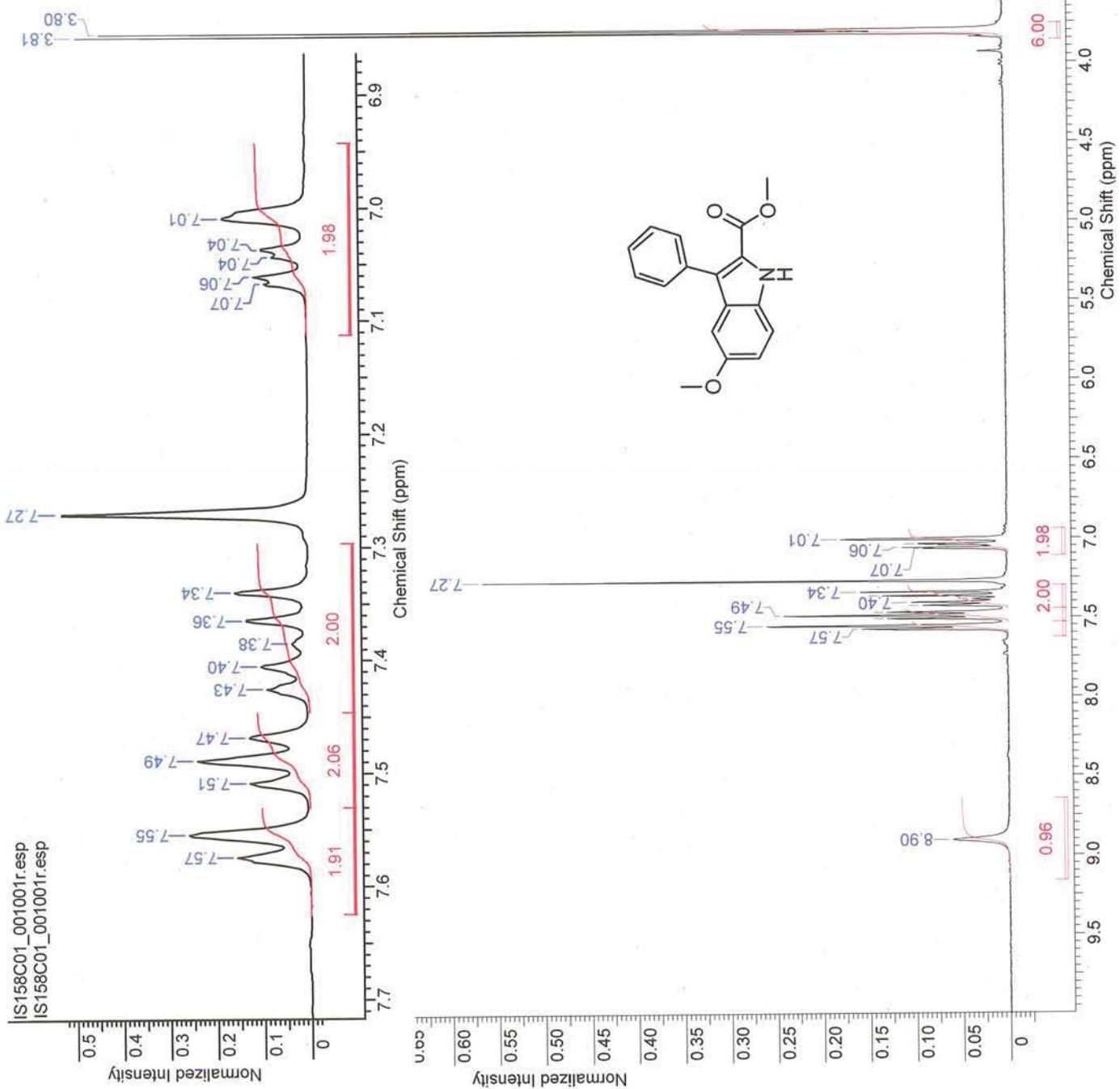
Charge = +1

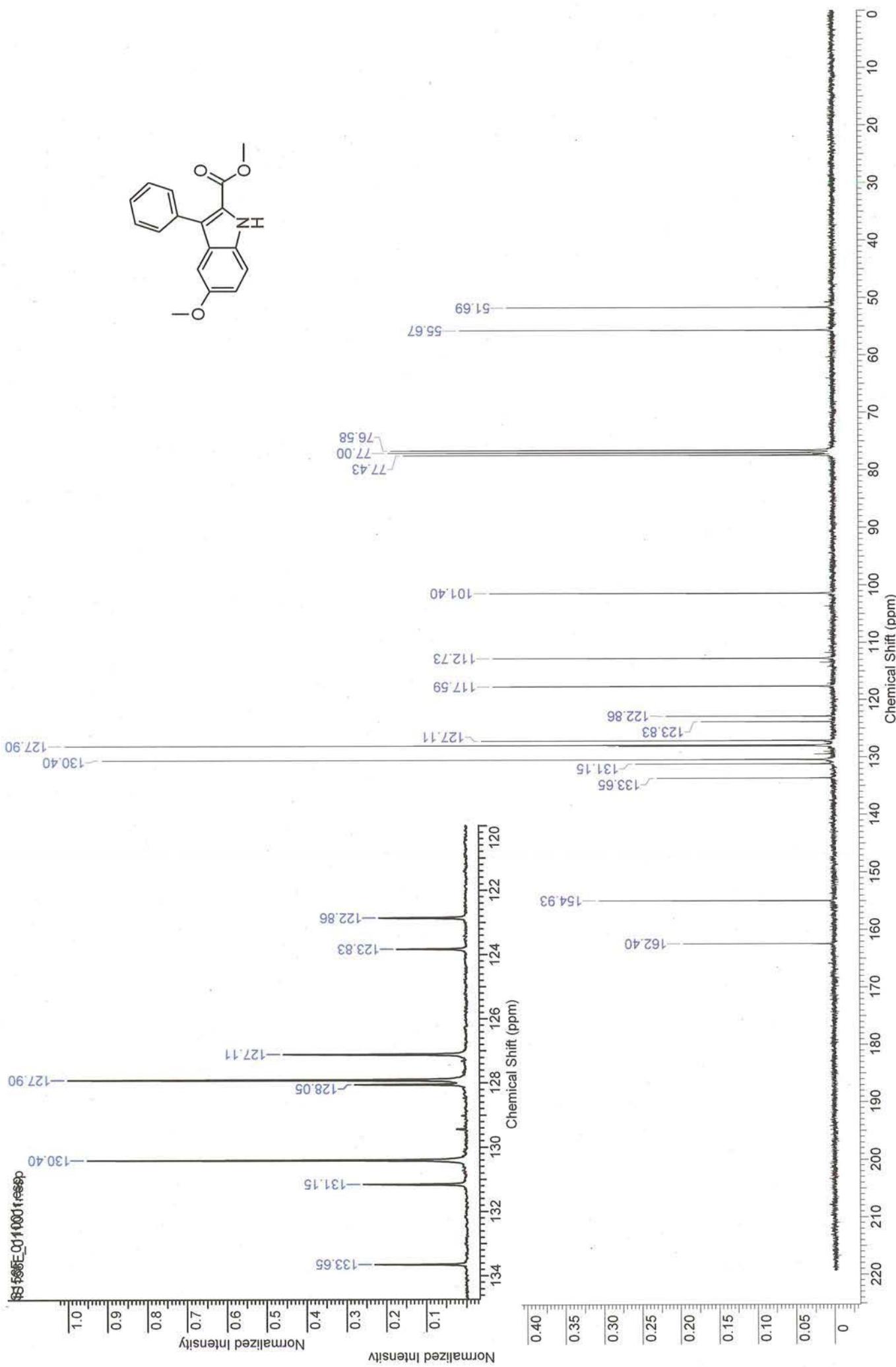
#	C	H	N	O	Na	mass	DBE	error
---	---	---	---	---	----	------	-----	-------

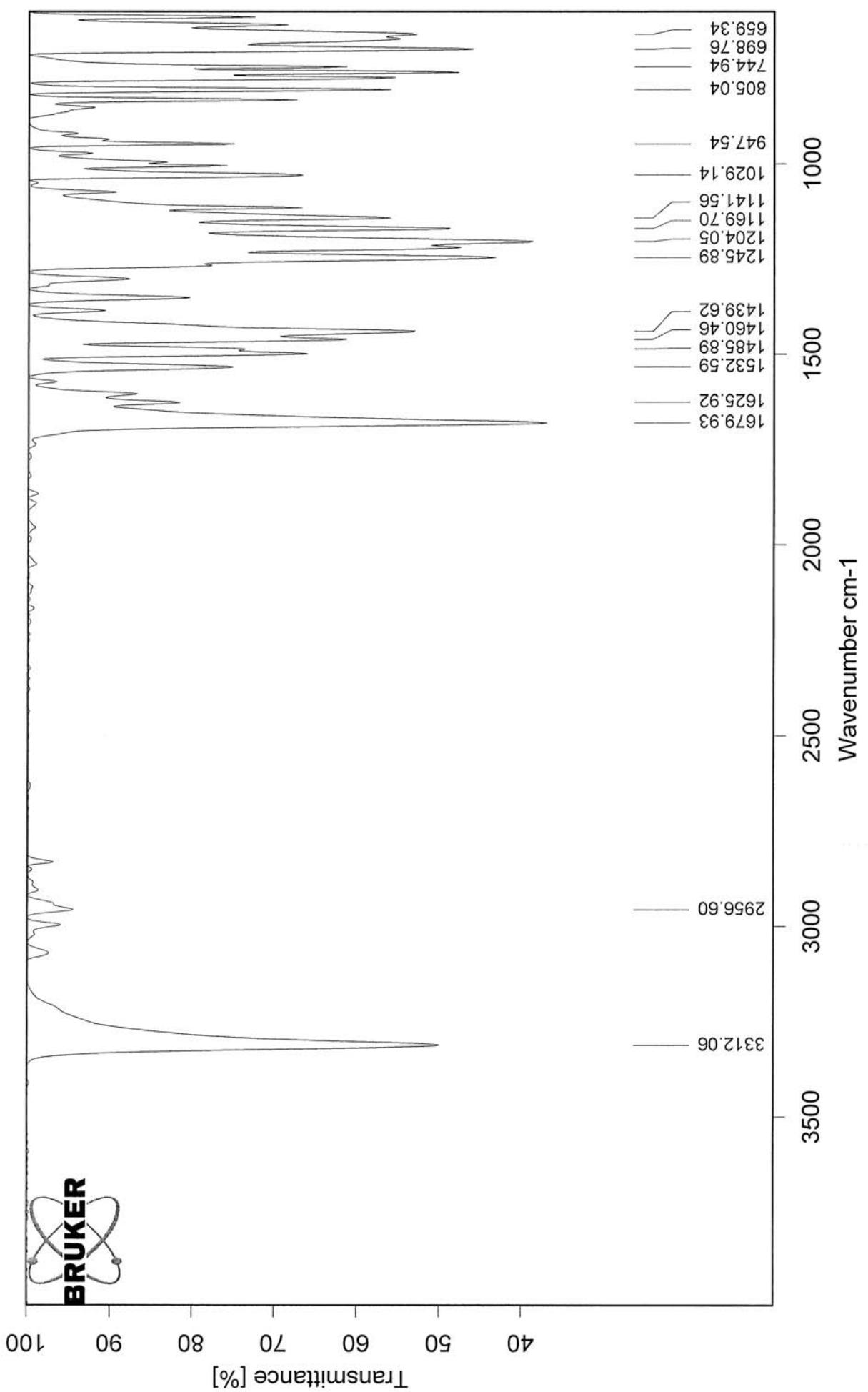
*** Mass Analysis for mass 304.0941460

1	17	15	1	3	1	304.0944144	10.5	2.684e-04
2	15	13	4	2	1	304.0930718	11.0	1.074e-03
3	17	12	4	2	0	304.0954771	14.0	1.331e-03
4	14	14	3	5	0	304.0927970	9.5	1.349e-03
5	5	19	3	10	1	304.0962649	-2.5	2.119e-03
6	19	14	1	3	0	304.0968197	13.5	2.674e-03
7	12	15	3	5	1	304.0903917	6.5	3.754e-03
8	11	16	2	8	0	304.0901169	5.0	4.029e-03
9	7	18	3	10	0	304.0986702	0.5	4.524e-03
10	8	17	4	7	1	304.0989450	2.0	4.799e-03

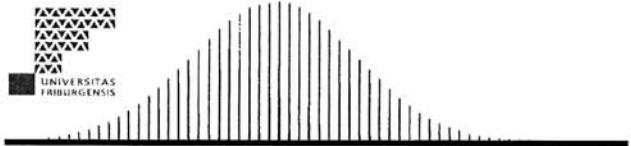








U:\IR_spectra\IS158F.0	IS158F	white solid	21/12/2010
------------------------	--------	-------------	------------



FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

ESI-MS: IS158.F

XMASS Mass Analysis for /Data/UNI_FR/ALIM5318_ESI/1/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

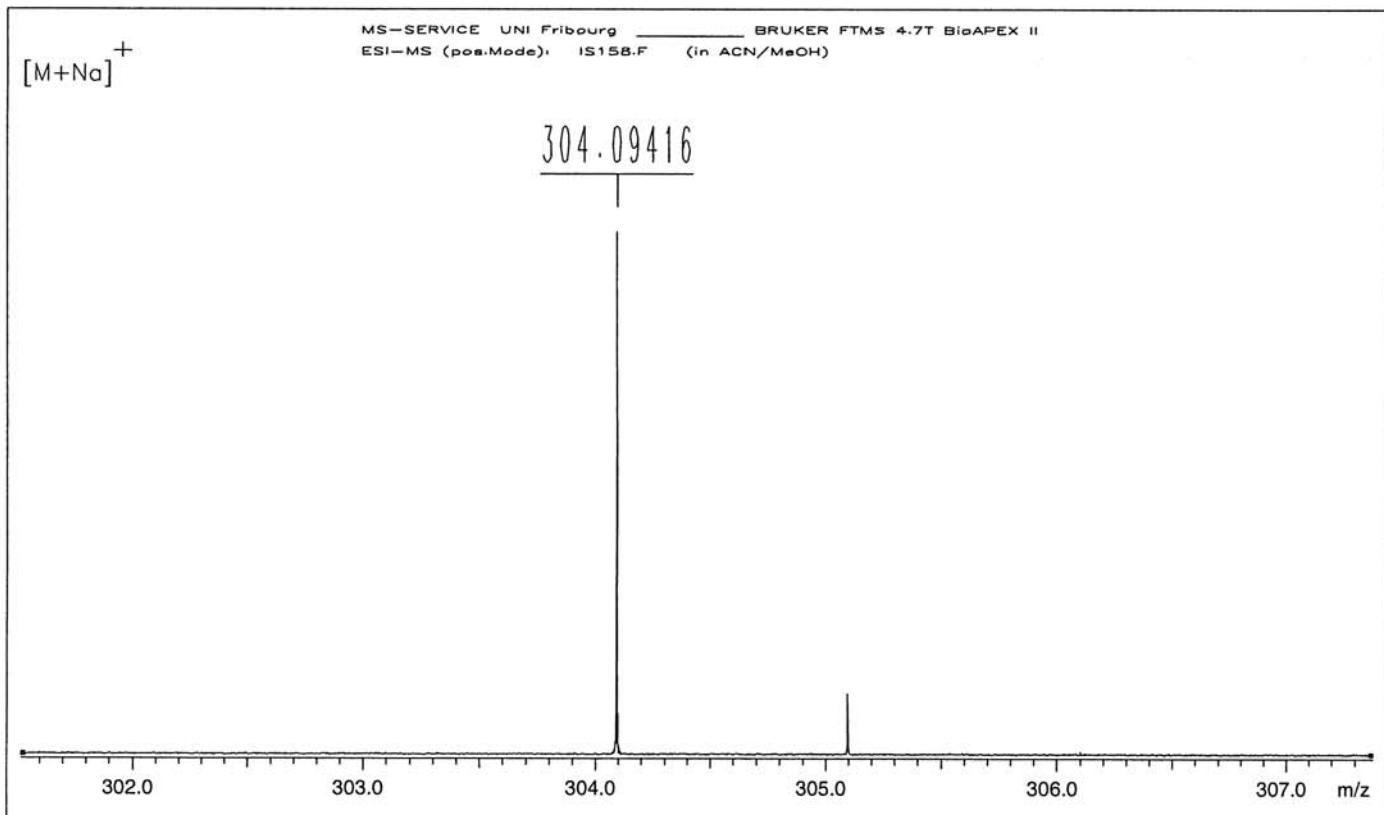
Ion mass = 304.0941630

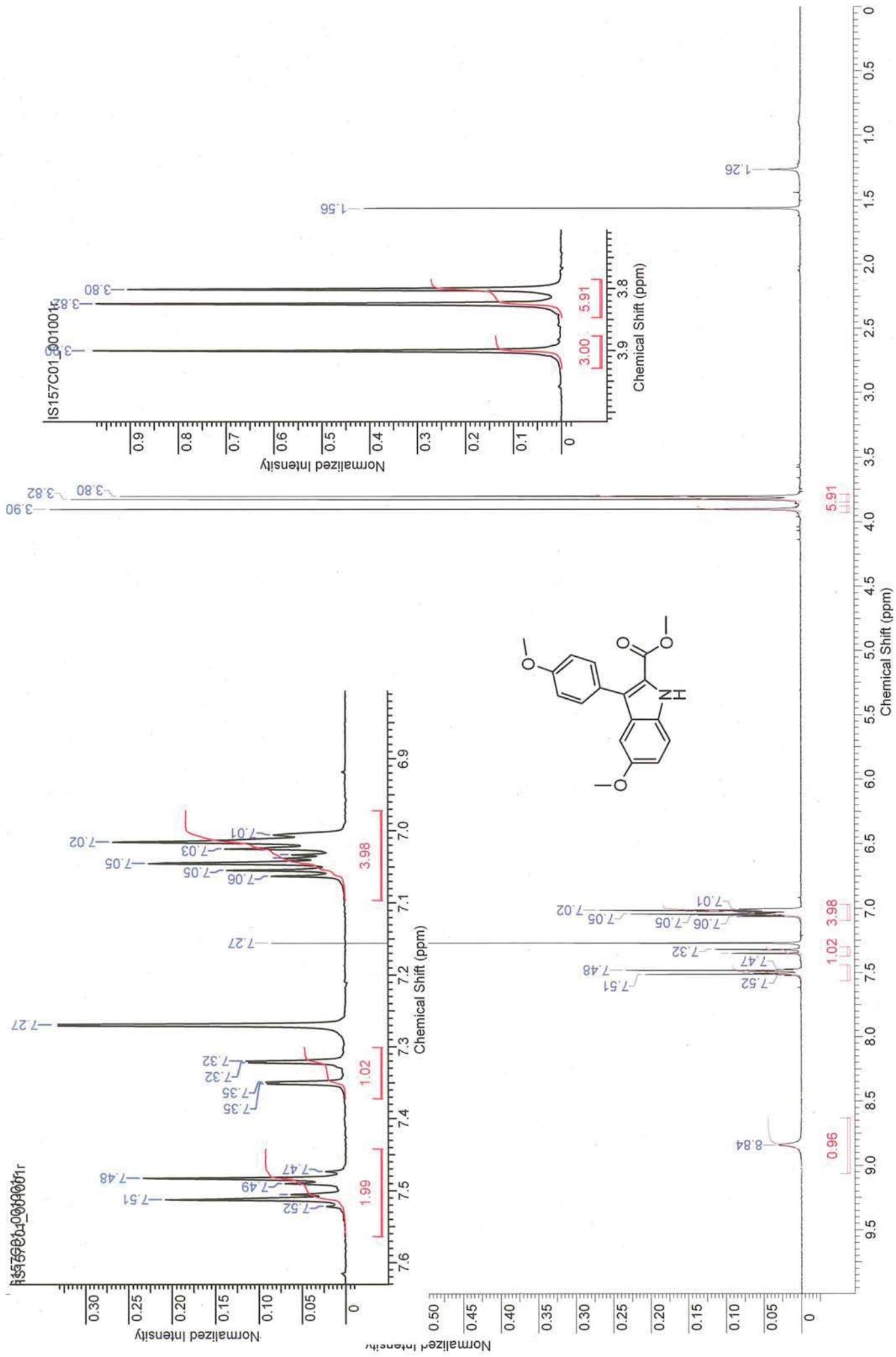
Charge = +1

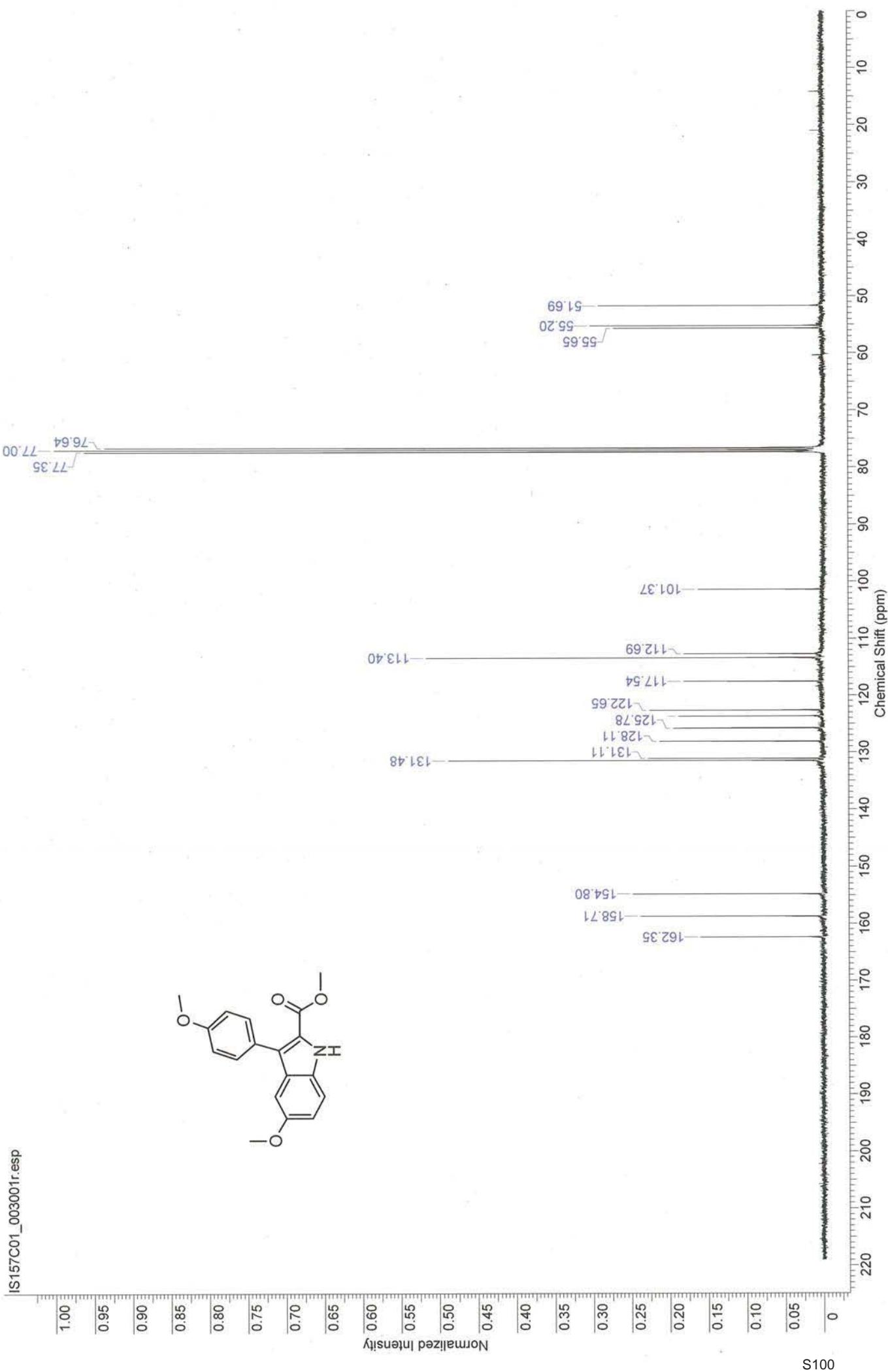
#	C	H	N	O	Na	mass	DBE	error
---	---	---	---	---	----	------	-----	-------

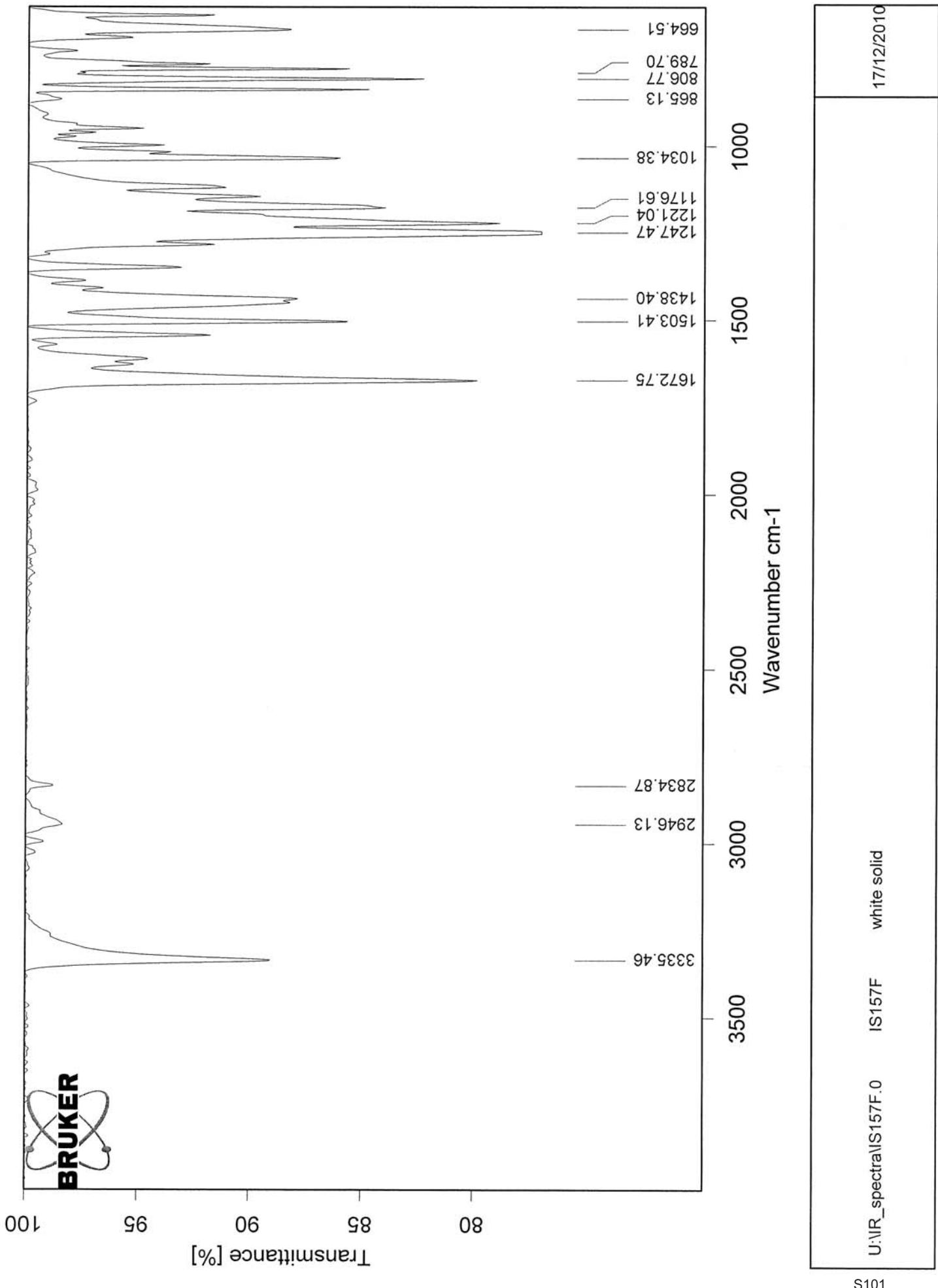
*** Mass Analysis for mass 304.0941630

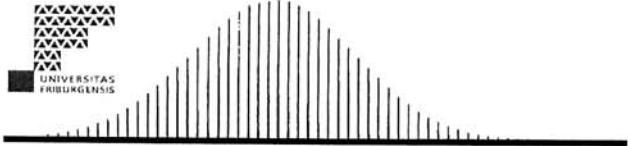
1	17	15	1	3	1	304.0944144	10.5	2.514e-04
2	15	13	4	2	1	304.0930718	11.0	1.091e-03
3	17	12	4	2	0	304.0954771	14.0	1.314e-03
4	14	14	3	5	0	304.0927970	9.5	1.366e-03
5	5	19	3	10	1	304.0962649	-2.5	2.102e-03
6	19	14	1	3	0	304.0968197	13.5	2.657e-03
7	12	12	6	4	0	304.0914543	10.0	2.709e-03
8	5	16	6	9	0	304.0973276	1.0	3.165e-03
9	12	15	3	5	1	304.0903917	6.5	3.771e-03
10	11	16	2	8	0	304.0901169	5.0	4.046e-03











FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

ESI-MS: IS157C01

XMASS Mass Analysis for /Data/UNI_FR/ALIM5301_ESI/2/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

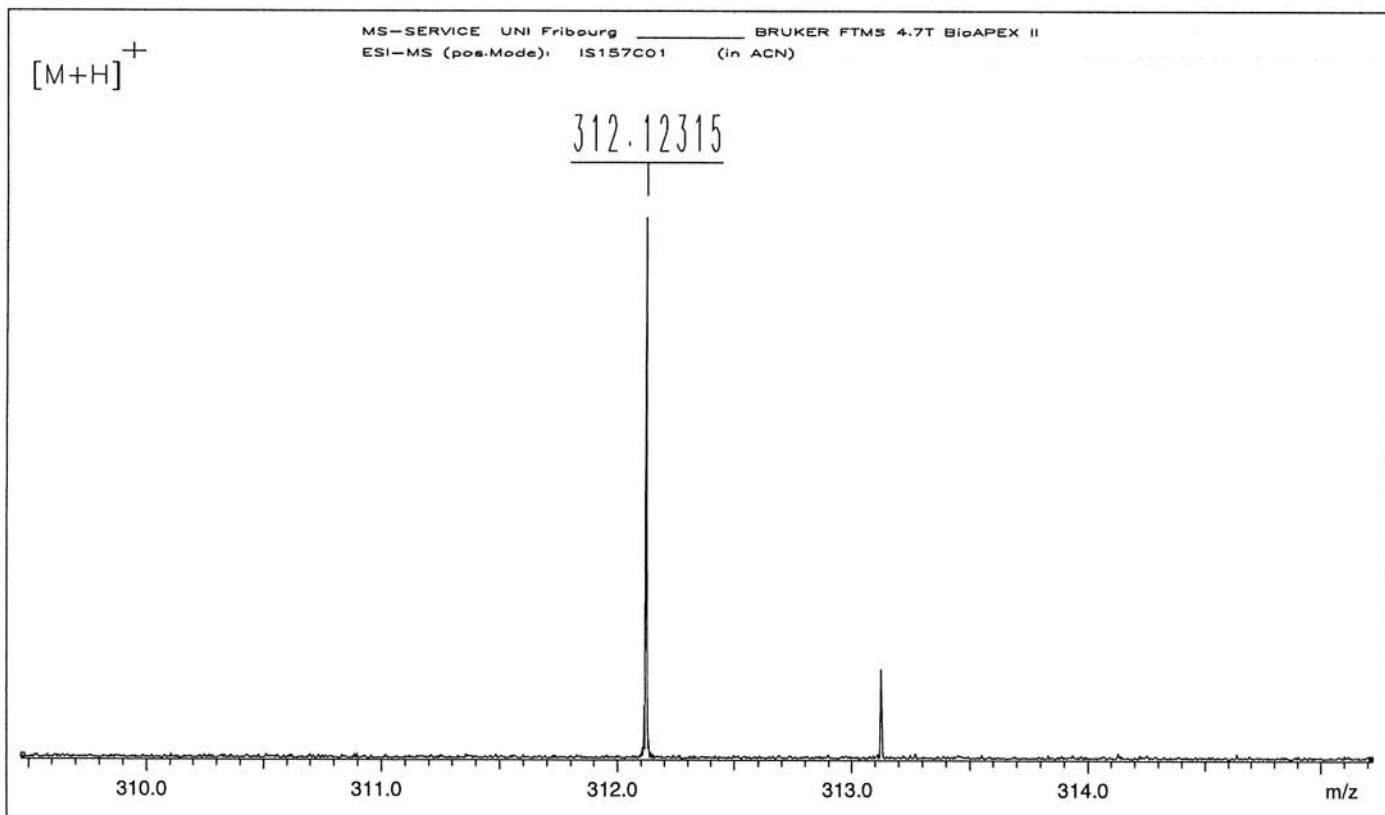
Ion mass = 312.1231460

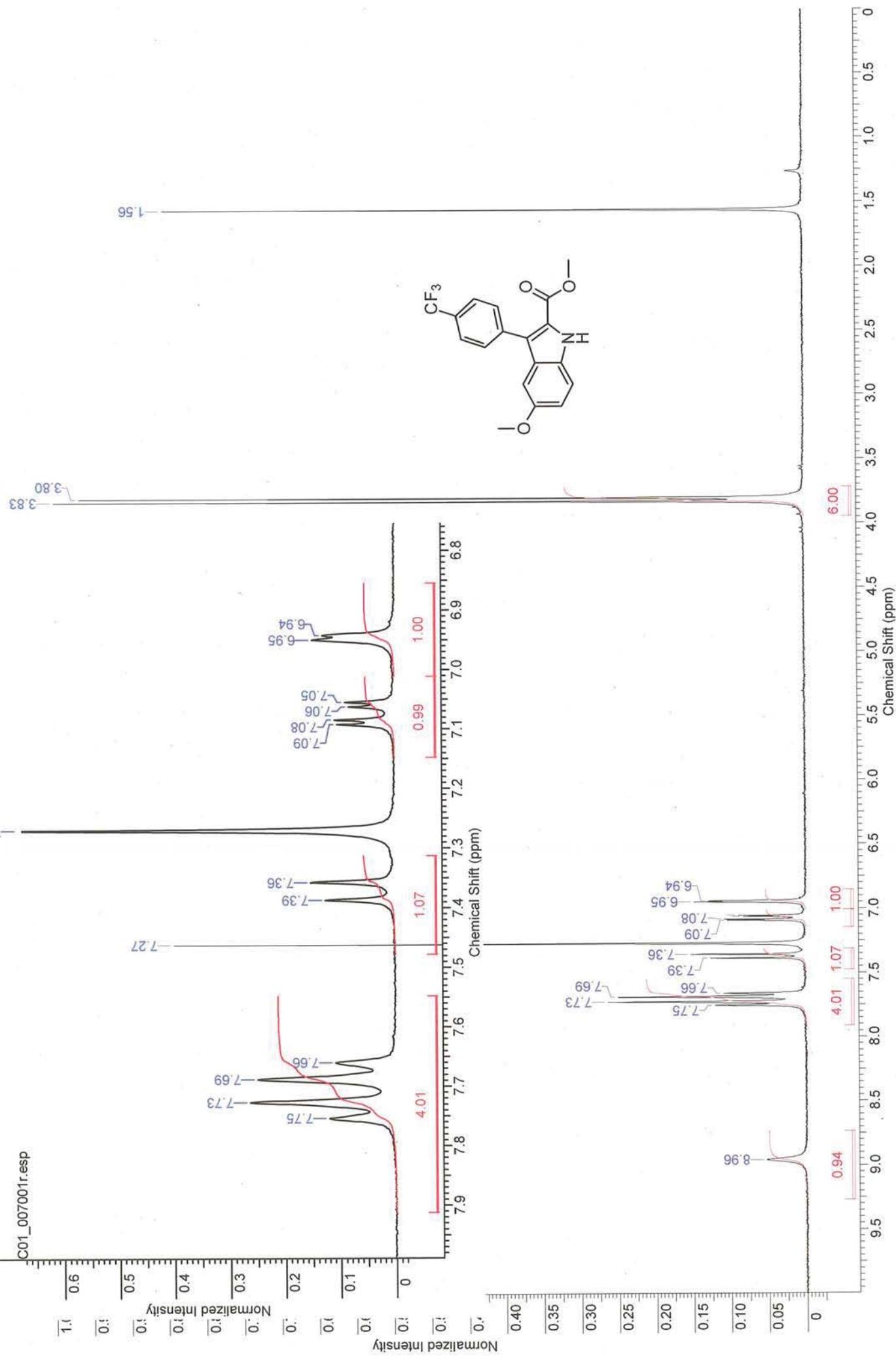
Charge = +1

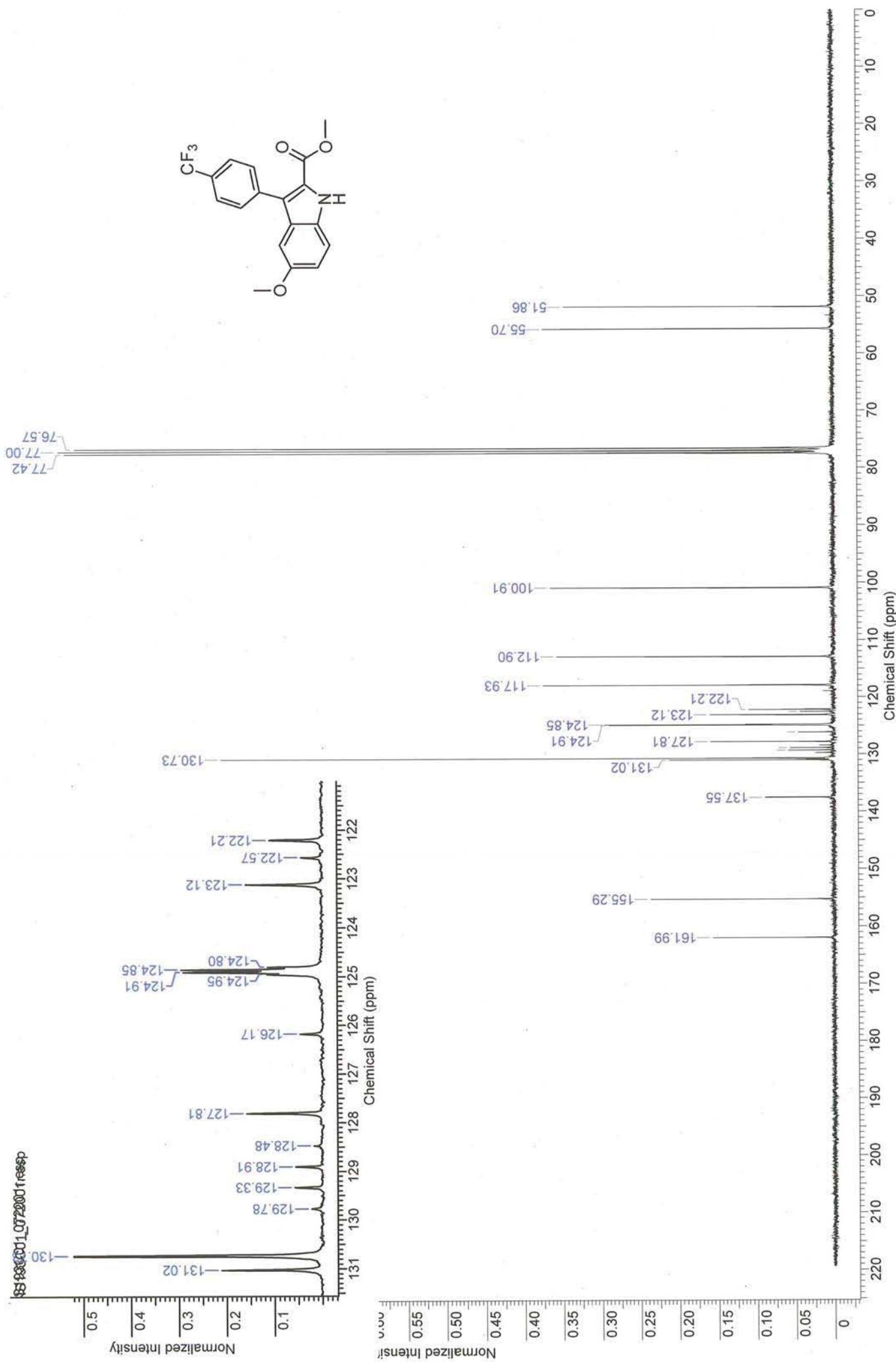
#	C	H	N	O	mass	DBE	error
---	---	---	---	---	------	-----	-------

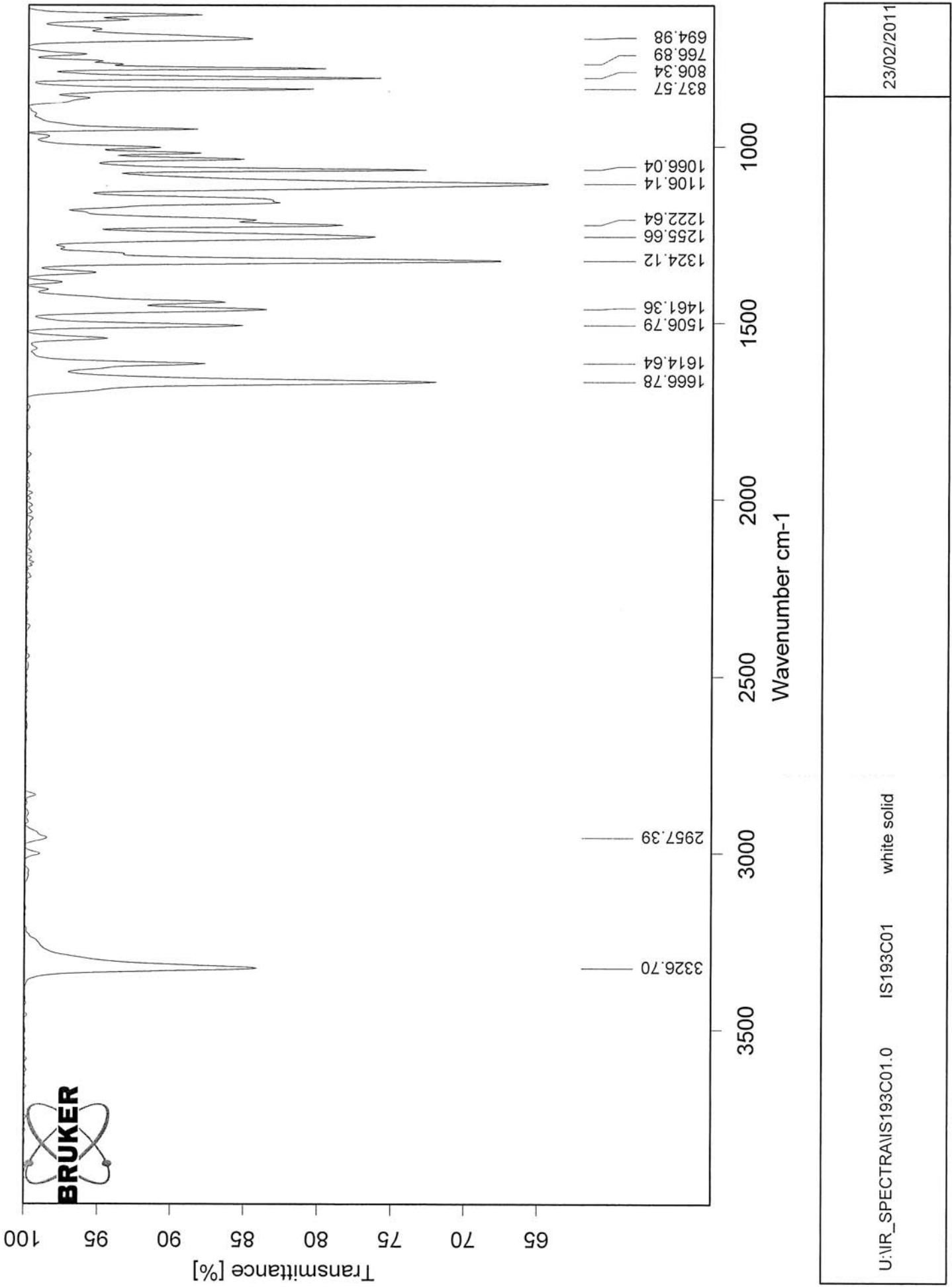
*** Mass Analysis for mass 312.1231460

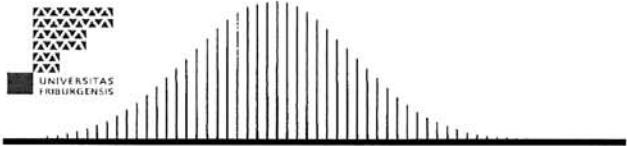
1	18	18	1	4	312.1230345	10.5	1.115e-04
2	16	16	4	3	312.1216918	11.0	1.454e-03
3	5	16	10	6	312.1248797	3.0	1.734e-03
4	21	16	2	1	312.1257146	15.0	2.569e-03
5	14	14	7	2	312.1203491	11.5	2.797e-03
6	7	18	7	7	312.1262224	2.5	3.076e-03
7	13	18	3	6	312.1190117	6.5	4.134e-03
8	12	12	10	1	312.1190065	12.0	4.140e-03
9	9	20	4	8	312.1275651	2.0	4.419e-03
10	11	16	6	5	312.1176691	7.0	5.477e-03











FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

ESI-MS: IS193C01

XMASS Mass Analysis for /Data/UNI_FR/ALIM5470_ESI/5/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

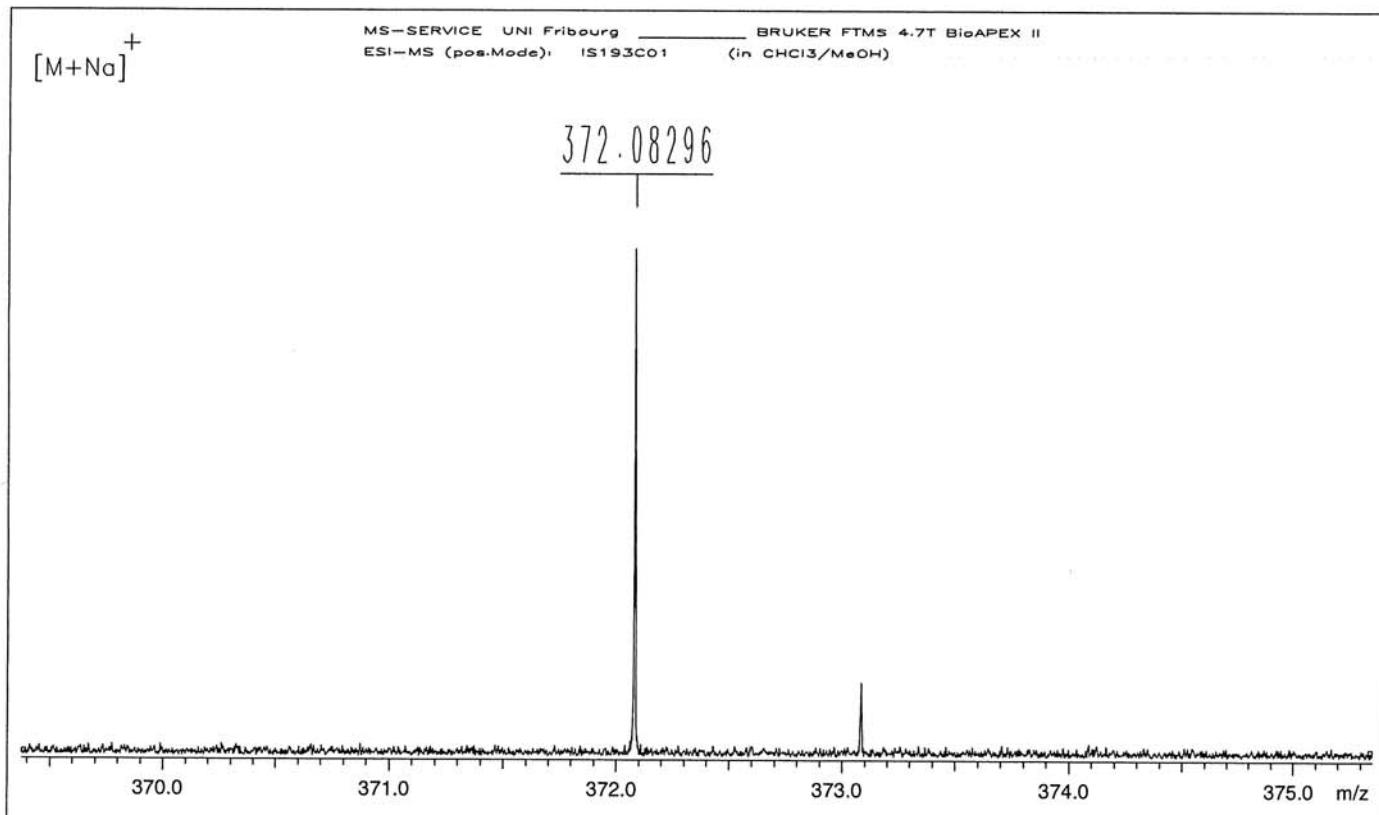
Ion mass = 372.0829640

Charge = +1

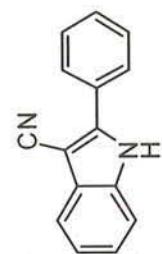
#	C	H	F	N	O	Na	mass	DBE	error
---	---	---	---	---	---	----	------	-----	-------

*** Mass Analysis for mass 372.0829640

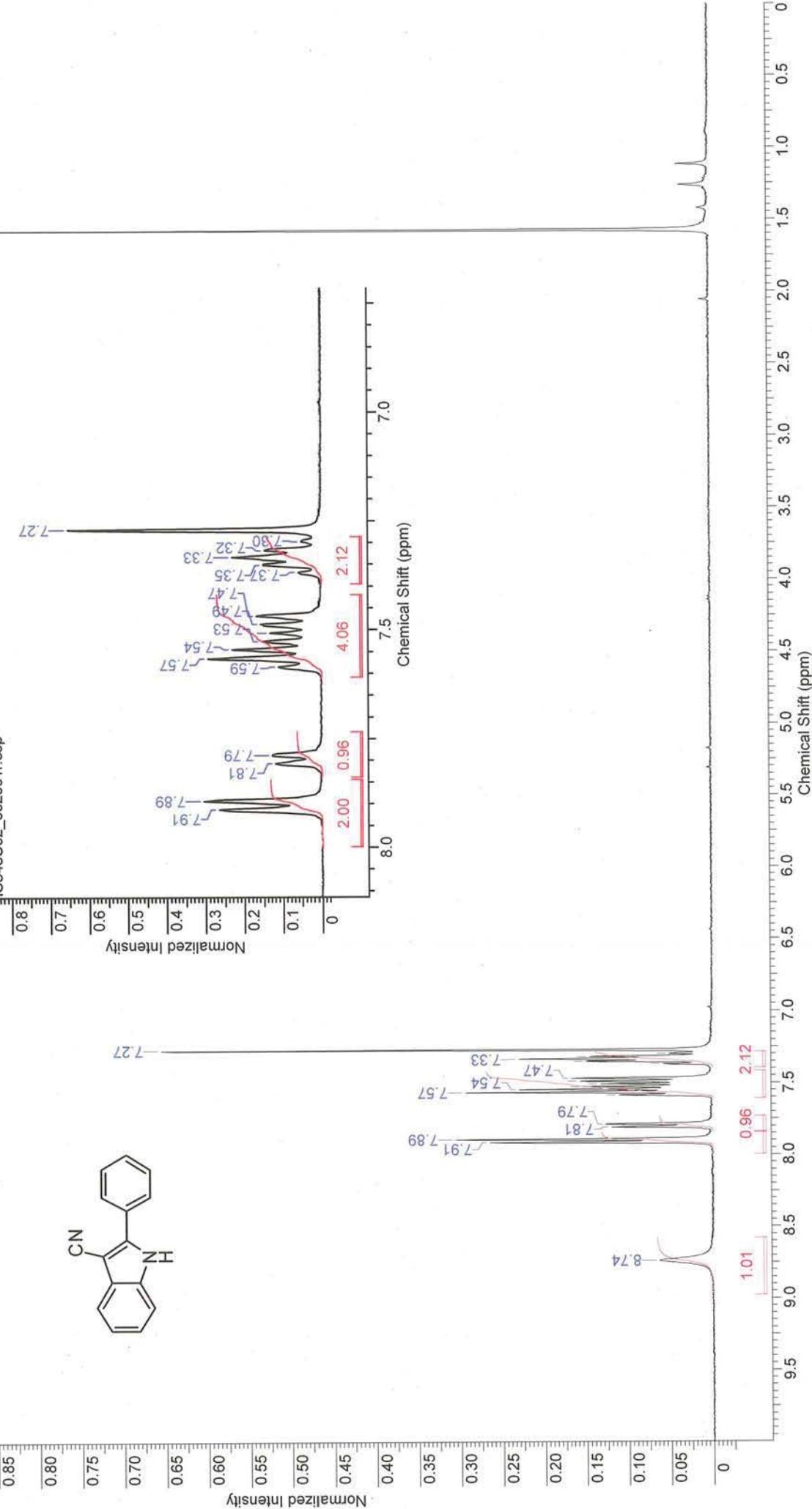
1	18	14	3	1	3	1	372.0817990	10.5	1.165e-03
2	20	13	3	1	3	0	372.0842043	13.5	1.240e-03
3	17	16	1	1	6	1	372.0853865	9.5	2.423e-03
4	15	13	3	3	5	0	372.0801816	9.5	2.782e-03
5	18	12	2	3	4	0	372.0790387	13.5	3.925e-03
6	19	15	1	1	6	0	372.0877918	12.5	4.828e-03
7	20	14	1	2	3	1	372.0880666	14.0	5.103e-03
8	13	14	3	3	5	1	372.0777763	6.5	5.188e-03
9	17	15	2	2	4	1	372.0892095	10.0	6.245e-03
10	16	13	2	3	4	1	372.0766334	10.5	6.331e-03



