Supplementary Data

Streptogramin A derivatives as mitochondrial translation inhibitors to suppress glioblastoma stem cells growth

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Contents: **Page(s):**

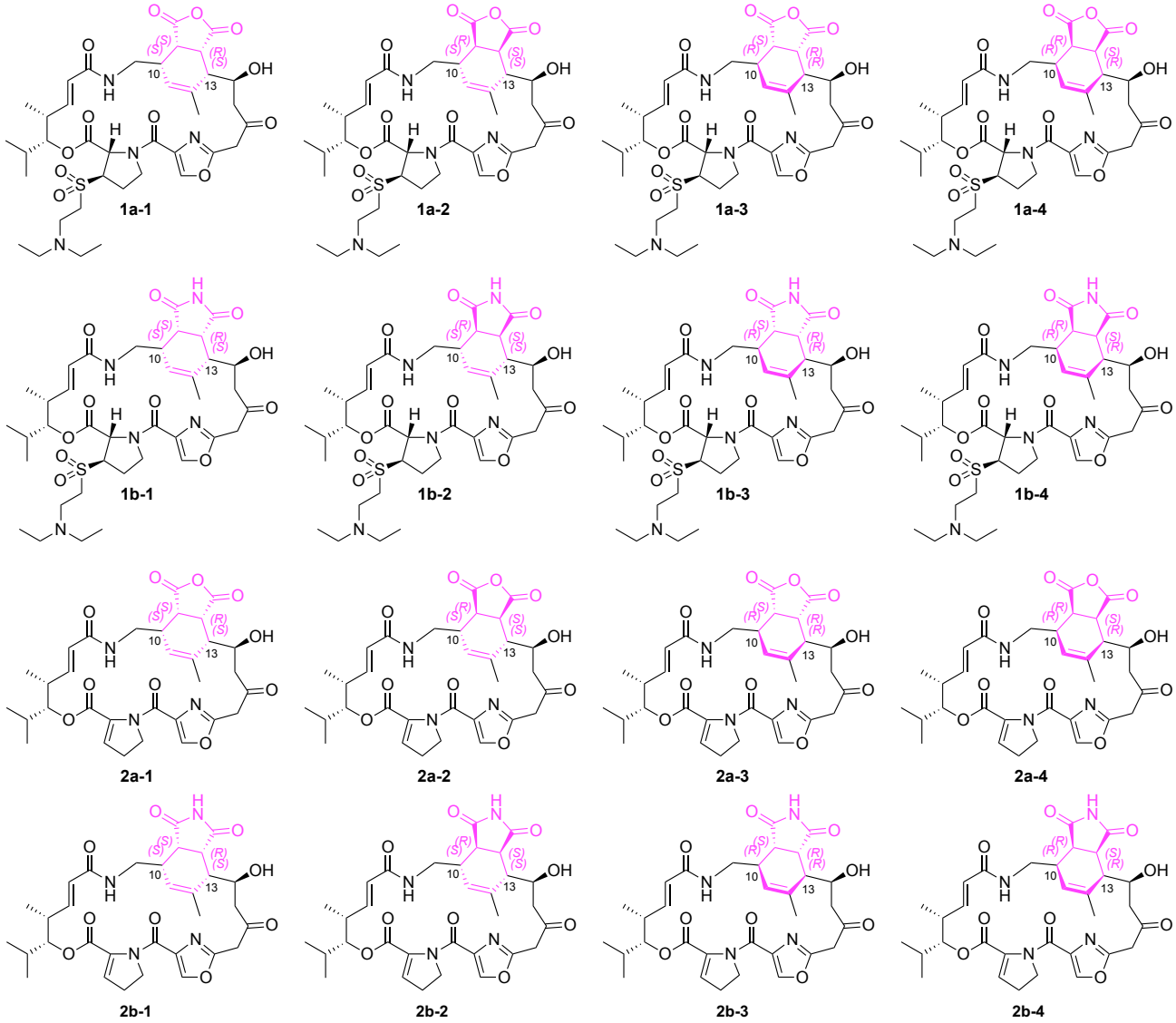


Fig. S1. The molecular structures of each of the four possible diastereomers generated by the Diels-Alder reactions between dalfopristin (**1**) or virginiamycin M1 (**2**) and maleic anhydride or maleimide as dienophiles. The stereochemistry of each possible diastereomer obtainable from the reaction is highlighted.

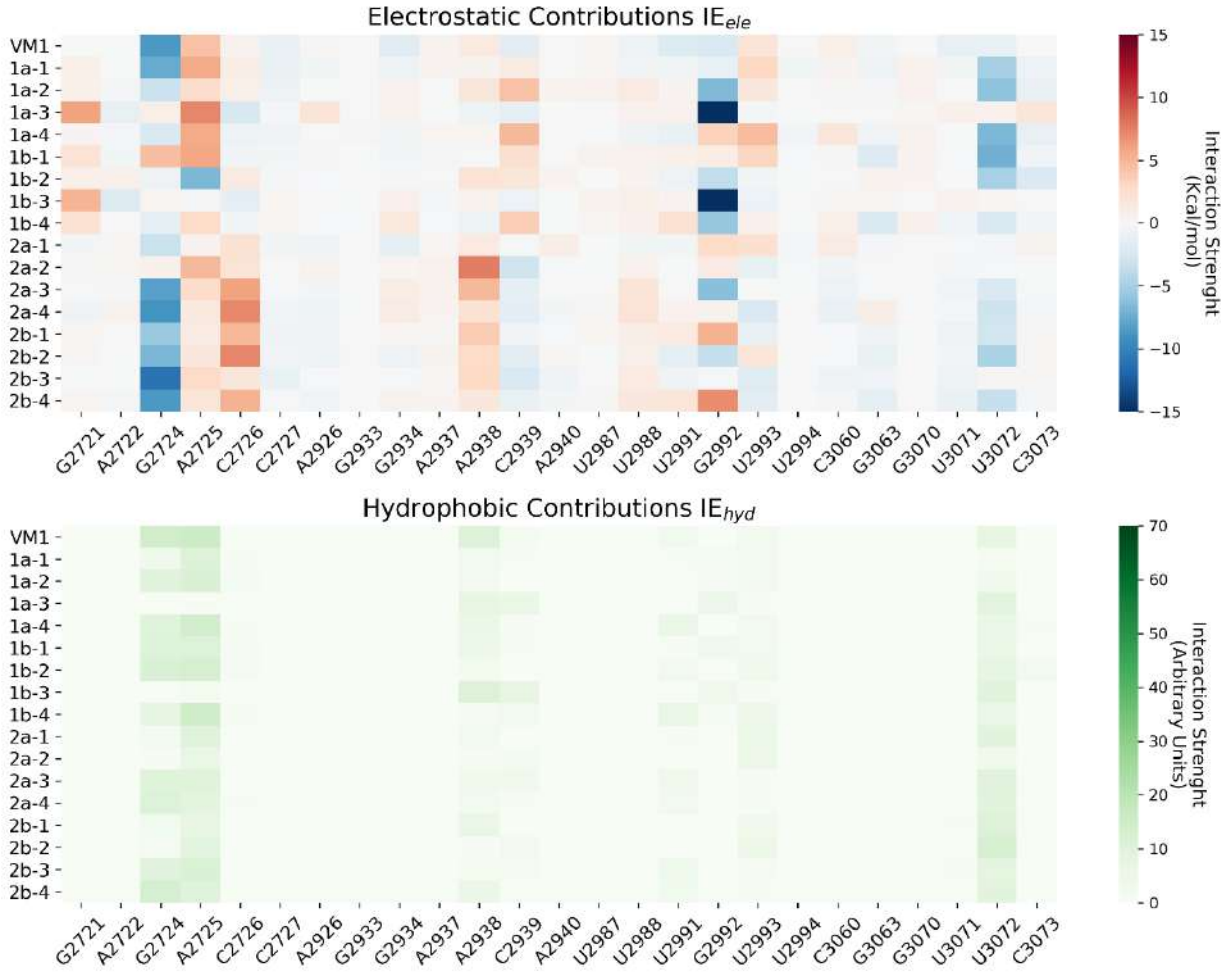


Fig. S2. Per-residue interaction energy decomposition performed on the docking-predicted binding mode for C10/C13 derivatives. VM1 (**2**) binding mode from structure 6I9R is also reported for reference. In detail, the upper panel reports a per-residue decomposition of electrostatic interaction energy: values (in Kcal/mol) are plotted according to the colour scale reported on the right, ranging from red (positive value, repulsive interaction) to black (negative value, attractive interaction). The lower panel, instead, reports a per-residue decomposition of hydrophobic interactions: values (in arbitrary/adimensional units) are plotted according to the color scale reported on the right, ranging from white (zero, absence of interaction) to dark teal (indicating a good hydrophobic interaction).

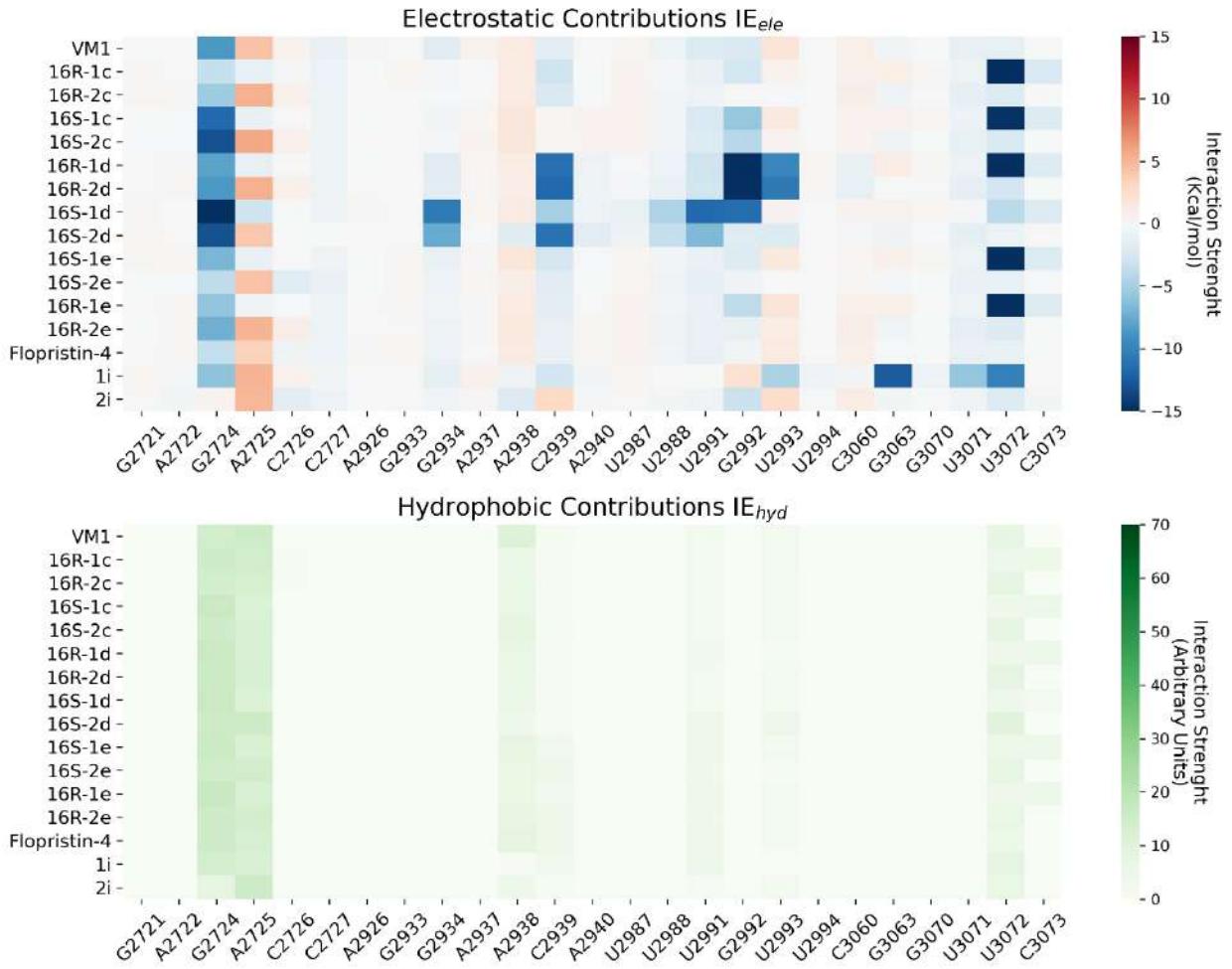


Fig. S3. Per-residue interaction energy decomposition performed on the docking-predicted binding mode for C16 derivatives. VM1 (**2**) binding mode from structure 6I9R is also reported for reference. In detail, the upper panel reports a per-residue decomposition of electrostatic interaction energy: values (in Kcal/mol) are plotted according to the colour scale reported on the right, ranging from red (positive value, repulsive interaction) to black (negative value, attractive interaction). The lower panel, instead, reports a per-residue decomposition of hydrophobic interactions: values (in arbitrary/adimensional units) are plotted according to the color scale reported on the right, ranging from white (zero, absence of interaction) to dark teal (indicating a good hydrophobic interaction).

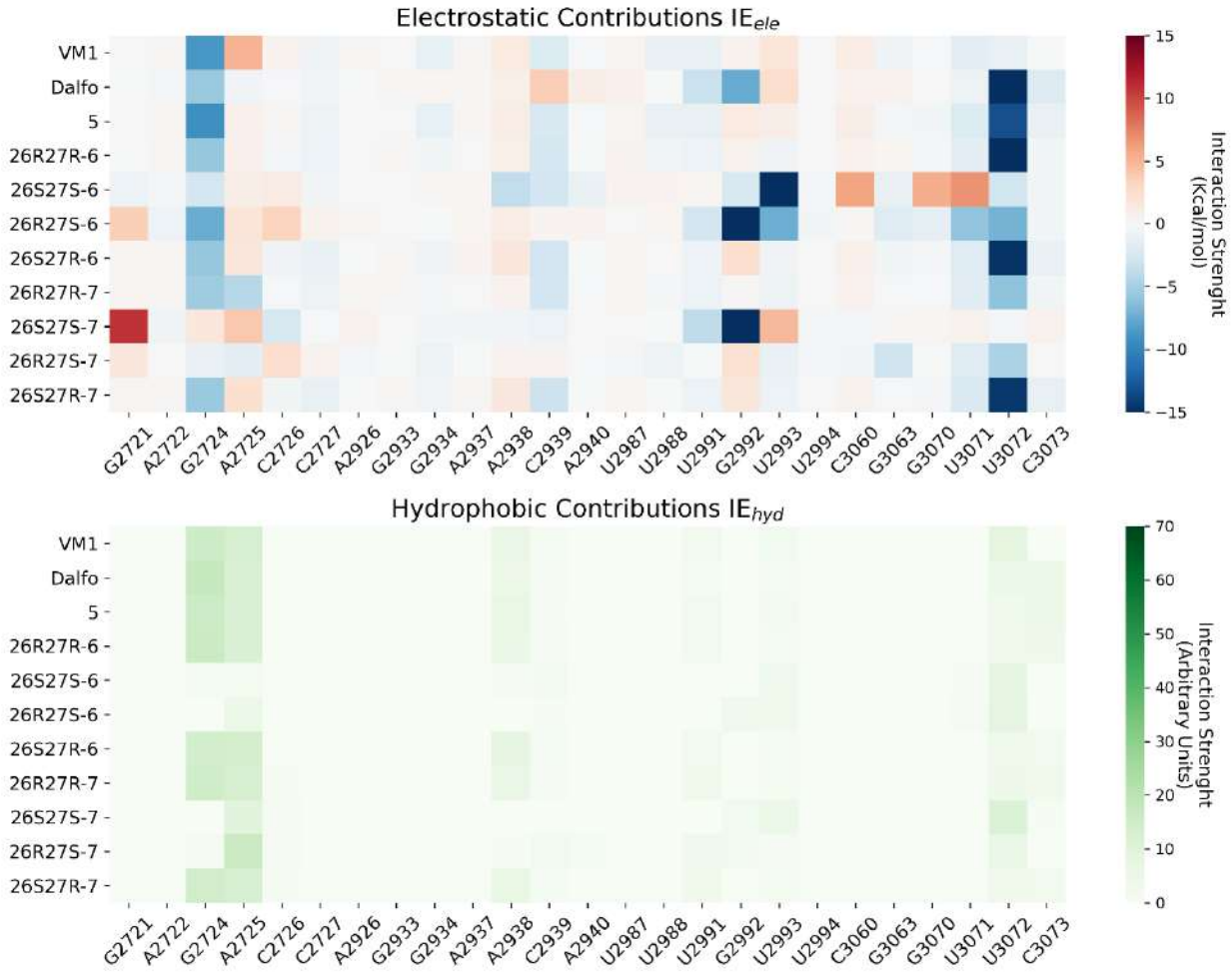


Fig. S4. Per-residue interaction energy decomposition performed on the docking-predicted binding mode for C26 derivatives. VM1 (**2**) binding mode from structure 6I9R is also reported for reference. In detail, the upper panel reports a per-residue decomposition of electrostatic interaction energy: values (in Kcal/mol) are plotted according to the colour scale reported on the right, ranging from red (positive value, repulsive interaction) to black (negative value, attractive interaction). The lower panel, instead, reports a per-residue decomposition of hydrophobic interactions: values (in arbitrary/adimensional units) are plotted according to the color scale reported on the right, ranging from white (zero, absence of interaction) to dark teal (indicating a good hydrophobic interaction).

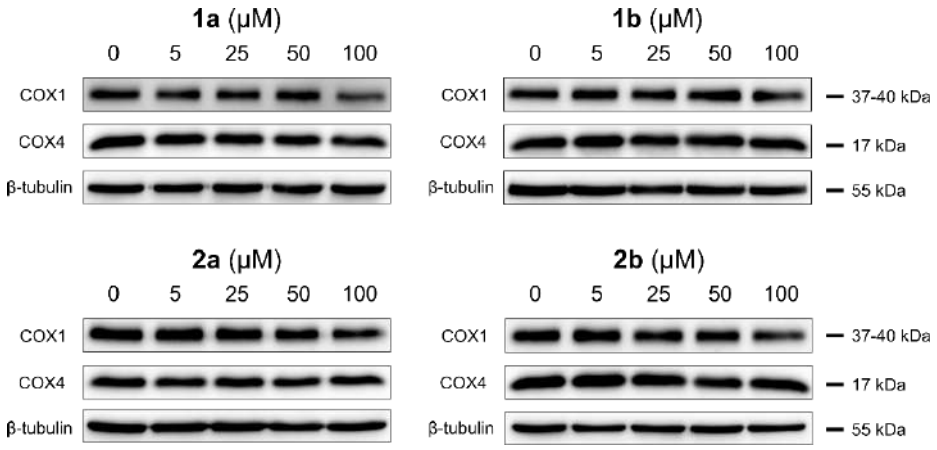


Fig. S5. Western blot analysis for the evaluation of mitochondrial translation inhibition of C10/C13 derivatives. C10/C13 derivatives do not inhibit mitochondrial translation. Immunoblotting on COMI cells after 48 hours of treatment with increasing concentrations of **1a**, **1b, 2a**, and **2c**, alone (*left column*) or in combination with Q (**3**) (*right column*). Effects on COX1, COX4 and beta tubulin proteins are shown. One representative result, n=3 biological replicates.

