

Supporting information

Macrocyclic peptidomimetics with antimicrobial activity: synthesis, bioassay, and molecular modeling studies

Mohamed A. Ibrahim,^{*a,b,c} Siva S. Panda,^{*a,d} Alexander A. Oliferenko,^a Polina V. Oliferenko,^a Adel S. Girgis,^e Mohamed Elagawany,^{a,f} F. Zehra Küçükbay,^{a,g} Chandramukhi S. Panda,^a Girinath G. Pillai,^{a,h} Ahmed Samir,ⁱ Kaido Tamm,^h C. Dennis Hall^a and Alan R. Katritzky^{a,†}

^a*Department of Chemistry, University of Florida, Gainesville, FL 32611-7200, USA. E-mail: drmmonem@yahoo.com, sspanda12@gmail.com; fax: +1-352-392-9199; Tel: +1-352-870-9288.*

^b*Department of Organic Chemistry, College of pharmacy, Misr University for Science and Technology, Al-Motamayez District, 6th of the October, P.O.Box: 77, Egypt*

^c*Department of Pharmaceutical Chemistry, College of Pharmacy, Almaarefa college, Riyadh, Kingdom of Saudia Arabia.*

^d*Department of Chemistry & Physics, Georgia Regents University, Augusta, GA 30912, USA*

^e*Pesticide Chemistry Department, National Research Centre, Dokki, Cairo 12622, Egypt.*

^f*Department of Pharmaceutical Chemistry, faculty of pharmacy, Damanhour University, Damanhour, Egypt*

^g*Department of Basic Pharmaceutical Sciences, İnönü University, 44280 Malatya, Turkey*

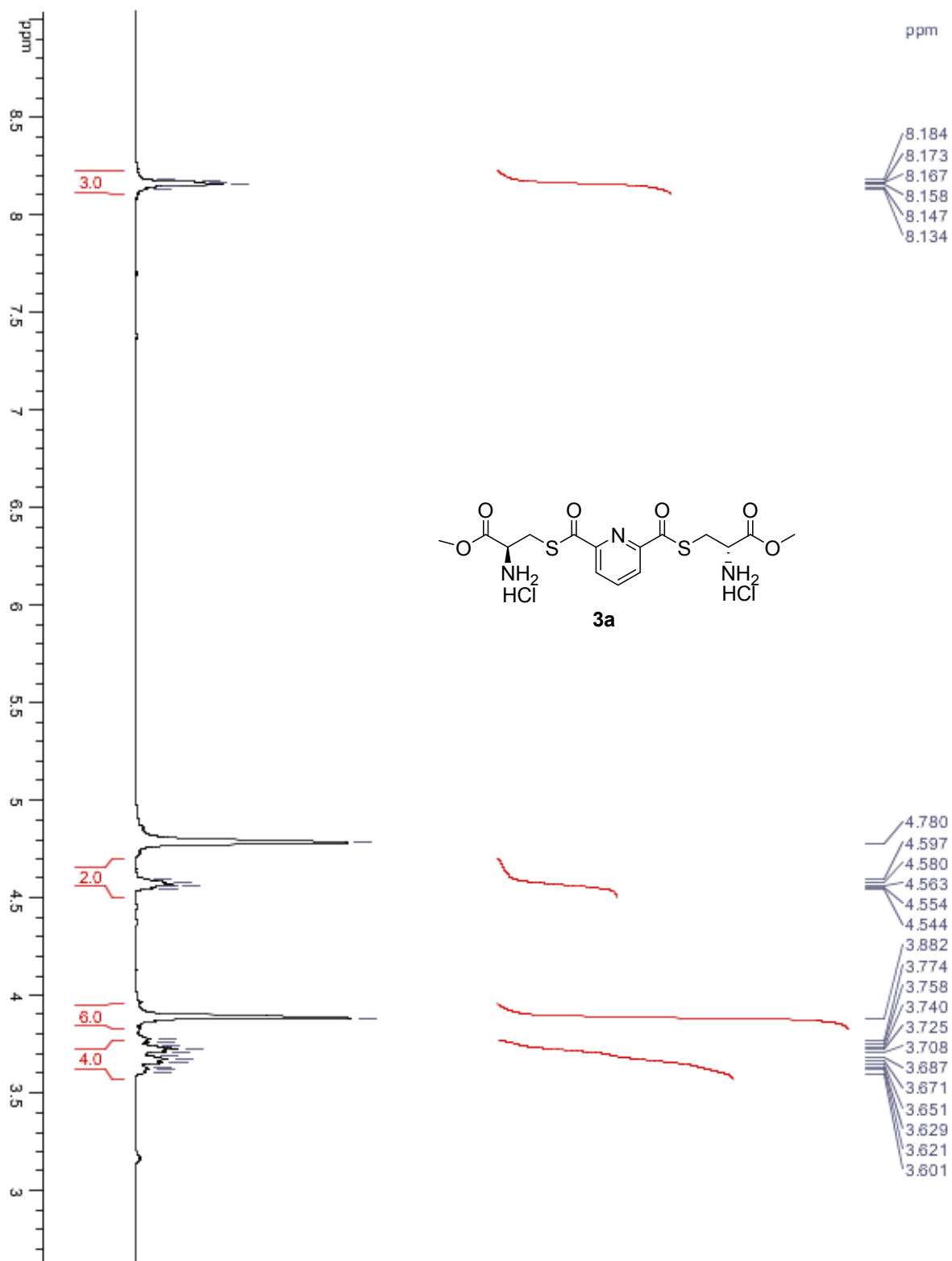
^h*Department of Chemistry, University of Tartu, 50411 Tartu, Estonia*

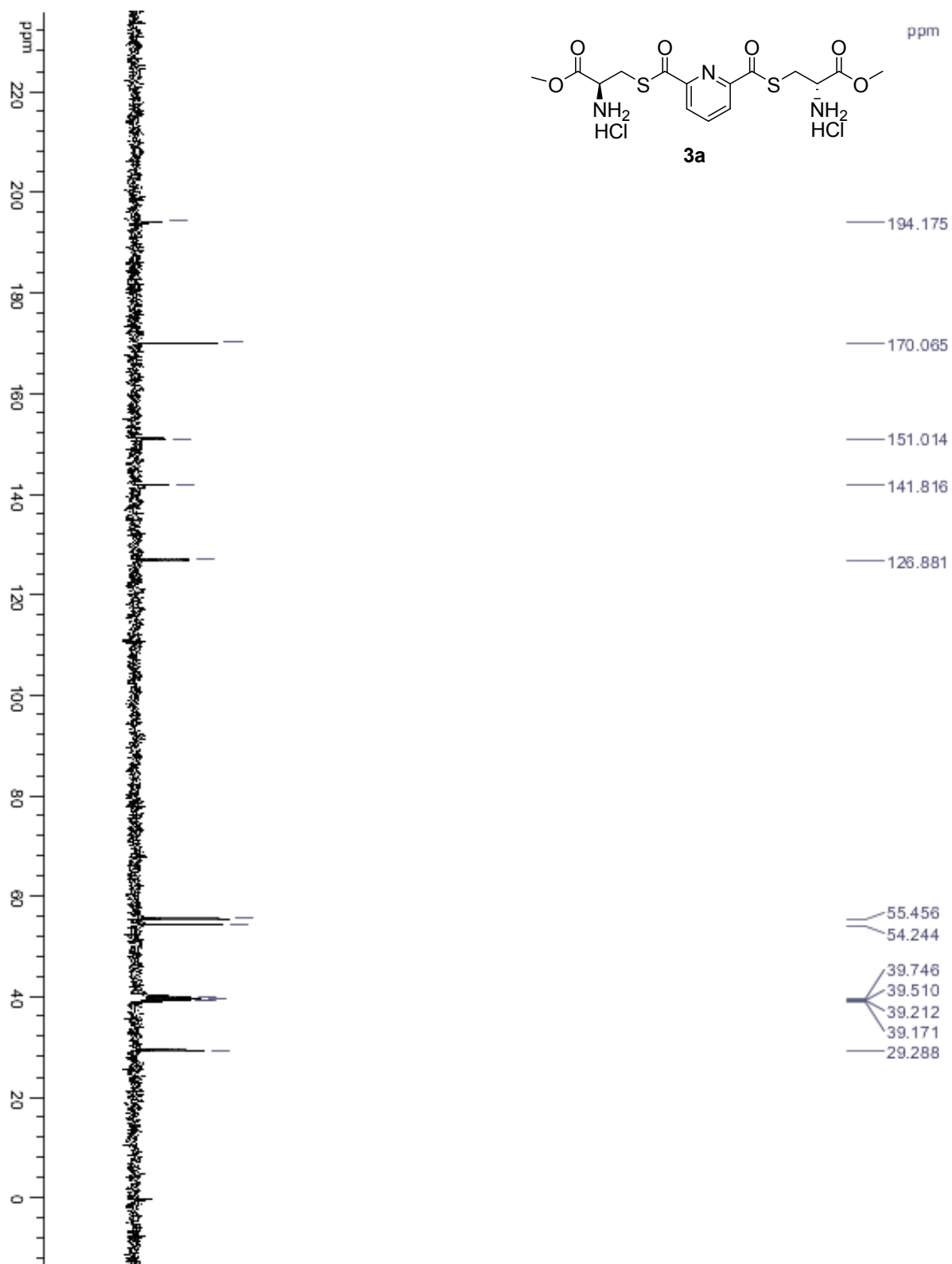
ⁱ*Department of Microbiology, Faculty of Veterinary Medicine, Cairo University, Cairo, Egypt*

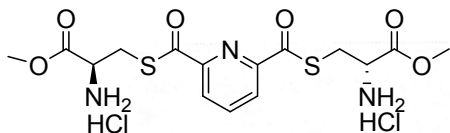
[†]*Deceased on February 10, 2014*

Table of contents

1. ^1H NMR, ^{13}C NMR, CHN/HRMS of all the compounds
2. **Table S1.** Molecular descriptor values of the BMLR-QSAR model for the antifungal active macrocyclic peptidomimetics.
3. **Figure S1.** (A) Constraint distances, and (B) constraint angle of the generated 3D-pharmacophore for the antifungal bio-active compounds **4a-4l**, and **5-7** against *C. albicans* which contains two hydrogen bonding acceptor (HBA-1, HBA-2, green), and one hydrogen bonding donor (HBD, purple).
4. **Figure S2.** 3D-pharmacophore mapped on the antifungal macrocyclic peptidomimetics **4a-4l**, and **5-7**.
5. **Figure S3.** Dose-response curves of the macrocyclic peptidomimetics **4a-l**, and **5-7** against HepG2 (liver) human tumor cell line.
6. **Figure S4.** Dose-response curves of the macrocyclic peptidomimetics **4a-l**, and **5-7** against HeLa (cervical) human tumor cell line.
7. **Figure S5.** Dose-response curves of the macrocyclic peptidomimetics **4a-l**, and **5-7** against HCT116 (colon) human tumor cell line.
8. **Figure S6.** Dose-response curves of the macrocyclic peptidomimetics **4a-l**, and **5-7** against PC3 (prostate) human tumor cell line.
9. **Figure S7.** Dose-response curves of the macrocyclic peptidomimetics **4a-l**, and **5-7** against MCF7 (breast) human tumor cell line.



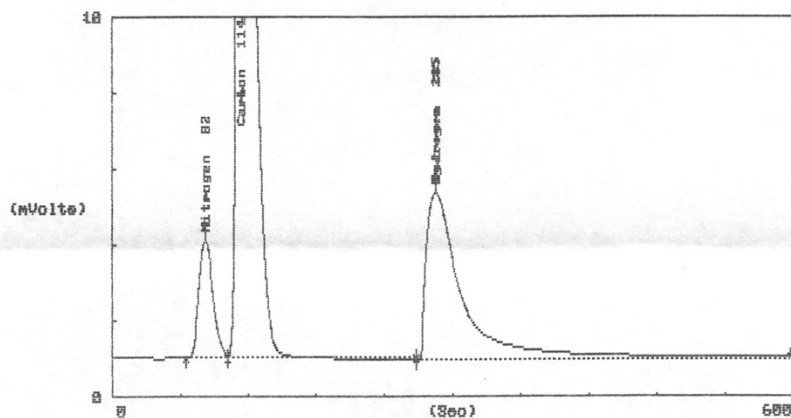




3a

EAGER 200 Stripchart

Sample Ident. : 20 Maaa6esterHCl Filename : 266420
 Analysed : 01-11-12 08:53:19 Printed : 01-11-2012 09:03:20



EAGER 200 Peak Integration Report

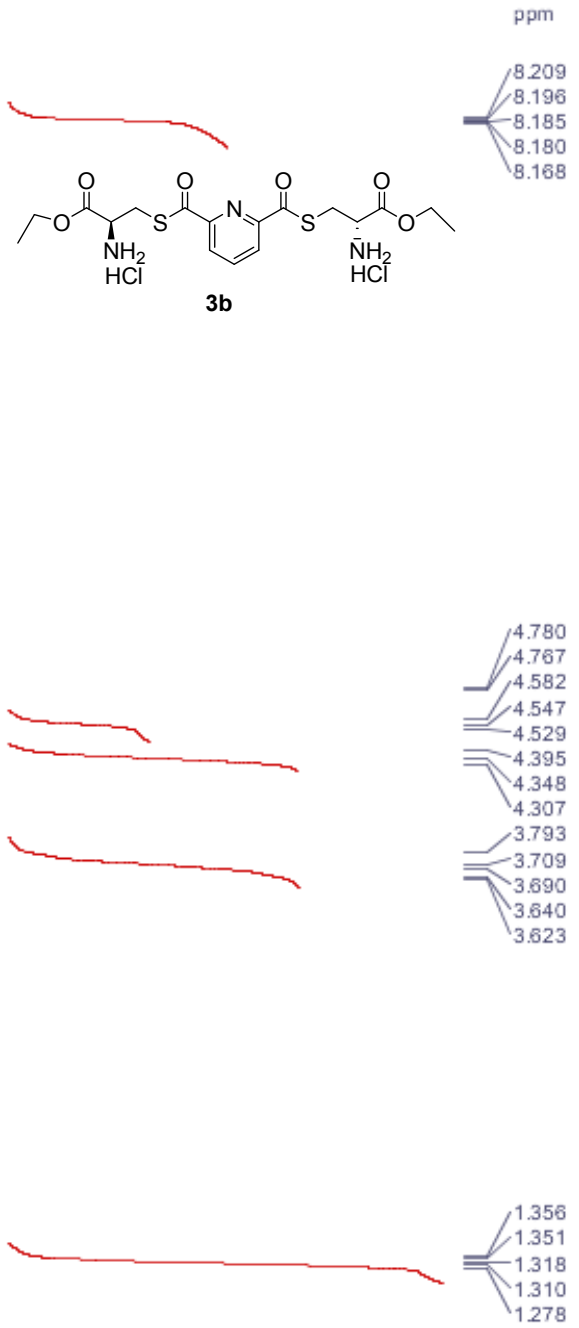
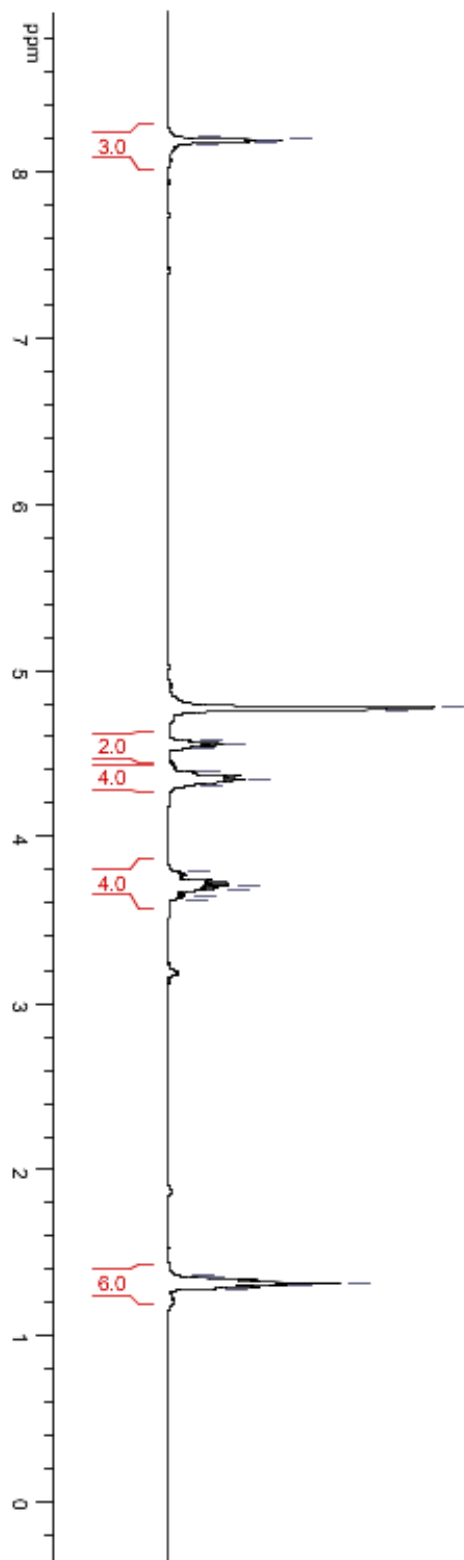
Instrument name : Instrument #1 Bline drift (fV):-4.7
 Company Name : U of Florida Operator Ident. : KOU
 Analysed : 01-11-12 08:53:19 Printed : 01-11-2012 09:03:21
 Sample Ident. : 20 Maaa6esterHCl Filename : 266420
 Sample Weight : 2.168 Calc.method: using 'K. Factors'

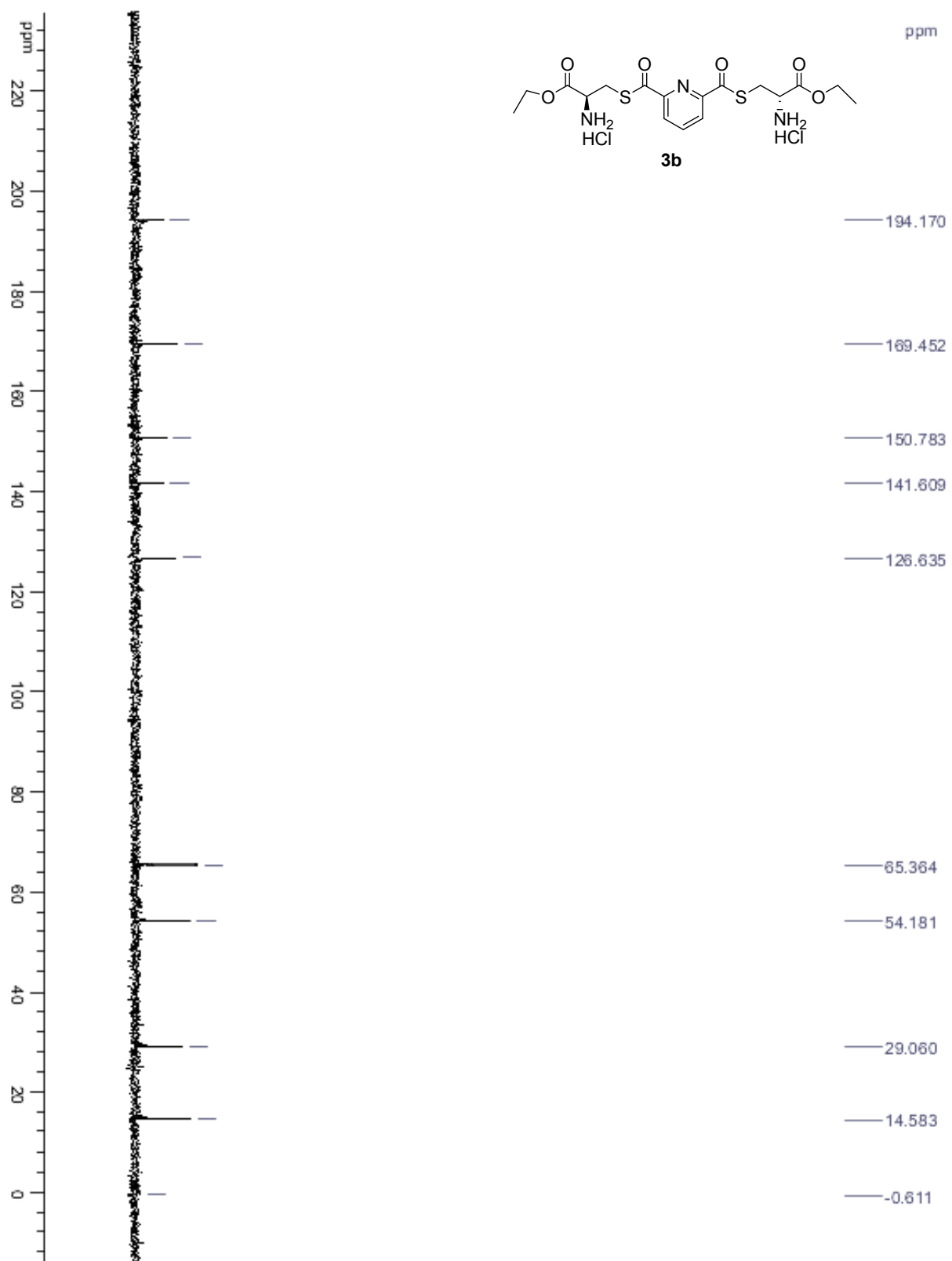
No. (#)	Type (#)	Start (Sec)	End (Sec)	Ret Time (Sec)	Height (fV)	Area (fV*Sec)	Area % (%)	Name
1	FU	65	102	82	3092.2	42236	6.08	Nitrogen
2	FU	102	268	114	33619.0	492721	70.94	Carbon
3	RS	268	598	285	4358.5	159626	22.98	Hydrogen
						694583	100.00	

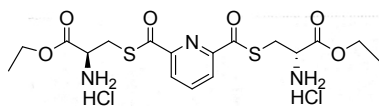
EAGER 200 Unk Report

Instrument name : Instrument #1 Bline drift (fV):-4.7
 Company Name : U of Florida Operator Ident. : KOU
 Analysed : 01-11-12 08:53:19 Printed : 01-11-2012 09:03:21
 Sample Ident. : 20 Maaa6esterHCl Filename : 266420
 Sample Weight : 2.168 Calc.method: using 'K. Factors'

Pk. (#)	Ret Time (Sec)	Area (fV*Sec)	Element % (%)	Area Ratio	Name
1	82	42236	8.828	.116658E+02	Nitrogen
2	114	492721	37.745	.100000E+01	Carbon
3	285	159626	4.569	.308673E+01	Hydrogen



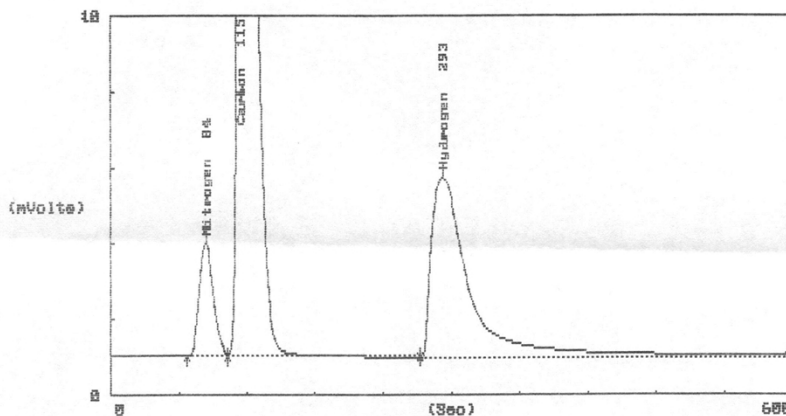




3b

EAGER 200 Stripchart

Sample Ident. : 20 maaa6ethyl Filename : 267120
 Analysed : 02-14-12 08:47:21 Printed : 02-14-2012 08:57:24



EAGER 200 Peak Integration Report

Instrument name : Instrument #1 Bline drift (fV): .9
 Company Name : U of Florida Operator Ident. : KOU
 Analysed : 02-14-12 08:47:21 Printed : 02-14-2012 08:57:24
 Sample Ident. : 20 maaa6ethyl Filename : 267120
 Sample Weight : 2.172 Calc.method: using 'K. Factors'

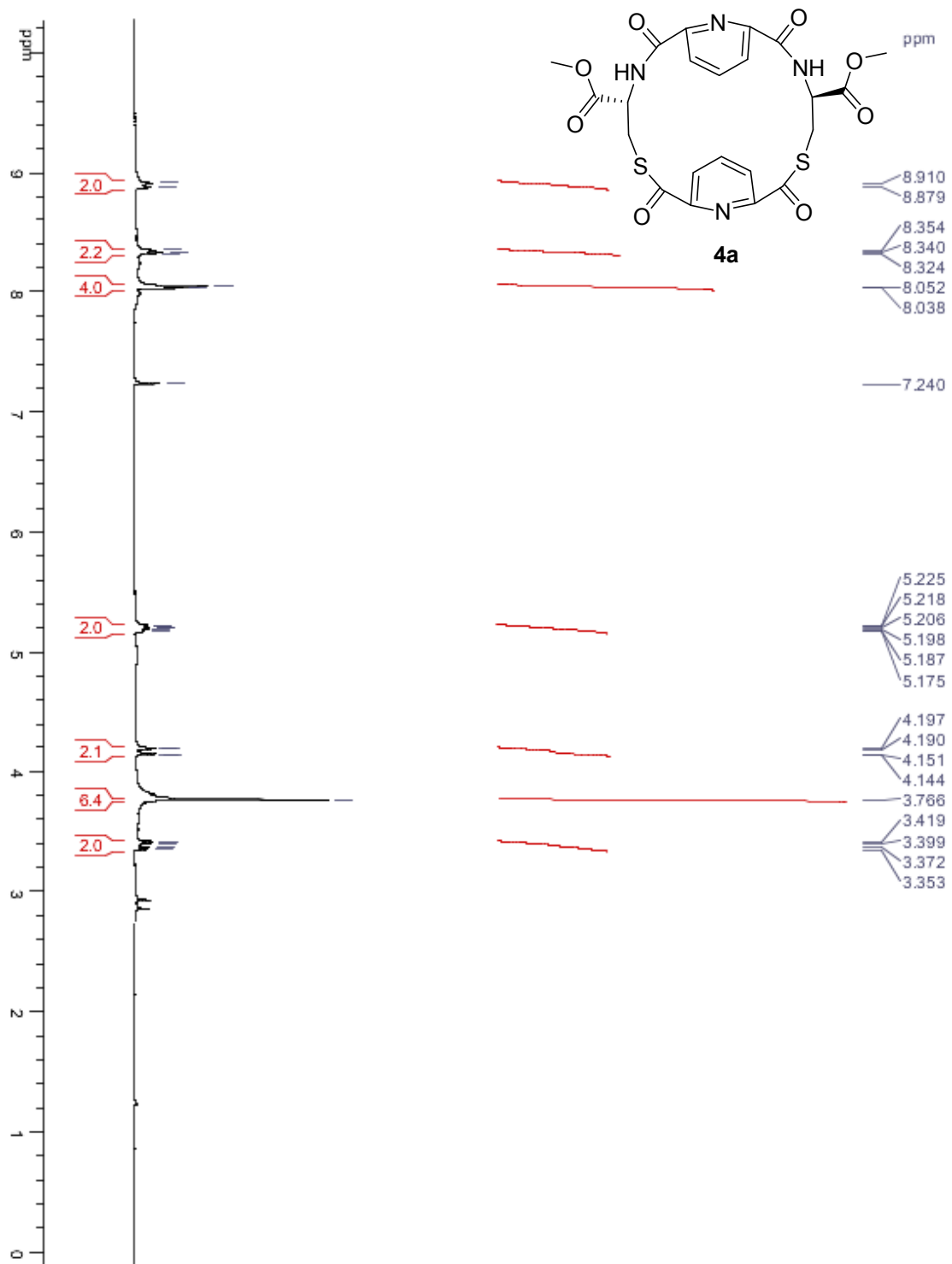
No. (#)	Type (#)	Start (Sec)	End (Sec)	Ret Time (Sec)	Height (fV)	Area (fV*Sec)	Area % (%)	Name
1	FU	67	103	84	2962.1	40633	5.34	Nitrogen
2	FU	103	272	115	36689.9	538909	70.78	Carbon
3	RS	272	597	293	4711.7	181833	23.88	Hydrogen

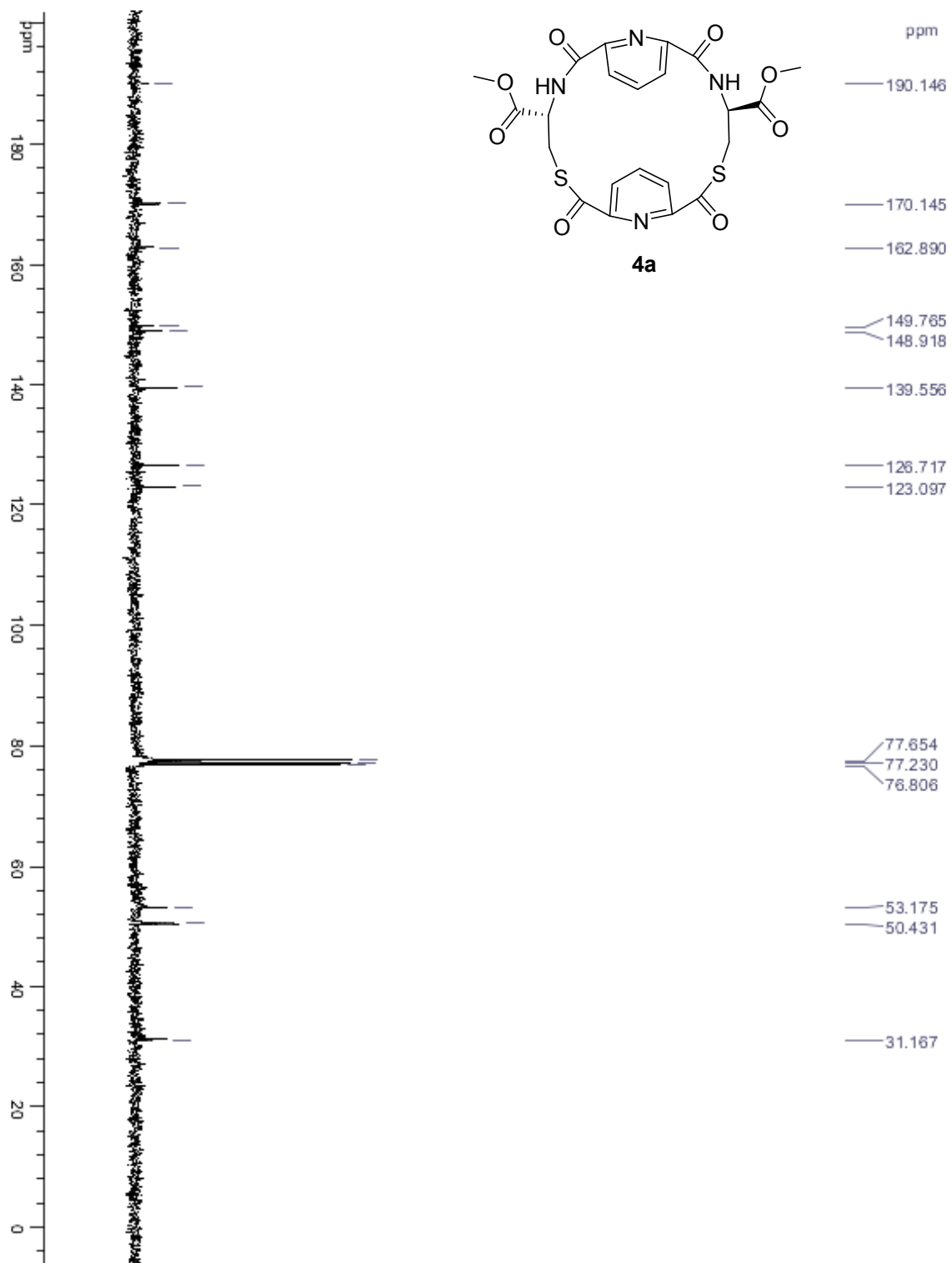
761375 100.00

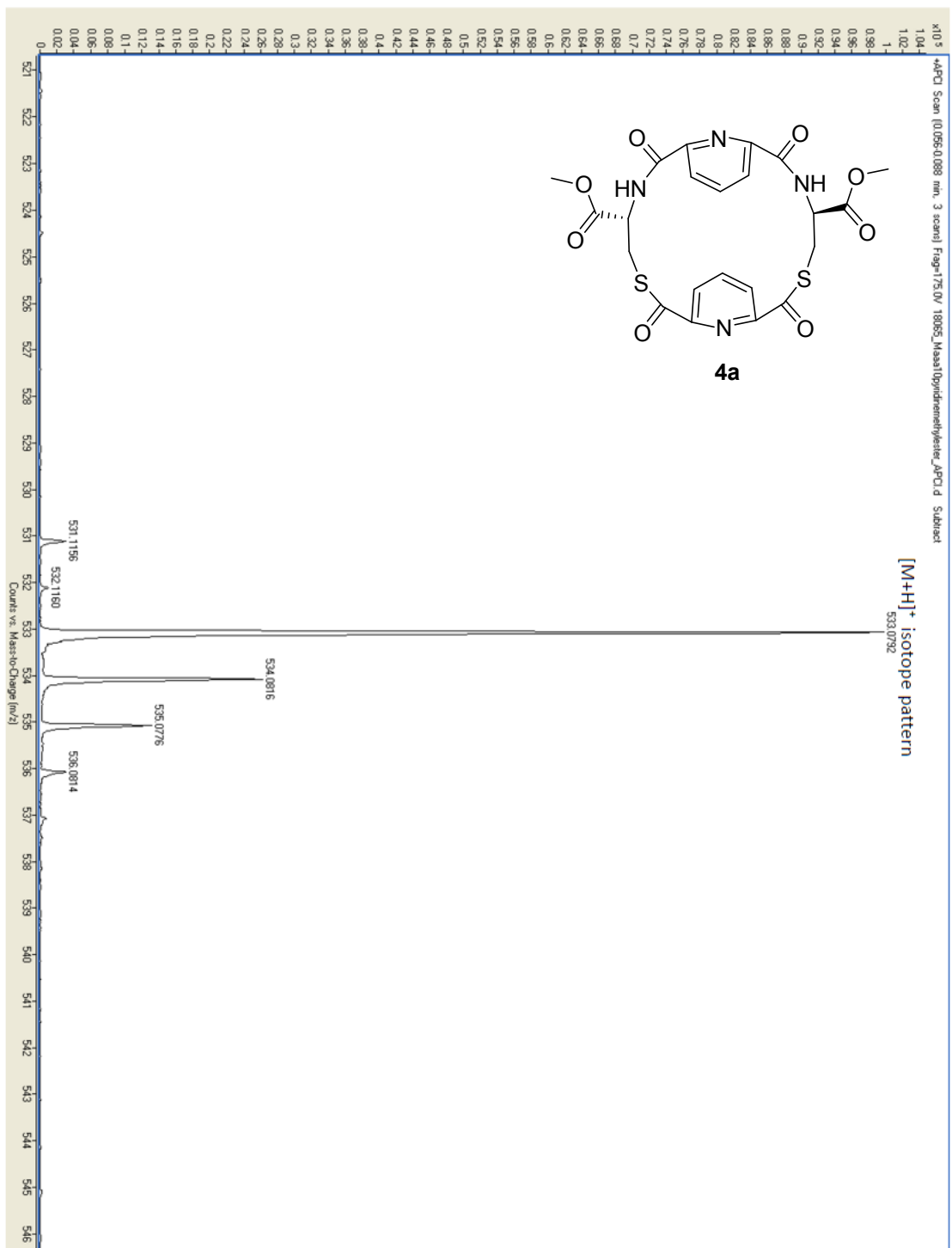
EAGER 200 Unk Report

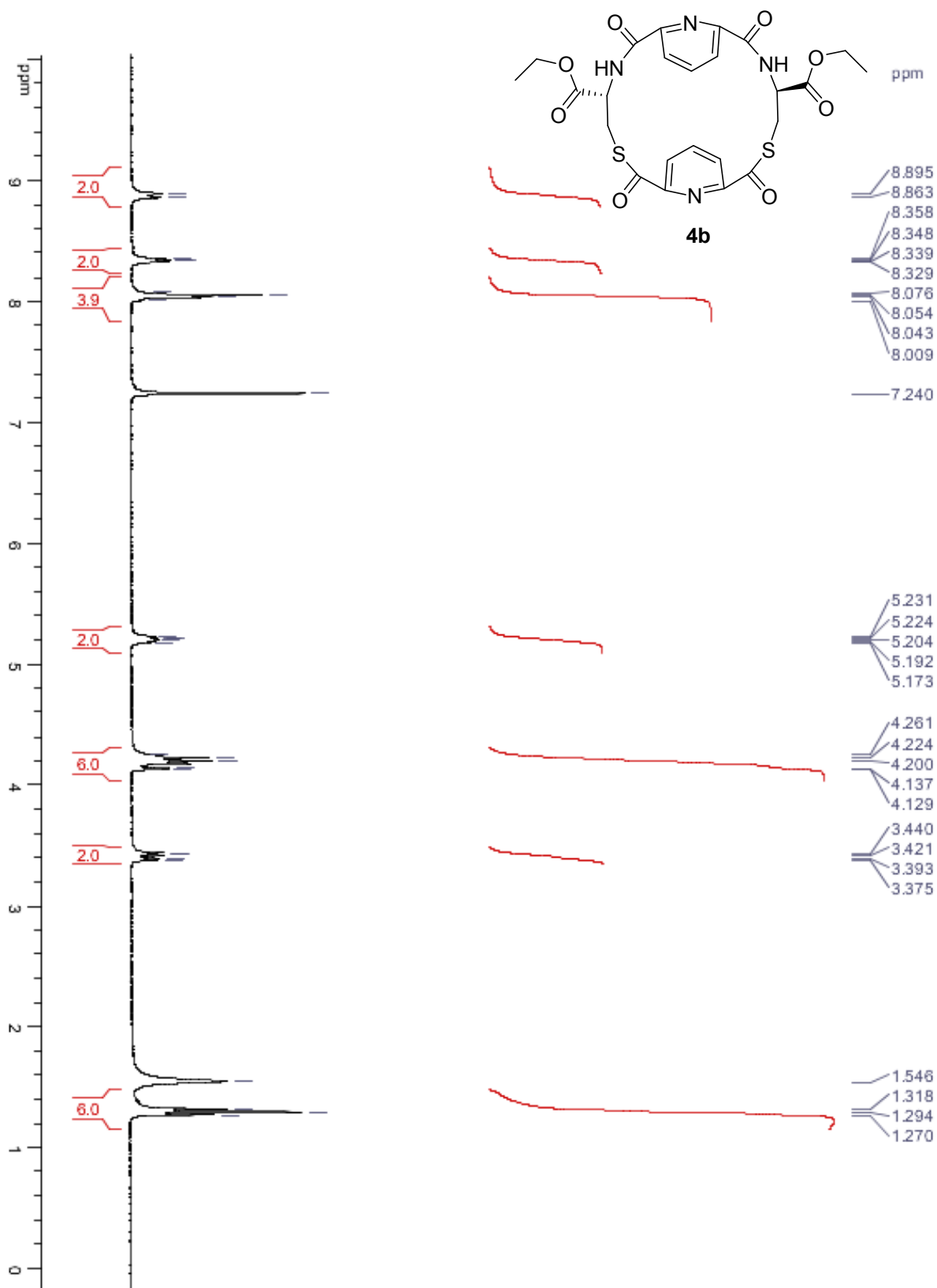
Instrument name : Instrument #1 Bline drift (fV): .9
 Company Name : U of Florida Operator Ident. : KOU
 Analysed : 02-14-12 08:47:21 Printed : 02-14-2012 08:57:24
 Sample Ident. : 20 maaa6ethyl Filename : 267120
 Sample Weight : 2.172 Calc.method: using 'K. Factors'

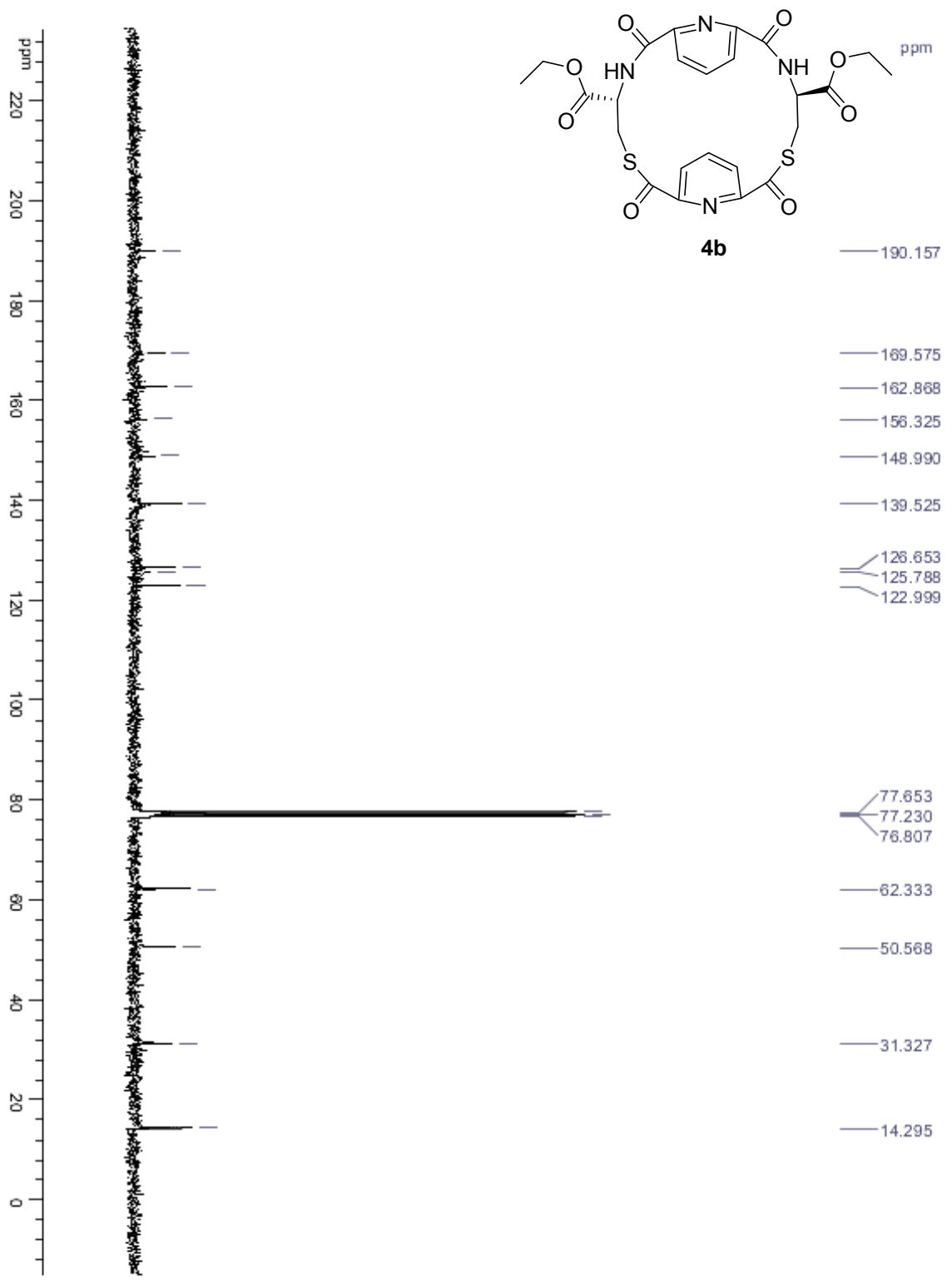
Pk. (#)	Ret Time (Sec)	Area (fV*Sec)	Element % (%)	Area Ratio	Name
1	84	40633	8.454	.132628E+02	Nitrogen
2	115	538909	40.999	.100000E+01	Carbon
3	293	181833	5.372	.296375E+01	Hydrogen