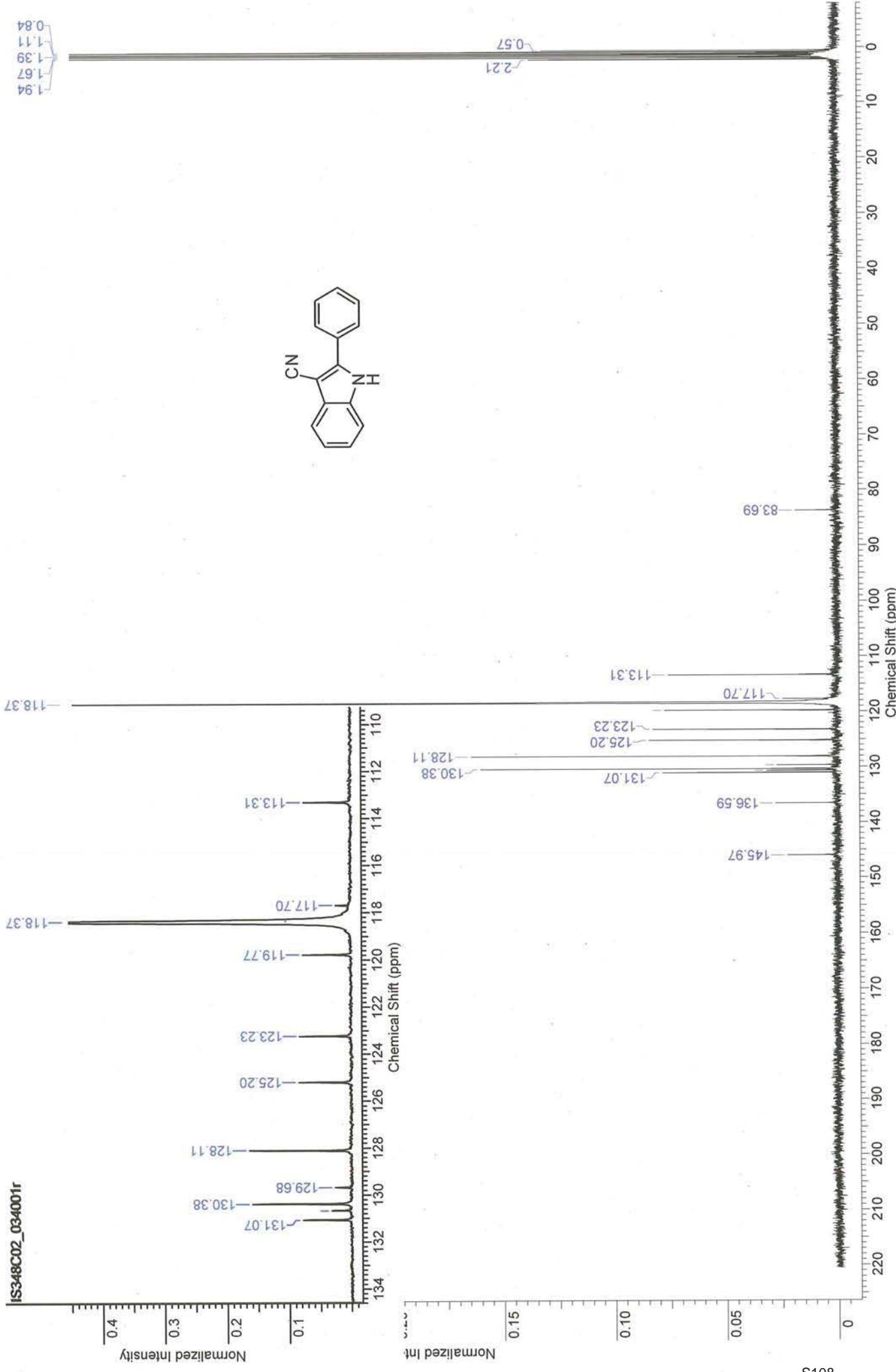
IS348C02_002001r.esp

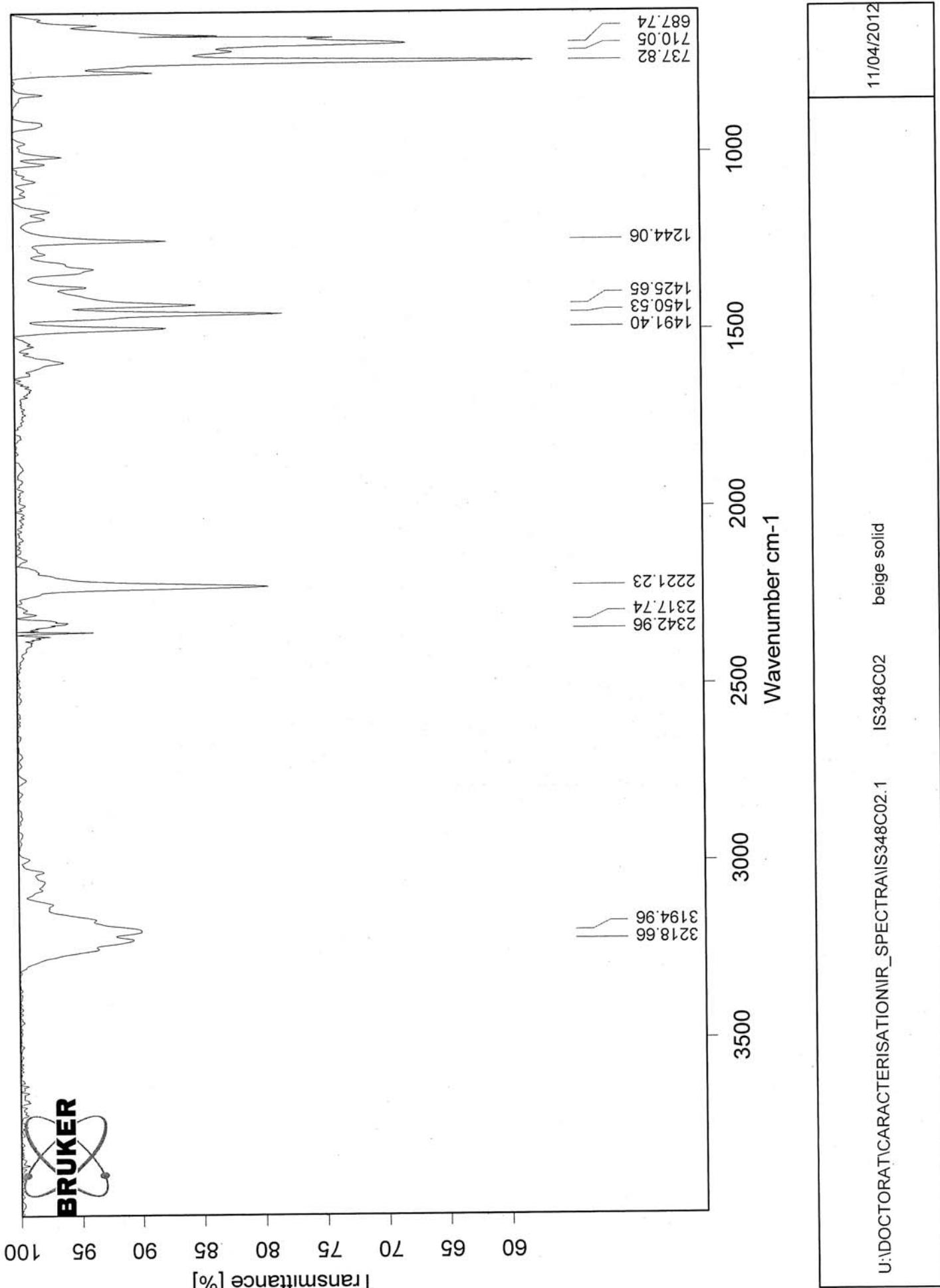


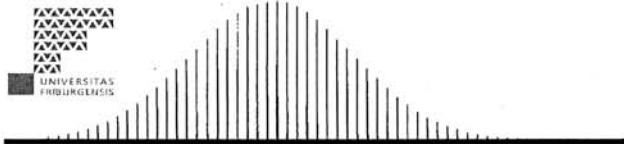
IS348C02_002001r.esp



IS348C02_034001r







FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

ESI-MS: IS348C01

XMASS Mass Analysis for /Data/UNI_FR/ALIM6535_ESI/10/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

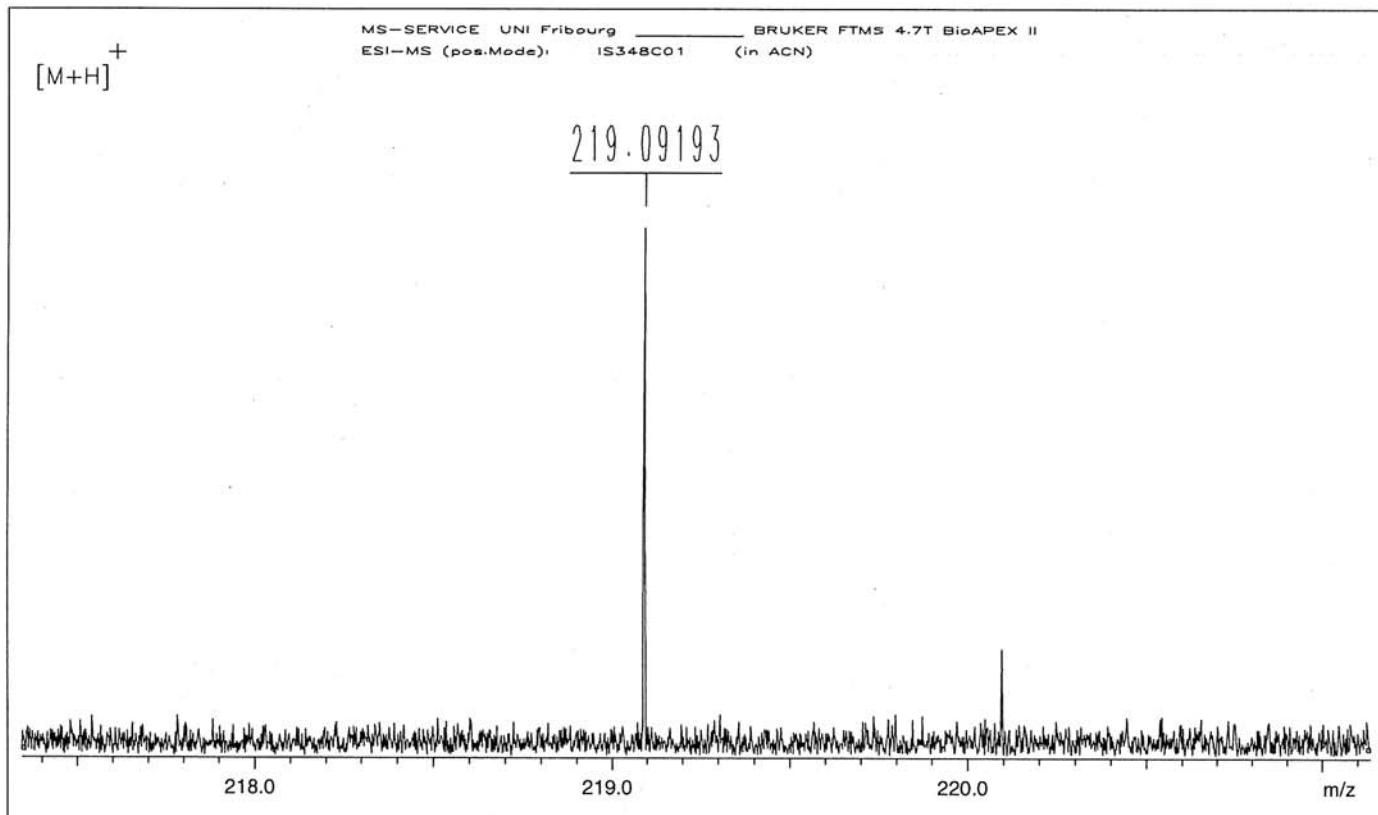
Ion mass = 219.0919320

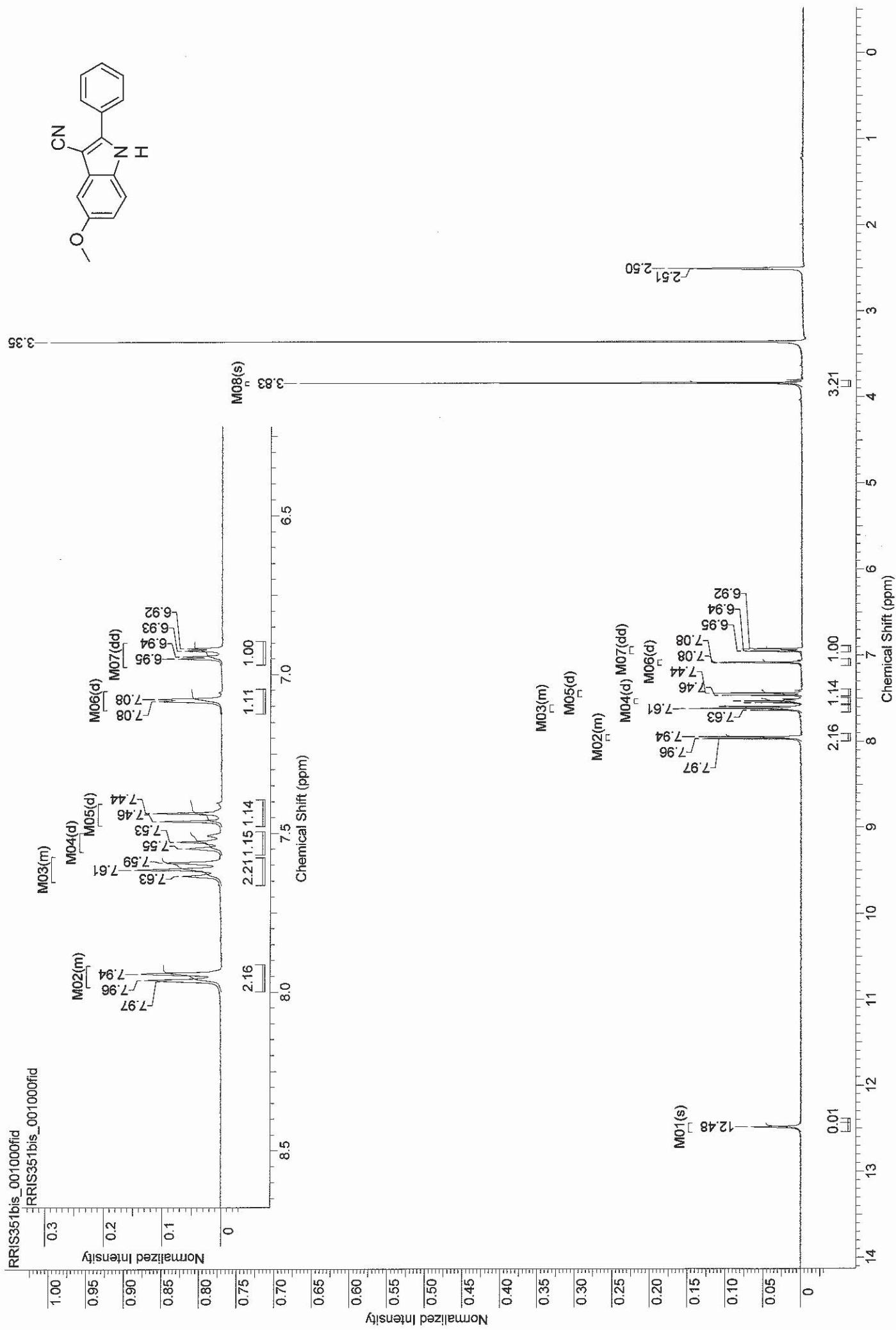
Charge = +1

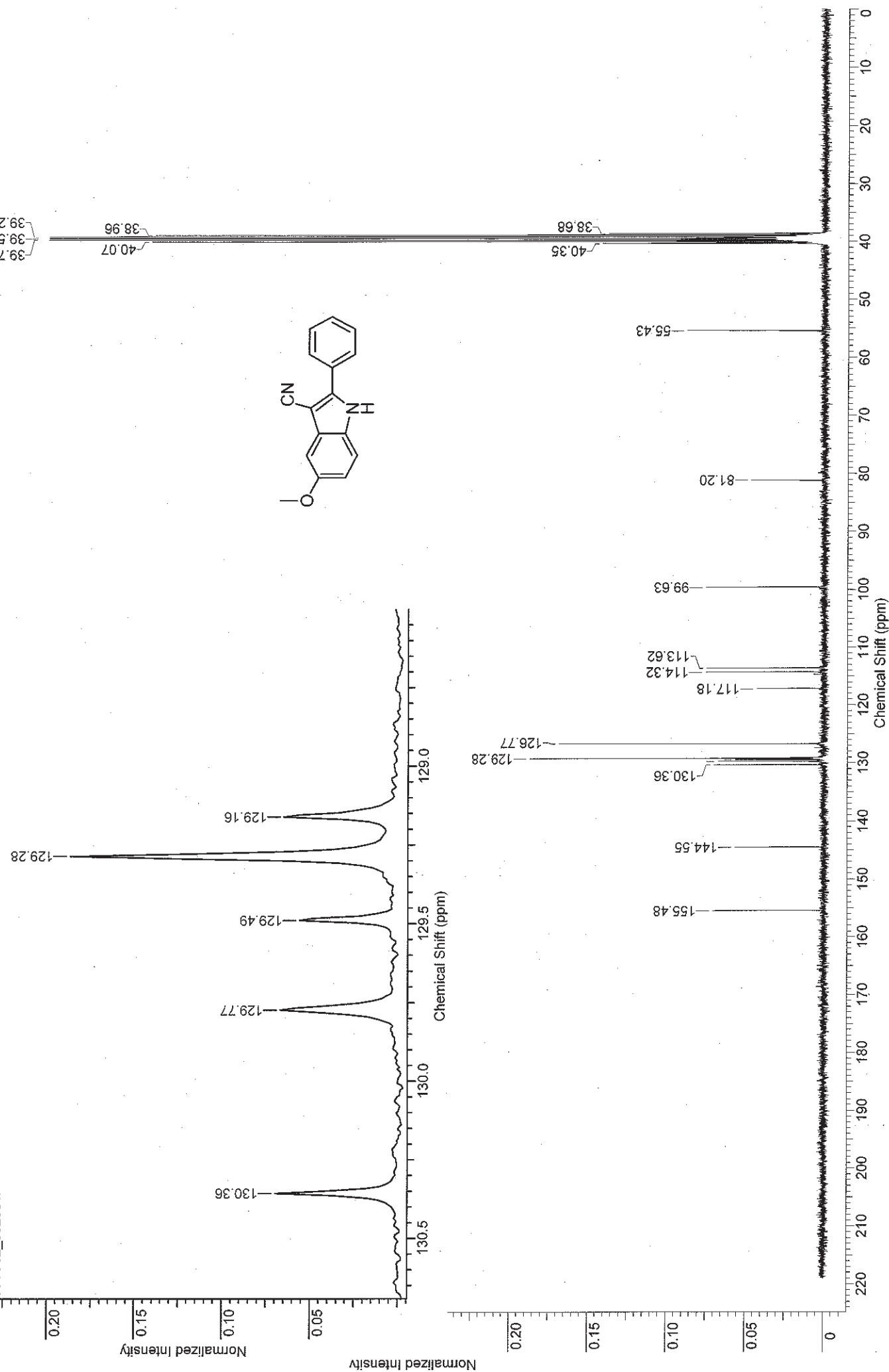
#	C	H	N	mass	DBE	error
---	---	---	---	------	-----	-------

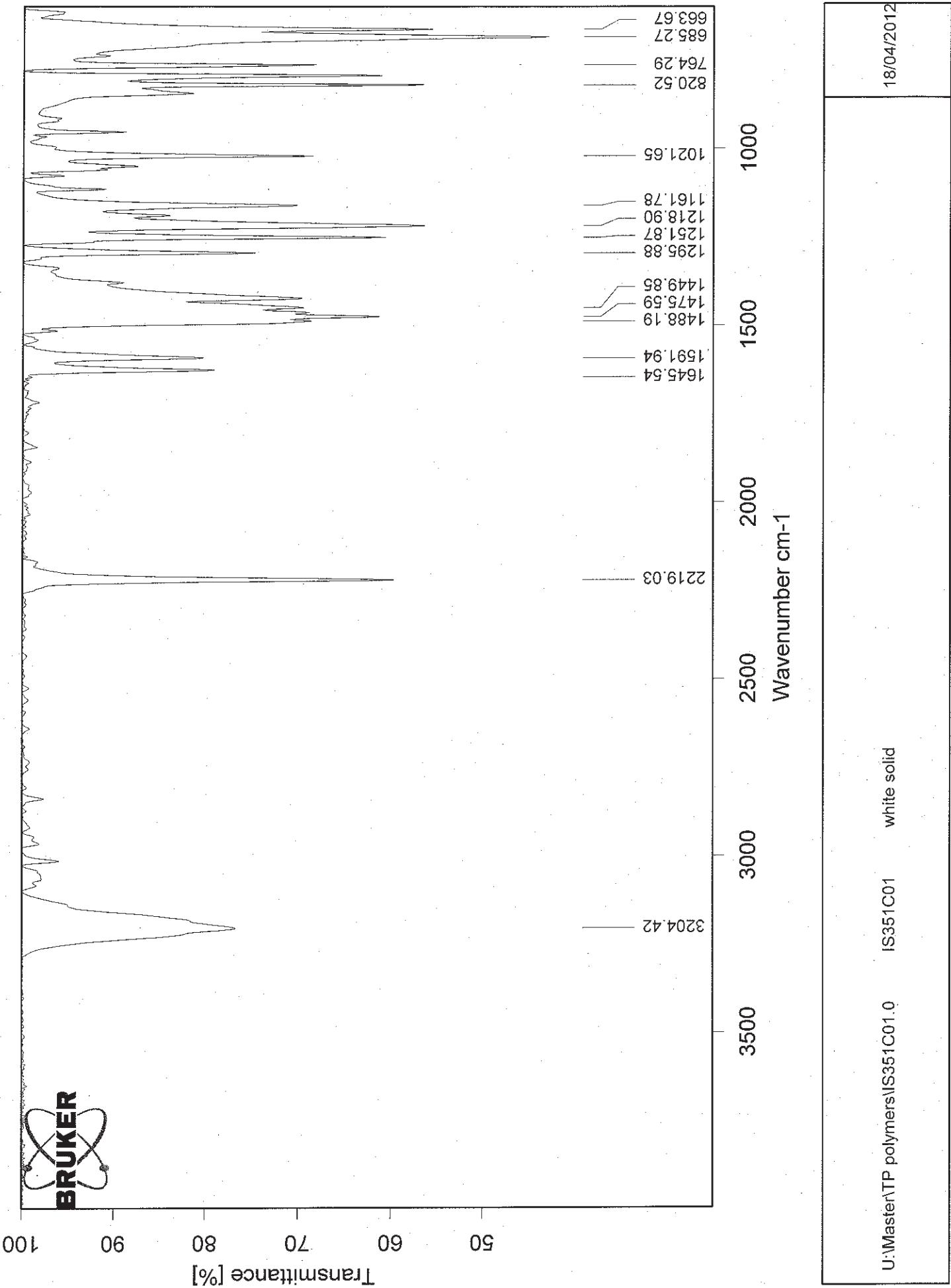
*** Mass Analysis for mass 219.0919320

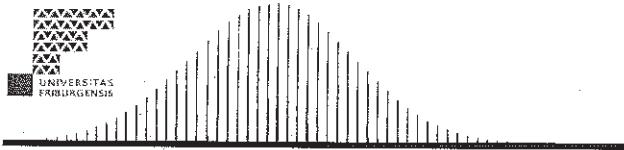
1	15	11	2	219.0916748	11.5	2.572e-04
2	7	9	9	219.0975428	8.0	5.611e-03
3	6	7	10	219.0849667	8.5	6.965e-03
4	16	13	1	219.1042508	11.0	1.232e-02
5	14	9	3	219.0790987	12.0	1.283e-02
6	8	11	8	219.1101188	7.5	1.819e-02
7	13	7	4	219.0665227	12.5	2.541e-02
8	9	13	7	219.1226949	7.0	3.076e-02
9	12	5	5	219.0539466	13.0	3.799e-02
10	10	15	6	219.1352709	6.5	4.334e-02











FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

ESI-MS: IS351C02

XMASS Mass Analysis for /Data/UNI_FR/ALIM6555_ESI/4/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

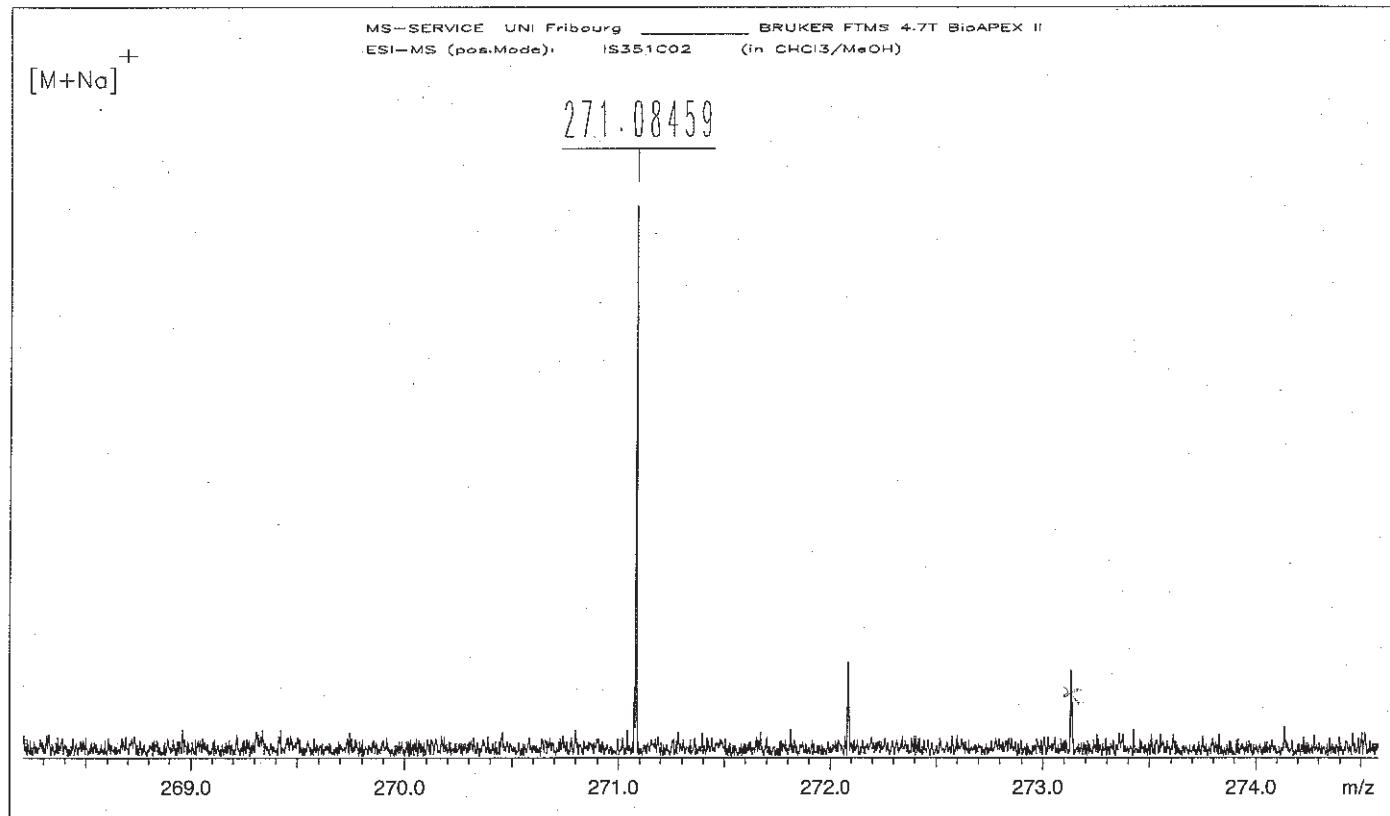
Ion mass = 271.0845900

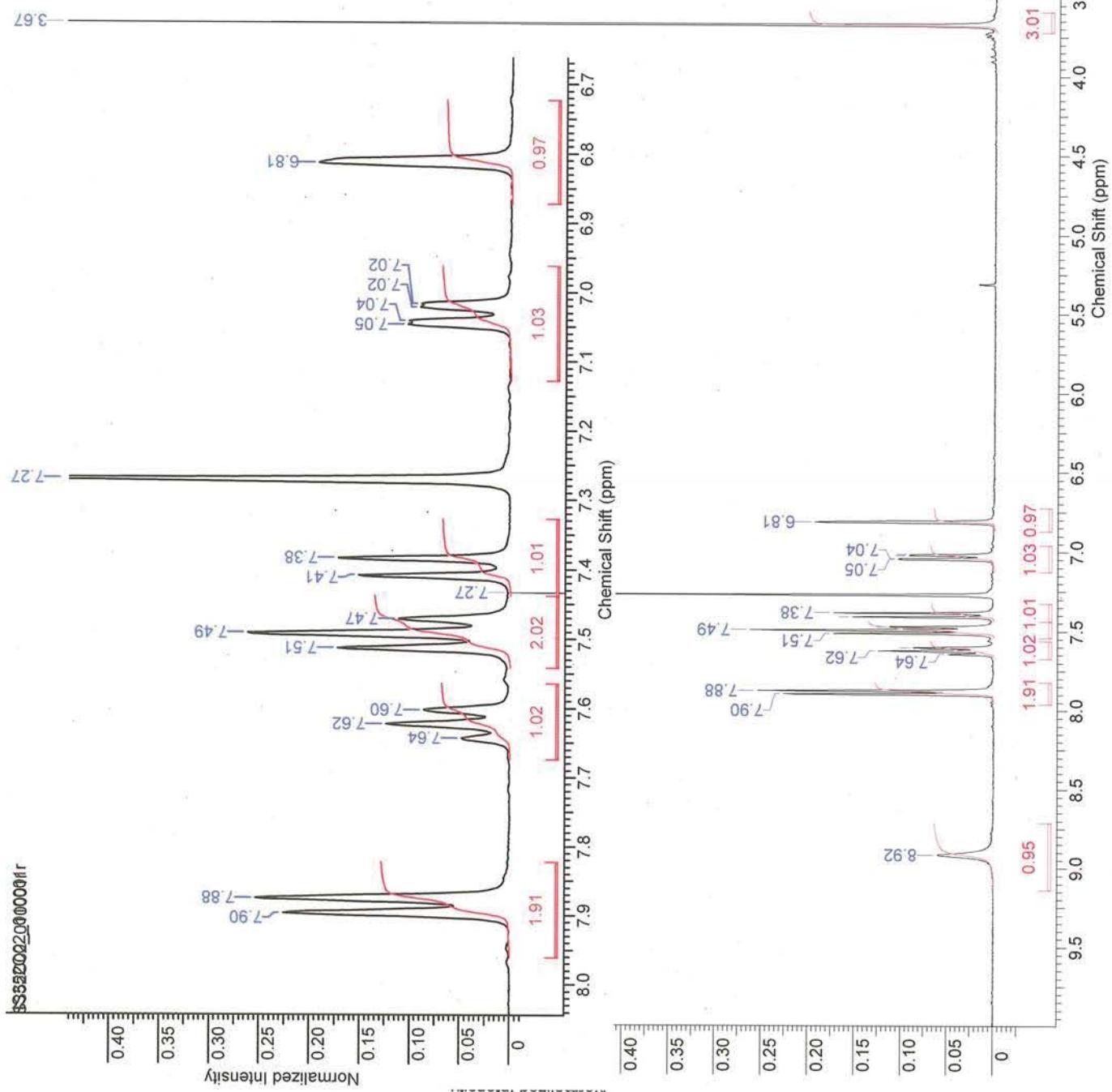
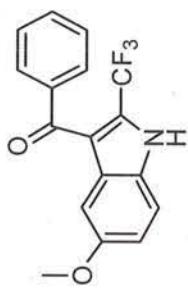
Charge = +1

#	C	H	N	O	Na	mass	DBE	error
---	---	---	---	---	----	------	-----	-------

*** Mass Analysis for mass 271.0845900

1	16	12	2	1	1	271.0841841	11.5	4.059e-04
2	15	13	1	4	0	271.0839093	10.0	6.807e-04
3	18	11	2	1	0	271.0865894	14.5	1.999e-03
4	13	11	4	3	0	271.0825667	10.5	2.023e-03
5	5	12	8	4	1	271.0873720	3.5	2.782e-03
6	6	18	1	9	1	271.0873773	-2.0	2.787e-03
7	13	14	1	4	1	271.0815040	7.0	3.086e-03
8	11	9	7	2	0	271.0812240	11.0	3.366e-03
9	6	15	4	8	0	271.0884399	1.5	3.850e-03
10	7	14	5	5	1	271.0887147	3.0	4.125e-03

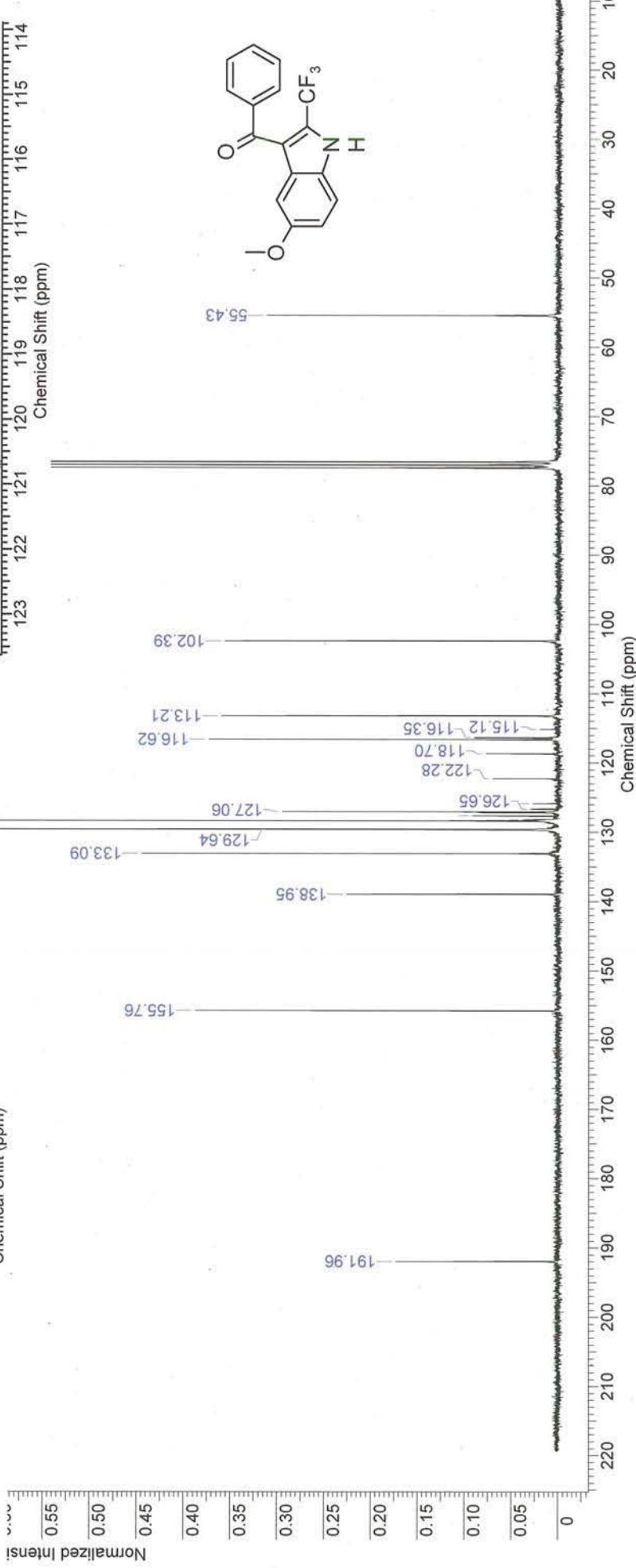
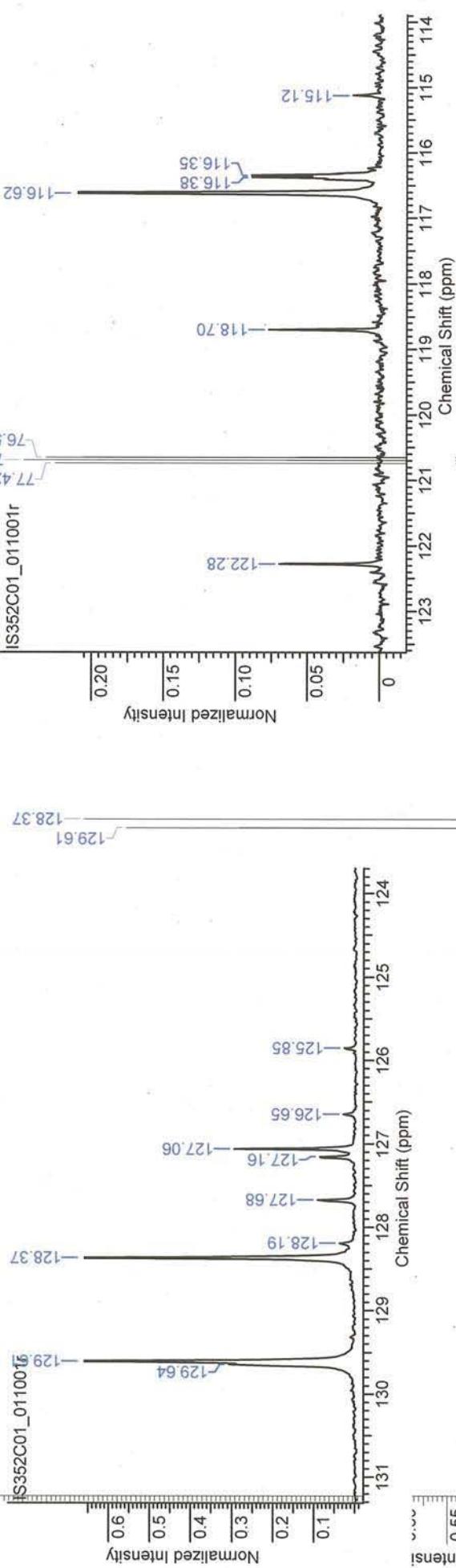


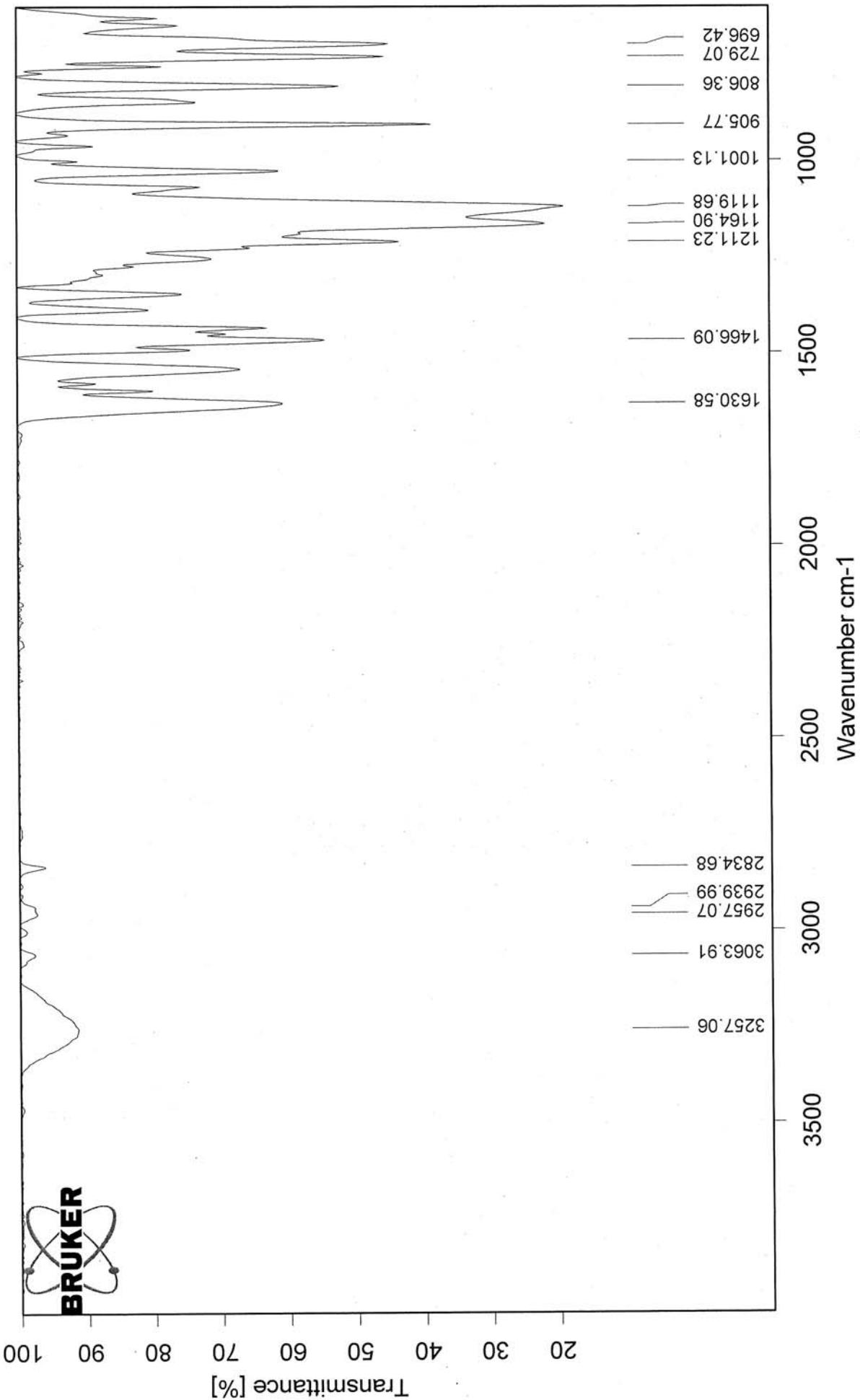


IS352C01_011001r

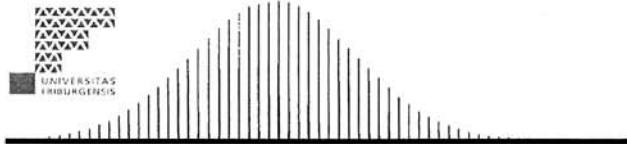
IS352C01_011001r

IS352C01_011001r





U:\DOCTORAT\CHARACTERISATION\IR_SPECTRA\IS352C01.0	IS352C01	brown liquid	19/04/2012
--	----------	--------------	------------



FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

ESI-MS: IS352C01

XMASS Mass Analysis for /Data/UNI_FR/ALIM6554_ESI/4/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

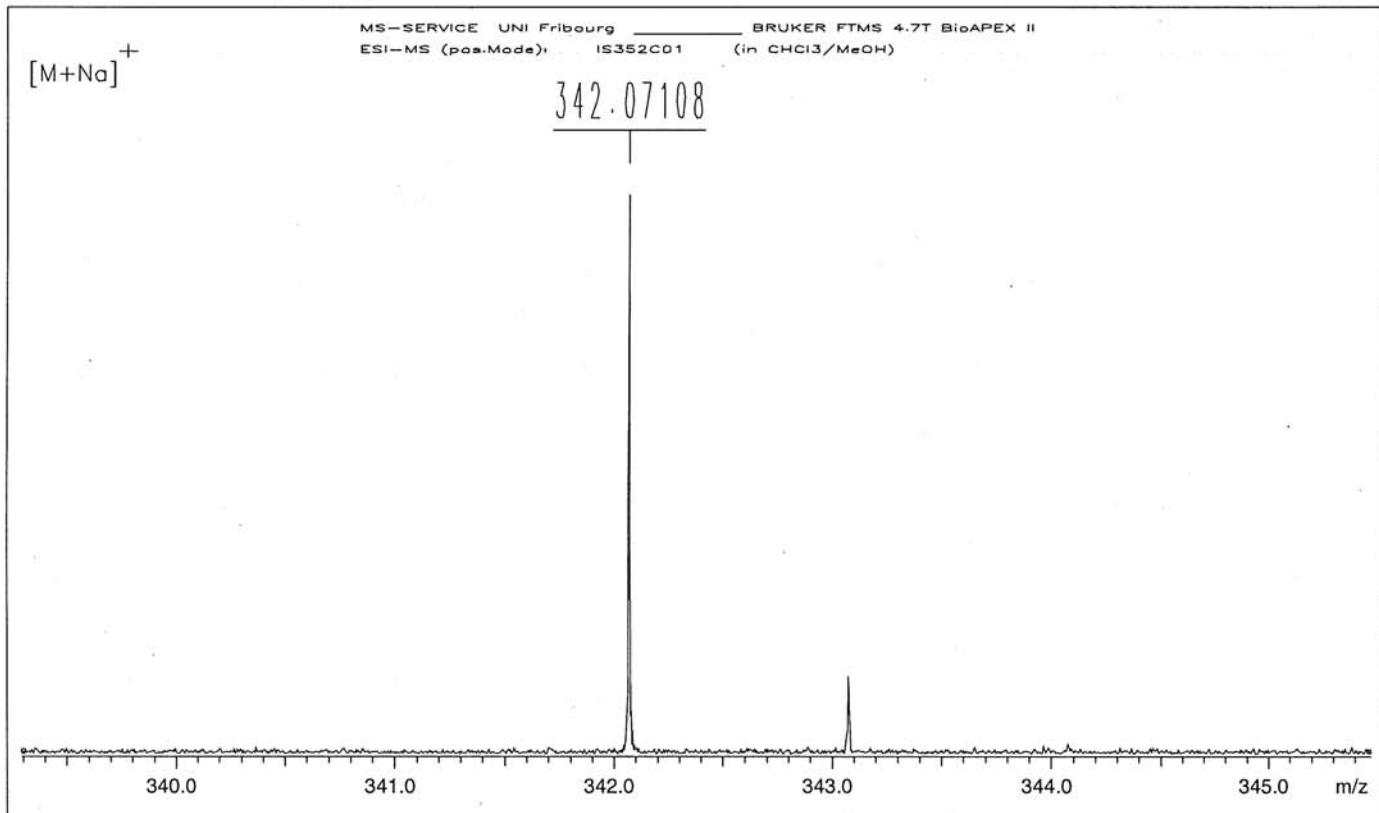
Ion mass = 342.0710770

Charge = +1

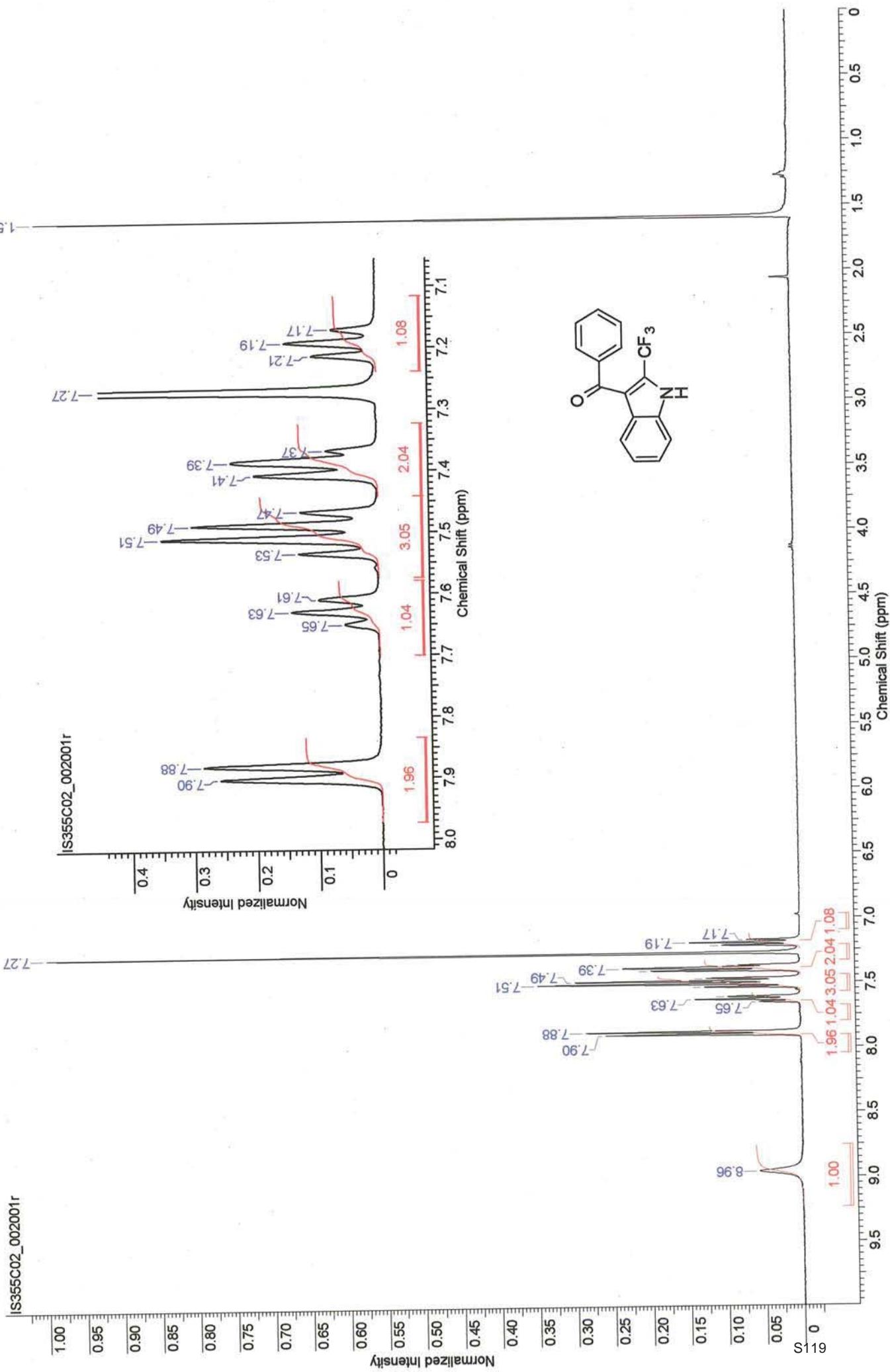
#	C	H	F	N	O	Na	mass	DBE	error
---	---	---	---	---	---	----	------	-----	-------