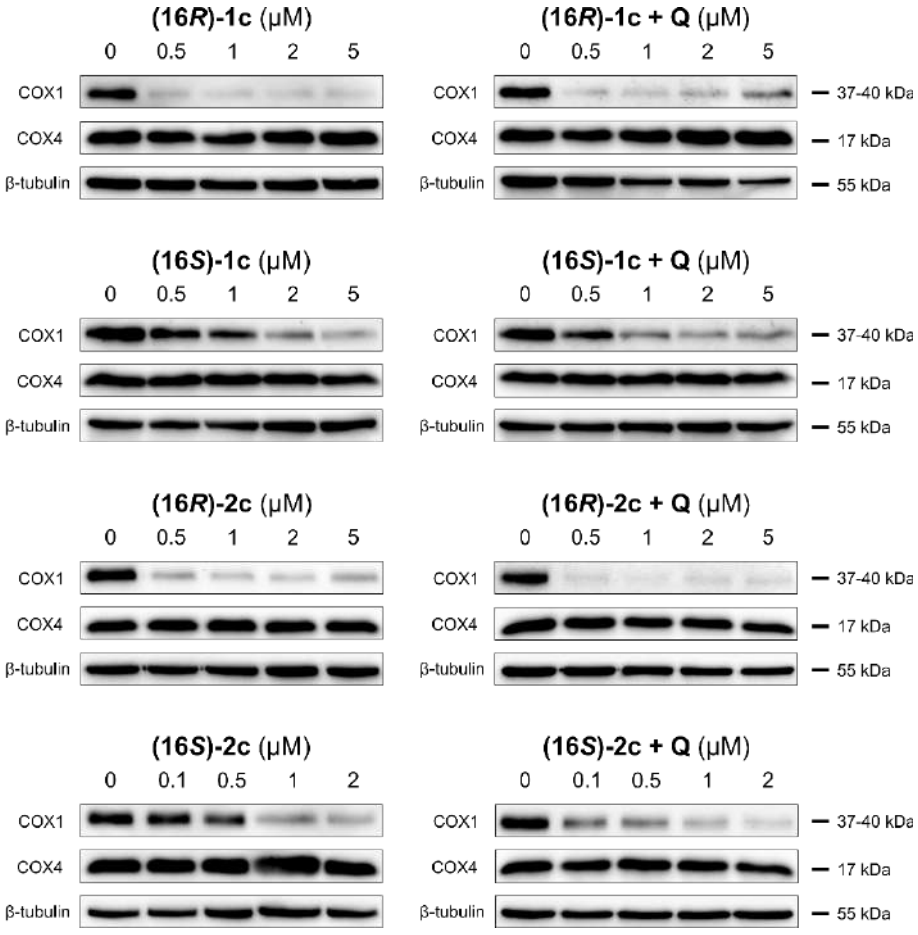


Fig. S6. Western blot analysis for the evaluation of mitochondrial translation inhibition of C16 hydroxyl derivatives. C16 hydroxyl derivatives inhibit mitochondrial translation. Immunoblotting on COMI cells after 48 hours of treatment with increasing concentrations of **(16*R*)-1c**, **(16*S*)-1c, (16*R*)-2c**, and **(16*S*)-2c**, alone (*left column*) or in combination with Q (**3**) (*right column*). Effects on COX1, COX4 and beta tubulin proteins are shown. One representative result, n=3 biological replicates.

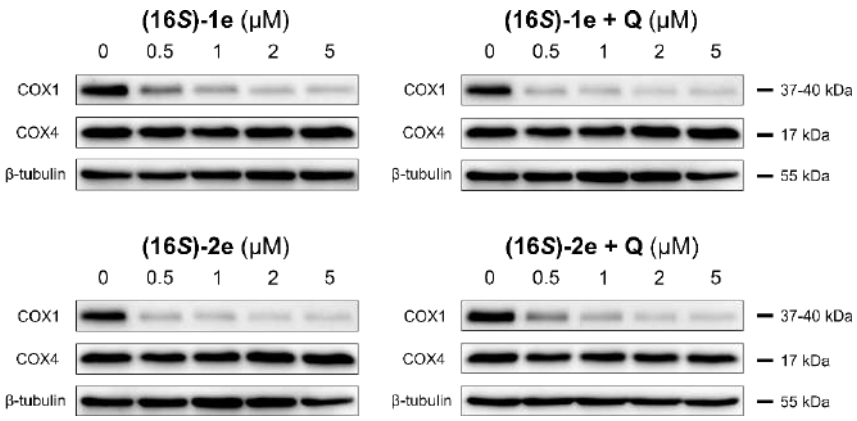


Fig. S7. Western blot analysis for the evaluation of mitochondrial translation inhibition of (16*S*) fluoro derivatives. (16*S*) fluoro derivatives inhibit mitochondrial translation. Immunoblotting on COMI cells after 48 hours of treatment with increasing concentrations of **(16*S*)-1e**, and **(16*S*)-2e**, alone (*left column*) or in combination with Q (**3**) (*right column*). Effects on COX1, COX4 and beta-tubulin proteins are shown. One representative result, n=3 biological replicates.

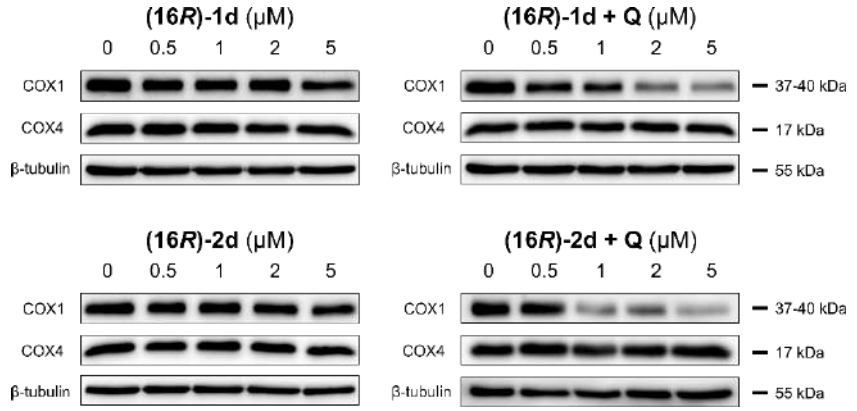


Fig. S8. Western blot analysis for the evaluation of mitochondrial translation inhibition of (16*R*) NHMe derivatives. (16*R*) NHMe derivatives do not inhibit mitochondrial translation. Immunoblotting on COMI cells after 48 hours of treatment with increasing concentrations of **(16*R*)-1d**, and **(16*R*)-2d**, alone (*left column*) or in combination with Q (**3**) (*right column*). Effects on COX1, COX4 and beta-tubulin proteins are shown. One representative result, n=3 biological replicates.

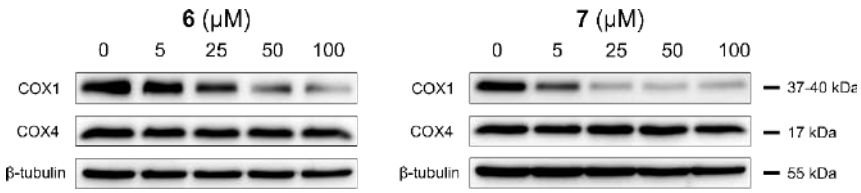


Fig. S9. Western blot analysis for the evaluation of mitochondrial translation inhibition of **6** and **7** derivatives. Compounds **6** and **7** inhibit mitochondrial translation only at high concentrations. Immunoblotting on COMI cells after 48 hours of treatment with increasing concentrations of **6** and **7**. Effects on COX1, COX4 and beta tubulin proteins are shown. One representative result, n=3 biological replicates.

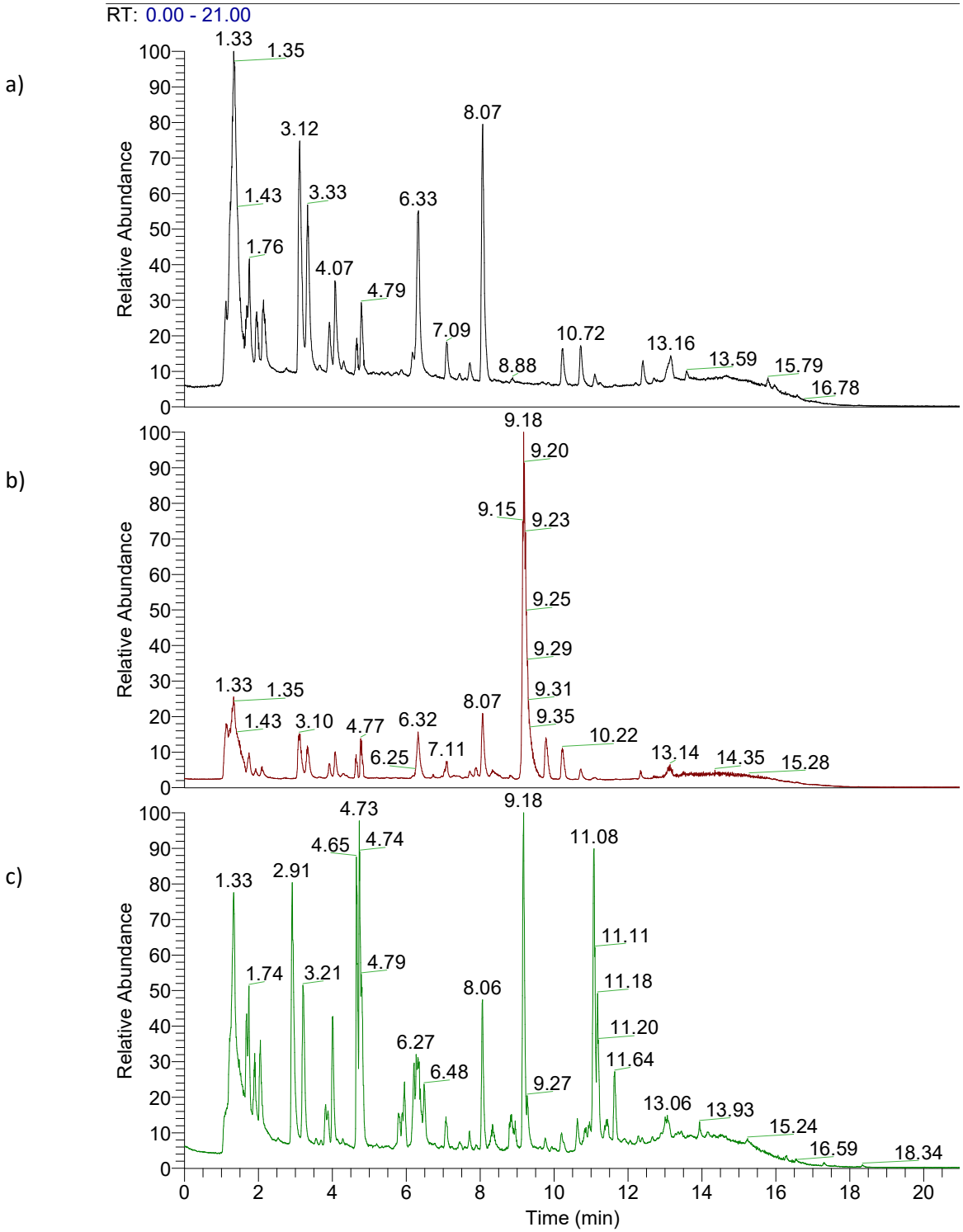


Fig. S10. UHPLC chromatograms of a) drug-free sample (blank); b) standard sample used for calibration curve of VM1 (**2**) (50 ng/mL); c) sample treated with VM1 (**2**). UHPLC conditions: Hypersil Gold C18 column (Thermofisher, 100 x 2.1 mm, particle size: 1.9 µm); linear gradient from 5 to 100% B (B: acetonitrile 0.1% of formic acid; A: water + 0.1% formic acid) over 15 min, followed by 4 min at 100%; flow 0.2 mL/min; MS spray voltage 3500 V; full scan at 120.000 FWHM (200 m/z); scan range 100-1000 m/z.

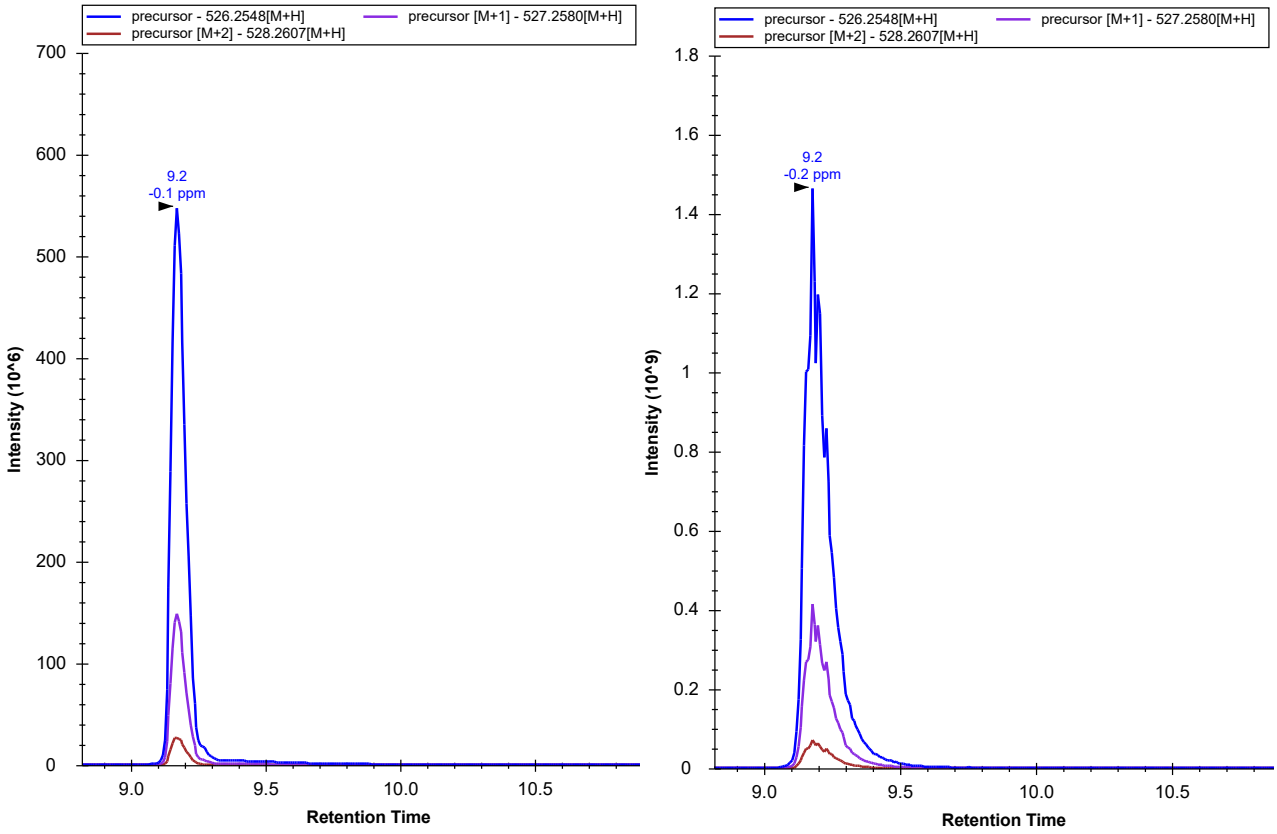


Fig. S11. Extracted ion chromatograms of the $[M+H]^+$ adducts (Skyline software) of a representative standard sample of VM1 (**2**) conc. 50 ng/mL (on the left) and sample treated with VM1 (**2**) (on the right). The peak area of the precursor ions were extracted from chromatograms for both standards and treated samples to build the standard curves of each compound and to compute concentration values of the compound in samples, respectively.

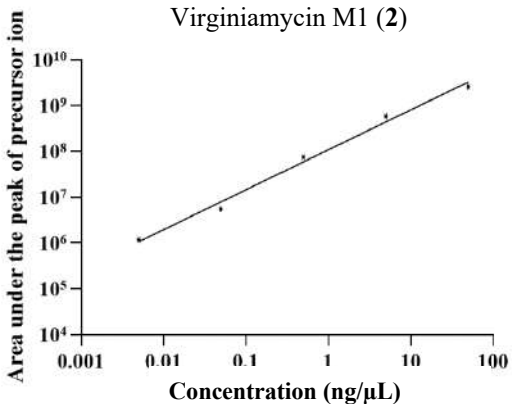


Fig. S12. Calibration curve for VM1 (**2**); $R^2 = 0.9917$. Five serial drug dilutions, from 0.005 to 50 ng/µL, were prepared in non-treated metabolite extracts resuspended in 5% acetonitrile/0.1% formic acid aqueous solution starting from a 10 mM stock of VM1 (**2**). 20 µL of each dilution plus a blank samples were loaded into the LC autosampler. After the analysis, the log-transformed value of the total area of precursor ions of each standard sample was plotted in function of the concentration to construct the standard curve. The error on concentrations was calculated as the physical error deriving from the weighting of the compound plus the error from pipetting. For each serial dilution performed, the error was propagated using the variance formula.

Table S1
UHPLC-MS retention time (min) returned by Skyline software for extracted peak of each compound.