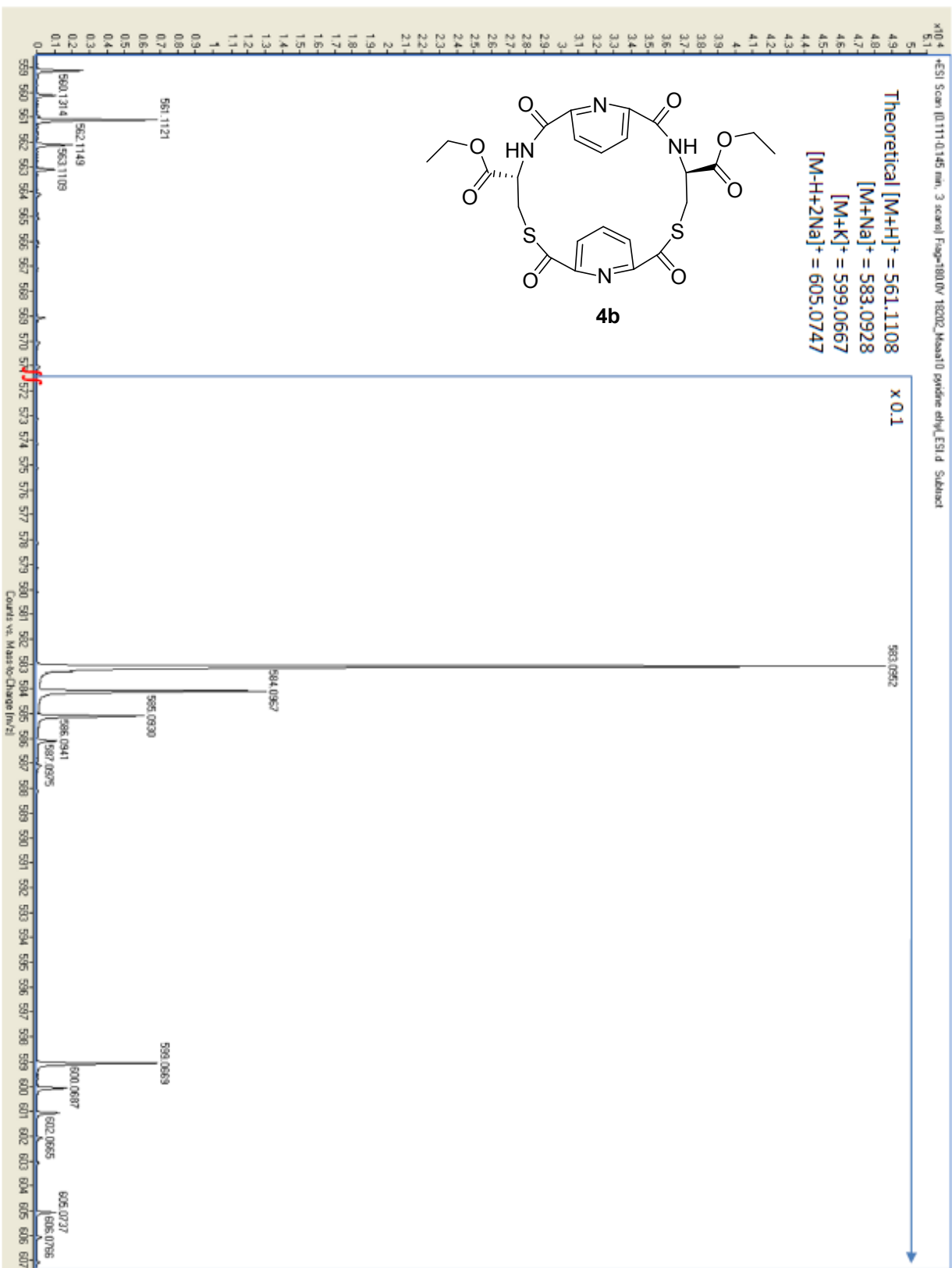


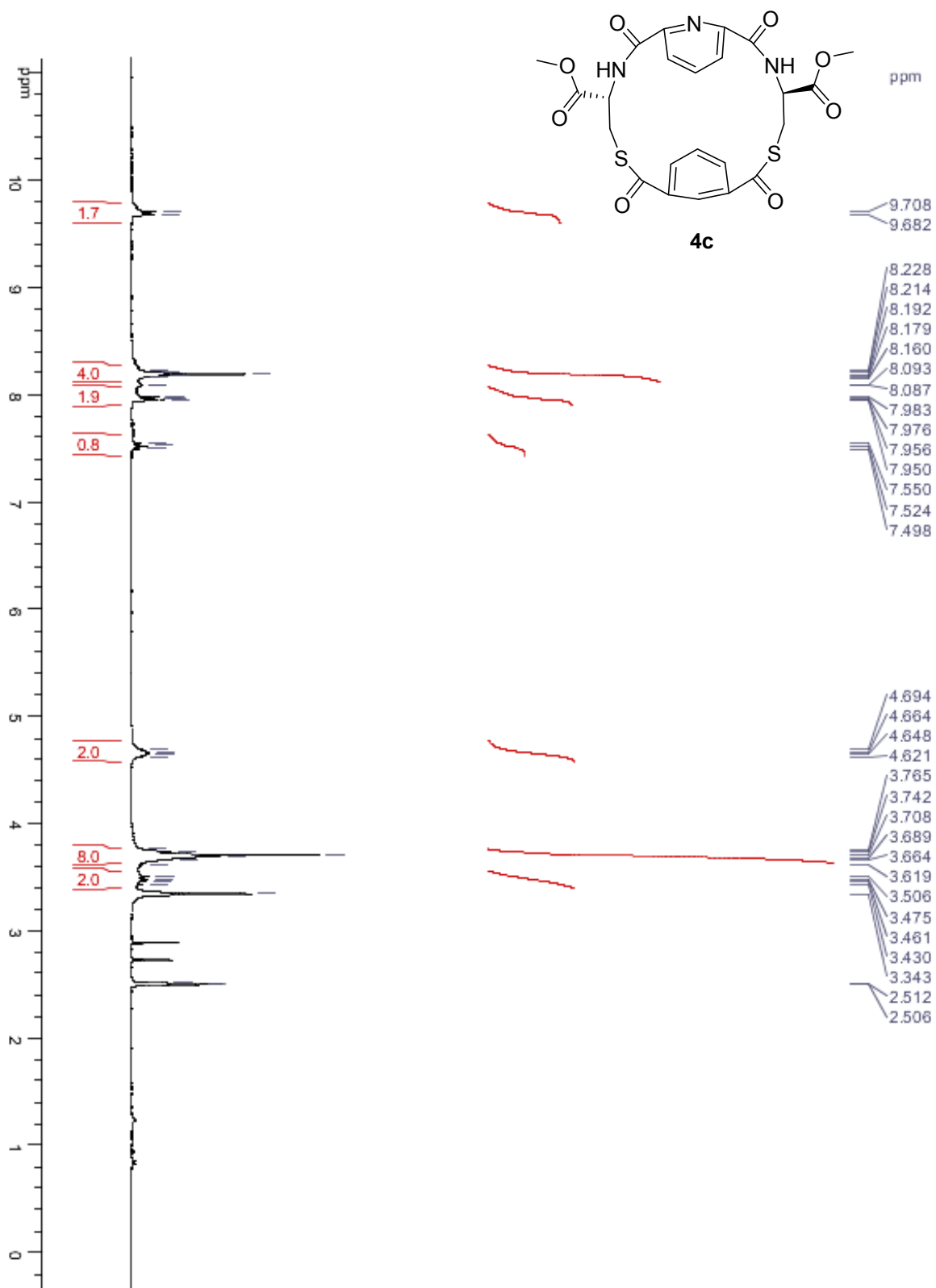


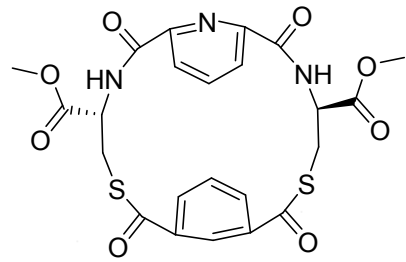




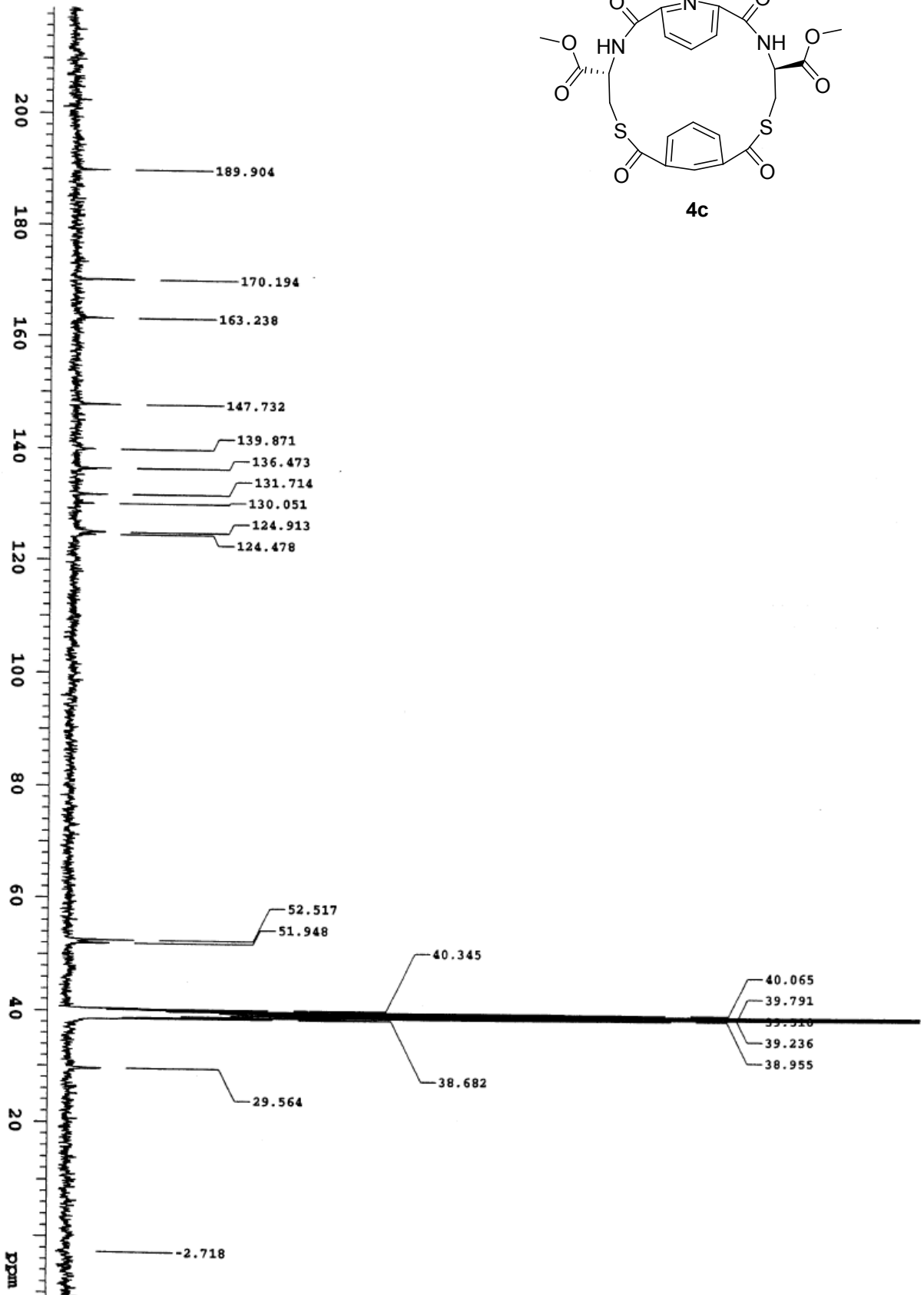


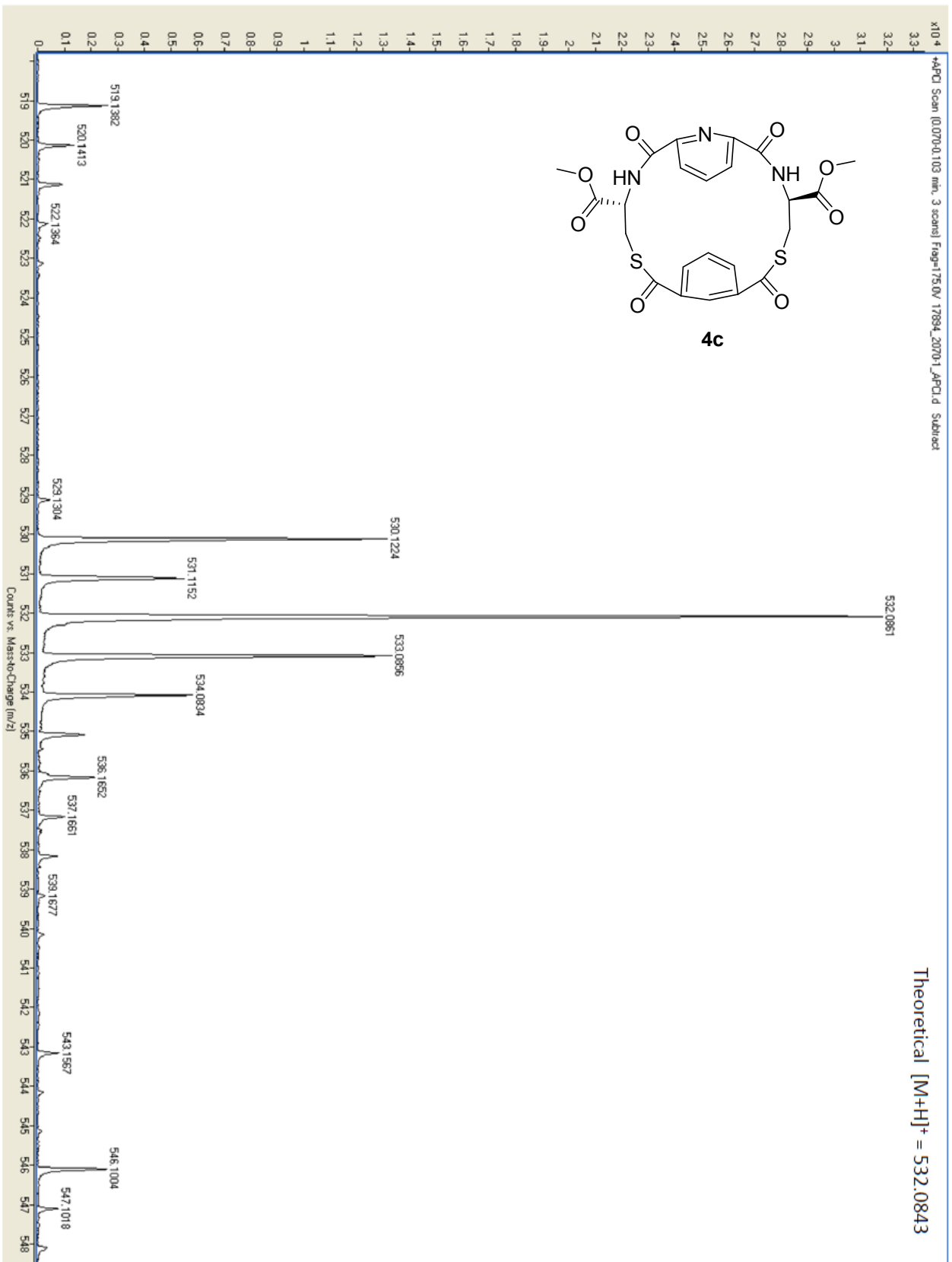


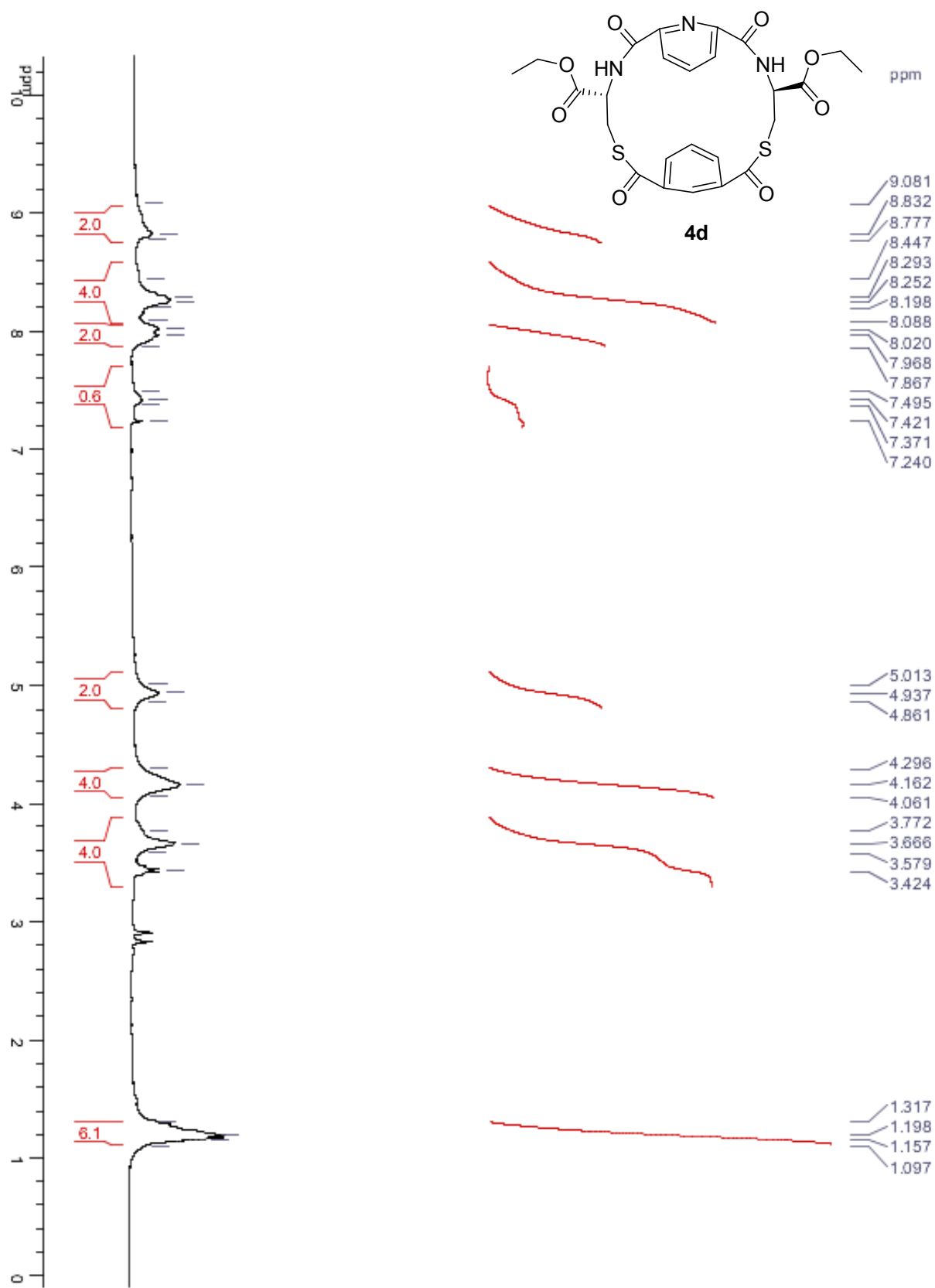


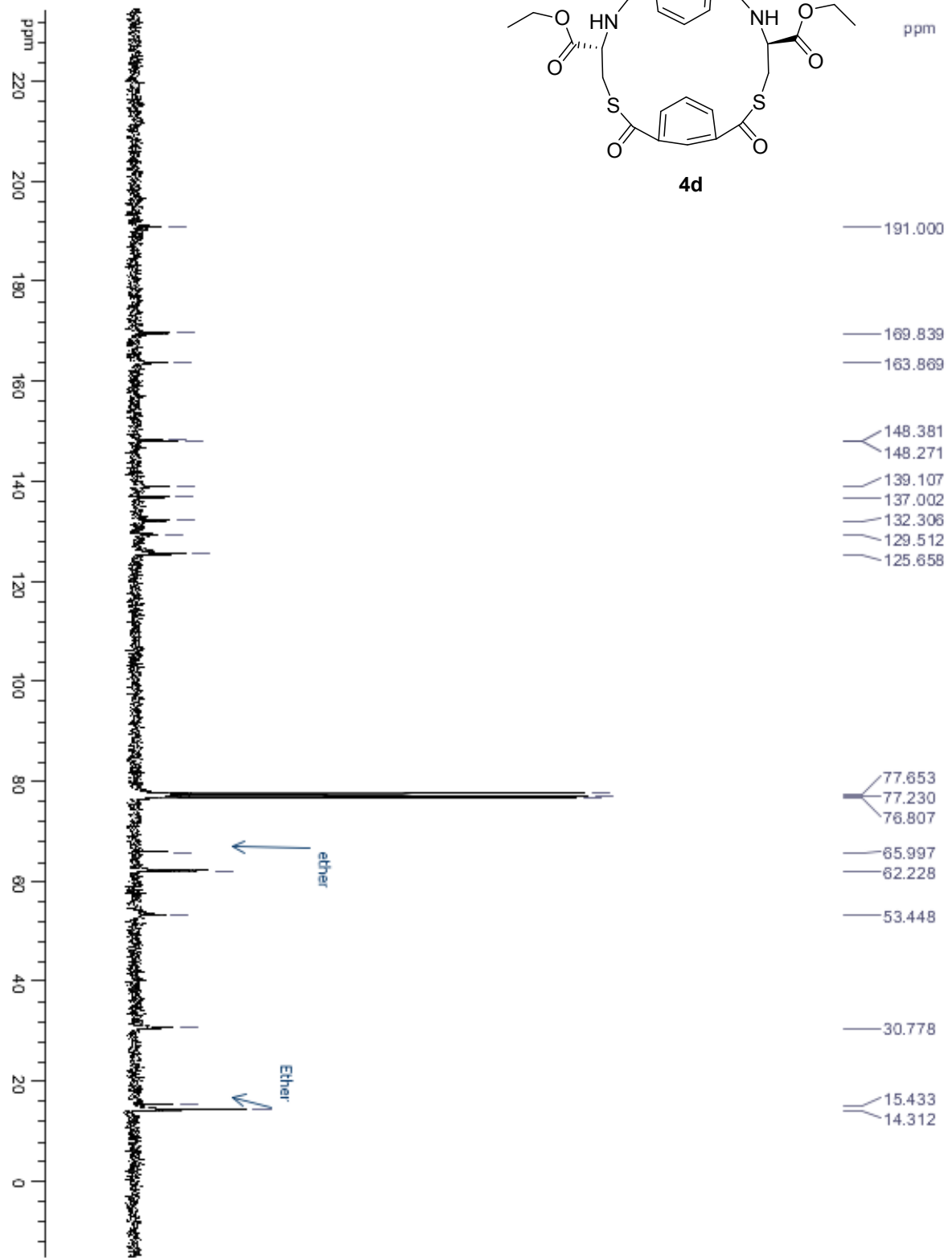
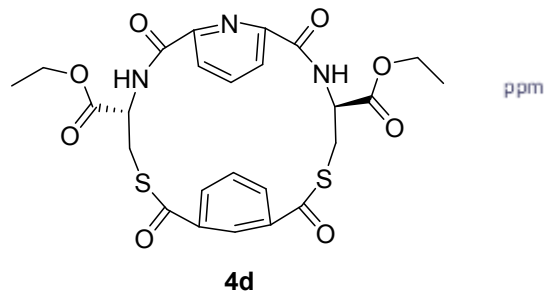


4c



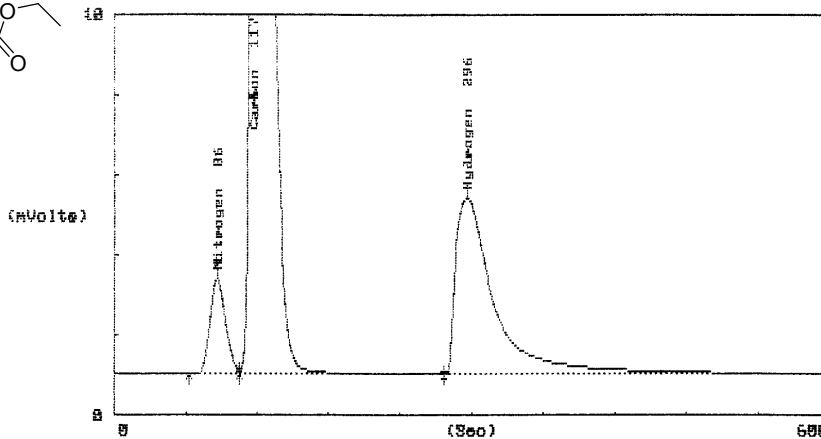
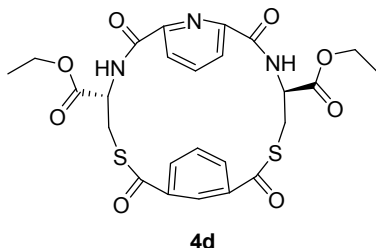






EAGER 200 Stripchart

Sample Ident. : 23 ^{phthalic} ~~mucoic~~ ~~phthalic~~ ethyl Filename : 269423
 Analysed : 04-25-12 08:12:19 Printed : 04-25-2012 08:22:21



EAGER 200 Peak Integration Report

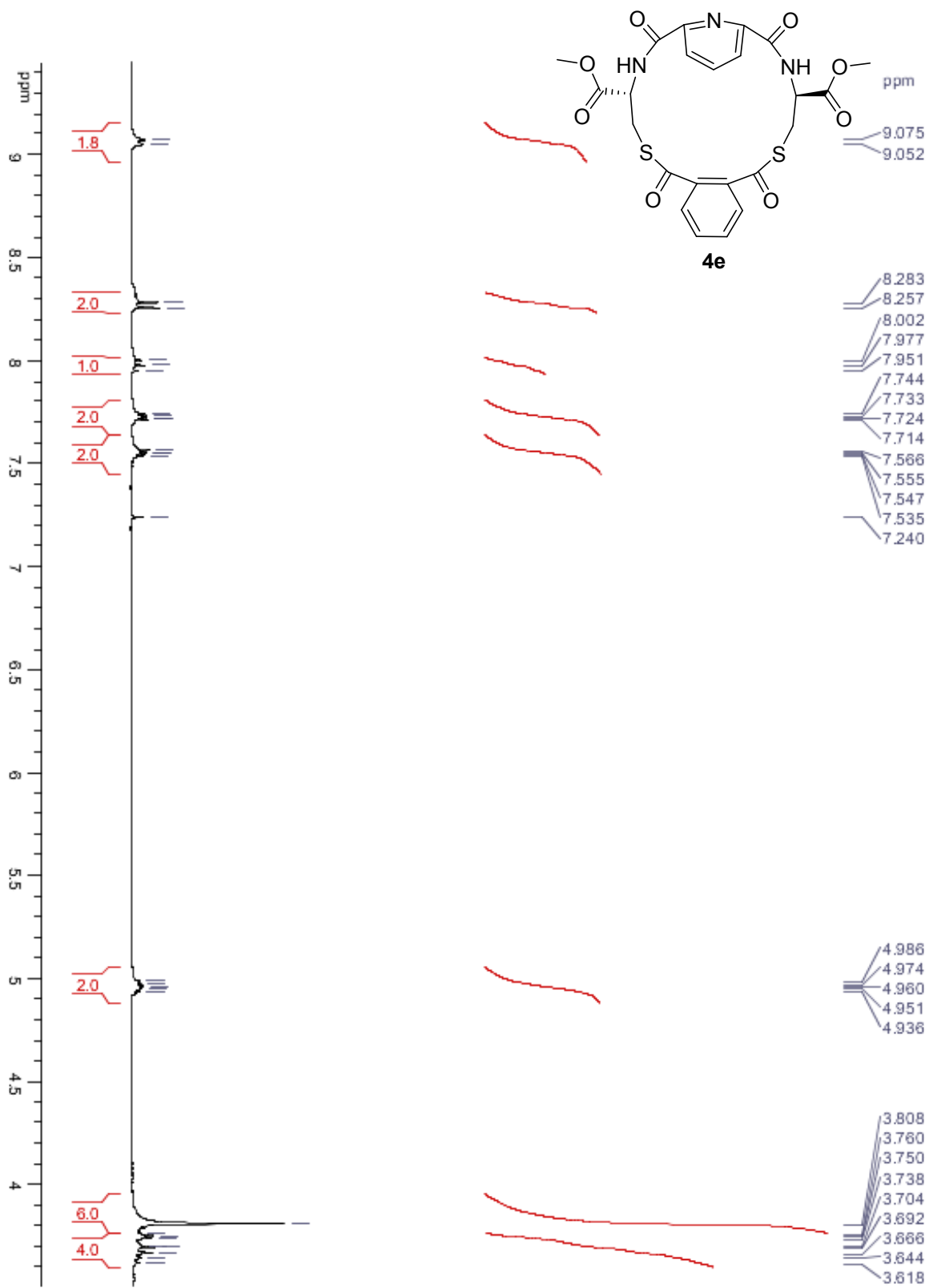
Instrument name : Instrument #1 Bline drift (fV): 6.8
 Company Name : U of Florida Operator Ident. : KOU
 Analysed : 04-25-12 08:12:19 Printed : 04-25-2012 08:22:22
 Sample Ident. : 23 1985-01 Filename : 269423
 Sample Weight : 2.111 Calc.method: using 'K. Factors'

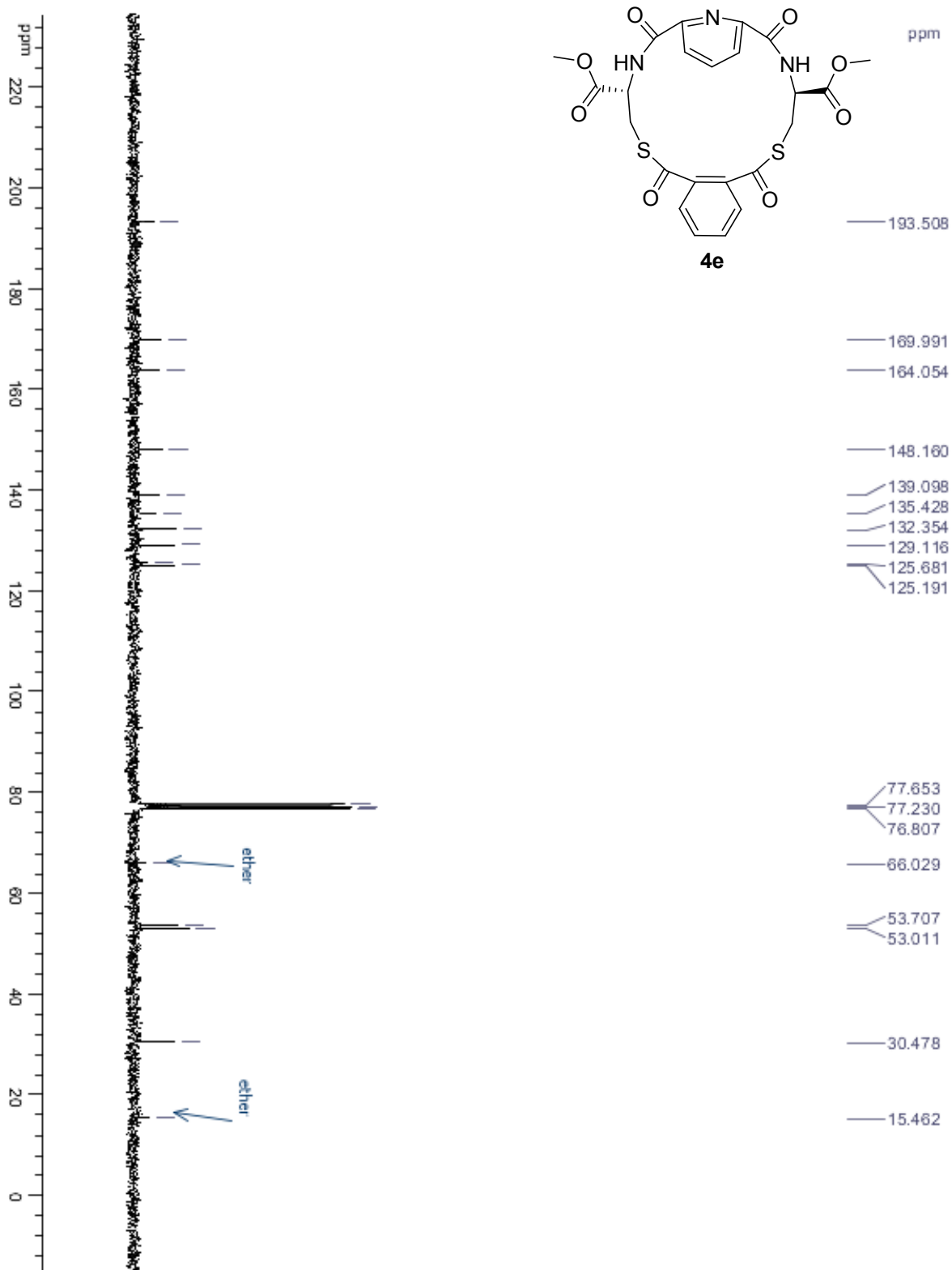
No. (#)	Type (#)	Start (Sec)	End (Sec)	Ret Time (Sec)	Height (fV)	Area (fV*Sec)	Area % (%)	Name
1	FU	63	105	86	2391.2	34425	3.82	Nitrogen
2	FU	105	277	117	44233.0	696815	77.23	Carbon
3	RS	277	597	296	4385.9	171012	18.95	Hydrogen
							902252	100.00

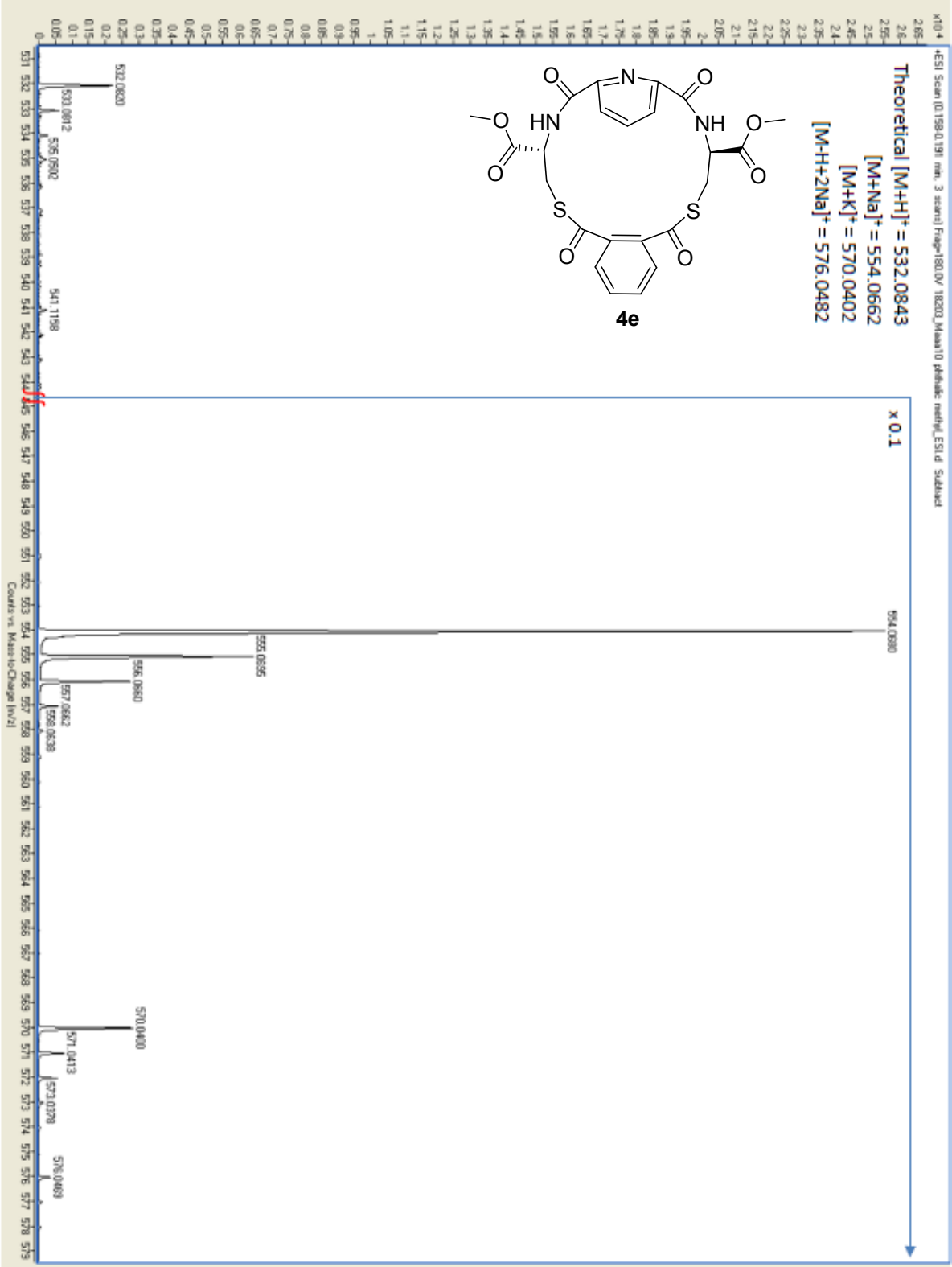
EAGER 200 Unk Report

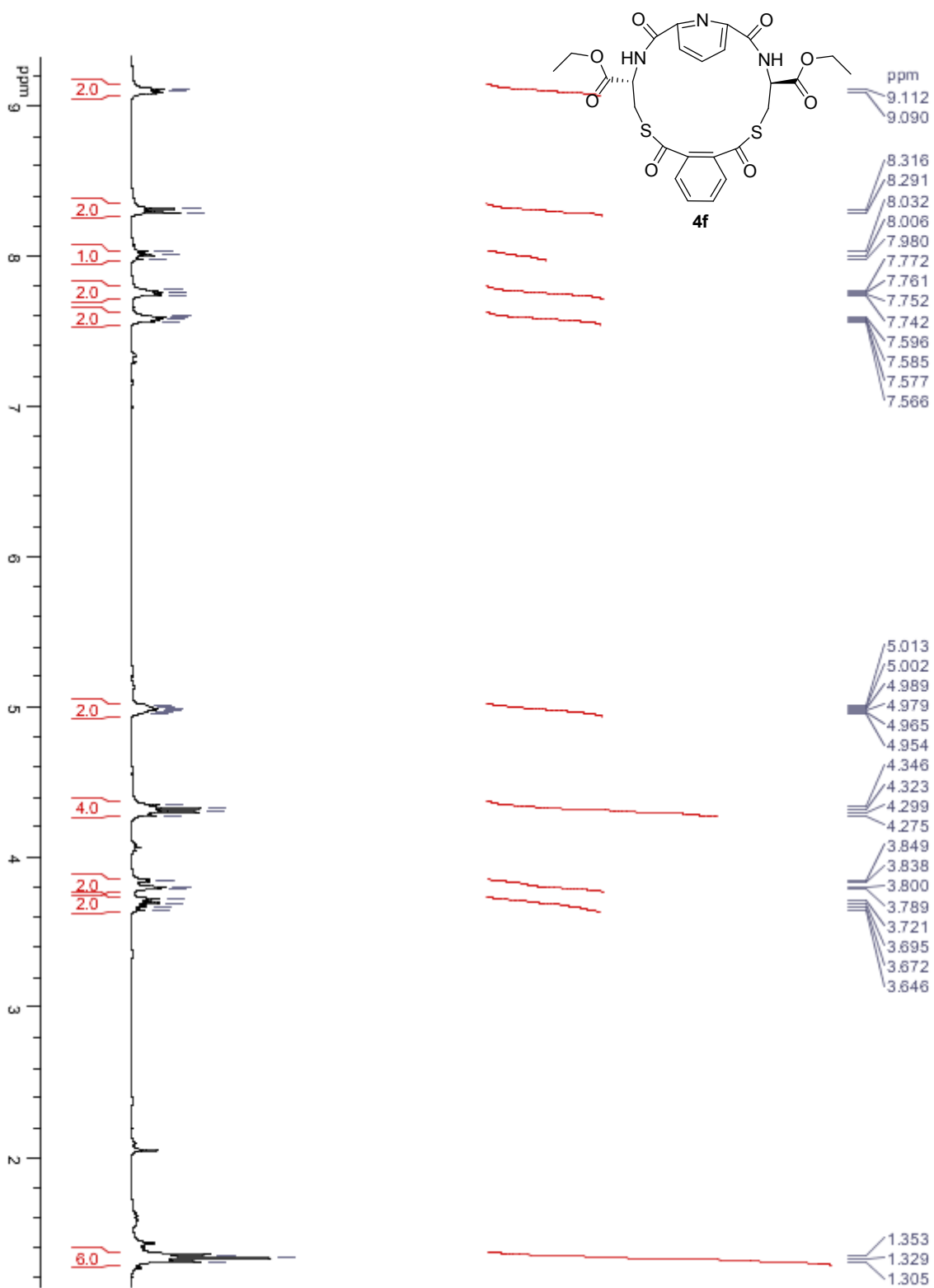
Instrument name : Instrument #1 Bline drift (fV): 6.8
 Company Name : U of Florida Operator Ident. : KOU
 Analysed : 04-25-12 08:12:19 Printed : 04-25-2012 08:22:22
 Sample Ident. : 23 1985-01 Filename : 269423
 Sample Weight : 2.111 Calc.method: using 'K. Factors'

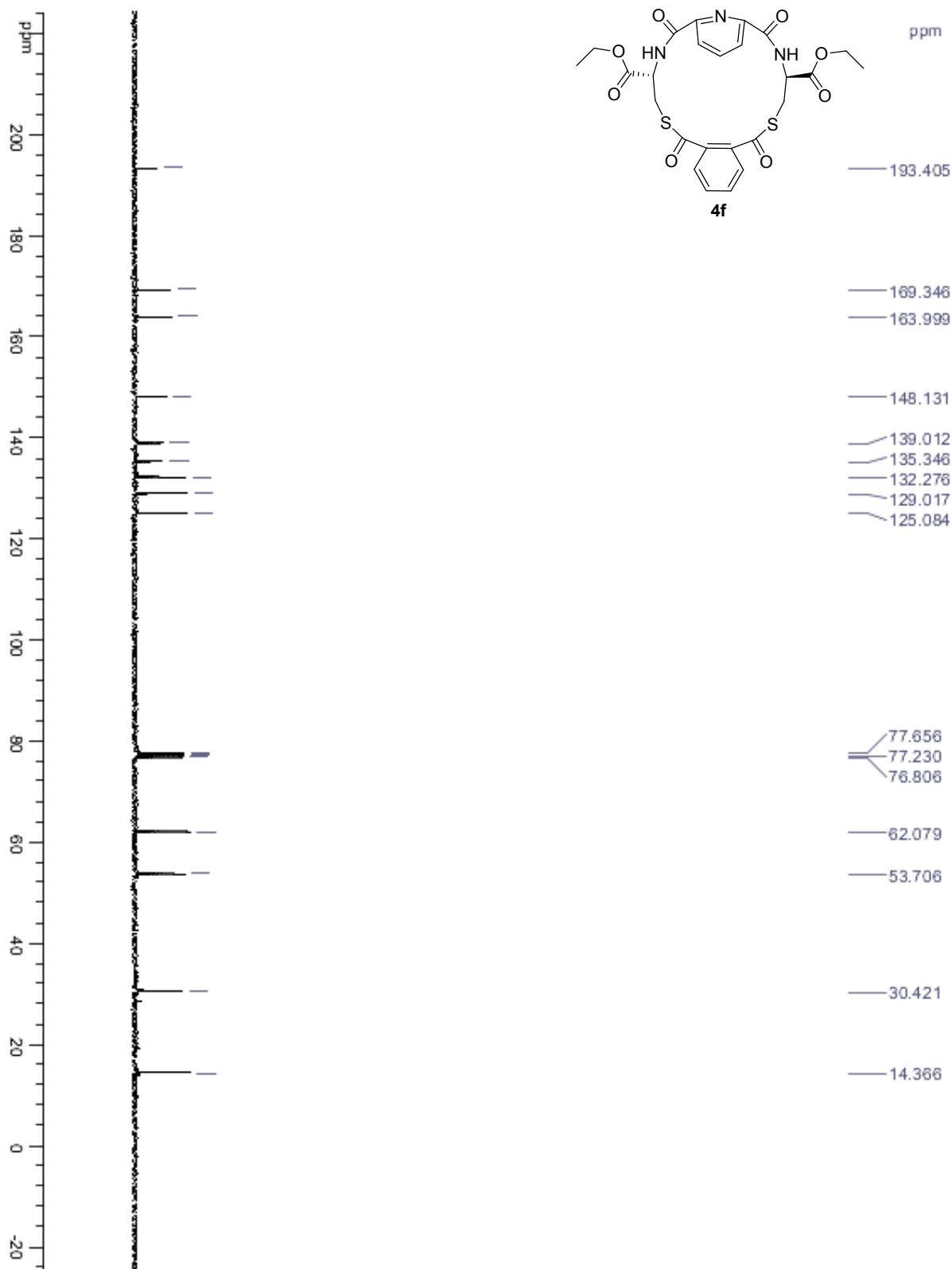
Pk. (#)	Ret Time (Sec)	Area (fV*Sec)	Element % (%)	Area Ratio	Name
1	86	34425	7.248	.202415E+02	Nitrogen
2	117	696815	53.546	.100000E+01	Carbon
3	296	171012	4.749	.407465E+01	Hydrogen