*** Mass Analysis for mass 342.0710770

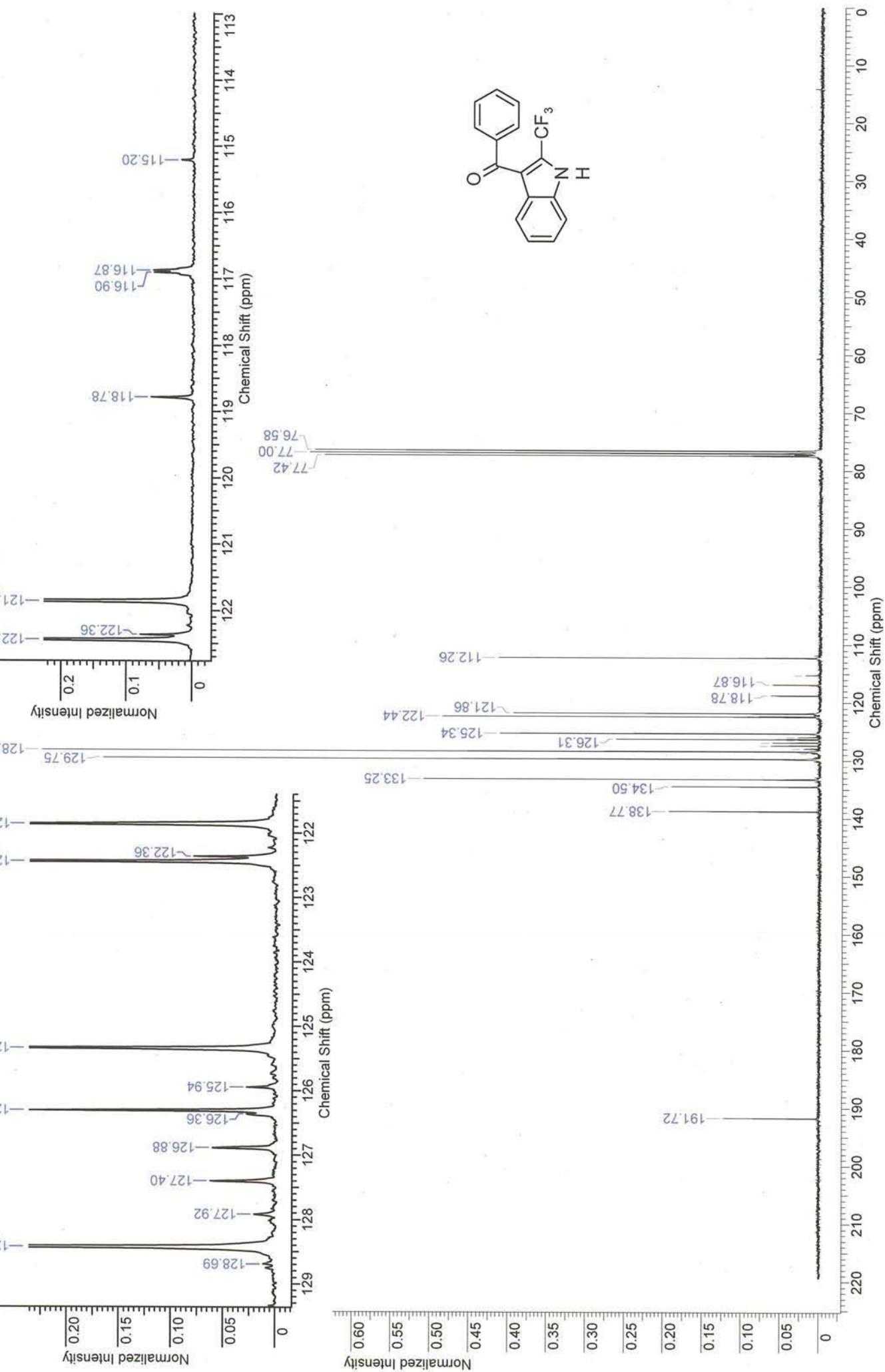
1	17	12	3	1	2	1	342.0712343	10.5	1.573e-04
2	11	12	4	3	5	0	342.0707597	5.5	3.173e-04
3	20	11	2	1	1	1	342.0700915	14.5	9.855e-04
4	14	13	4	1	3	1	342.0723772	6.5	1.300e-03
5	22	10	2	1	1	0	342.0724968	17.5	1.420e-03
6	14	11	3	3	4	0	342.0696169	9.5	1.460e-03
7	11	14	5	1	4	1	342.0735200	2.5	2.443e-03
8	19	11	3	1	2	0	342.0736396	13.5	2.563e-03
9	17	10	2	3	3	0	342.0684740	13.5	2.603e-03
10	9	13	4	3	5	1	342.0683544	2.5	2.723e-03

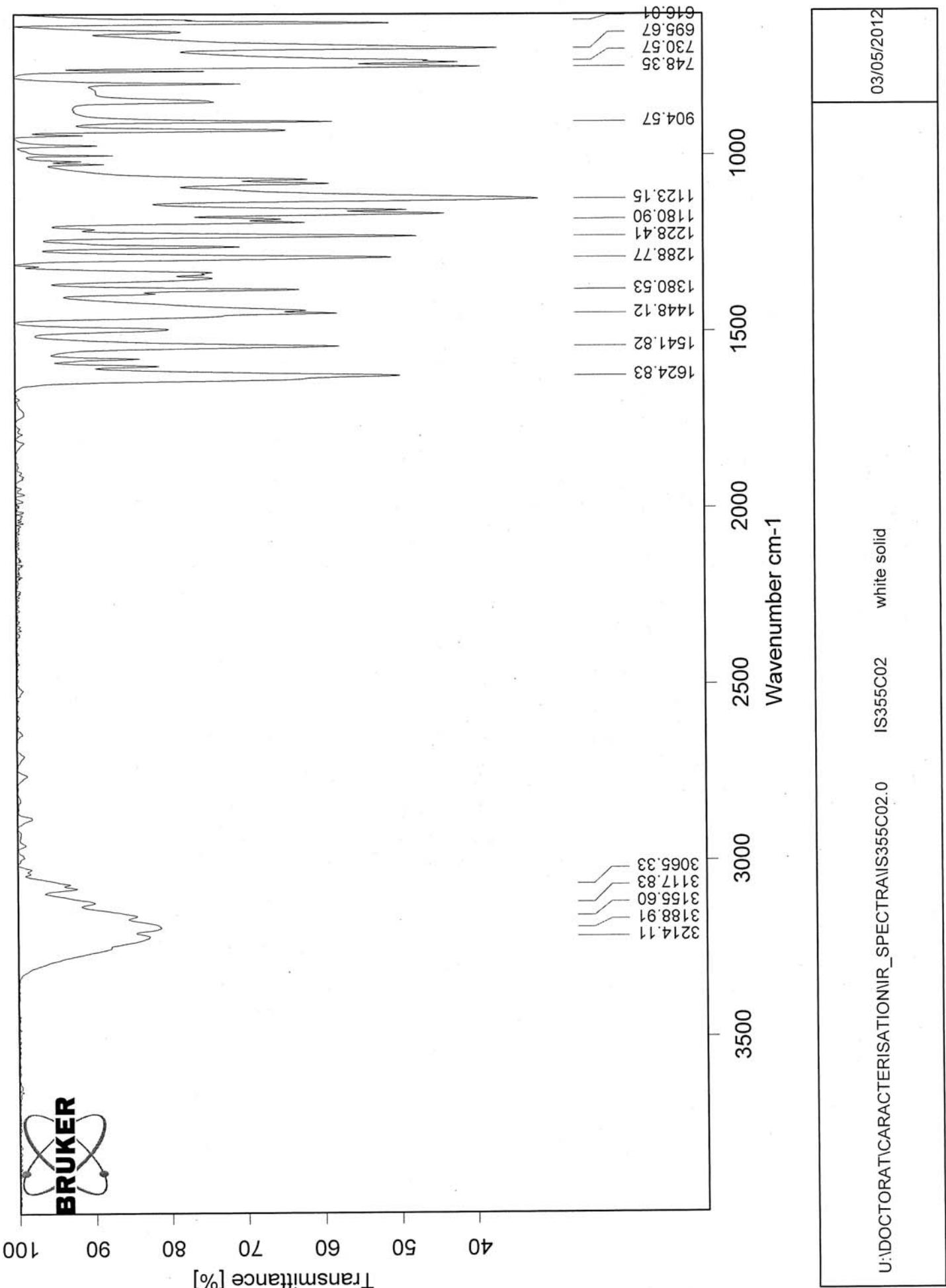


03.05.2012 10:22:15



J3355C02_002001
J3355C02_002001

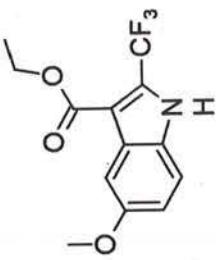
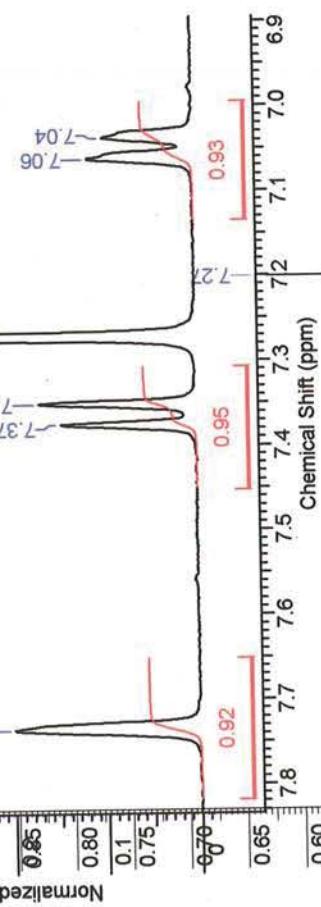


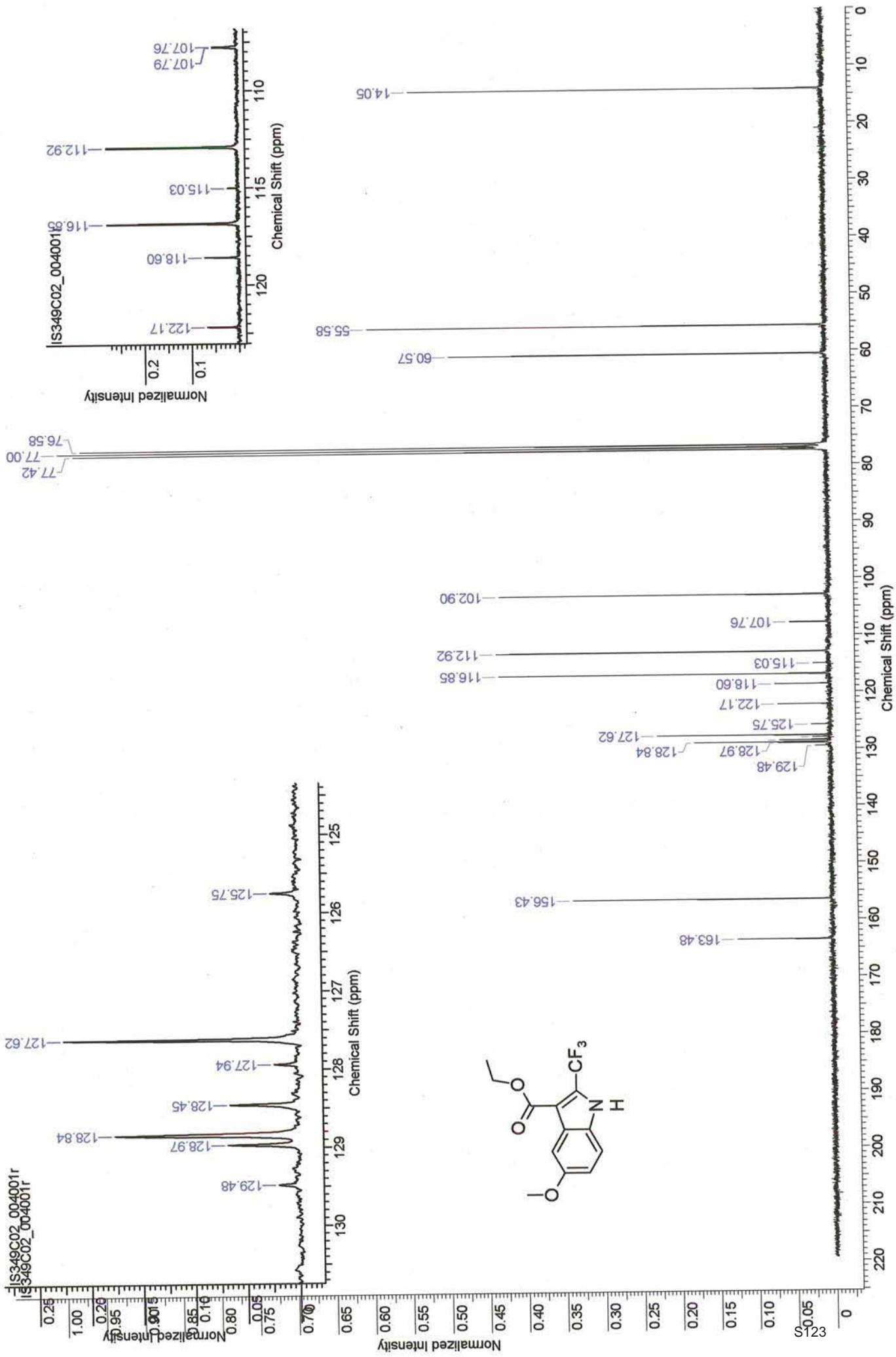


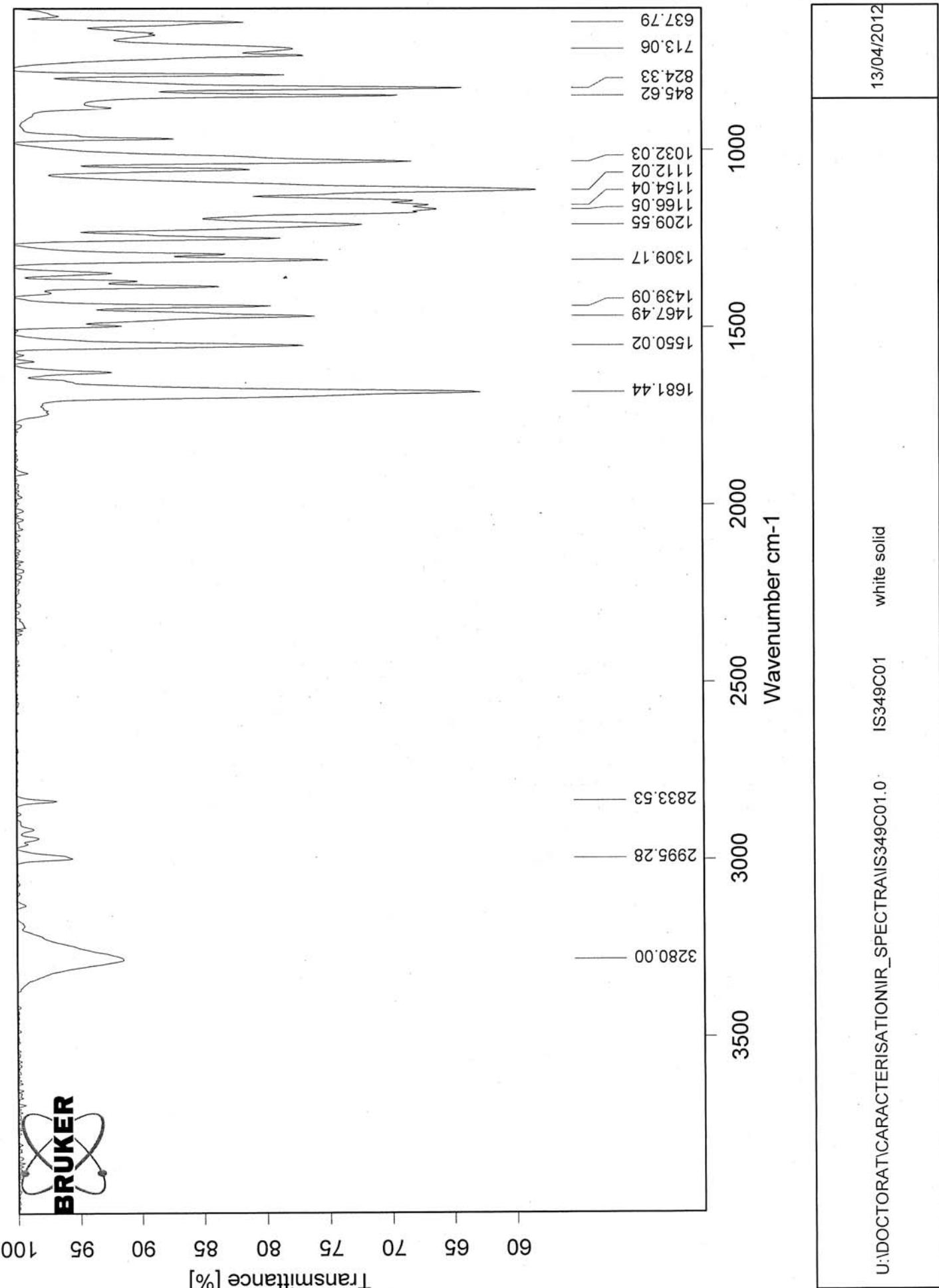
03.05.2012 07:09:37

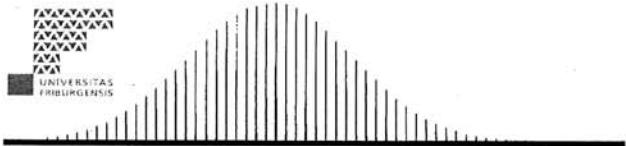
IS349C01_001001r.esp

IS349C01_001001r.esp









FTMS 4.7T BioAPEX II MS-Service UNI-Fribourg

ESI-MS: IS349C01

XMASS Mass Analysis for /Data/UNI_FR/ALIM6545_ESI/1/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

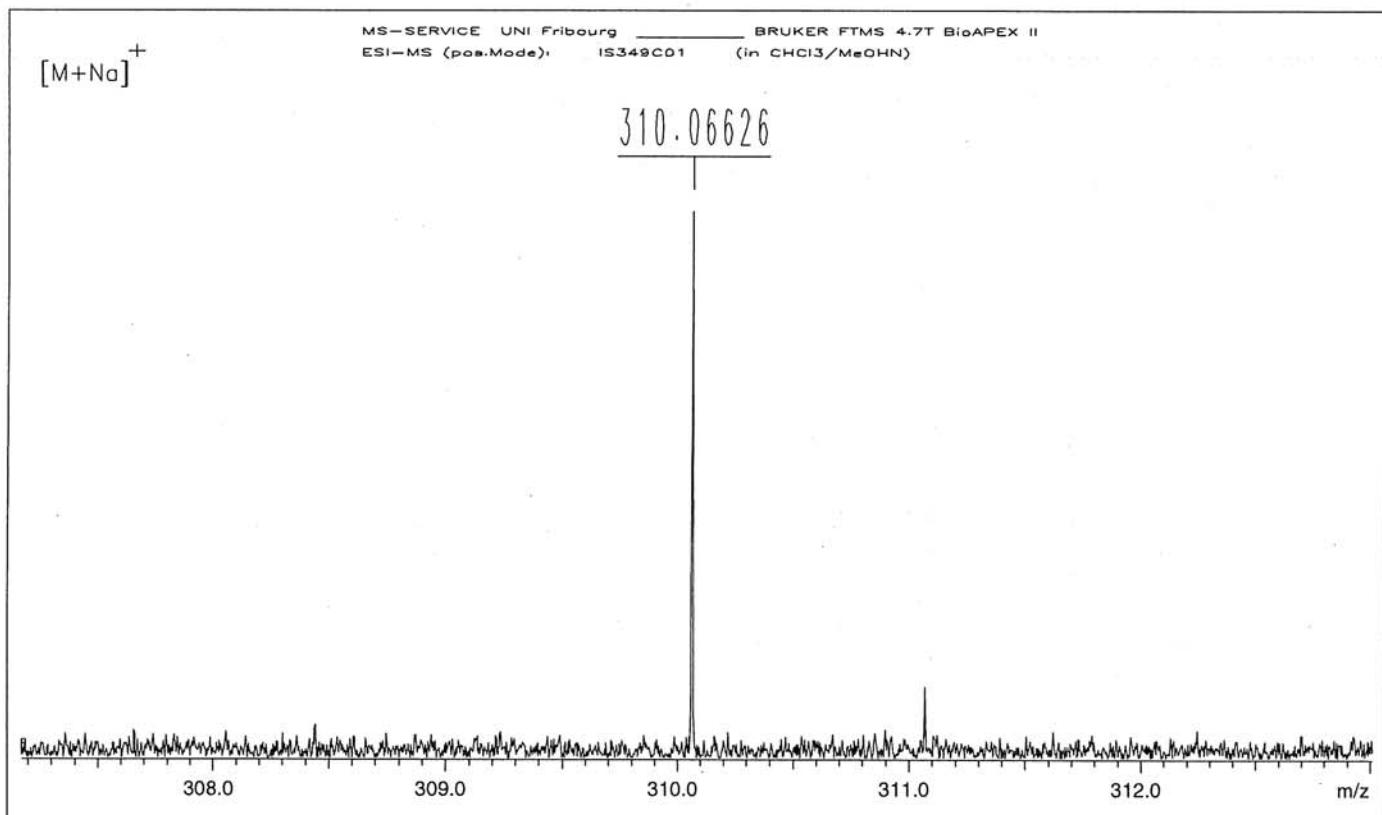
Ion mass = 310.0662590

Charge = +1

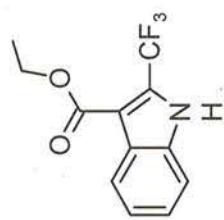
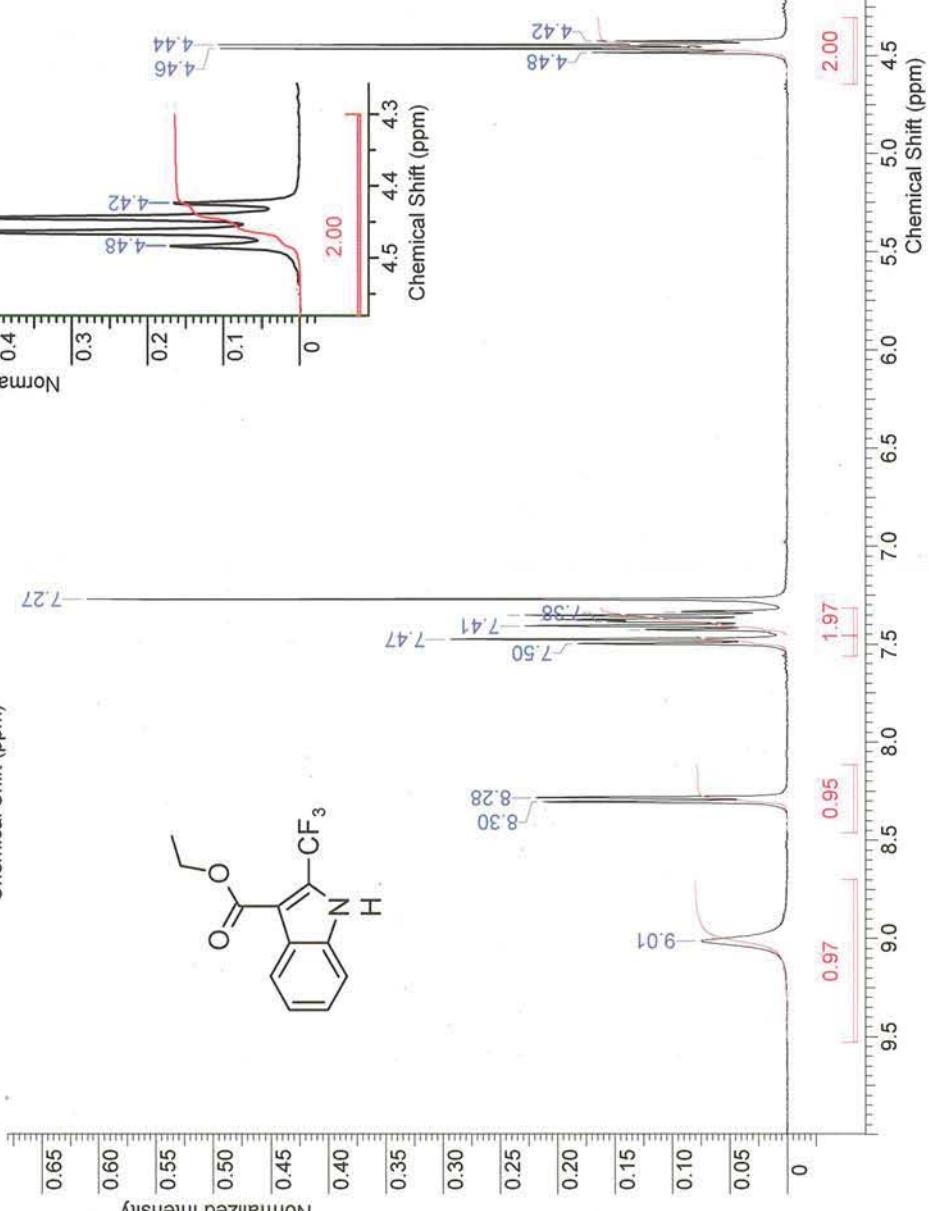
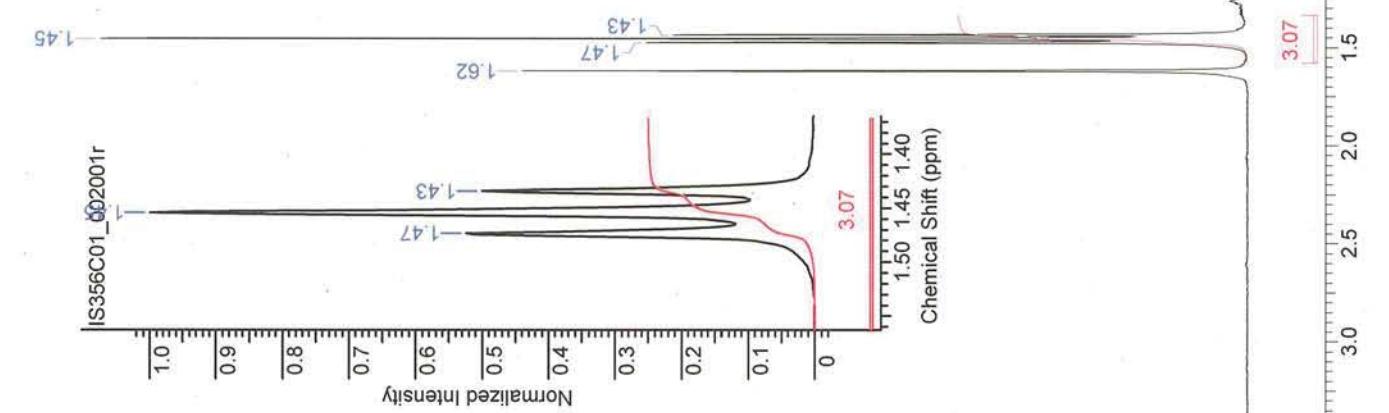
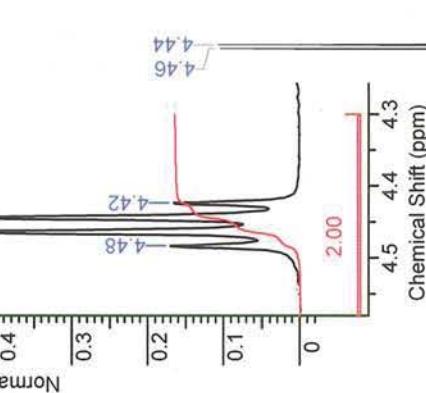
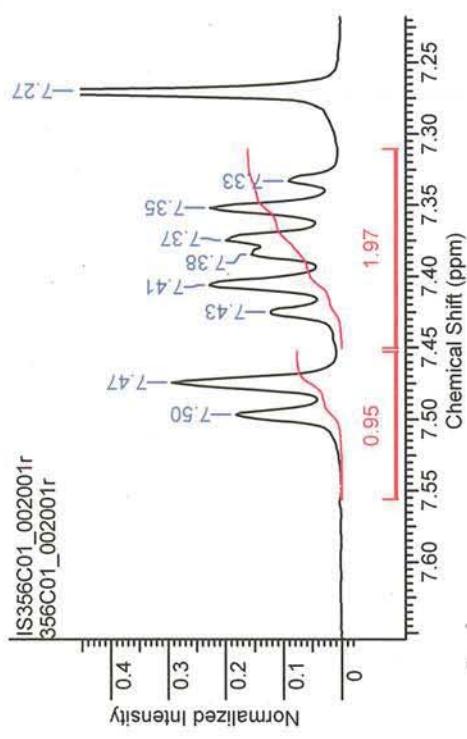
#	C	H	F	N	O	Na	mass	DBE	error
---	---	---	---	---	---	----	------	-----	-------

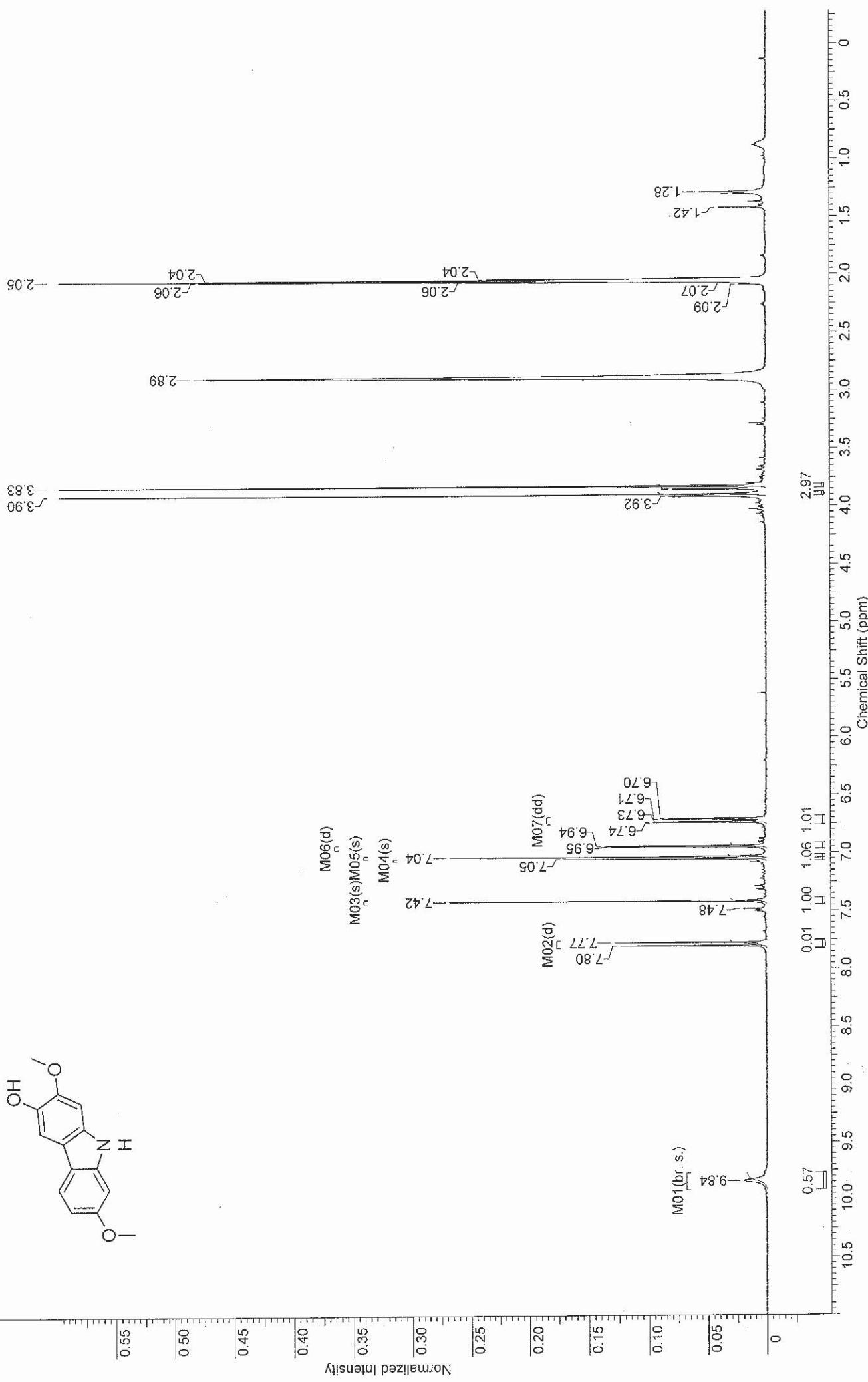
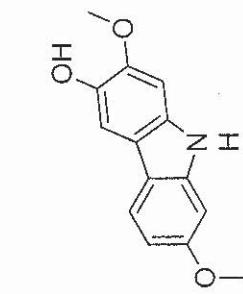
*** Mass Analysis for mass 310.0662590

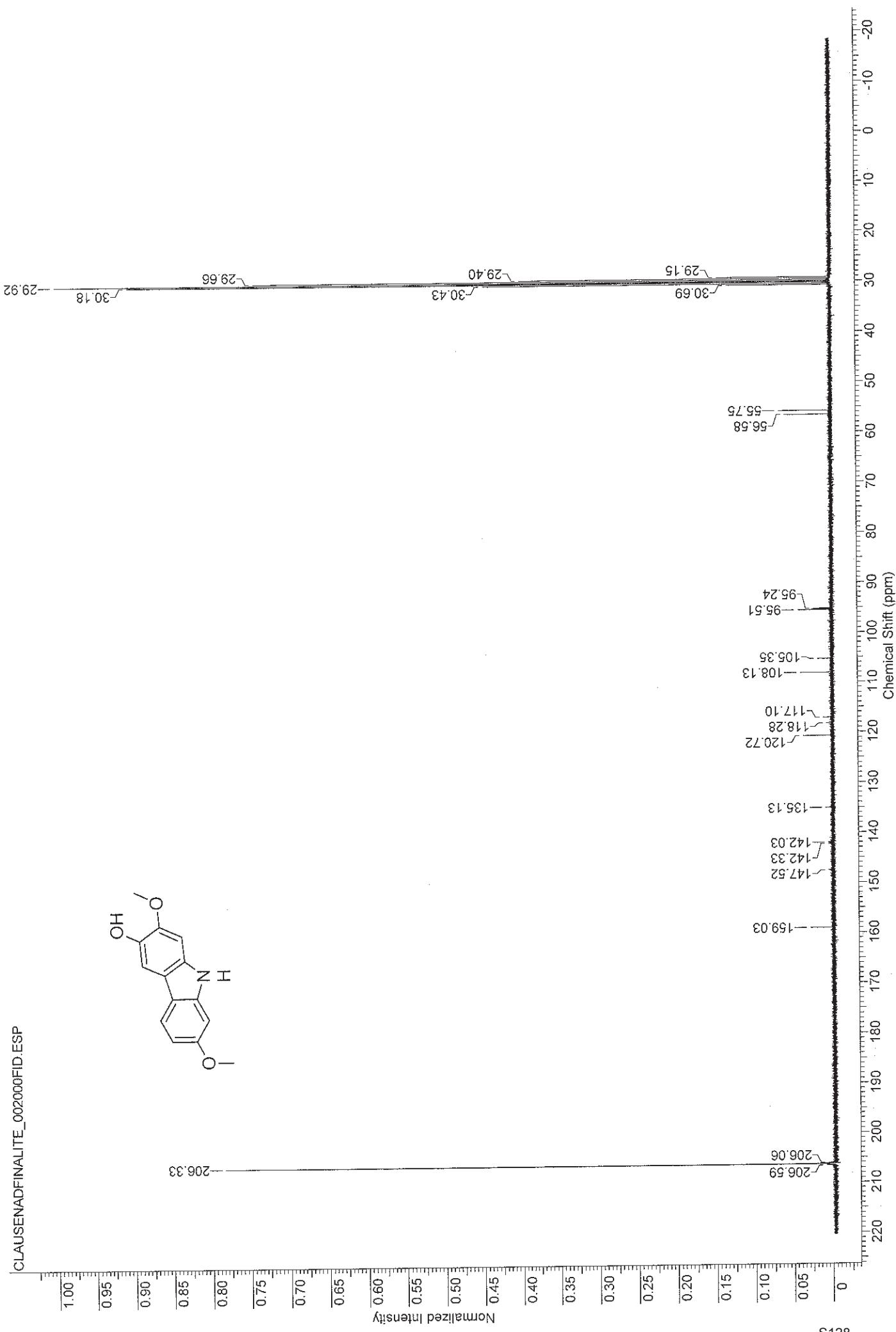
1	13	12	3	1	3	1	310.0661489	6.5	1.101e-04
2	13	10	6	1	1	0	310.0661096	6.5	1.494e-04
3	16	8	2	4	1	0	310.0660687	14.0	1.903e-04
4	8	11	4	4	3	1	310.0659491	3.0	3.099e-04
5	7	12	4	3	6	0	310.0656744	1.5	5.846e-04
6	5	12	5	4	4	1	310.0670920	-1.0	8.330e-04
7	13	9	3	4	2	0	310.0672116	10.0	9.526e-04
8	10	13	4	1	4	1	310.0672918	2.5	1.033e-03
9	18	10	2	1	2	0	310.0674114	13.5	1.152e-03
10	16	11	2	1	2	1	310.0650061	10.5	1.253e-03

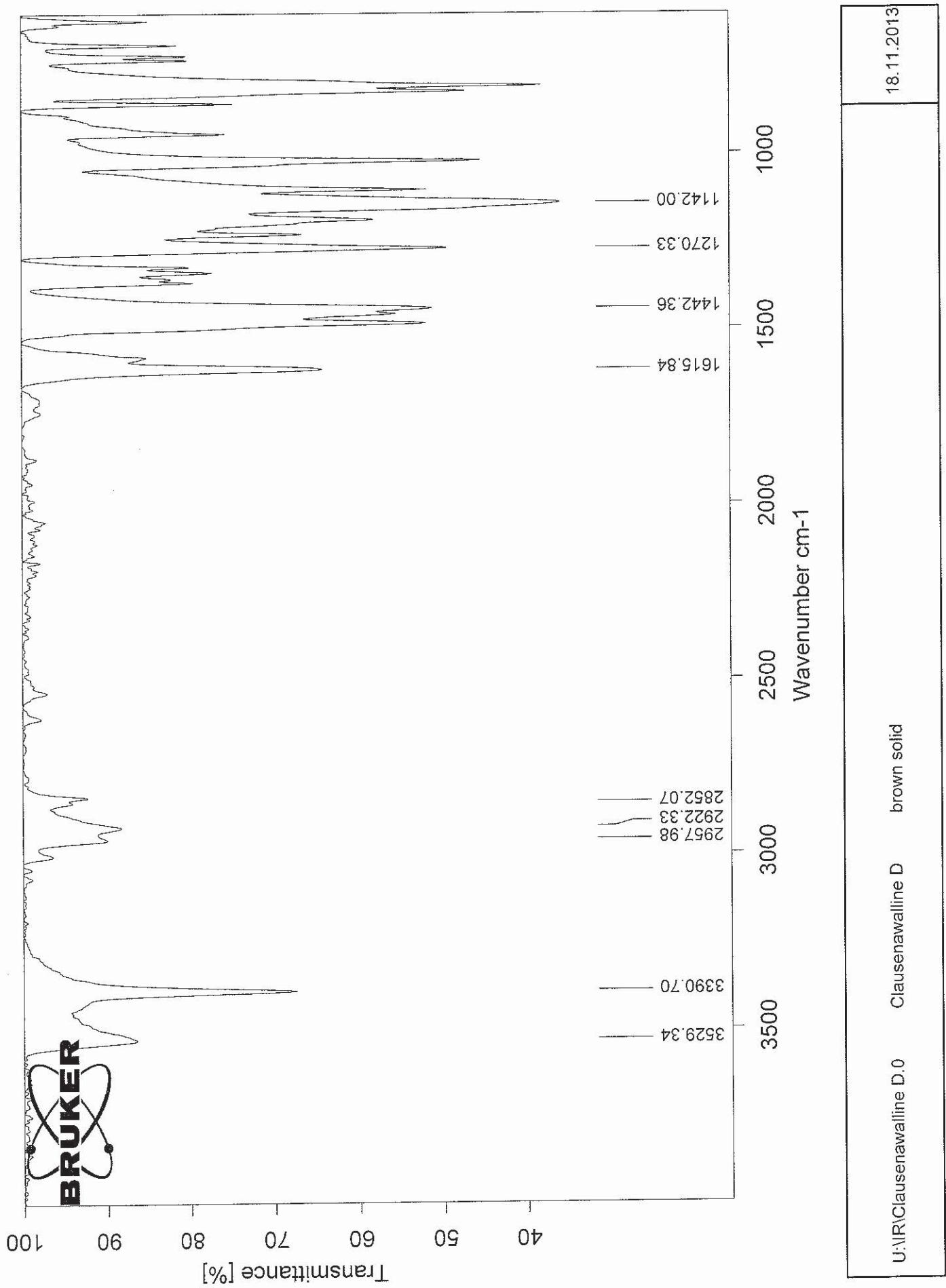


IS356C01_002001r
IS356C01_002001r









ESI-MS: RR073A

XMASS Mass Analysis for /Data/UNI_FR/REMY7660_ESI/1/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

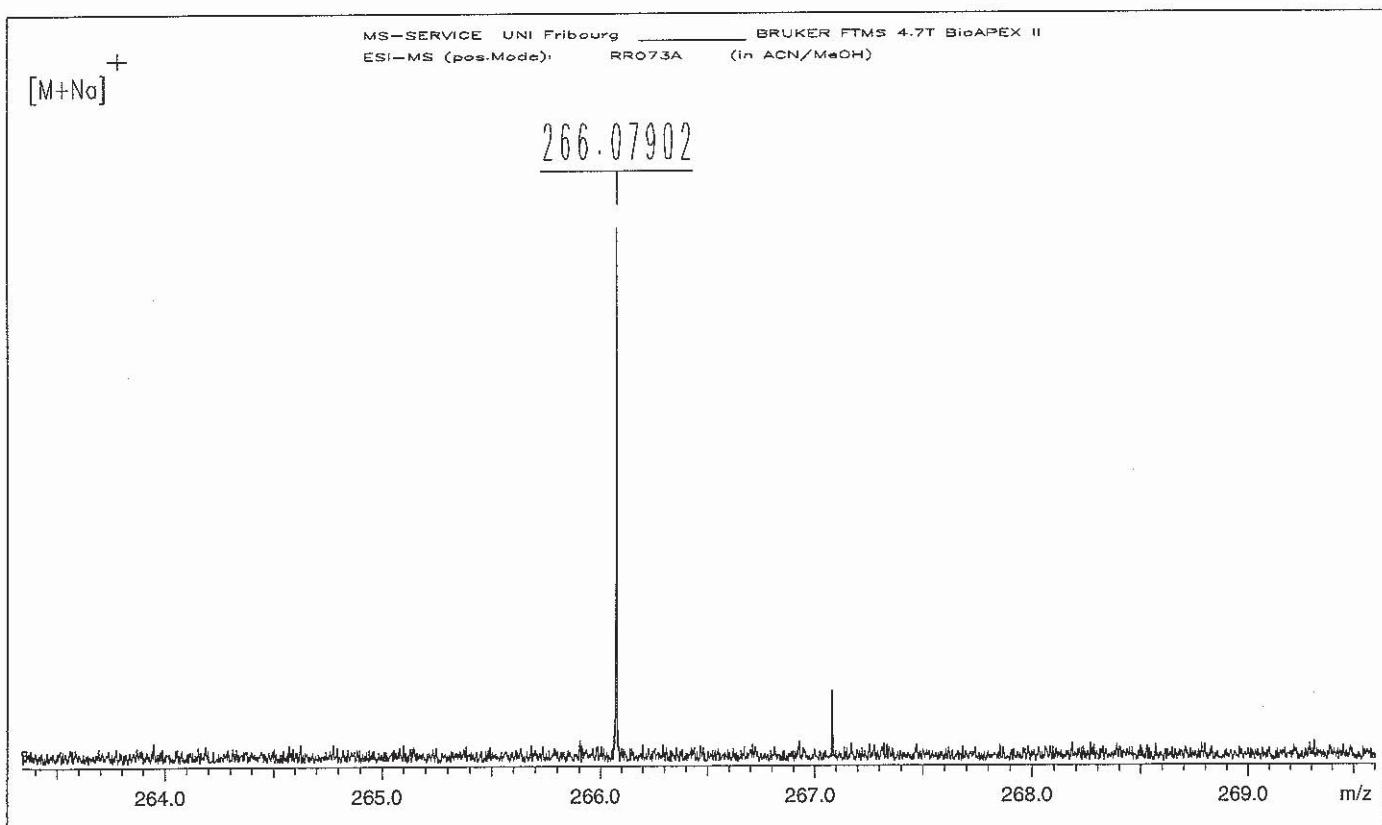
Ion mass = 266.0790200

Charge = +1

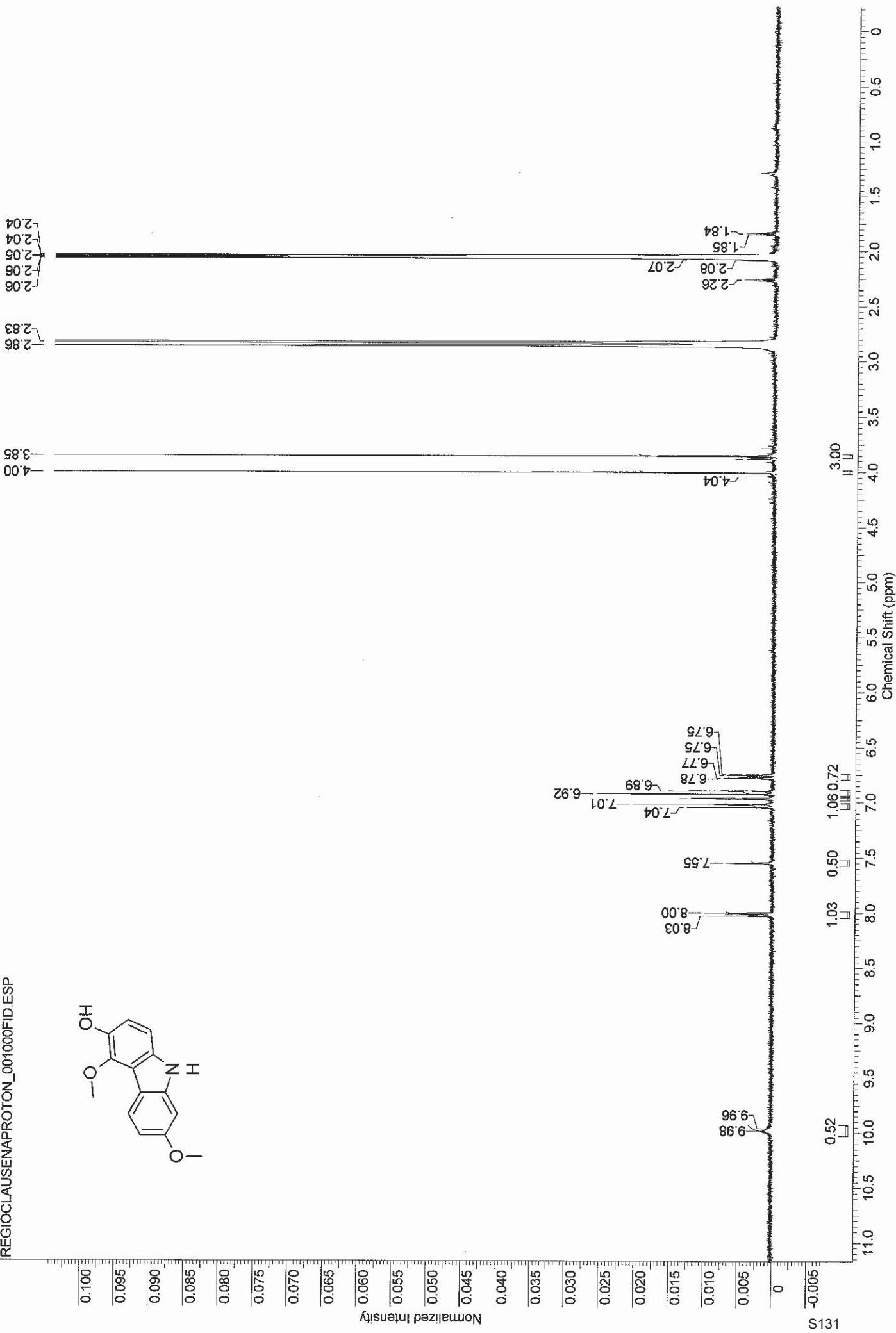
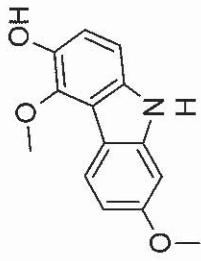
#	C	H	N	O	Na	mass	DBE	error
---	---	---	---	---	----	------	-----	-------