	RT		RT
D (1)	6.9	**VM1 (2)**	9.2
1a	6.6	**2a**	8.6
(16*R*)-1c	6.6	**(16*R*)-2c**	8.3
(16*R*)-1e	7.3	**(16*R*)-2e**	9.3
5	7.0		
6	6.2		

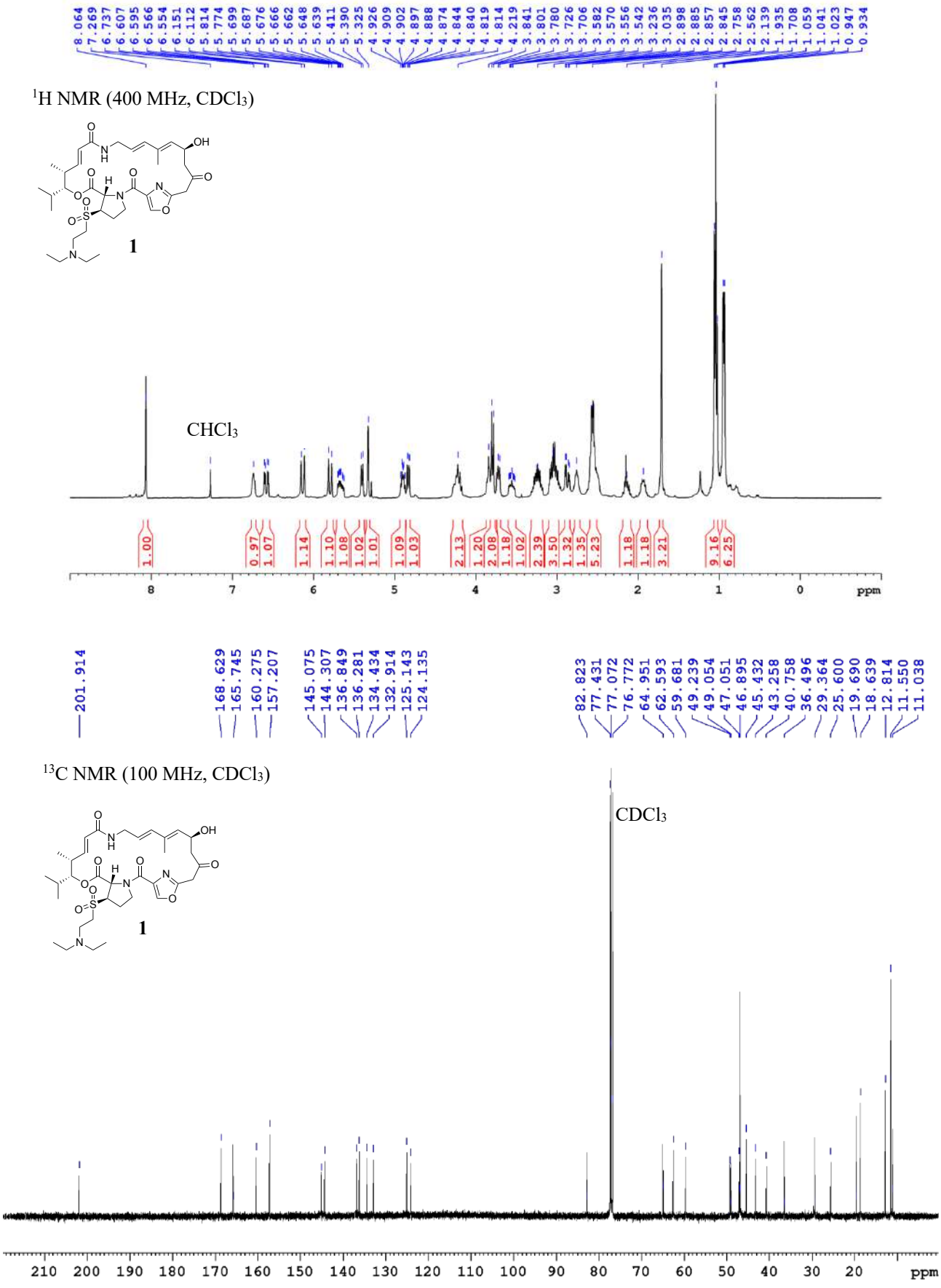


Fig. S13. ^{1}H NMR and ^{13}C NMR spectra of compound **1**.

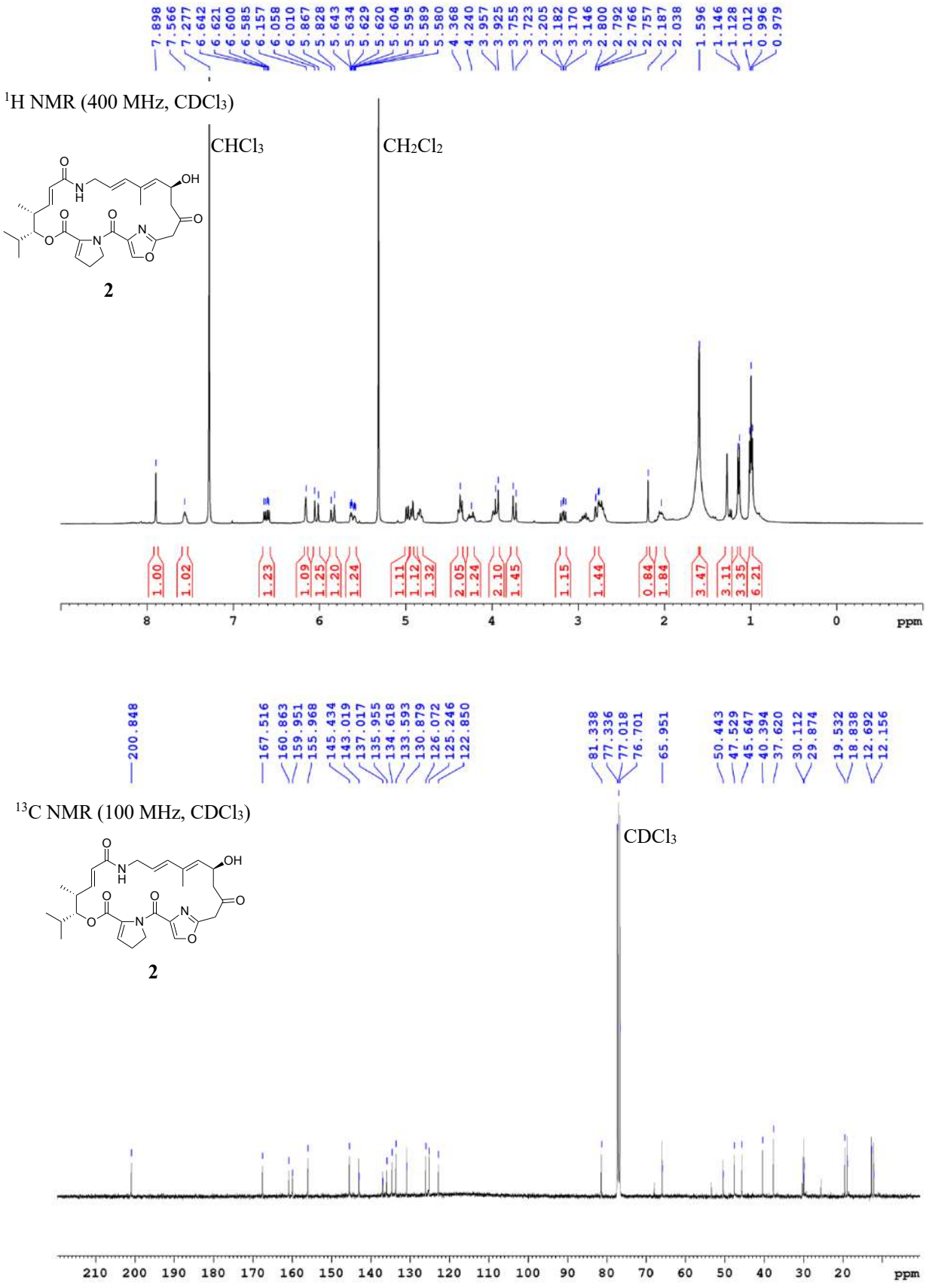


Fig. S14. ^{1}H NMR and ^{13}C NMR spectra of compound **2**.

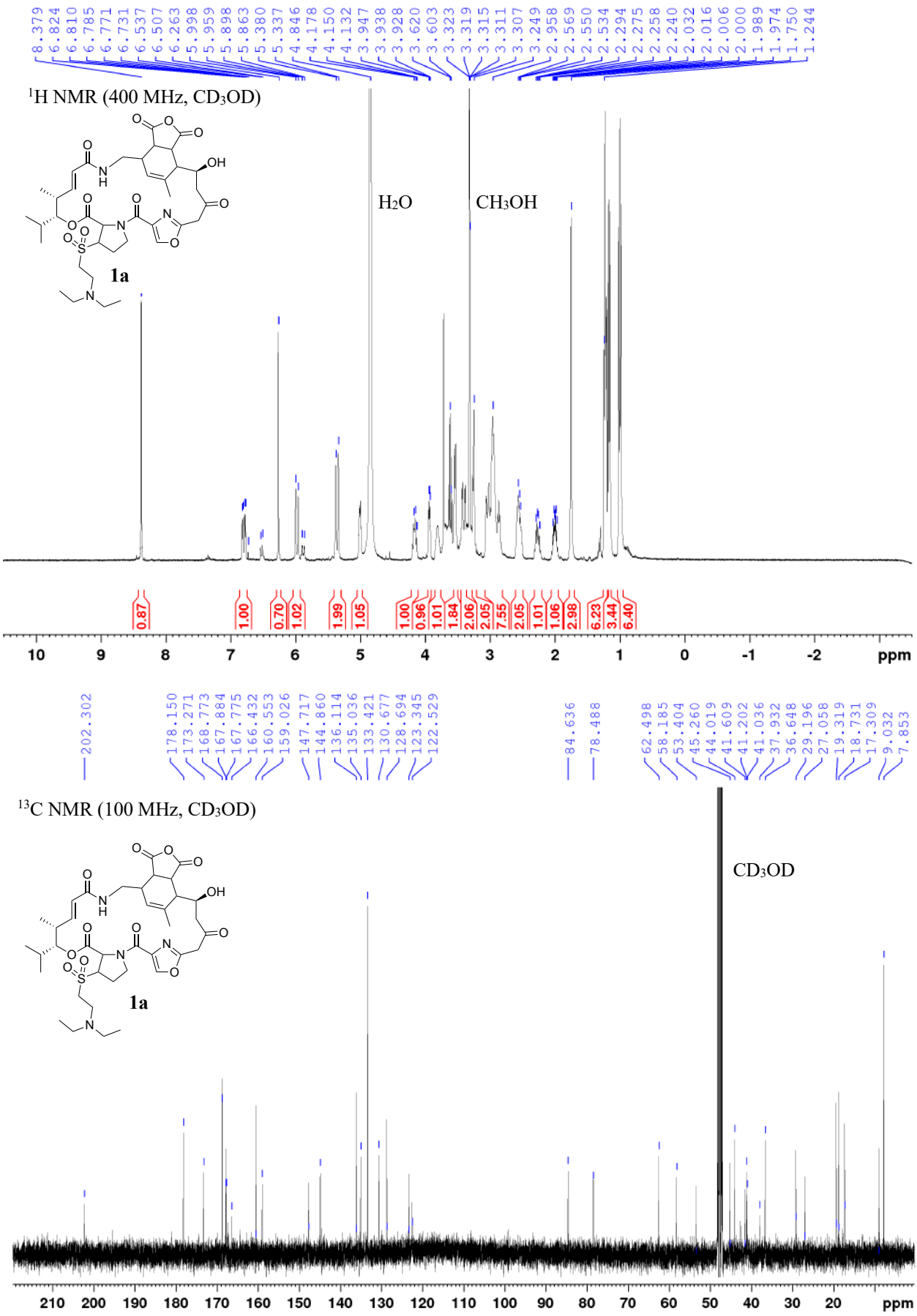


Fig. S15. ^{1}H NMR and ^{13}C NMR spectra of compound **1a**.

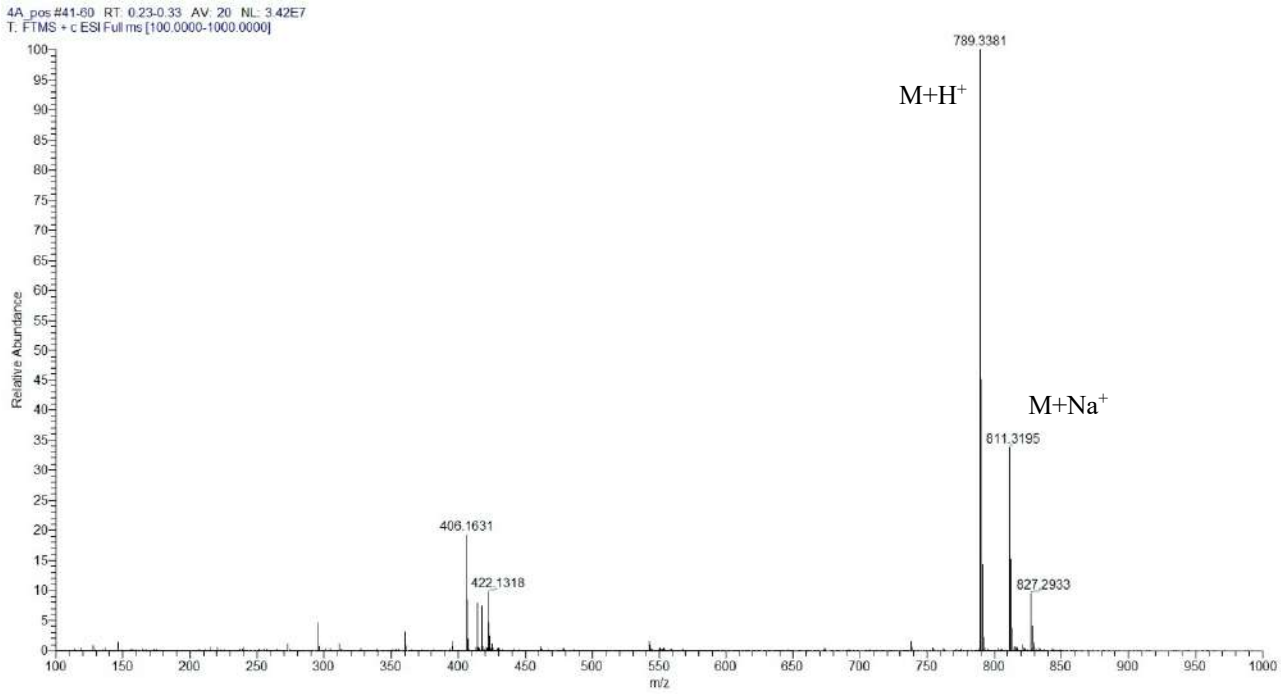
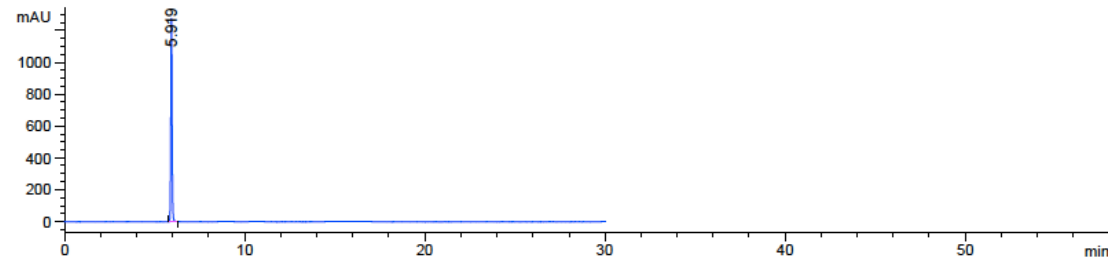


Fig. S16. High resolution ESI(+) MS spectrum of compound **1a**.



Signal 1: DAD1 A, Sig=210,16 Ref=off

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	5.919	BB	0.0771	6367.26514	1277.52002	100.0000
Totals :				6367.26514	1277.52002	

Fig. S17. HPLC chromatogram of compound **1a** (RP18 column, isocratic water/acetonitrile 30:70 + 1% trifluoroacetic acid).

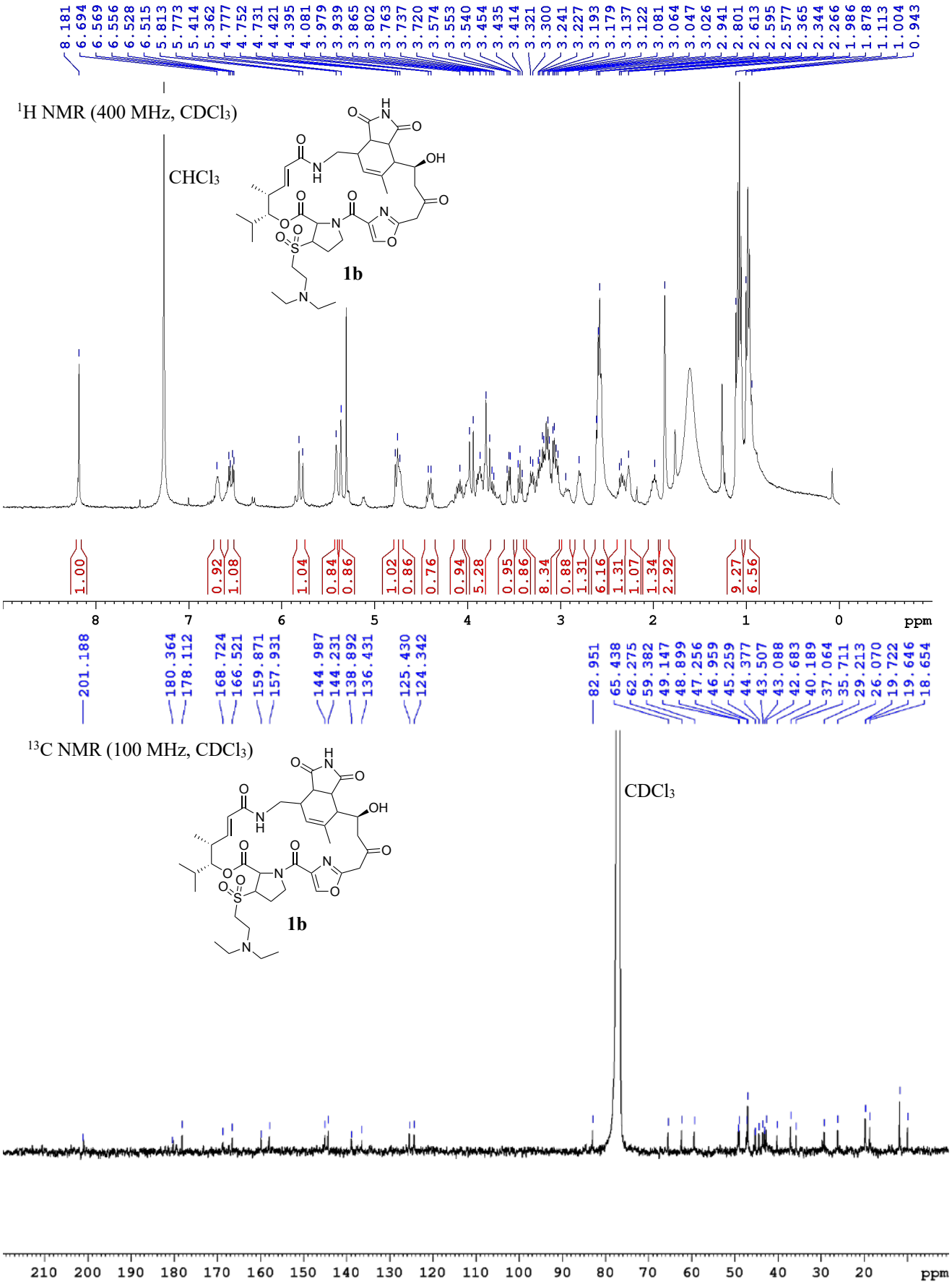


Fig. S18. ^{1}H NMR and ^{13}C NMR spectra of compound **1b**.