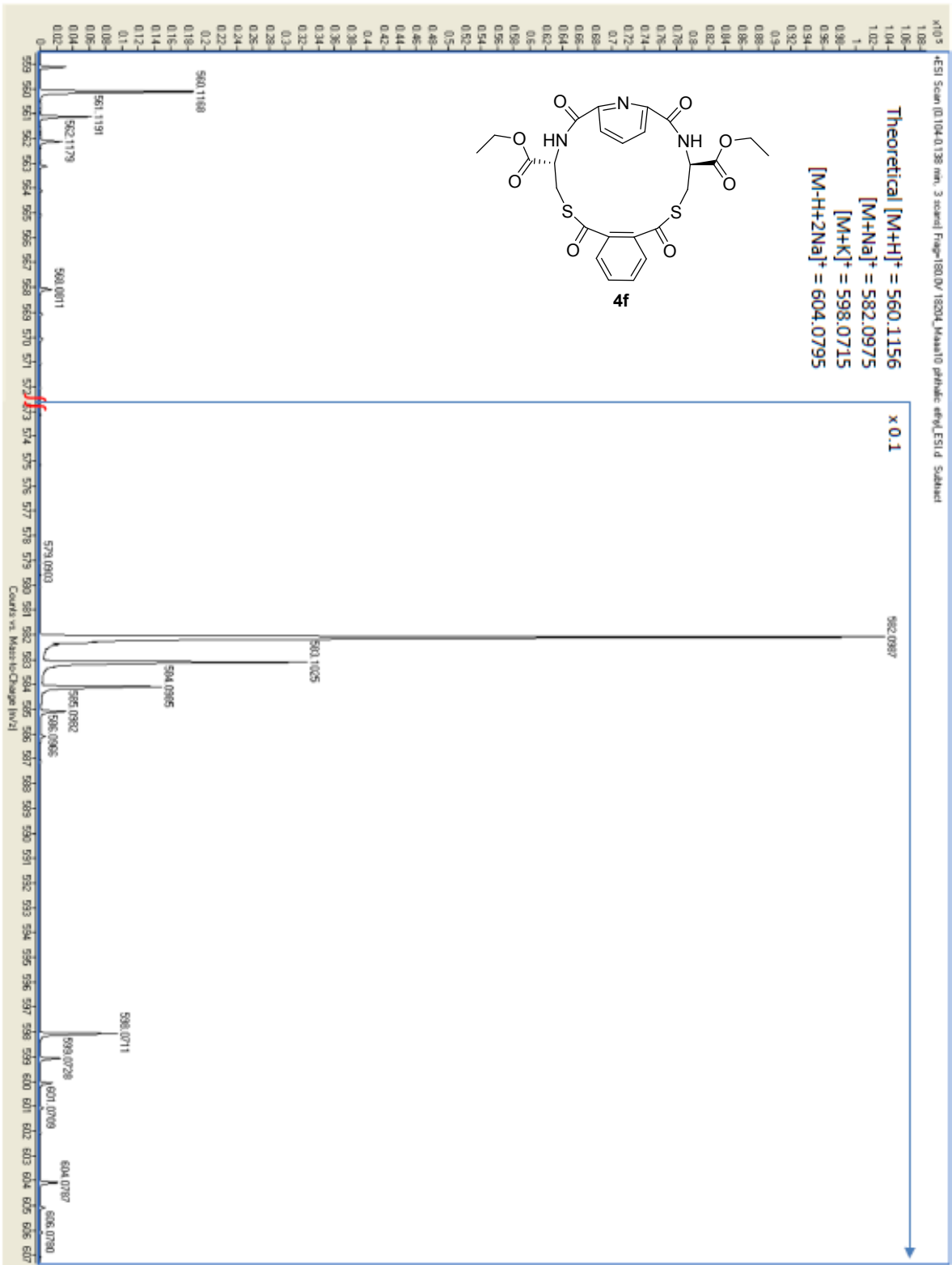


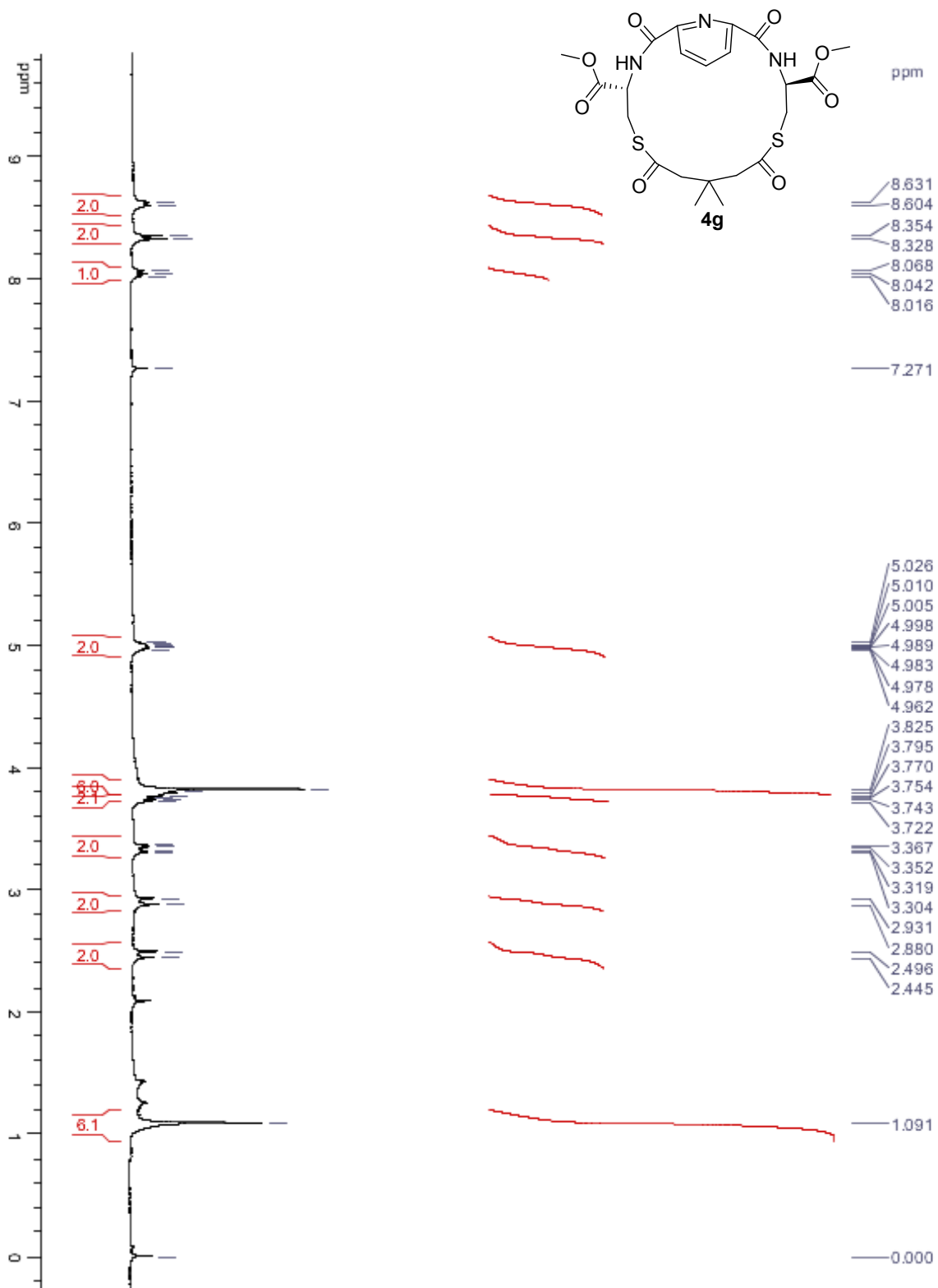


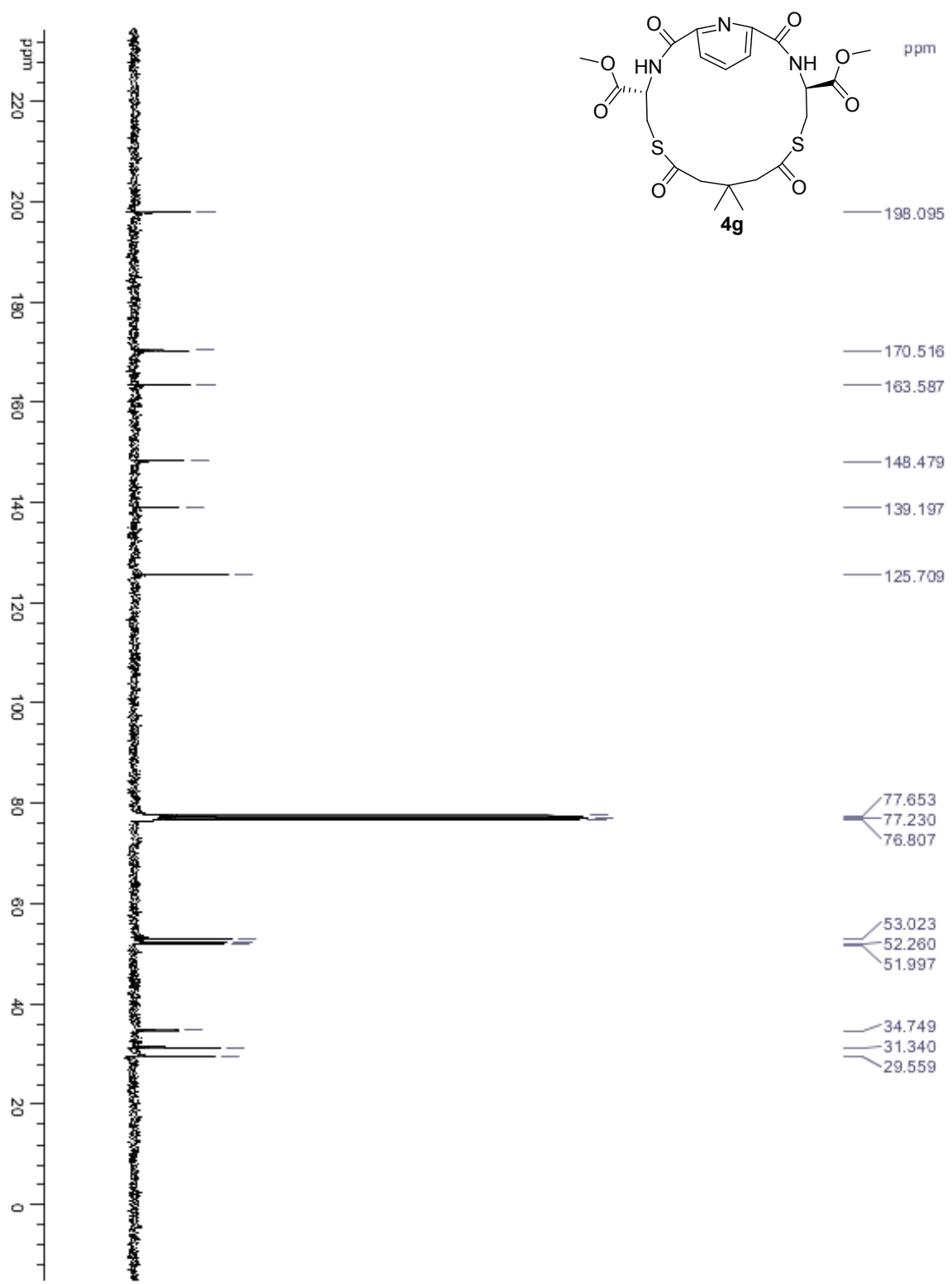


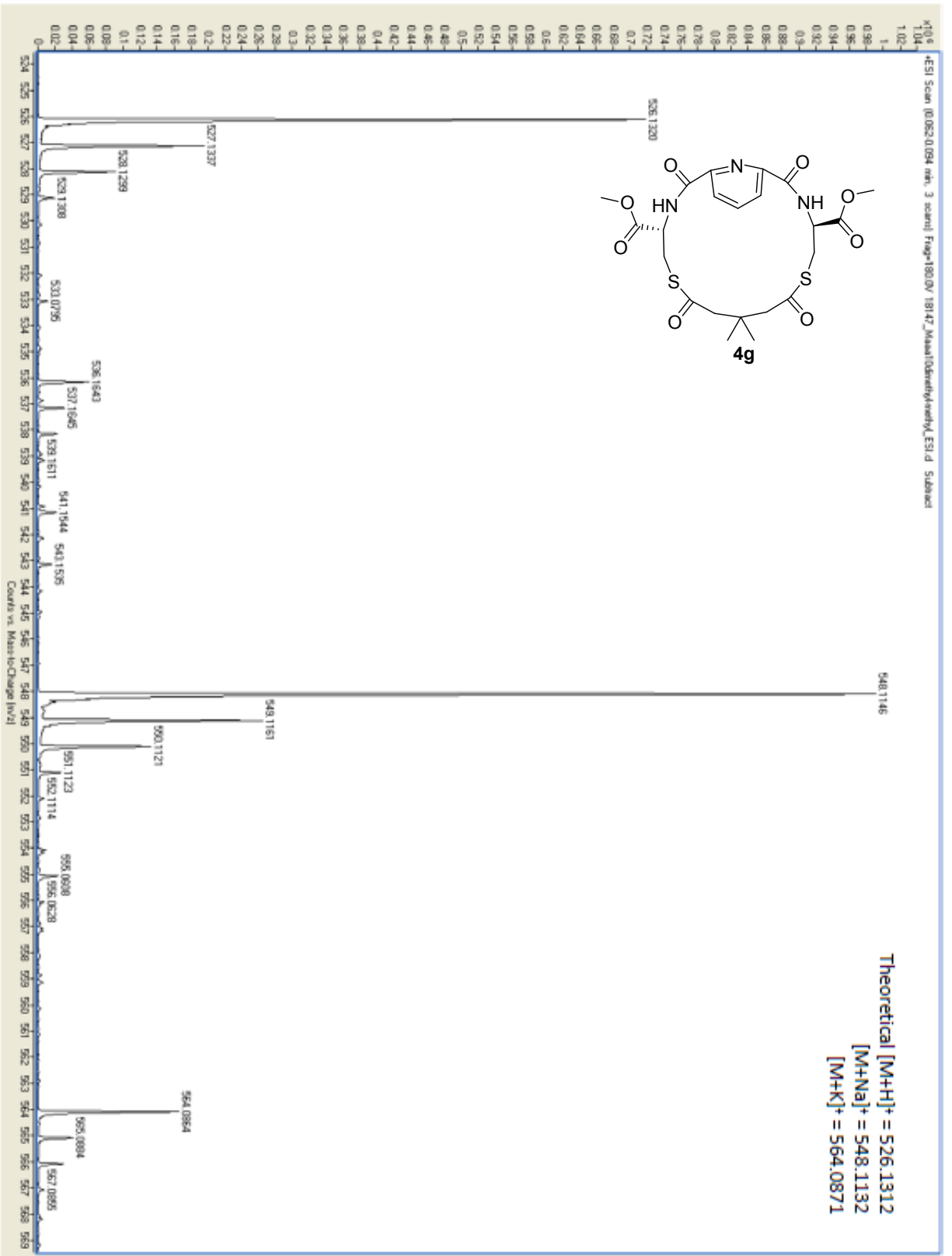


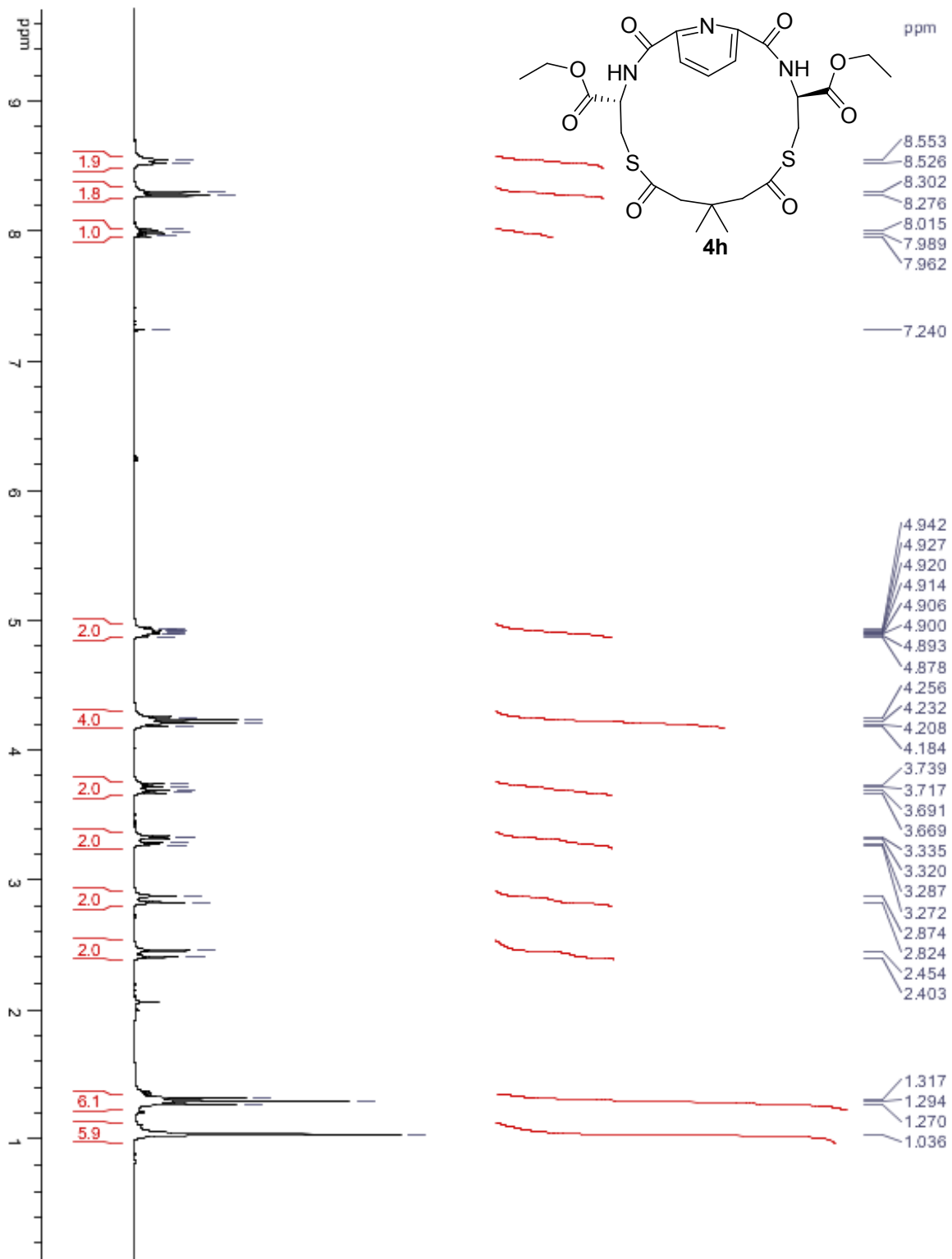


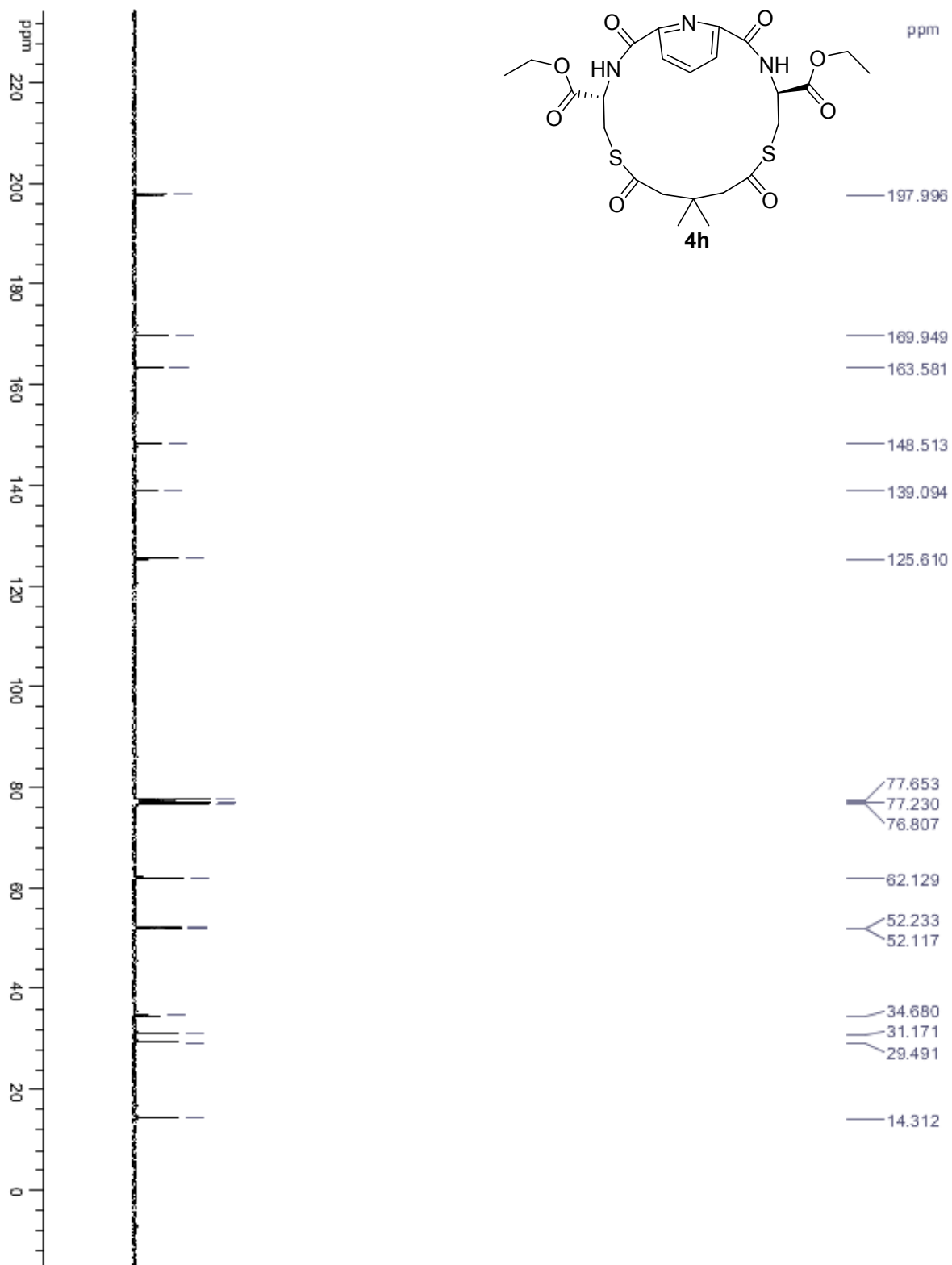


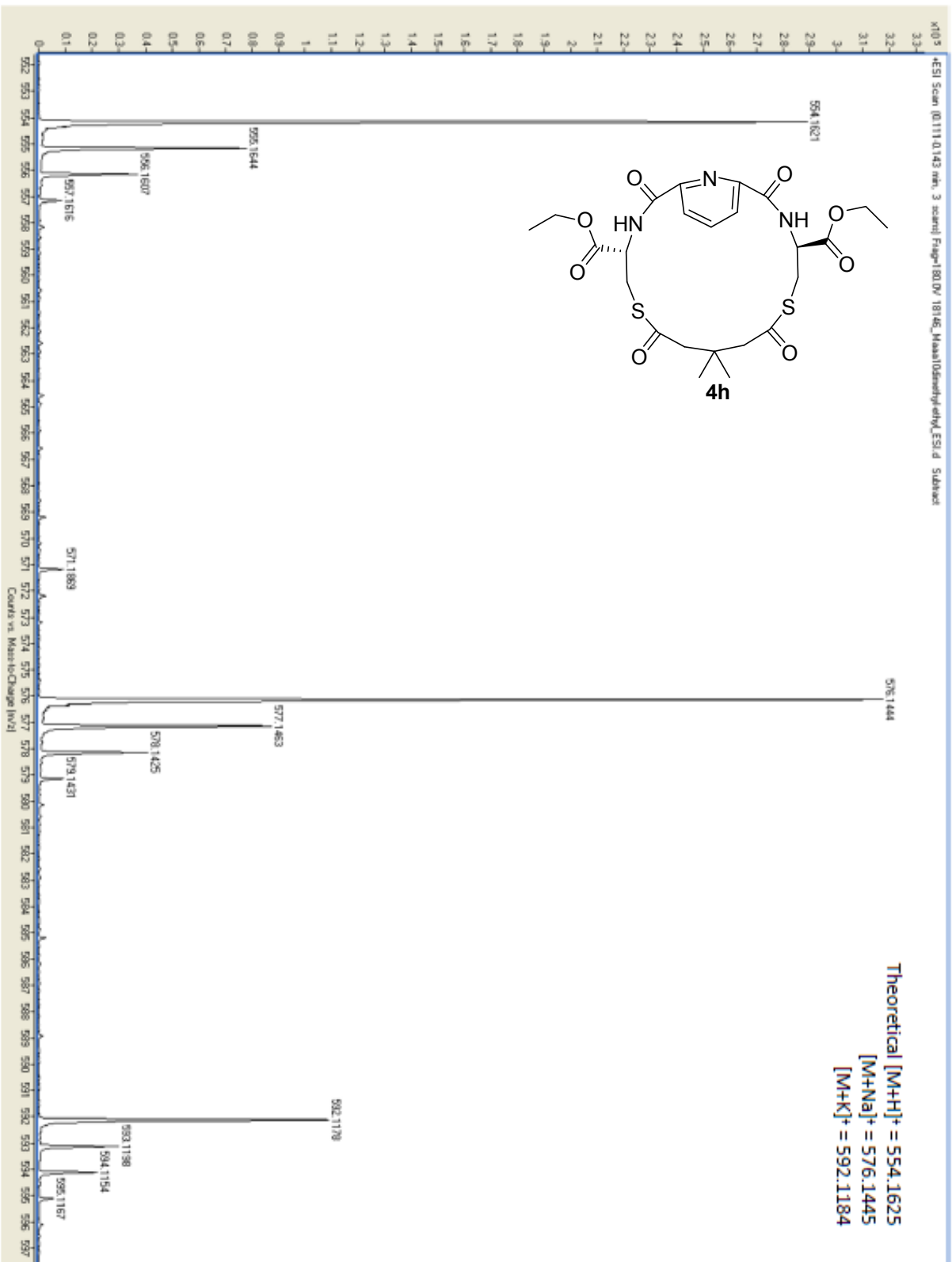


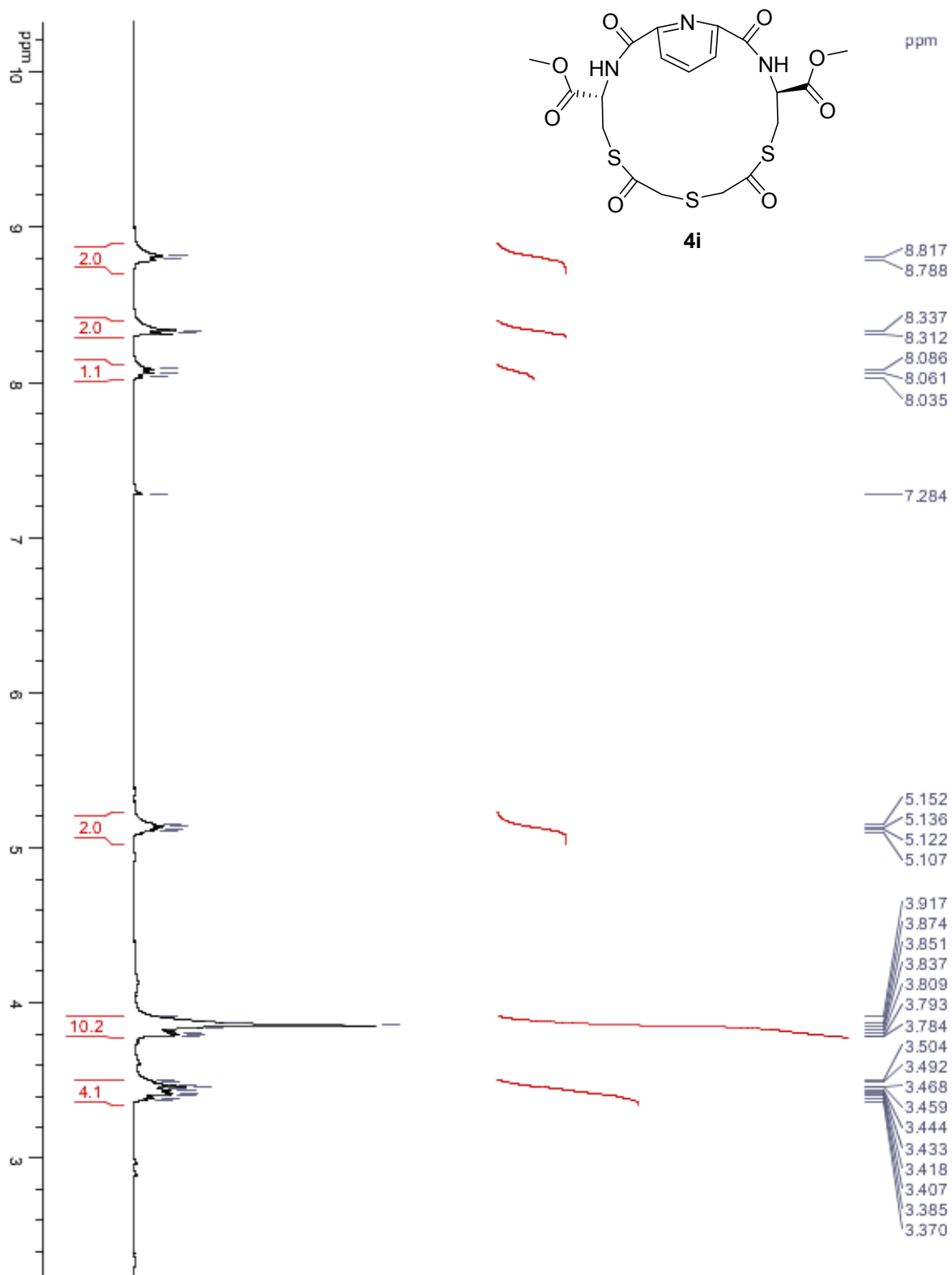


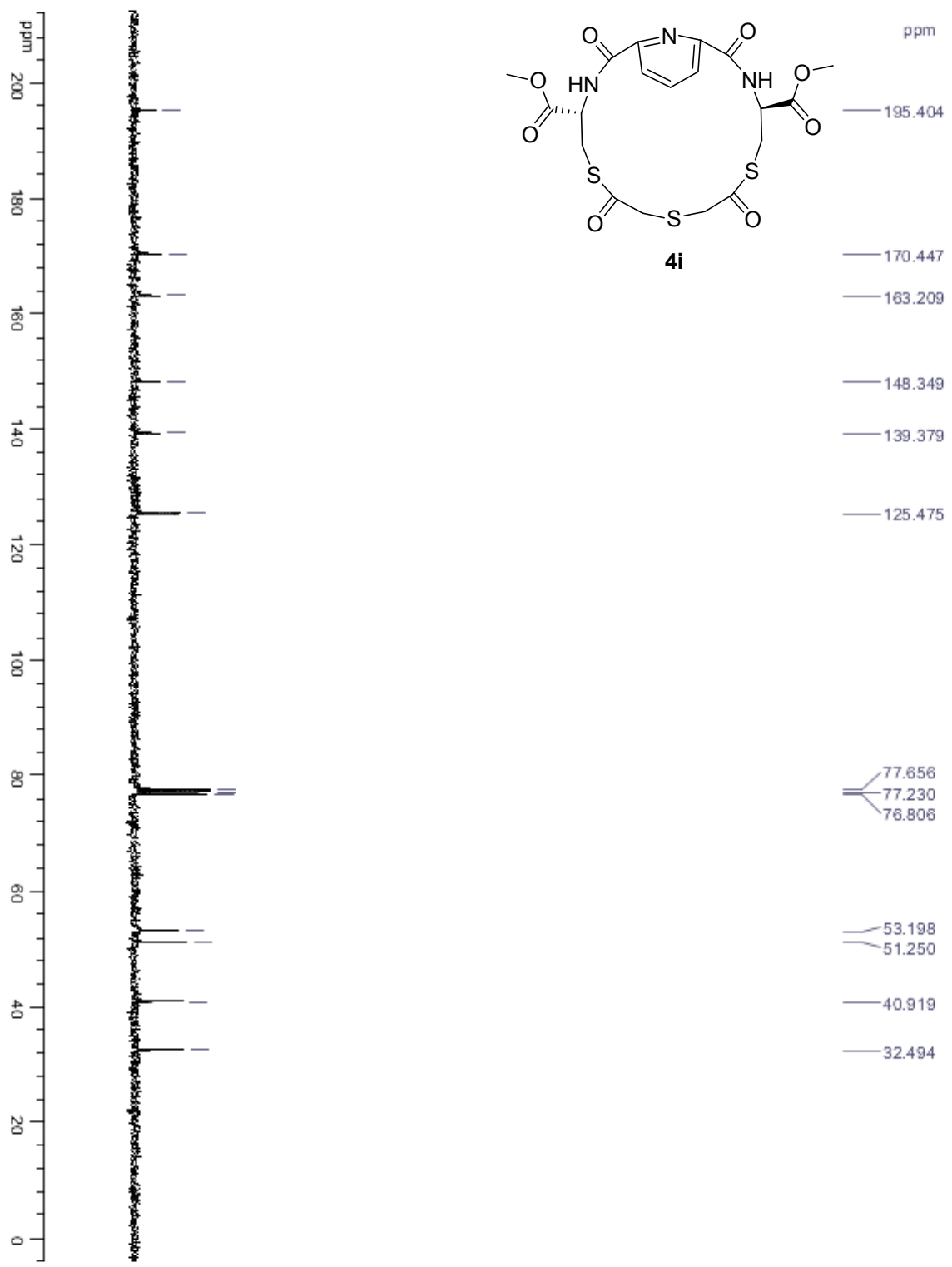


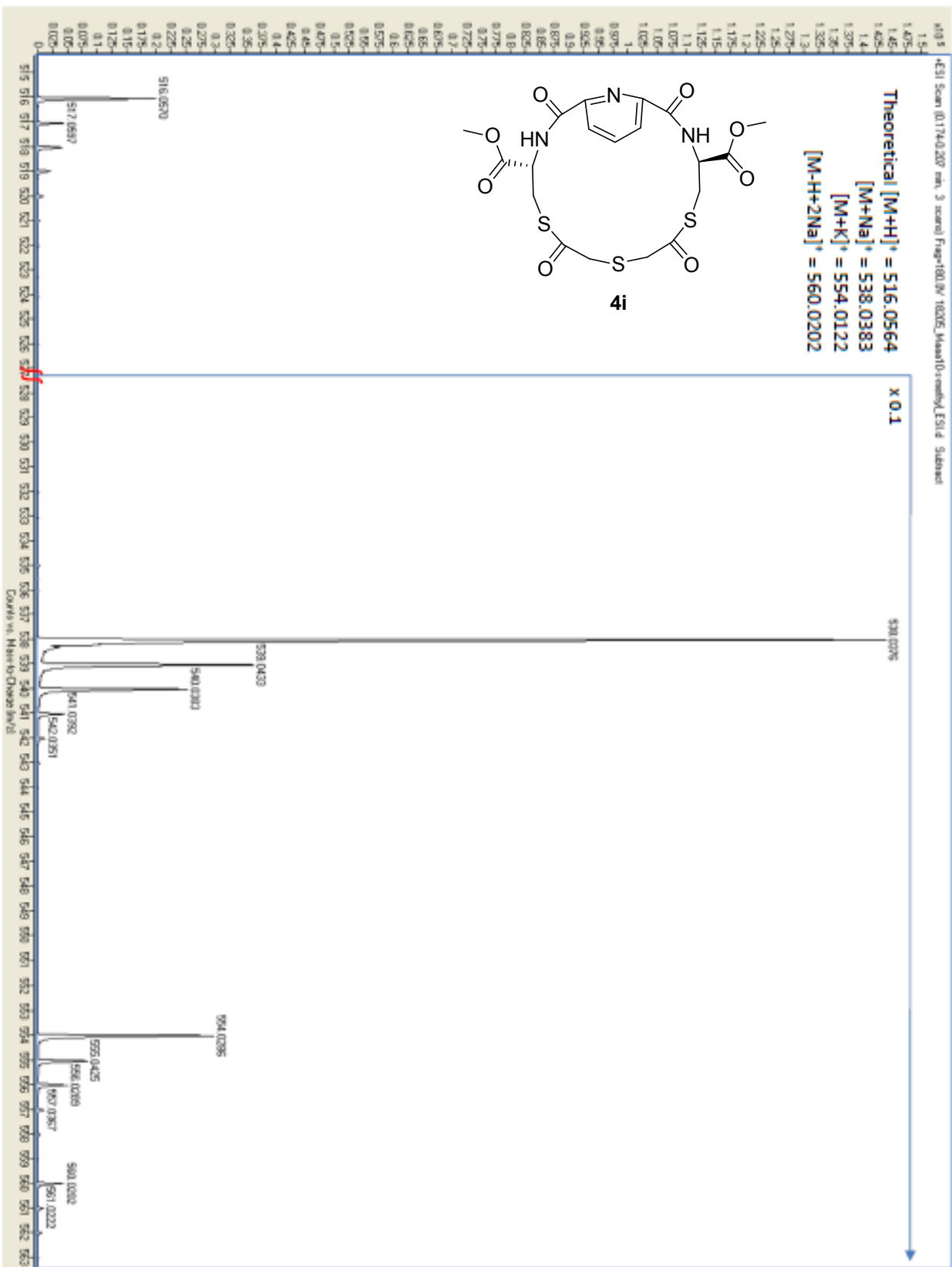


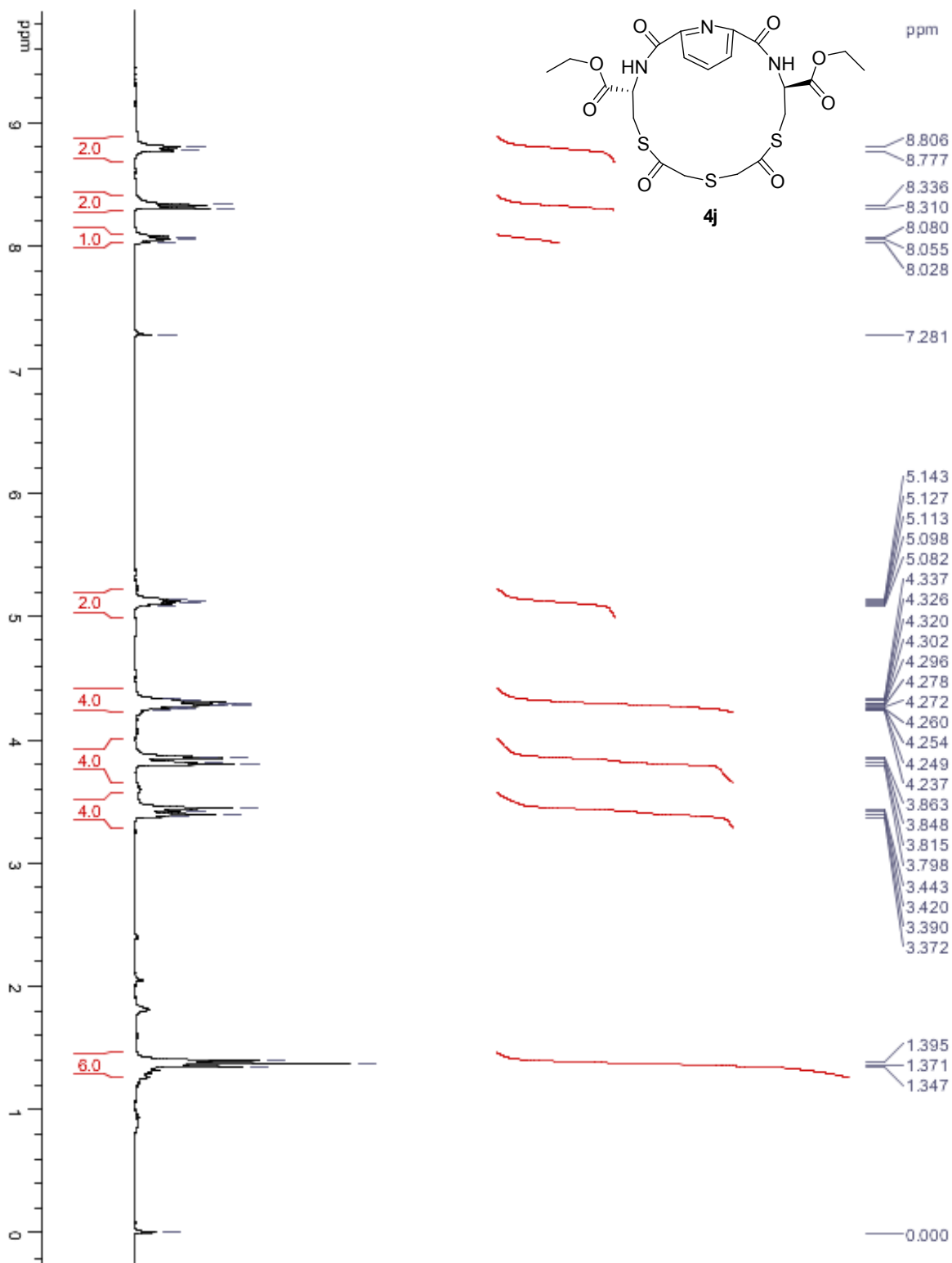


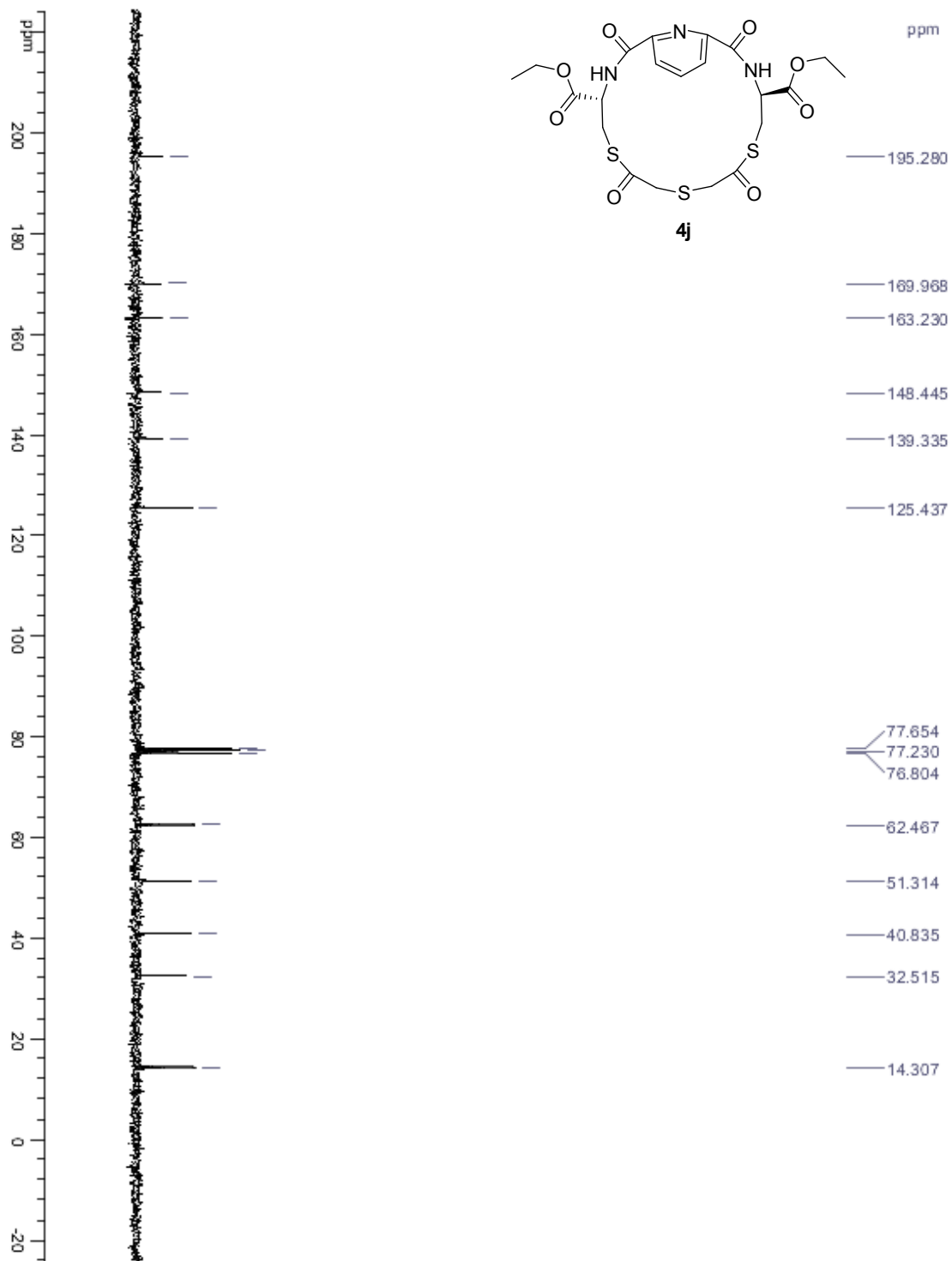


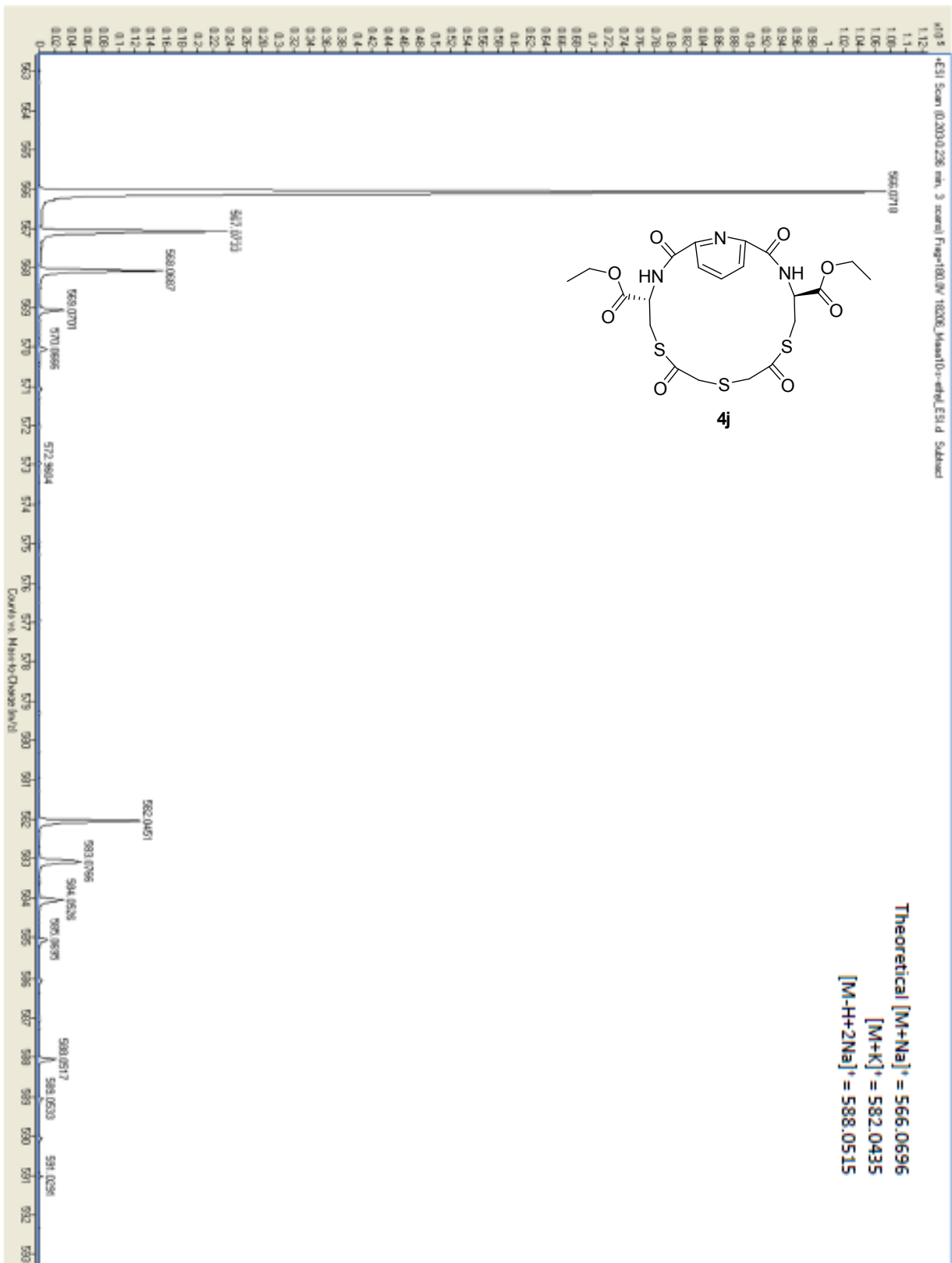


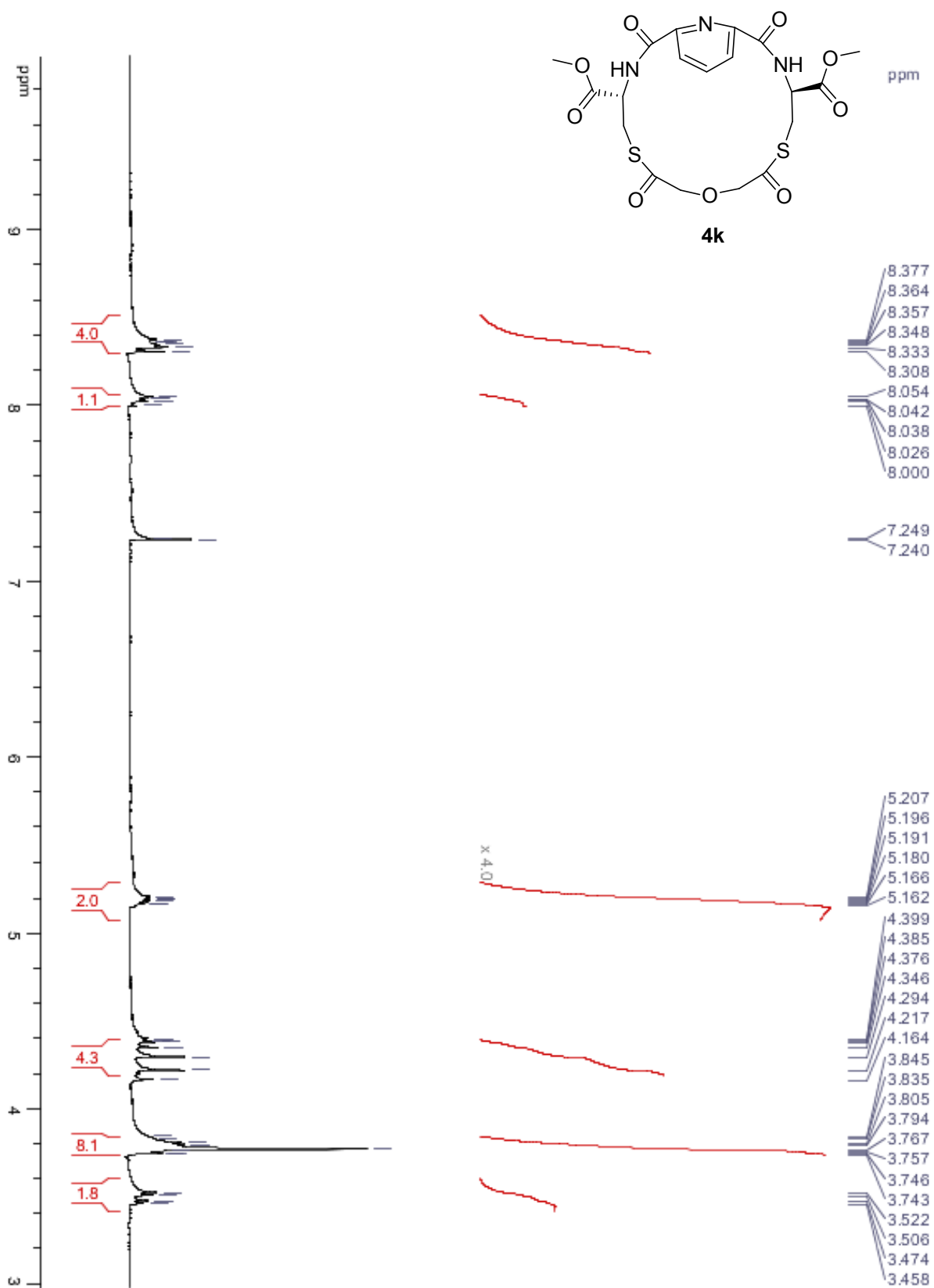


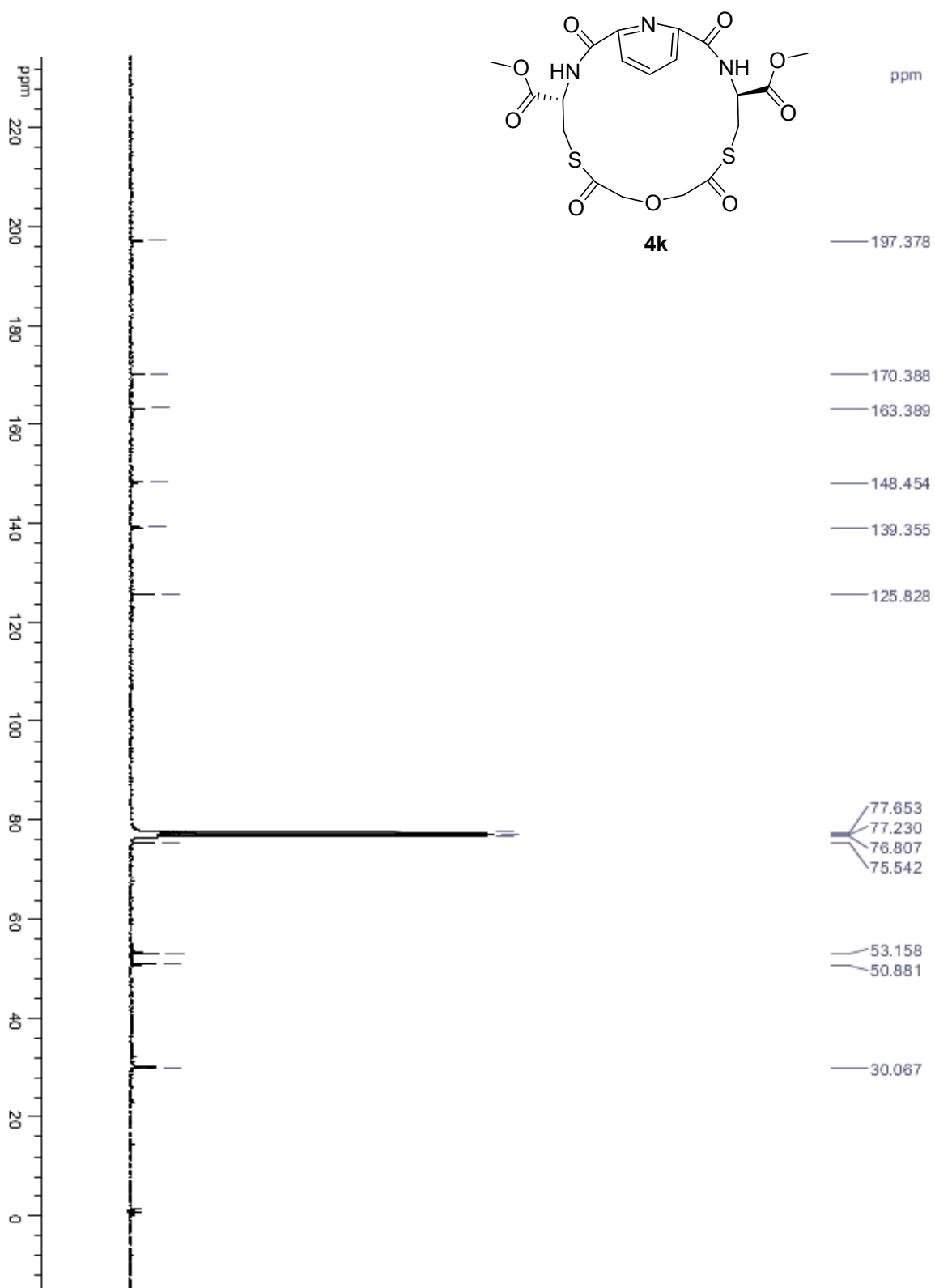


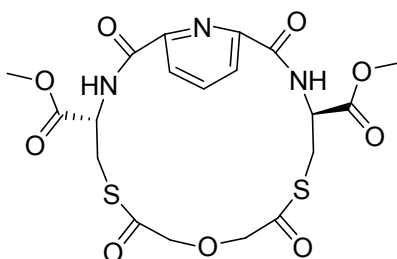






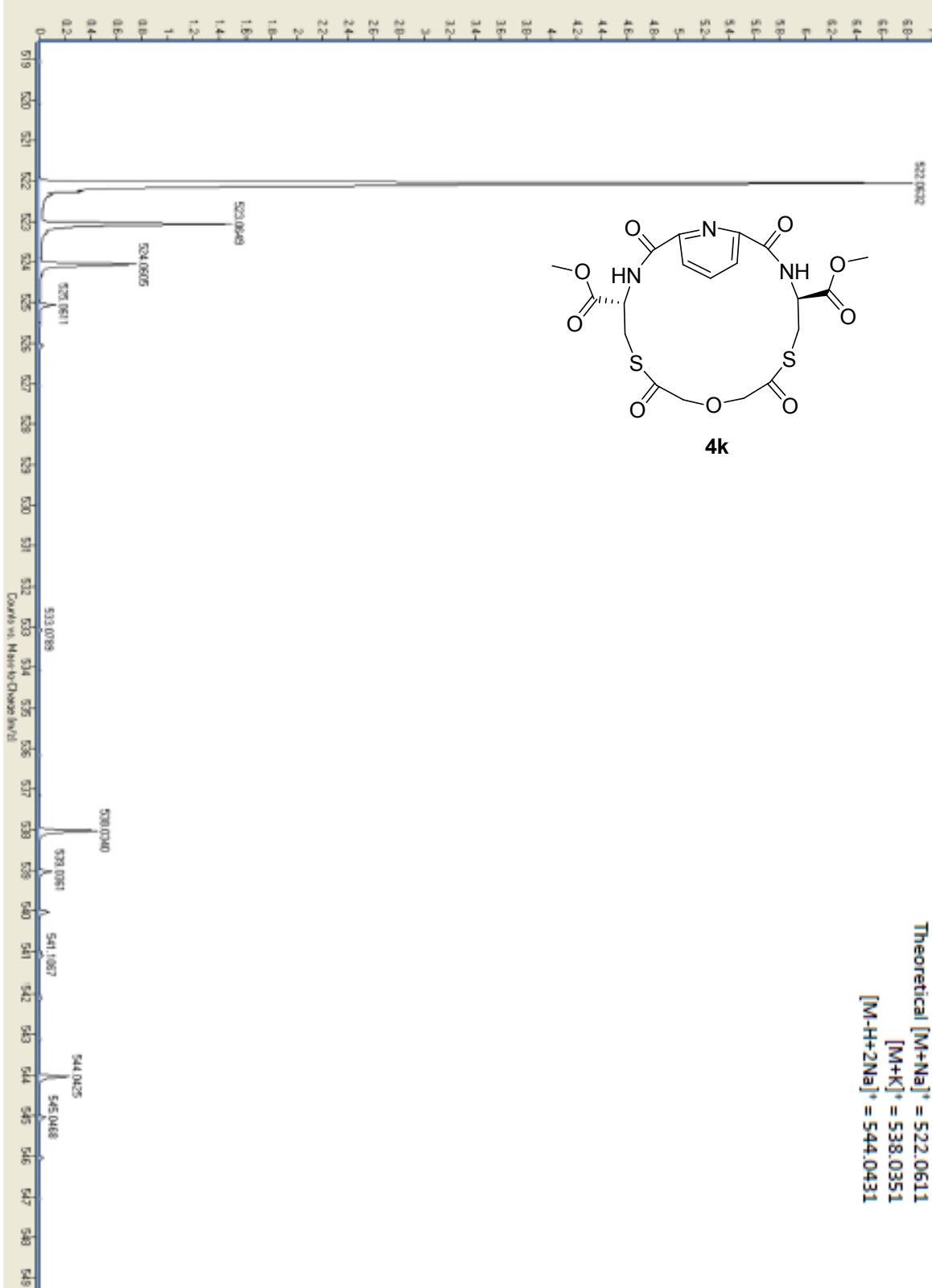


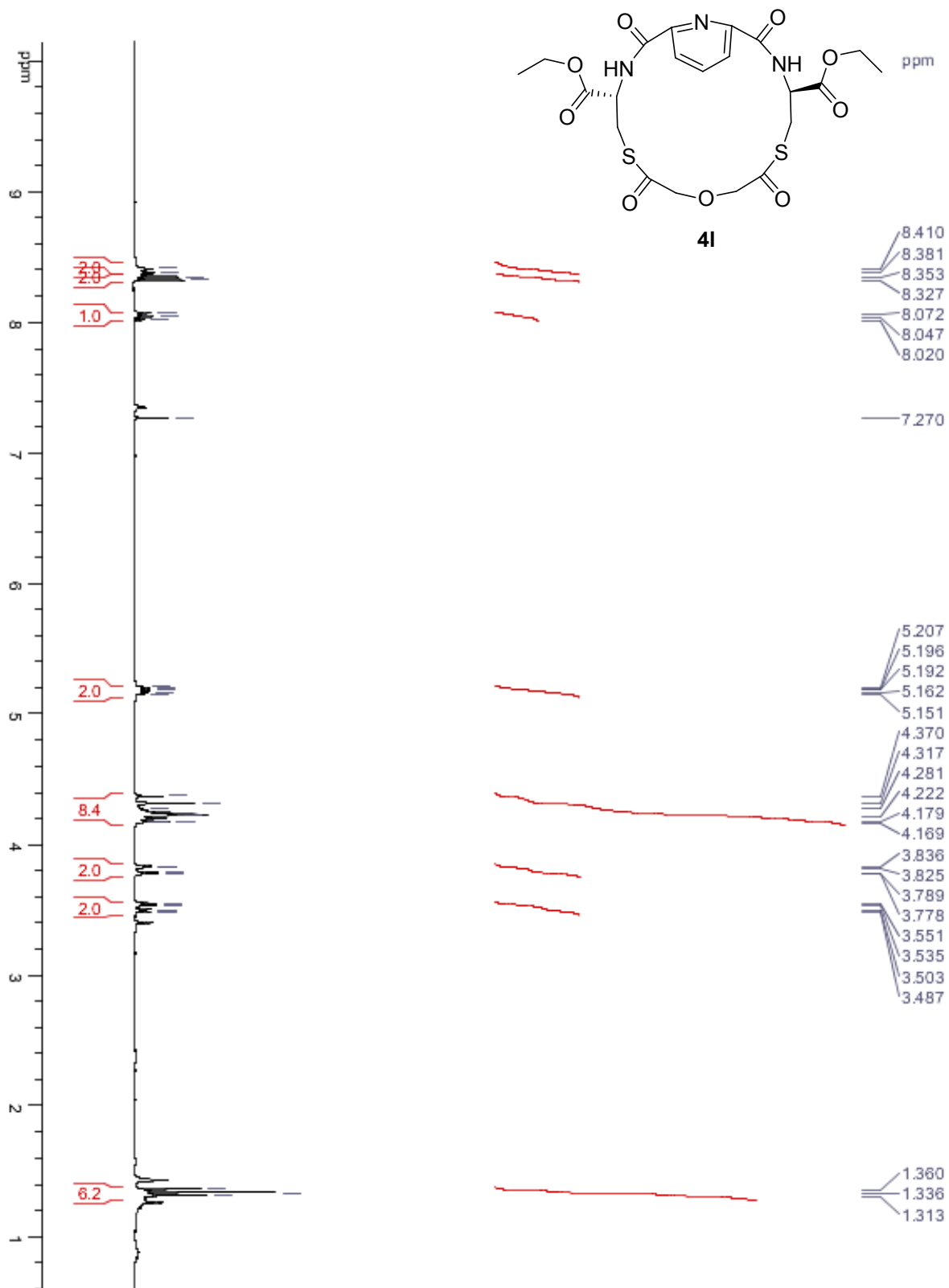


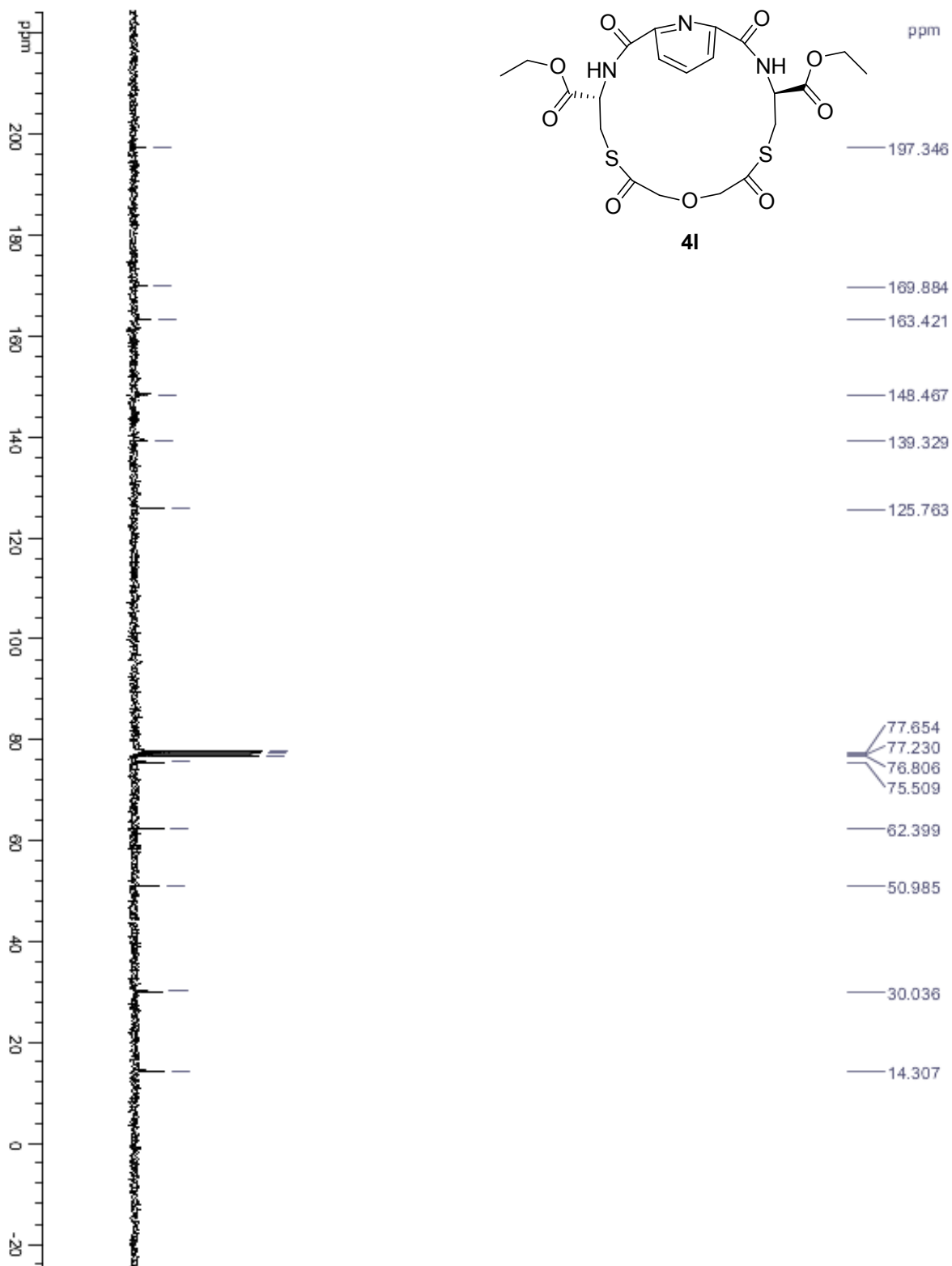


4k

Theoretical $[M+Na]^+$ = 522.0611
 $[M+K]^+$ = 538.0351
 $[M-H+2Na]^+$ = 544.0431







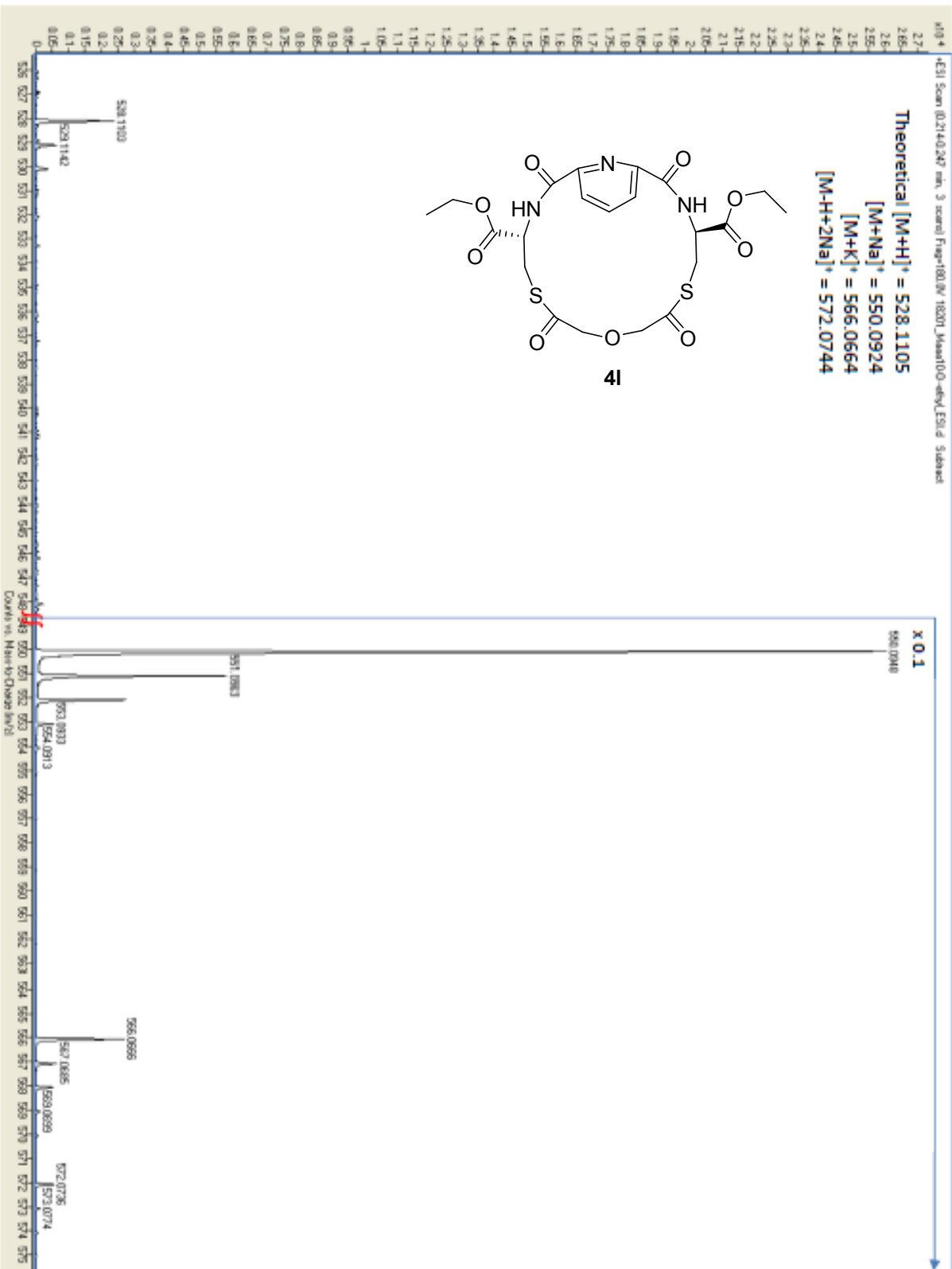


Table S1. Molecular descriptor values of the BMLR-QSAR model for the antifungal active macrocyclic peptidomimetics.

Entry	Compd.	Descriptors*		
		D ₁	D ₂	D ₃
1	4a	0.0043	-1153.12	19.5355
2	4b	0.0046	-1209.1	19.4839
3	4c	0.00623	-1264.3	19.5096
4	4d	0.00455	-1460.3	19.5223