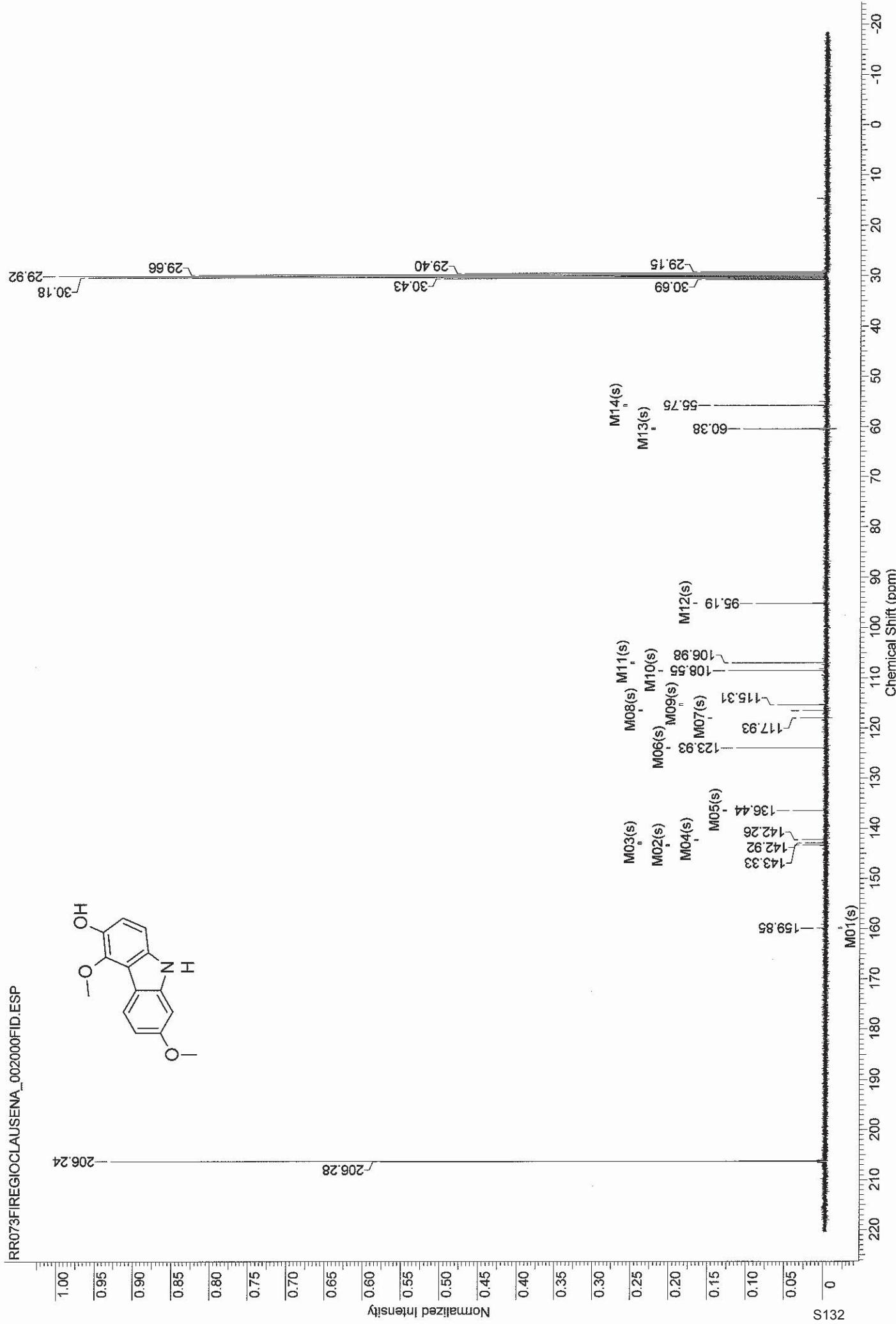
*** Mass Analysis for mass 266.0790200

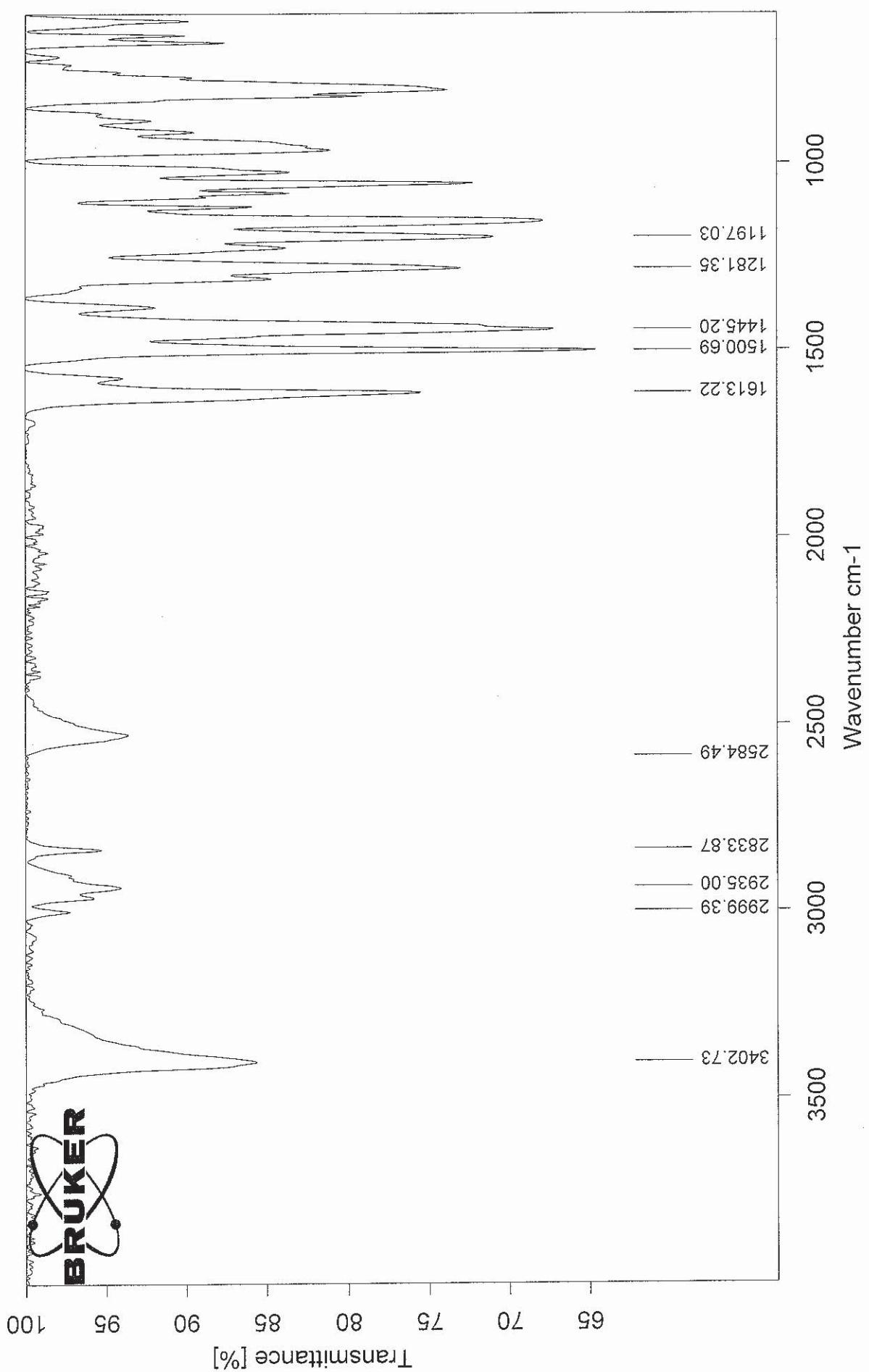
1	14	13	1	3	1	266.0787644	8.5	2.556e-04
2	12	8	7	1	0	266.0784843	12.5	5.357e-04
3	14	10	4	2	0	266.0798270	12.0	8.070e-04
4	12	11	4	2	1	266.0774217	9.0	1.598e-03
5	11	12	3	5	0	266.0771469	7.5	1.873e-03
6	16	12	1	3	0	266.0811697	11.5	2.150e-03
7	10	9	7	1	1	266.0760790	9.5	2.941e-03
8	9	10	6	4	0	266.0758043	8.0	3.216e-03
9	5	15	4	7	1	266.0832949	0.0	4.275e-03
10	9	13	3	5	1	266.0747416	4.5	4.278e-03



REGIOCLAUSENAPROTON_001000FID.ESP







U:\NIR\4,7-dimethoxy-9H-carbazol-3-ol 4,7-dimethoxy-9H-carbazol-3-ol brown solid

18.11.2013

ESI-MS: RR073B

XMASS Mass Analysis for /Data/UNI_FR/REMY7721_ESI/1/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

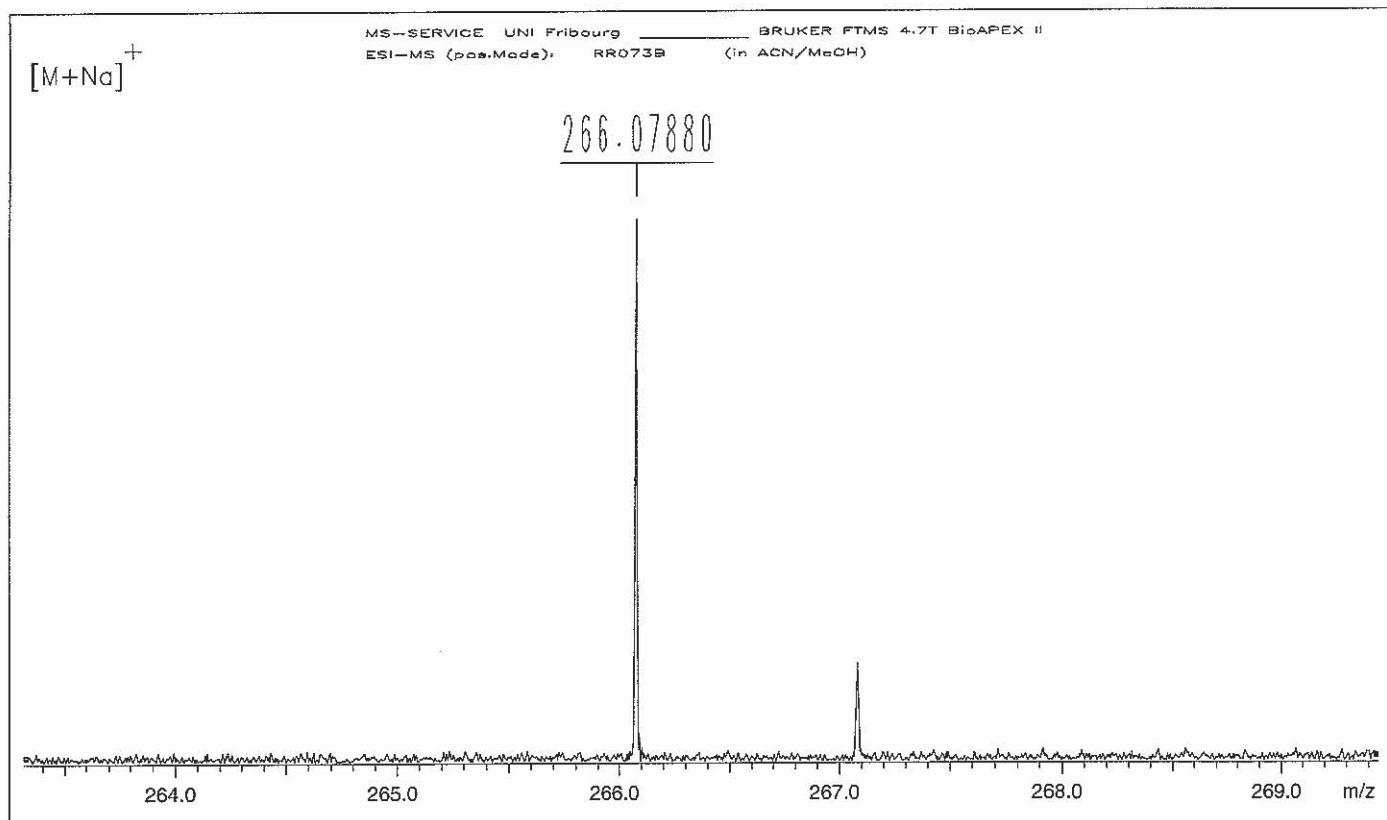
Ion mass = 266.0788000

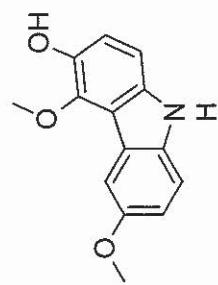
Charge = +1

C H N O Na mass DBE error

*** Mass Analysis for mass 266.0788000

1	14	13	1	3	1	266.0787644	8.5	3.562e-05
2	14	10	4	2	0	266.0798270	12.0	1.027e-03
3	12	11	4	2	1	266.0774217	9.0	1.378e-03
4	11	12	3	5	0	266.0771469	7.5	1.653e-03
5	16	12	1	3	0	266.0811697	11.5	2.370e-03
6	9	13	3	5	1	266.0747416	4.5	4.058e-03
7	5	15	4	7	1	266.0832949	0.0	4.495e-03
8	6	12	5	7	0	266.0731242	3.5	5.676e-03
9	7	14	4	7	0	266.0857002	3.0	6.900e-03
10	8	13	5	4	1	266.0859750	4.5	7.175e-03





0.40

0.35

0.30

0.25

0.20

0.15

0.10

0.05

0

Normalized Intensity

M05(m)

M03(dd)

M04(m)

7.06

7.09

6.98

7.34

7.37

7.70

7.71

M02(d, p)

7.74

7.77

7.80

7.83

7.96

8.00

8.03

8.06

8.10

8.13

8.16

8.20

8.23

8.26

8.29

8.32

8.35

8.38

8.41

8.44

8.47

8.50

8.53

8.56

8.59

8.62

8.65

8.68

8.71

8.74

8.77

8.80

8.83

8.86

8.89

8.92

8.95

8.98

9.01

9.04

9.07

9.10

9.13

9.16

9.19

9.22

9.25

9.28

9.31

9.34

9.37

9.40

9.43

9.46

9.49

9.52

9.55

9.58

9.61

9.64

9.67

9.70

9.73

9.76

9.79

9.82

9.85

9.88

9.91

9.94

9.97

10.00

10.03

10.06

10.09

10.12

10.15

10.18

10.21

10.24

10.27

10.30

10.33

10.36

10.39

10.42

10.45

10.48

10.51

10.54

10.57

10.60

10.63

10.66

10.69

10.72

10.75

10.78

10.81

10.84

10.87

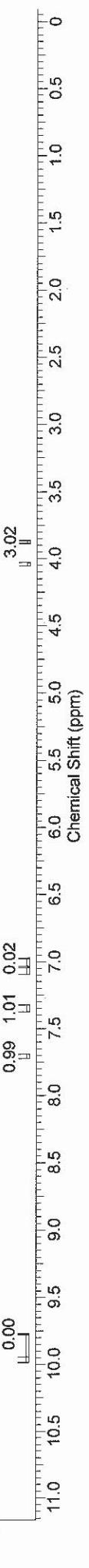
10.90

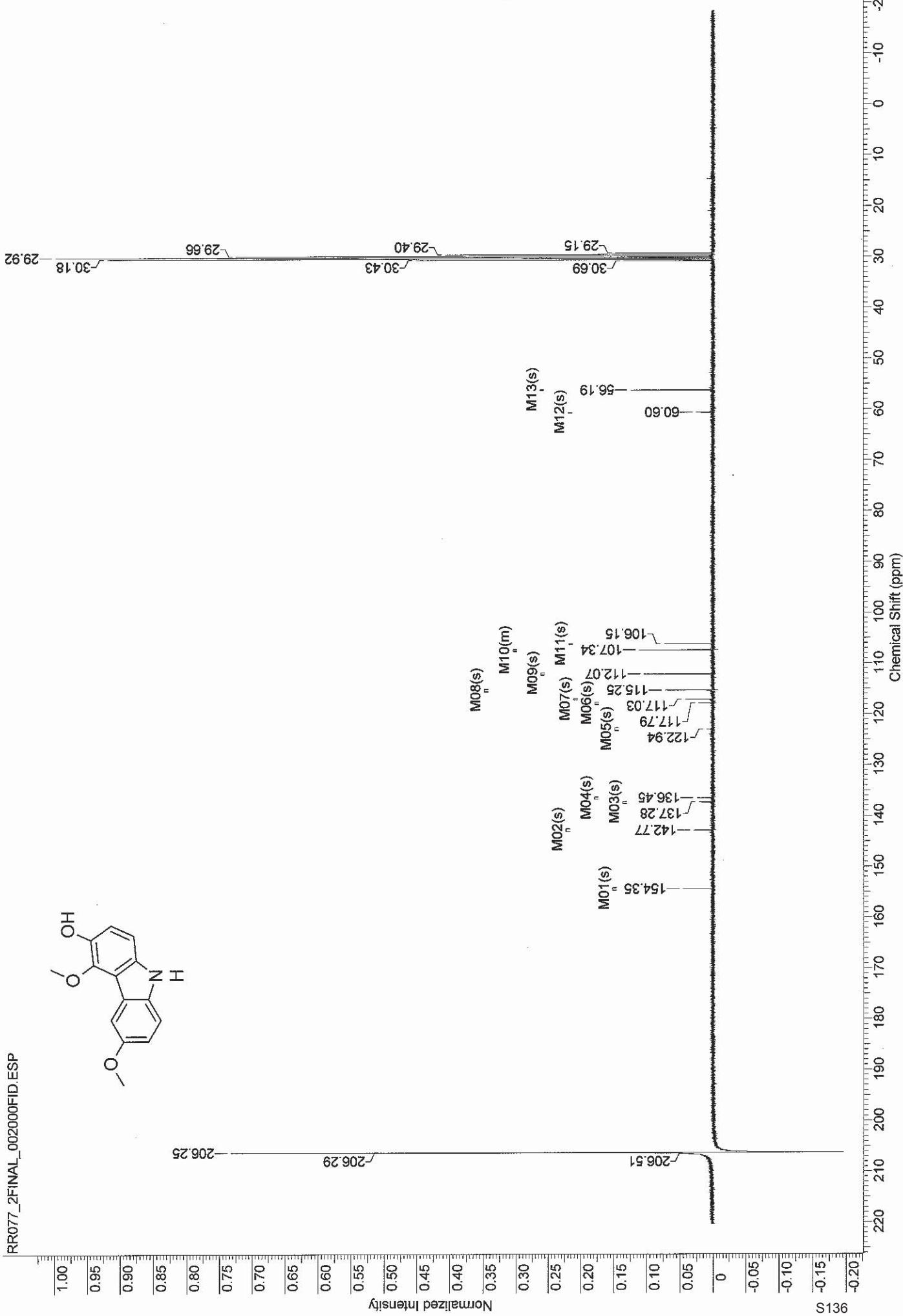
10.93

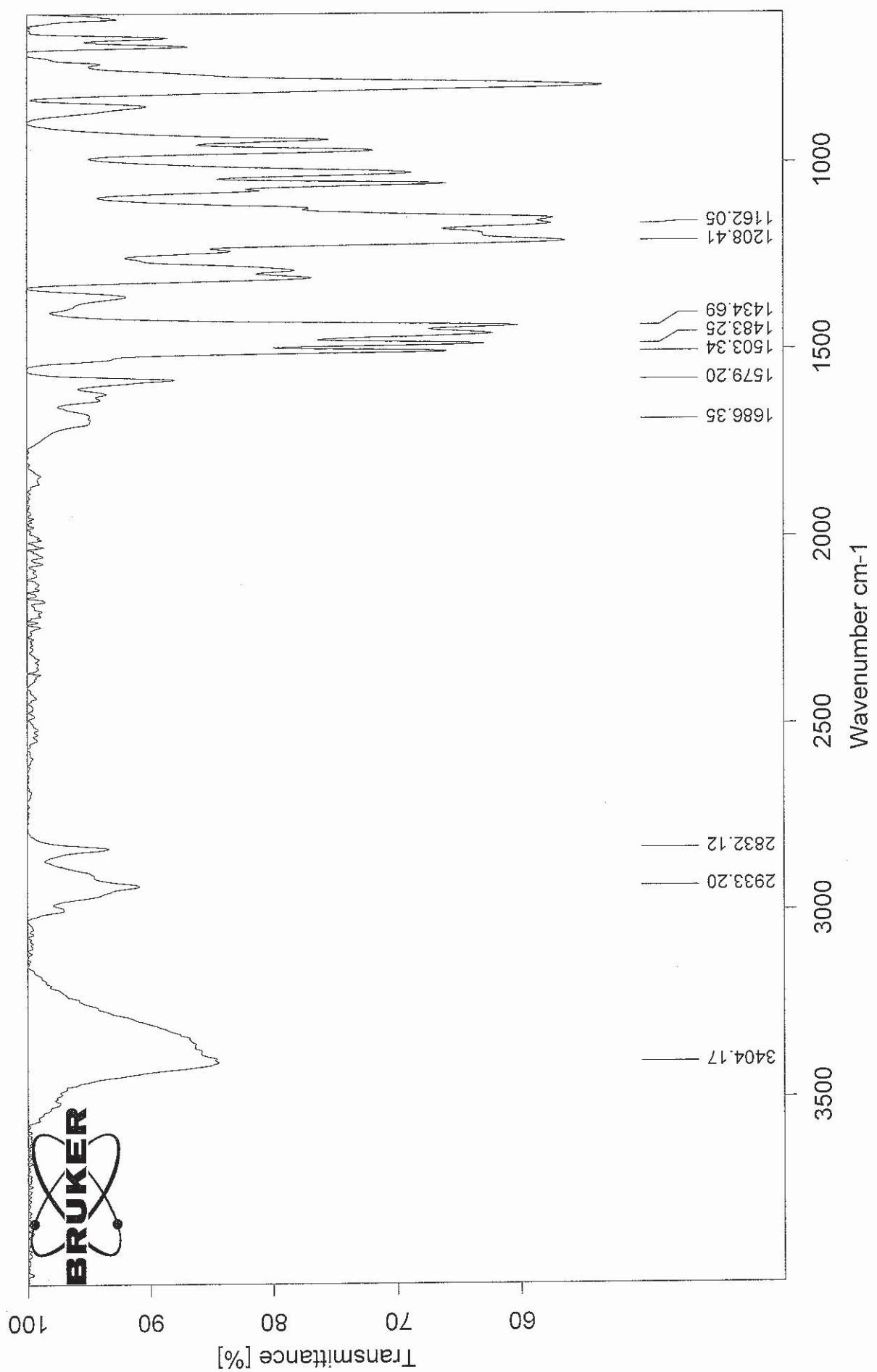
10.96

10.99

11.02







18.11.2013

U:\IR\4,6-dimethoxy-9H-carbazol-3-ol 4,6-dimethoxy-9H-carbazol-3-ol brown solid



ESI-MS: RR077B

XMASS Mass Analysis for /Data/UNI_FR/REMY7722_ESI/1/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

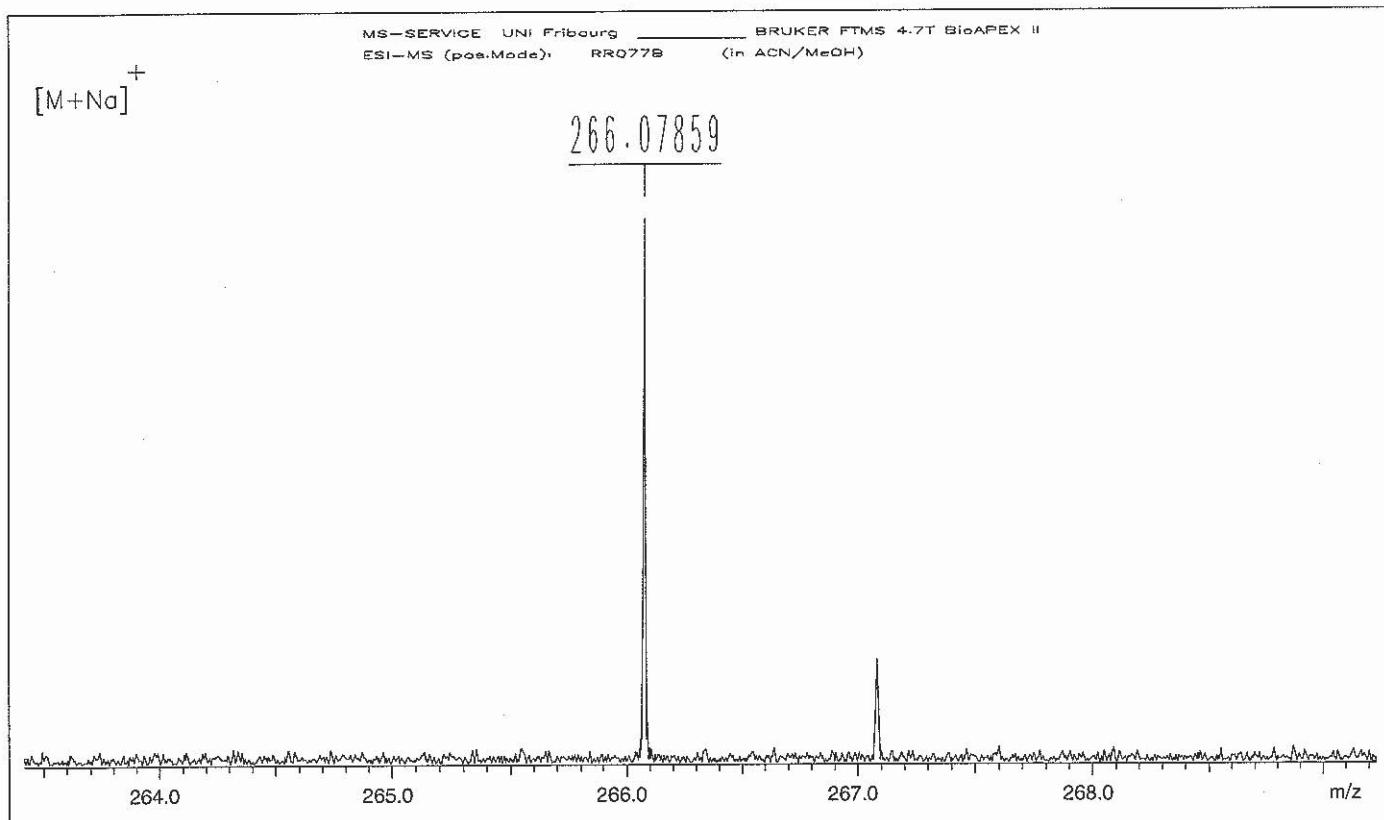
Ion mass = 266.0785860

Charge = +1

C H N O Na mass DBE error

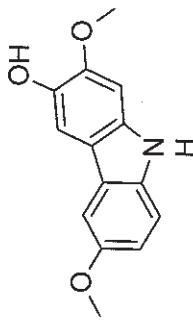
*** Mass Analysis for mass 266.0785860

1	14	13	1	3	1	266.0787644	8.5	1.784e-04
2	12	11	4	2	1	266.0774217	9.0	1.164e-03
3	14	10	4	2	0	266.0798270	12.0	1.241e-03
4	11	12	3	5	0	266.0771469	7.5	1.439e-03
5	16	12	1	3	0	266.0811697	11.5	2.584e-03
6	9	13	3	5	1	266.0747416	4.5	3.844e-03
7	5	15	4	7	1	266.0832949	0.0	4.709e-03
8	6	12	5	7	0	266.0731242	3.5	5.462e-03
9	7	14	4	7	0	266.0857002	3.0	7.114e-03
10	8	13	5	4	1	266.0859750	4.5	7.389e-03



M01(s)

M02(s)



3.92

3.85

3.83

-2.81

-2.06 -2.04 -2.05

-2.78
-2.780.01
μl

-2.06 -2.04 -2.05

-2.78
-2.78

M03(dd)

M06(m)

M05(s)

M04(s)

M07(s)

M08(d)

M09(d)

6.91
6.90
6.88
6.877.00
7.047.28
7.31
7.48
7.50
7.490.99
0.99
0.99
0.99
0.997.5
8.0
8.5
9.0
9.5
10.0
10.5

M09(br. s.)

0.89
0.899.5
10.0
10.5

Normalized Intensity

Normalized Intensity

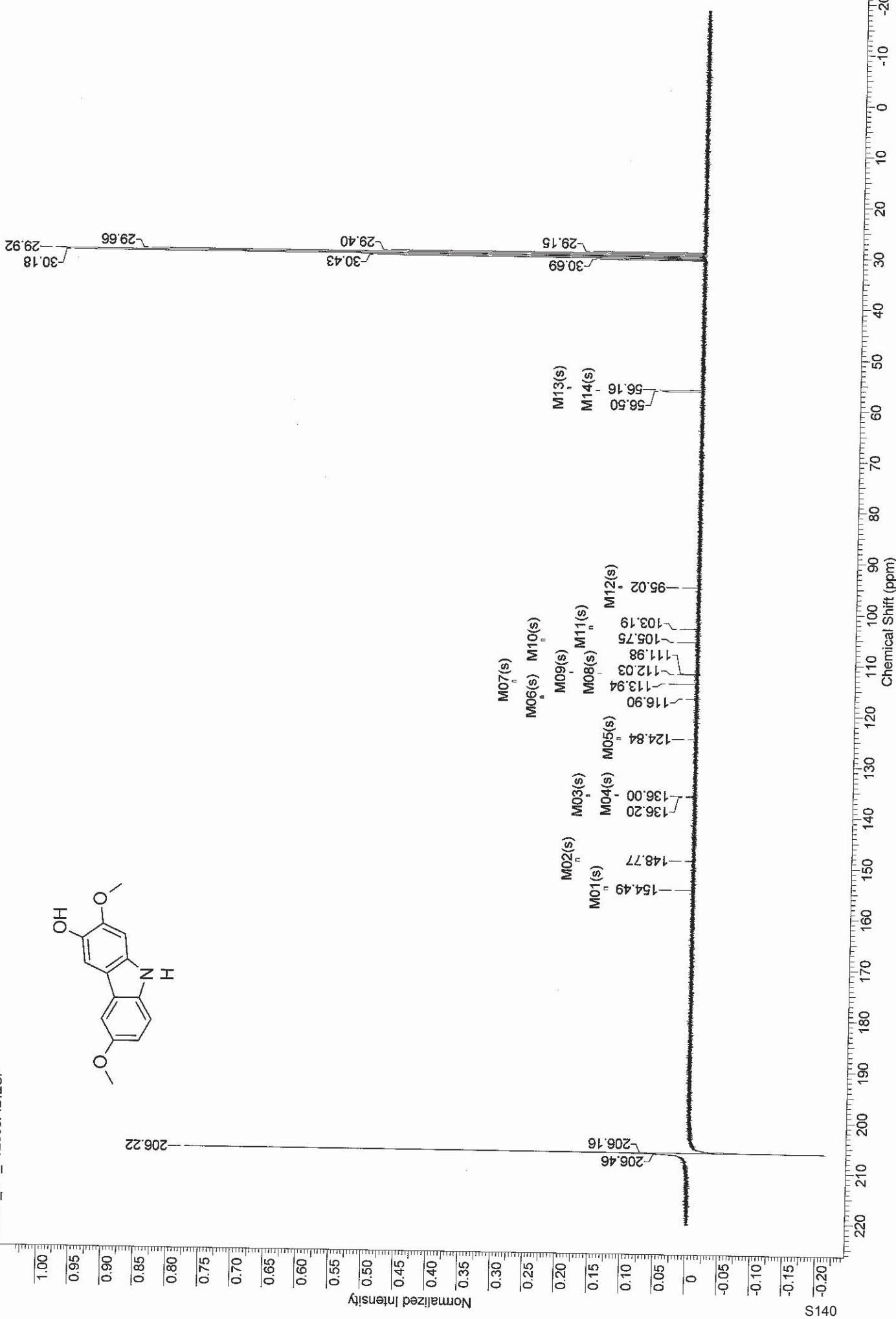
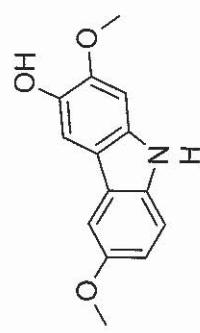
S139

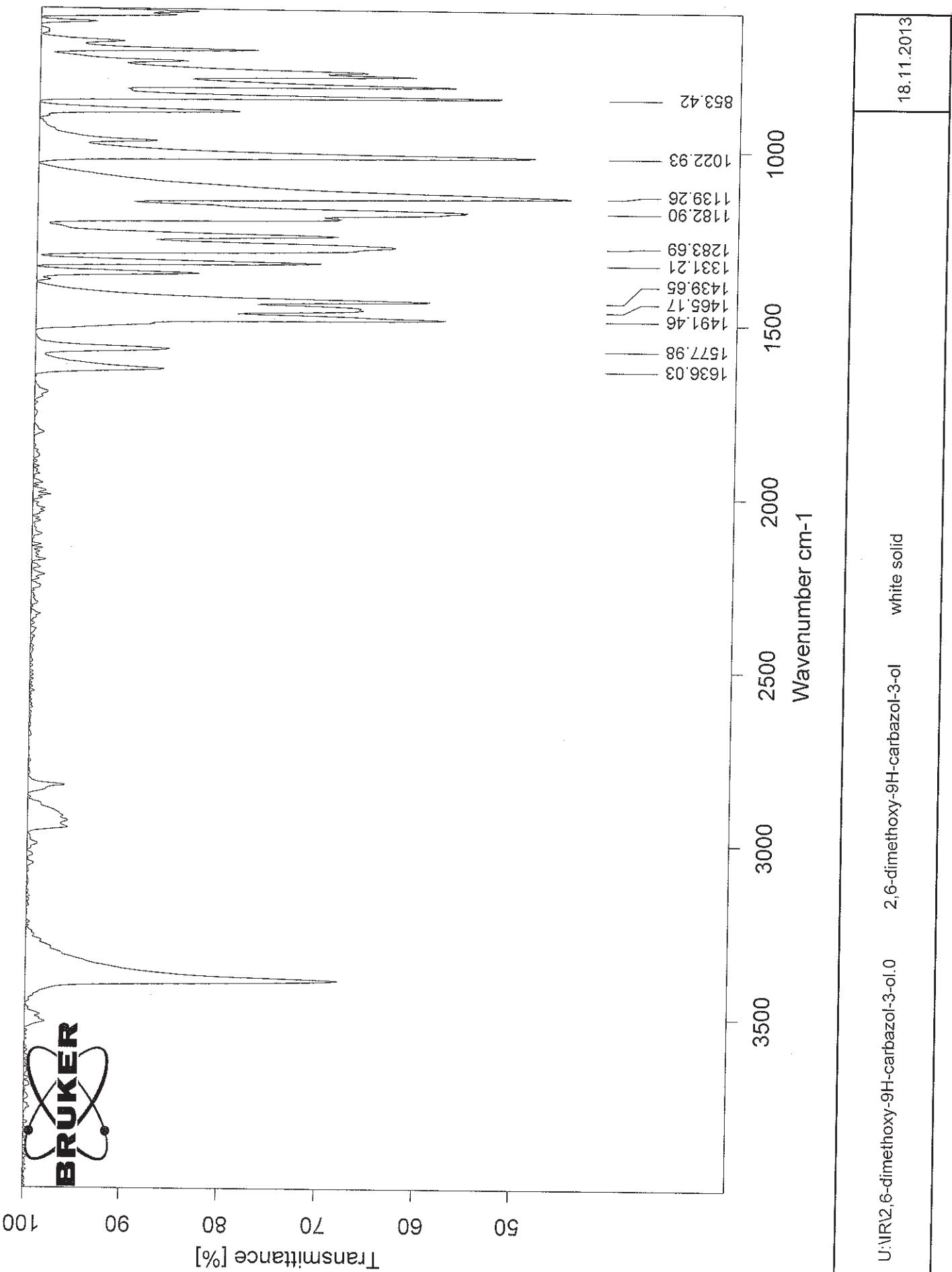
0
0.5
1.0
1.5
2.0
2.5
3.0
3.5
4.0
4.5
5.0-2.0
-1.5
-1.0
-0.5
0

Chemical Shift (ppm)

6.0
5.5
5.0
4.5
4.0
3.5
3.0
2.5
2.0
1.5
1.0
0.5
0

Chemical Shift (ppm)







ESI-MS: RR077A

XMASS Mass Analysis for /Data/UNI_FR/REMY7661_ESI/1/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

Ion mass = 266.0790090

Charge = +1

#	C	H	N	O	Na	mass	DBE	error
---	---	---	---	---	----	------	-----	-------

*** Mass Analysis for mass 266.0790090

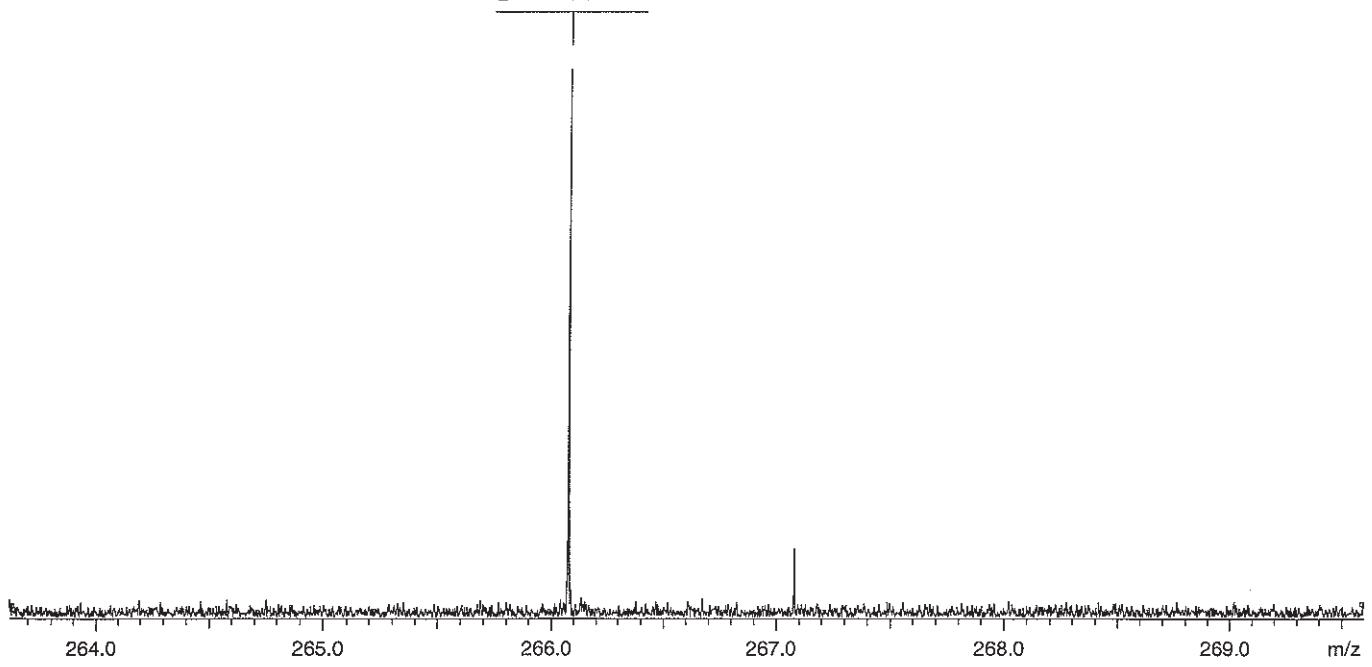
1	14	13	1	3	1	266.0787644	8.5	2.446e-04
2	12	8	7	1	0	266.0784843	12.5	5.247e-04
3	14	10	4	2	0	266.0798270	12.0	8.180e-04
4	12	11	4	2	1	266.0774217	9.0	1.587e-03
5	11	12	3	5	0	266.0771469	7.5	1.862e-03
6	16	12	1	3	0	266.0811697	11.5	2.161e-03
7	10	9	7	1	1	266.0760790	9.5	2.930e-03
8	9	10	6	4	0	266.0758043	8.0	3.205e-03
9	9	13	3	5	1	266.0747416	4.5	4.267e-03
10	5	15	4	7	1	266.0832949	0.0	4.286e-03

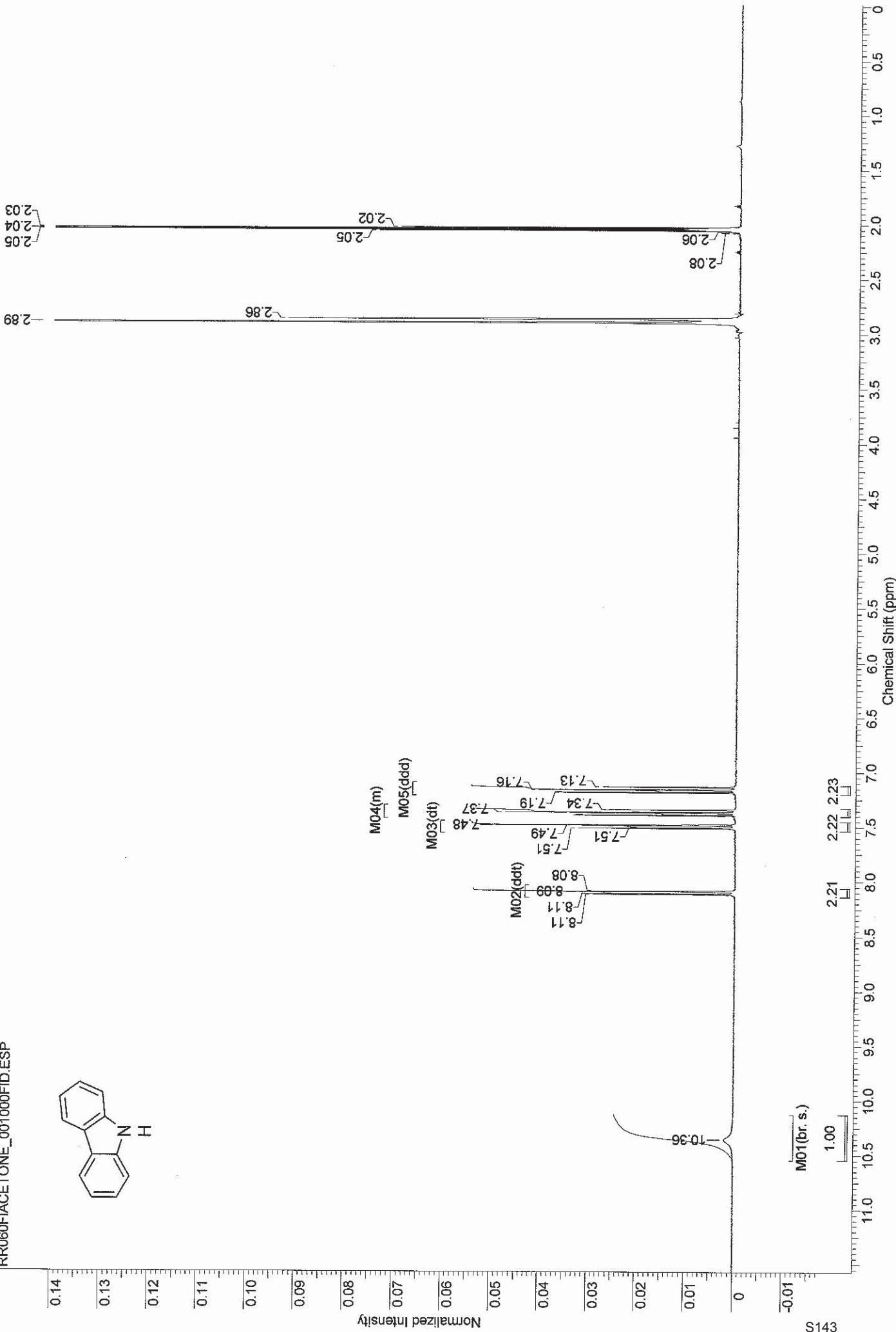
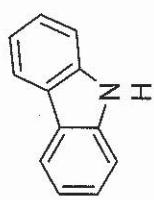
MS-SERVICE UNI Fribourg BRUKER FTMS 4.7T BioAPEX II
ESI-MS (pos.Mode) RR077A (in ACN/MeOH)

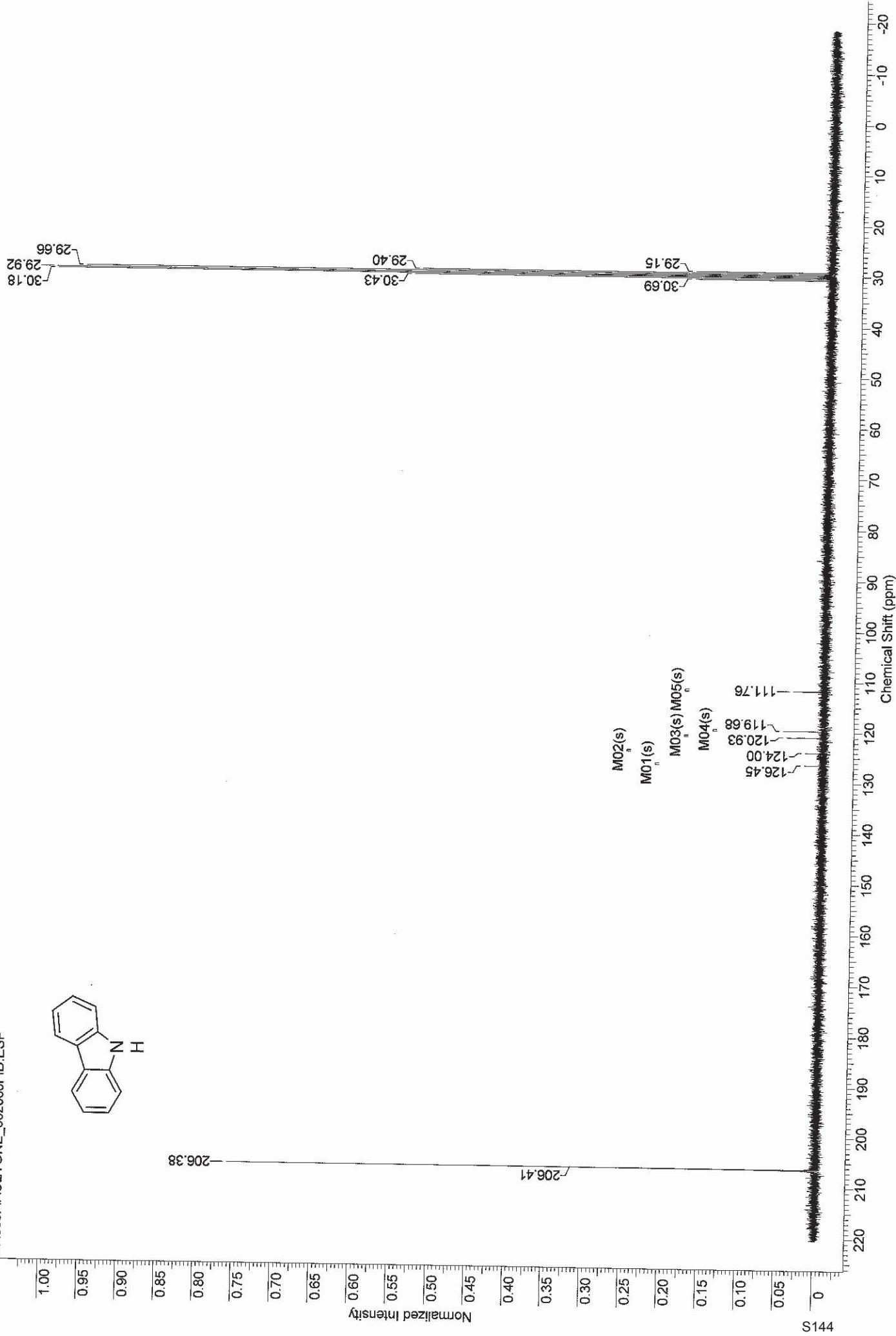
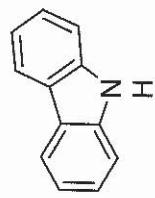
+

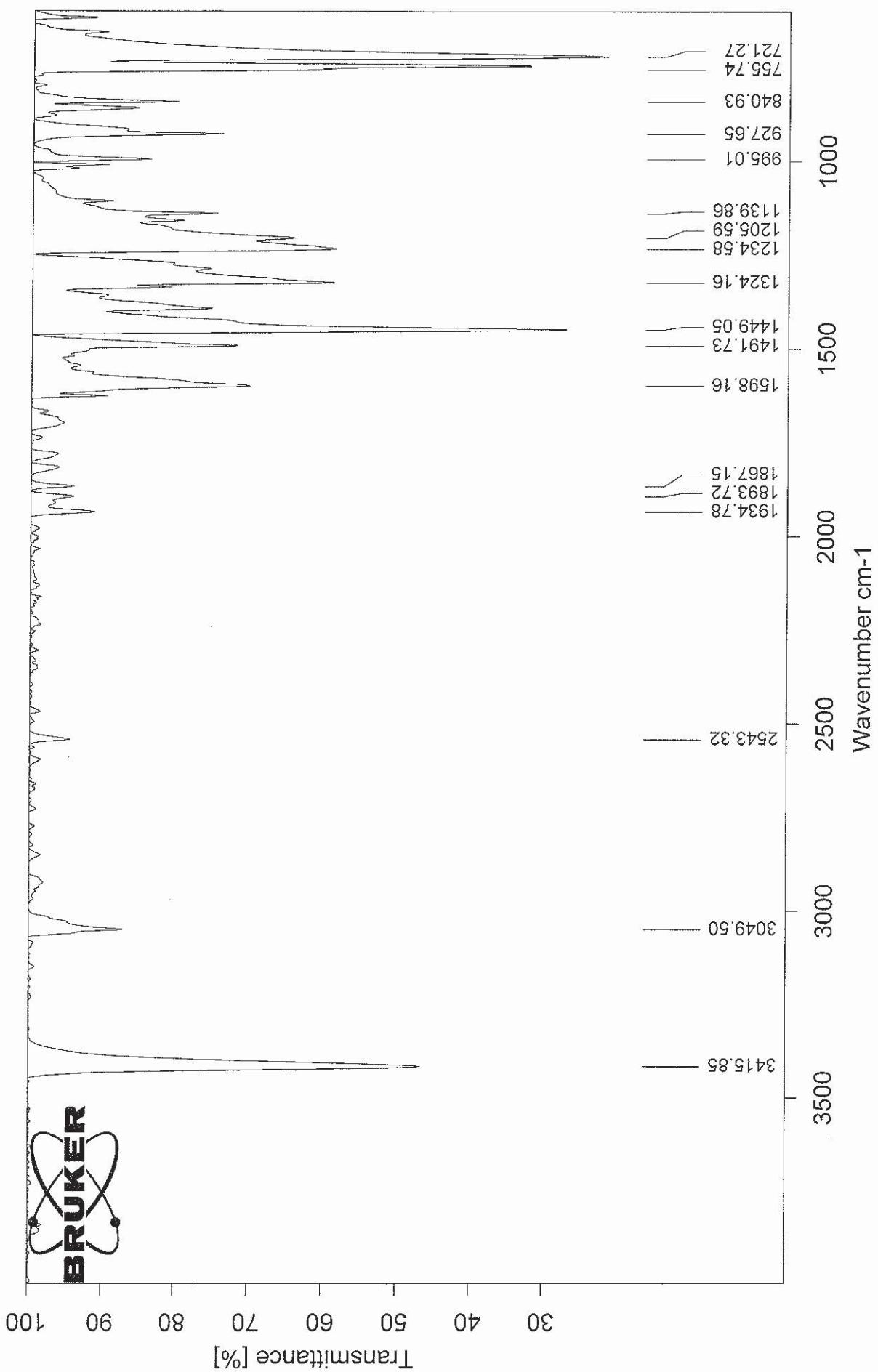
[M+Na]⁺

266.07901









U:\My Documents\polymer science lab\RR061IR.0 RR061IR

24.04.2013

D:\Data\MS_SERVICE_2013\REMY7739_MALDI0_G9\1\1SRef

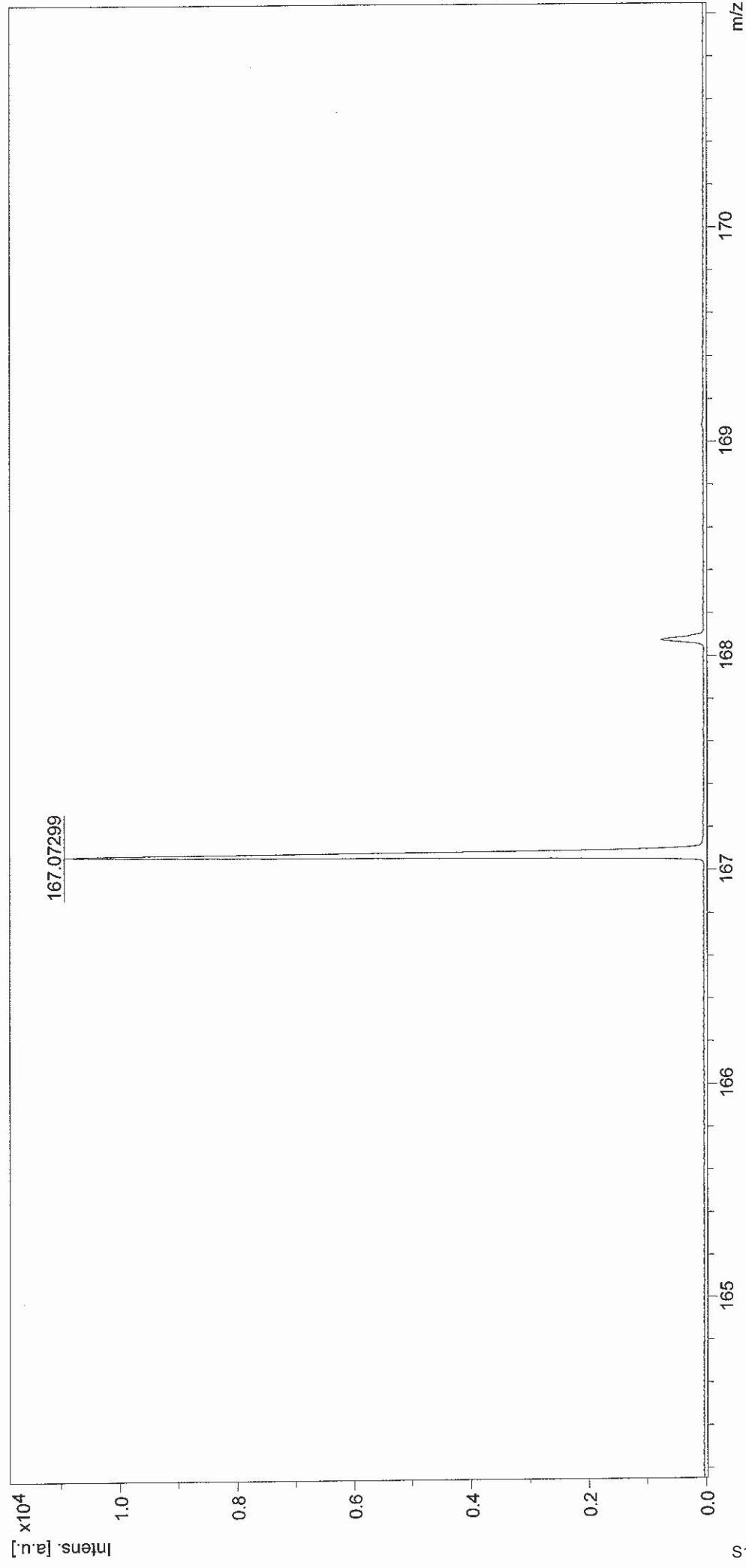
Acquisition method name D:\Methods\MS_SERVICE_2013\RP_150-2000.Da.par

Comment 1
Comment 2

RR061M in DCTB

Instrument type: MALDI-TOF

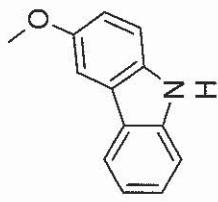
BRUKER ultraflexXtreme



S146

Bruker Daltonics flexAnalysis
printed: 11/22/2013 8:24:06 AM

M08(s)



1.00

0.95

0.90

0.85

0.80

0.75

0.70

0.65

0.60

0.55

0.50

0.45

0.40

0.35

0.30

Normalized Intensity

0.30

0.25

0.20

0.15

0.10

0.05

0

M06(m)

M07(m)

M04(m)

M05(dd)

7.27

M03(d)

7.41

7.42

7.10

7.22

7.09

7.07

7.06

M01(br. s.)