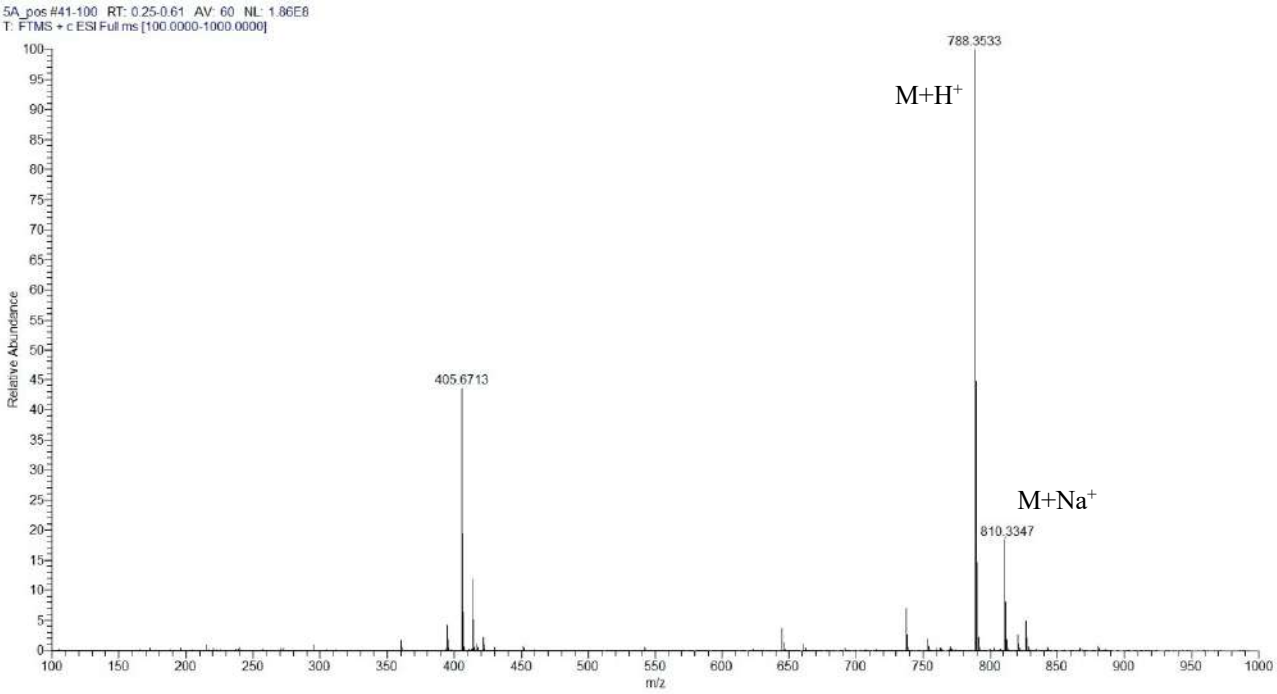
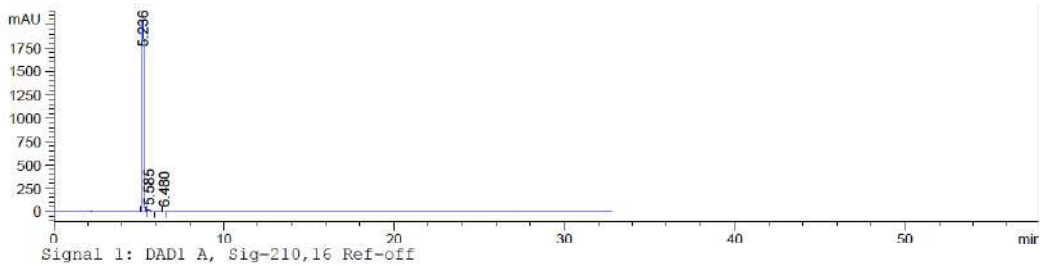


Fig. S19. High resolution ESI(+) MS spectrum of compound **1b**.



Signal 1: DAD1 A, Sig=210,16 Ref=off

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	5.236	BV	0.1163	1.48404e4	2053.51514	98.4657
2	5.585	VB	0.1549	212.02127	20.12010	1.4068
3	6.480	BV	0.0805	19.21877	3.47546	0.1275
Totals :				1.50717e4	2077.11069	

Fig. S20. HPLC chromatogram of compound **1b** (RP18 column, isocratic water/acetonitrile 30:70 + 1% trifluoroacetic acid).

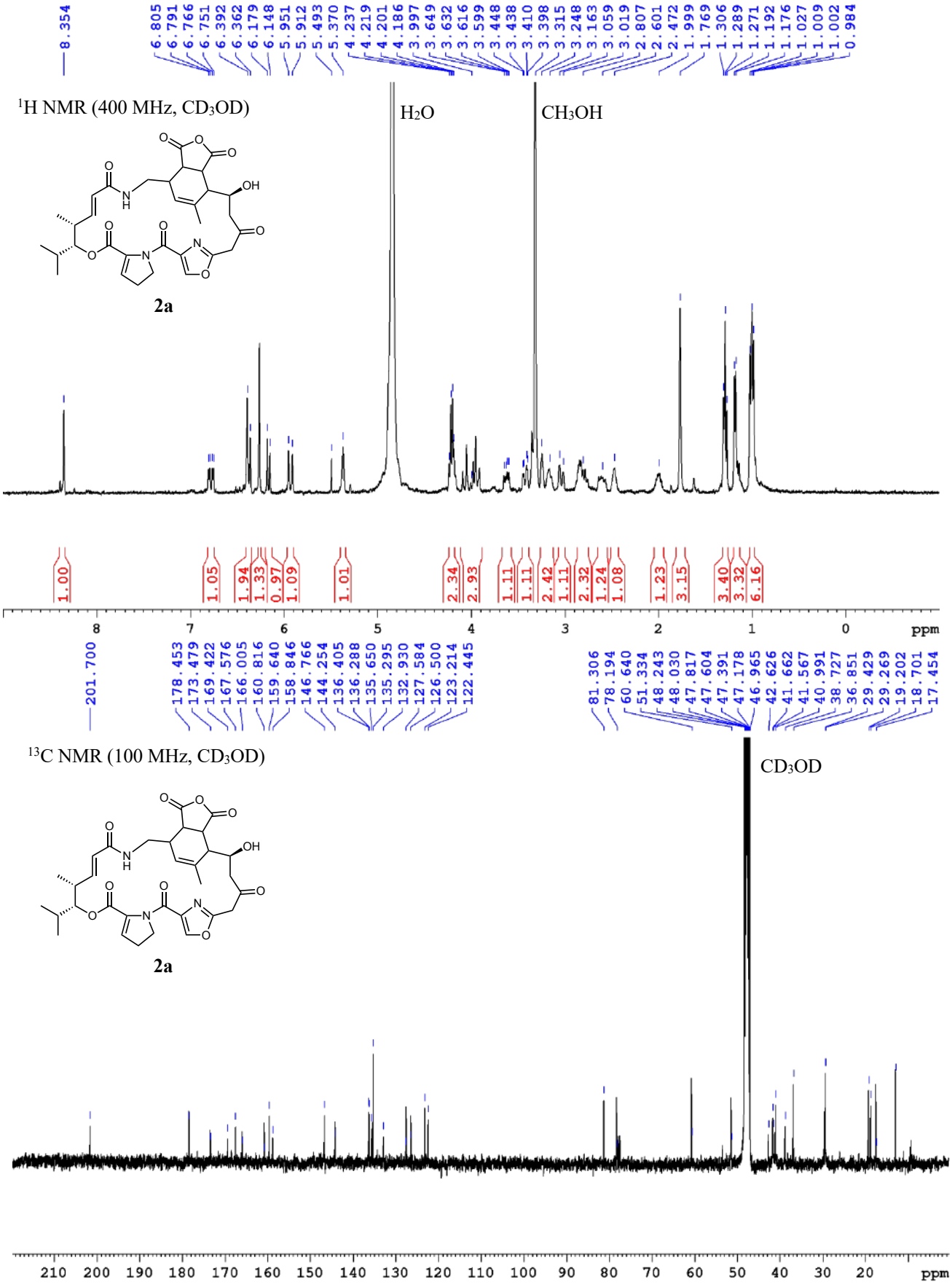


Fig. S21. ^{1}H NMR and ^{13}C NMR spectra of compound **2a**.

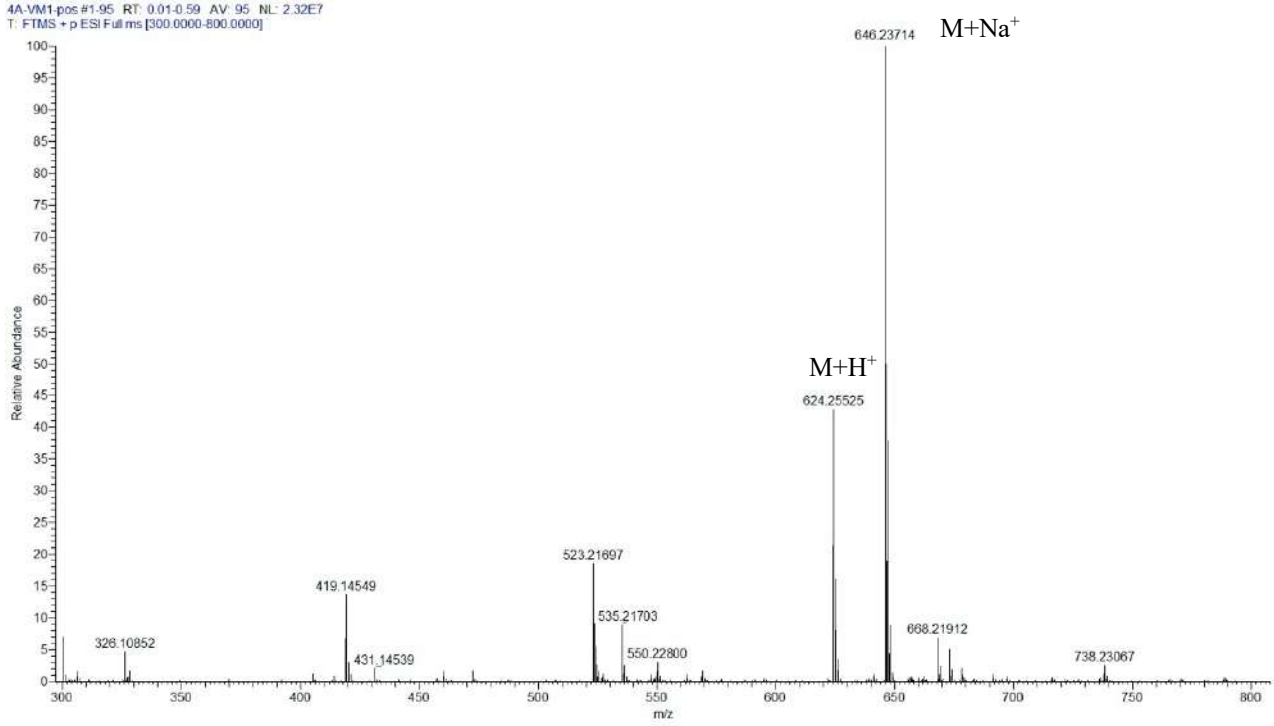
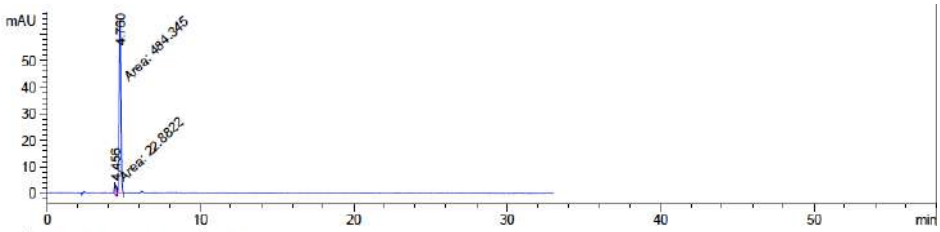


Fig. S22. High resolution ESI(+) MS spectrum of compound **2a**.



Signal 2: DAD1 B, Sig=254,16 Ref=off

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	4.456	MM	0.0997	22.88224	3.82557	4.5112
2	4.760	MM	0.1241	484.34534	65.03886	95.4888
Totals :				507.22758	68.86443	

Fig. S23. HPLC chromatogram of compound **2a** (RP18 column, gradient water/acetonitrile 30:70 to 80:20).

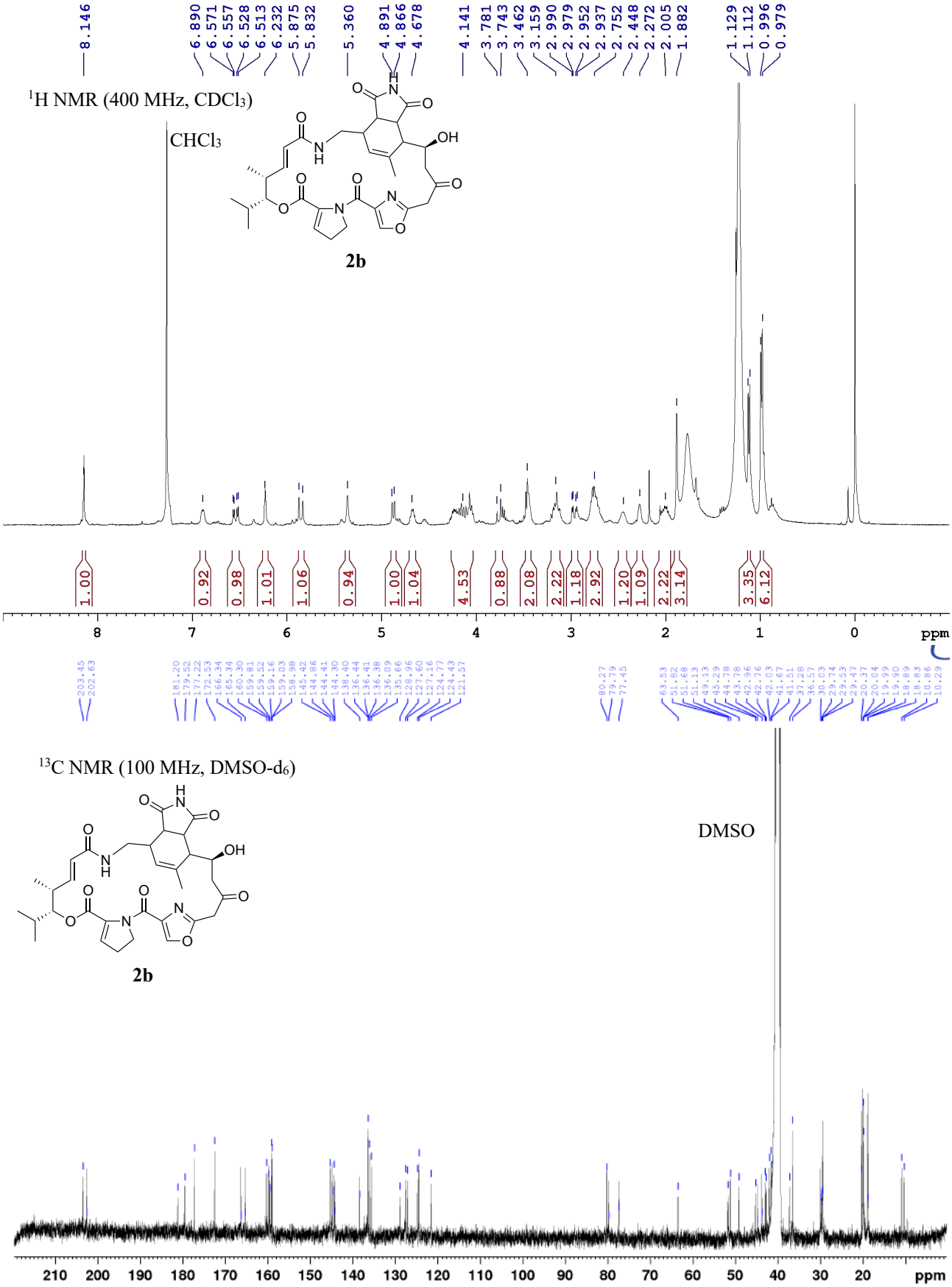


Fig. S24. ^{1}H NMR and ^{13}C NMR spectra of compound **2b**.

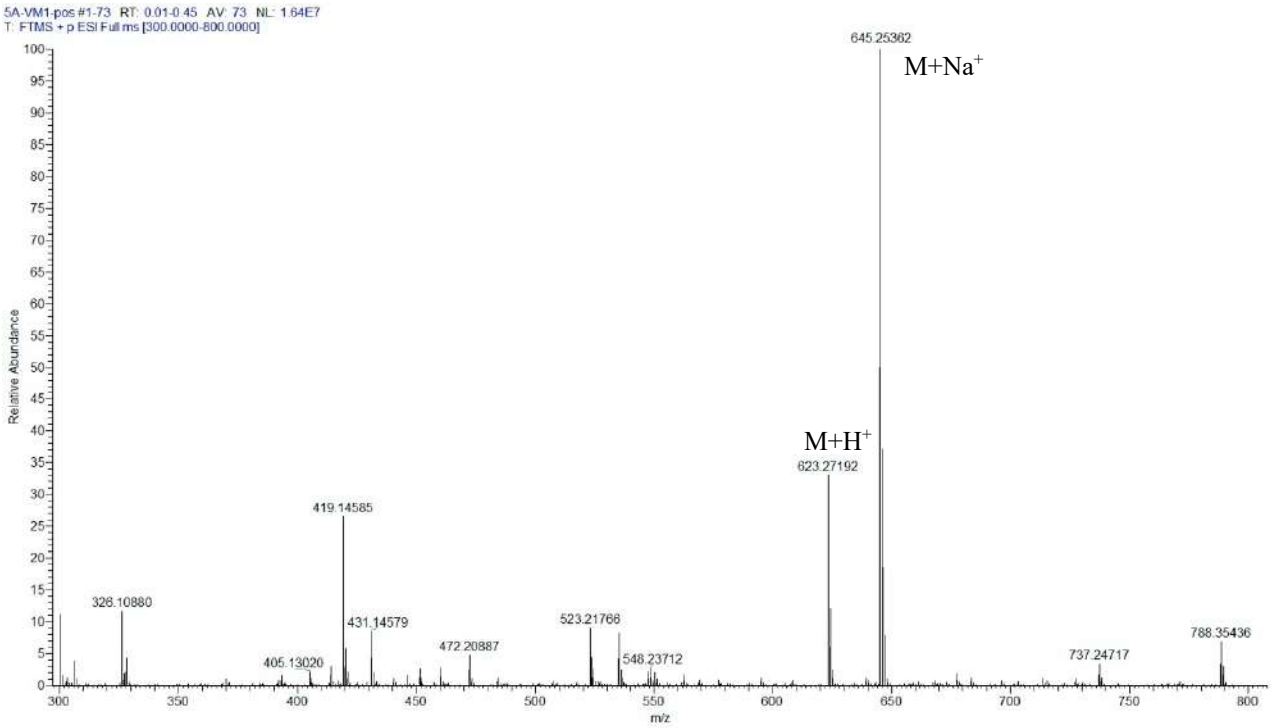
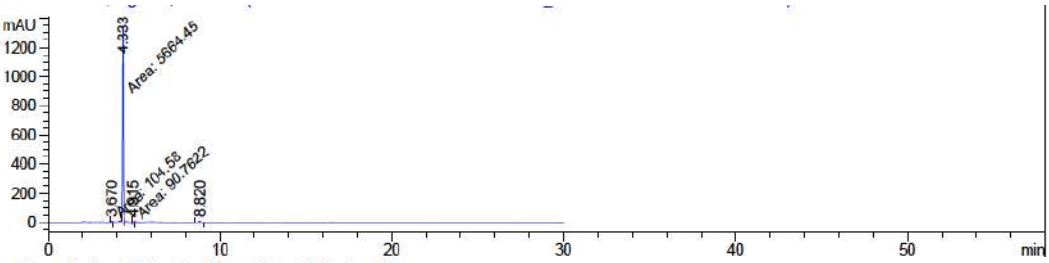


Fig. S25. High resolution ESI(+) MS spectrum of compound **2b**.



Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	3.670	MM	0.1238	104.58009	14.07907	1.7640
2	4.333	MM	0.0697	5664.45410	1355.36768	95.5474
3	4.915	MM	0.0677	90.76216	19.46412	1.5310
4	8.820	VV	0.1244	68.62528	8.30189	1.1576

Fig. S26. HPLC chromatogram of compound **2b** (RP18, gradient water/acetonitrile 30:70 to 80:20).

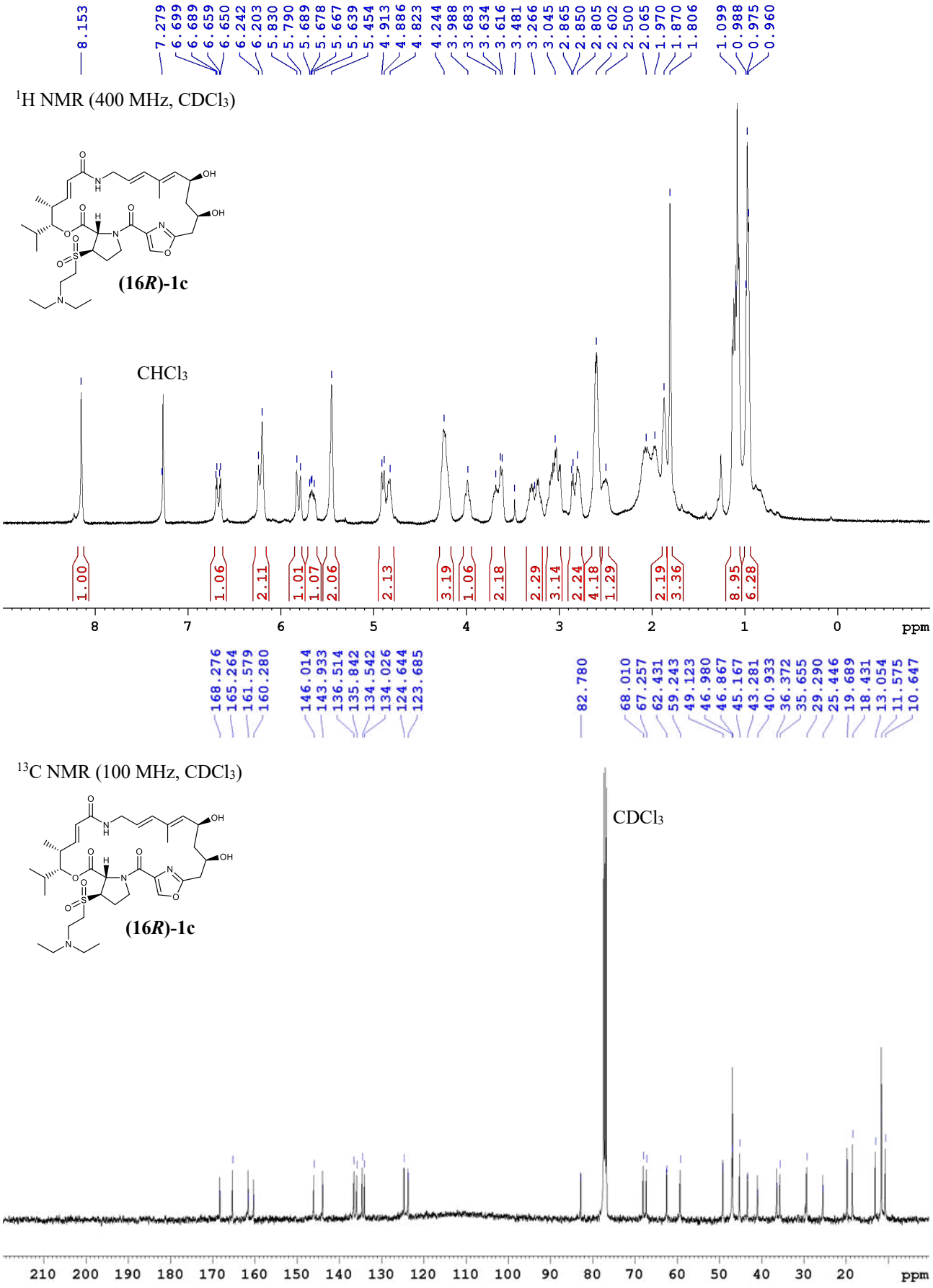


Fig. S27. ^{1}H NMR and ^{13}C NMR spectra of compound **(16*R*)-1c**.

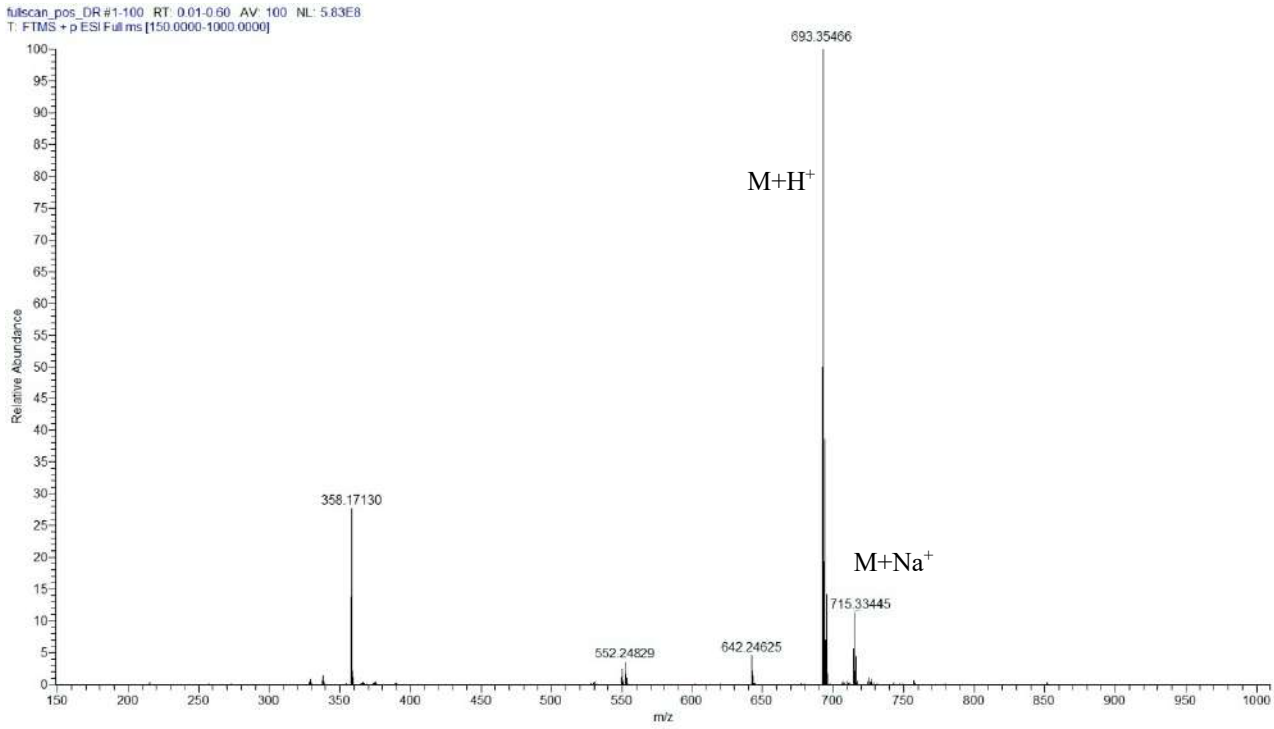
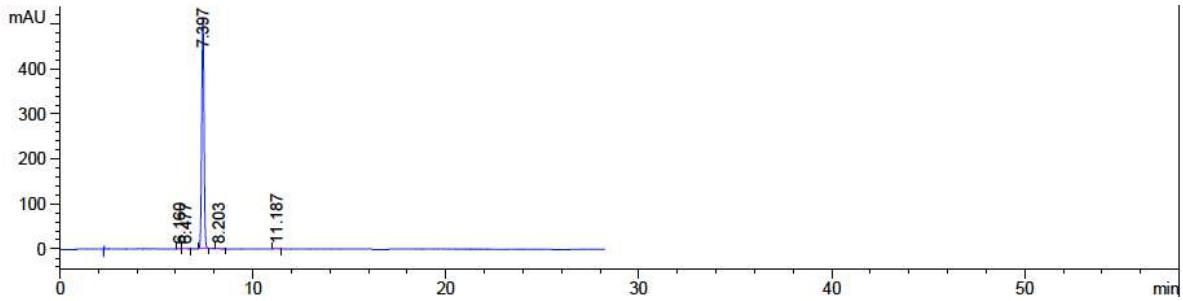


Fig. S28. High resolution ESI(+) MS spectrum of compound **(16*R*)-1c**.



Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	6.160	BV	0.1187	12.32514	1.27571	0.2819
2	6.477	VB	0.1331	15.07991	1.36264	0.3449

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
3	7.397	BB	0.1313	4309.53516	511.31335	98.5622
4	8.203	VB	0.1648	17.57674	1.48805	0.4020
5	11.187	BB	0.1654	17.88631	1.44462	0.4091

Fig. S29. HPLC chromatogram of compound **(16*R*)-1c** (RP18, isocratic water/acetonitrile 30:70 + 1% trifluoroacetic acid).

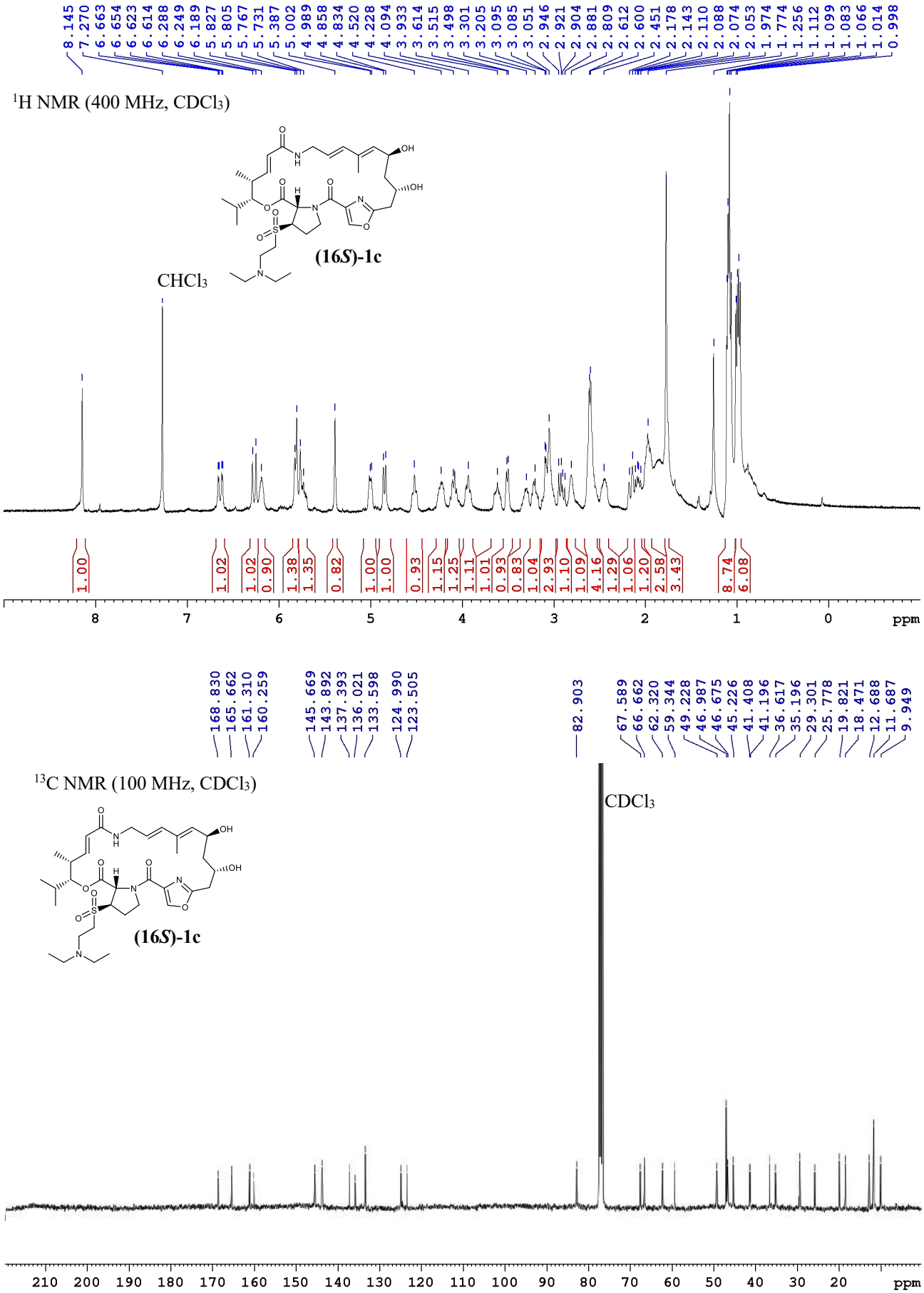


Fig. S30. ^{1}H NMR and ^{13}C NMR spectra of compound **(16*S*)-1c**.

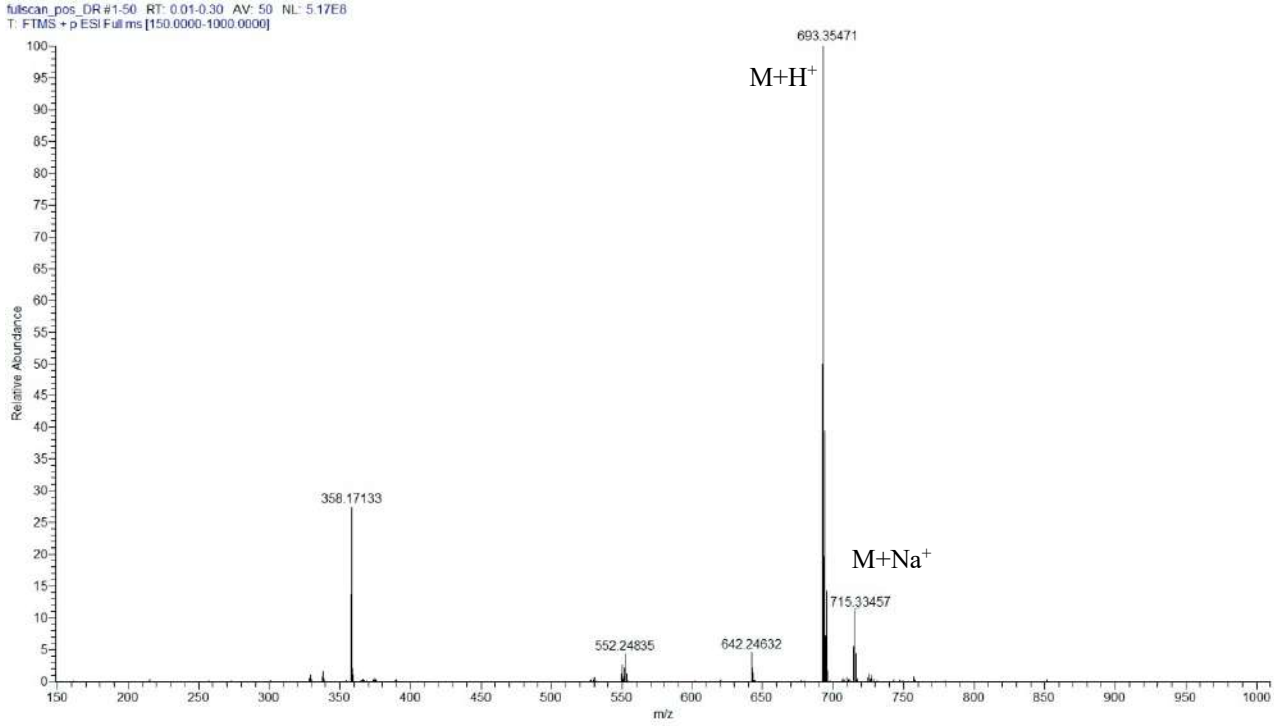
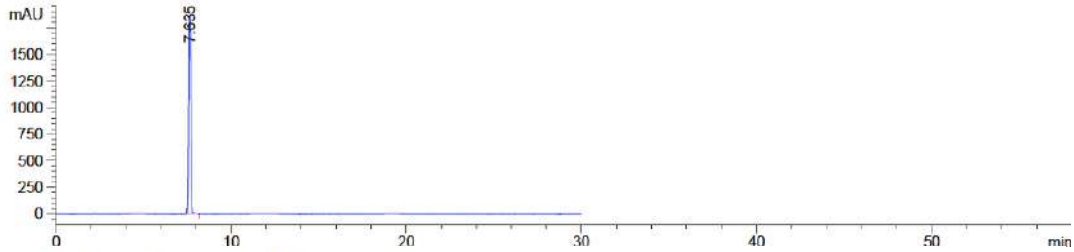


Fig. S31. High resolution ESI(+) MS spectrum of compound **(16*S*)-1c**.



Signal 1: DAD1 A, Sig=210,16 Ref=off

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	7.635	BB	0.1196	1.40293e4	1868.48645	100.0000
Totals :				1.40293e4	1868.48645	

Fig. S32. HPLC chromatogram of compound **(16*S*)-1c** (RP18, isocratic water/acetonitrile 30:70 + 1% trifluoroacetic acid).