5	4e	0.00944	-1268.36	19.5329
6	4f	0.0091	-1300.97	19.4882
7	4g	0.00238	-1221.38	19.468
8	4h	0.00213	-1061.68	19.4667
9	4i	0.00559	-1047.9	19.4659
10	4j	0.00159	-1017.77	19.4645
11	4k	0.00755	-934.668	19.5165
12	4l	0.00235	-1179.24	19.4717
13	5	0.01884	-1221.8	19.5034
14	6	0.02522	-1232.66	19.5044
15	7	0.01537	-900.929	19.472

*D₁ = FHDCA Fractional HDCA (HDCA/TMSA) (MOPAC PC), D₂ = WNSA-2 Weighted PNSA (PNSA2*TMSA/1000) (MOPAC PC), D₃ = Max. total interaction for bond C-C.

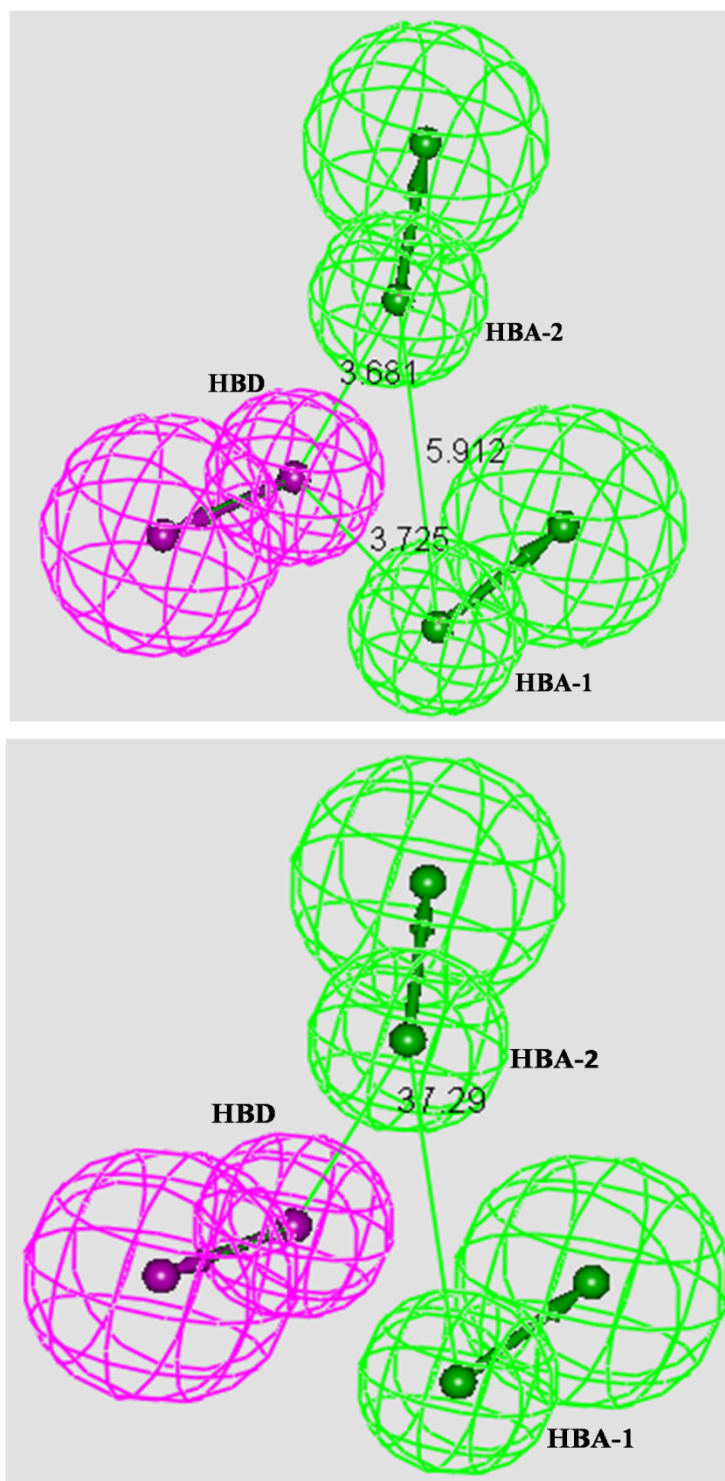
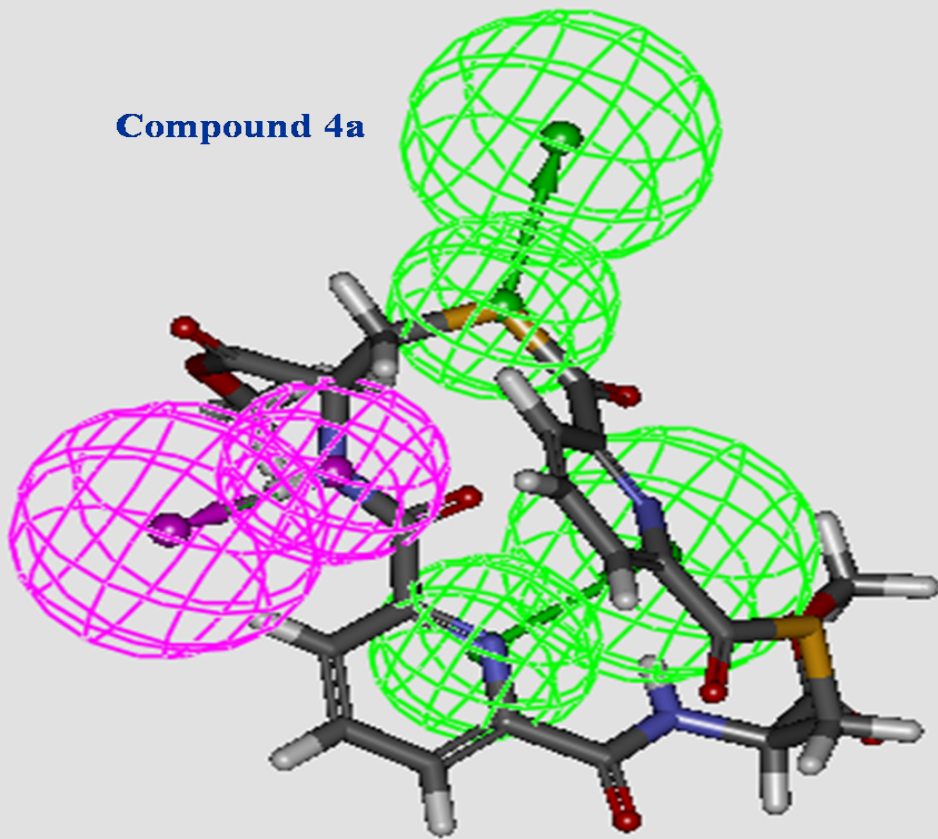
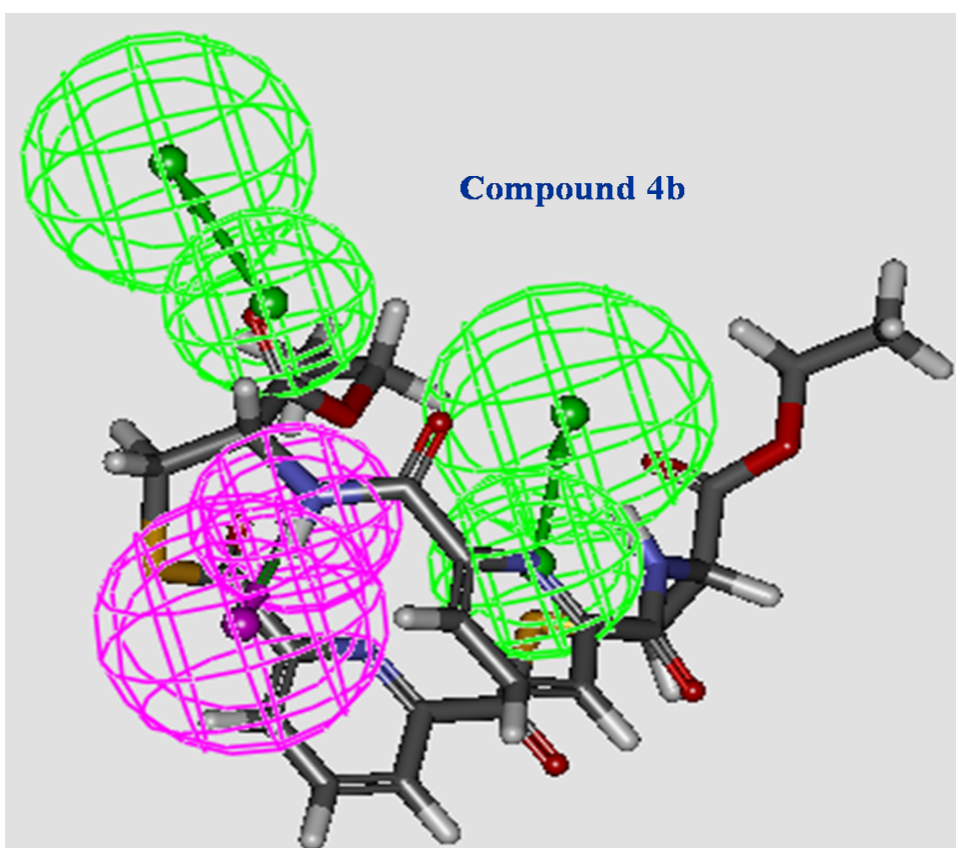


Figure S1. (A) Constraint distances, and (B) constraint angle of the generated 3D-pharmacophore for the antifungal bio-active compounds **4a-4l**, and **5-7** against *C. albicans* which contains two hydrogen bonding acceptor (HBA-1, HBA-2, green), and one hydrogen bonding donor (HBD, purple).

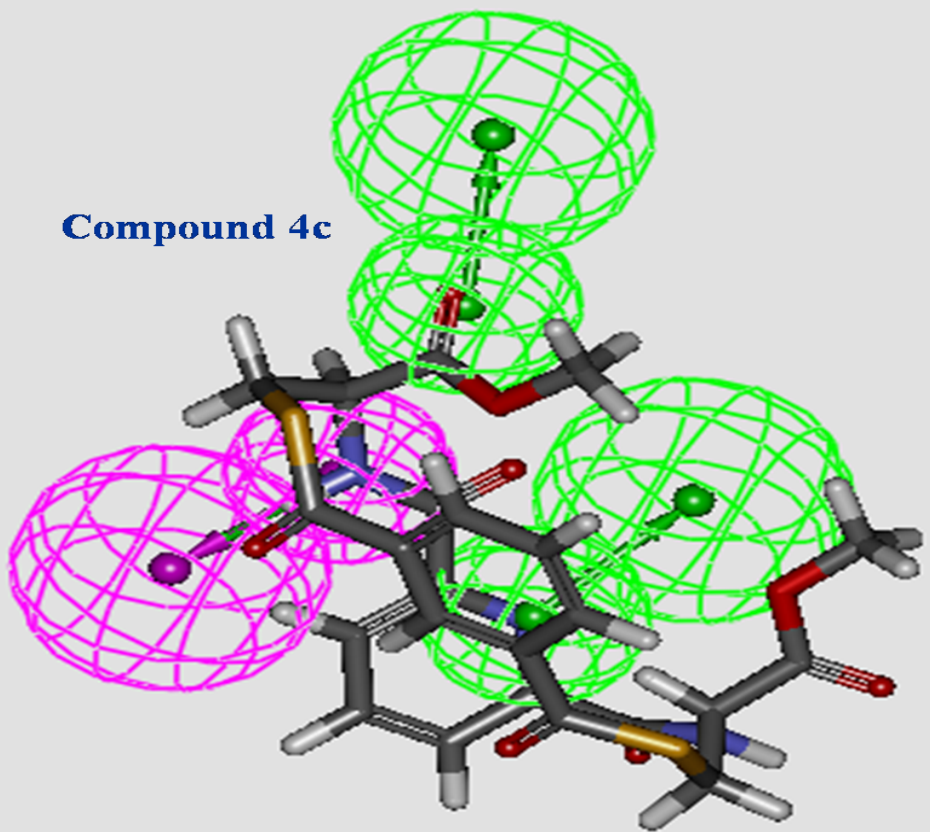
Compound 4a



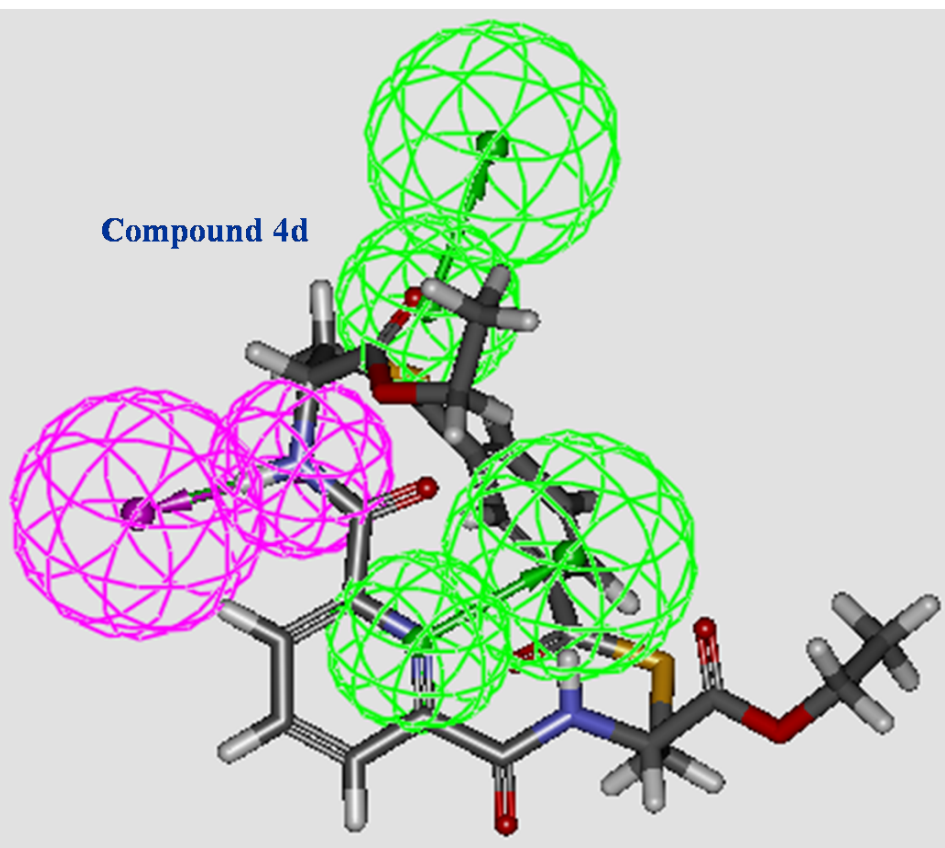
Compound 4b



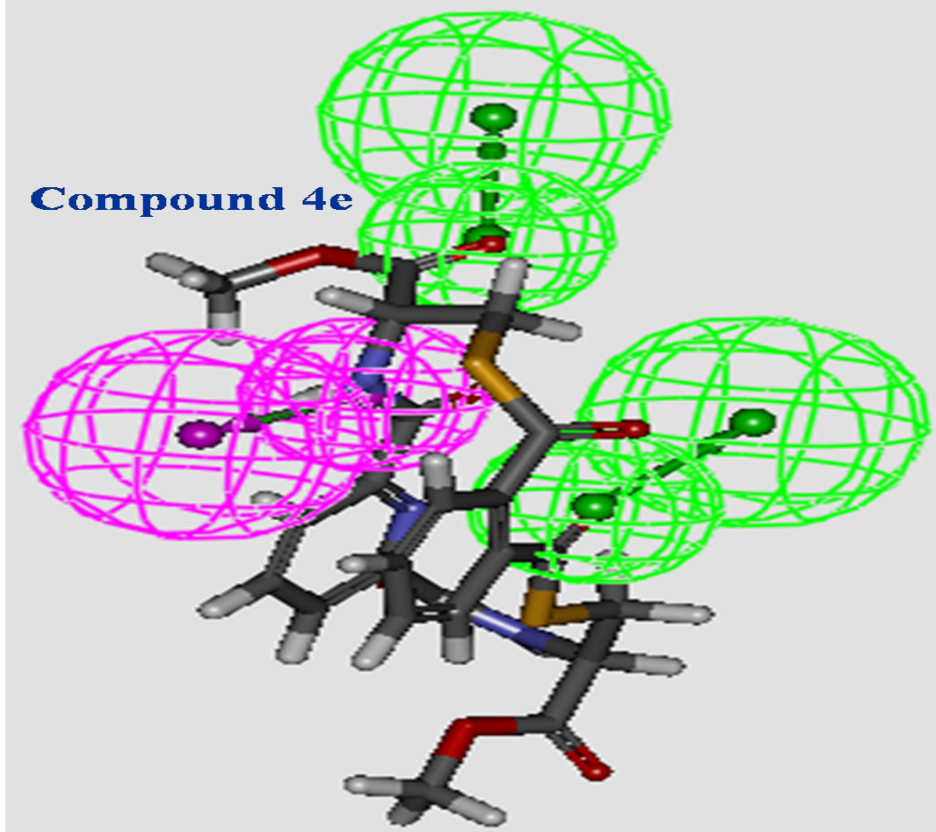
Compound 4c



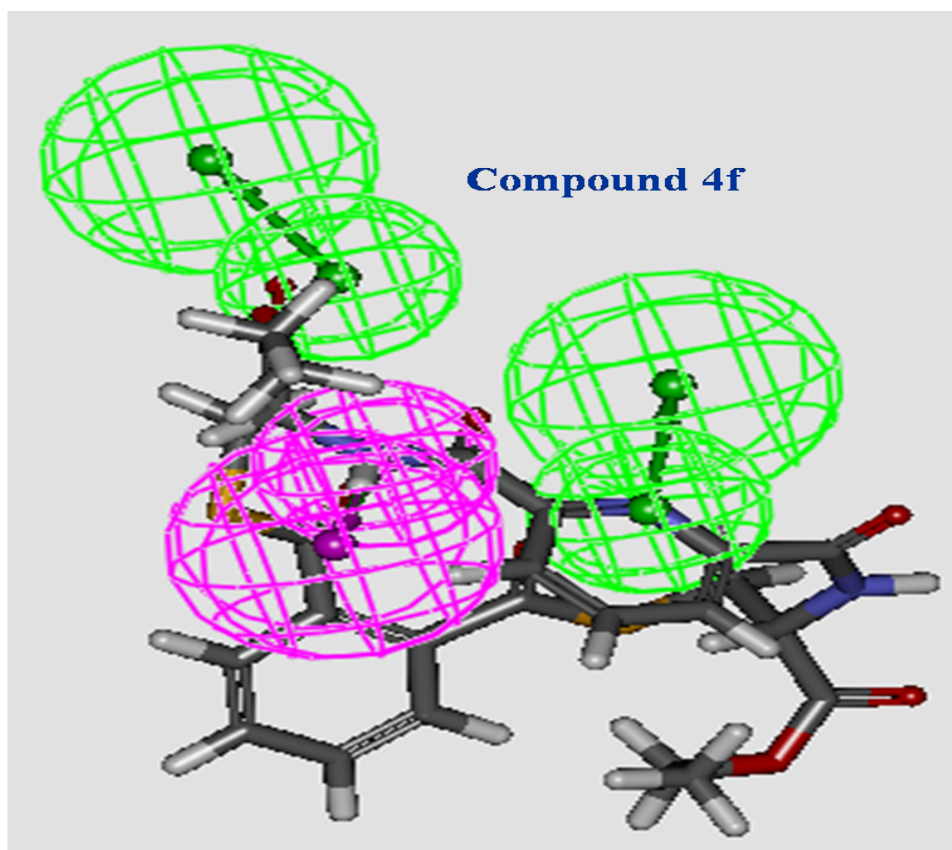
Compound 4d

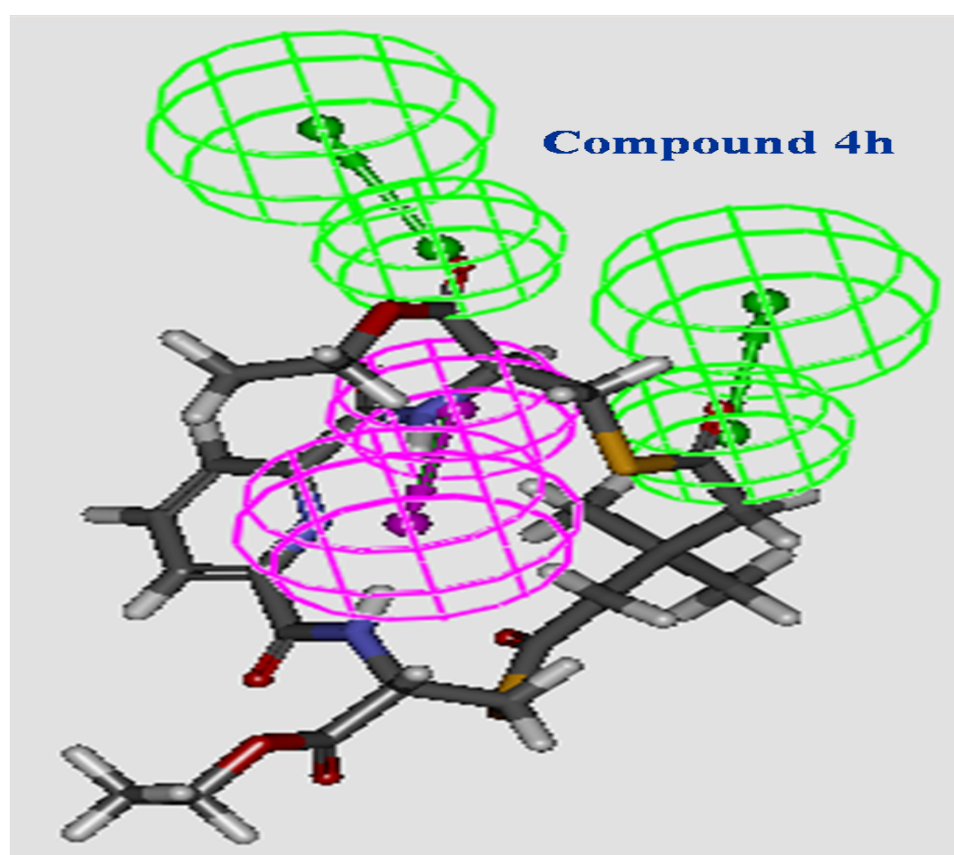
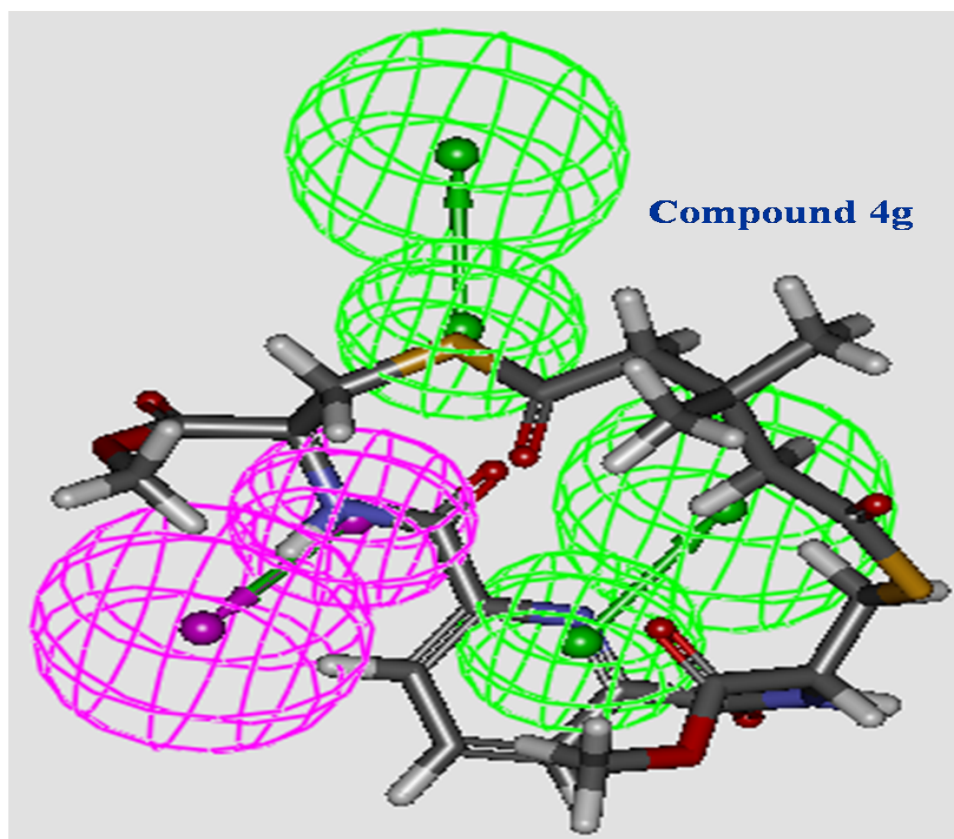