7.58

7.57

7.42

8.03

8.04

8.06

8.06

8.06

8.06

M02(br. s.)

8.03

8.04

8.05

8.06

8.07

8.08

8.09

8.10

8.11

8.12

8.12

0.96

0.97

0.98

0.99

2.92

2.06

1.28

1.56

3.95

0

0.5

1.0

1.5

2.0

2.5

3.0

3.5

4.0

4.5

5.0

5.5

6.0

6.5

7.0

7.5

8.0

8.5

9.0

9.5

10.0

S147

0

0.5

1.0

1.5

2.0

2.5

3.0

3.5

4.0

4.5

5.0

5.5

6.0

6.5

7.0

7.5

8.0

8.5

9.0

9.5

10.0

0

0.5

1.0

1.5

2.0

2.5

3.0

3.5

4.0

4.5

5.0

5.5

6.0

6.5

7.0

7.5

8.0

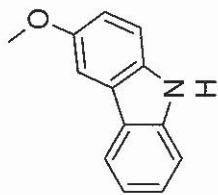
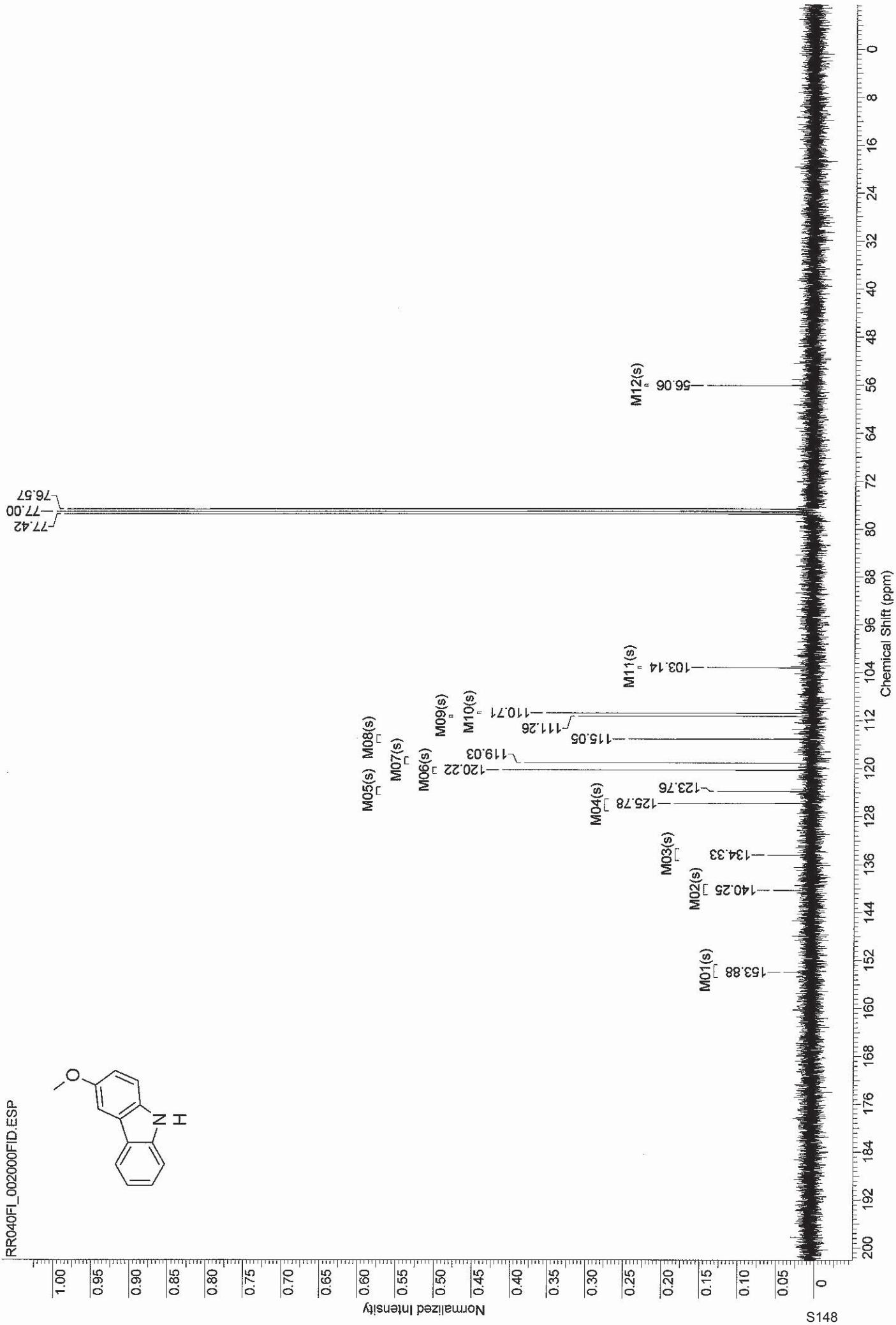
8.5

9.0

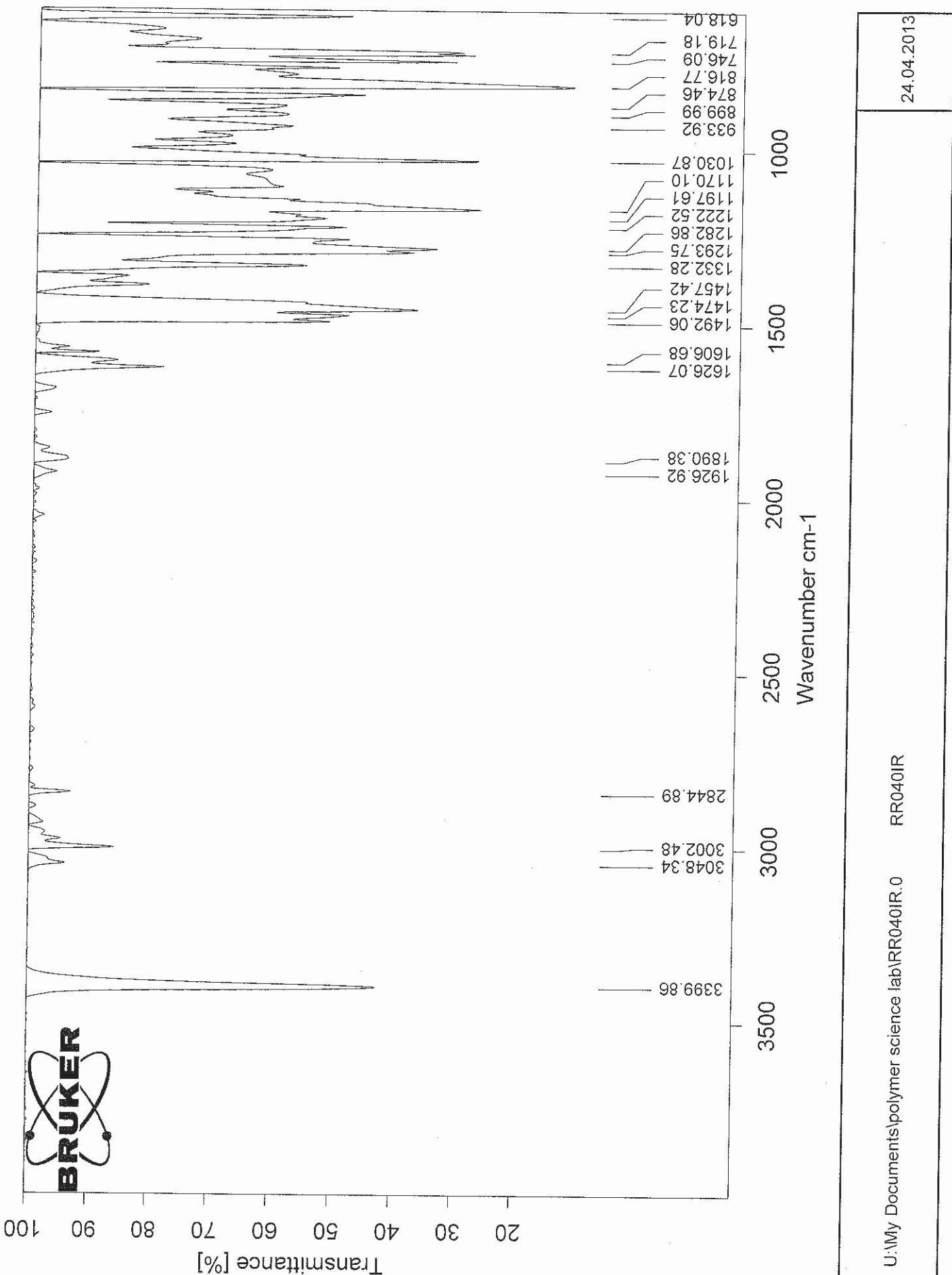
9.5

10.0

0



RR040FI_00020000FID.ESP



D:\Data\MS_SERVICE_2013\REMY7734_MALDI\0_B9\2\1SRef

Acquisition method name

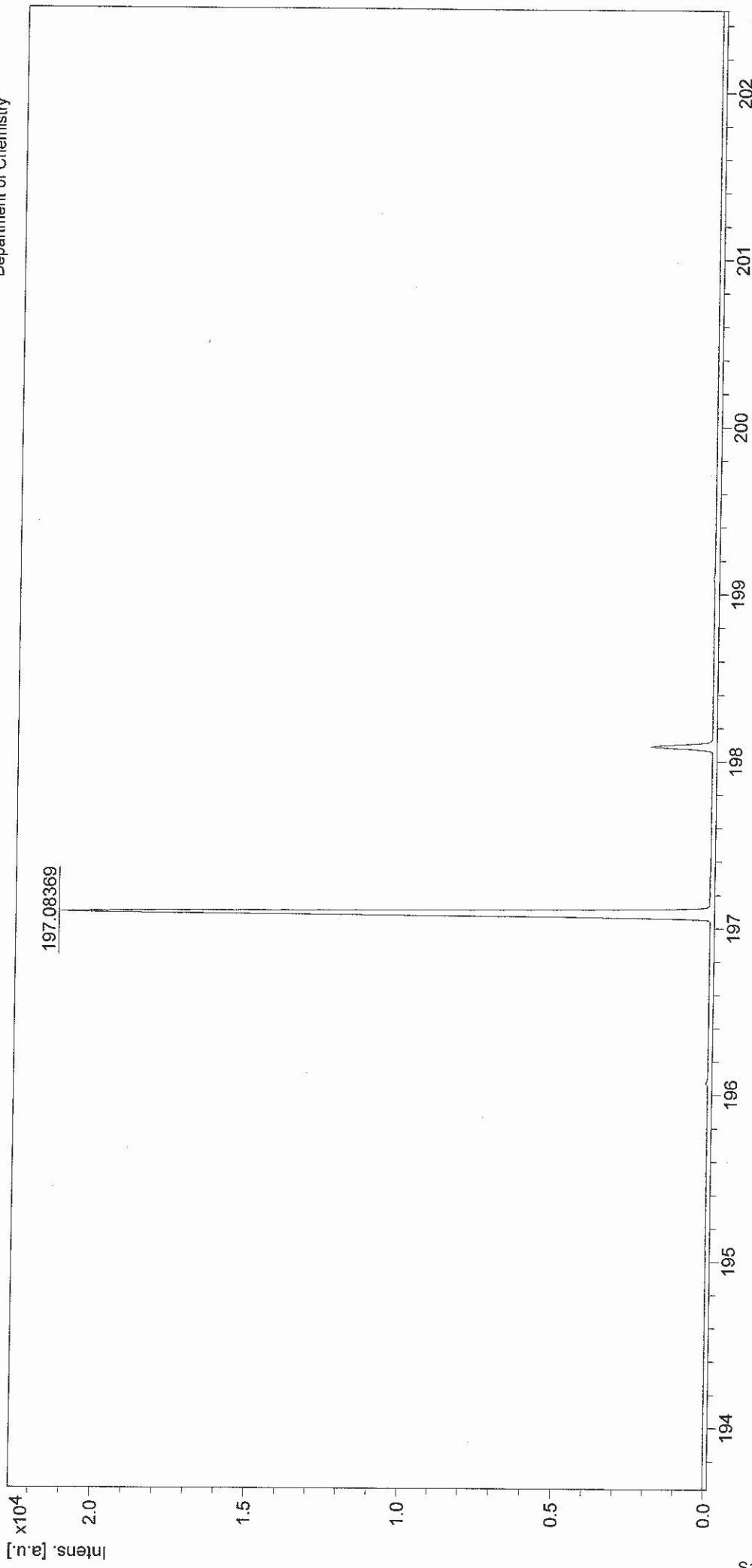
D:\Methods\MS_SERVICE_2013\RP_150-2000_Da.par

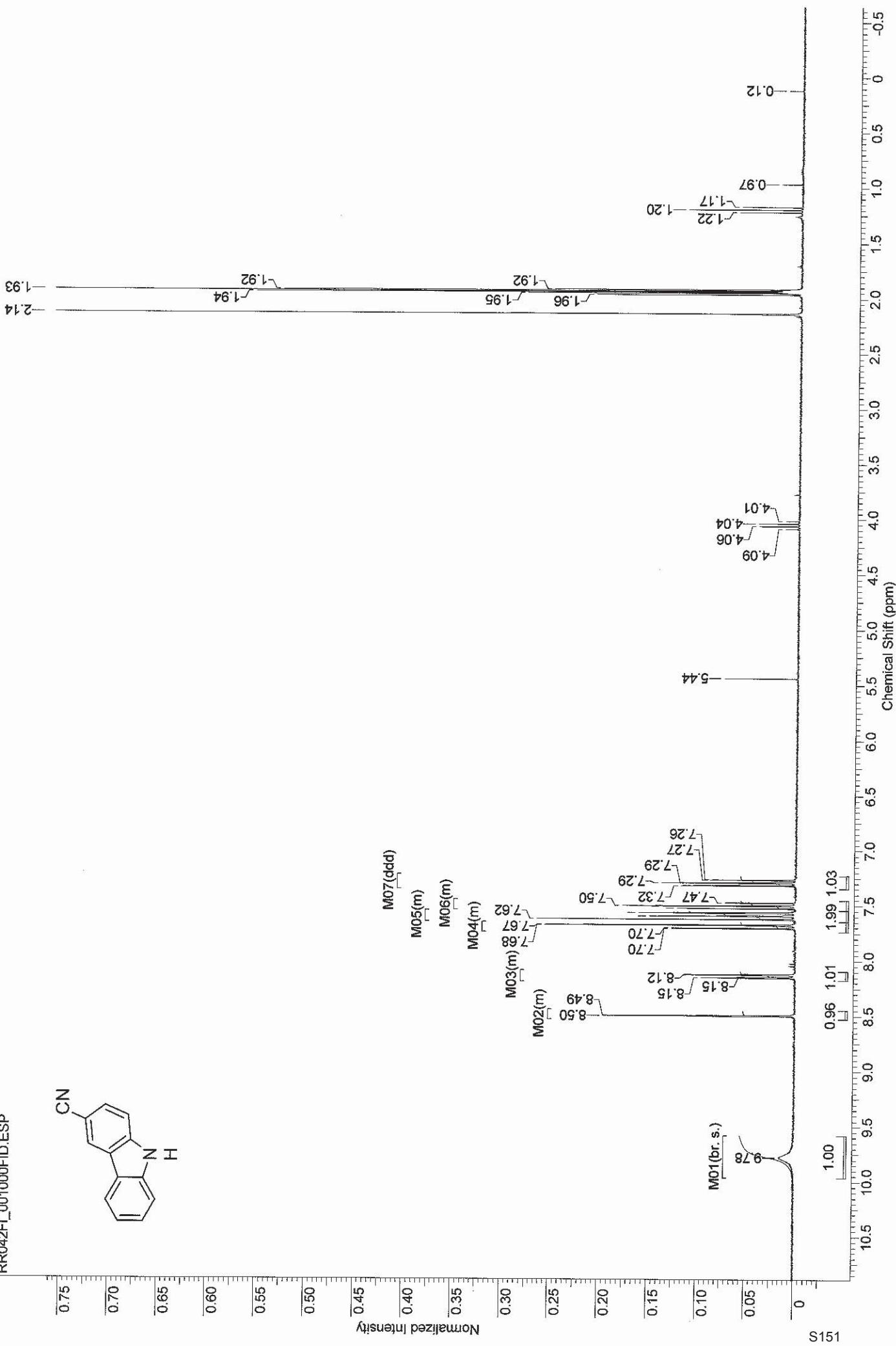
Comment 1
Comment 2

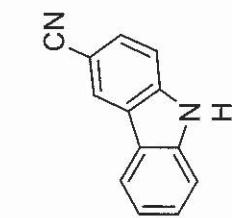
RR040M in DCTB

Instrument type: MALDI-TOF BRUKER ultraflexXtreme

Intens. [a.u.]
 $\times 10^4$







M07(br. s.)
M06(s)
M05(s)

118.38

M03(s)
M01(s)
M02(s)
M04(s)
M08(s)
M09(s)

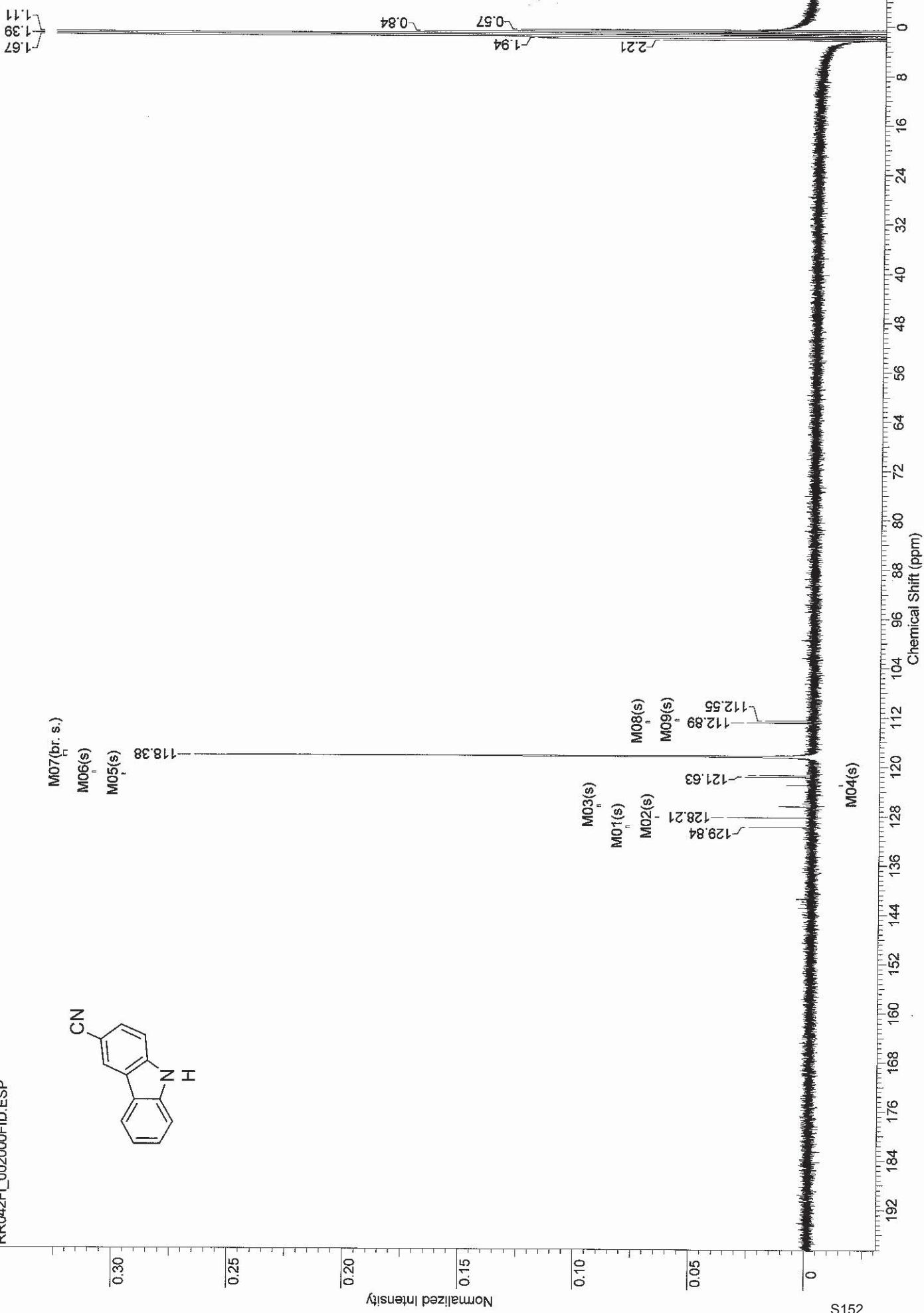
121.63

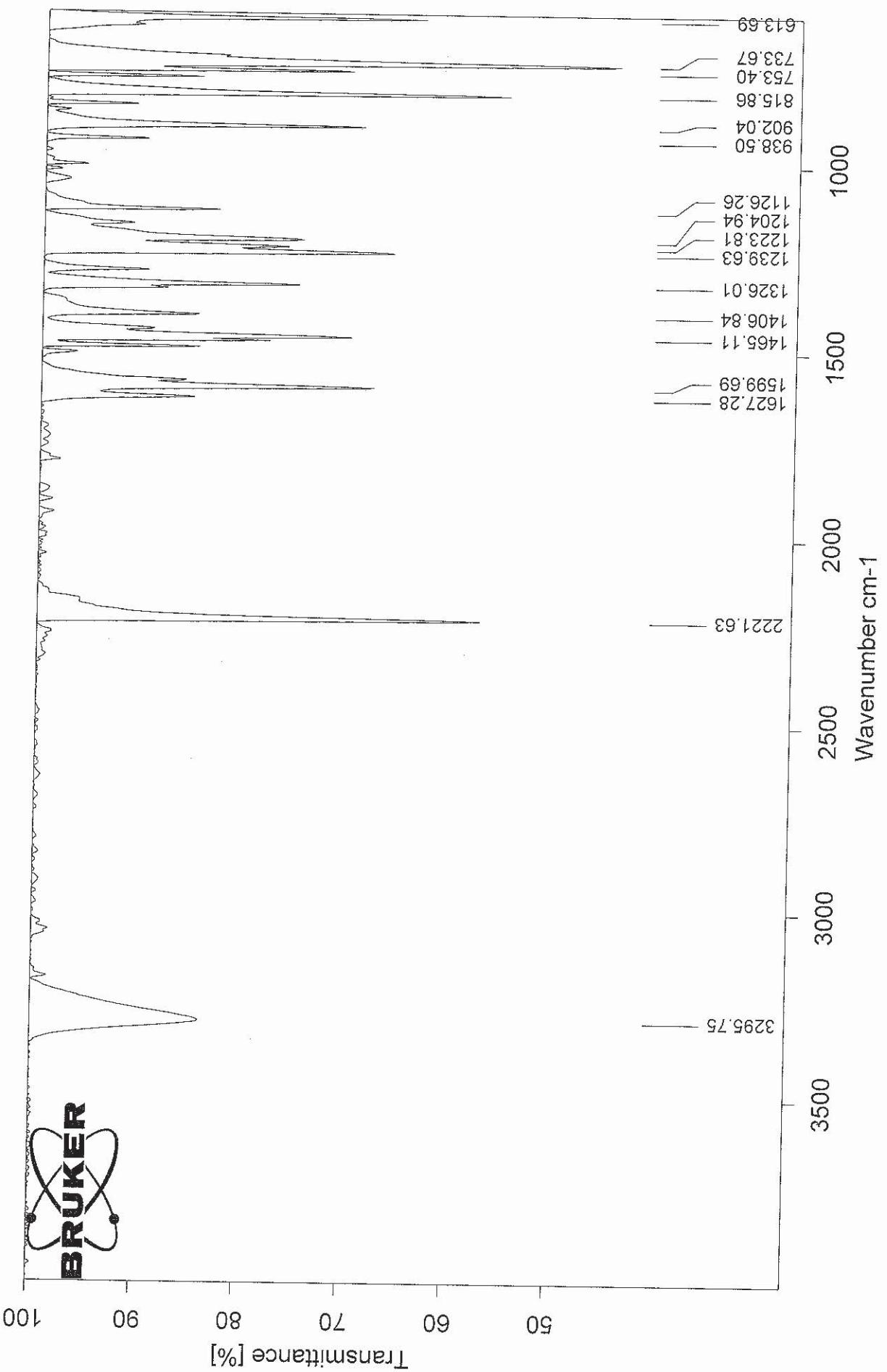
128.27

129.84

112.55

112.69





U:\My Documents\polymer science lab\RR042\IR.0

RR042\IR

24.04.2013



ESI-MS: RR042M

XMASS Mass Analysis for /Data/UNI_FR/REMY7737_ESI/2/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

Ion mass = 215.0574740

Charge = +1

#	C	H	N	Na	mass	DBE	error
---	---	---	---	----	------	-----	-------

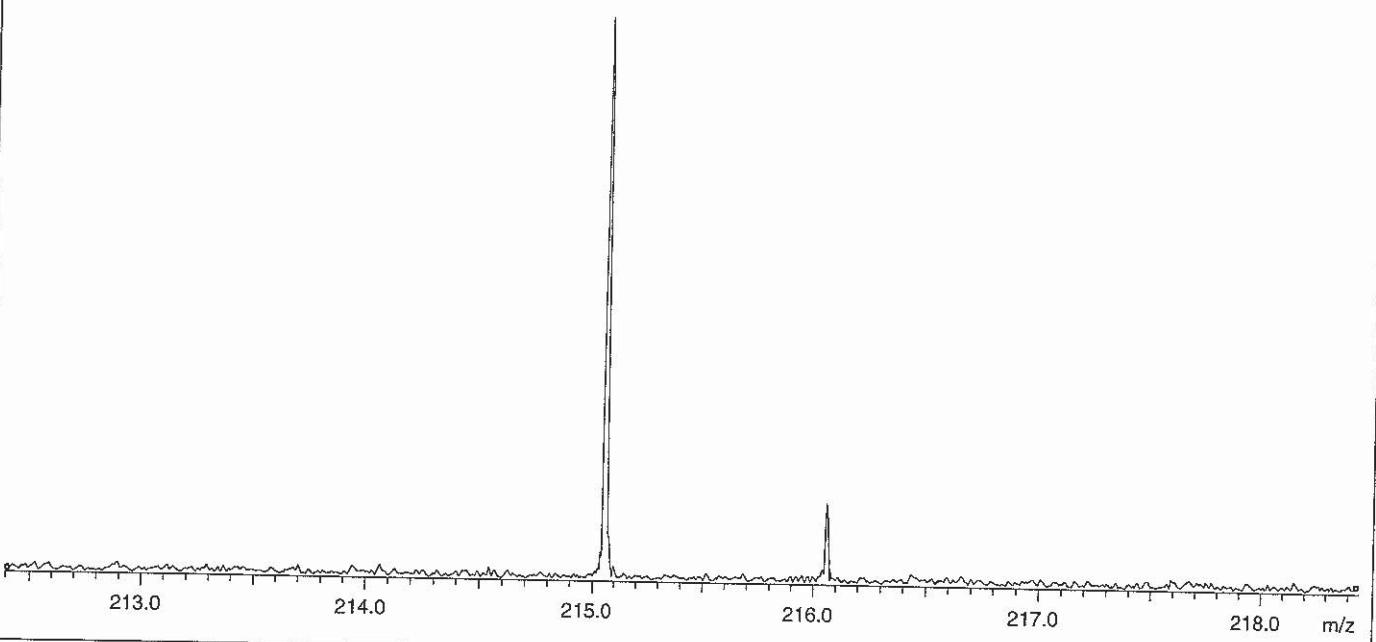
*** Mass Analysis for mass 215.0574740

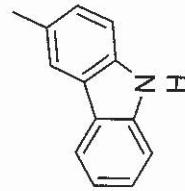
1	13	8	2	1	215.0579694	10.5	4.954e-04
2	15	7	2	0	215.0603747	13.5	2.901e-03
3	6	3	10	0	215.0536666	10.5	3.807e-03
4	5	6	9	1	215.0638373	7.0	6.363e-03
5	7	5	9	0	215.0662426	10.0	8.769e-03
6	14	5	3	0	215.0477986	14.0	9.675e-03
7	12	6	3	1	215.0453933	11.0	1.208e-02
8	14	10	1	1	215.0705454	10.0	1.307e-02
9	16	9	1	0	215.0729507	13.0	1.548e-02
10	6	8	8	1	215.0764134	6.5	1.894e-02

[M+Na]⁺

MS-SERVICE UNI Fribourg BRUKER FTMS 4.7T BioAPEX II
ESI-MS (pos.Mode): RR042M
(in MeOH/CHCl₃)

215.05747





M07(m)

M05(m)

M04(m)

M06(m)

M09(m)

M08(s)

-2.56

-7.42

-7.44

-7.29

-7.26

-7.23

-7.22

-7.21

-7.20

M02(br. s.)

M01(dq)

M03(td)

M04

M05

M06

M07

M08

M09

M10

M11

M12

M13

M14

M15

M16

M17

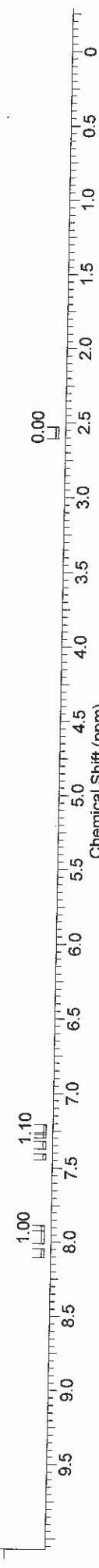
M18

M19

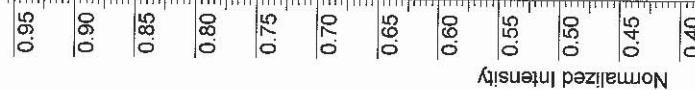
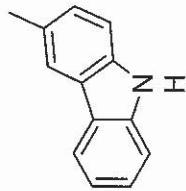
M20

Normalized Intensity

S155



77.42
77.00
76.58



M01(m)

M03(s)

M05(s)

M04(s) M09(s)

M08(s)

M10(s)

M11(s)

M10

M11

M2

M03(s)

M02(s)

M01

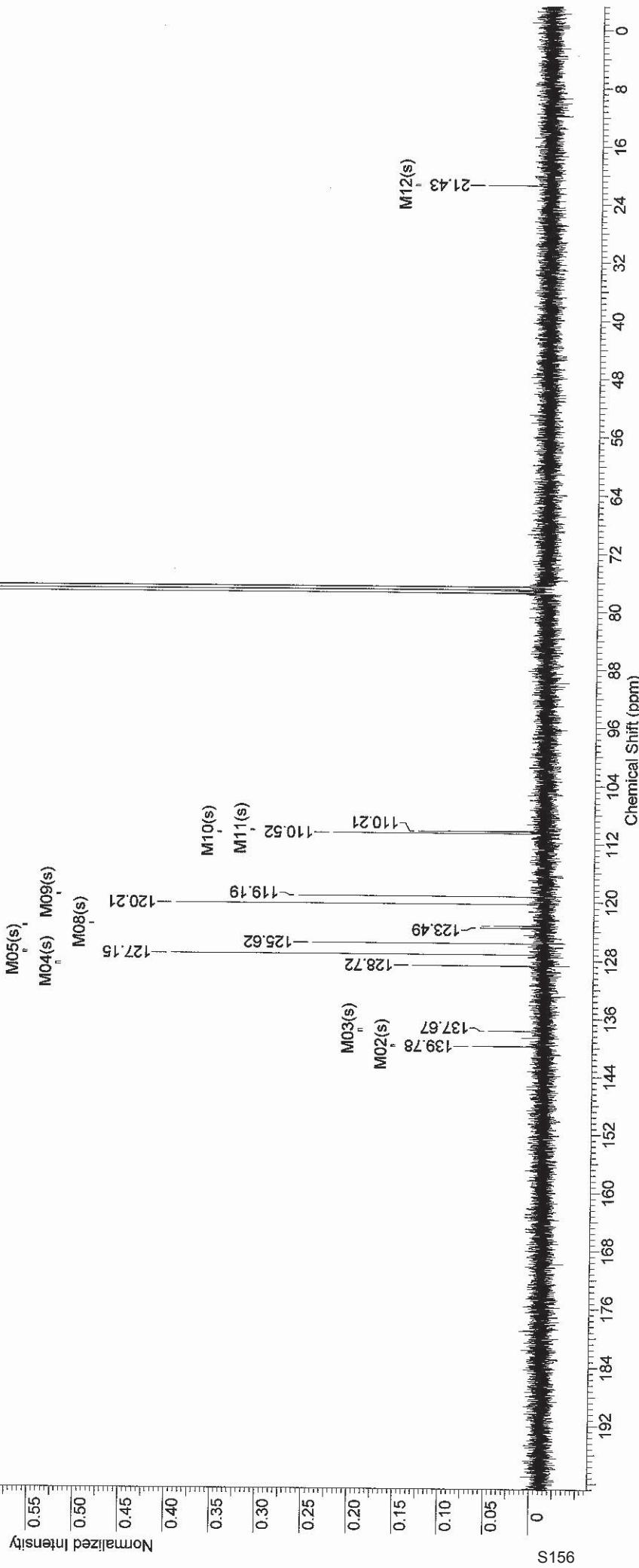
M02

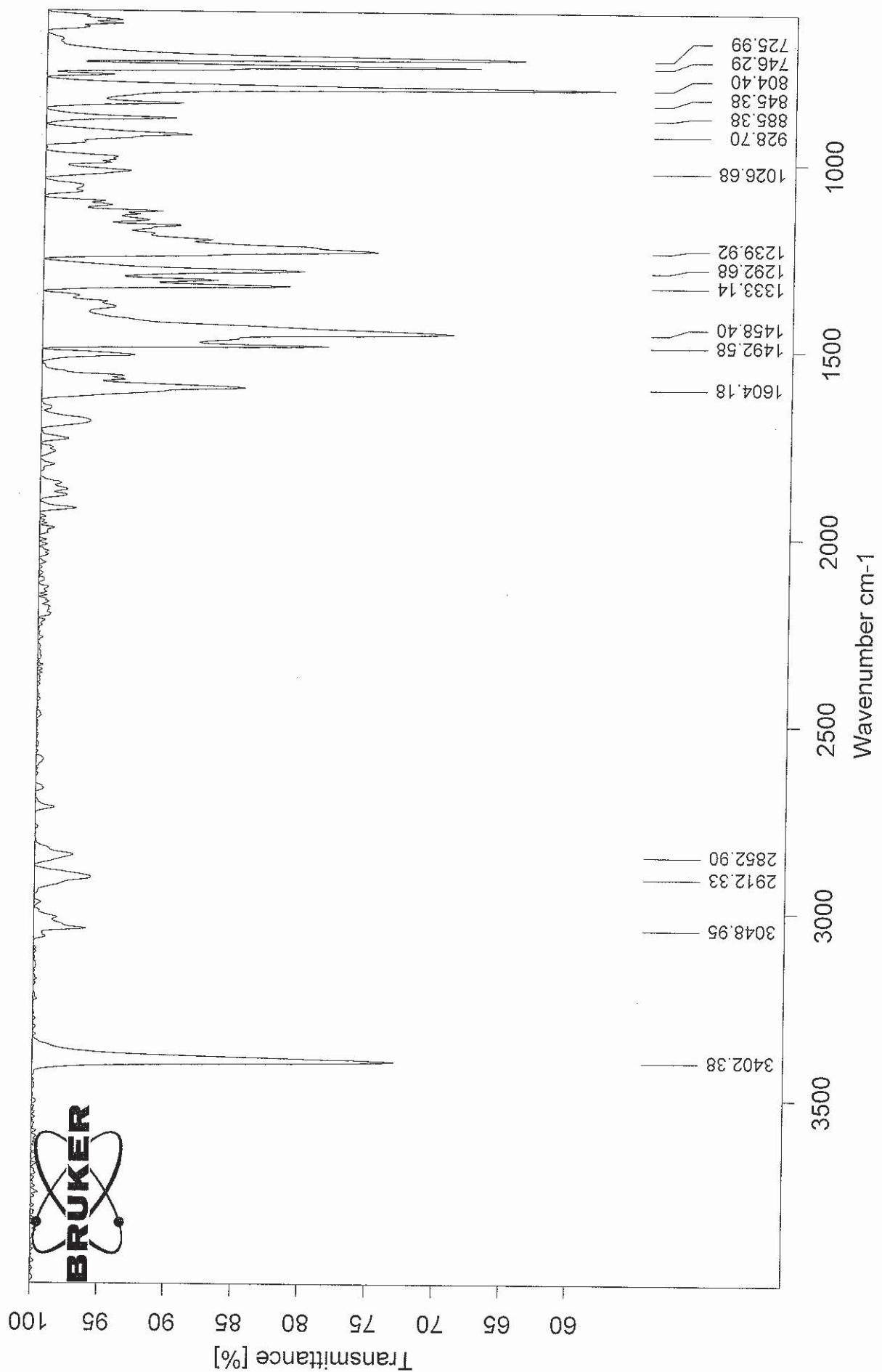
M12(s)

M13

M14

S156





U:\My Documents\polymer science lab\RR083IR.0

RR083IR

24.04.2013

D:\Data\MS_SERVICE_2013\REMY7738_MALDI\0_F9\1\1SRef

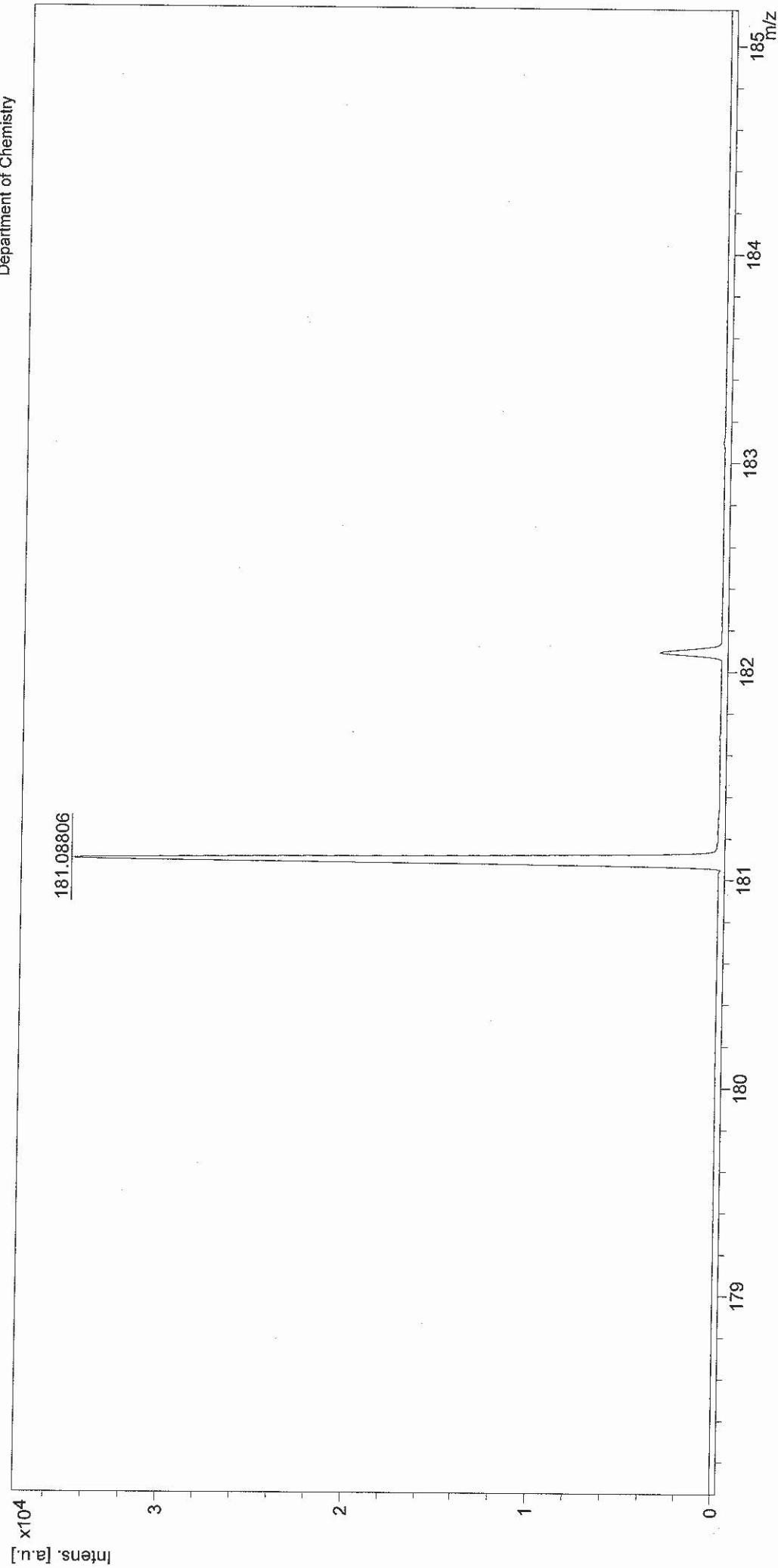
Acquisition method name

D:\Methods\MS_SERVICE_2013\RP_150-2000_Da.par

Comment 1
Comment 2

RR083M in DCTB

Instrument type: MALDI-TOF BRUKER ultraflexXtreme

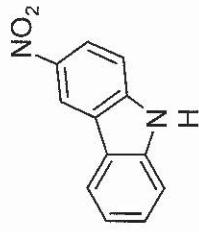


M08(m)

M07(m)

M06(dd)

M05(m)



7.27

0.85

0.80

0.75

0.70

0.65

0.60

0.55

0.50

0.45

0.40

0.35

0.30

0.25

0.20

0.15

0.10

0.05

0

Normalized Intensity

M02(m)

M03(d)

M04(m)

M01(d)

M02(m)

M03(d)

M04(m)

M05(m)

M06(dd)

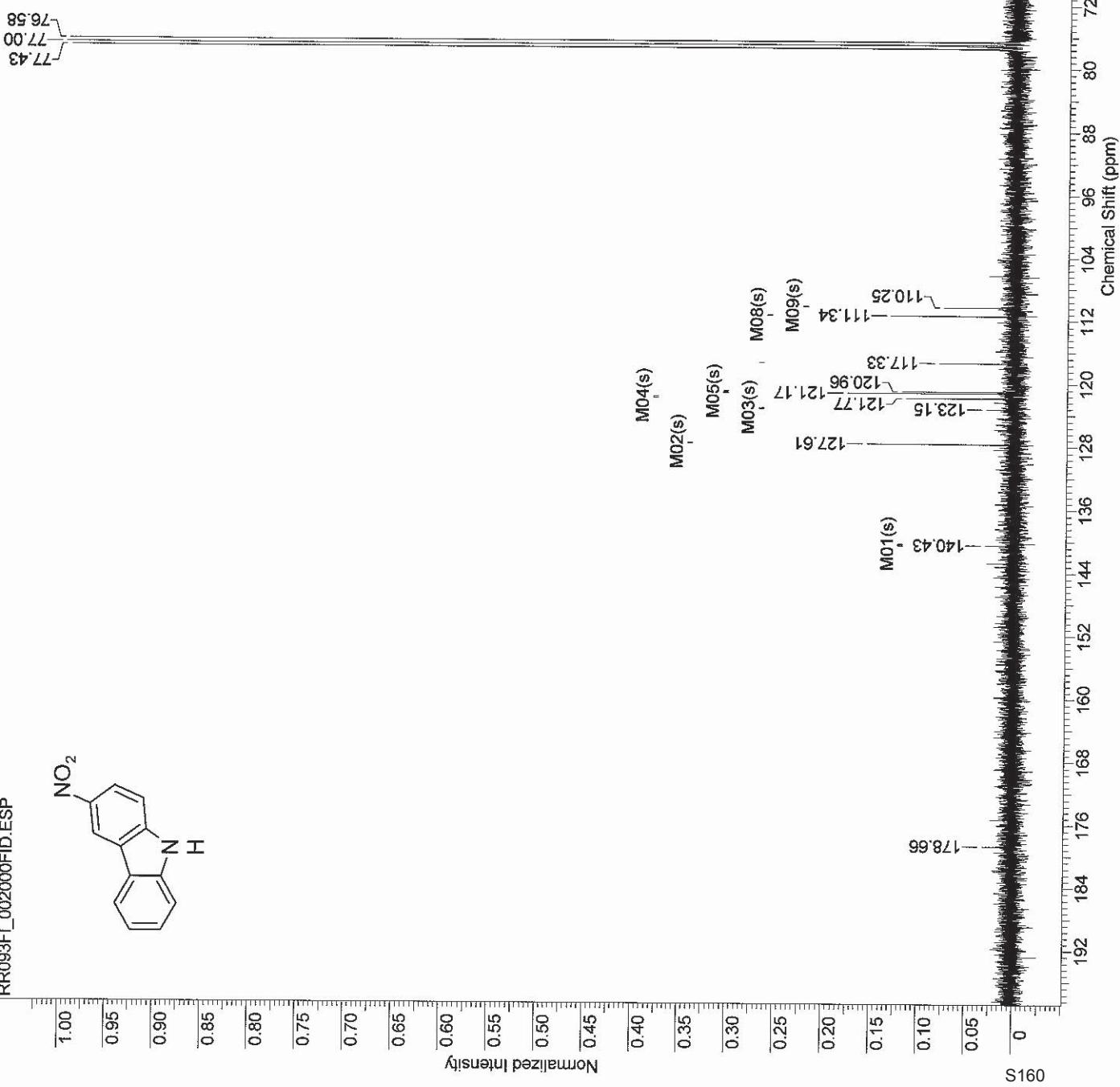
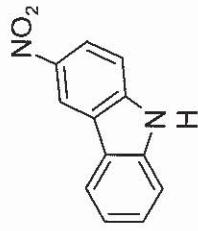
M07(m)

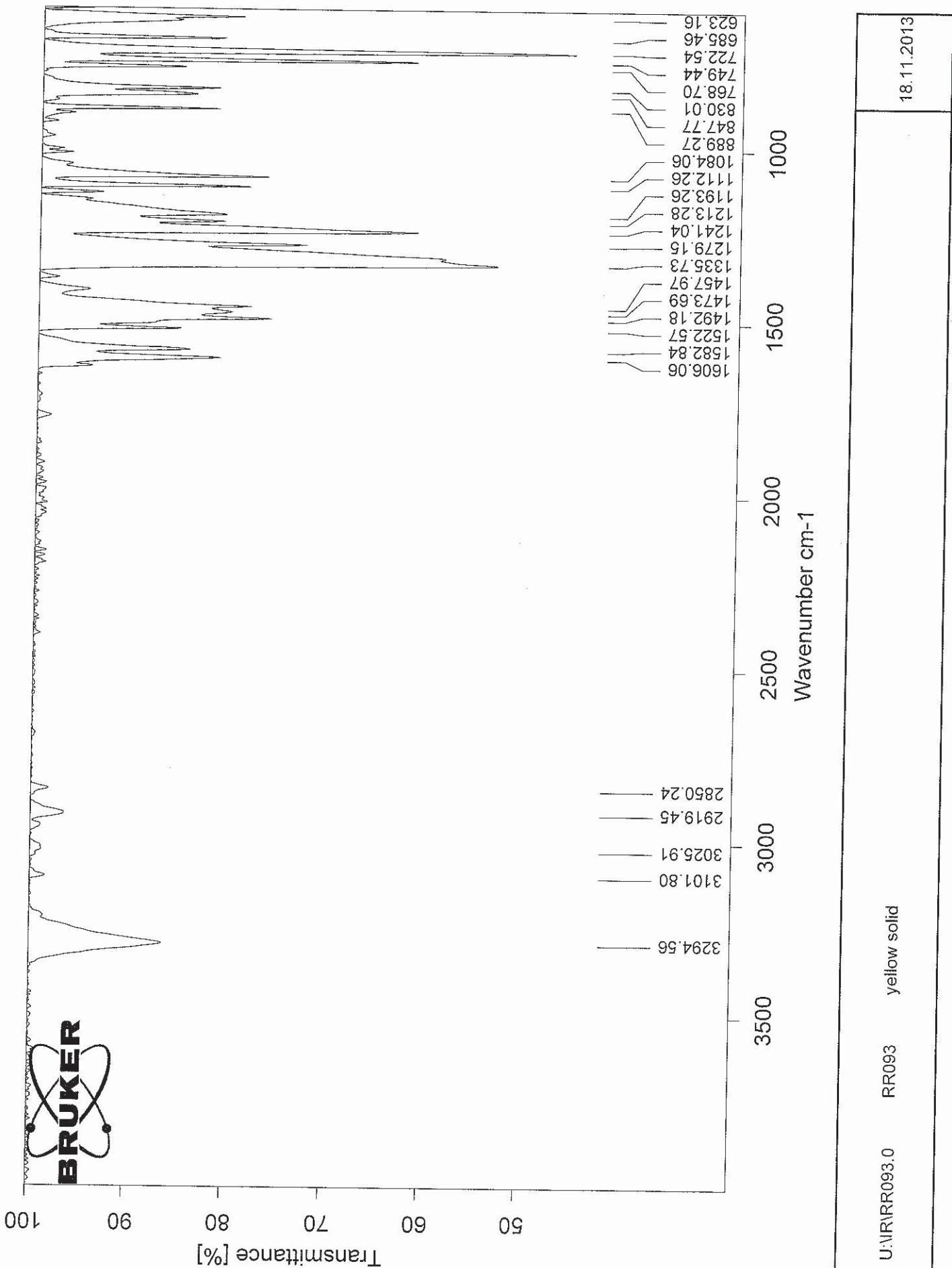
M08(m)

M09(m)

S159









ESI-MS: RR093M

XMASS Mass Analysis for /Data/UNI_FR/REMY7736_ESI/1/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