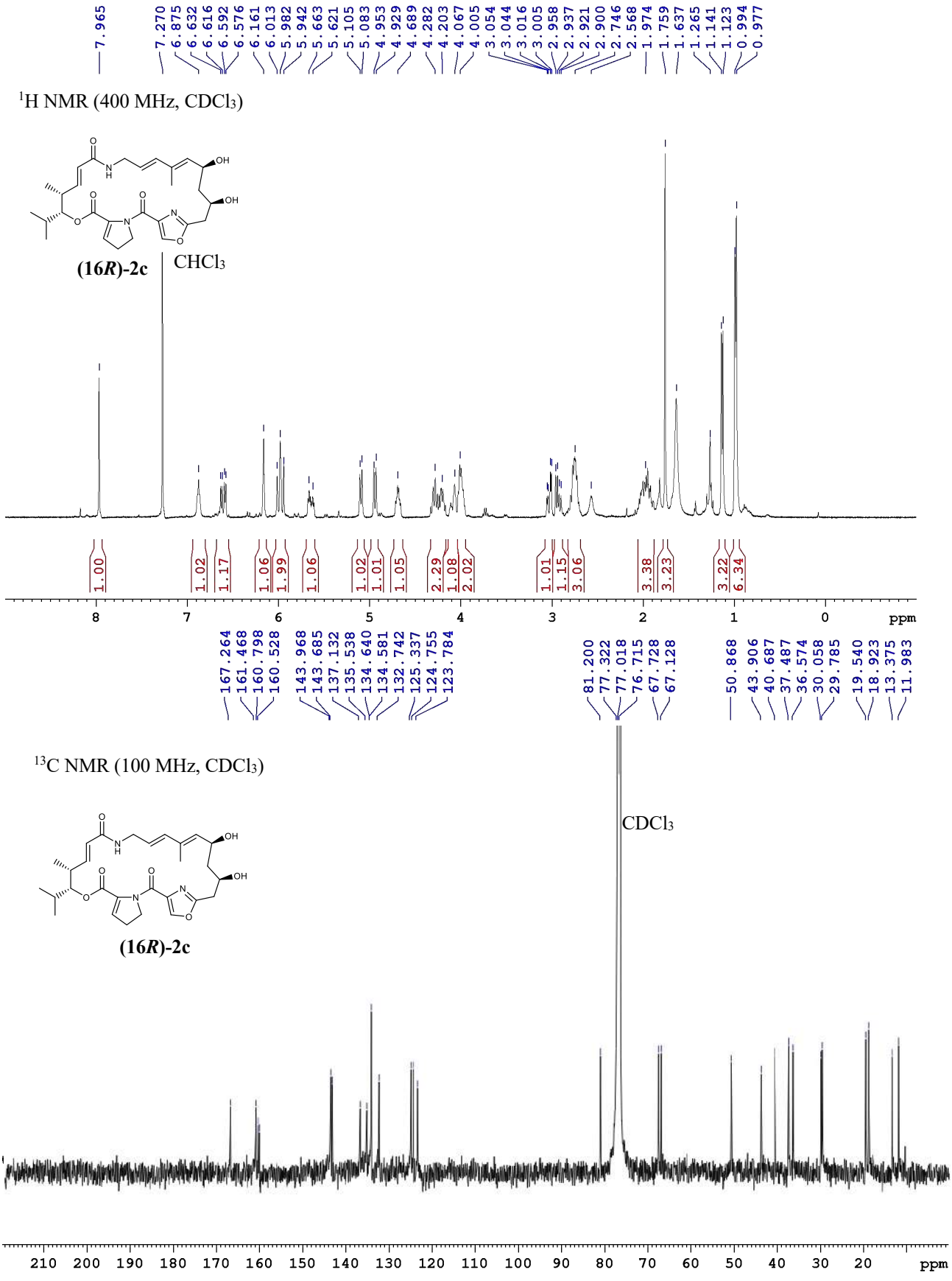


Fig. S33. ^{1}H NMR and ^{13}C NMR spectra of compound **(16*R*)-2c**.

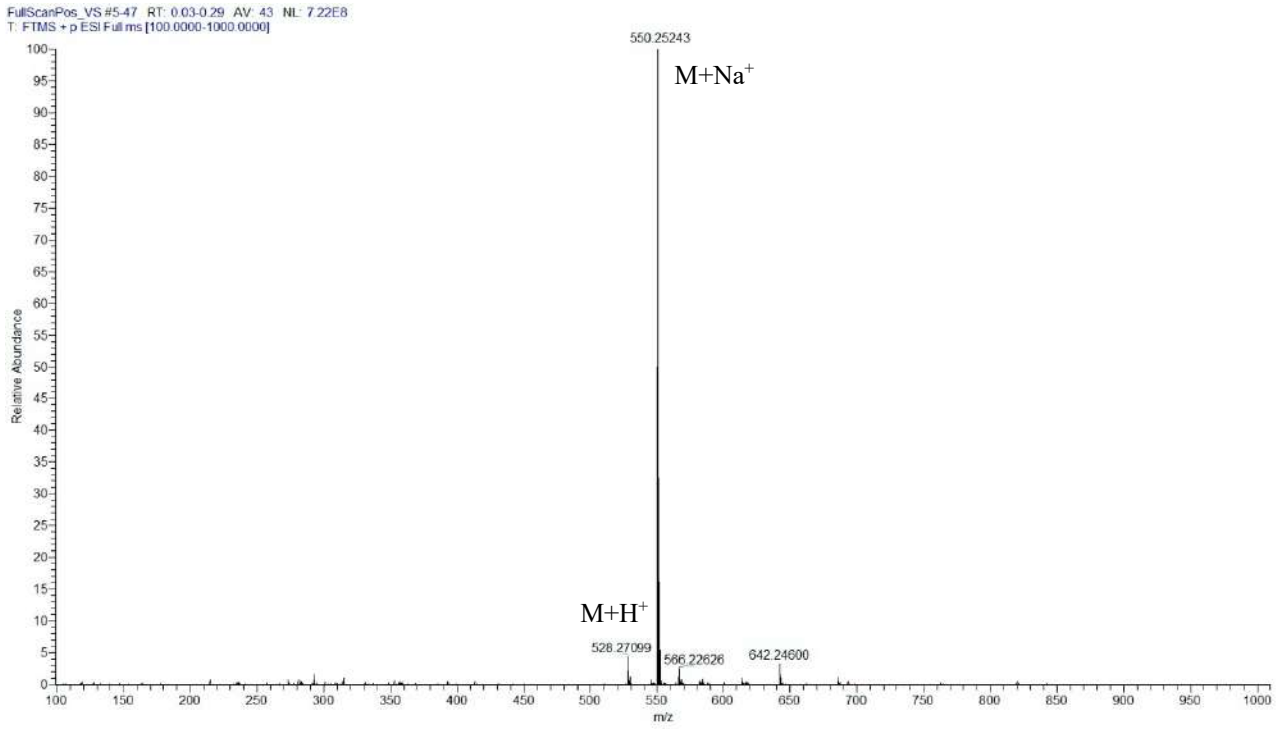
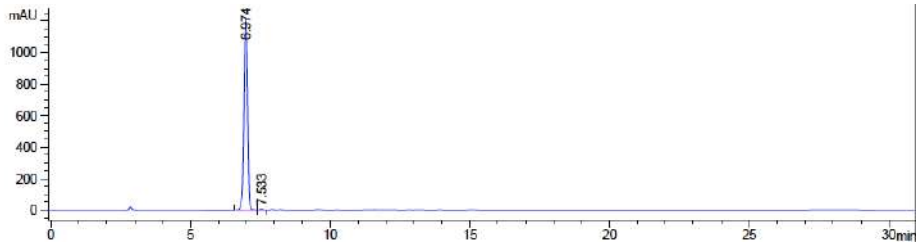


Fig. S34. High resolution ESI(+) MS spectrum of compound **(16*R*)-2c**.



Signal 2: DAD1 B, Sig=254,16 Ref=off

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	6.974	BV	0.1502	1.11277e4	1222.59326	99.5675
2	7.533	VV	0.1482	48.33984	5.41012	0.4325
Totals :				1.11760e4	1228.00338	

Fig. S35. HPLC chromatogram of compound **(16*R*)-2c** (RP18, gradient water/acetonitrile 30:70 to 80:20).

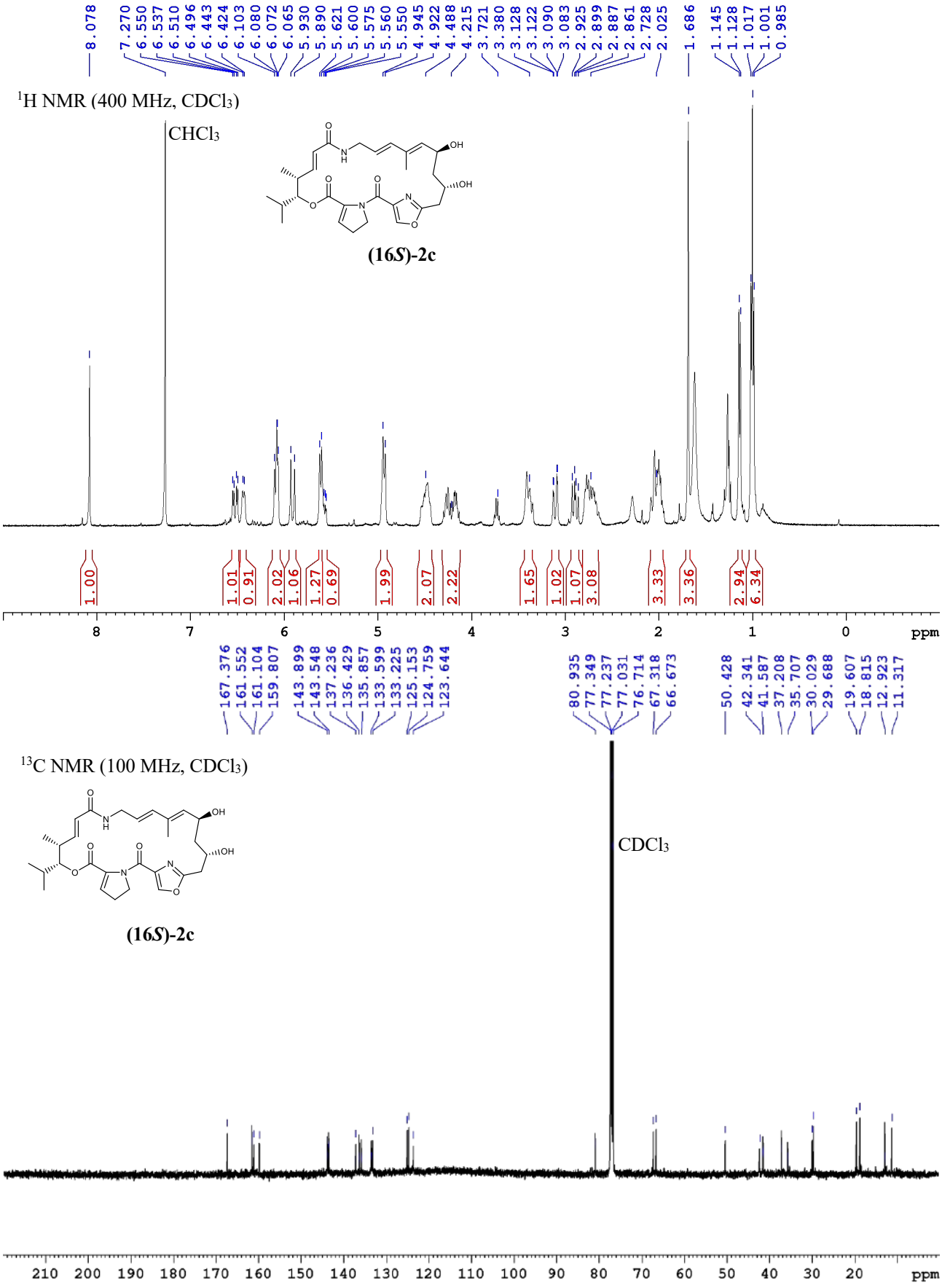


Fig. S36. ^{1}H NMR and ^{13}C NMR spectra of compound **(16*S*)-2c**.

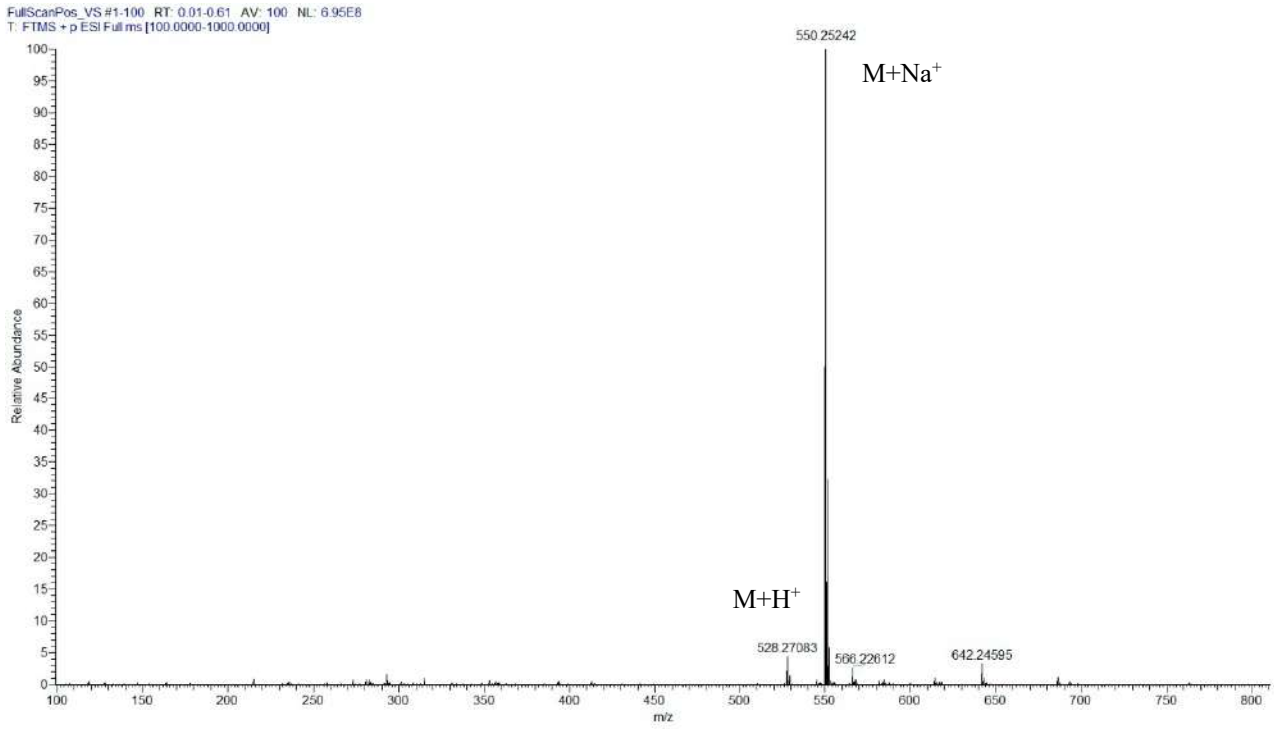
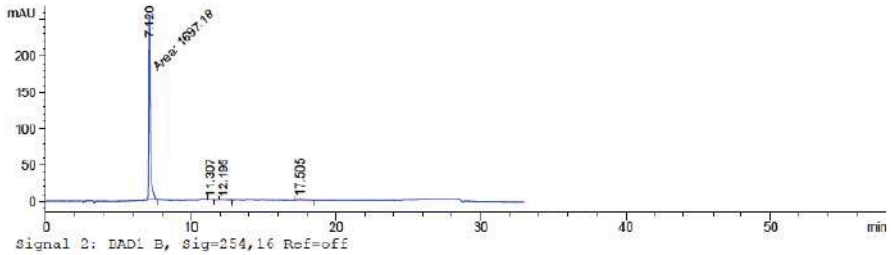


Fig. S37. High resolution ESI(+) MS spectrum of compound **(16*S*)-2c**.



Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	7.120	MM	0.1113	1697.18445	254.08295	96.0207
2	11.307	VV	0.2850	20.30526	8.61591e-1	1.1488
3	12.195	VV	0.4419	22.56616	6.21085e-1	1.2767
4	17.505	BB	0.2892	27.46401	1.36215	1.5538
Totals :				1767.51987	256.92777	

Fig. S38. HPLC chromatogram of compound **(16*S*)-2c** (RP18, gradient water/acetonitrile 30:70 to 80:20).

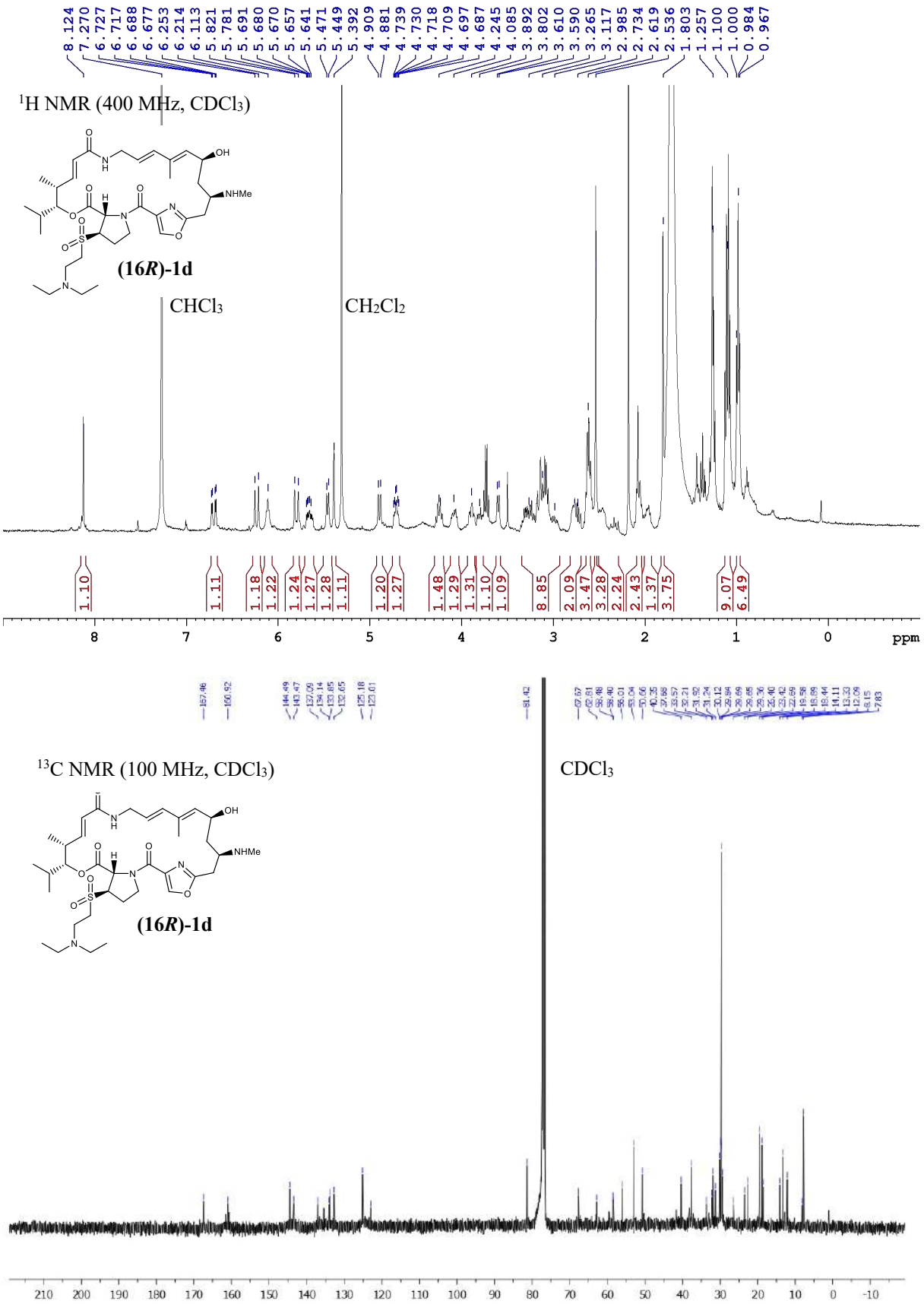


Fig. S39. ^{1}H NMR and ^{13}C NMR spectra of compound **(16*R*)-1d**.