Compound 4e

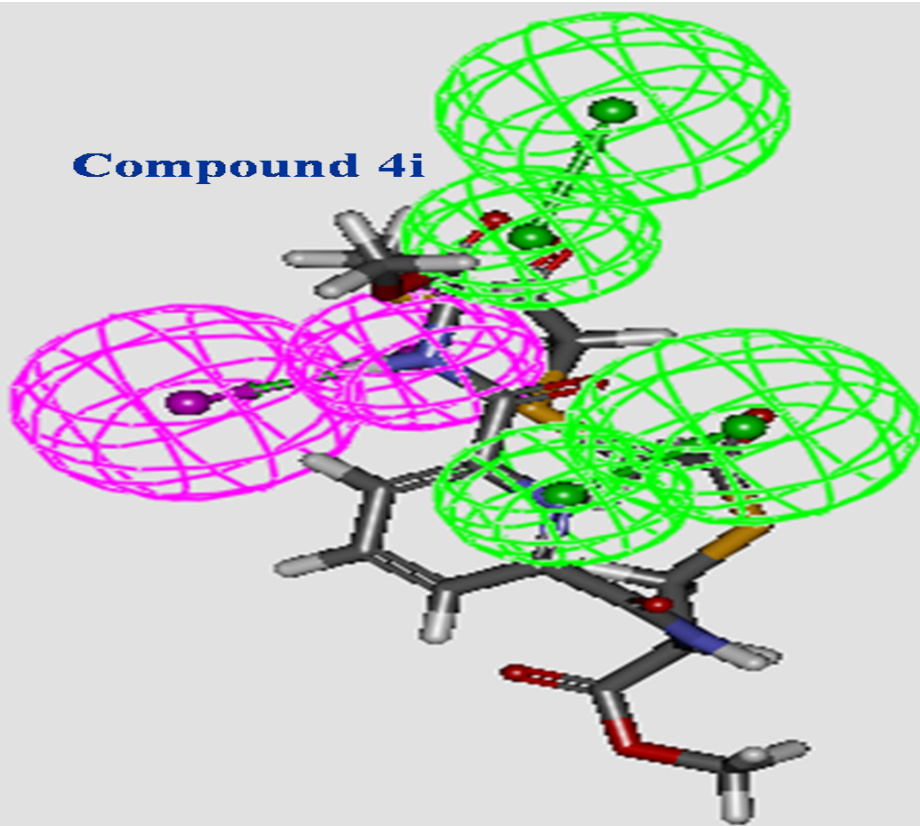


Compound 4f

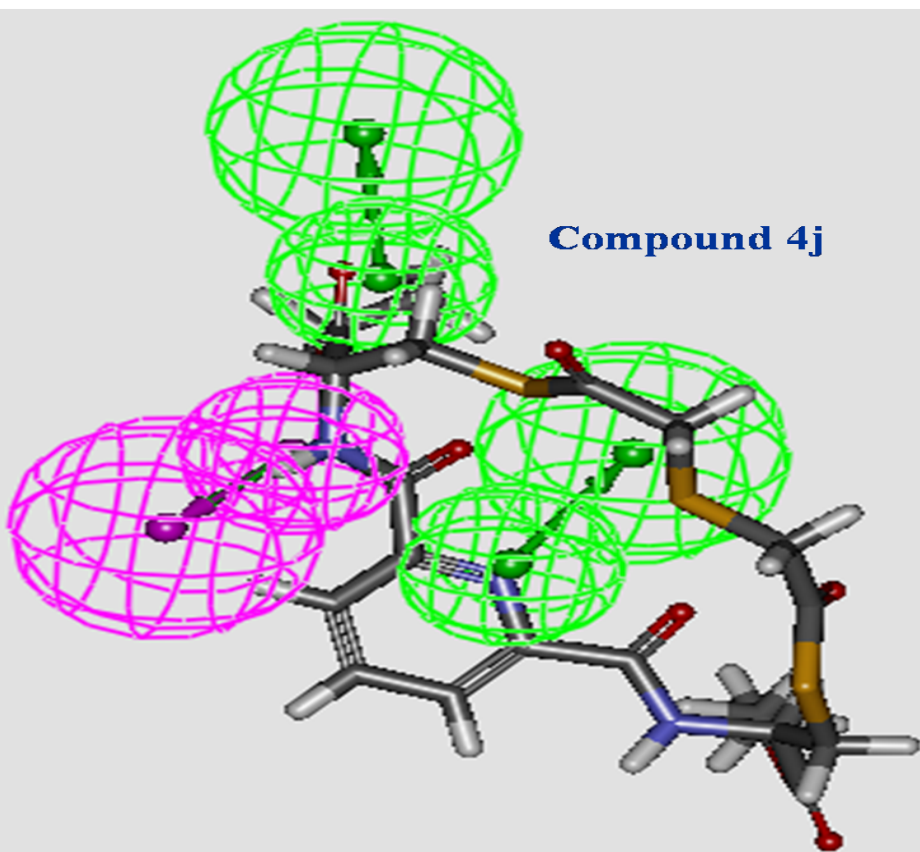




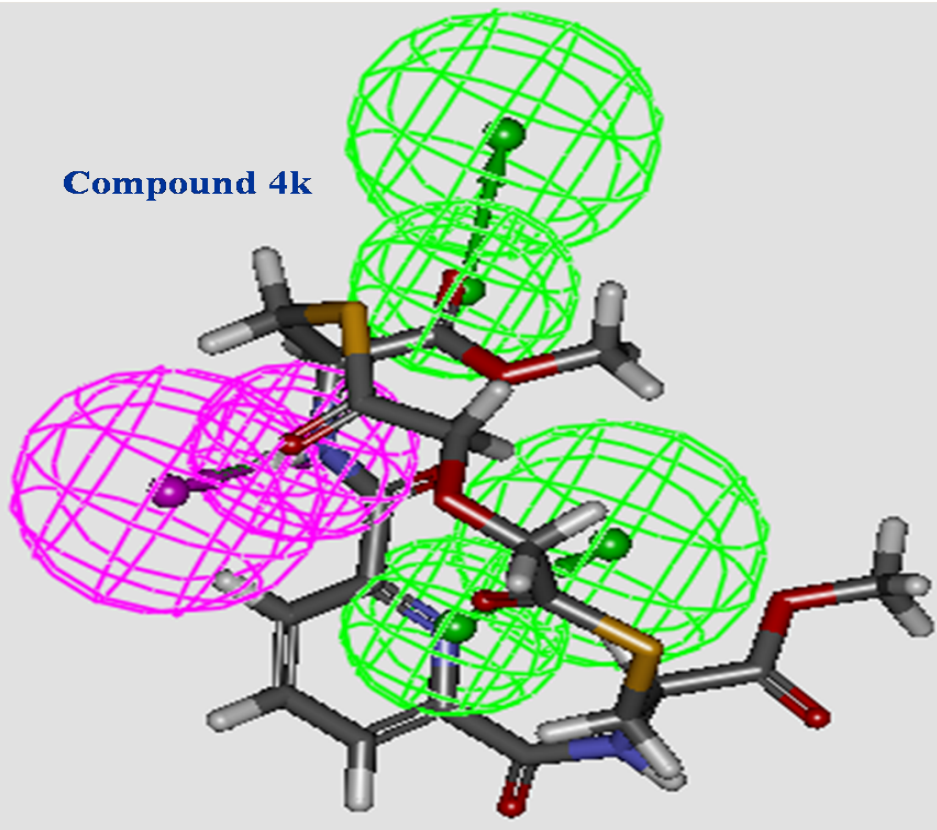
Compound 4i



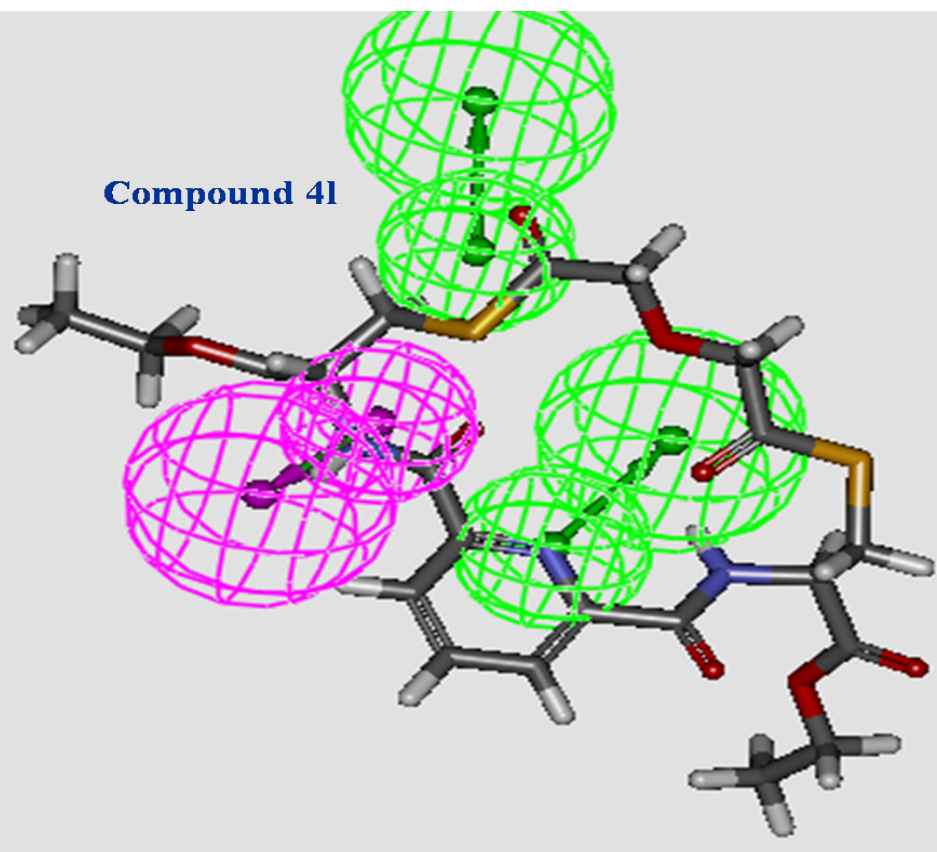
Compound 4j



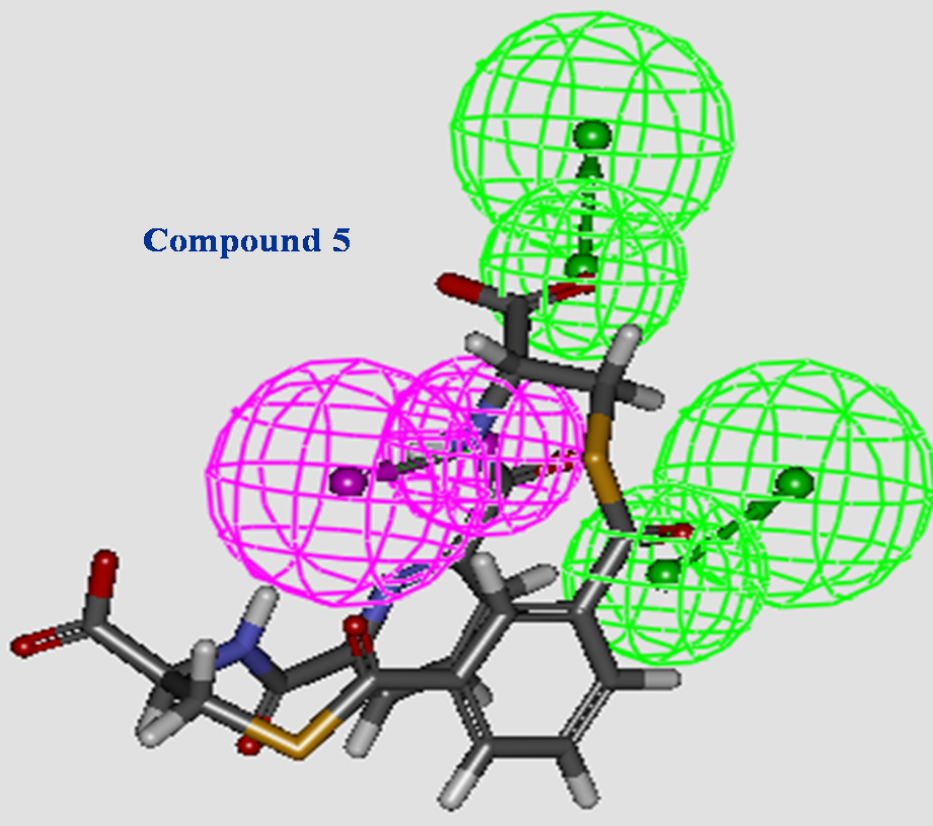
Compound 4k



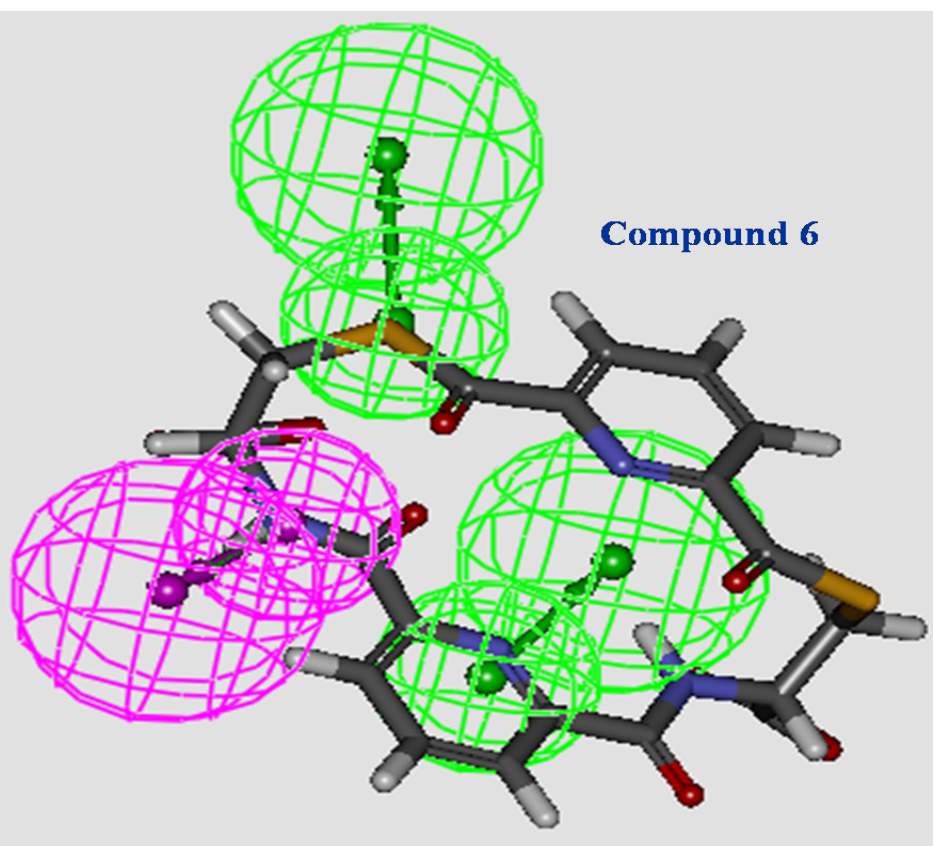
Compound 4l



Compound 5



Compound 6



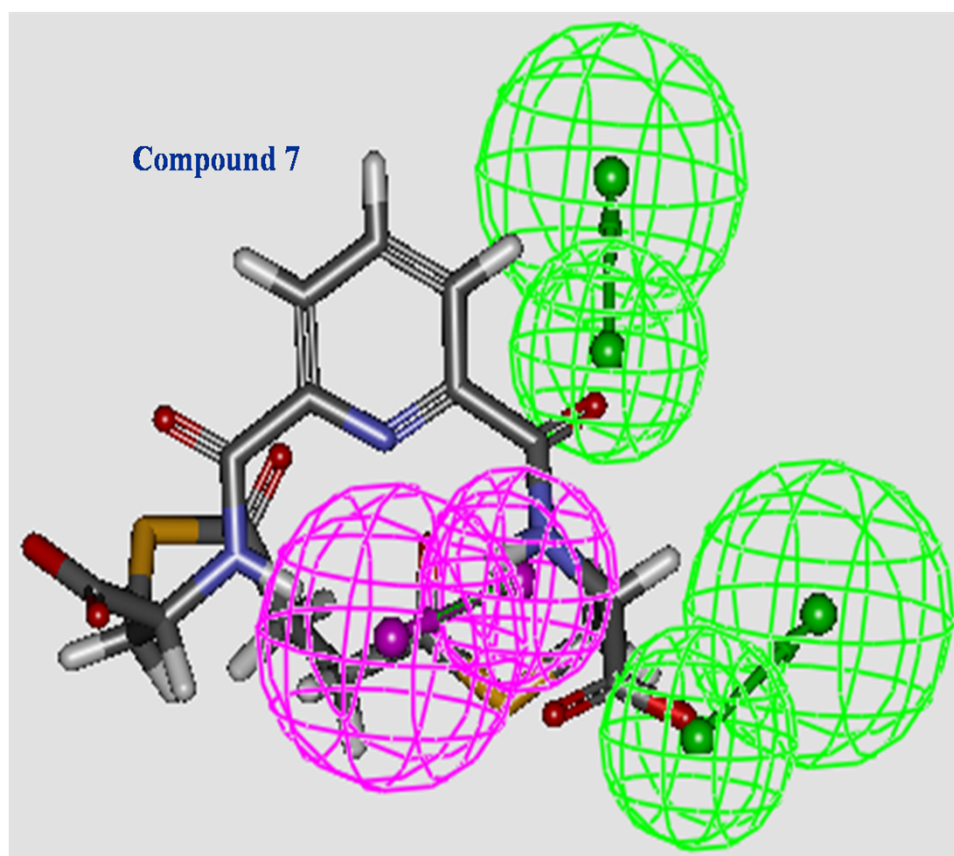
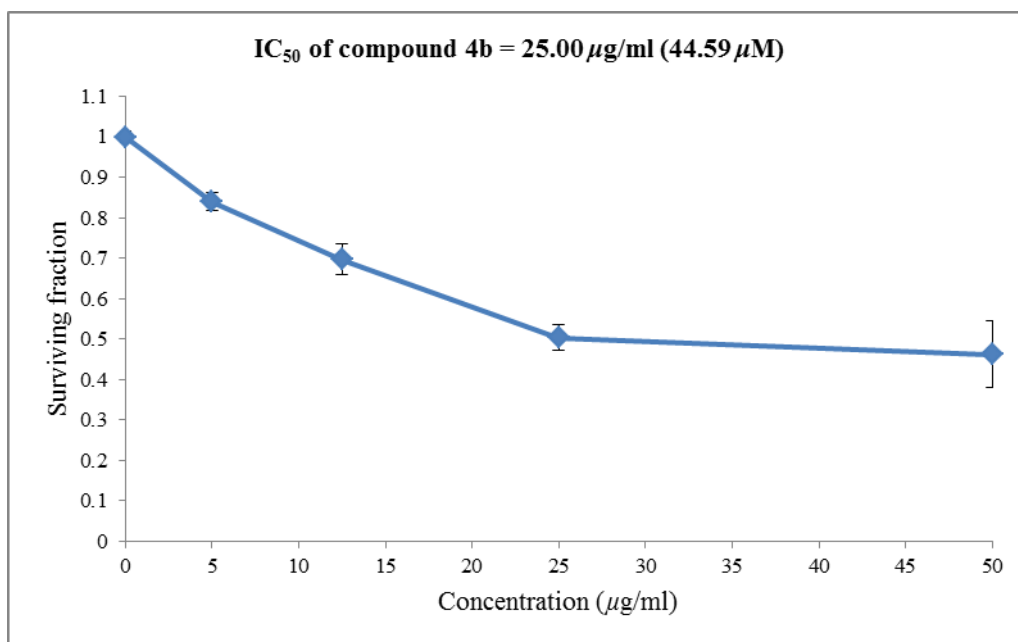
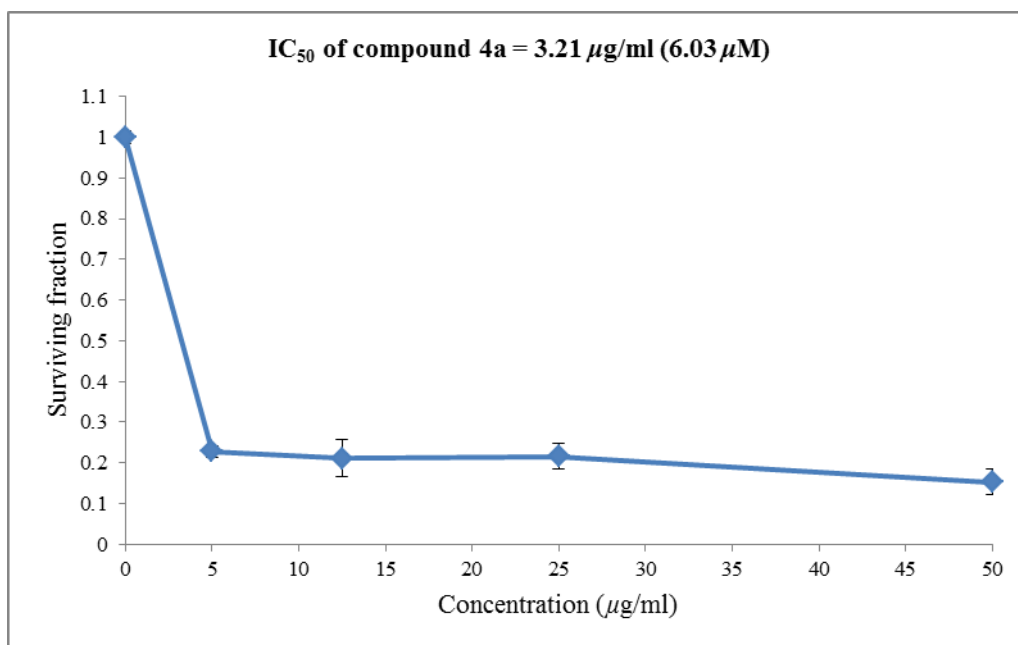
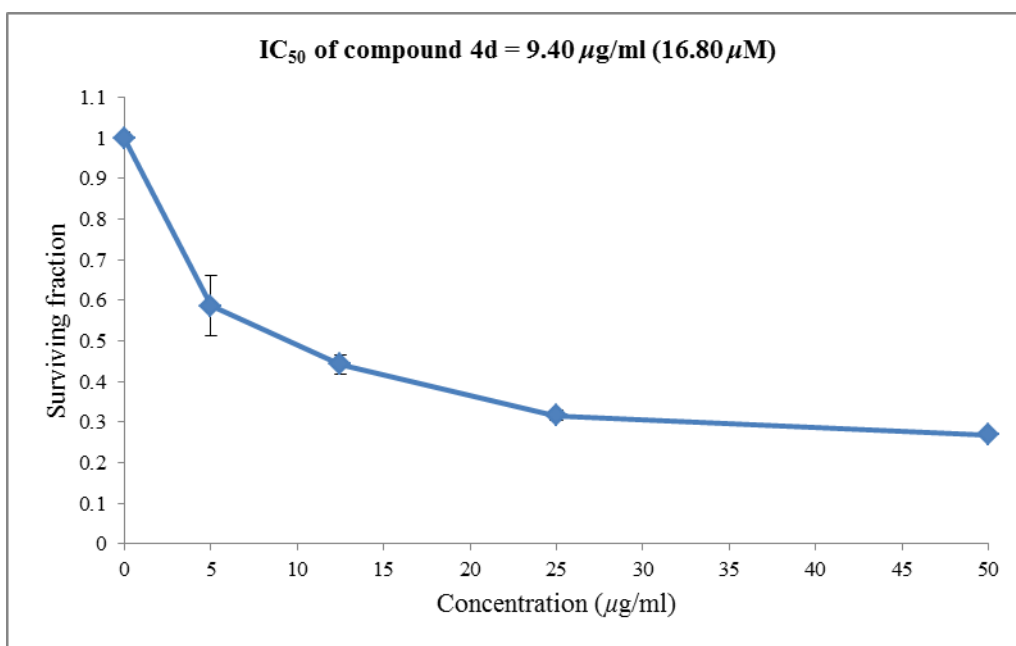
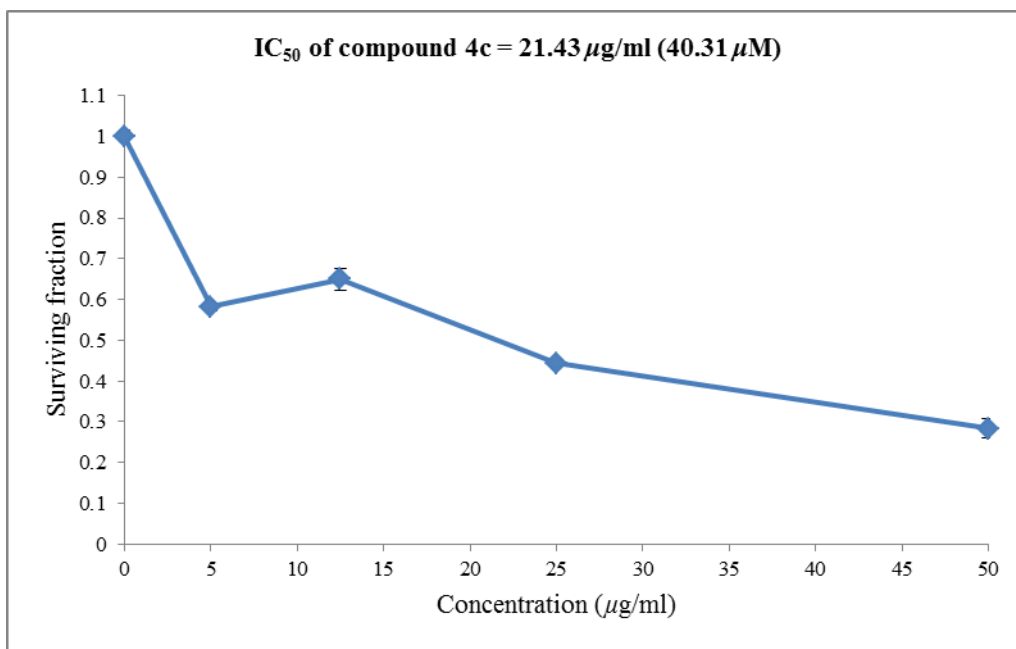
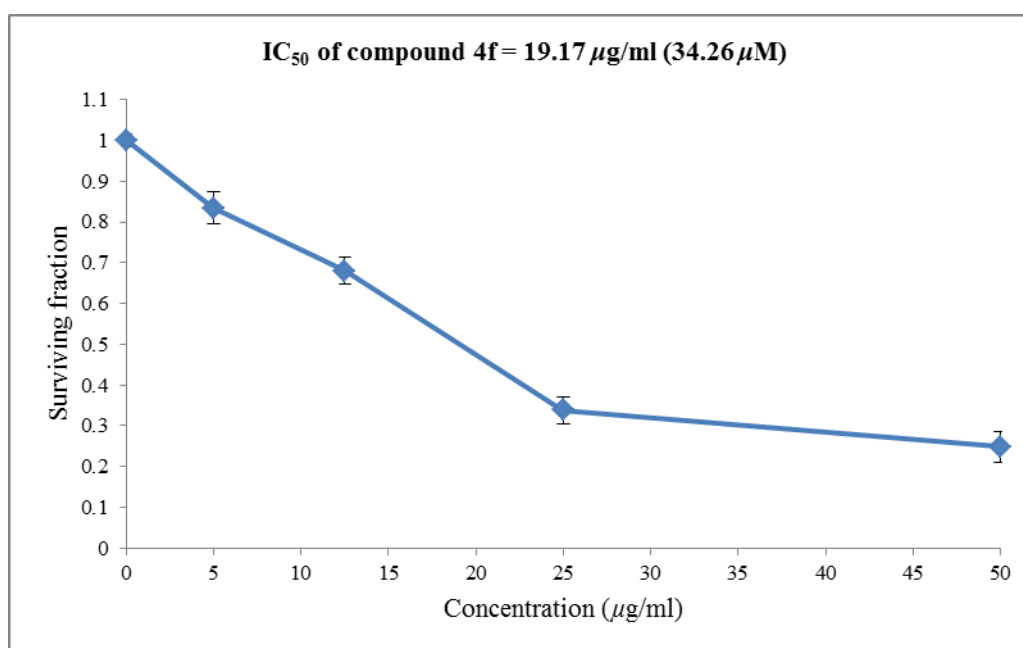
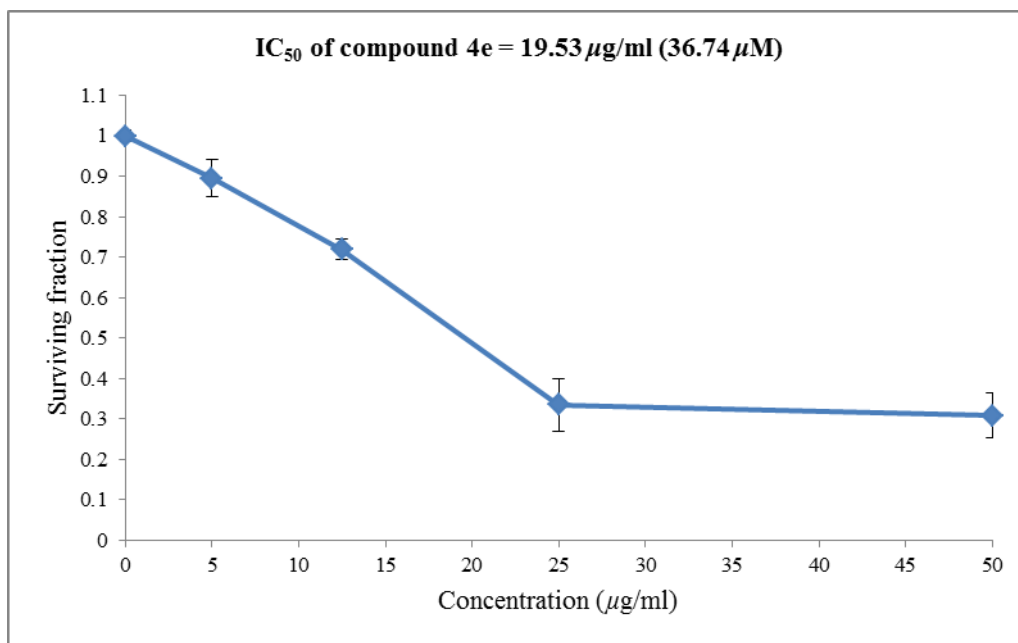
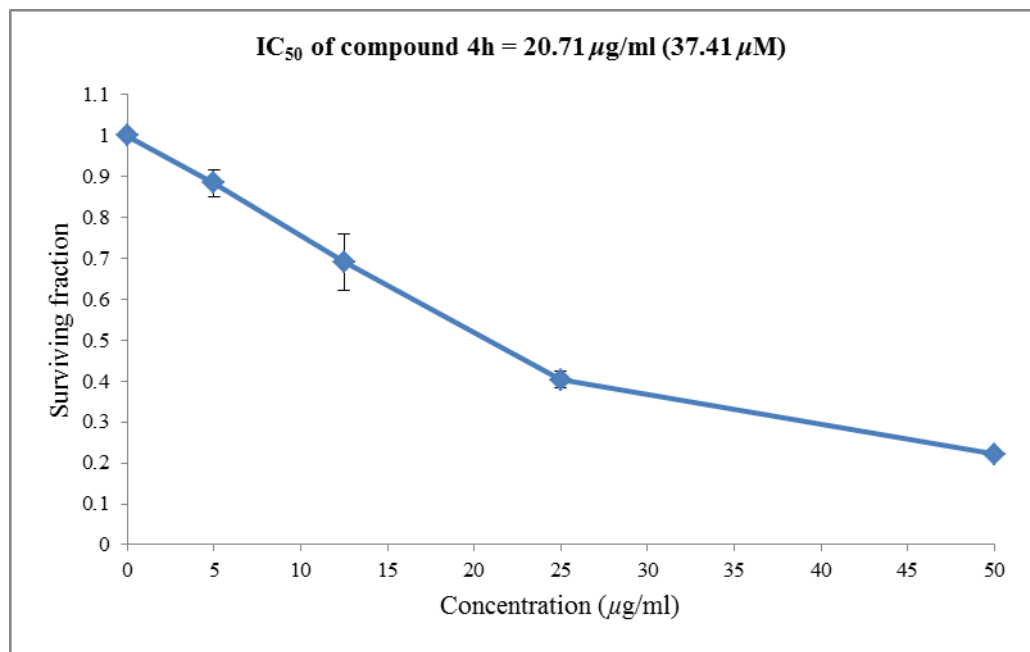
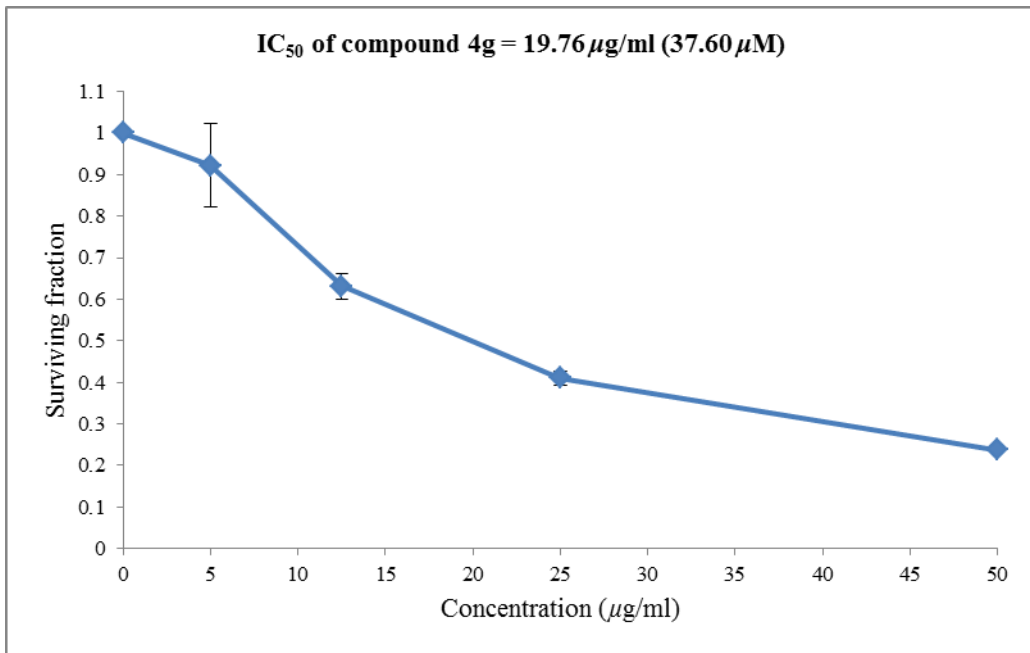


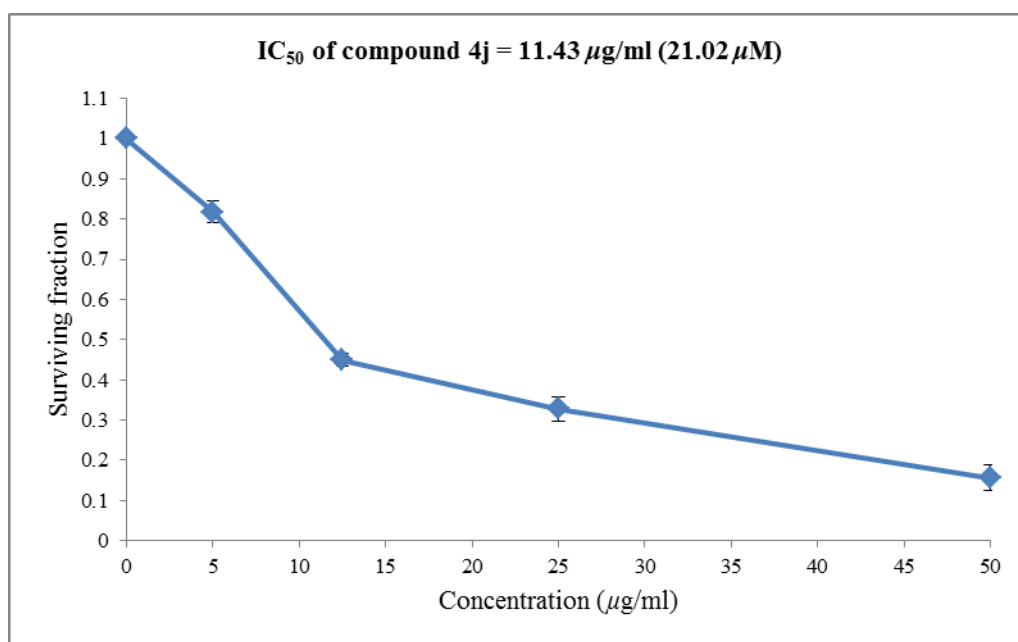
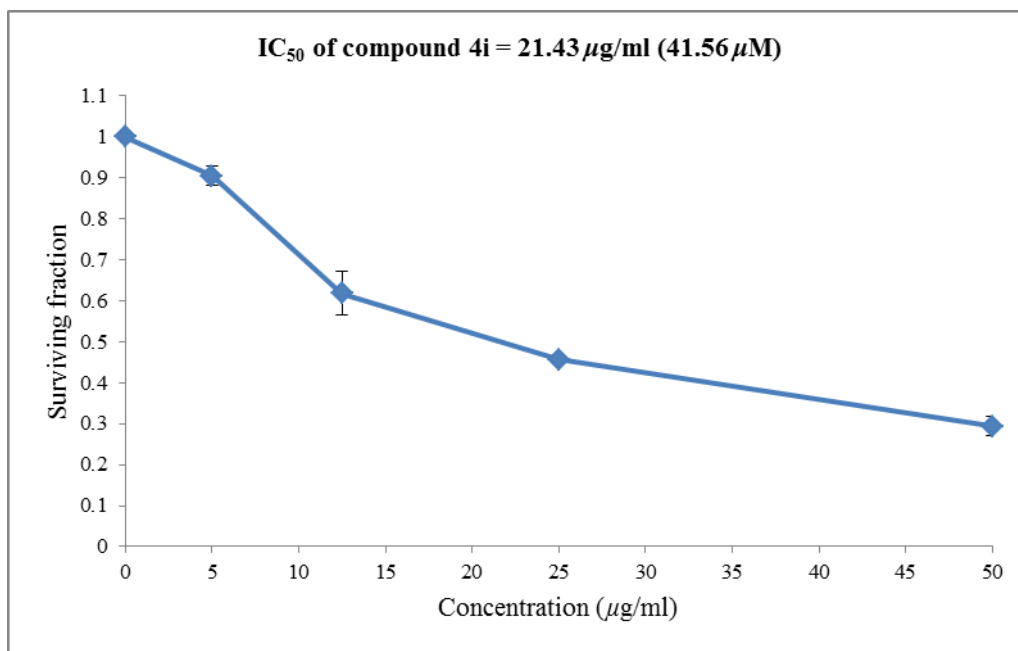
Figure S2. 3D-pharmacophore mapped on the antifungal macrocyclic peptidomimetics **4a-4l**, and **5-7**.