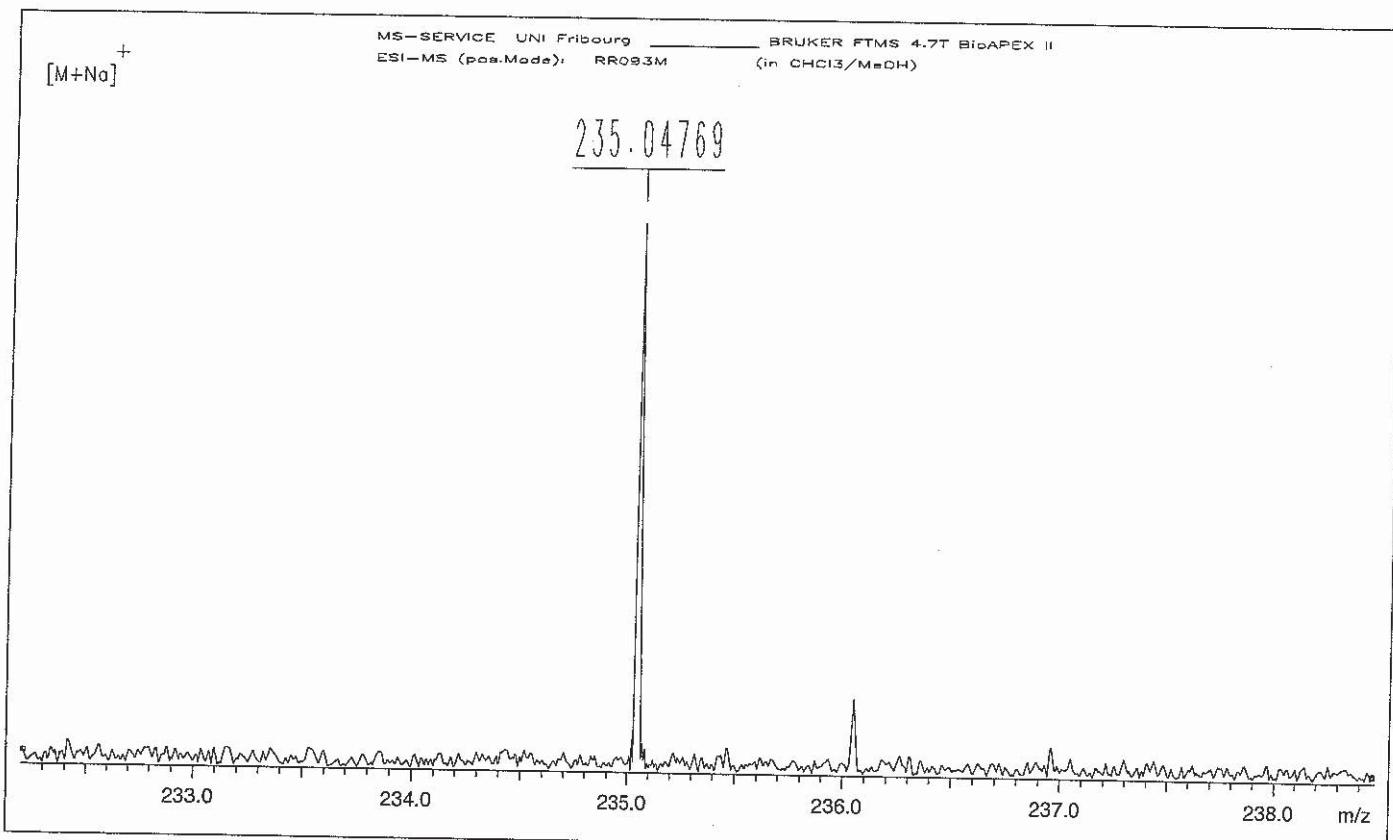
Ion mass = 235.0476920

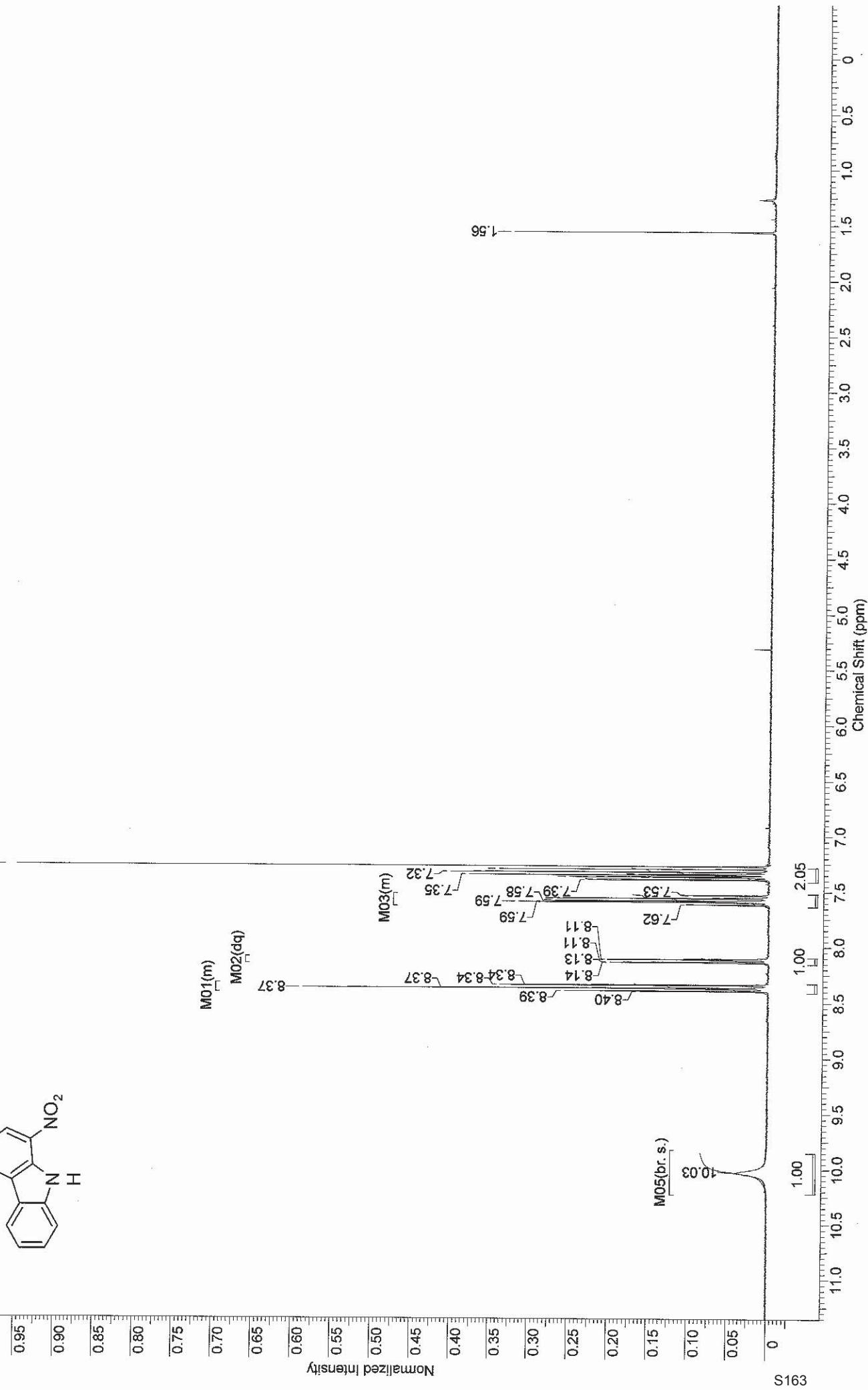
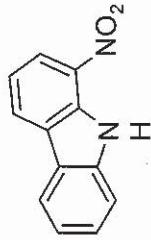
Charge = +1

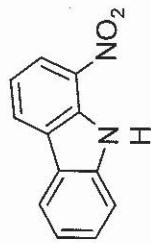
#	C	H	N	O	Na	mass	DBE	error
---	---	---	---	---	----	------	-----	-------

*** Mass Analysis for mass 235.0476920

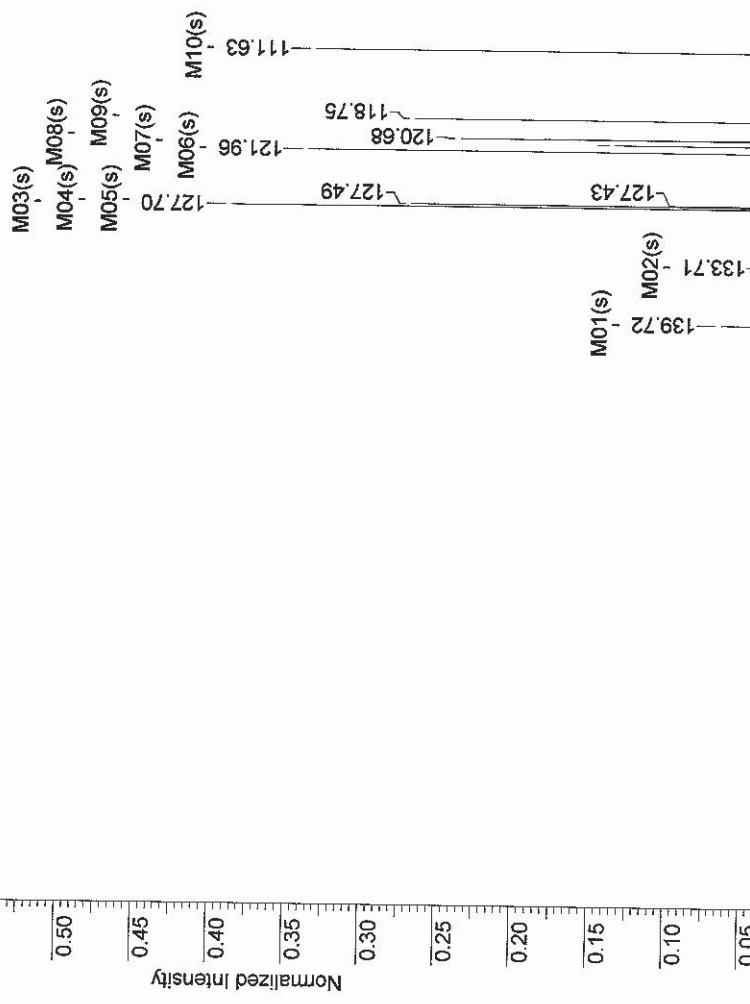
1	12	8	2	2	1	235.0477986	9.5	1.066e-04
2	11	9	1	5	0	235.0475238	8.0	1.682e-04
3	12	5	5	1	0	235.0488612	13.0	1.169e-03
4	10	6	5	1	1	235.0464559	10.0	1.236e-03
5	9	7	4	4	0	235.0461812	8.5	1.511e-03
6	14	7	2	2	0	235.0502039	12.5	2.512e-03
7	9	10	1	5	1	235.0451185	5.0	2.573e-03
8	7	5	7	3	0	235.0448385	9.0	2.854e-03
9	7	8	4	4	1	235.0437759	5.5	3.916e-03
10	6	9	3	7	0	235.0435011	4.0	4.191e-03

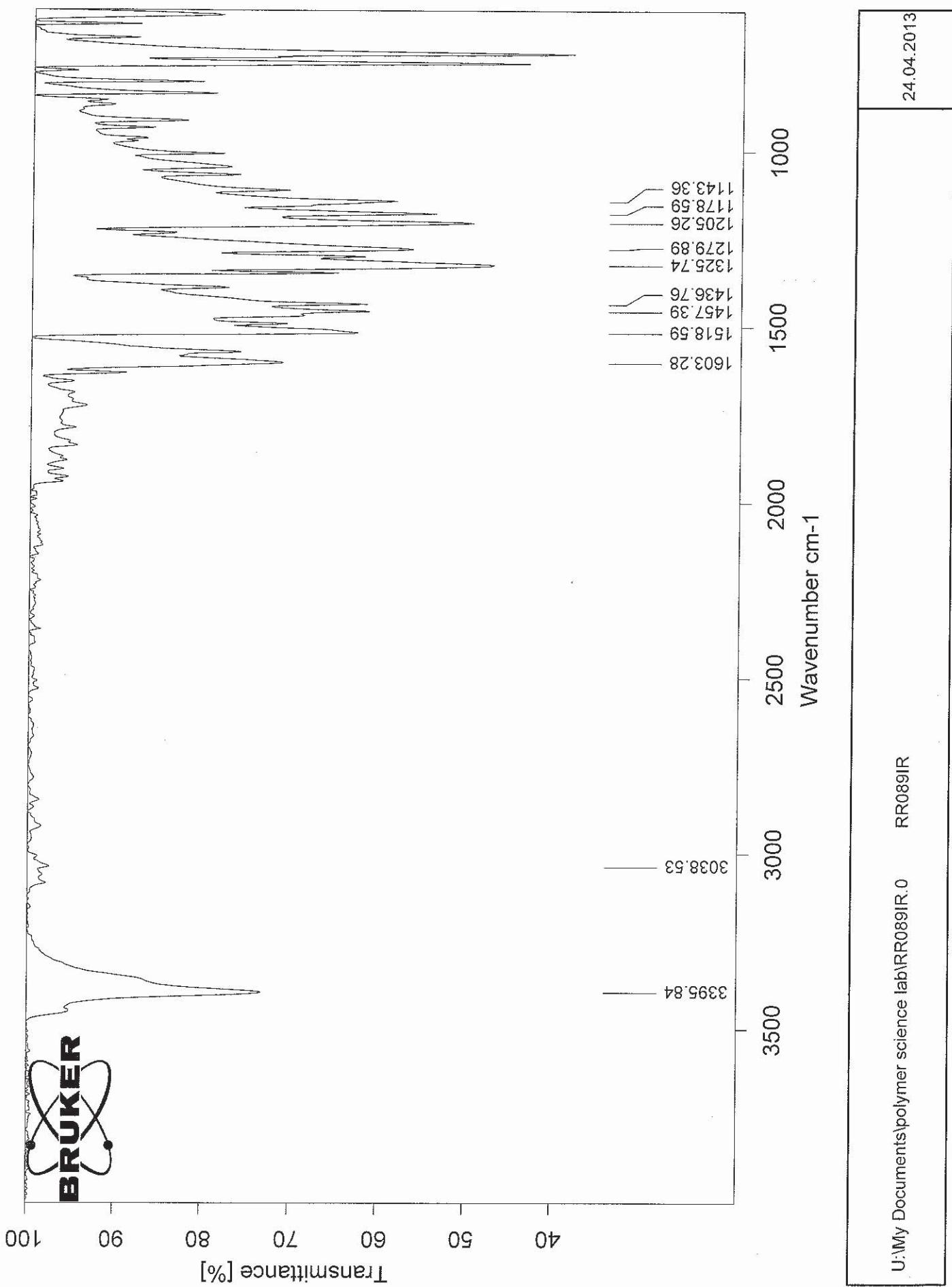






77.42
77.00
76.58





U:\My Documents\polymer science lab\RR089IR.0

RR089IR

24.04.2013



ESI-MS: RR0893M

XMASS Mass Analysis for /Data/UNI_FR/REMY7735_ESI/1/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

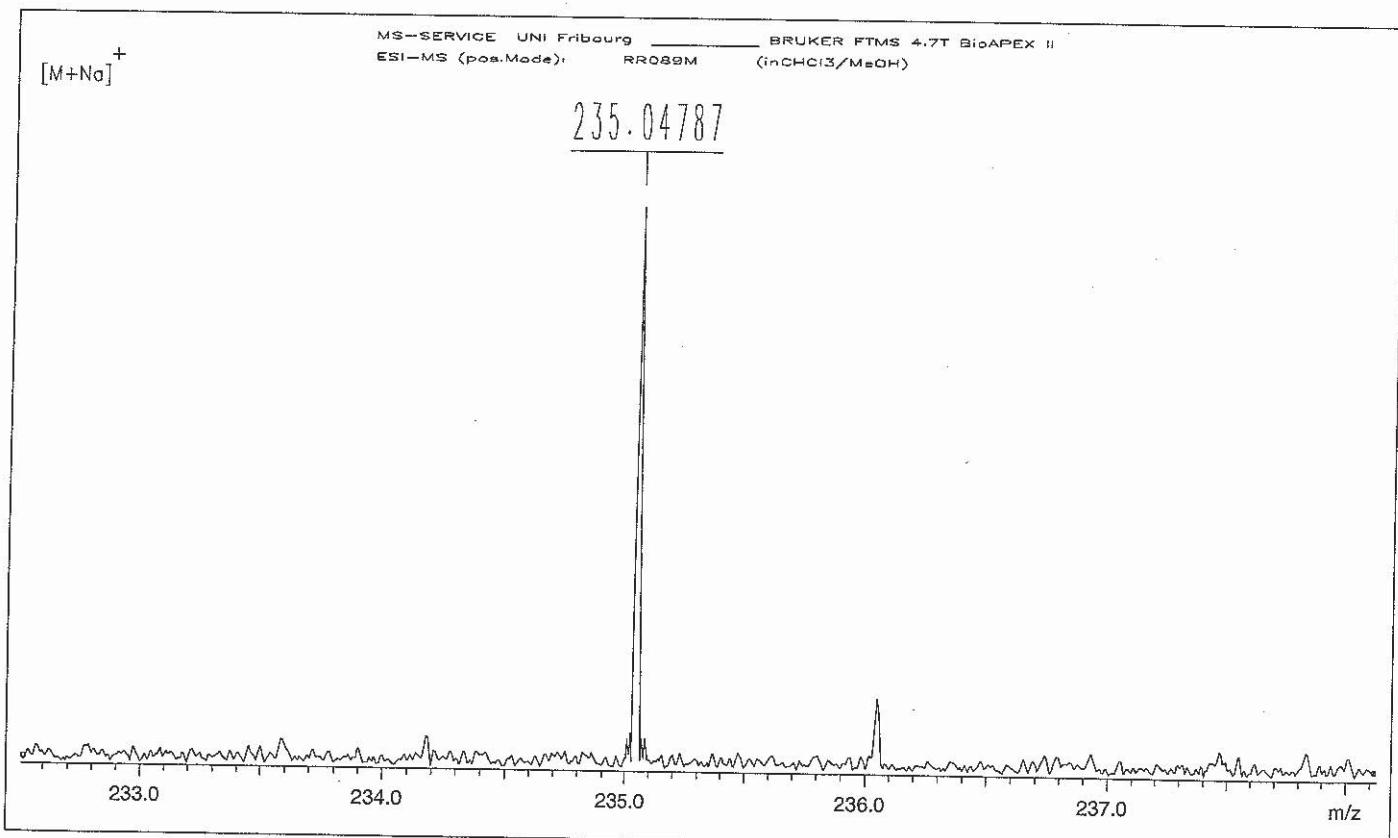
Ion mass = 235.0478660

Charge = +1

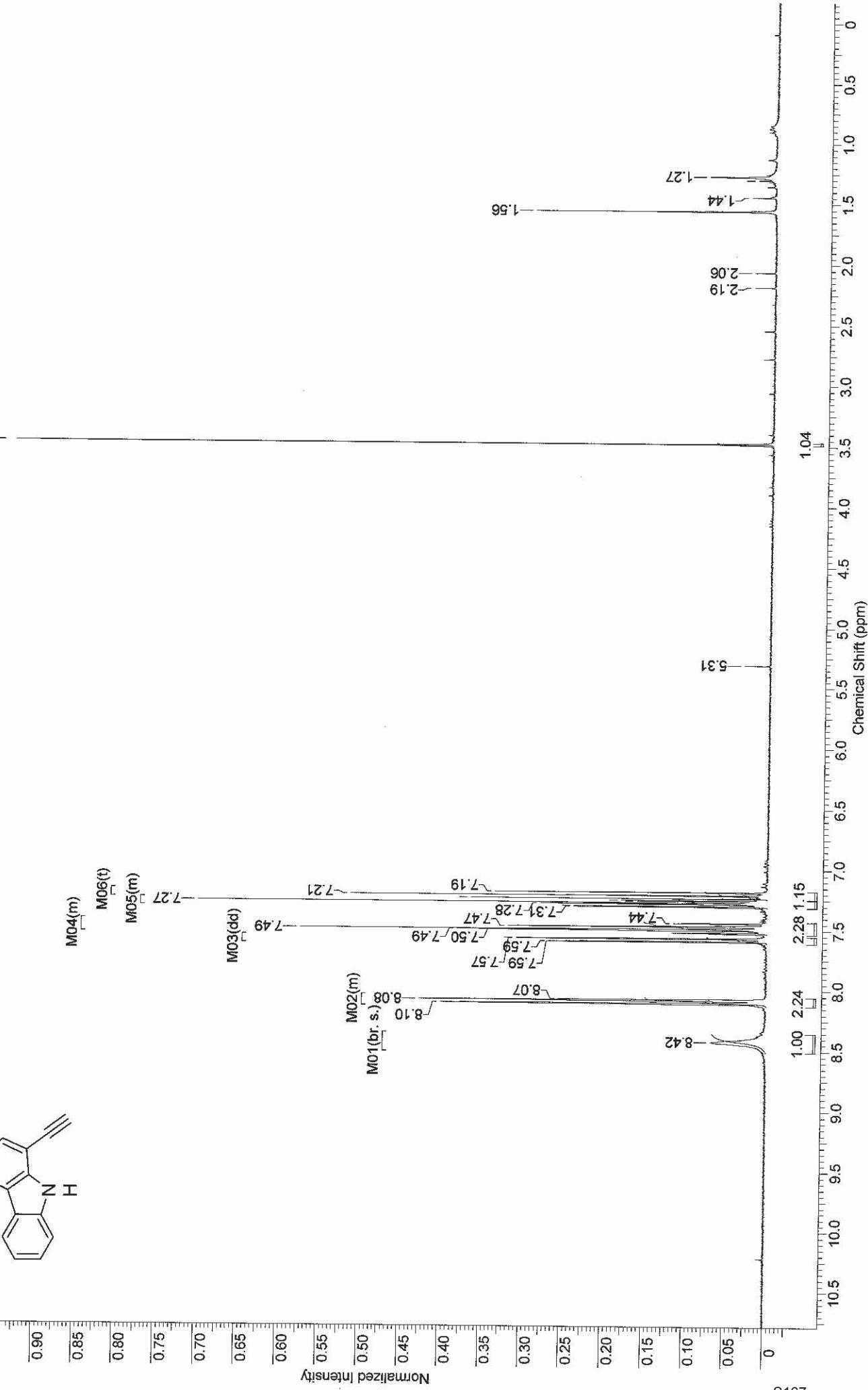
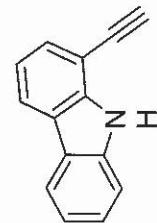
#	C	H	N	O	Na	mass	DBE	error
---	---	---	---	---	----	------	-----	-------

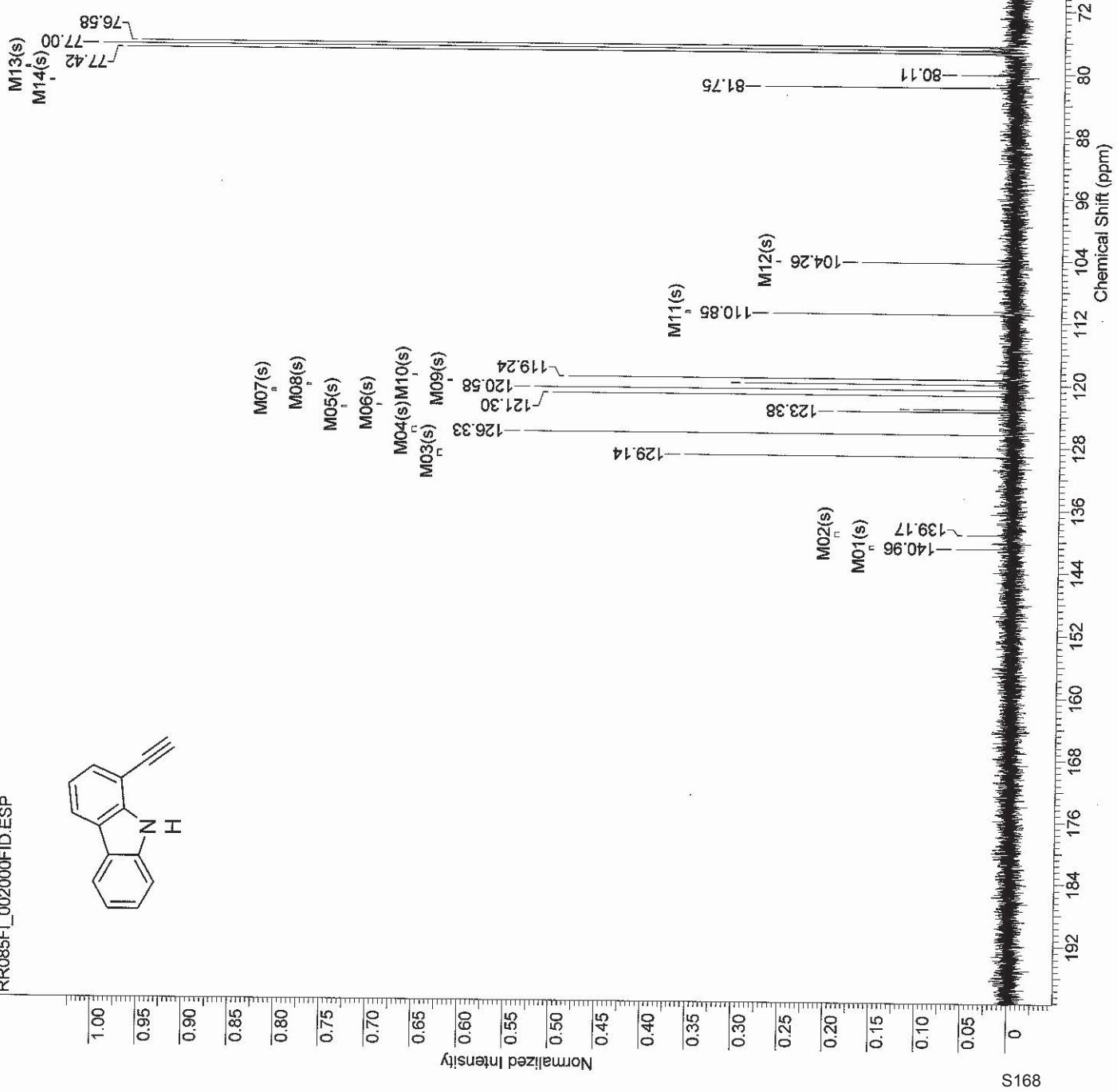
*** Mass Analysis for mass 235.0478660

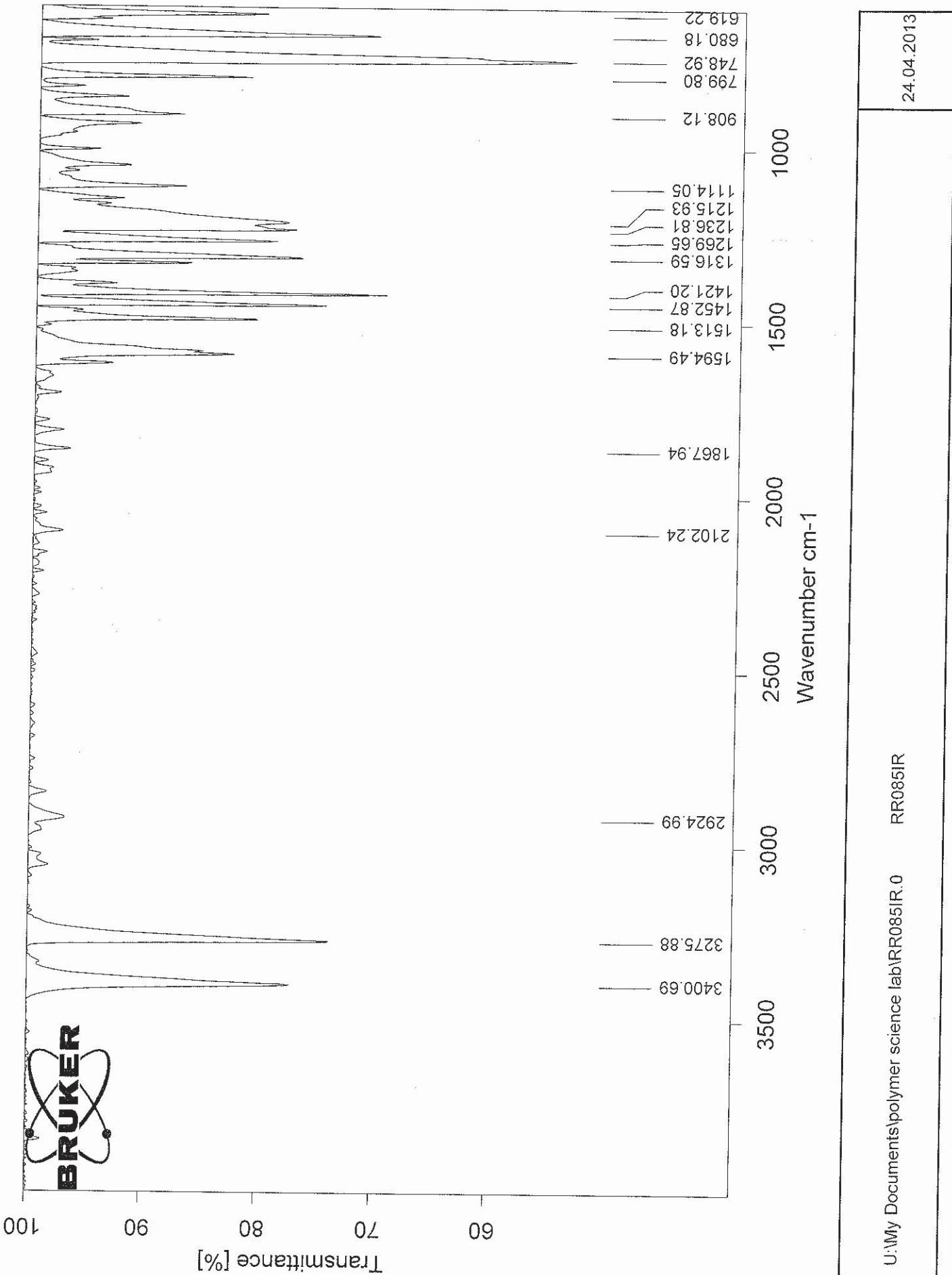
1	12	8	2	2	1	235.0477986	9.5	6.740e-05
2	11	9	1	5	0	235.0475238	8.0	3.422e-04
3	12	5	5	1	0	235.0488612	13.0	9.952e-04
4	10	6	5	1	1	235.0464559	10.0	1.410e-03
5	9	7	4	4	0	235.0461812	8.5	1.685e-03
6	14	7	2	2	0	235.0502039	12.5	2.338e-03
7	9	10	1	5	1	235.0451185	5.0	2.747e-03
8	7	5	7	3	0	235.0448385	9.0	3.028e-03
9	7	8	4	4	1	235.0437759	5.5	4.090e-03
10	6	9	3	7	0	235.0435011	4.0	4.365e-03



M07(s)







C:\Xcalibur\..\REMY7813_EI

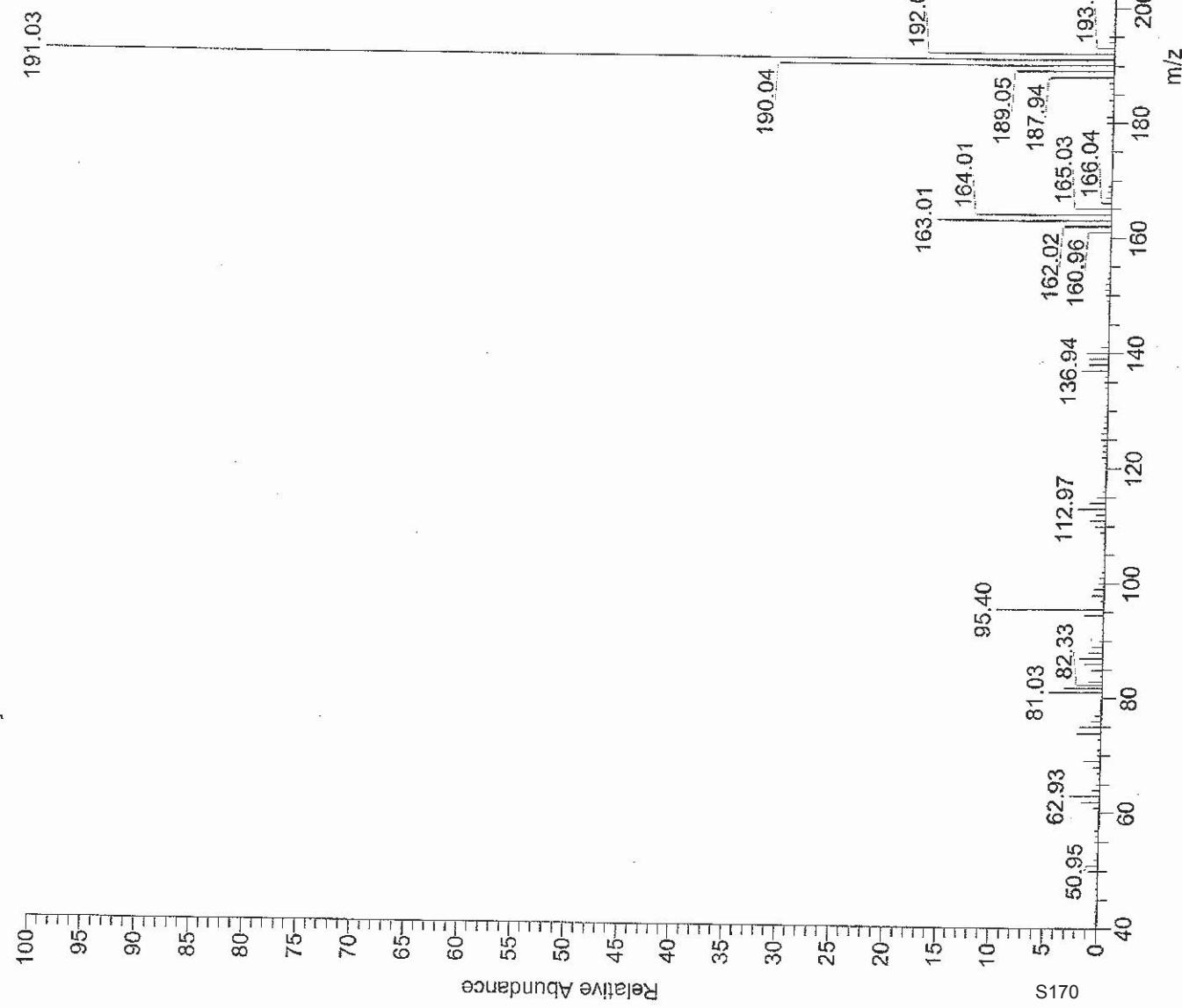
REMY7813_EI #51-65 RT: 0.55-0.69 AV: 15 NL: 7.33E8
T: + c Full ms [40.00-650.00]

12/20/2013 9:35:29 AM

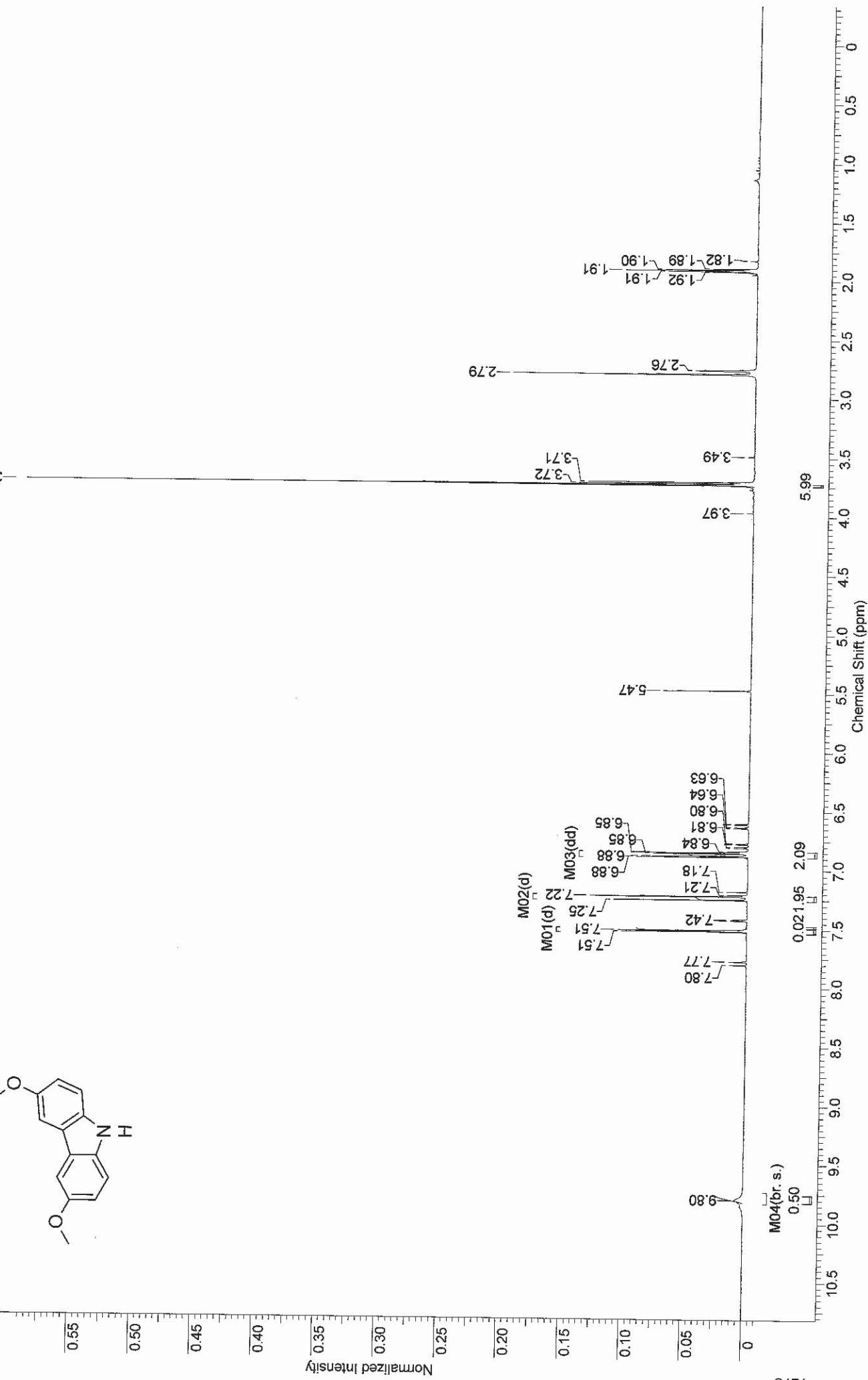
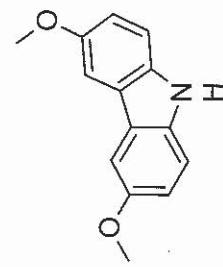
RR085M

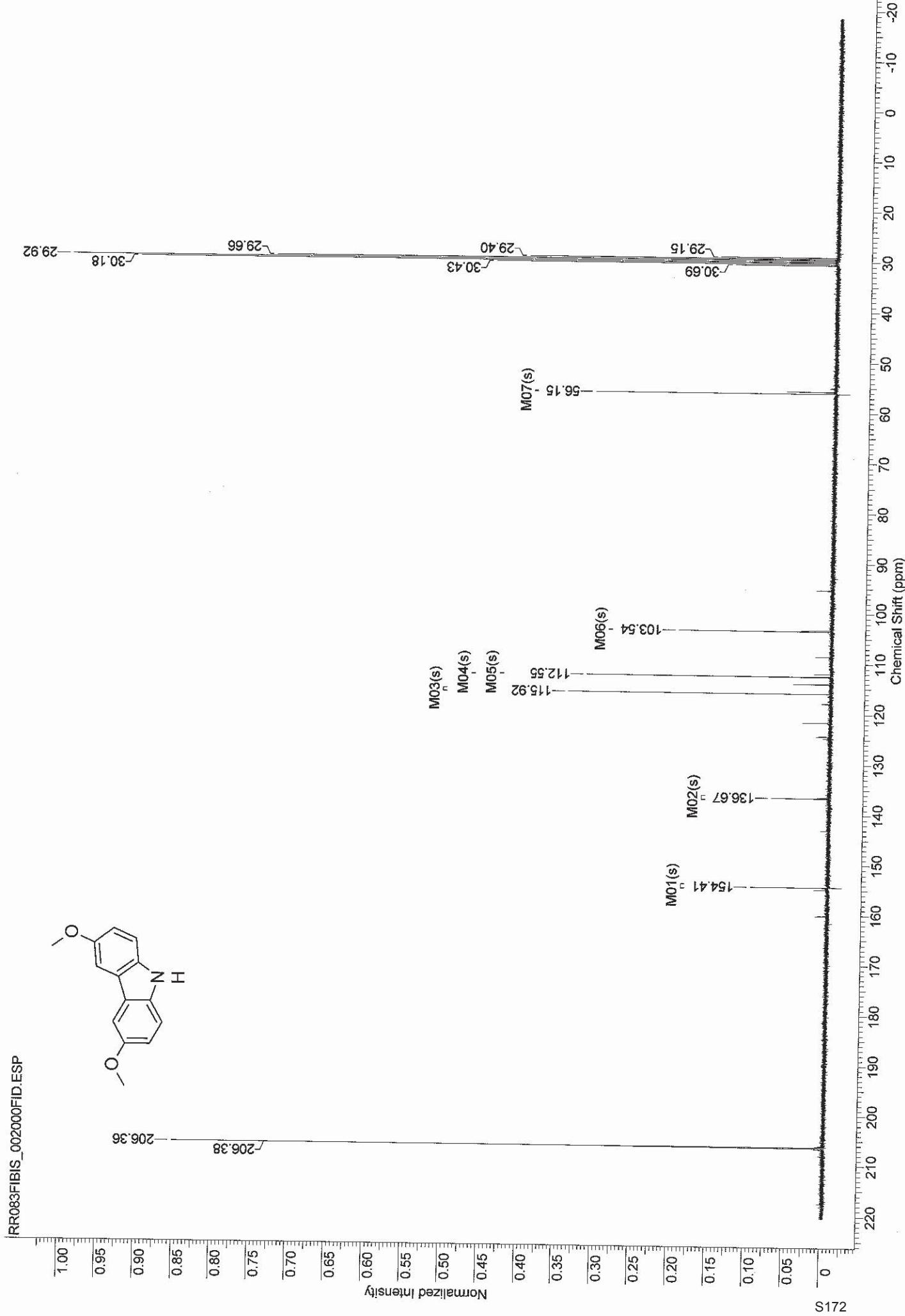
100
95
90
85
80
75
70
65
60
55
50
45
40
35
30
25
20
15
10
5
0

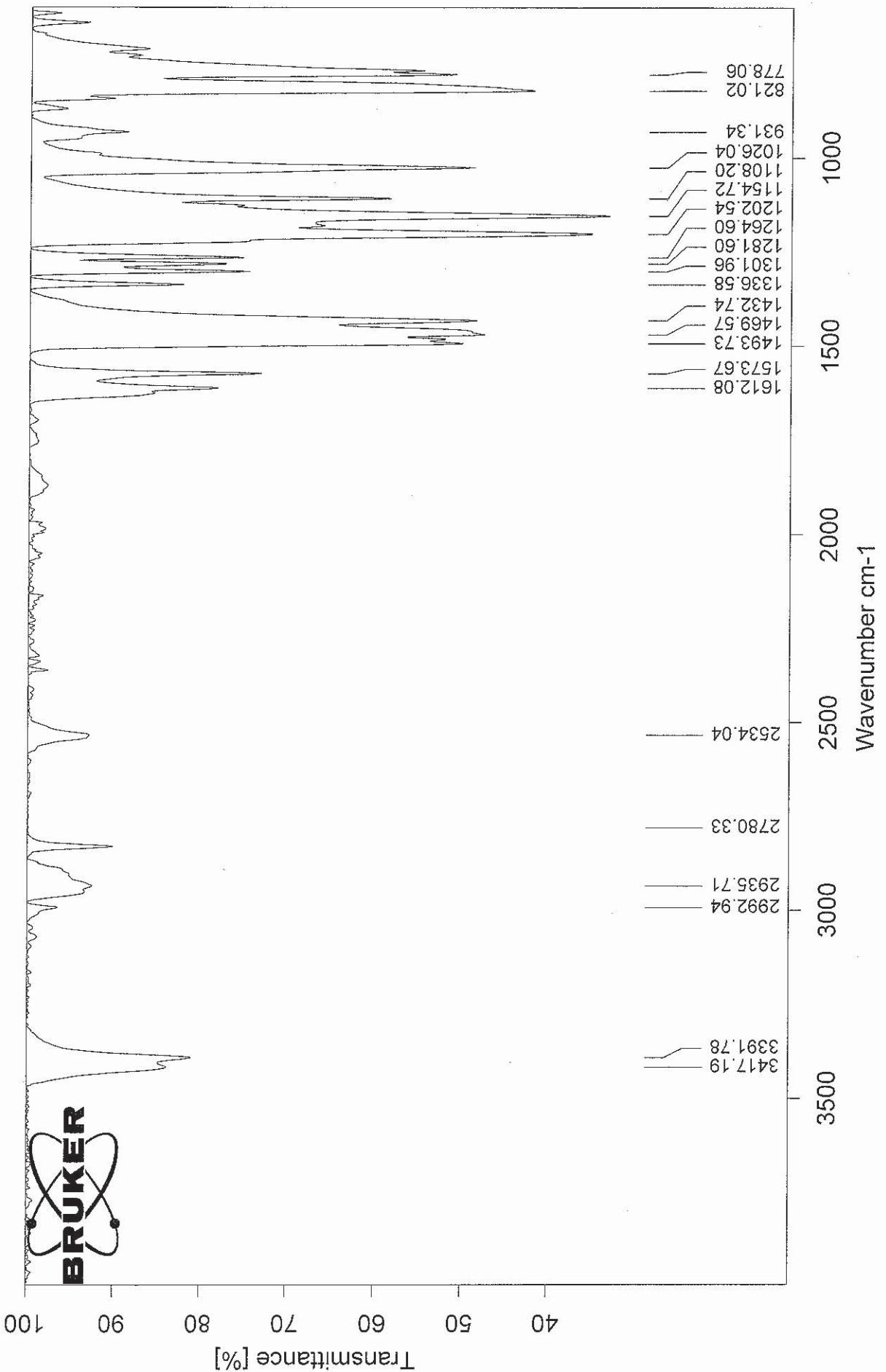
Relative Abundance



M05(s)







U:\IR\RR083D.0	RR083D	brown solid	10.12.2013
----------------	--------	-------------	------------

REMY7812_EI #61-82 RT: 0.67-0.89 AV: 22 NL: 3.95E8
T: +c Full ms [40.00-650.00]

100
95
90
85
80
75
70
65
60
55
50
45
40
35
30
25
20
15
10
5
0

227.06

211.98

228.10

184.01

229.12
240.06

185.05
197.02

141.00

82.75

42.81
46.82

S174

142.01

125.98

113.89

86.89

74.94

62.94

46.82

84.79

113.03

139.92

142.01

154.02

169.00

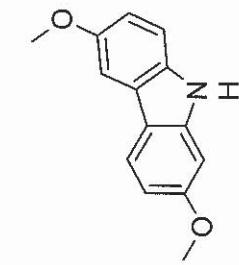
185.05

197.02

Relative Abundance

340
320
300
280
260
240
220
200
180
160
140
120
100
80
60
40

M07(s)



M08(s)