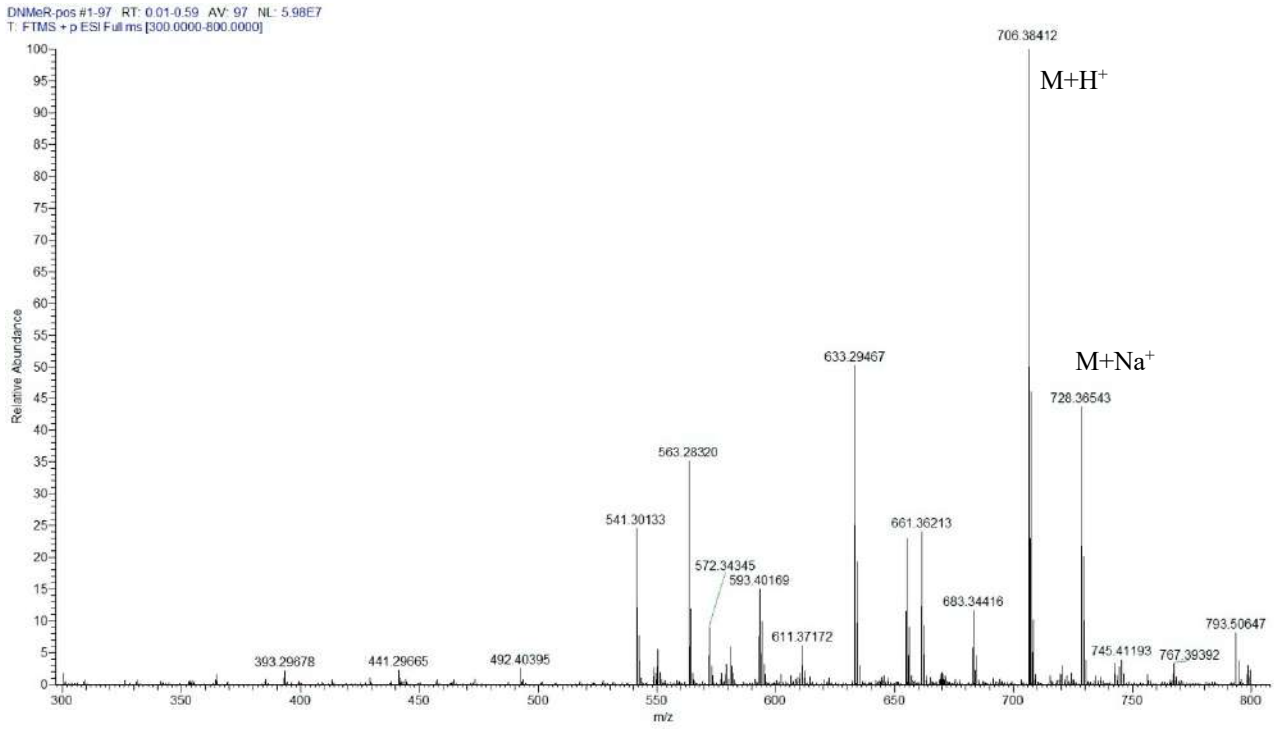
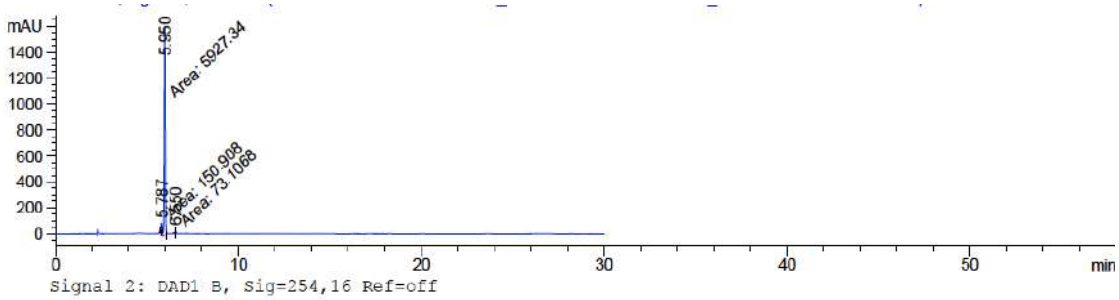


Fig. S40. High resolution ESI(+) MS spectrum of compound **(16*R*)-1d**.



Signal 2: DAD1 B, Sig=254,16 Ref=off

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	5.787	MM	0.0381	150.90755	66.02733	2.4532
2	5.950	MM	0.0612	5927.34229	1615.42639	96.3583
3	6.550	MM	0.0388	73.10684	31.42169	1.1885
Totals :				6151.35667	1712.87541	

Fig. S41. HPLC chromatogram of compound **(16*R*)-1d** (RP18, isocratic water/acetonitrile 30:70 + 1% trifluoroacetic acid).

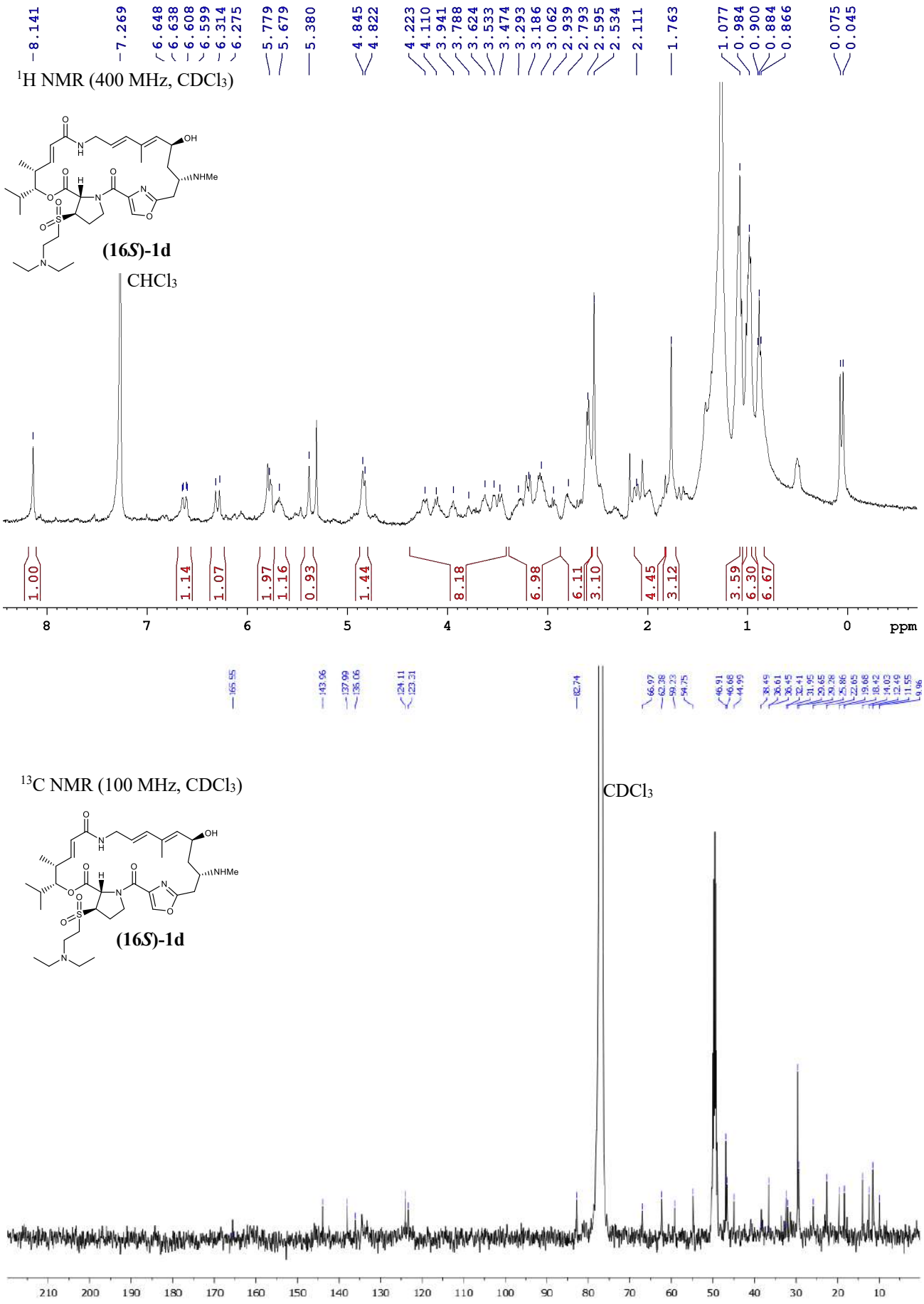


Fig. S42. ^{1}H NMR and ^{13}C NMR spectra of compound **(16*S*)-1d**.

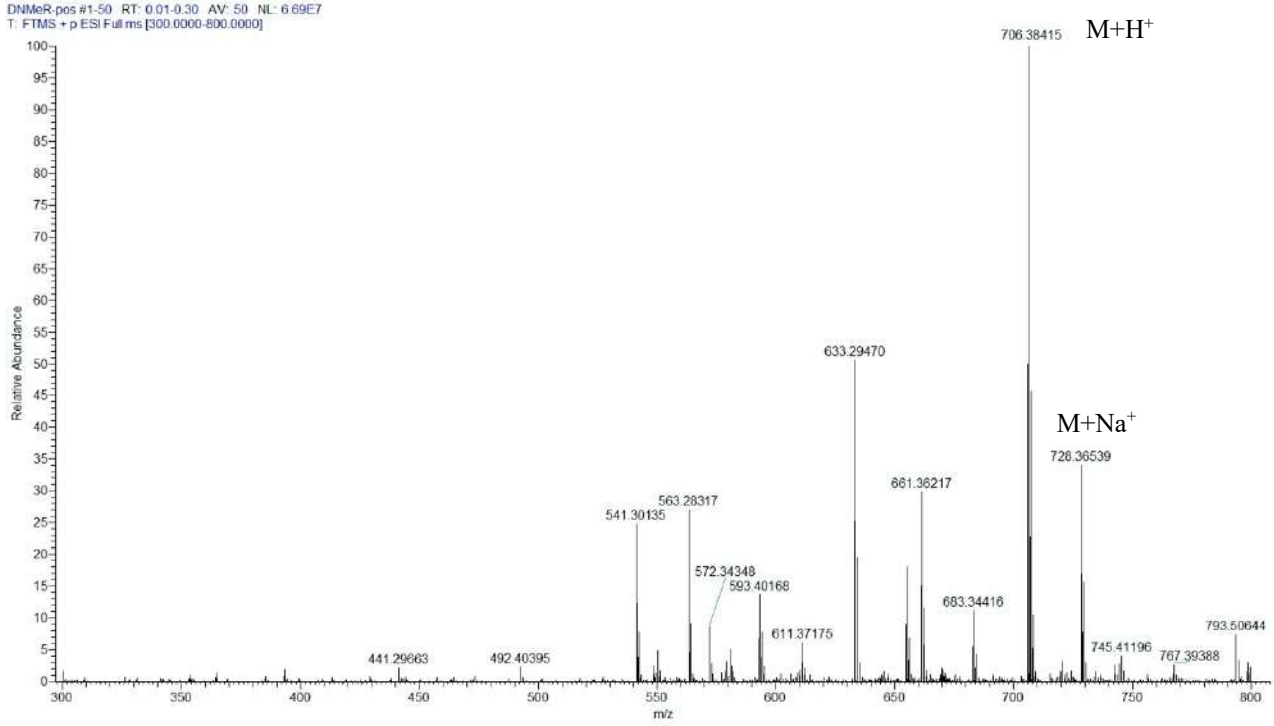
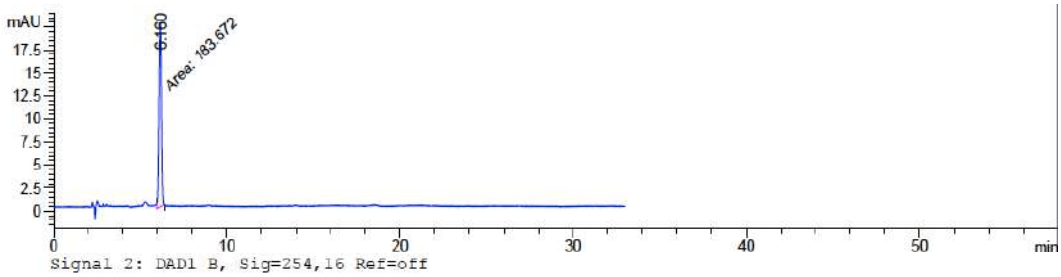


Fig. S43. High resolution ESI(+) MS spectrum of compound **(16*S*)-1d**.



Signal 2: DAD1 B, Sig=254,16 Ref=off

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	6.160	MM	0.1529	183.67215	20.02200	100.0000
Totals :				183.67215	20.02200	

Fig. S44. HPLC chromatogram of compound **(16*S*)-1d** (RP18, isocratic water/acetonitrile 30:70 + 1% trifluoroacetic acid).

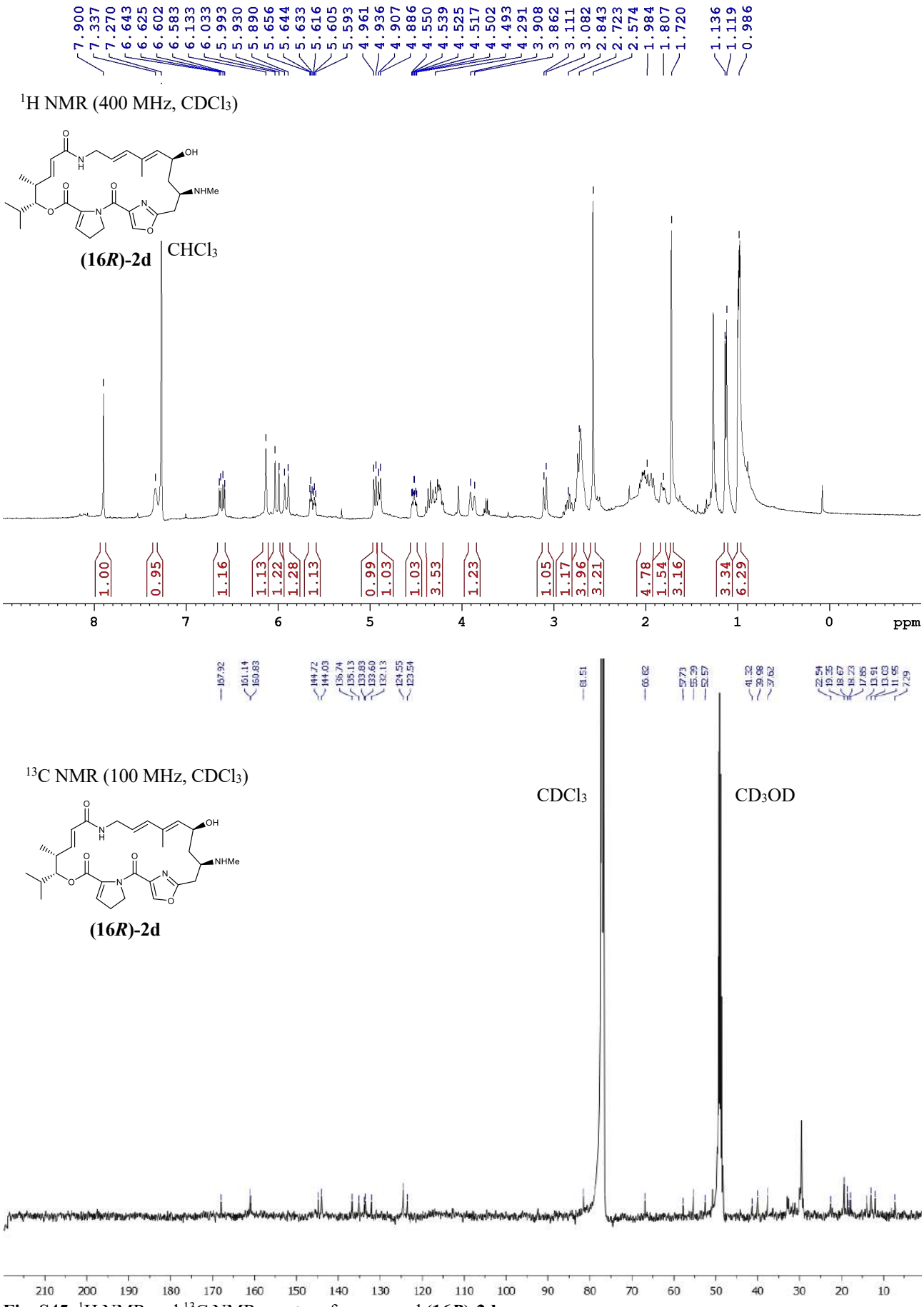


Fig. S45. ^{1}H NMR and ^{13}C NMR spectra of compound **(16*R*)-2d**.