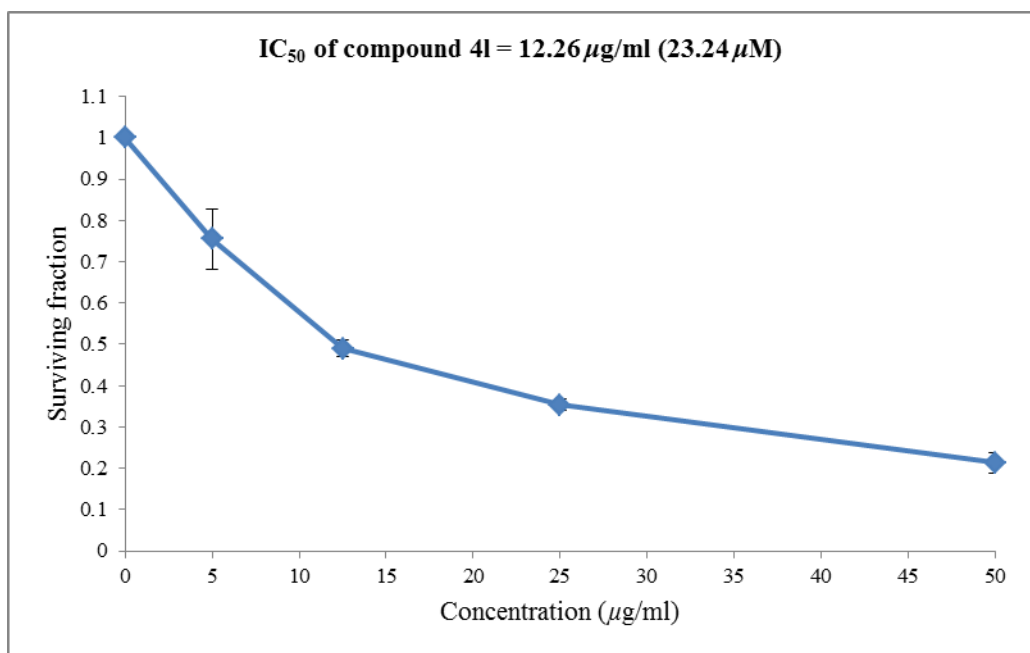
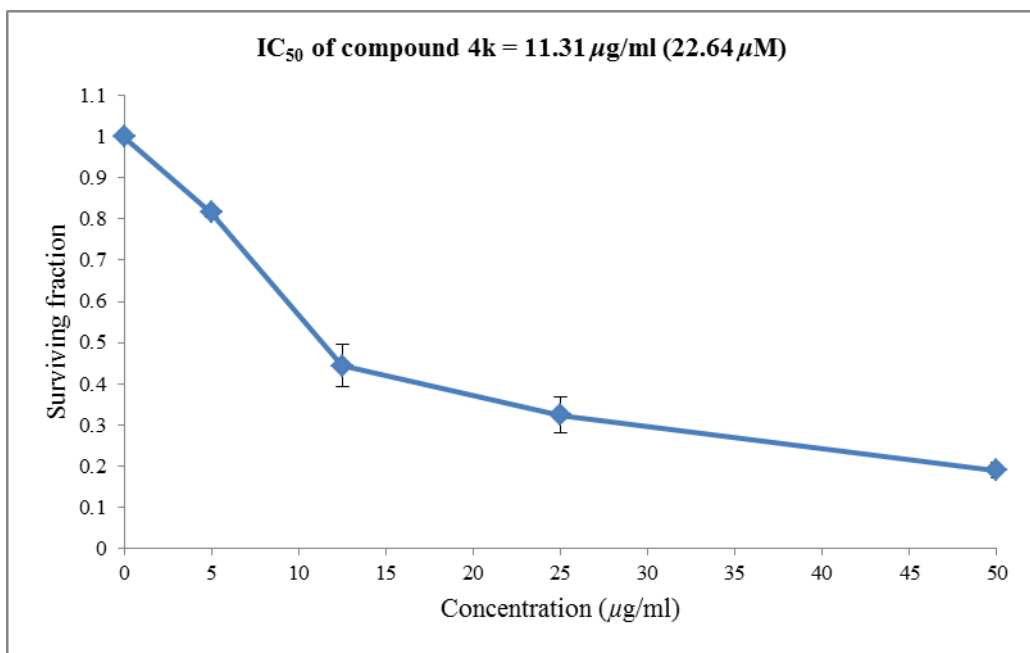


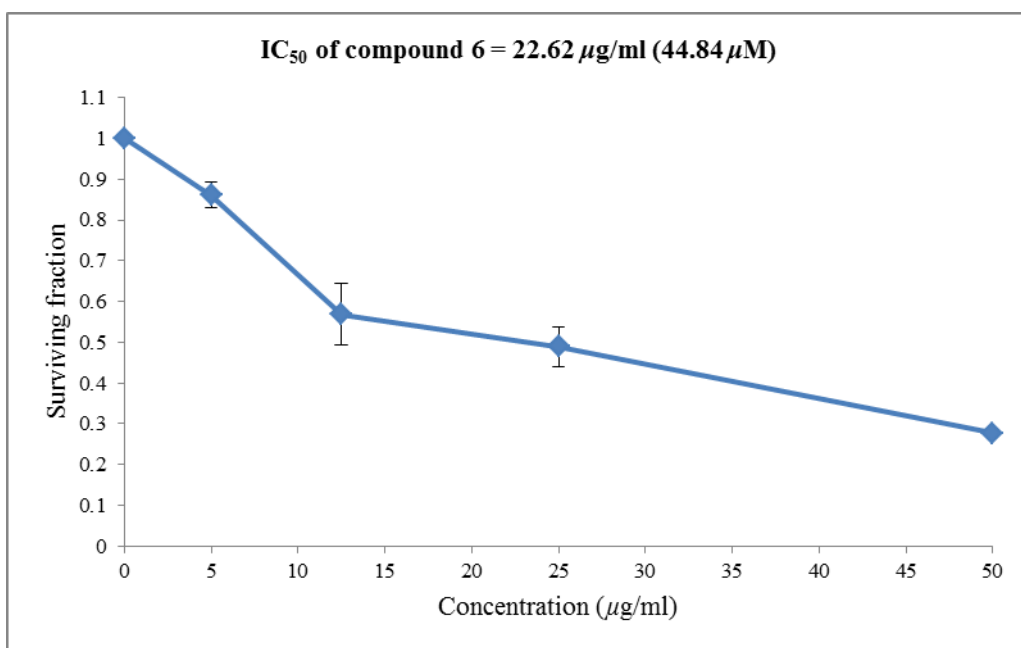
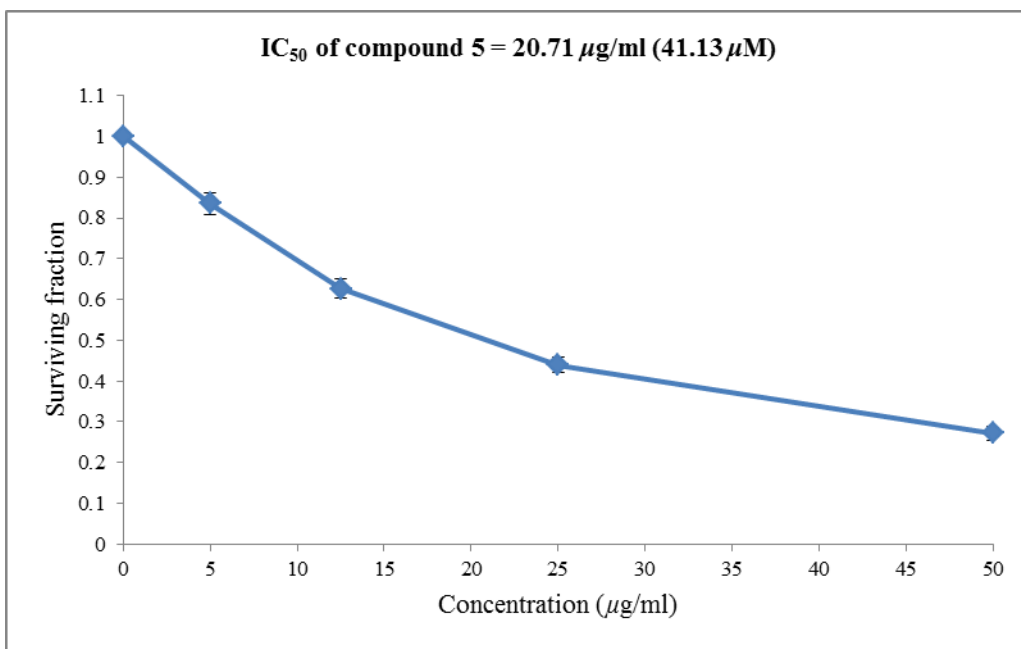












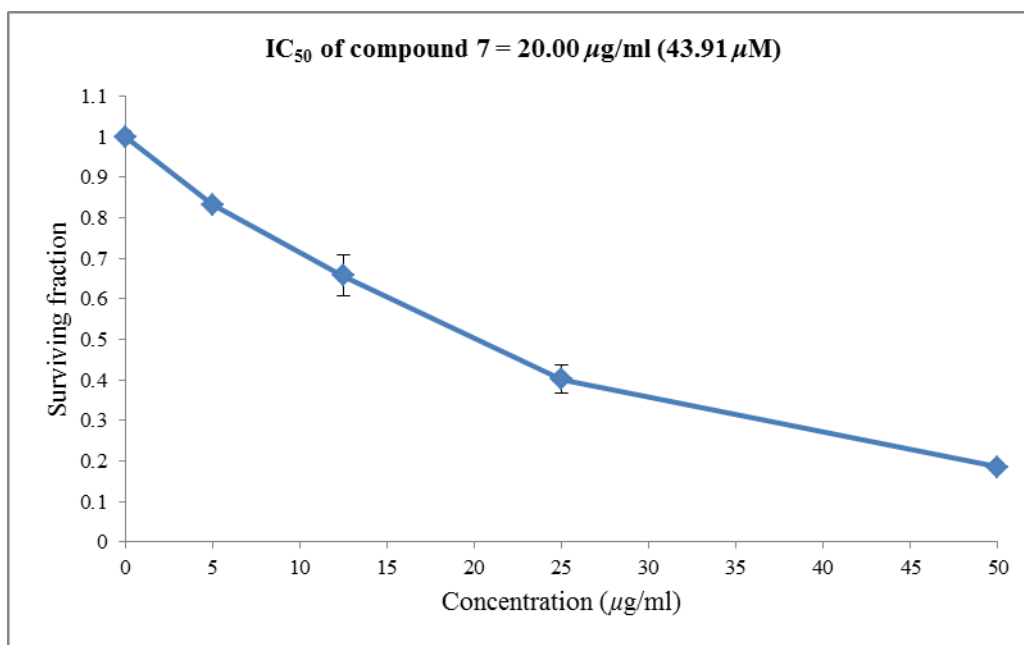
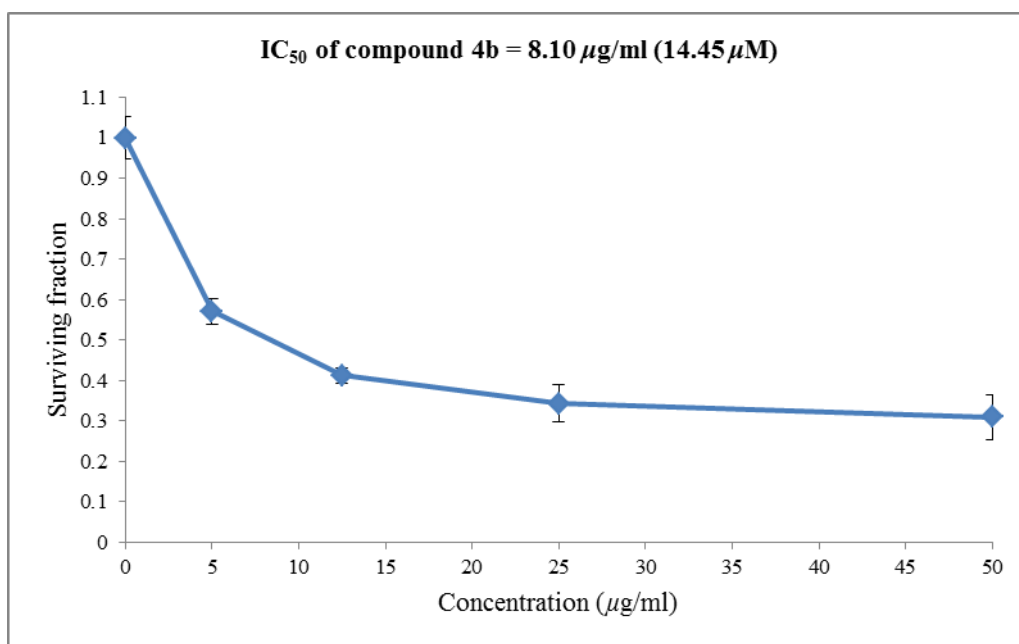
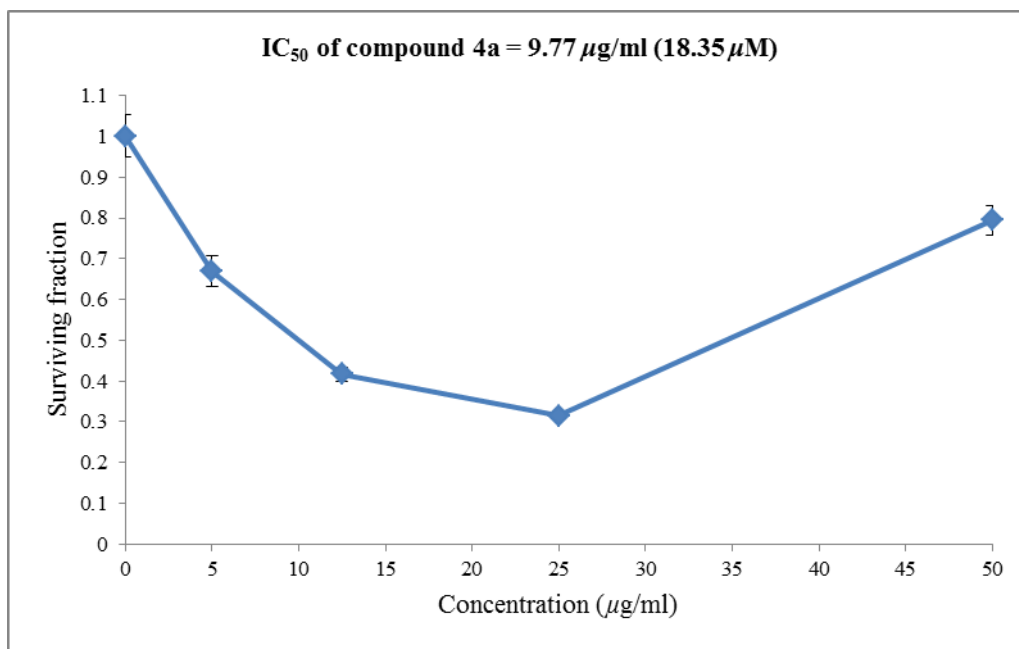
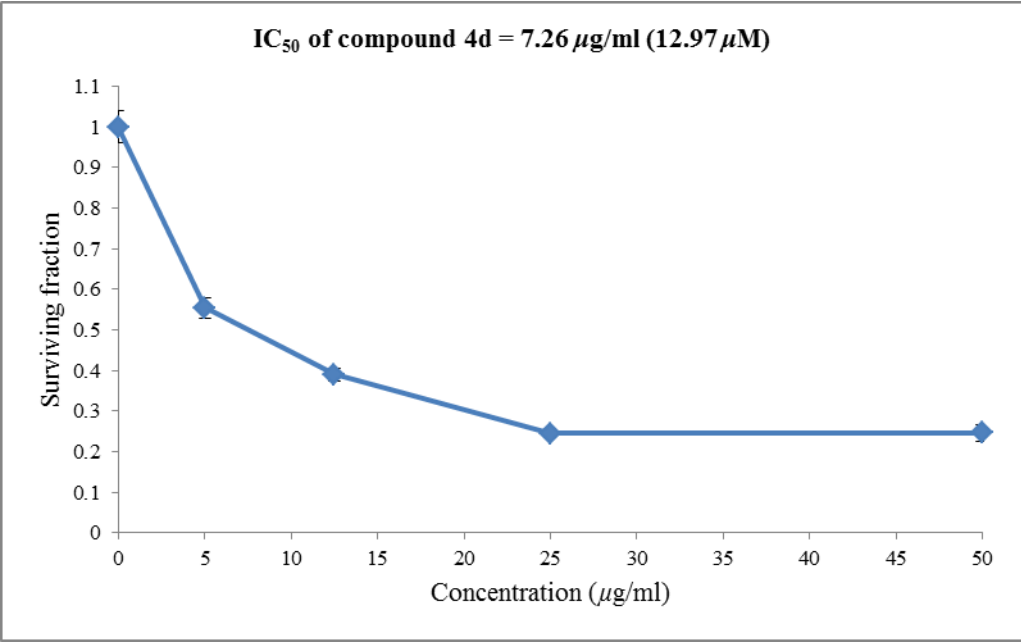
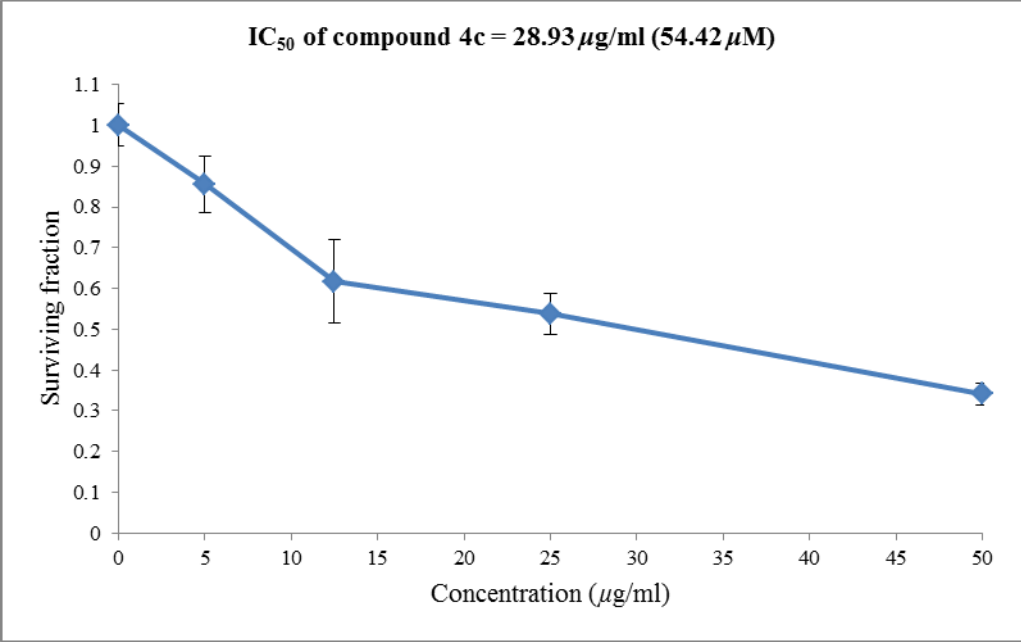
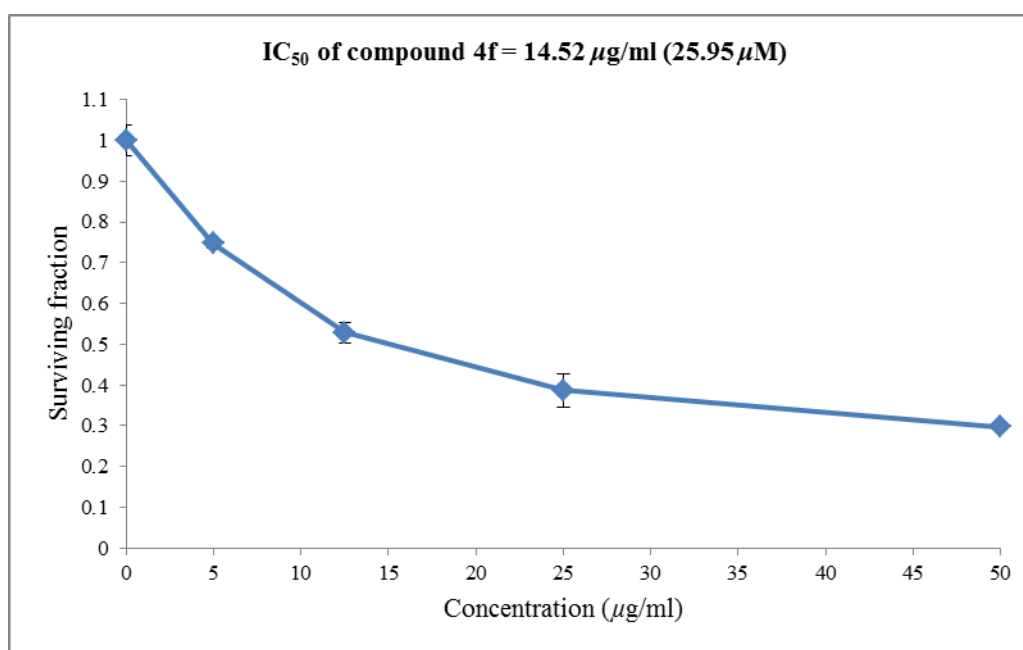
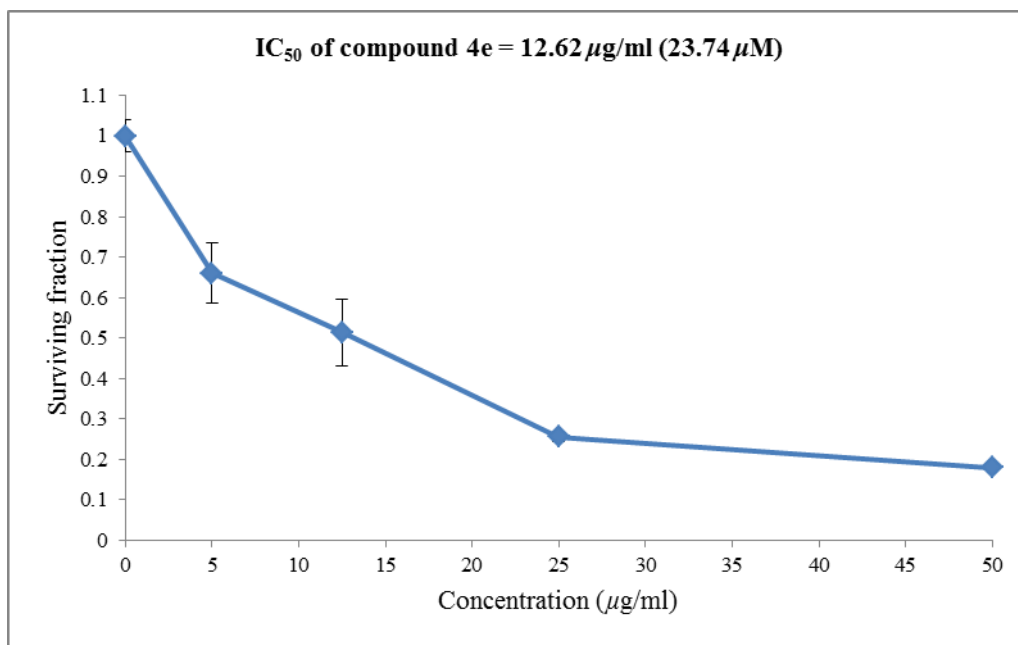
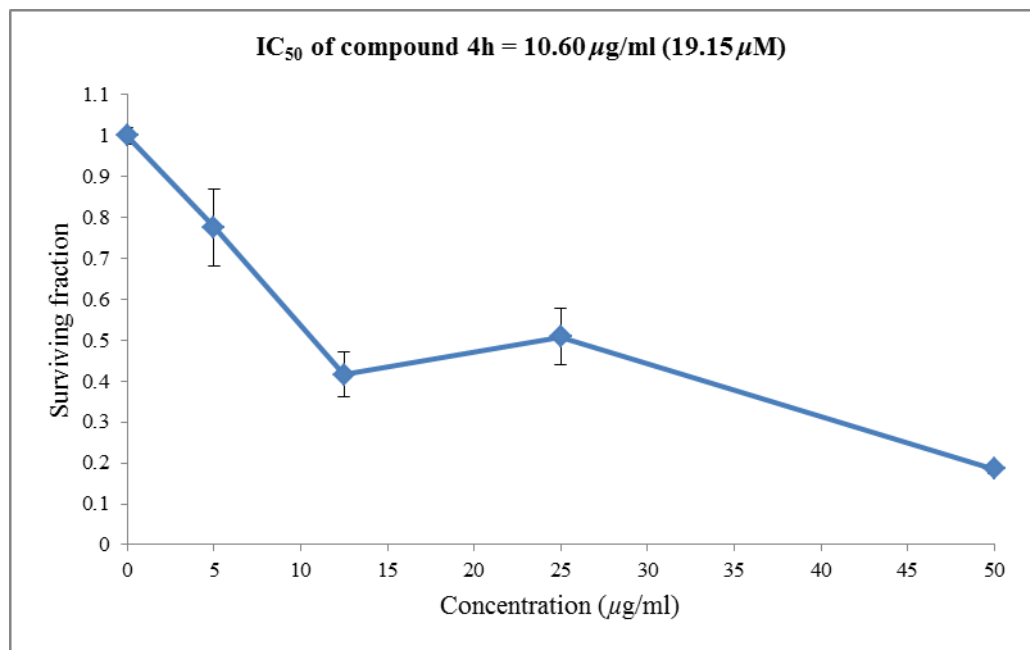
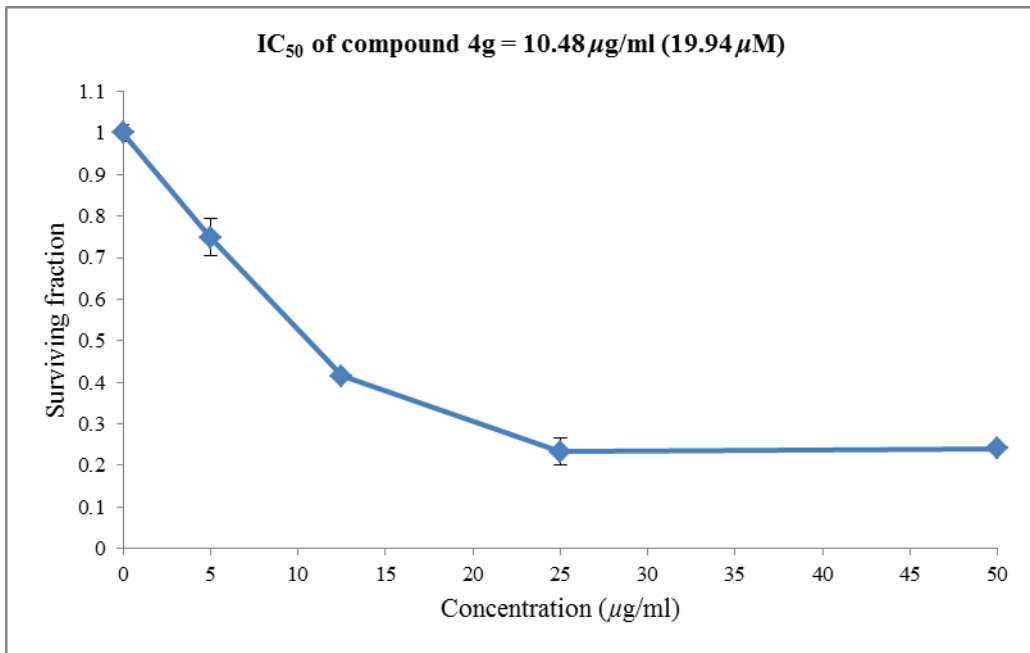


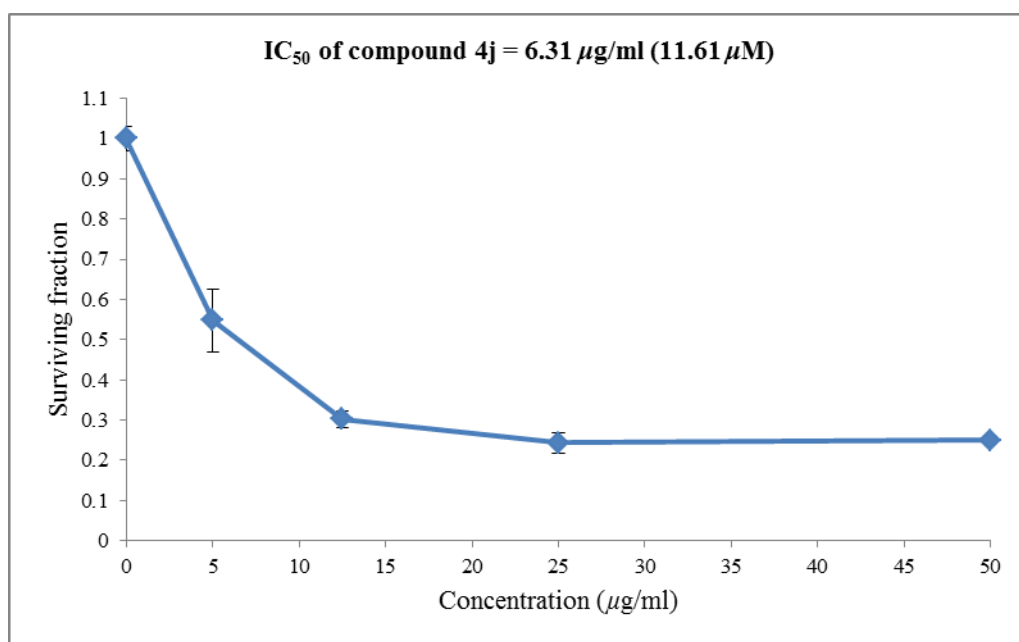
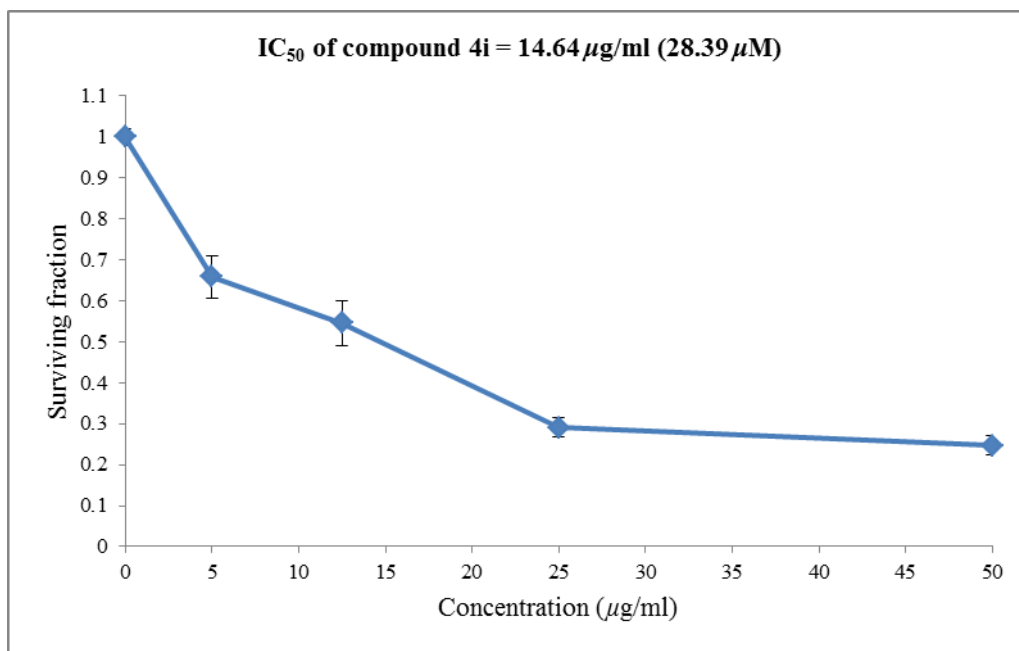
Figure S3. Dose-response curves of the macrocyclic peptidomimetics **4a-1**, and **5-7** against HepG2 (liver) human tumor cell line.

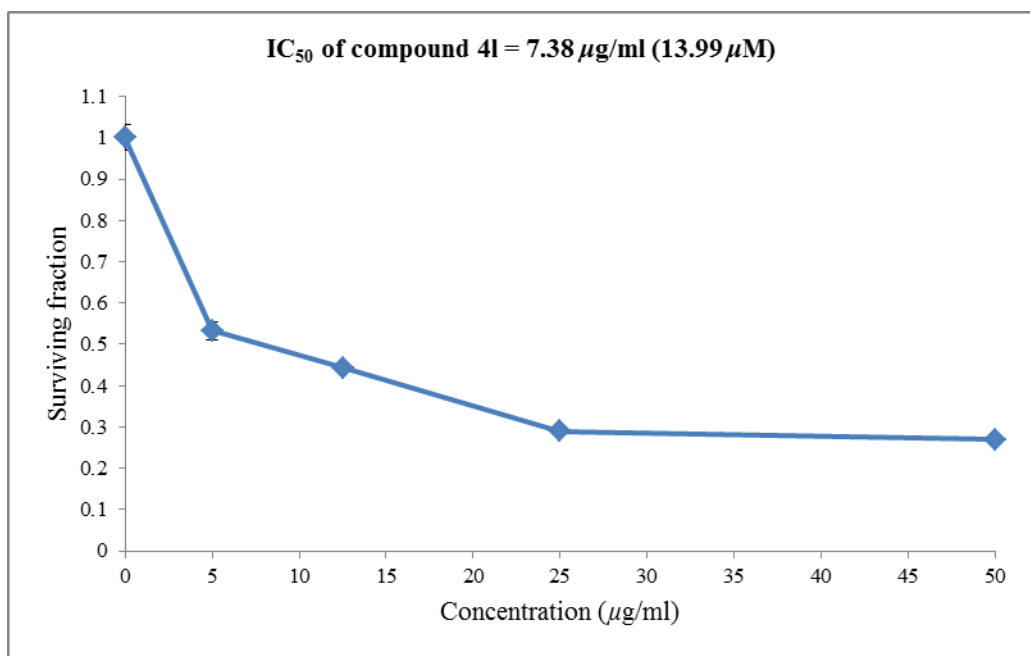
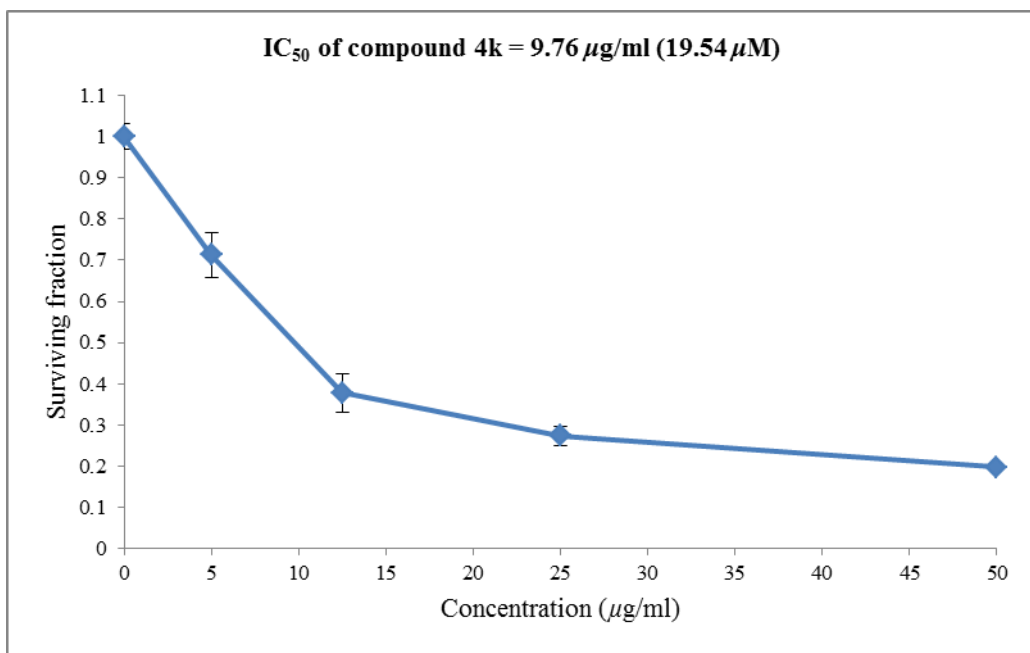


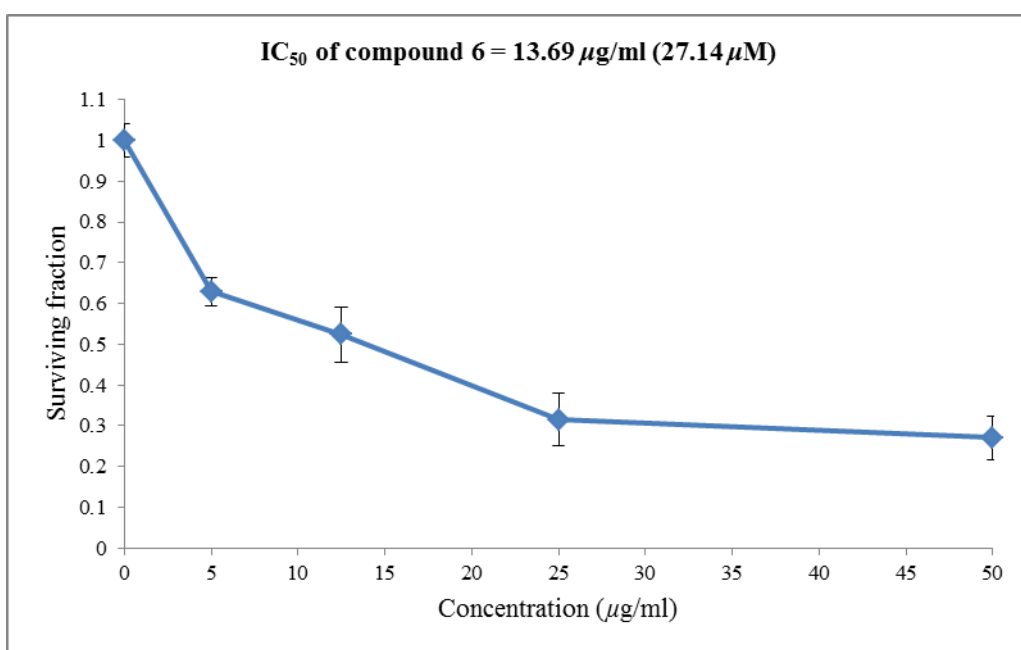
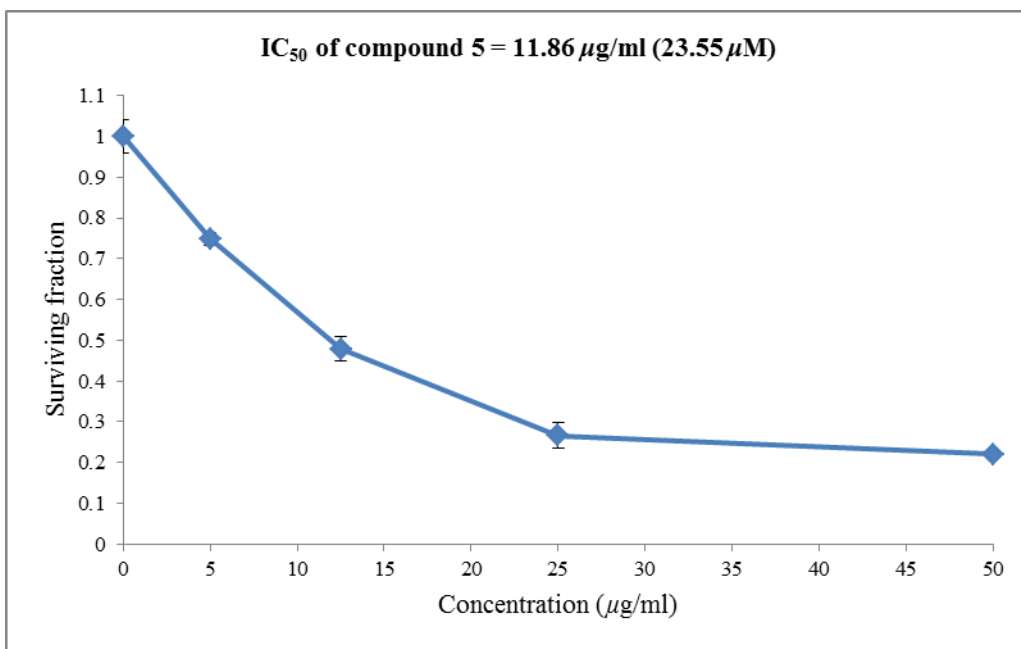