3.85

3.86

3.88

3.91

3.88

3.73

2.90

2.94

2.06

2.06

2.04

2.04

2.05

2.05

2.06

2.06

2.04

2.04

2.05

2.05

2.06

2.06

2.04

2.04

2.05

2.05

2.06

2.06

2.04

2.04

2.05

2.05

2.06

2.06

2.04

2.04

2.05

2.05

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

2.06

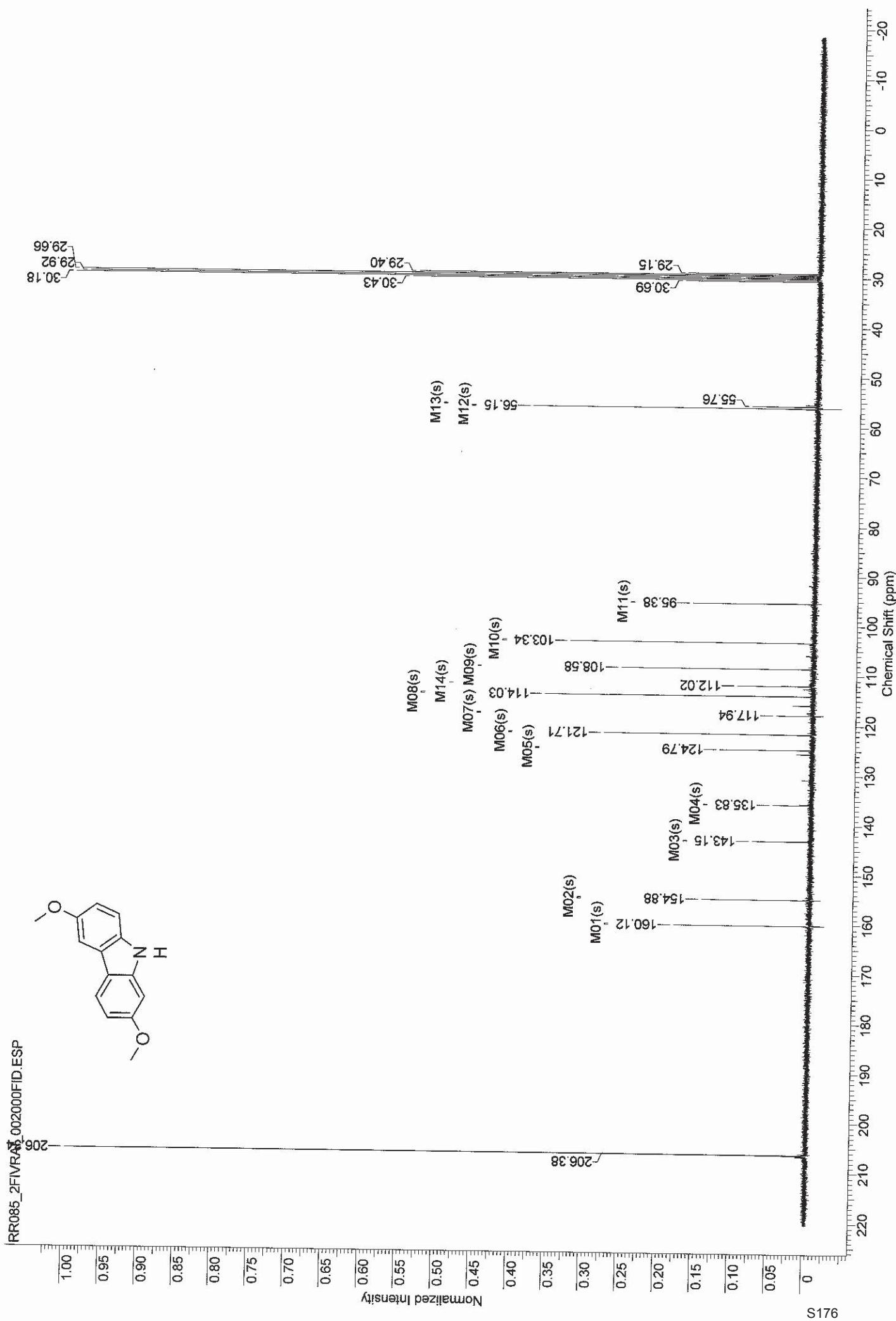
2.06

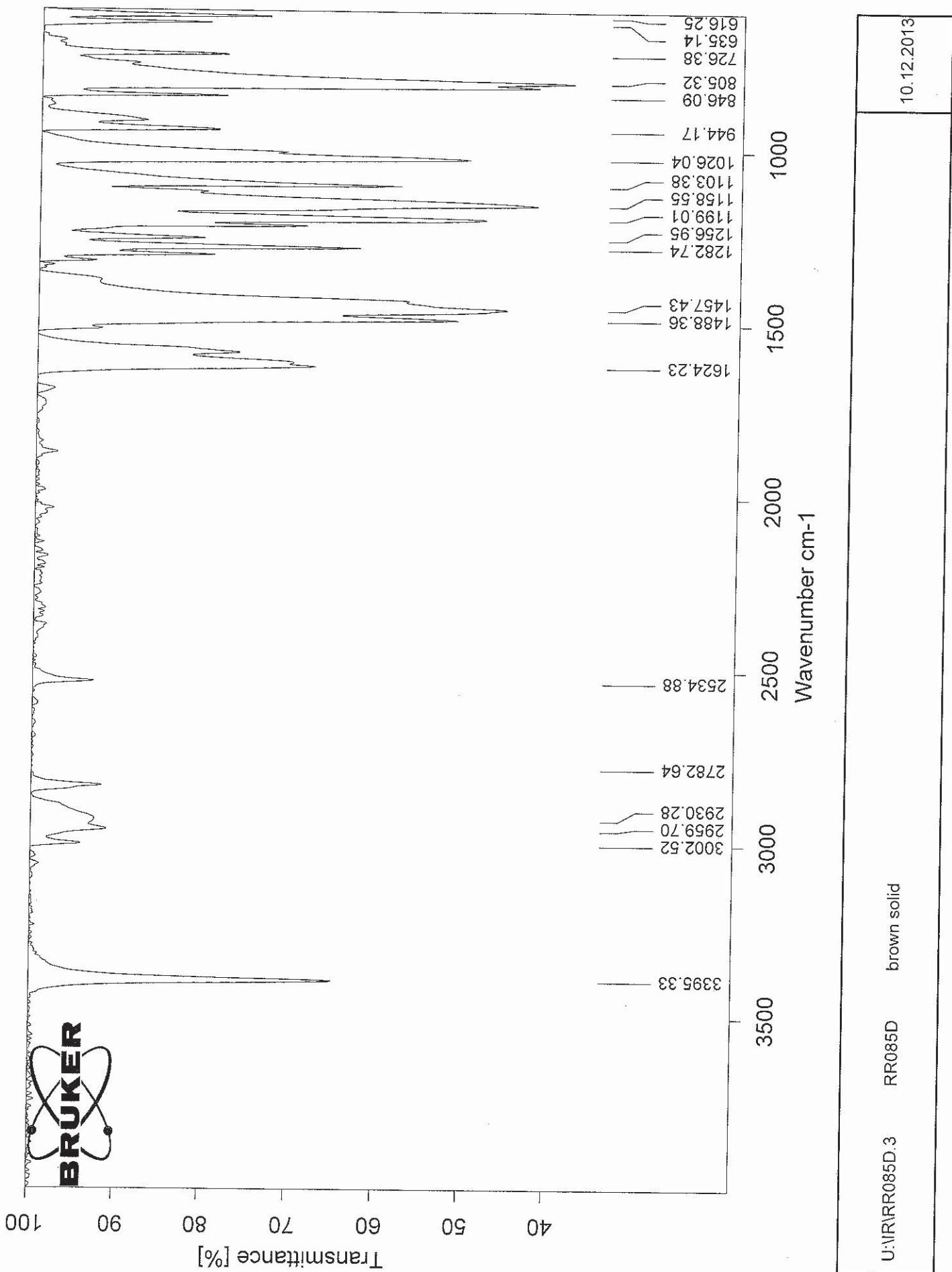
2.06

2.06

2.06

2.06



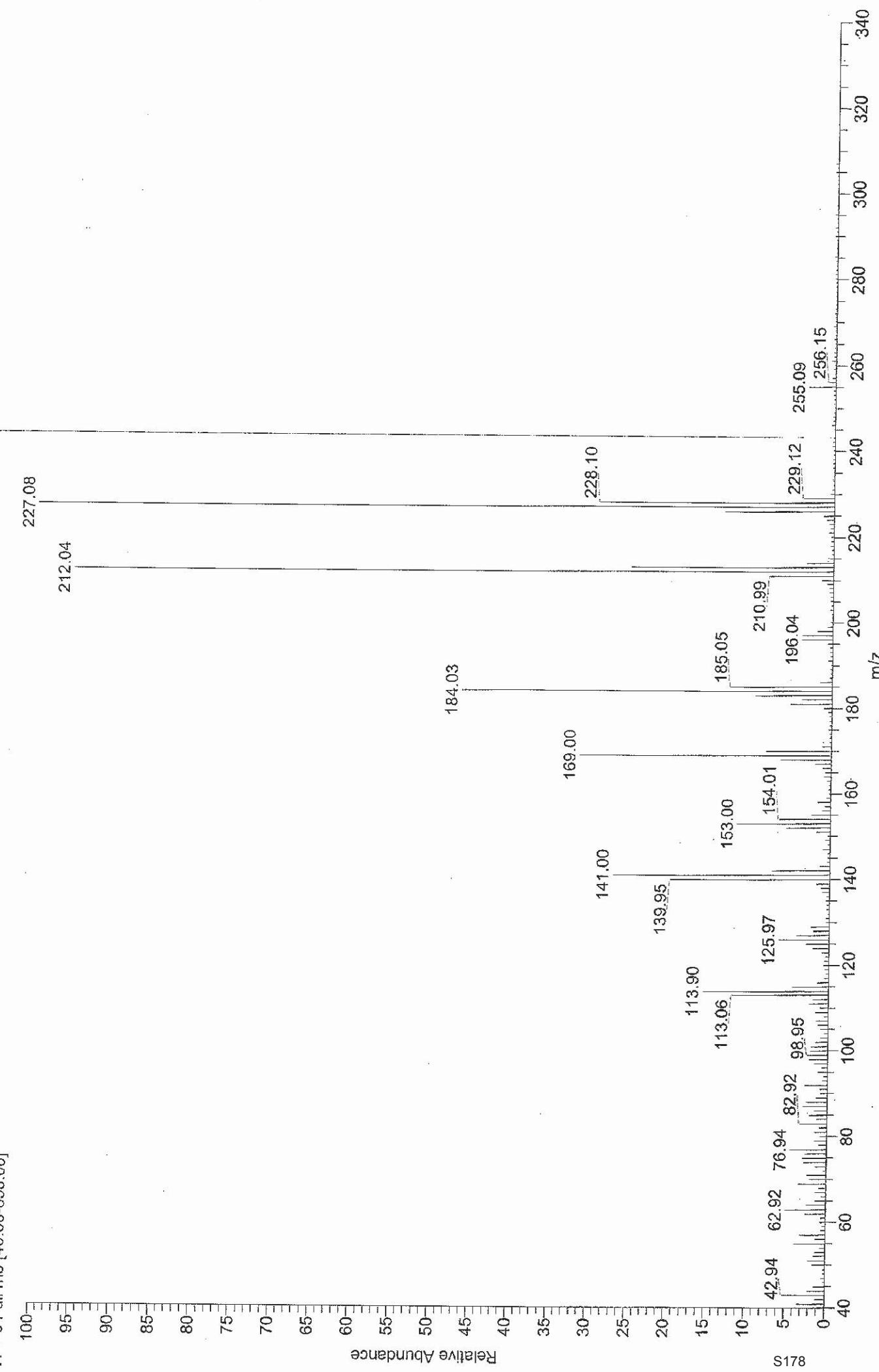


C:\Xcalibur\...\REMY7811_EI

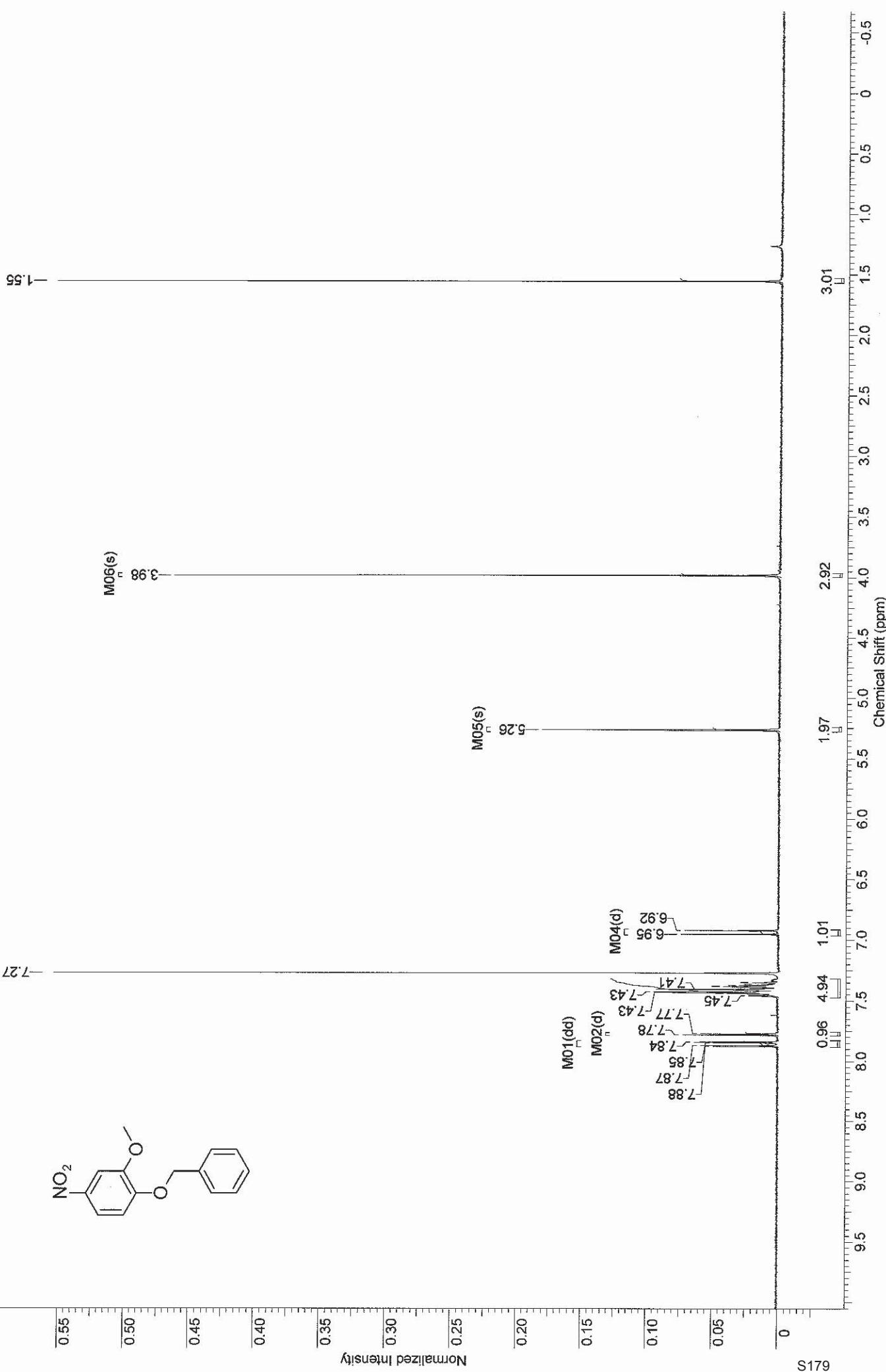
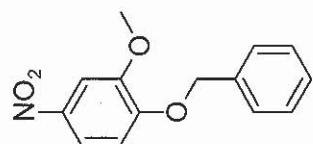
REMY7811_EI #75-93 RT: 0.81-1.00 AV: 19 NL: 5.67E7
T: + c Full ms [40.00-650.00]

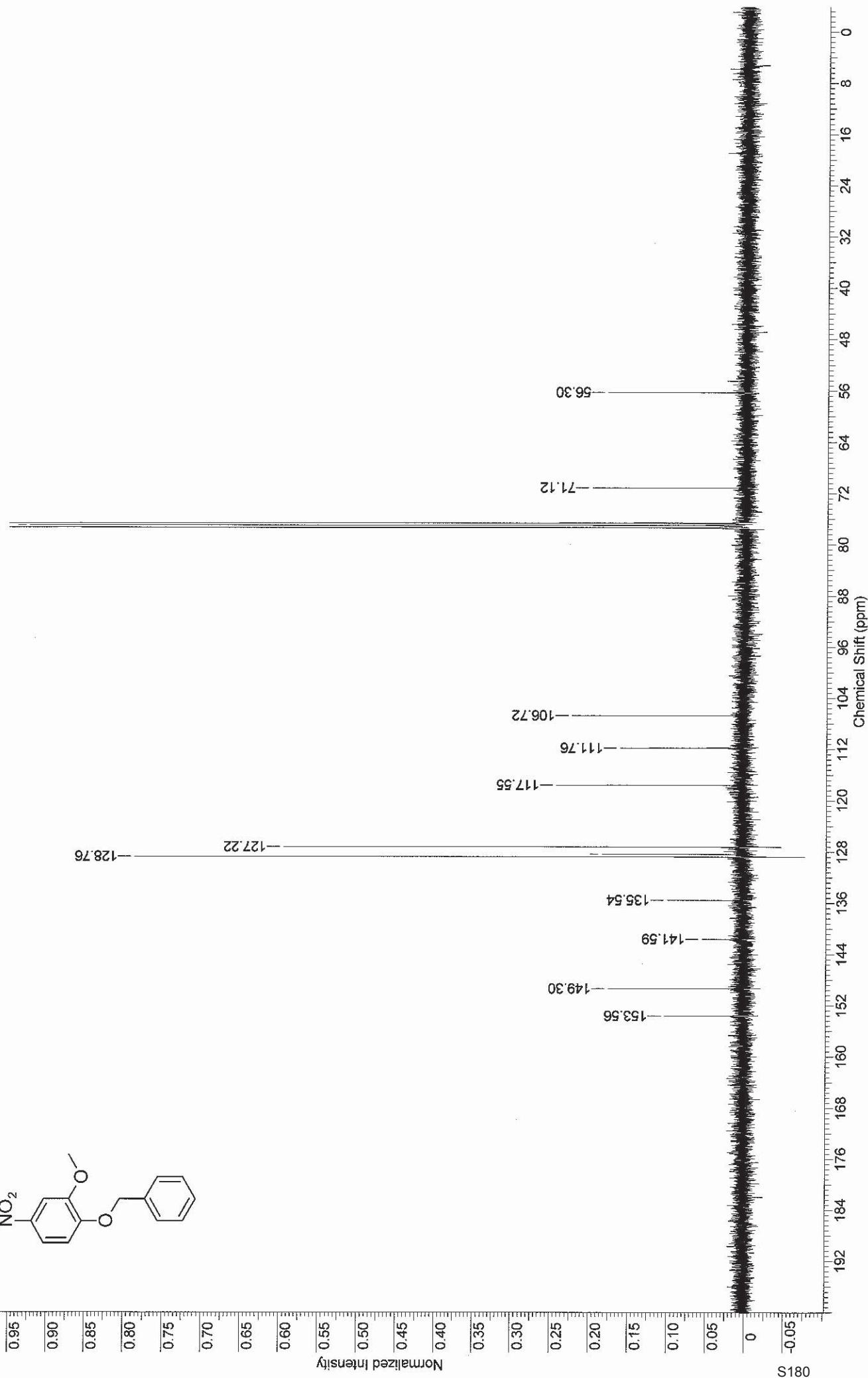
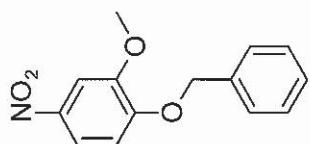
12/20/2013 8:57:39 AM

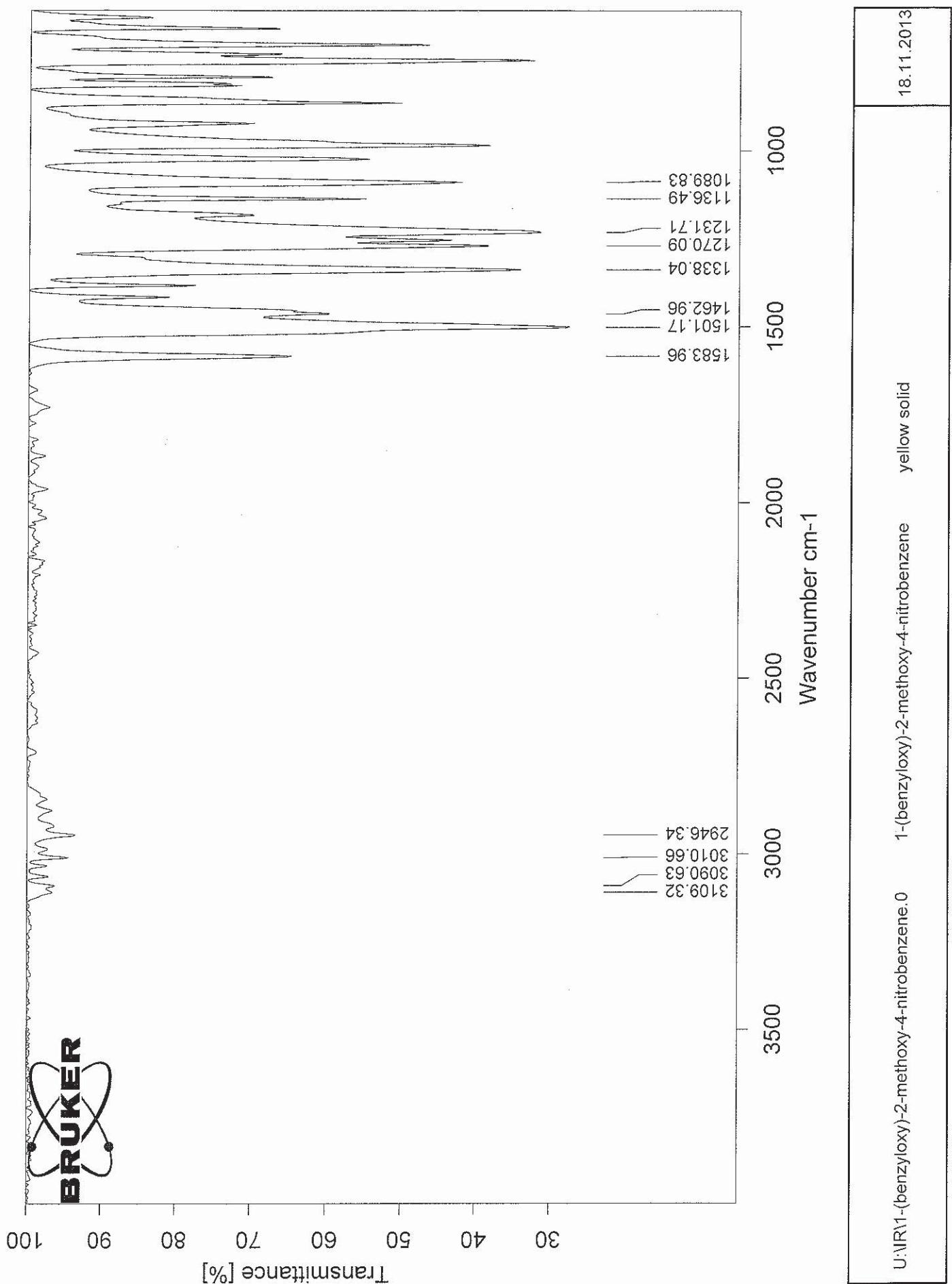
RR085D



S178







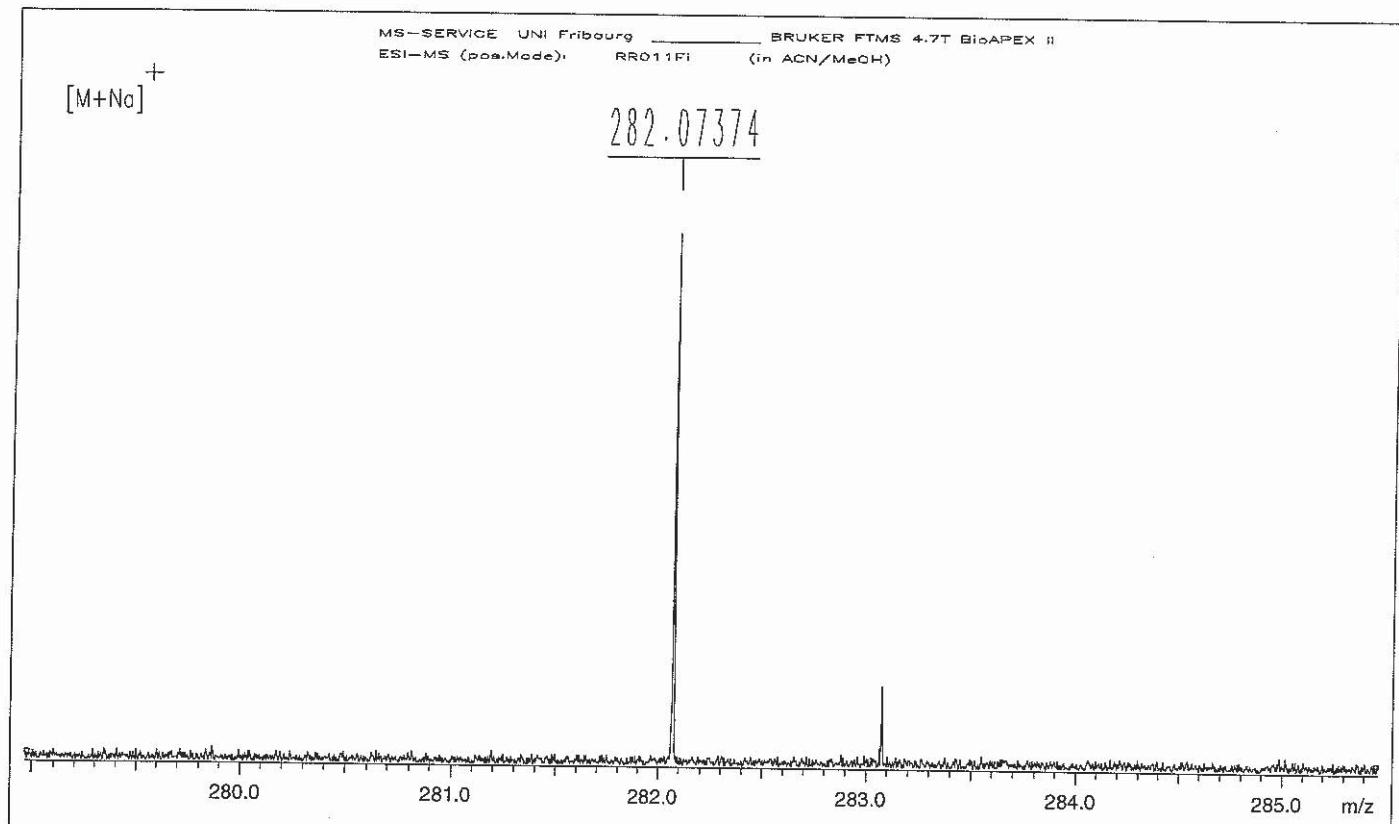
ESI-MS: RR011Fi

XMASS Mass Analysis Constraints

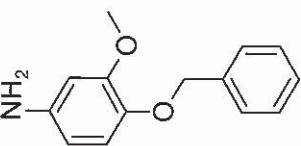
Ion mass = 282.0737410

Charge = +1

#	C	H	N	O	Na	mass	DBE	error
*** Mass Analysis for mass 282.0737410								
1	14	13	1	4	1	282.0736790	8.5	6.200e-05
2	12	8	7	2	0	282.0733990	12.5	3.420e-04
3	14	10	4	3	0	282.0747416	12.0	1.001e-03
4	12	11	4	3	1	282.0723363	9.0	1.405e-03
5	11	12	3	6	0	282.0720616	7.5	1.679e-03
6	10	6	10	1	0	282.0720563	13.0	1.685e-03
7	16	12	1	4	0	282.0760843	11.5	2.343e-03
8	17	11	2	1	1	282.0763591	13.0	2.618e-03
9	10	9	7	2	1	282.0709937	9.5	2.747e-03
10	9	10	6	5	0	282.0707189	8.0	3.022e-03



M01(s)



3.89

M02(s)

5.13

M03(m)

6.56

M05(m)

7.38

7.42

7.43

7.45

7.48

7.51

7.54

7.58

7.61

7.64

7.68

7.71

7.74

7.77

7.80

7.83

7.86

7.89

7.92

7.95

7.98

8.01

8.04

8.07

8.10

8.13

8.16

8.19

8.22

8.25

8.28

8.31

8.34

8.37

8.40

8.43

8.46

8.49

8.52

8.55

8.58

8.61

8.64

8.67

8.70

8.73

8.76

8.79

8.82

8.85

8.88

8.91

8.94

8.97

9.00

9.03

9.06

9.09

9.12

9.15

9.18

9.21

9.24

9.27

9.30

9.33

9.36

9.39

9.42

9.45

9.48

9.51

9.54

9.57

9.60

9.63

9.66

9.69

9.72

9.75

9.78

9.81

9.84

9.87

9.90

9.93

9.96

9.99

10.02

3.00

2.03

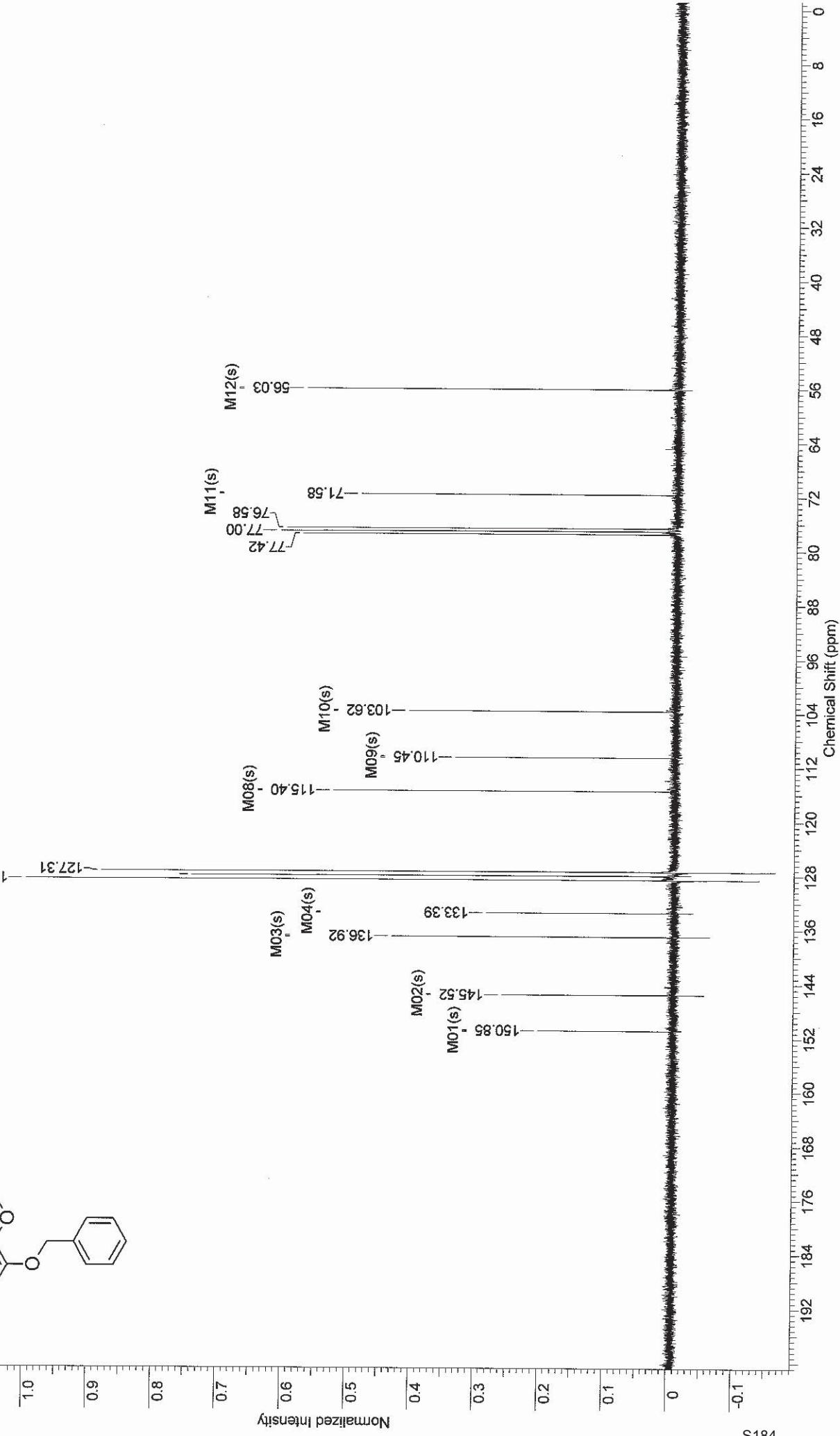
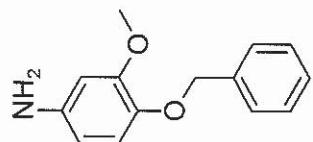
1.06

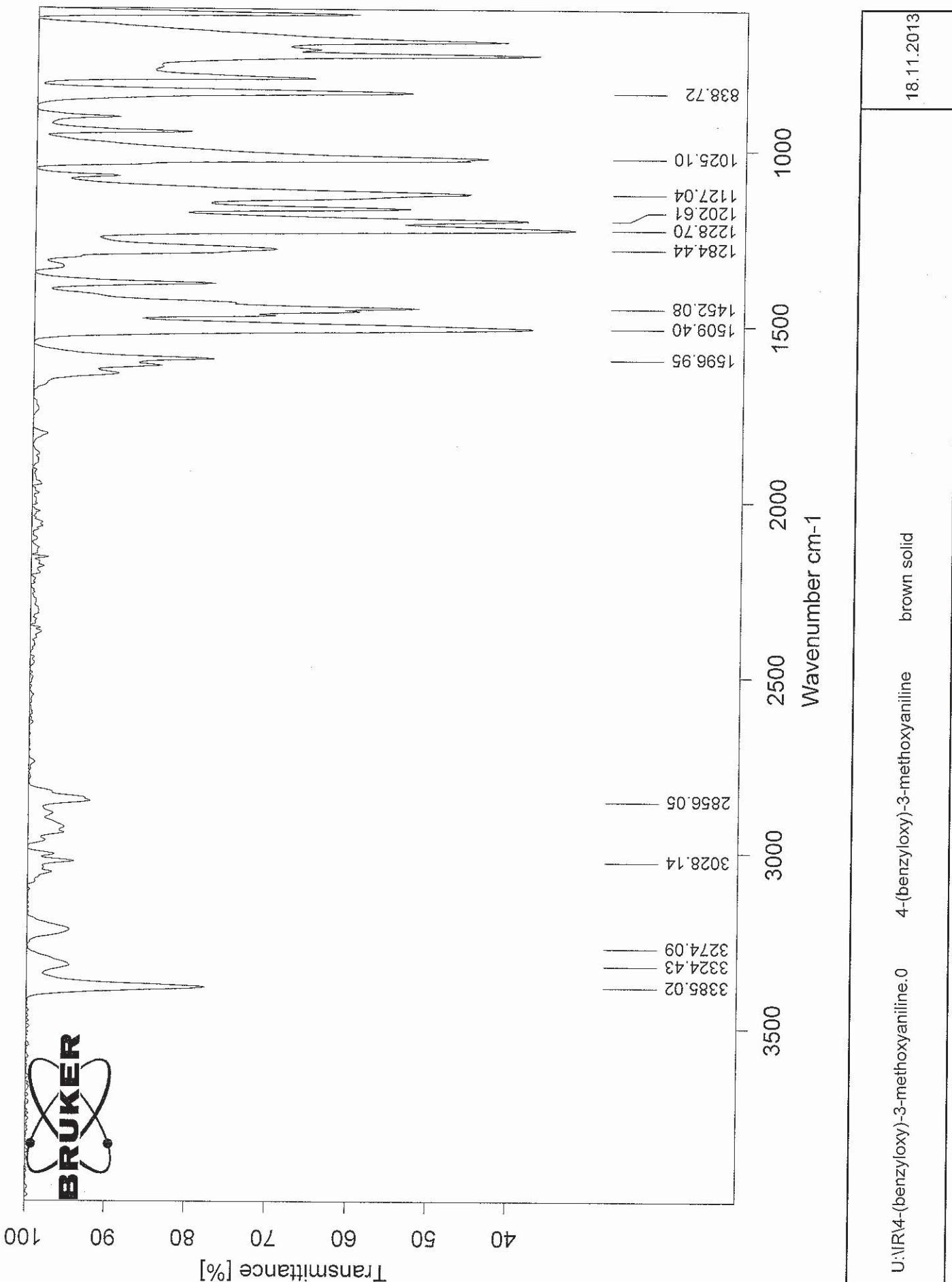
0.01

-0.01

0
0.5
1.0
1.5
2.0
2.5
3.0
3.5
4.0
4.5
5.0
5.5
6.0
6.5
7.0
7.5
8.0
8.5
9.0
9.5
10.0

M06(s)
M07(s)
M05(s)







ESI-MS: RR013FI

XMASS Mass Analysis for /Data/UNI_FR/REMY7651_ESI/1/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

Ion mass = 230.1176290

Charge = +1

#	C	H	N	O	mass	DBE	error
---	---	---	---	---	------	-----	-------

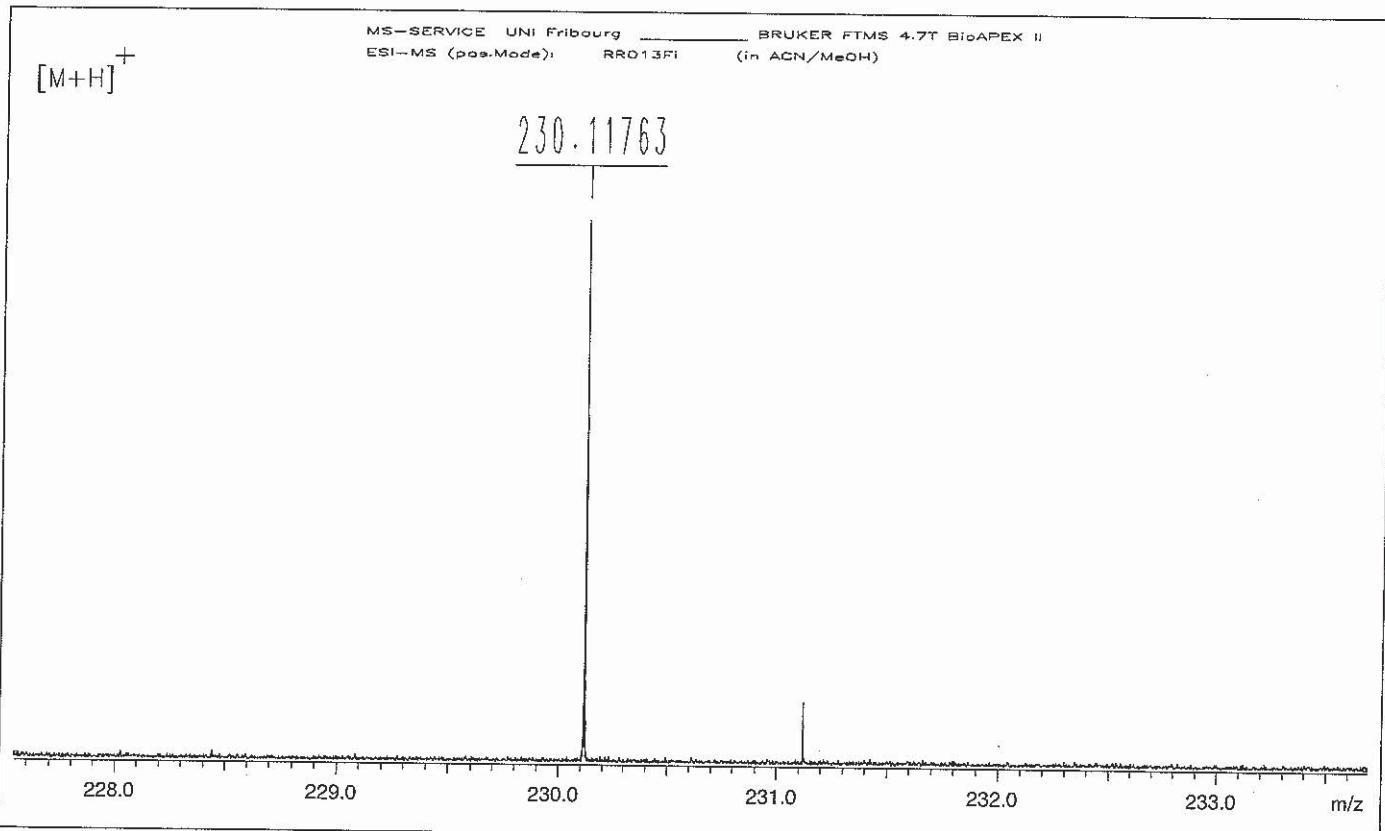
*** Mass Analysis for mass 230.1176290

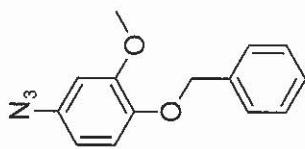
1	14	16	1	2	230.1175552	7.5	7.382e-05
2	12	14	4	1	230.1162125	8.0	1.416e-03
3	9	16	3	4	230.1135324	3.5	4.097e-03
4	5	18	4	6	230.1220858	-1.0	4.457e-03
5	7	14	6	3	230.1121898	4.0	5.439e-03
6	6	14	8	2	230.1234232	4.0	5.794e-03
7	7	20	1	7	230.1234284	-1.5	5.799e-03
8	6	18	2	7	230.1108524	-1.0	6.777e-03
9	5	12	9	2	230.1108471	4.5	6.782e-03
10	8	16	5	3	230.1247658	3.5	7.137e-03

MS-SERVICE UNI Fribourg BRUKER FTMS 4.7T BioAPEX II
ESI-MS (pos.-Mode): RR013FI (in ACN/MeOH)

[M+H]⁺

230.11763





0.45
0.40
0.35
0.30
0.25
0.20
0.15
0.10
0.05
0

Normalized Intensity

2.76
1.84
0.94
0.95
0.96
7.27

7.43 7.44 7.46

M02(d)
M03(d)
M04(dd)

5.31

6.15
6.16
6.18
6.19

6.32

6.33

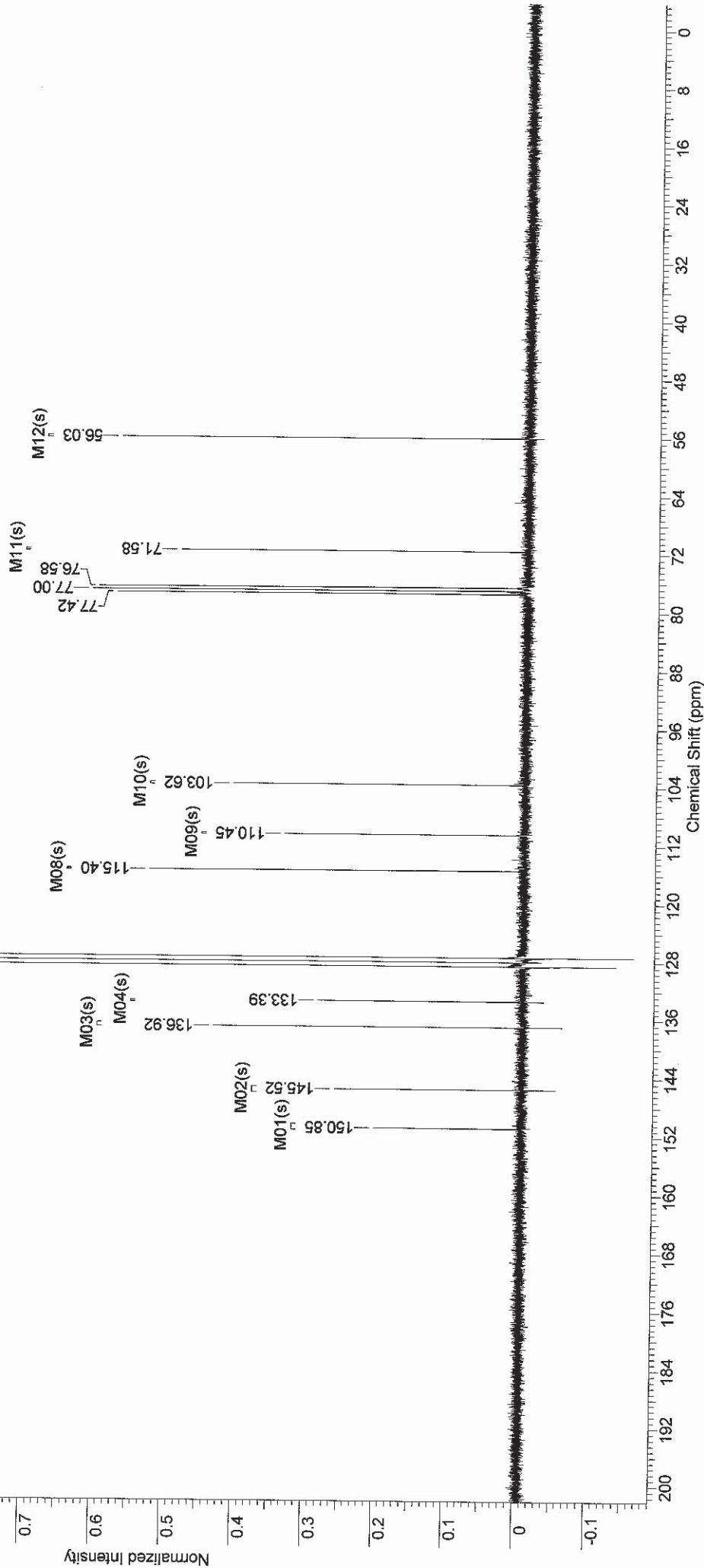
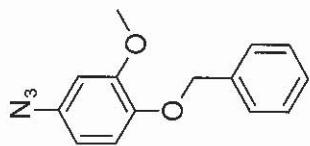
6.71

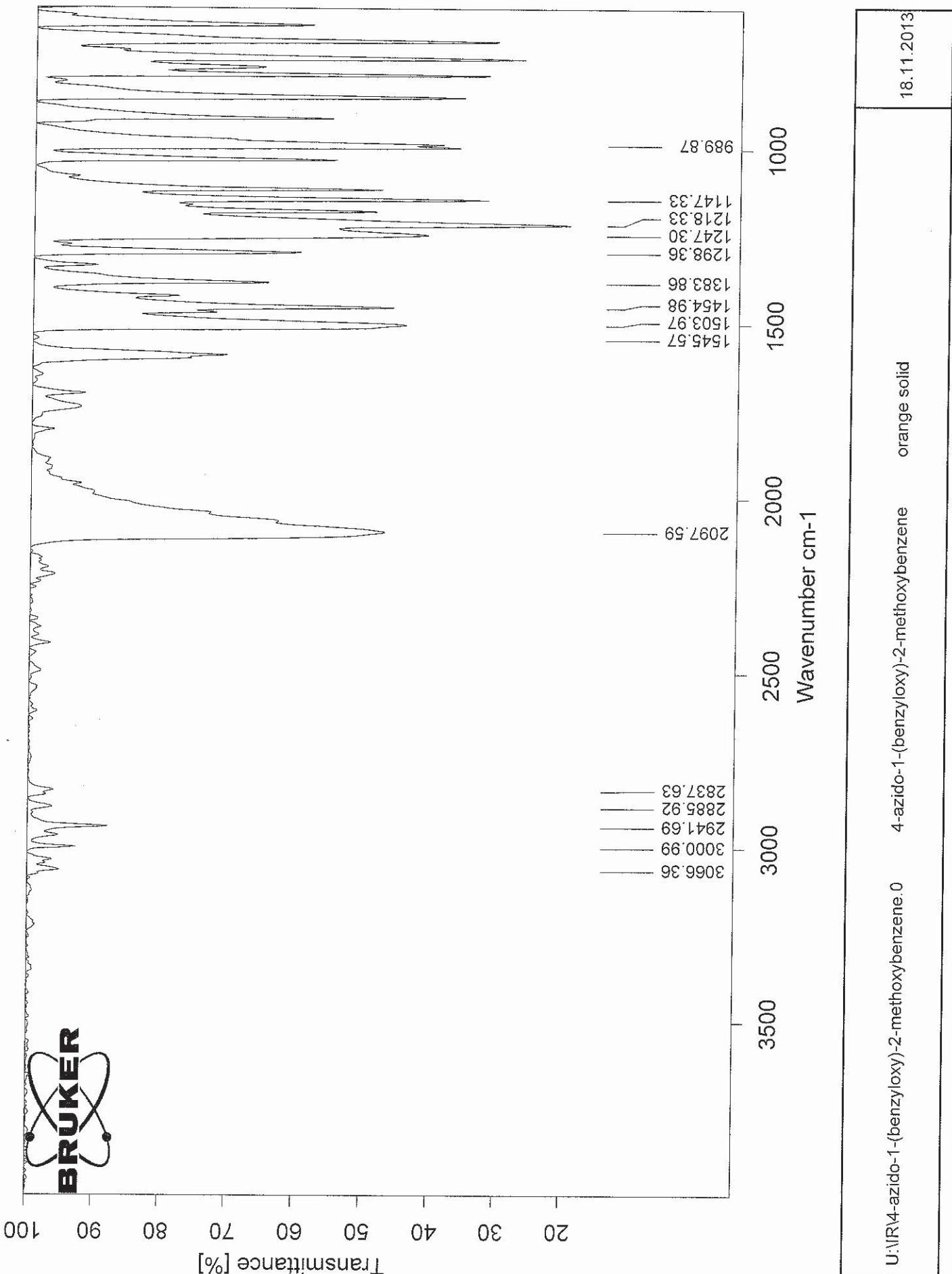
6.73

9.5 9.0 8.5 8.0 7.5 7.0 6.5 6.0 5.5 5.0 4.5 4.0 3.5 3.0 2.5 2.0 1.5 1.0 0.5 0

RR022FL_002000fid

M07(s)
M05(s)
M06(s)





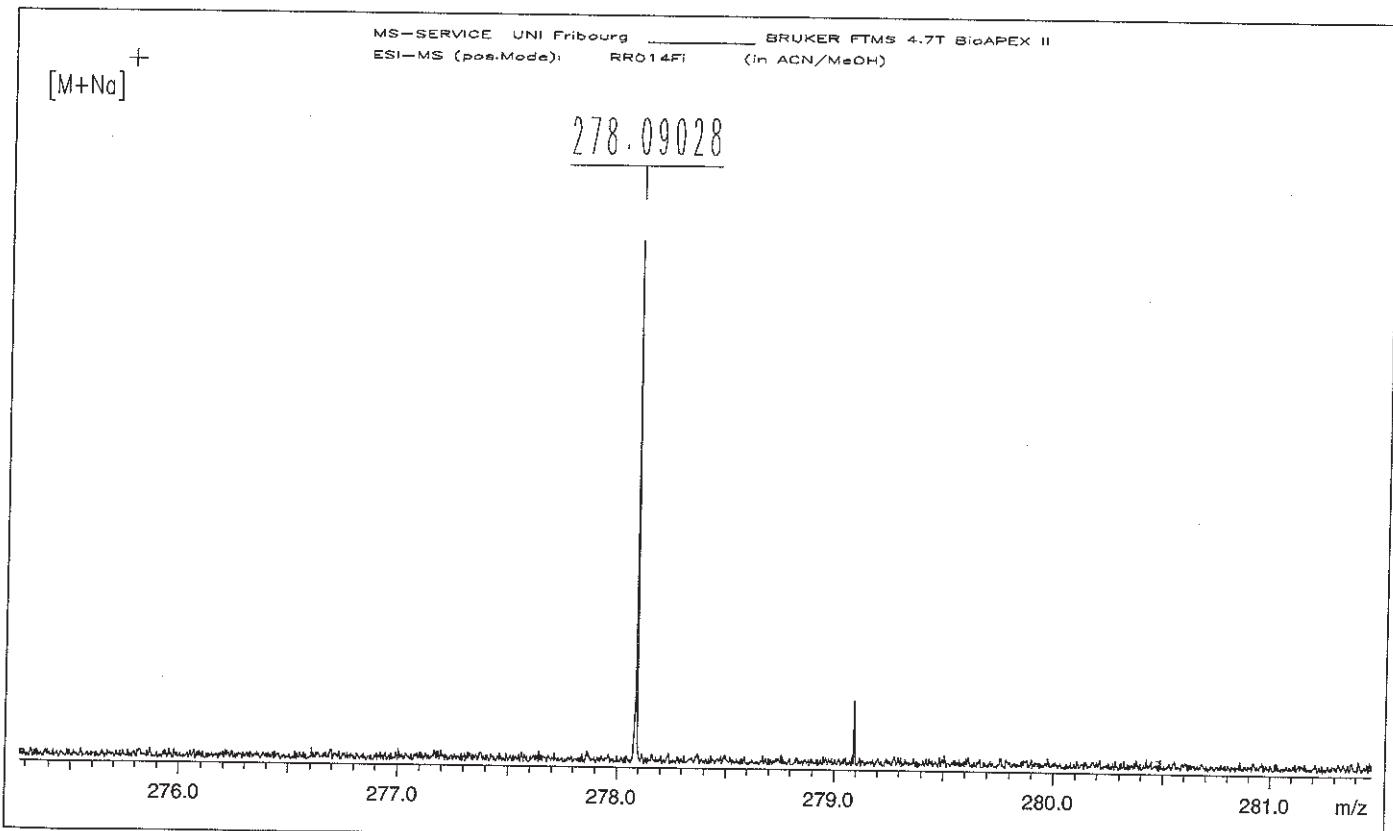
ESI-MS: RR014Fi

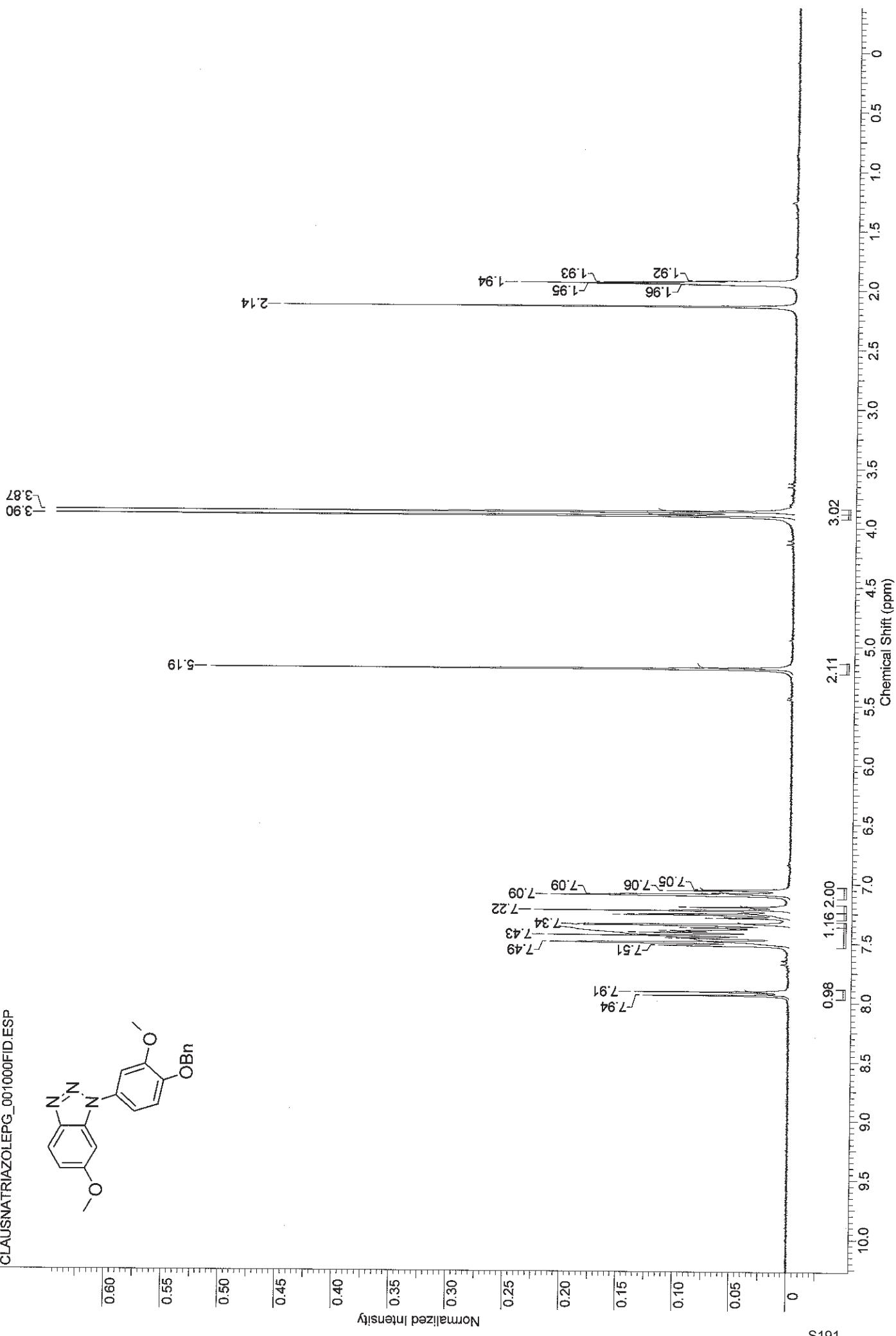
XMASS Mass Analysis for /Data/UNI_FR/REMY7654_ESI/1/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

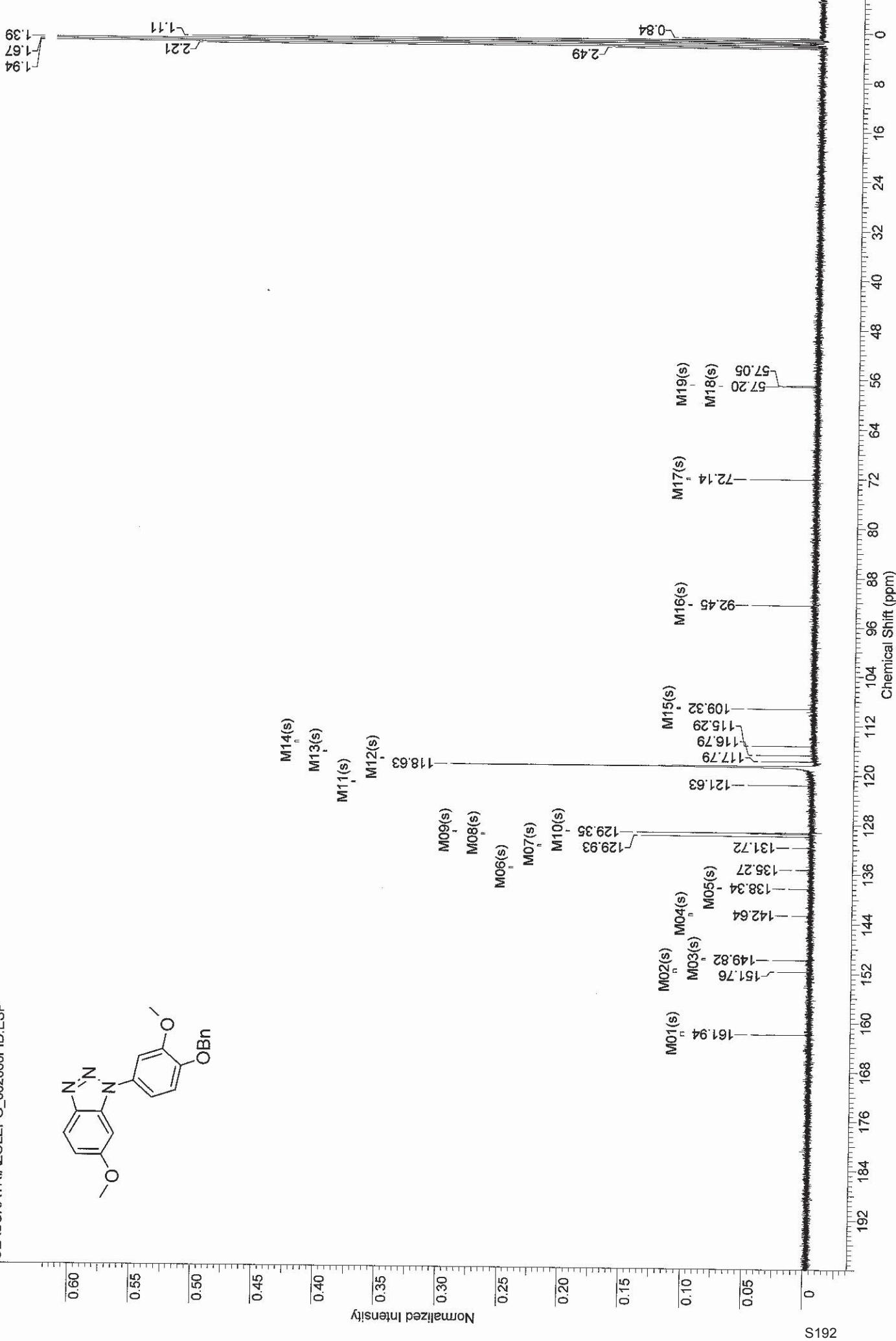
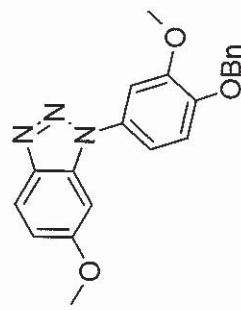
Ion mass = 278.0902750