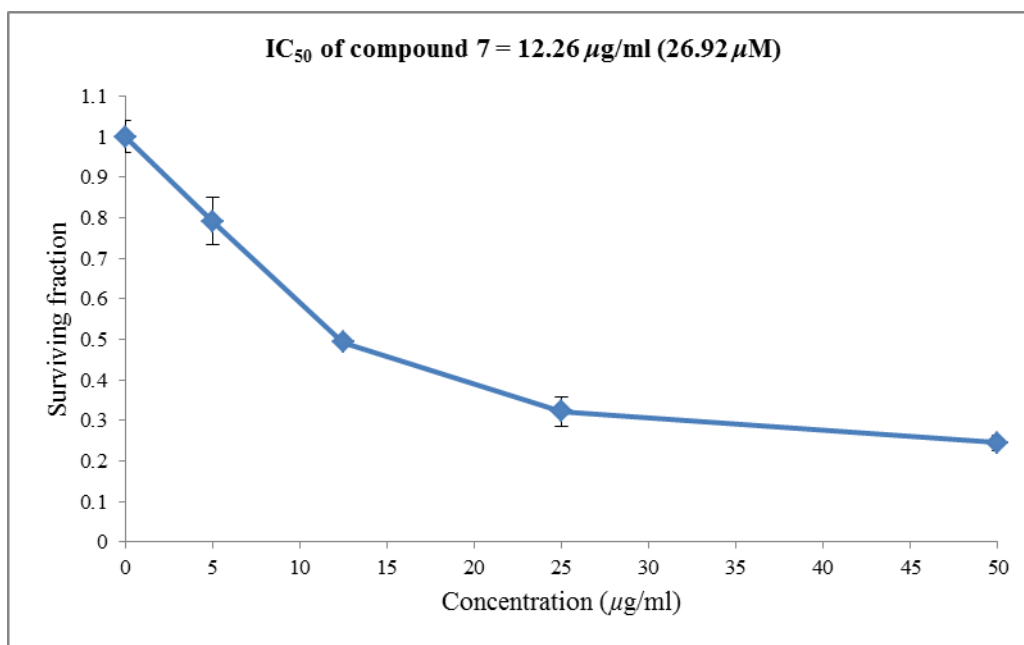
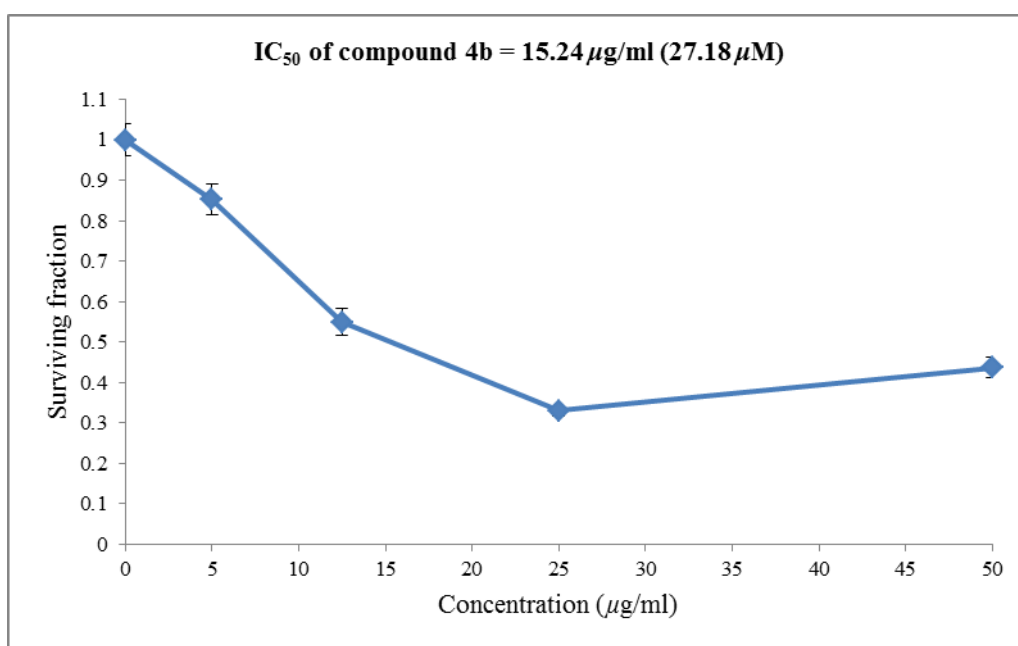
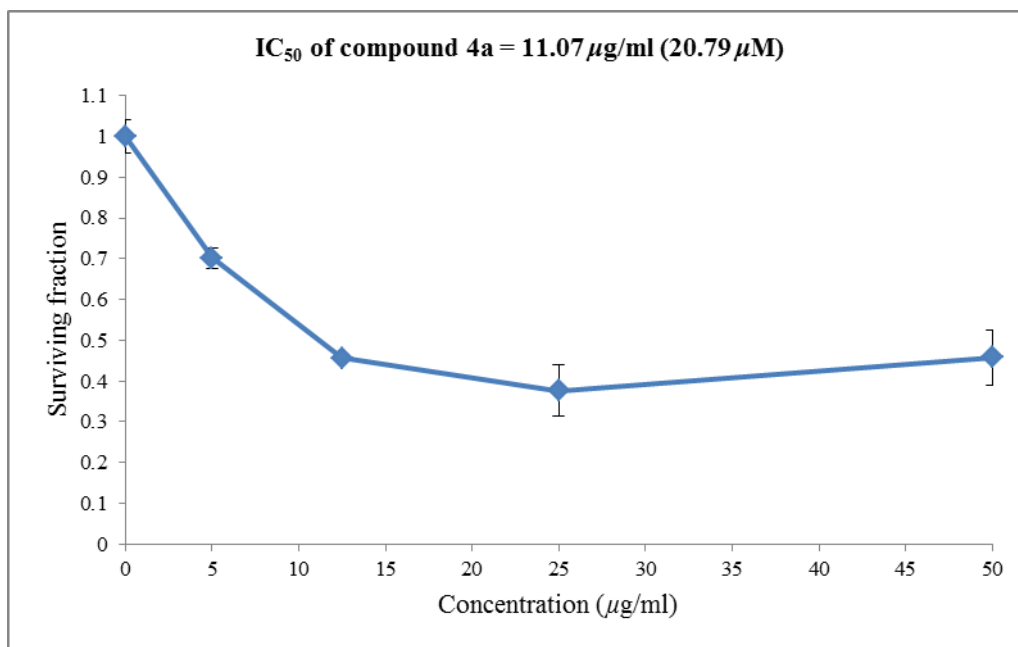
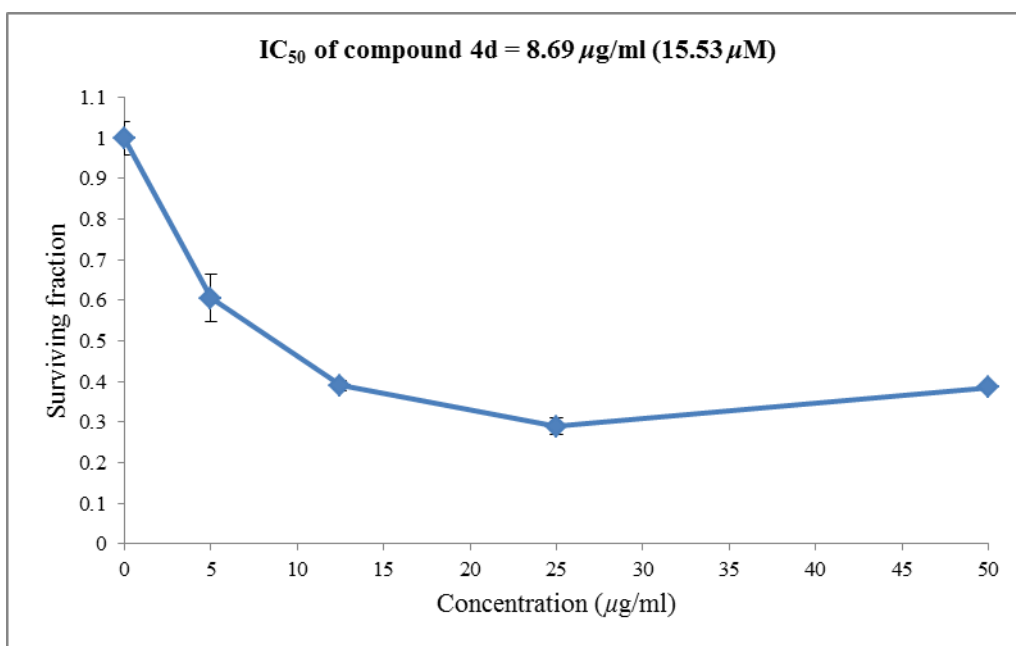
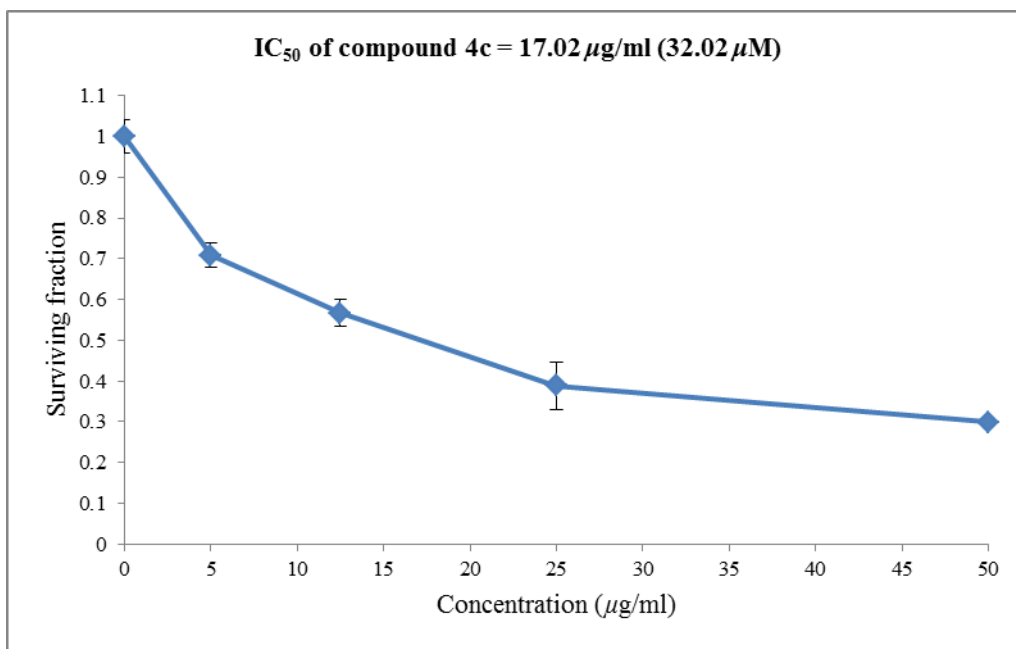
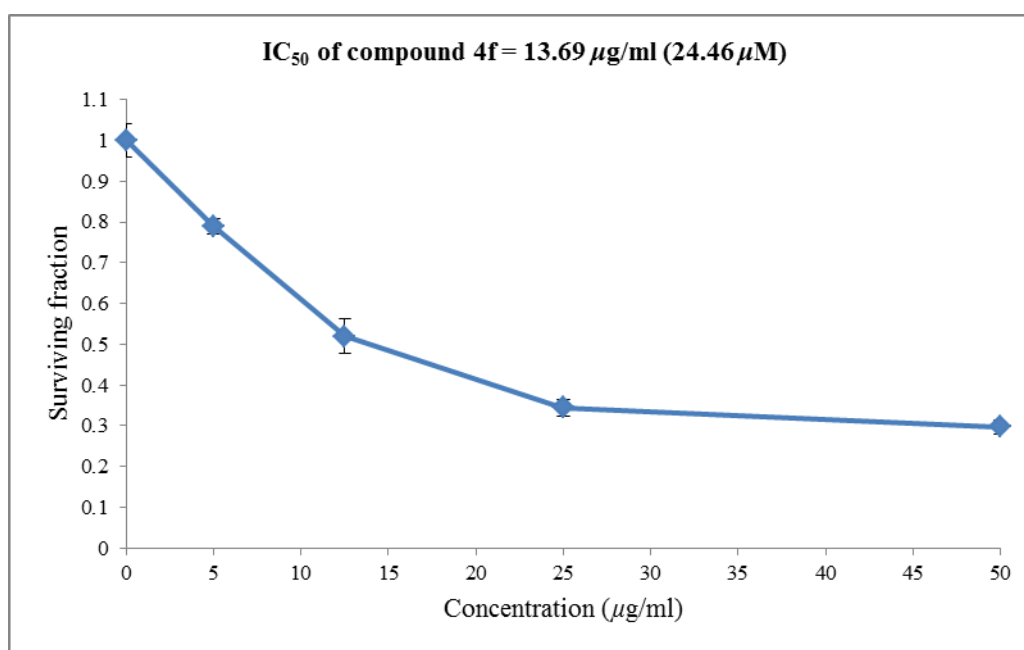
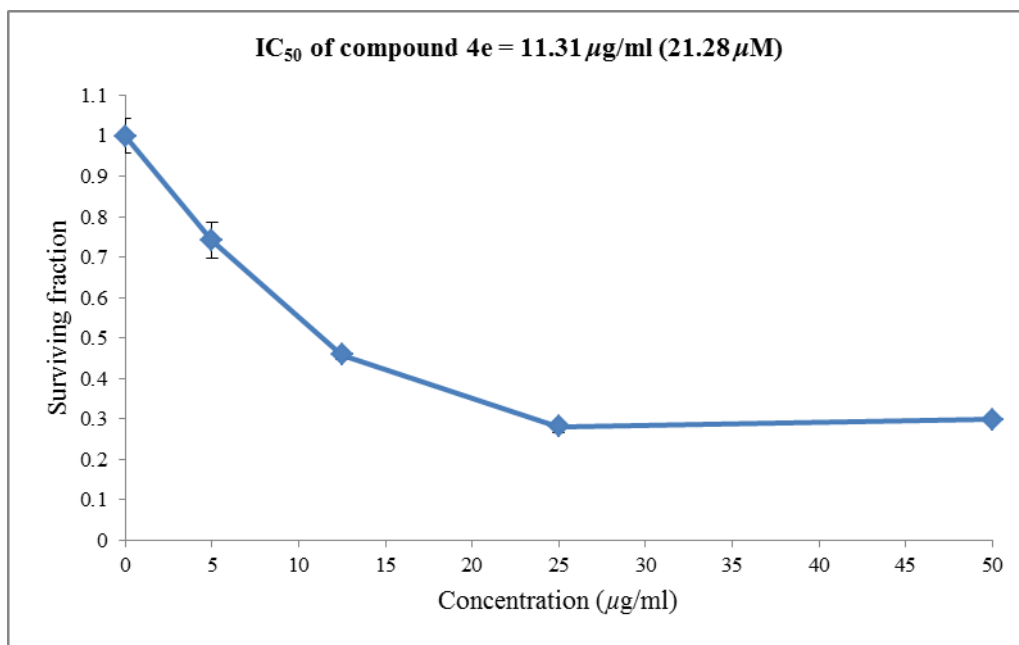
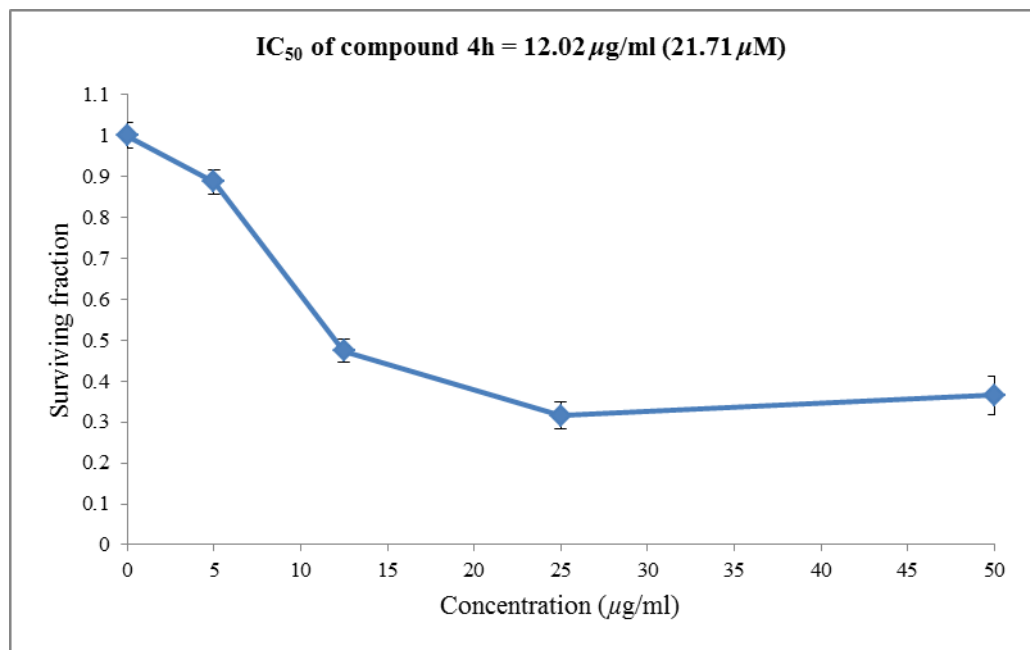
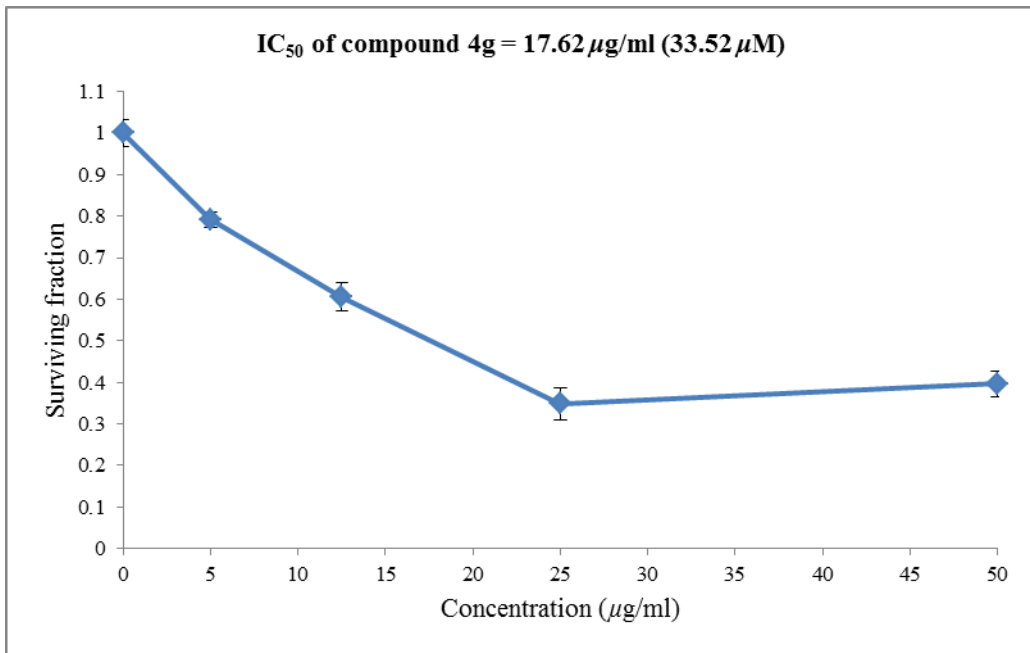


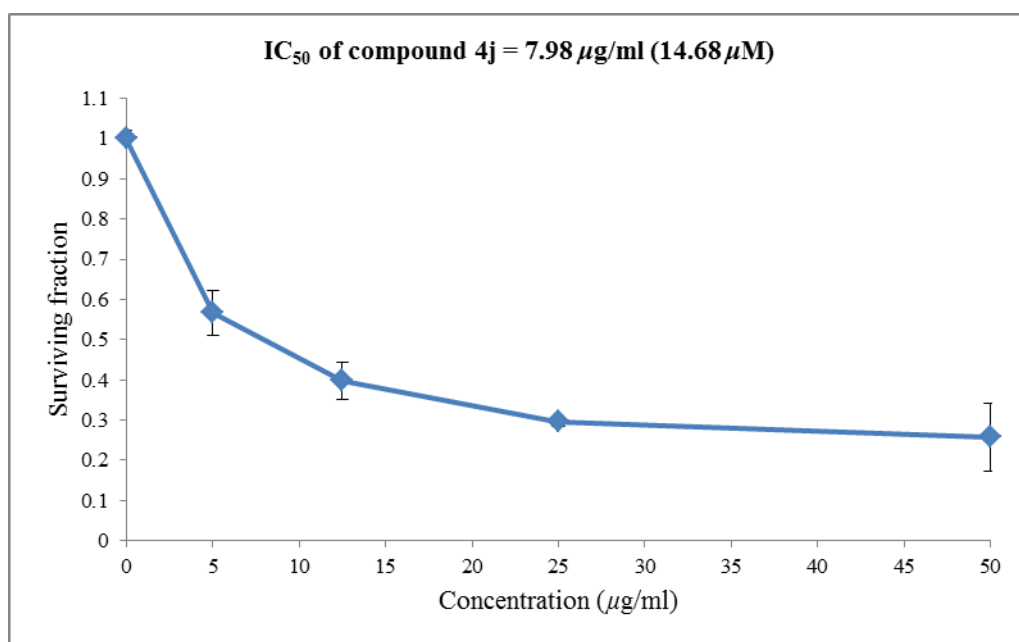
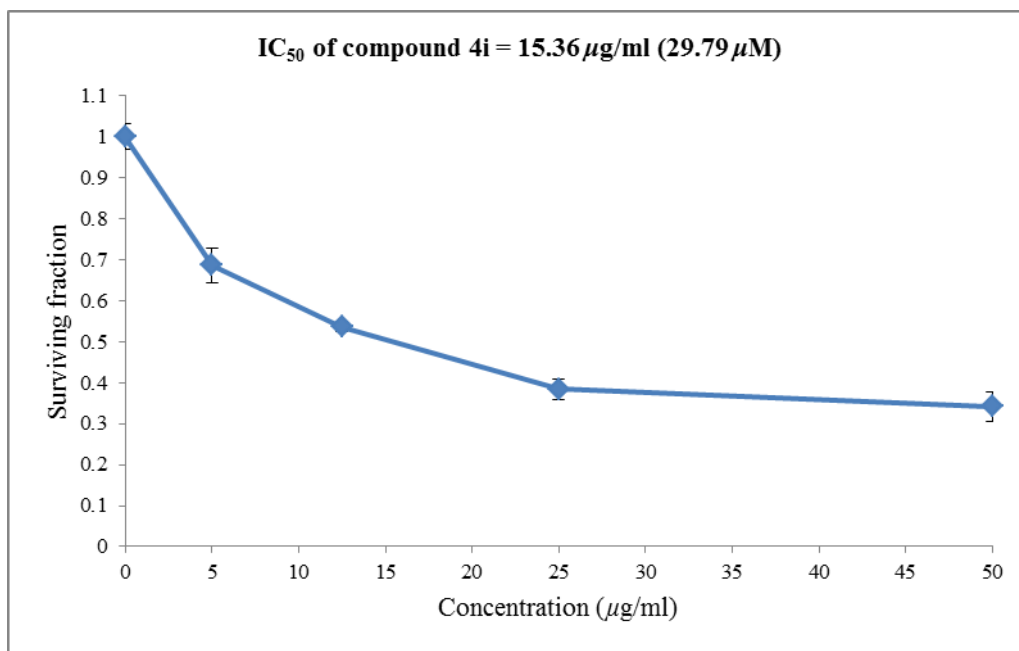
Figure S4. Dose-response curves of the macrocyclic peptidomimetics **4a-1**, and **5-7** against HeLa (cervical) human tumor cell line.

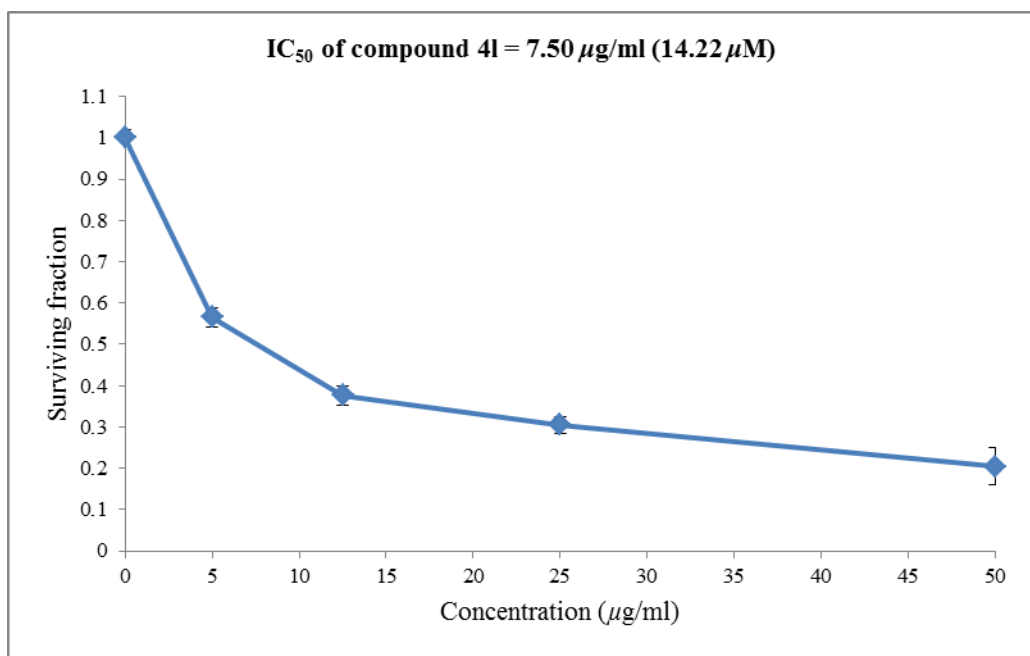
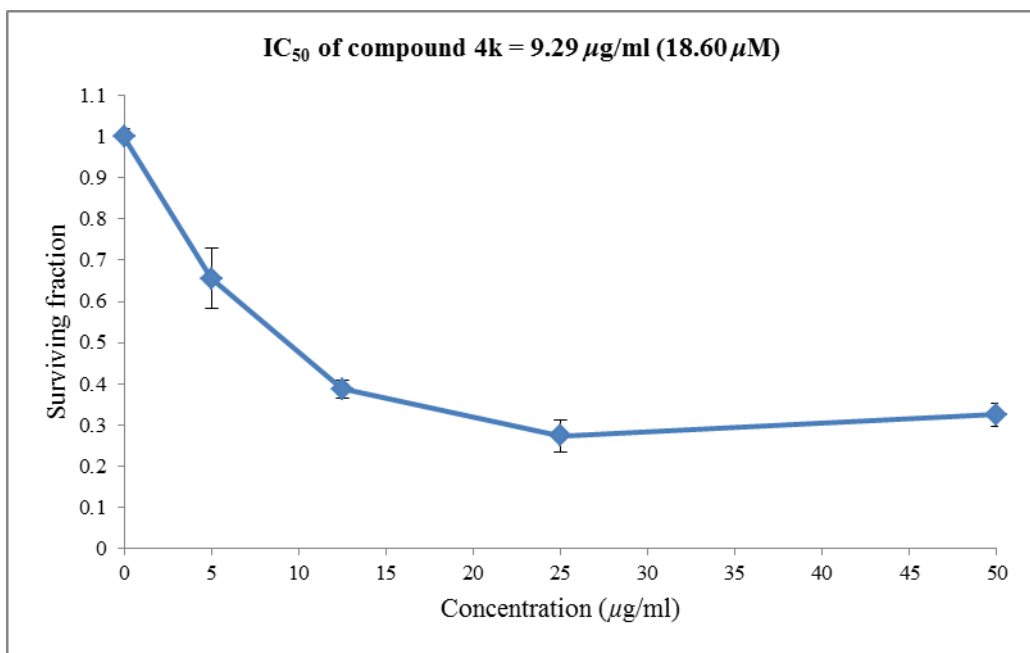


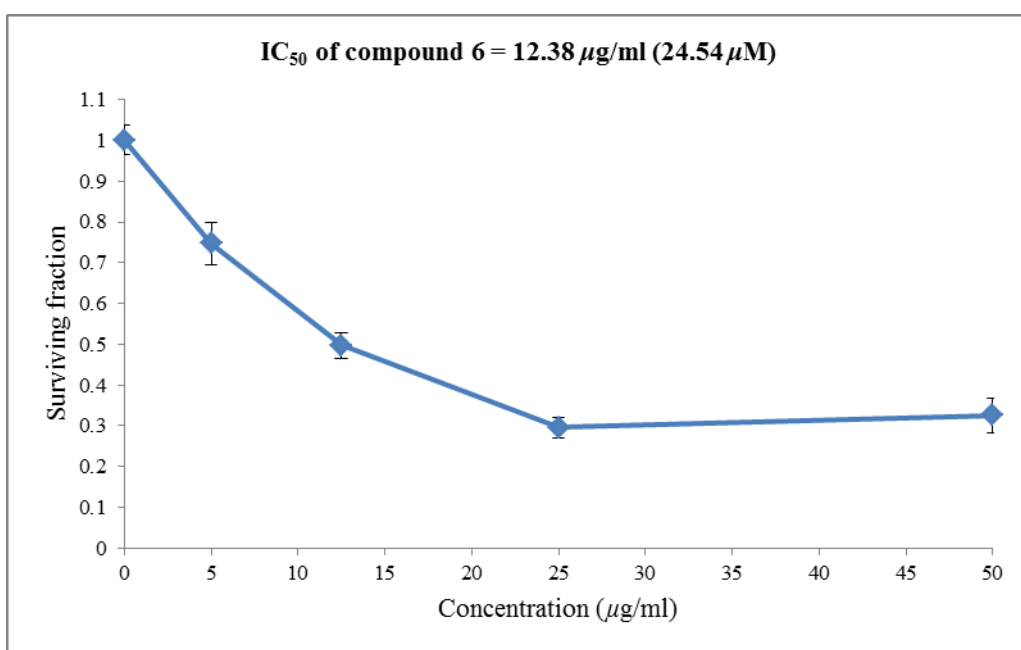
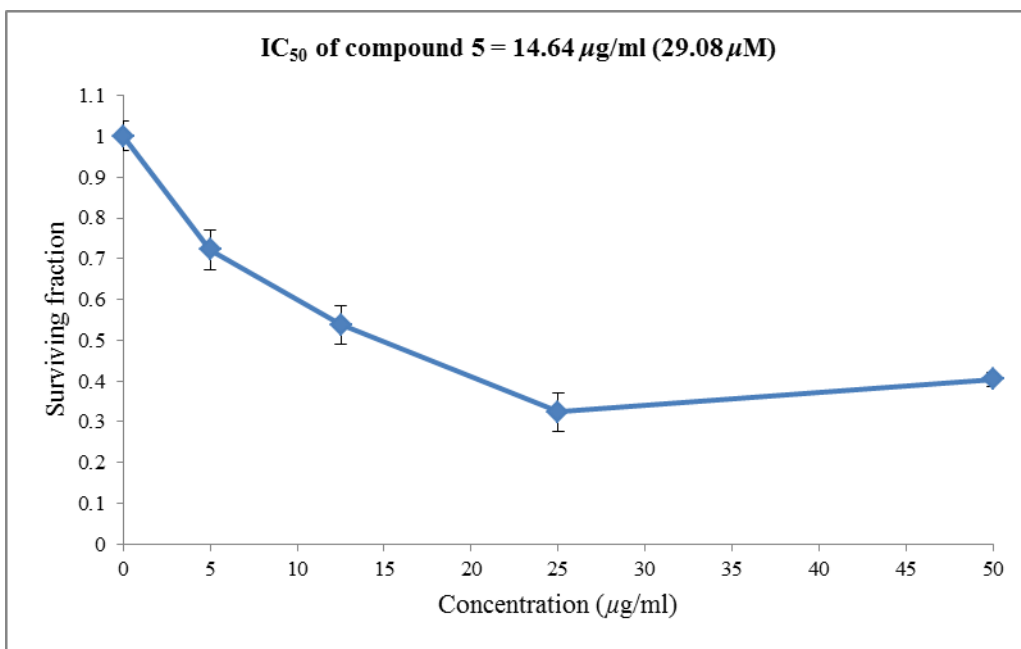












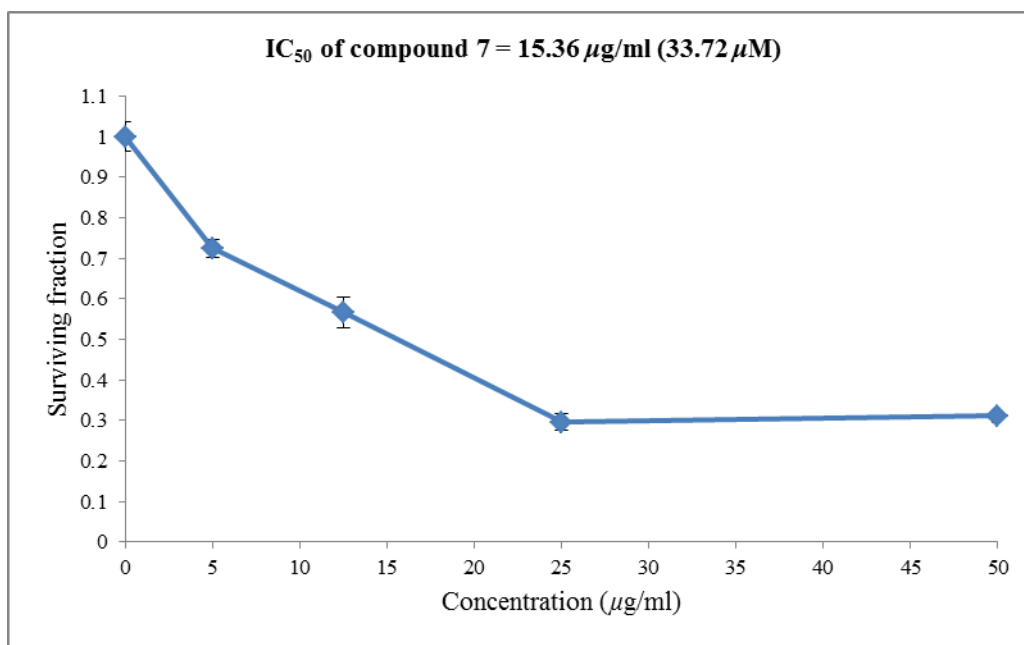
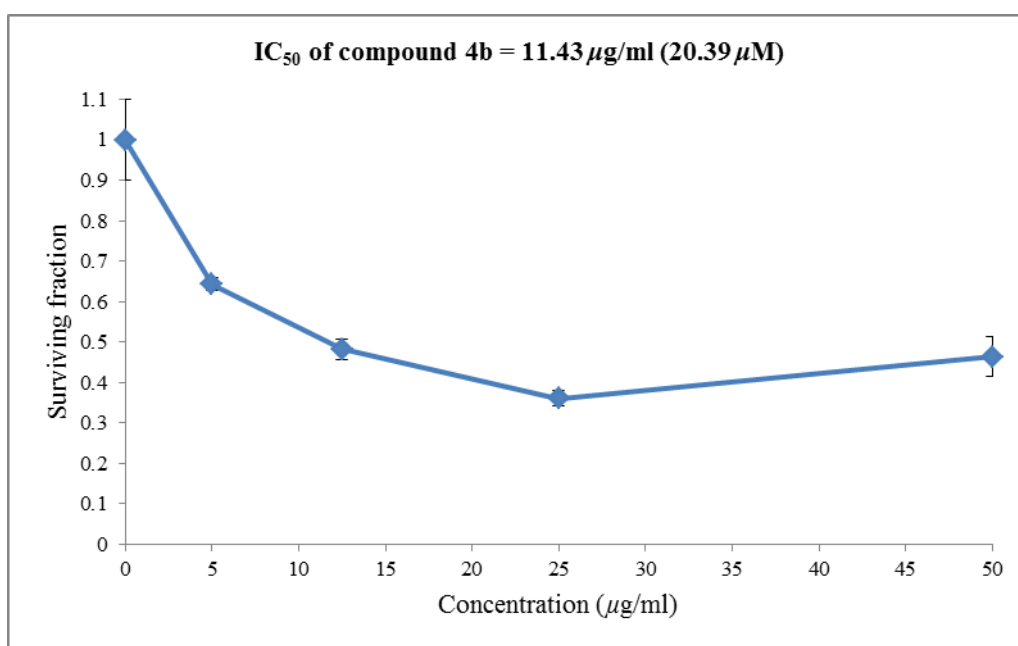
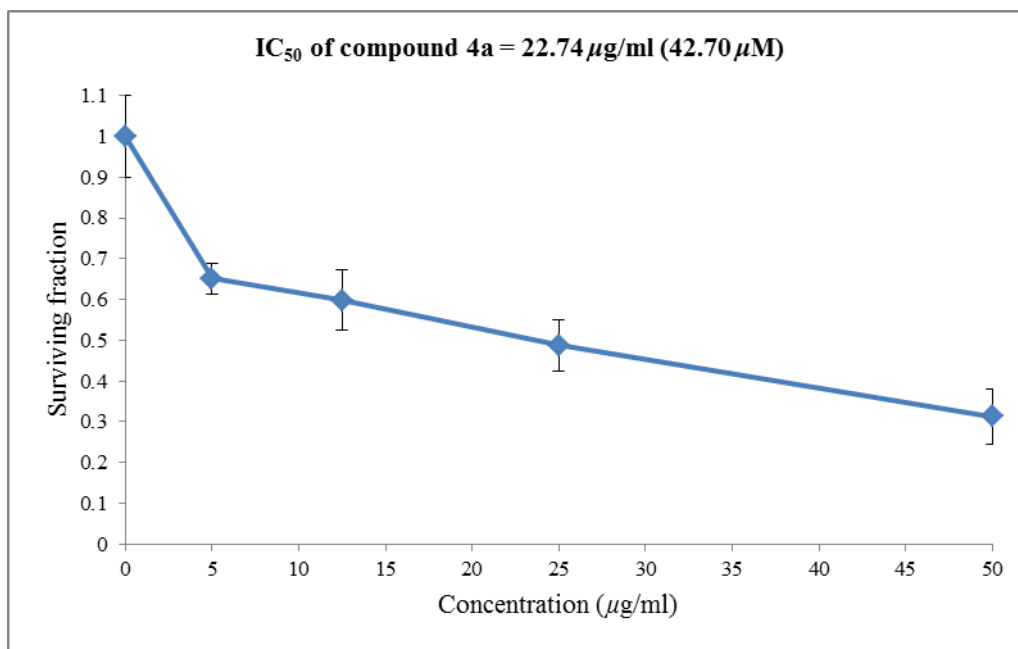
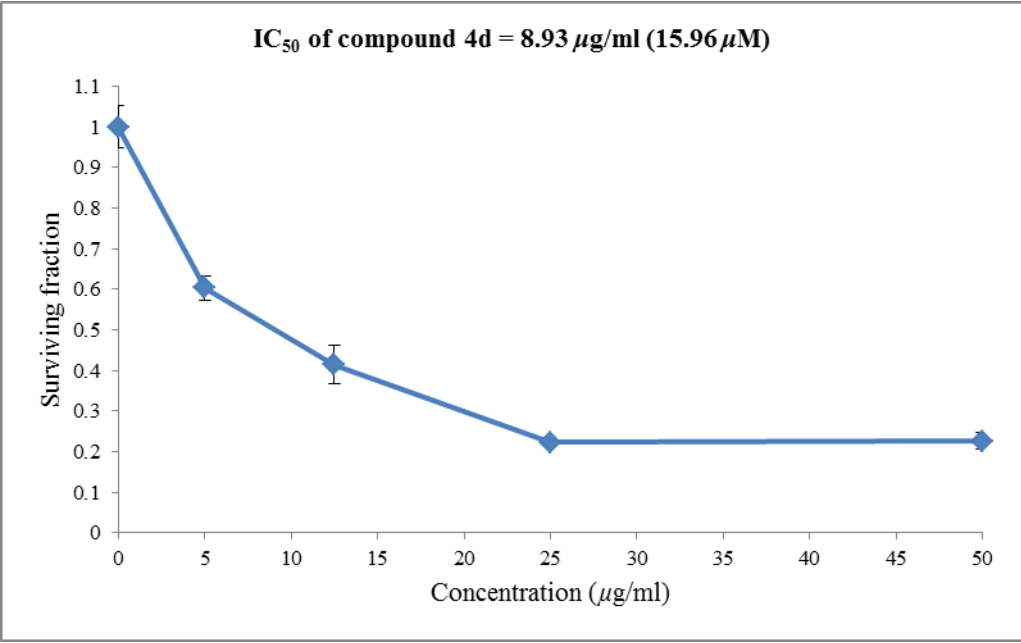
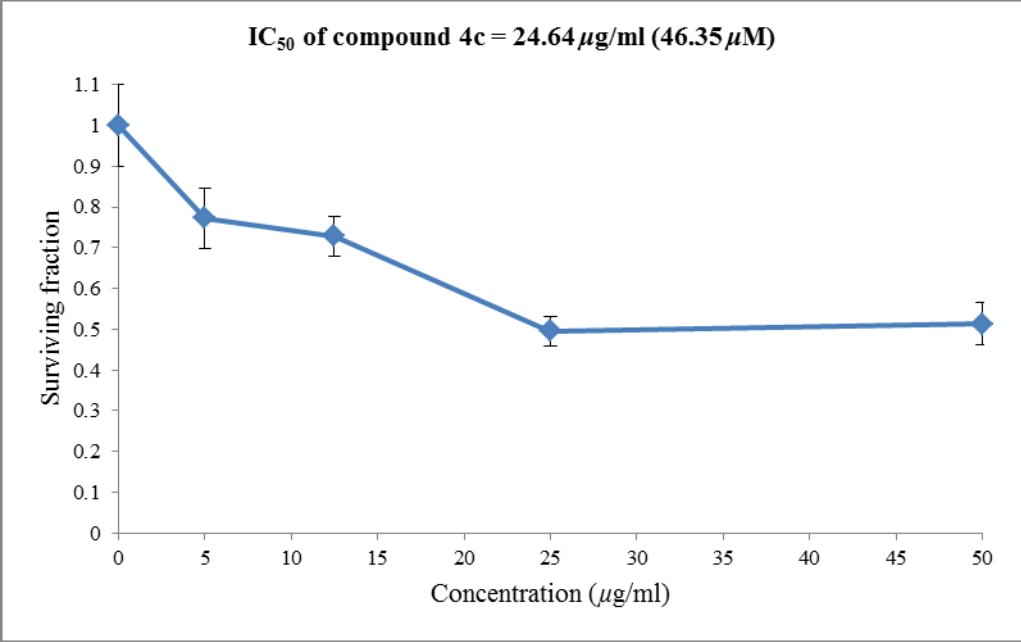
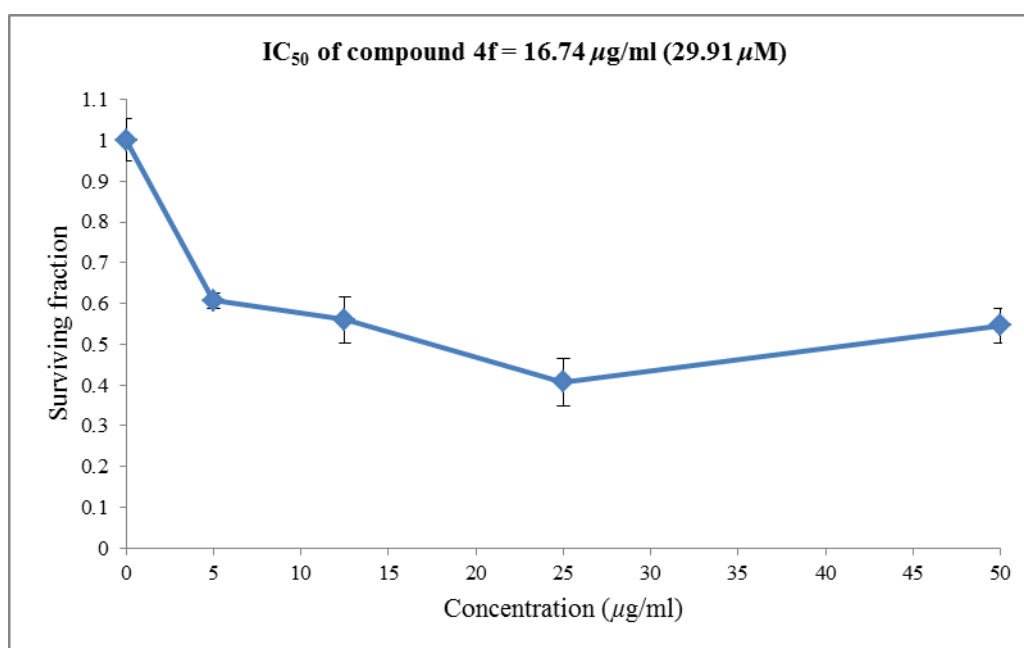
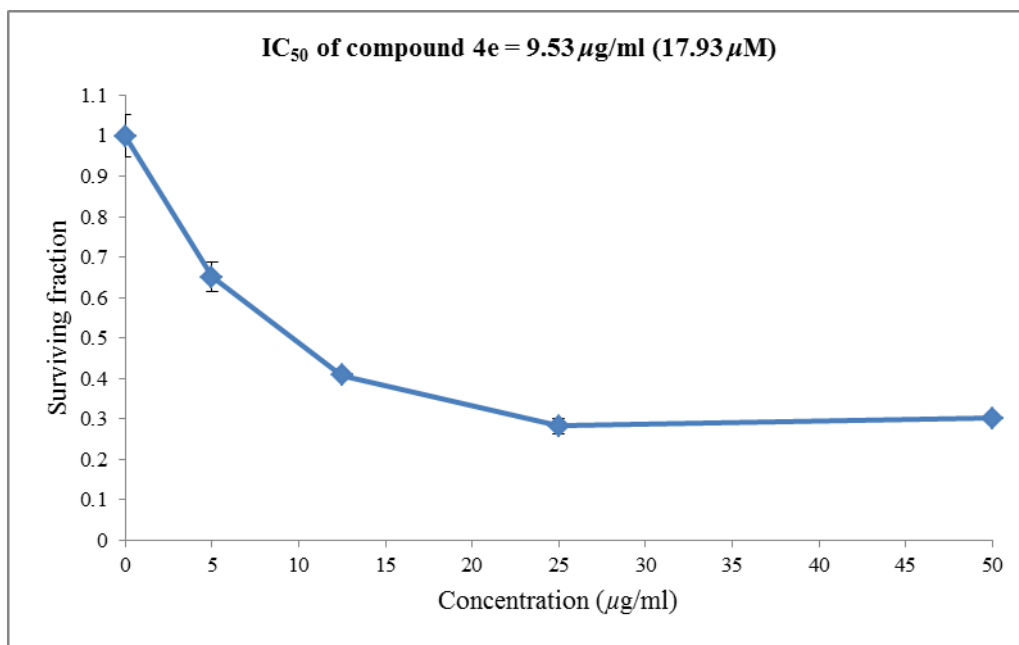
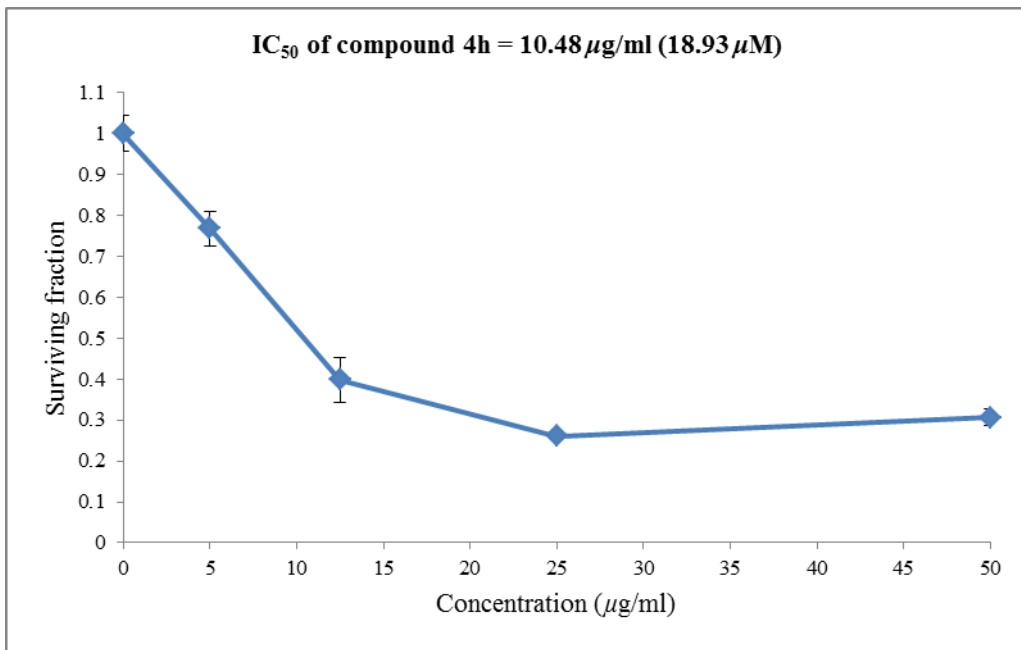
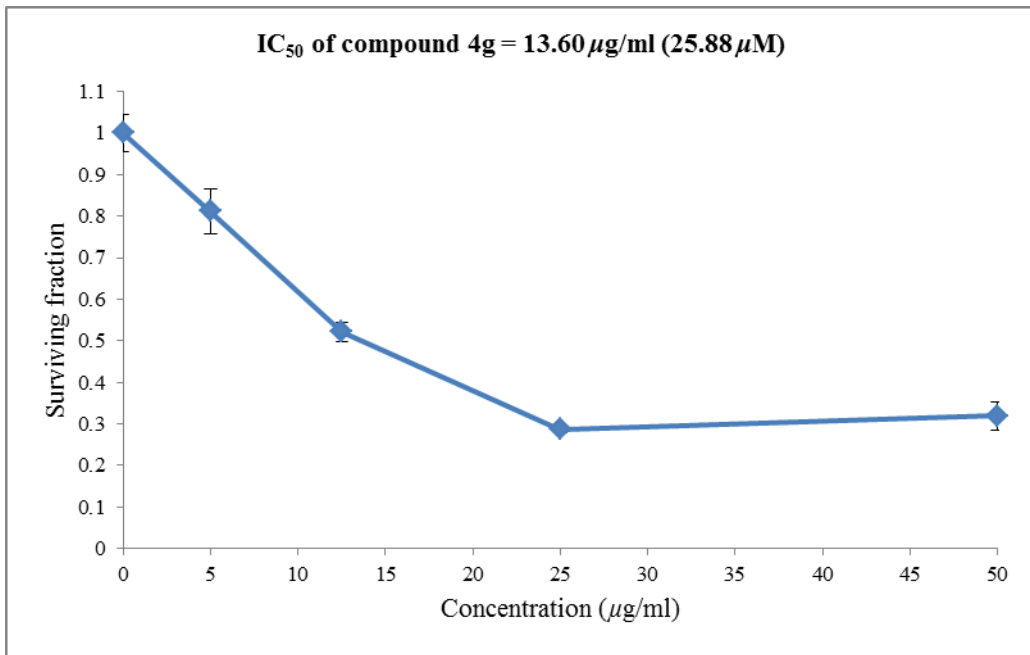


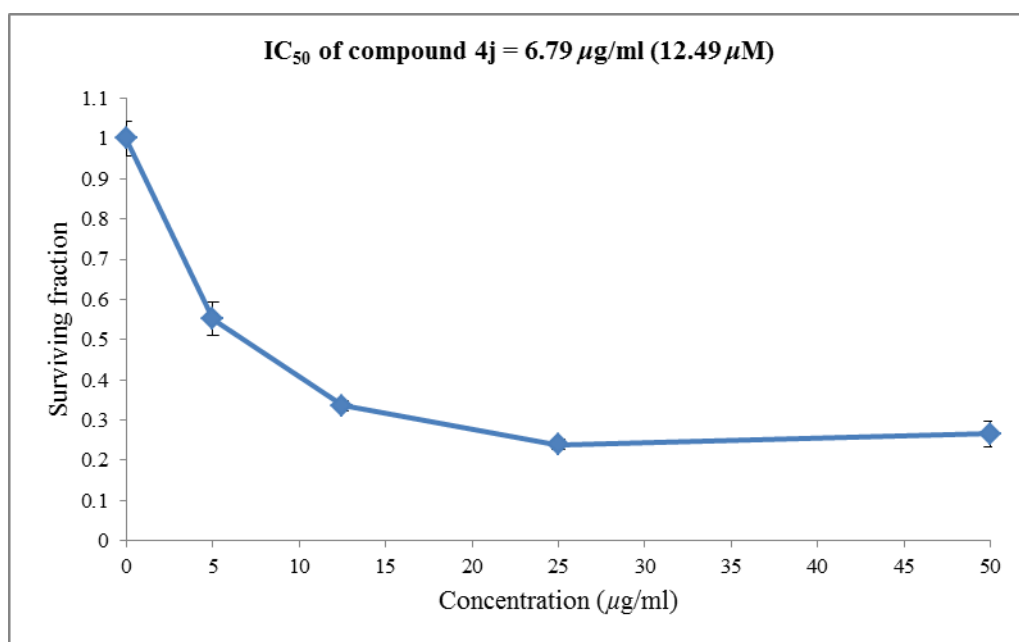
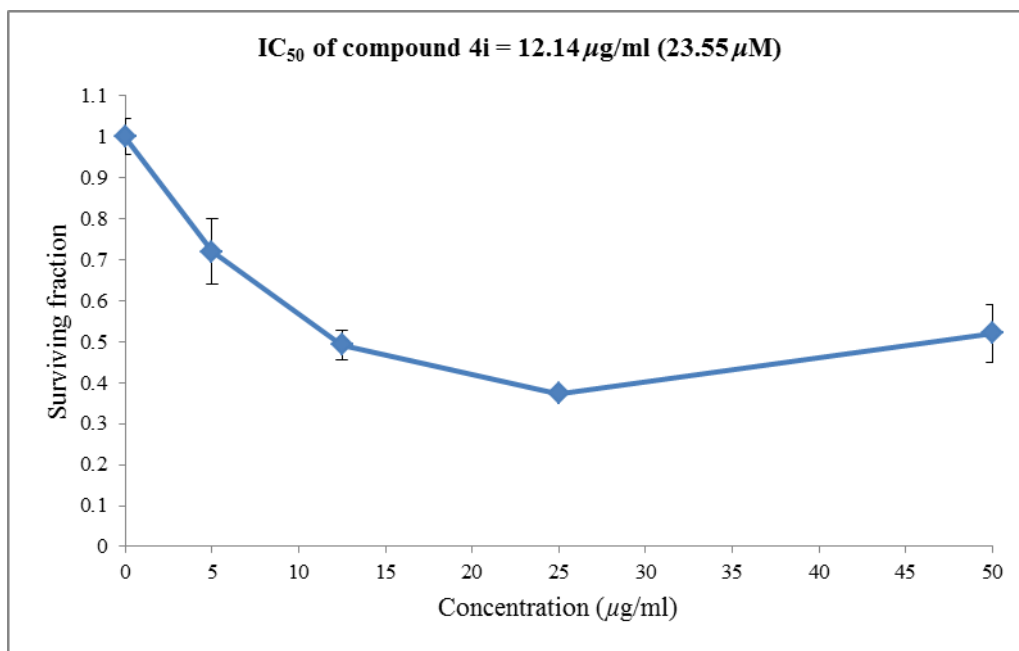
Figure S5. Dose-response curves of the macrocyclic peptidomimetics **4a-1**, and **5-7** against HCT116 (colon) human tumor cell line.