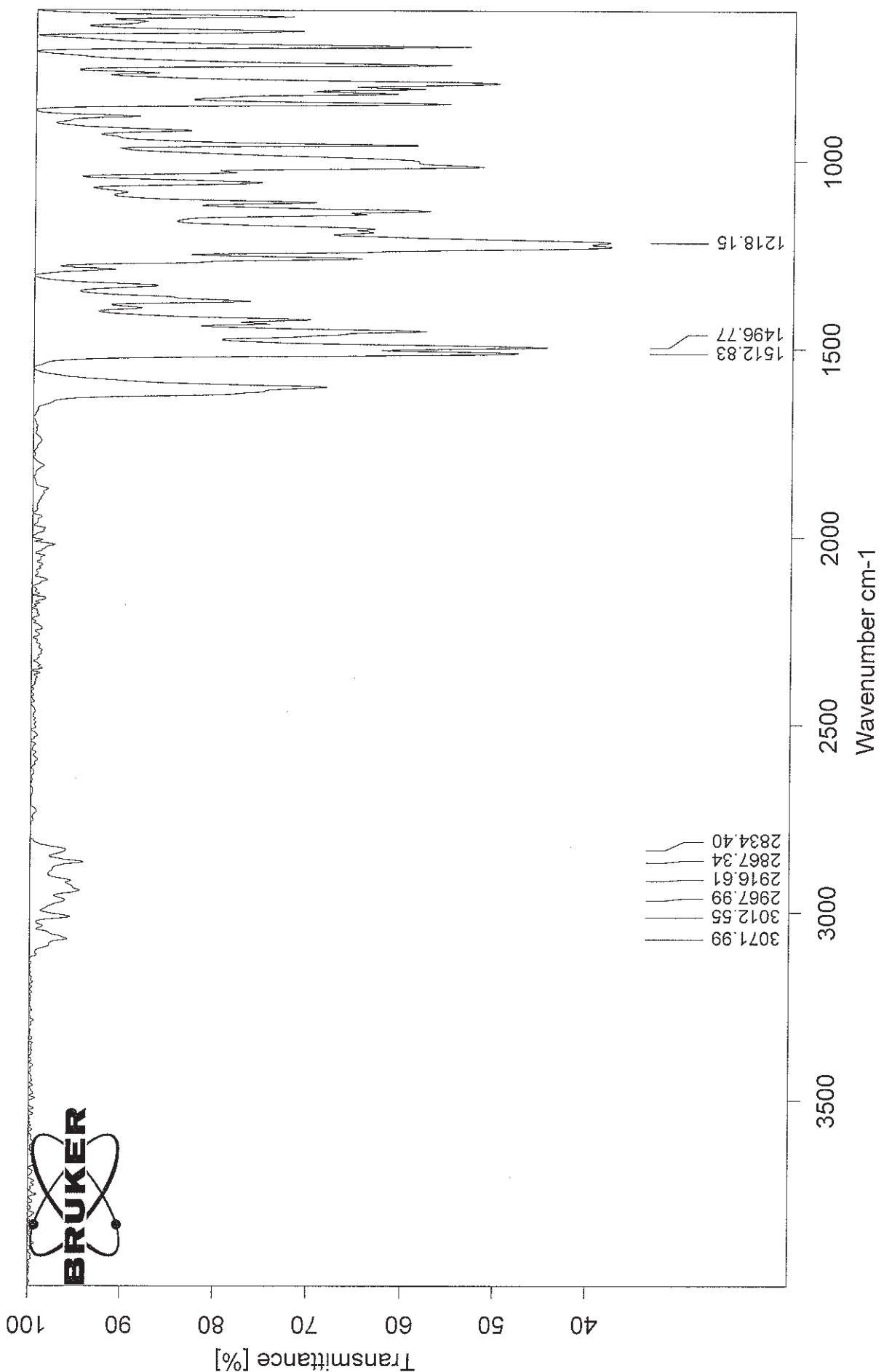
Charge = +1

#	C	H	N	O	Na	mass	DBE	error
*** Mass Analysis for mass 278.0902750								
1	14	13	3	2	1	278.0899978	9.5	2.772e-04
2	13	14	2	5	0	278.0897230	8.0	5.520e-04
3	14	10	6	1	0	278.0910604	13.0	7.854e-04
4	12	11	6	1	1	278.0886551	10.0	1.620e-03
5	11	12	5	4	0	278.0883803	8.5	1.895e-03
6	16	12	3	2	0	278.0924031	12.5	2.128e-03
7	11	15	2	5	1	278.0873177	5.0	2.957e-03
8	10	16	1	8	0	278.0870429	3.5	3.232e-03
9	9	10	8	3	0	278.0870376	9.0	3.237e-03
10	5	15	6	6	1	278.0945283	1.0	4.253e-03









U:\MR\1-(4-(benzyloxy)-3-methoxyphenyl)-6-methoxy-1Hbenzo[d][1,2,3]triazole.0 1-(4-(benzyloxy)-3-methoxyphenyl)-6-methoxy-1Hbenzo[d][1,2,3]triazole[1,2,3] 18.11.2013

ESI-MS: RR042Fi

XMASS Mass Analysis Constraints

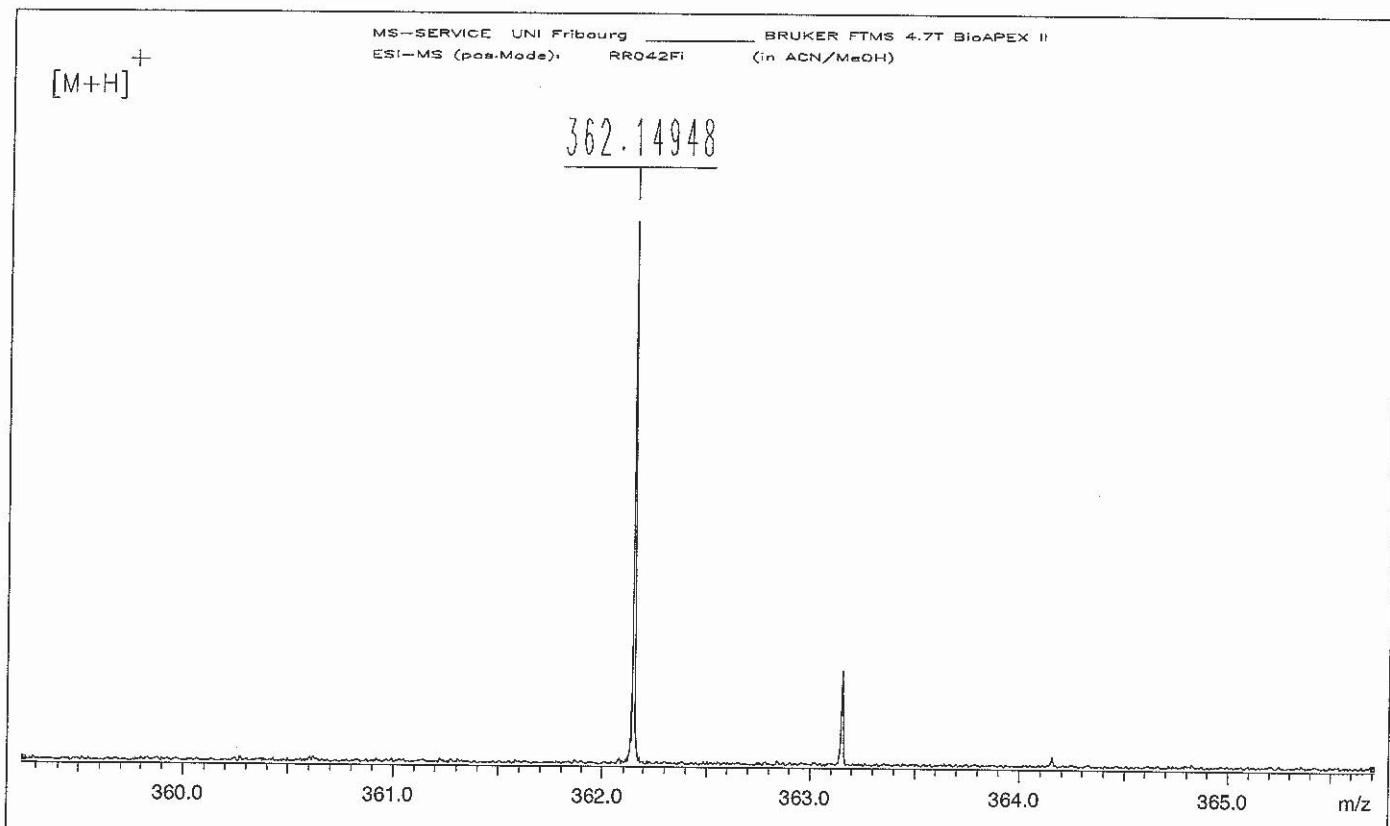
Ion mass = 362.1494770

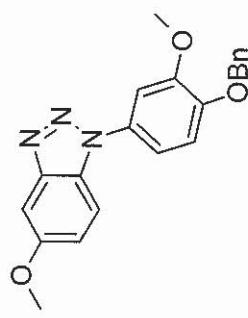
Charge = +1

#	C	H	N	O	mass	DBE	error
---	---	---	---	---	------	-----	-------

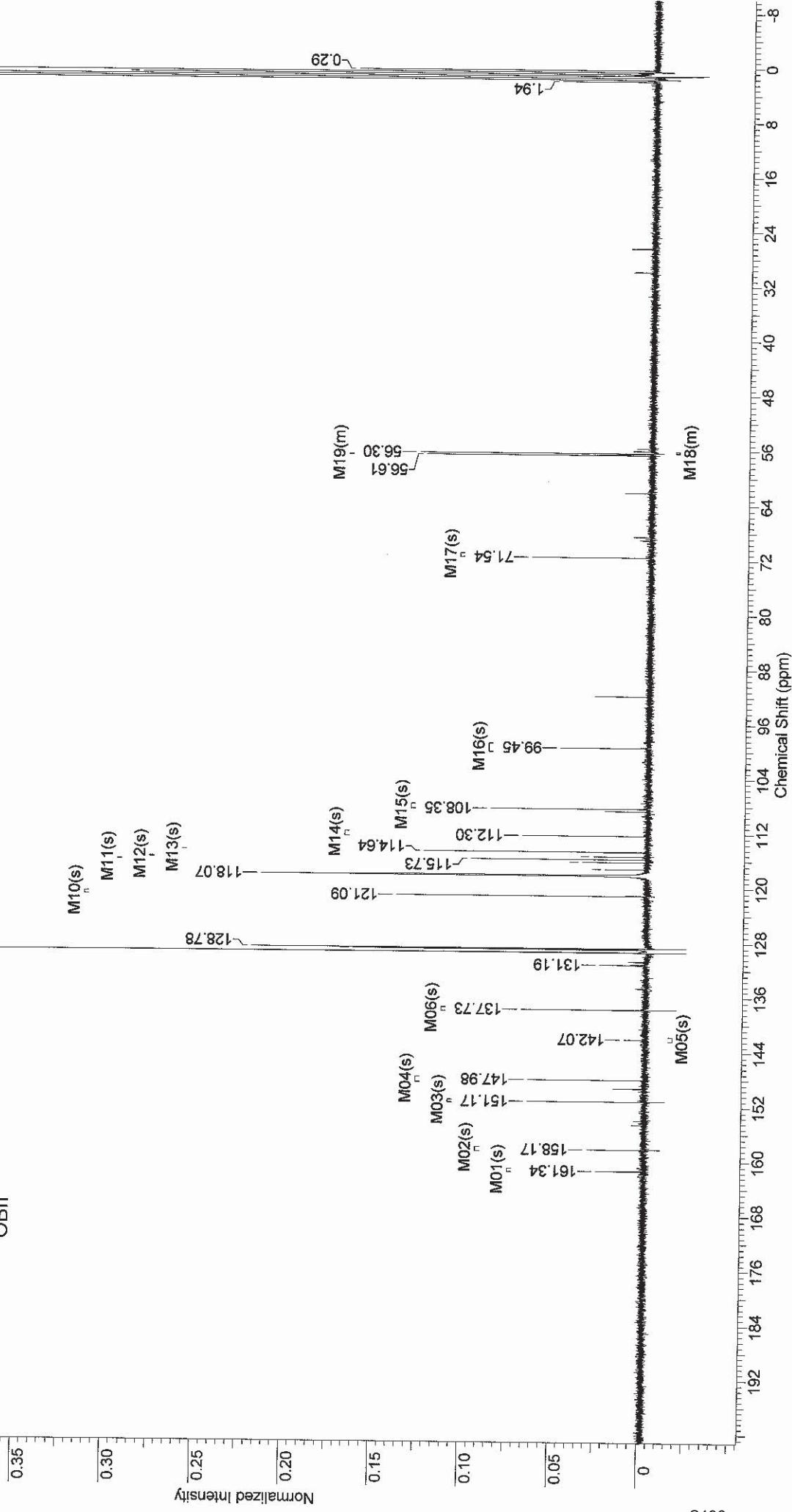
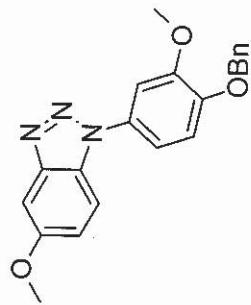
*** Mass Analysis for mass 362.1494770

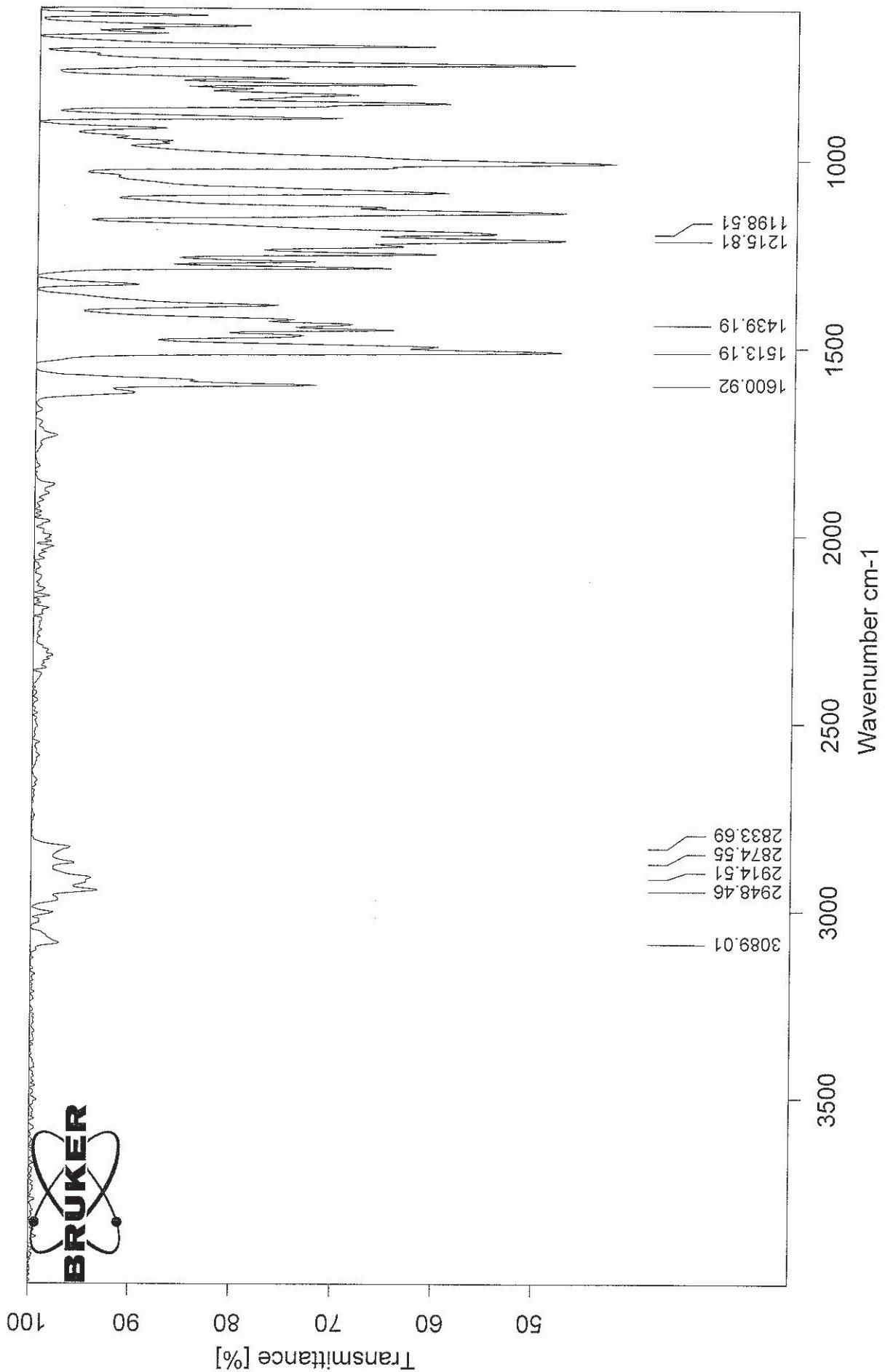
1	21	20	3	3	362.1499179	13.5	4.409e-04
2	19	18	6	2	362.1485753	14.0	9.017e-04
3	7	22	8	9	362.1504258	1.0	9.488e-04
4	18	22	2	6	362.1472379	9.0	2.239e-03
5	17	16	9	1	362.1472326	14.5	2.244e-03
6	9	24	5	10	362.1517684	0.5	2.291e-03
7	16	20	5	5	362.1458952	9.5	3.582e-03
8	10	20	9	6	362.1531058	5.5	3.629e-03
9	15	24	1	9	362.1445578	4.5	4.919e-03
10	14	18	8	4	362.1445525	10.0	4.924e-03





M07(s)
M08(s)
M09(s)





U:\IR\1-(4-(benzyloxy)-3-methoxyphenyl)-5-methoxy-1Hbenzo[d][1,2,3]triazole.0 1-(4-(benzyloxy)-3-methoxyphenyl)-5-methoxy-1Hbenzo[d][1,2,3]triazole.1
18.11.2013

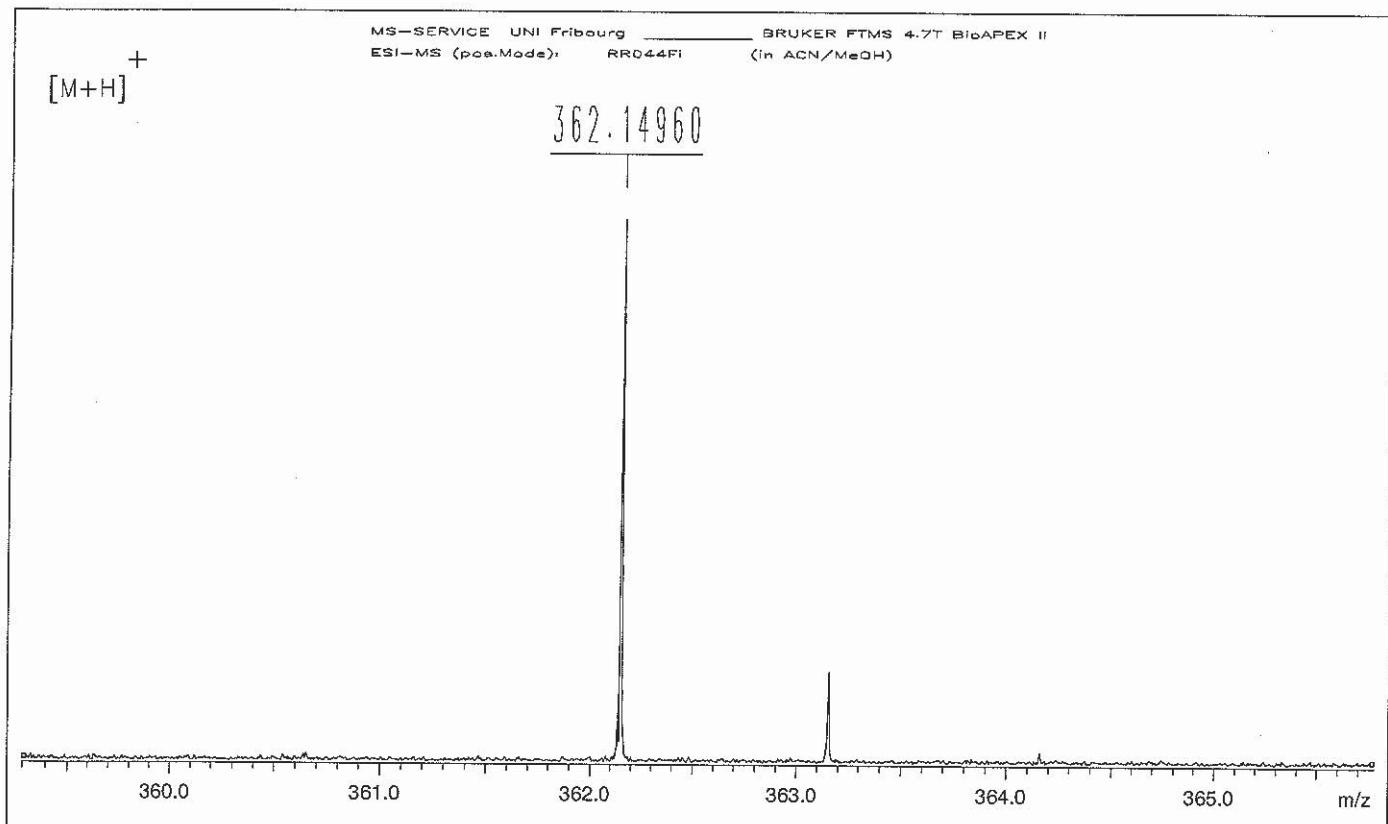
ESI-MS: RR044Fi

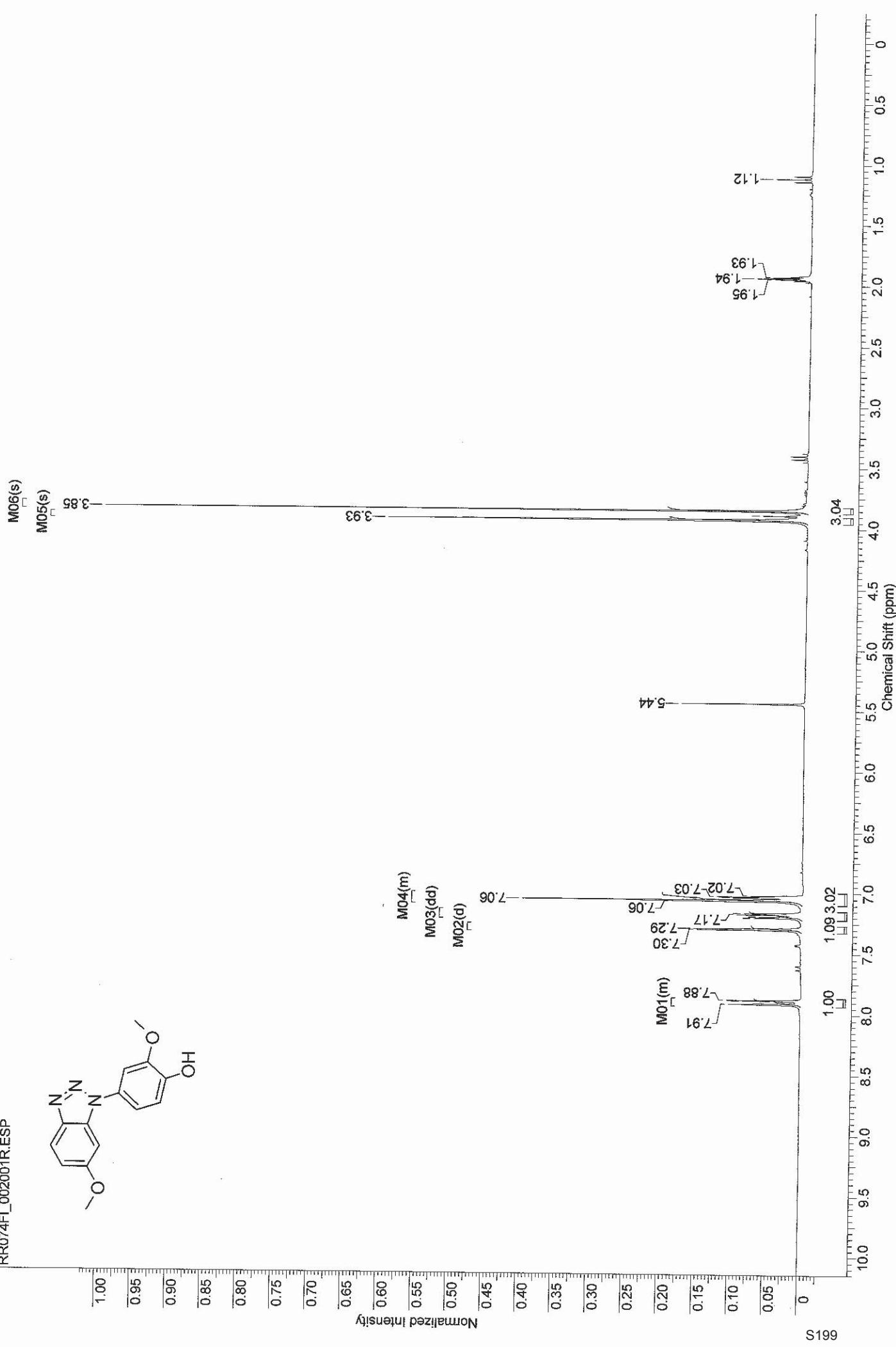
XMASS Mass Analysis for /Data/UNI_FR/REMY7649_ESI/3/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

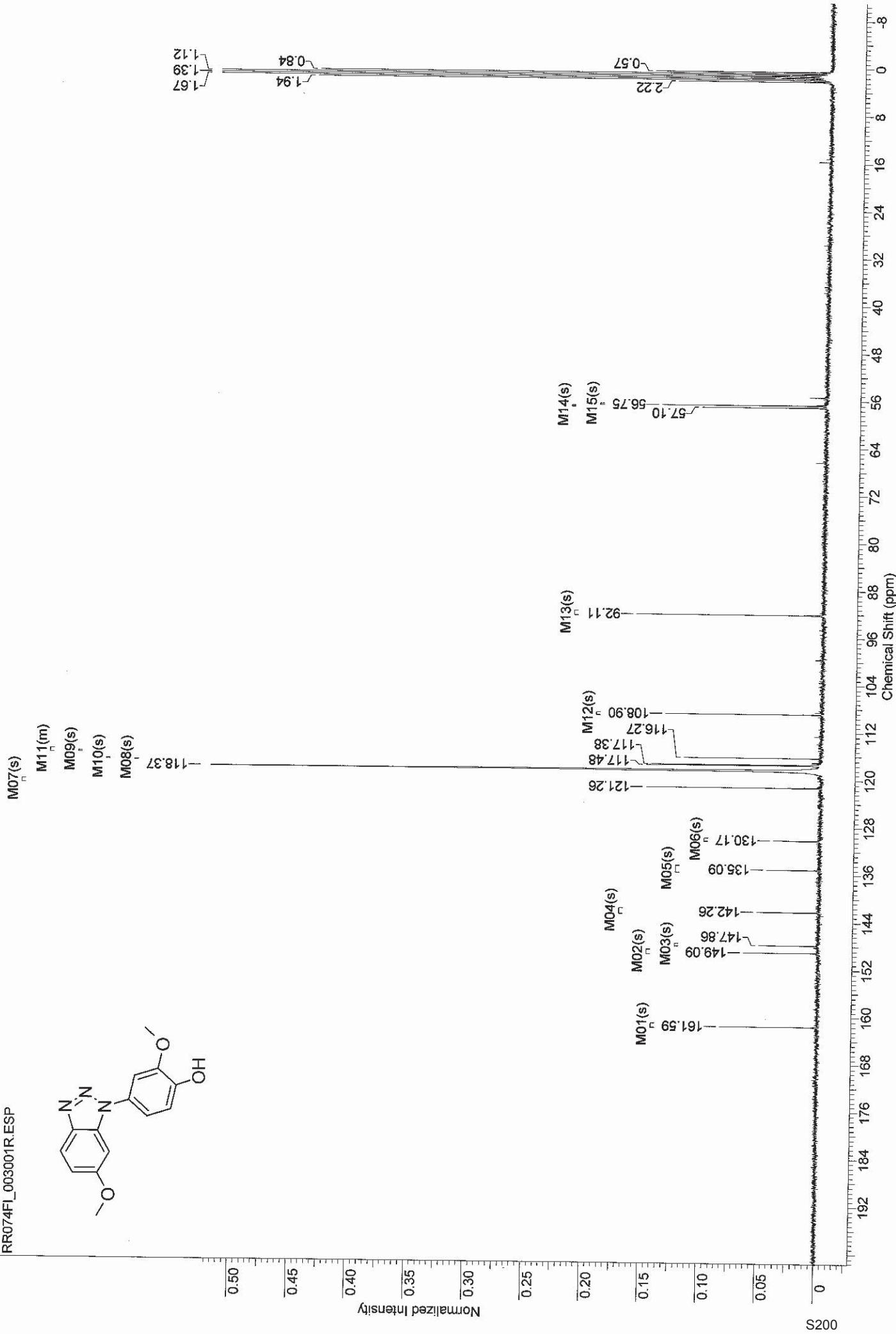
Ion mass = 362.1496000

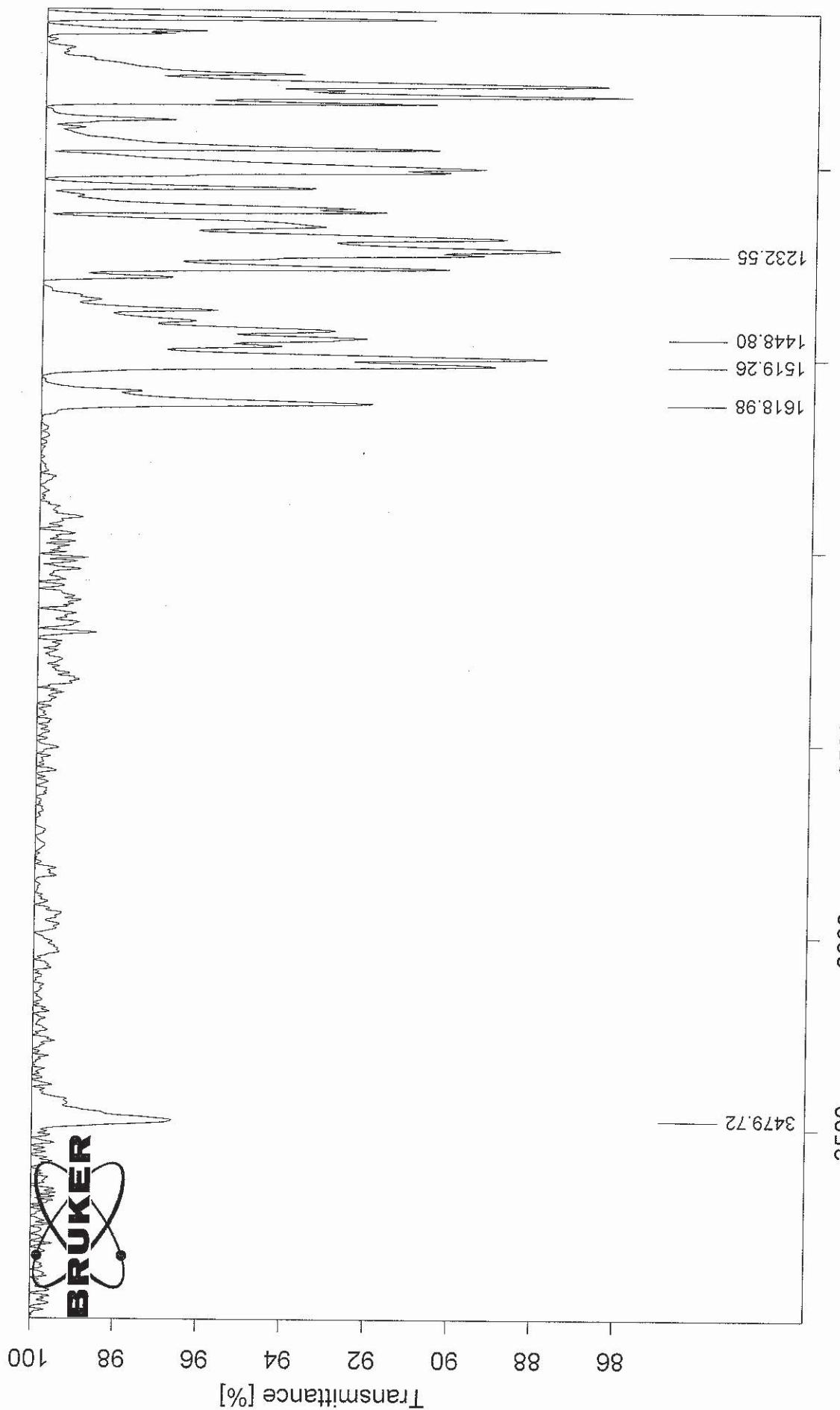
Charge = +1

#	C	H	N	O	mass	DBE	error
*** Mass Analysis for mass 362.1496000							
1	21	20	3	3	362.1499179	13.5	3.179e-04
2	7	22	8	9	362.1504258	1.0	8.258e-04
3	19	18	6	2	362.1485753	14.0	1.025e-03
4	9	24	5	10	362.1517684	0.5	2.168e-03
5	18	22	2	6	362.1472379	9.0	2.362e-03
6	17	16	9	1	362.1472326	14.5	2.367e-03
7	10	20	9	6	362.1531058	5.5	3.506e-03
8	16	20	5	5	362.1458952	9.5	3.705e-03
9	12	22	6	7	362.1544485	5.0	4.849e-03
10	15	24	1	9	362.1445578	4.5	5.042e-03









U:\IR\2-methoxy-4-(6-methoxy-1H-benzo[d][1,2,3]triazol-1-yl)phenol.0

2-methoxy-4-(6-methoxy-1H-benzo[d][1,2,3]triazol-1-yl)phenol

brow
18.11.2013

ESI-MS: RR074Fi

XMASS Mass Analysis for /Data/UNI_FR/REMY7652_ESI/3/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

Ion mass = 272.1028240

Charge = +1

#	C	H	N	O	mass	DBE	error
---	---	---	---	---	------	-----	-------

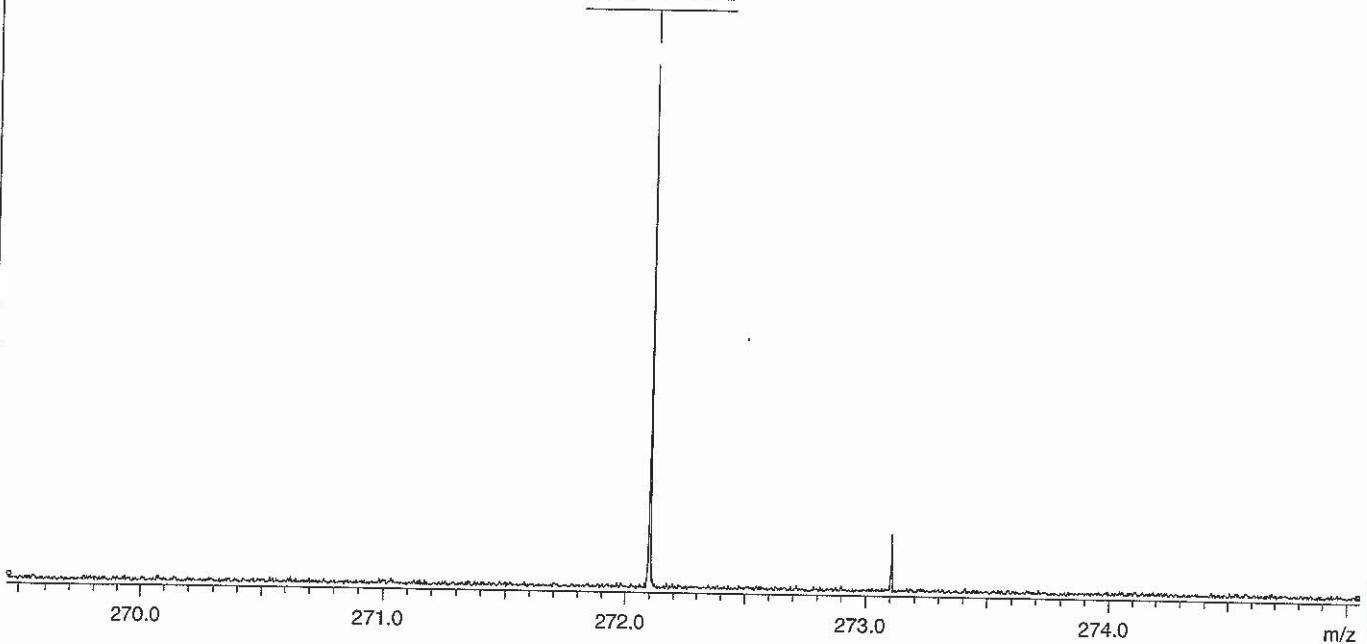
*** Mass Analysis for mass 272.1028240

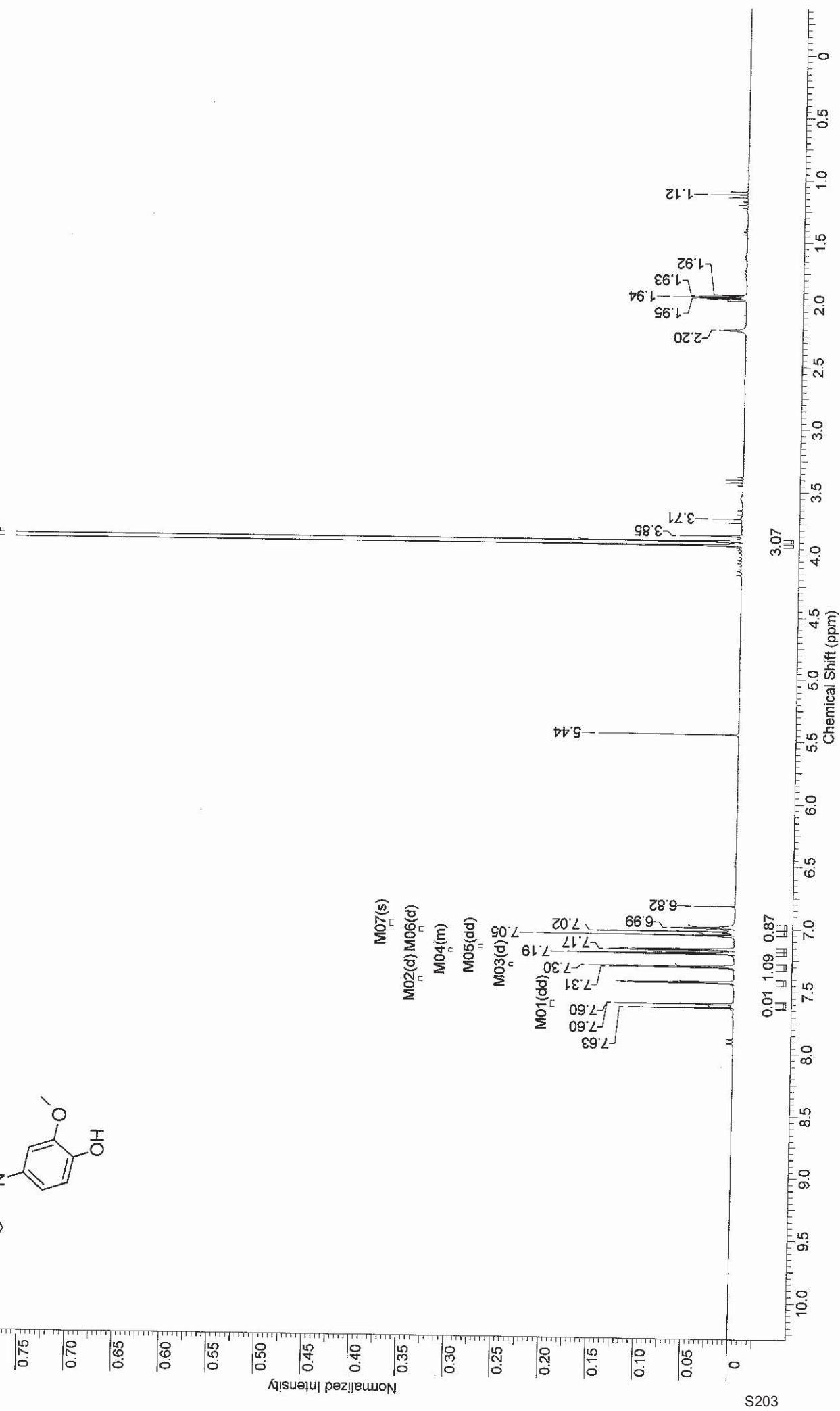
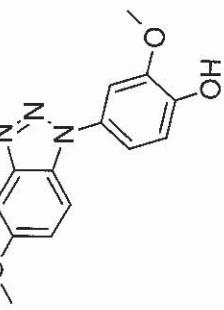
1	14	14	3	3	272.1029678	9.5	1.438e-04
2	12	12	6	2	272.1016251	10.0	1.199e-03
3	11	16	2	6	272.1002877	5.0	2.536e-03
4	10	10	9	1	272.1002824	10.5	2.542e-03
5	9	14	5	5	272.0989450	5.5	3.879e-03
6	19	14	1	1	272.1069905	13.5	4.166e-03
7	5	16	6	7	272.1074983	1.0	4.674e-03
8	8	18	1	9	272.0976076	0.5	5.216e-03
9	7	12	8	4	272.0976023	6.0	5.222e-03
10	6	12	10	3	272.1088357	6.0	6.012e-03

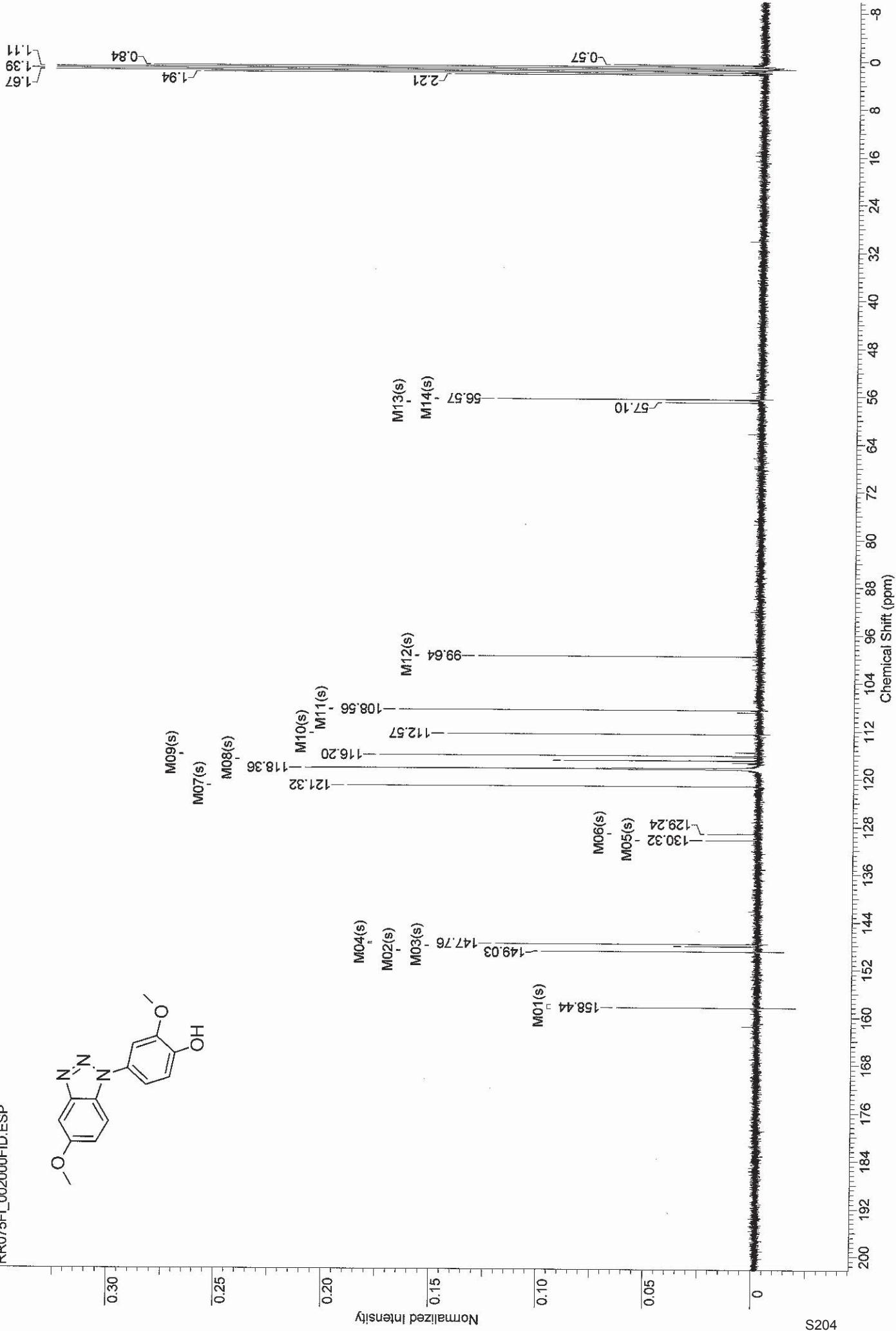
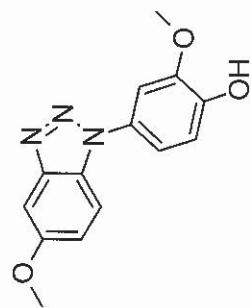
+
[M+H]

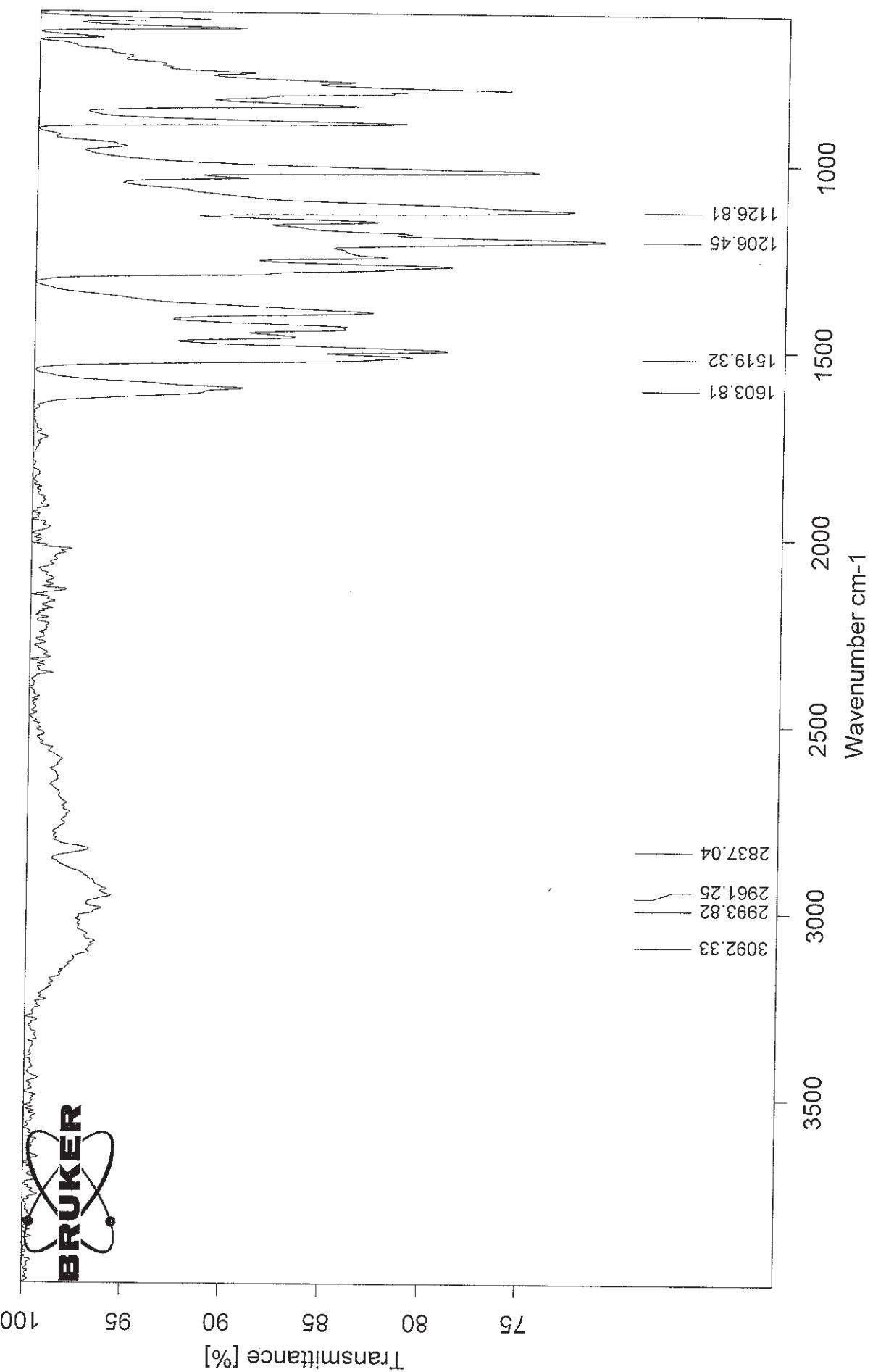
MS-SERVICE UNI Fribourg BRUKER FTMS 4.7T BioAPEX II
ESI-MS (pos.Mode), RR074Fi (in ACN/MeOH)

272.10282



M08(s)
M09(s)





U:\IR\2-methoxy-4-(5-methoxy-1H-benzo[d][1,2,3]triazol-1-yl)phenol.0

2-methoxy-4-(5-methoxy-1H-benzo[d][1,2,3]triazol-1-yl)phenol
brow 18.11.2013



ESI-MS: RR075FI

XMASS Mass Analysis for /Data/UNI_FR/REMY7653_ESI/4/pdata/1/massanal.res:
XMASS Mass Analysis Constraints

Ion mass = 272.1028760

Charge = +1

#	C	H	N	O	mass	DBE	error
---	---	---	---	---	------	-----	-------

*** Mass Analysis for mass 272.1028760

1	14	14	3	3	272.1029678	9.5	9.175e-05
2	12	12	6	2	272.1016251	10.0	1.251e-03
3	11	16	2	6	272.1002877	5.0	2.588e-03
4	10	10	9	1	272.1002824	10.5	2.594e-03
5	9	14	5	5	272.0989450	5.5	3.931e-03
6	19	14	1	1	272.1069905	13.5	4.114e-03
7	5	16	6	7	272.1074983	1.0	4.622e-03
8	8	18	1	9	272.0976076	0.5	5.268e-03
9	7	12	8	4	272.0976023	6.0	5.274e-03
10	6	12	10	3	272.1088357	6.0	5.960e-03