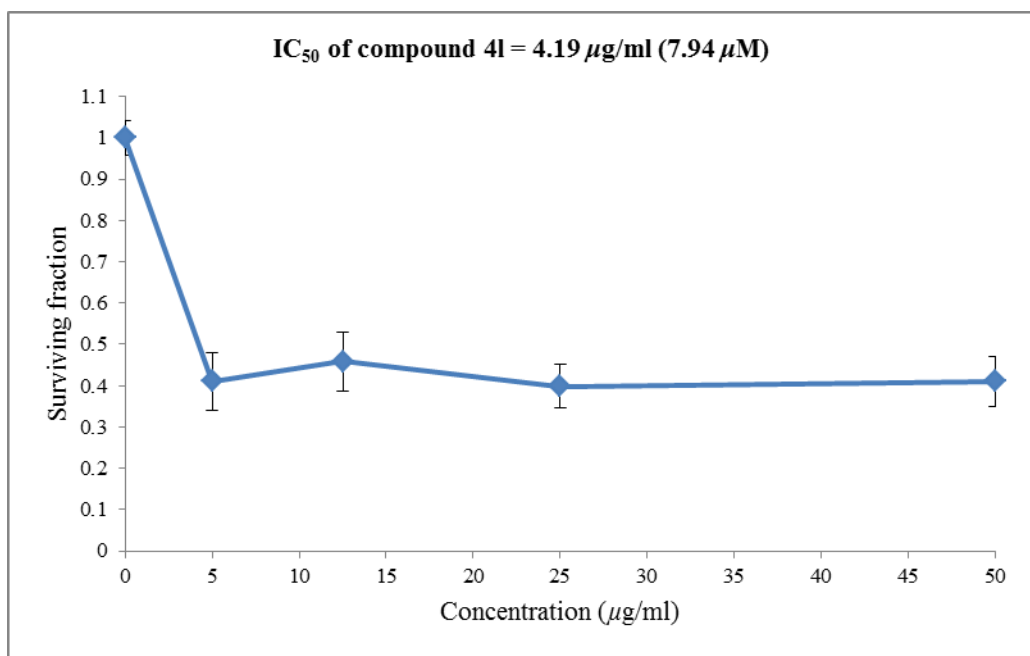
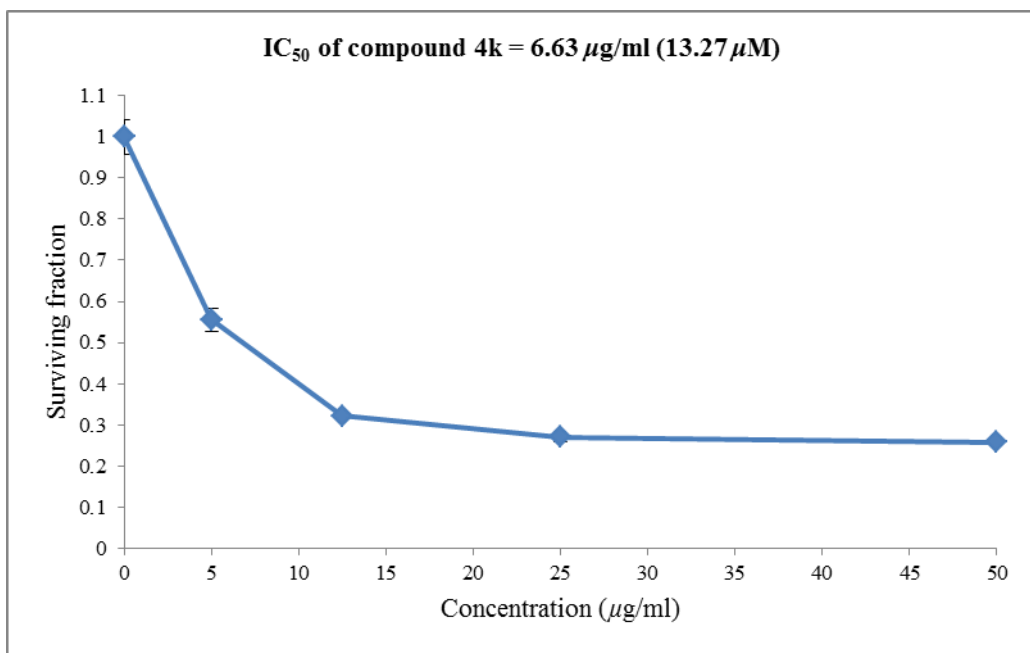


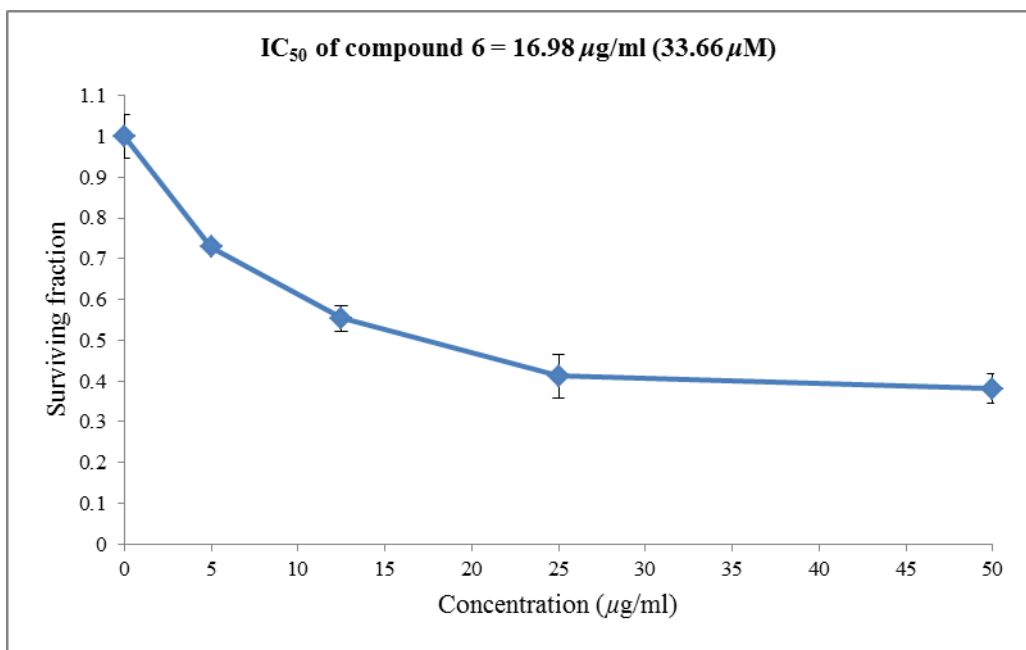
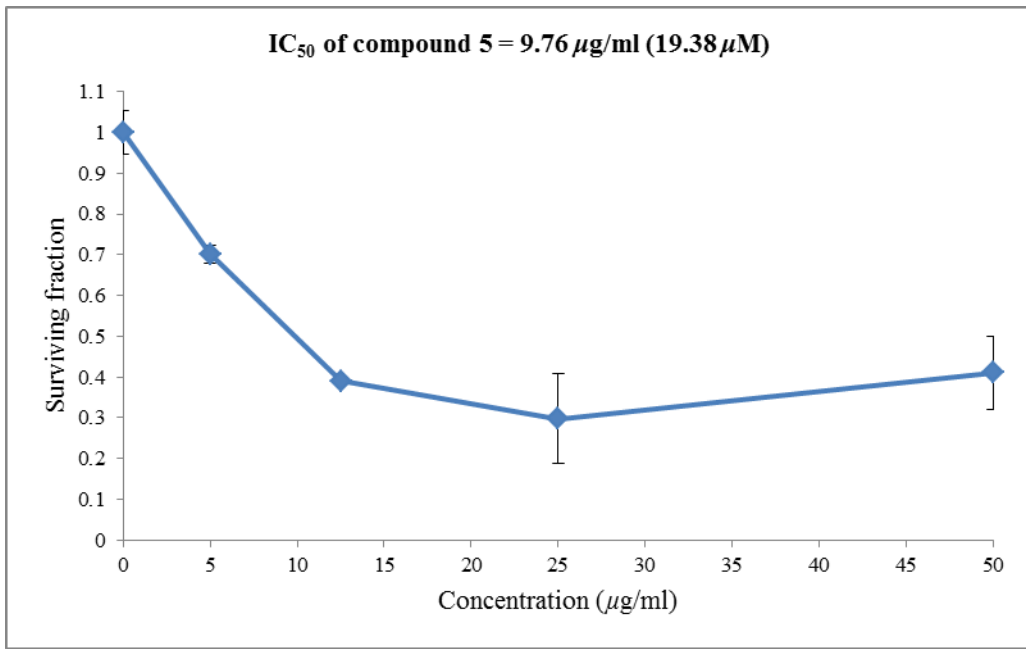












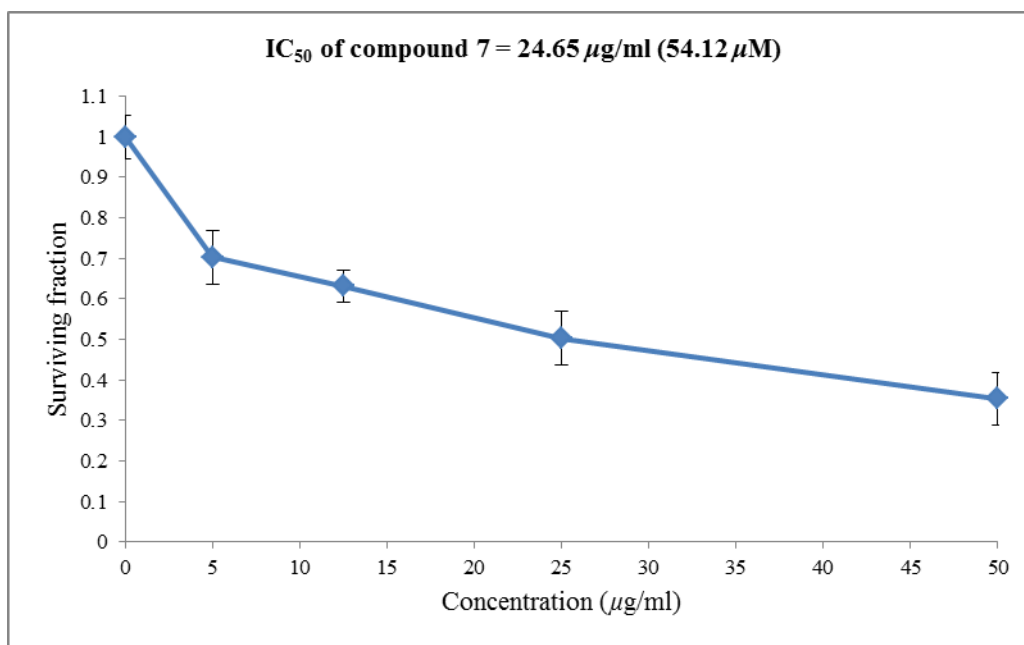
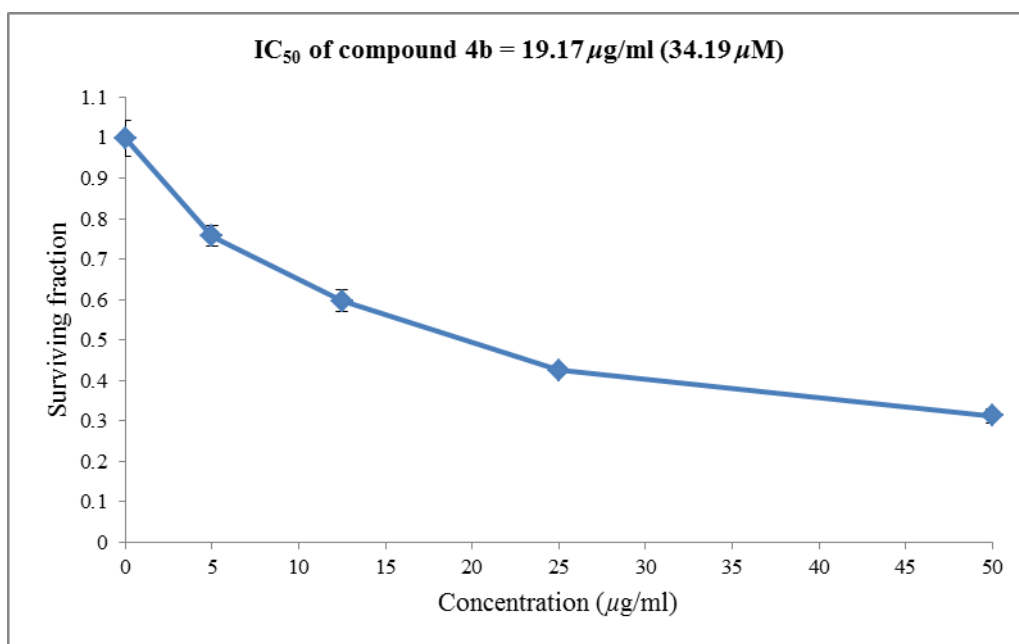
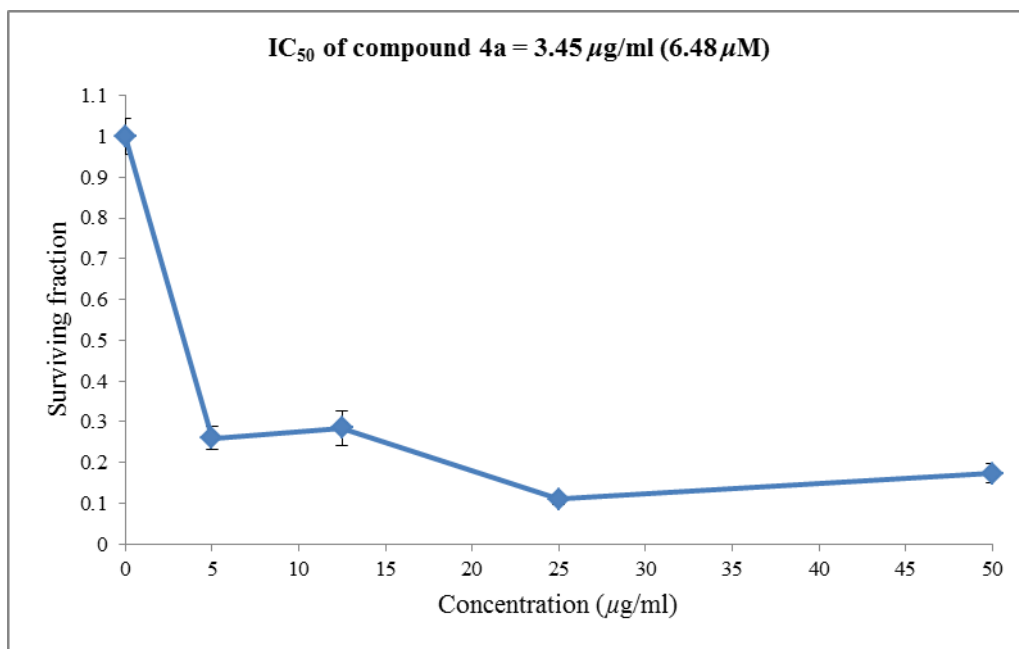
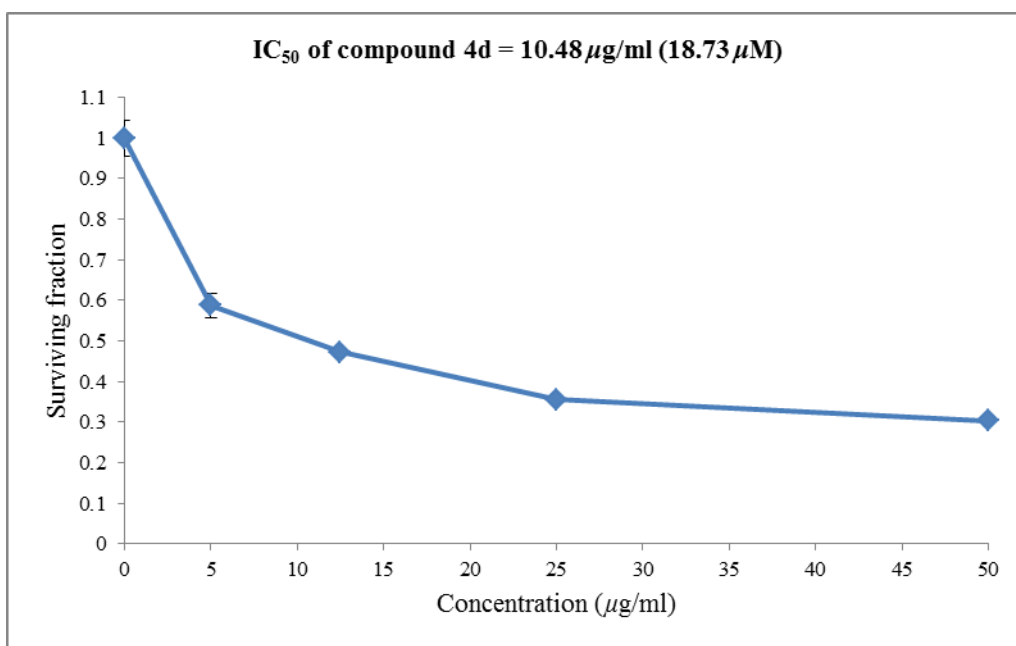
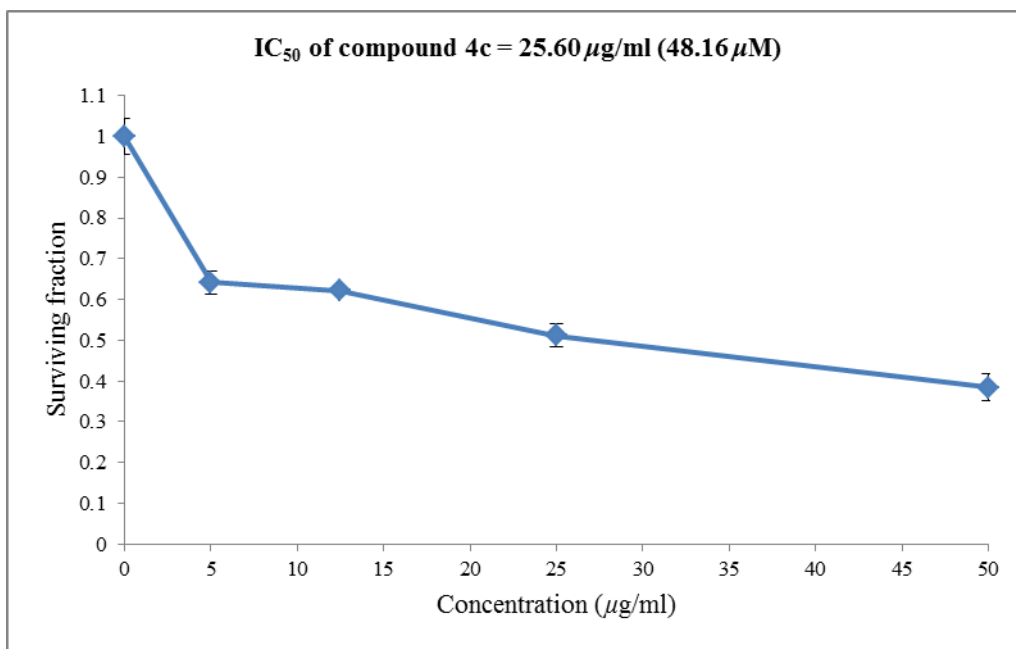
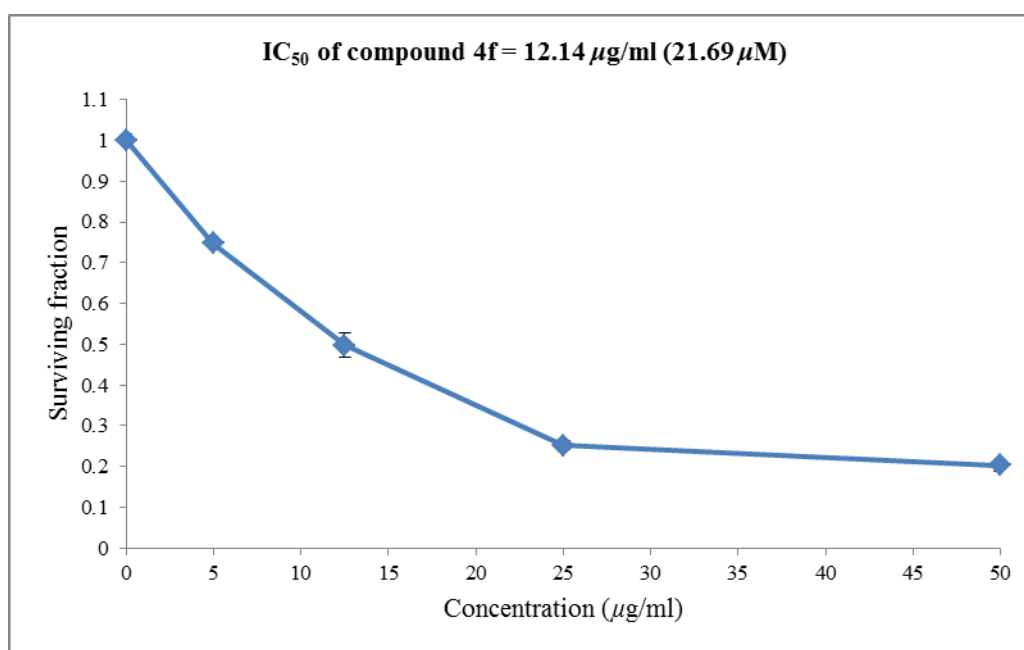
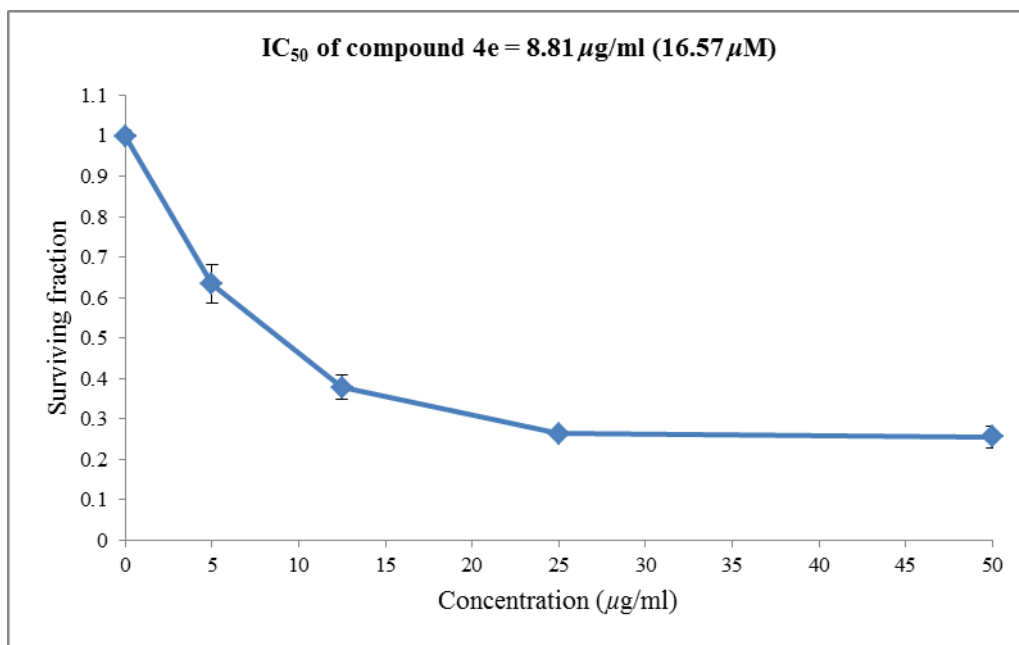
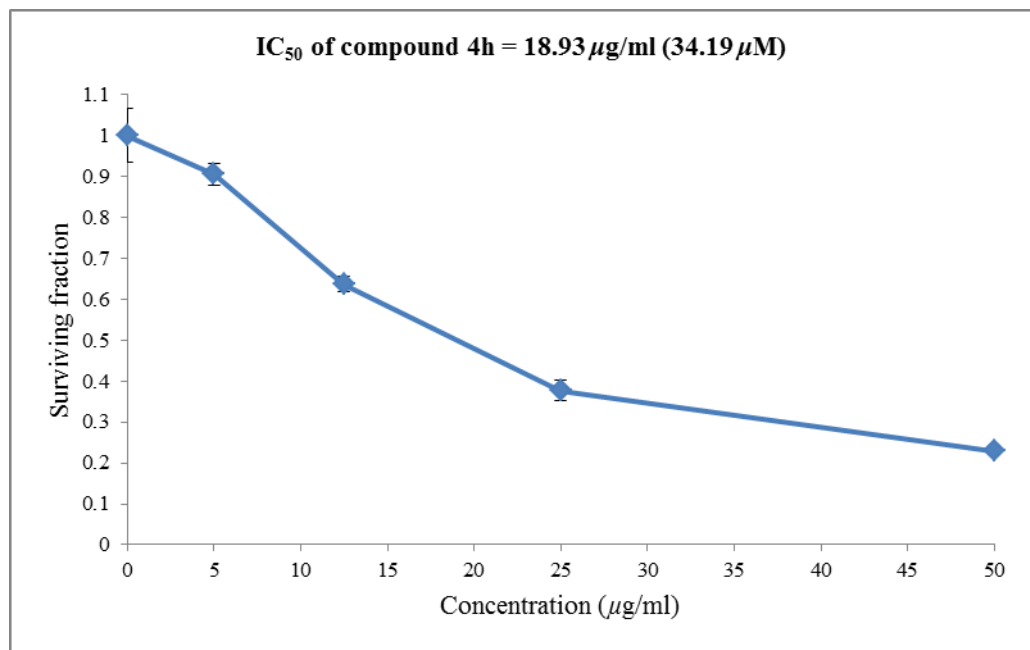
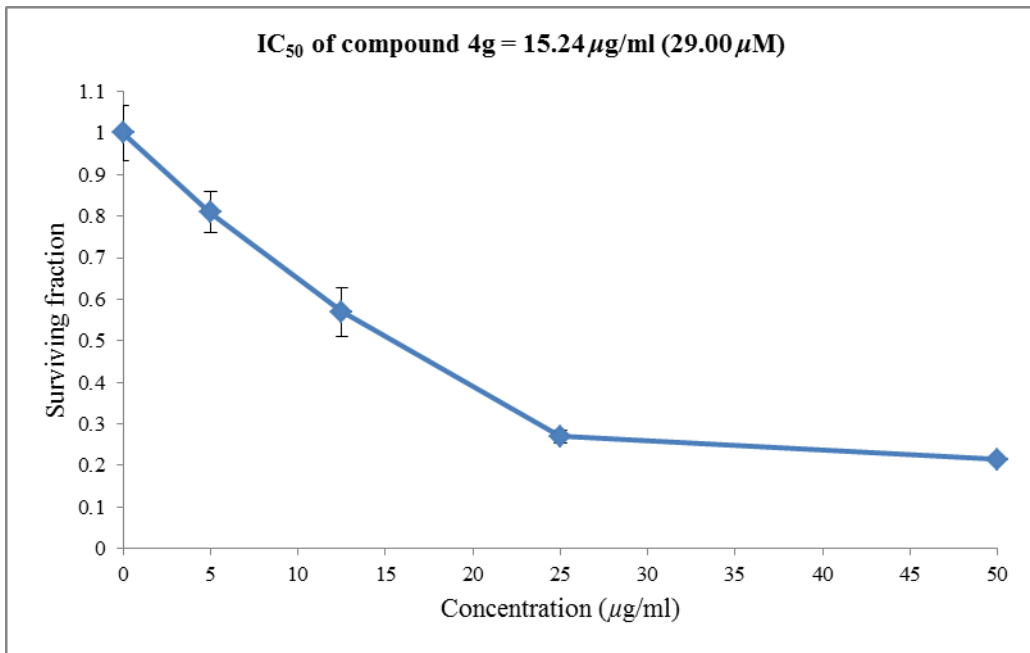


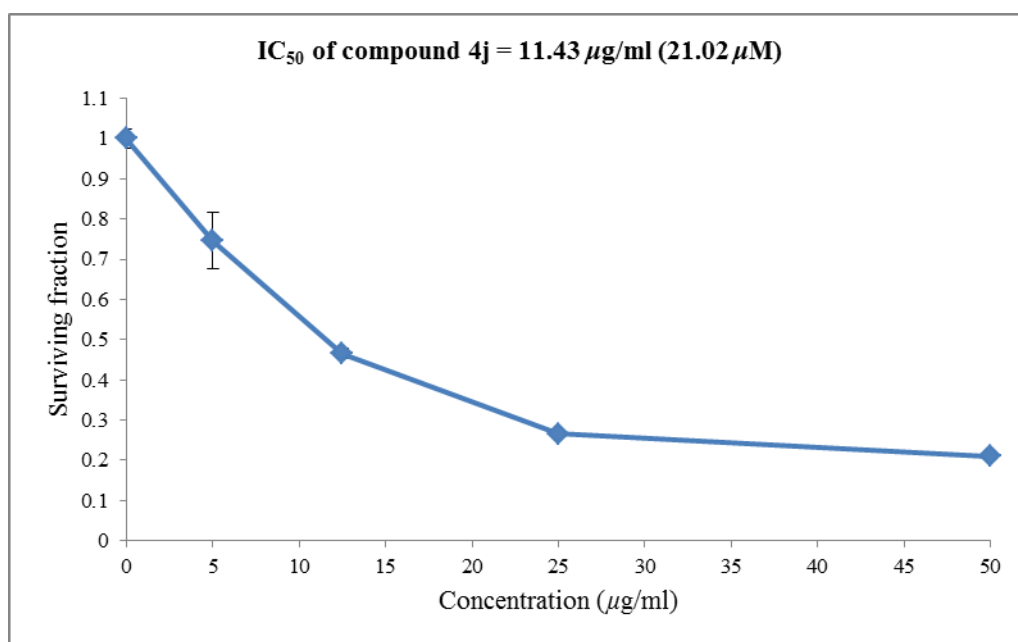
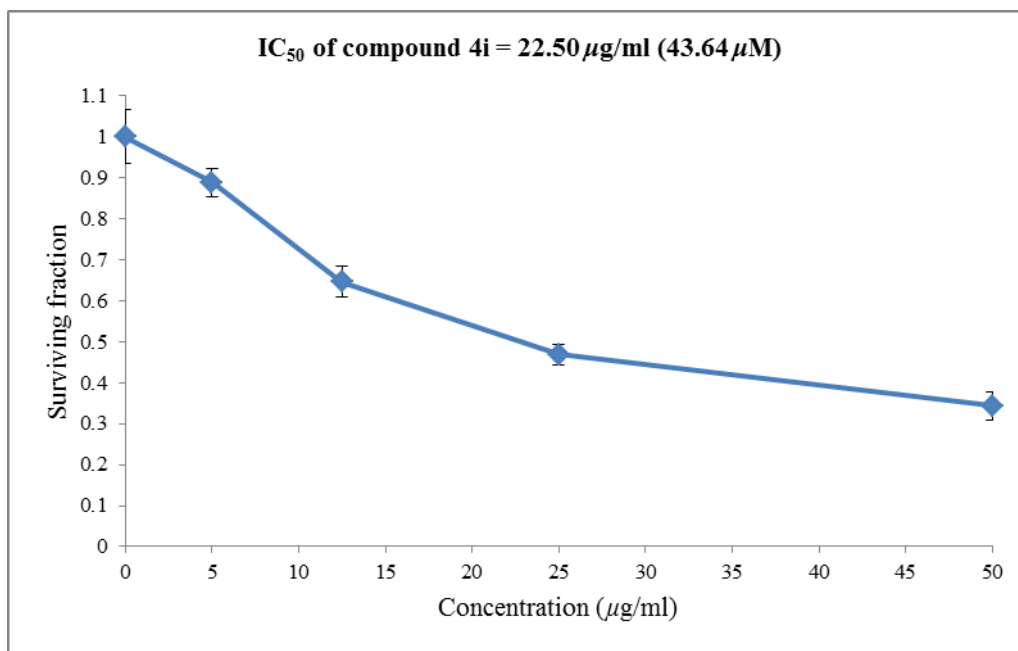
Figure S6. Dose-response curves of the macrocyclic peptidomimetics **4a-1**, and **5-7** against PC3 (prostate) human tumor cell line.

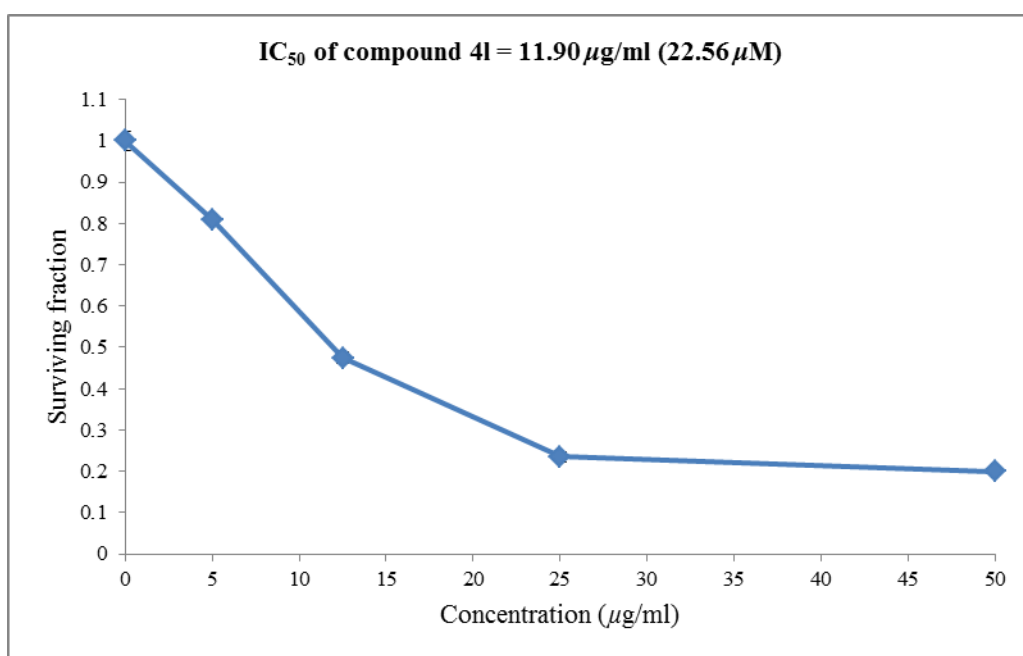
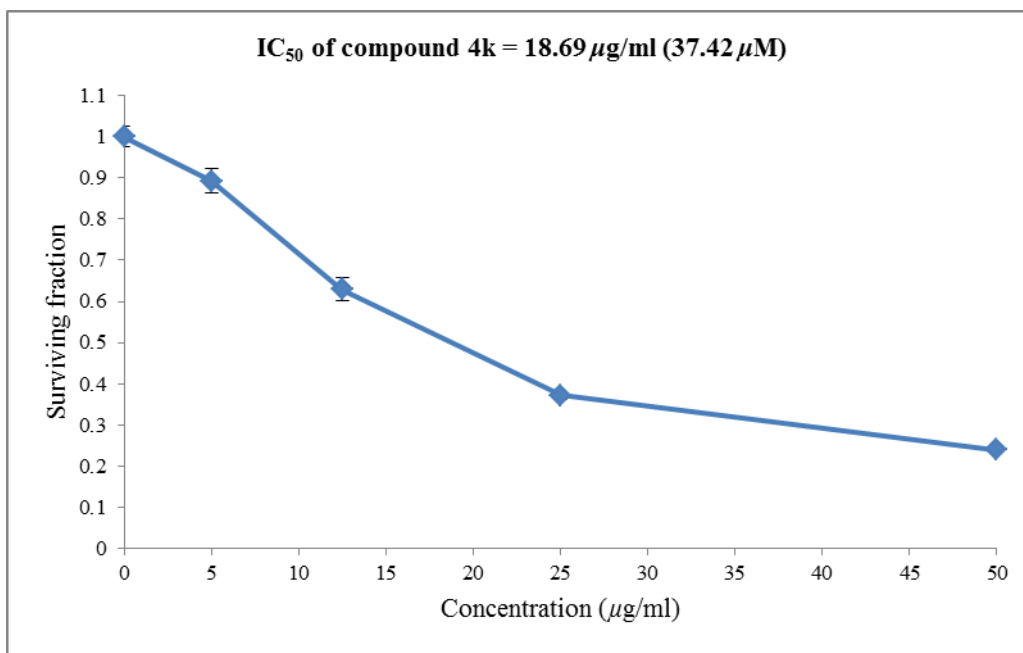


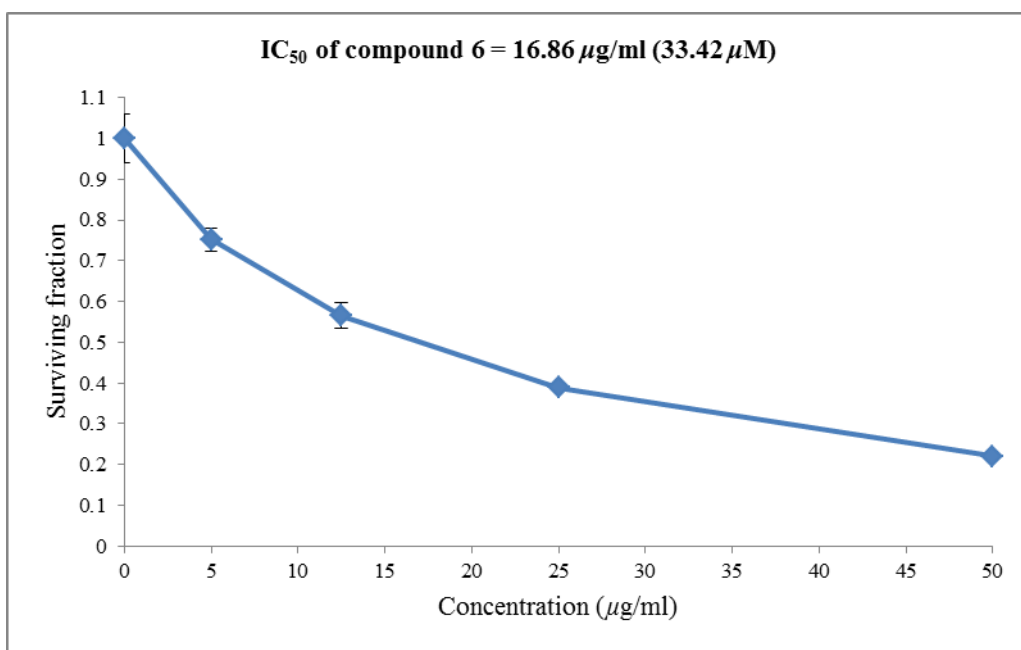
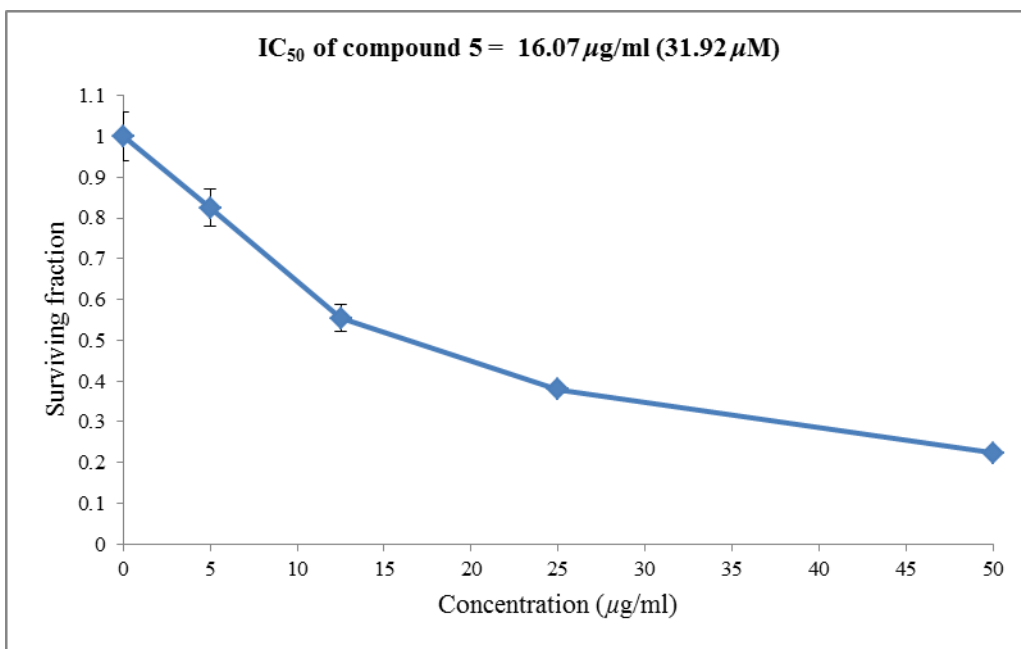












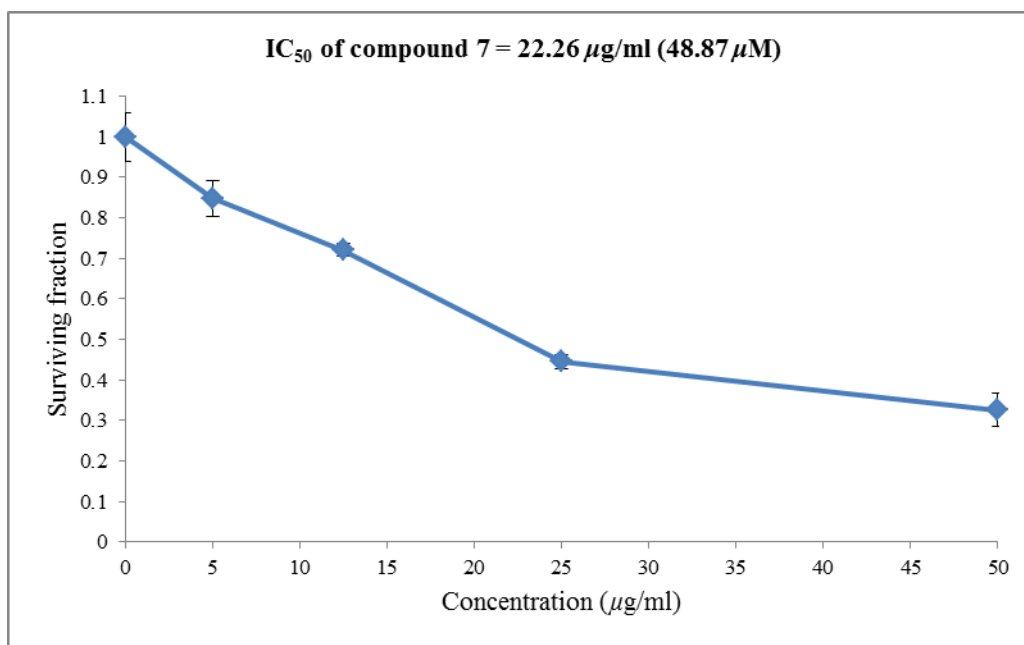


Figure S7. Dose-response curves of the macrocyclic peptidomimetics **4a-1**, and **5-7** against MCF7 (breast) human tumor cell line.