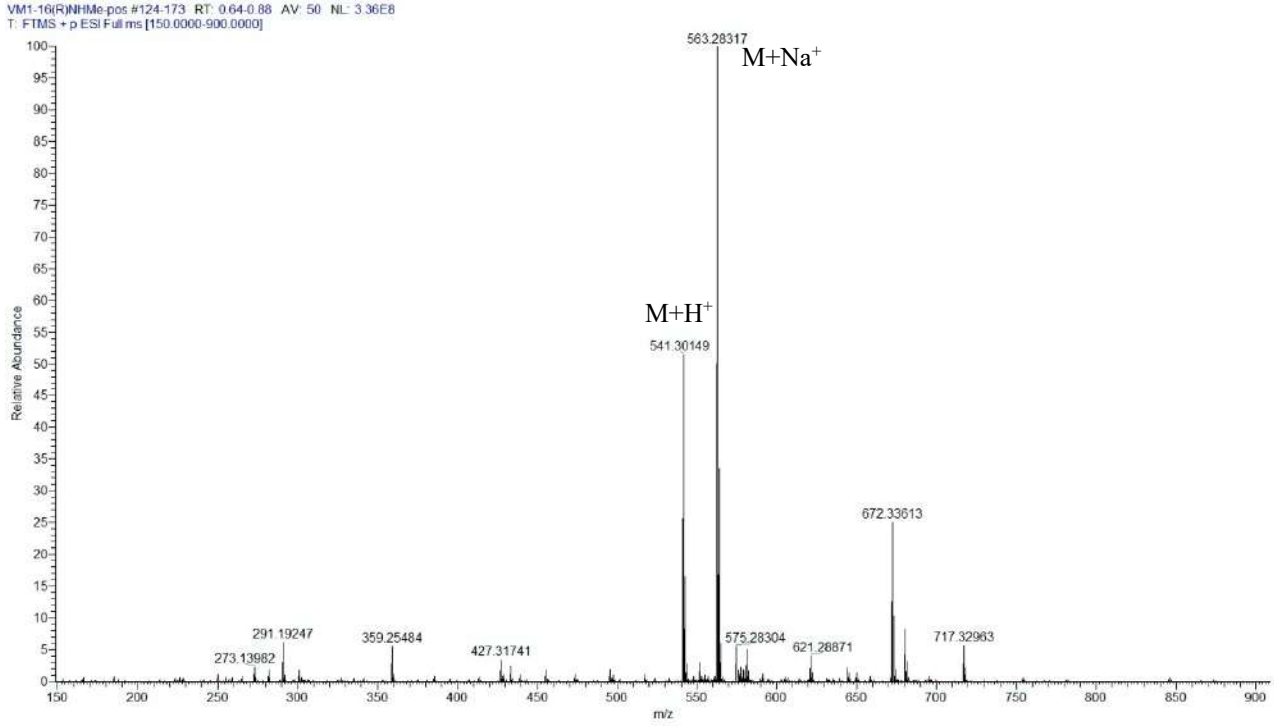
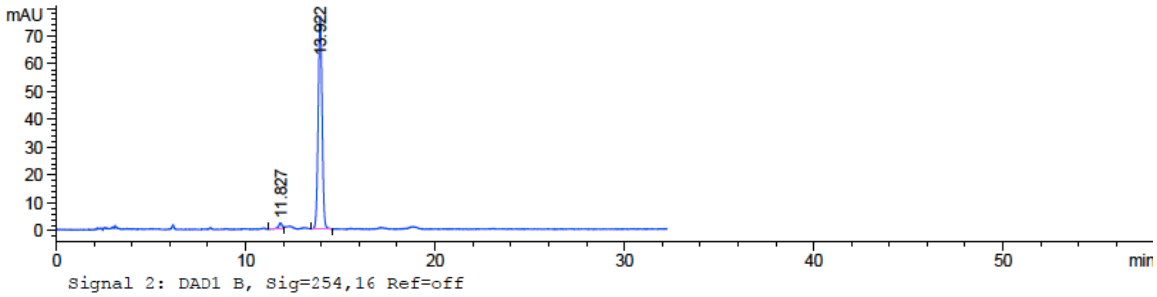


Fig. S46. High resolution ESI(+) MS spectrum of compound **(16*R*)-2d**.



Signal 2: DAD1 B, Sig=254,16 Ref=off

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	11.827	VV	0.2259	35.25024	2.28614	3.0621
2	13.922	VV	0.2262	1115.93811	76.48539	96.9379
Totals :				1151.18835	78.77153	

Fig. S47. HPLC chromatogram of compound **(16*R*)-2d** (RP18, isocratic water/acetonitrile 30:70 + 1% trifluoroacetic acid).

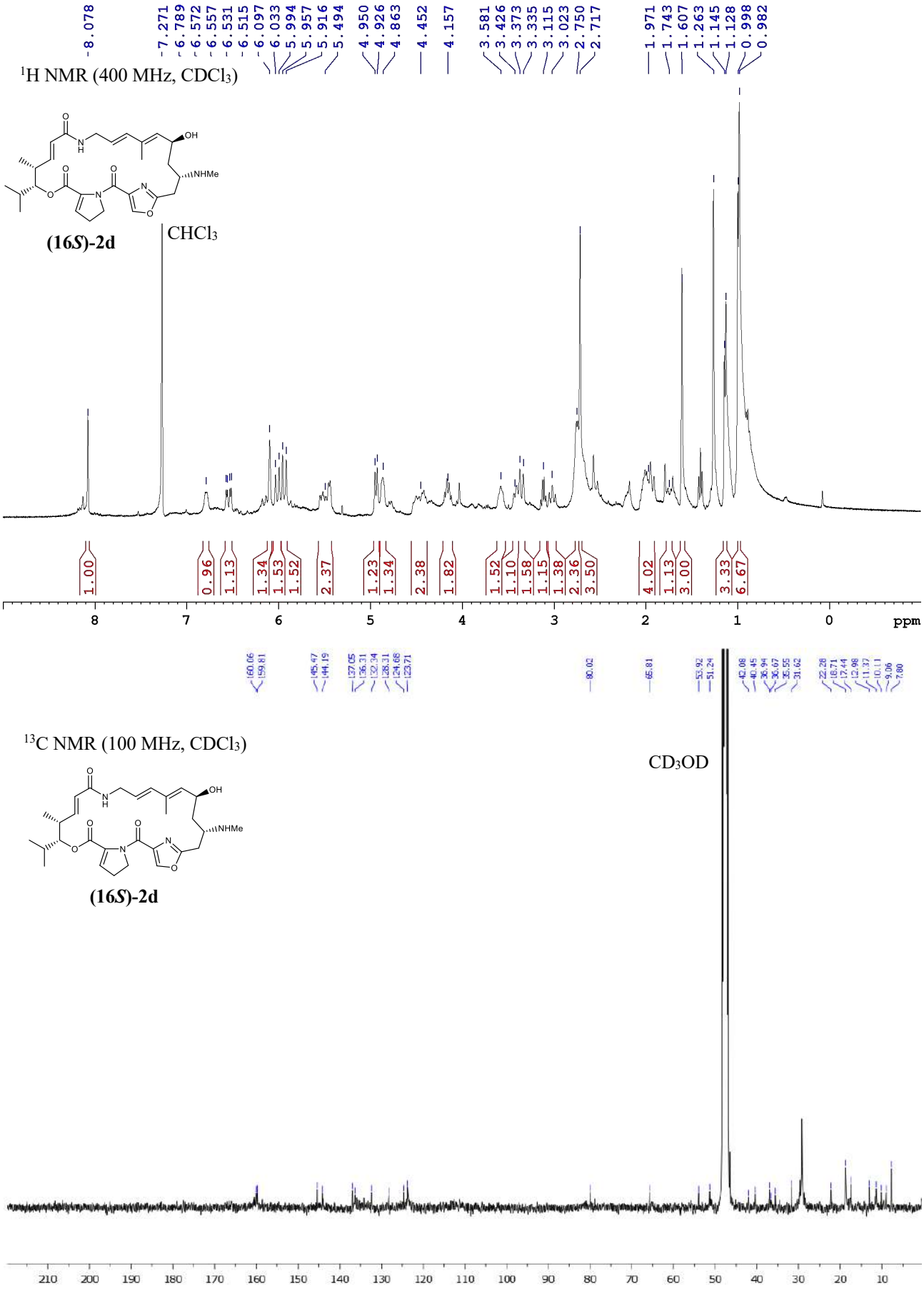


Fig. S48. ^{1}H NMR and ^{13}C NMR spectra of compound **(16*S*)-2d**.

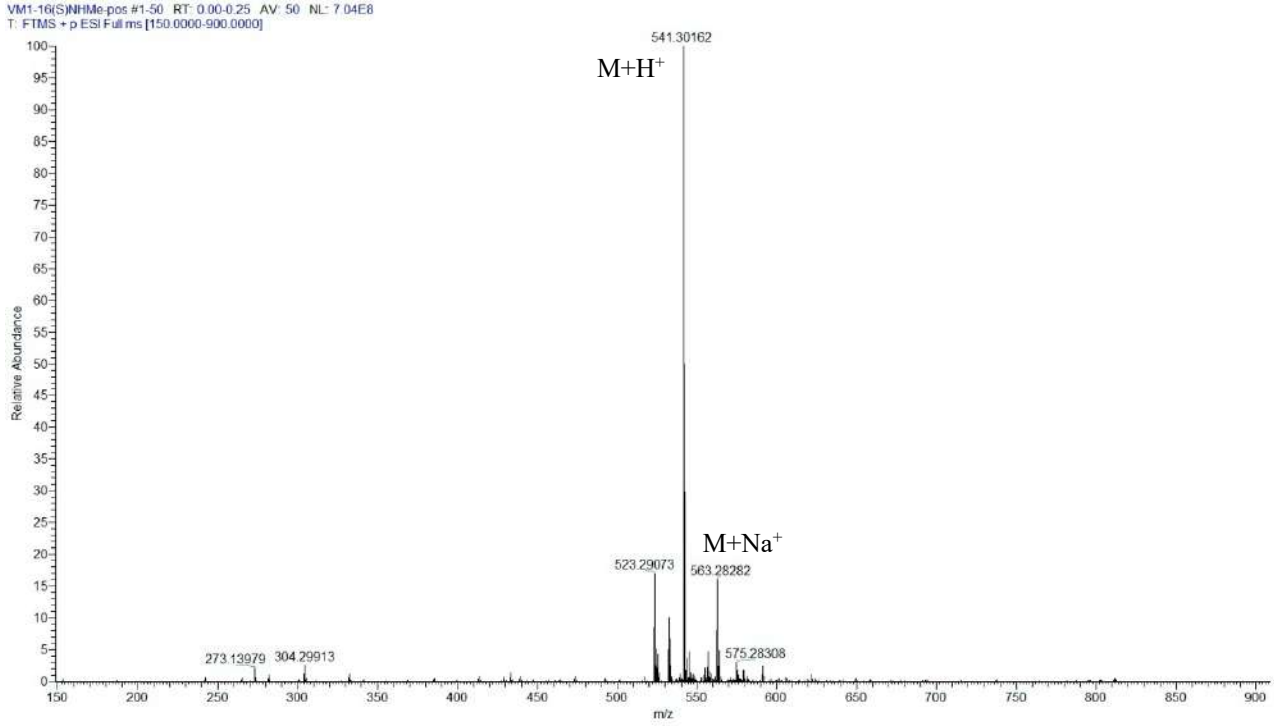
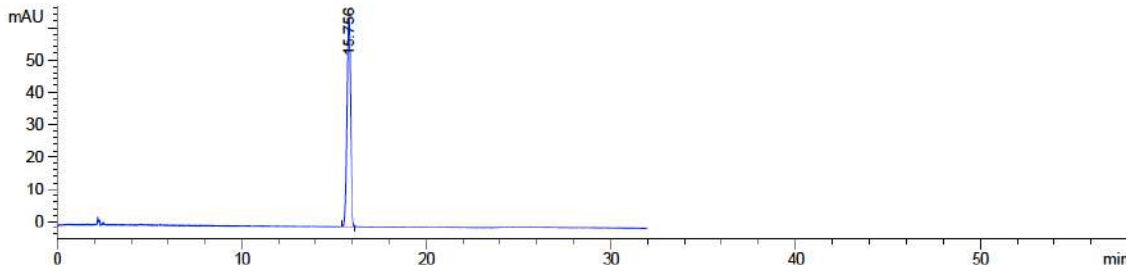


Fig. S49. High resolution ESI(+) MS spectrum of compound **(16*S*)-2d**.



Signal 1: DAD1 A, Sig=210,16 Ref=off

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	15.756	BB	0.2148	899.62329	64.46941	100.0000
Totals :				899.62329	64.46941	

Fig. S50. HPLC chromatogram of compound **(16*S*)-2d** (RP18, isocratic water/acetonitrile 30:70 + 1% trifluoroacetic acid).

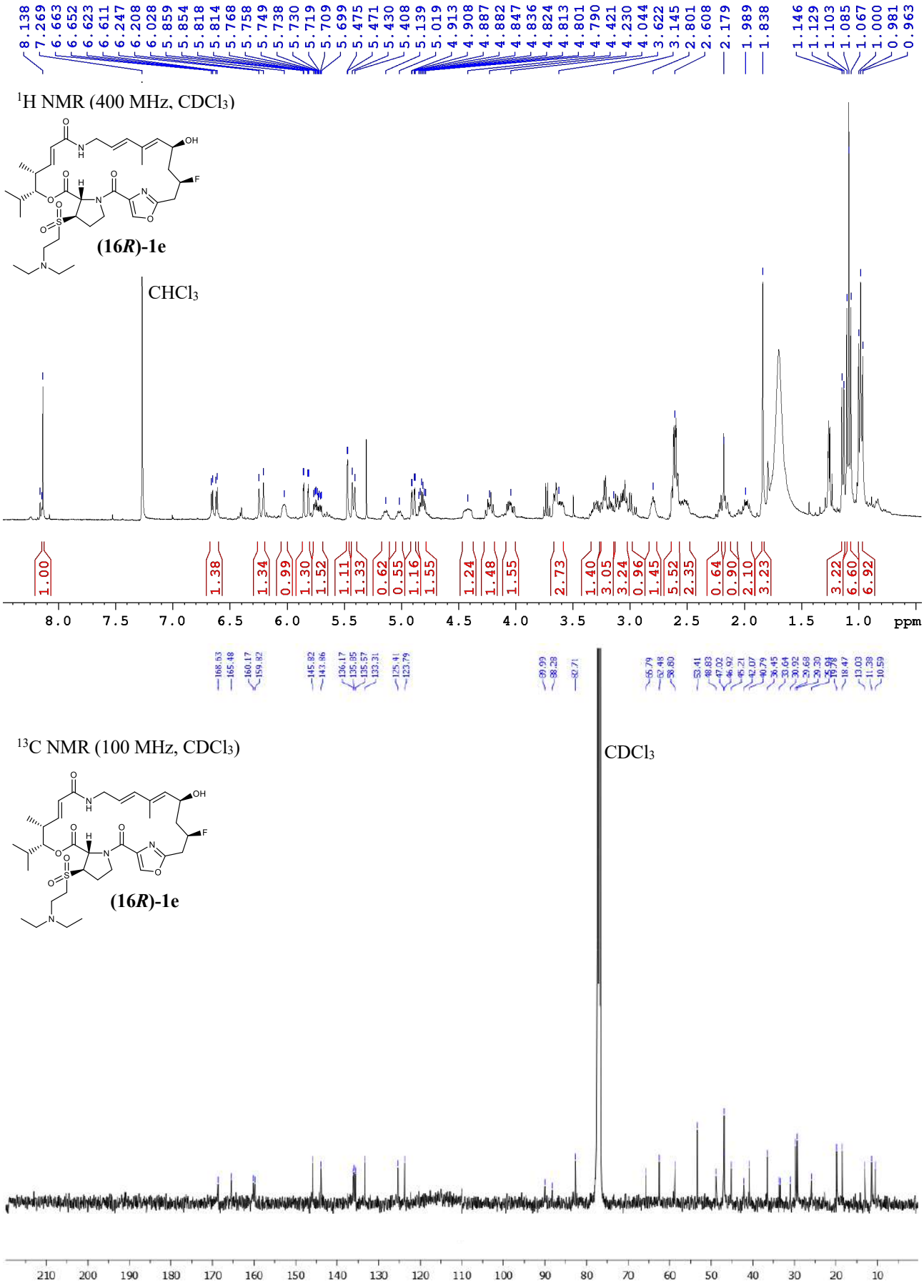


Fig. S51. ^{1}H NMR and ^{13}C NMR spectra of compound **(16*R*)-1e**.

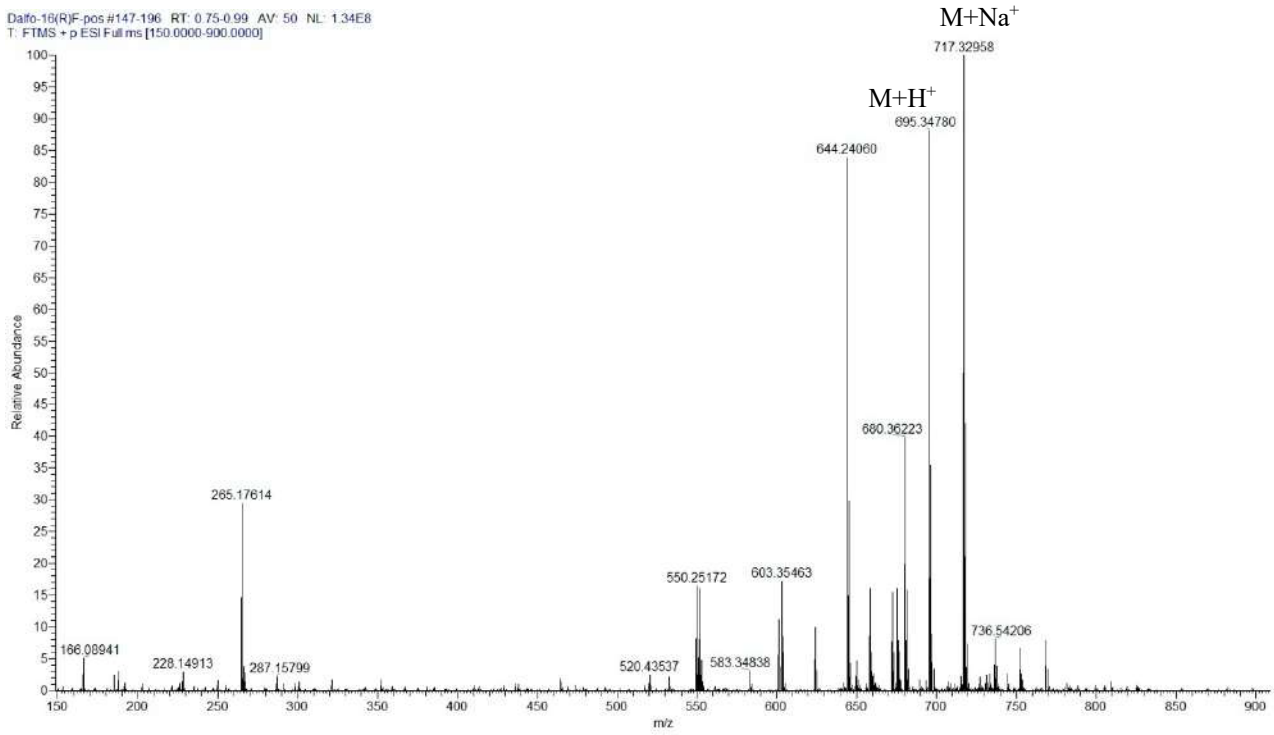
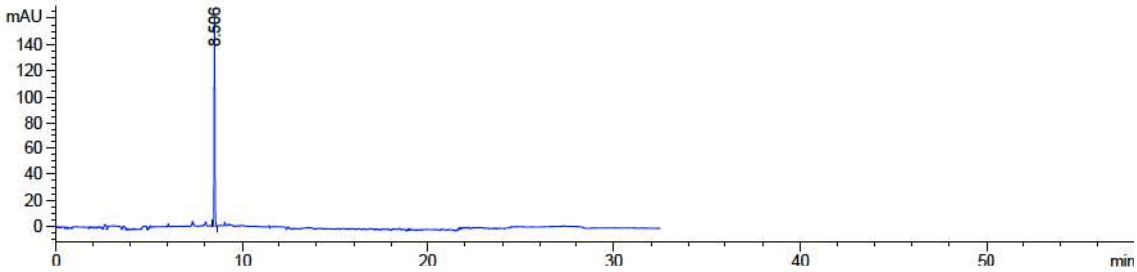


Fig. S52. High resolution ESI(+) MS spectrum of compound **(16*R*)-1e**.



Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	8.506	VV	0.0460	462.79251	158.88995	100.0000
Totals :				462.79251	158.88995	

Fig. S53. HPLC spectrum of compound **(16*R*)-1e** (RP18, isocratic water/acetonitrile 30:70 + 1% trifluoroacetic acid).

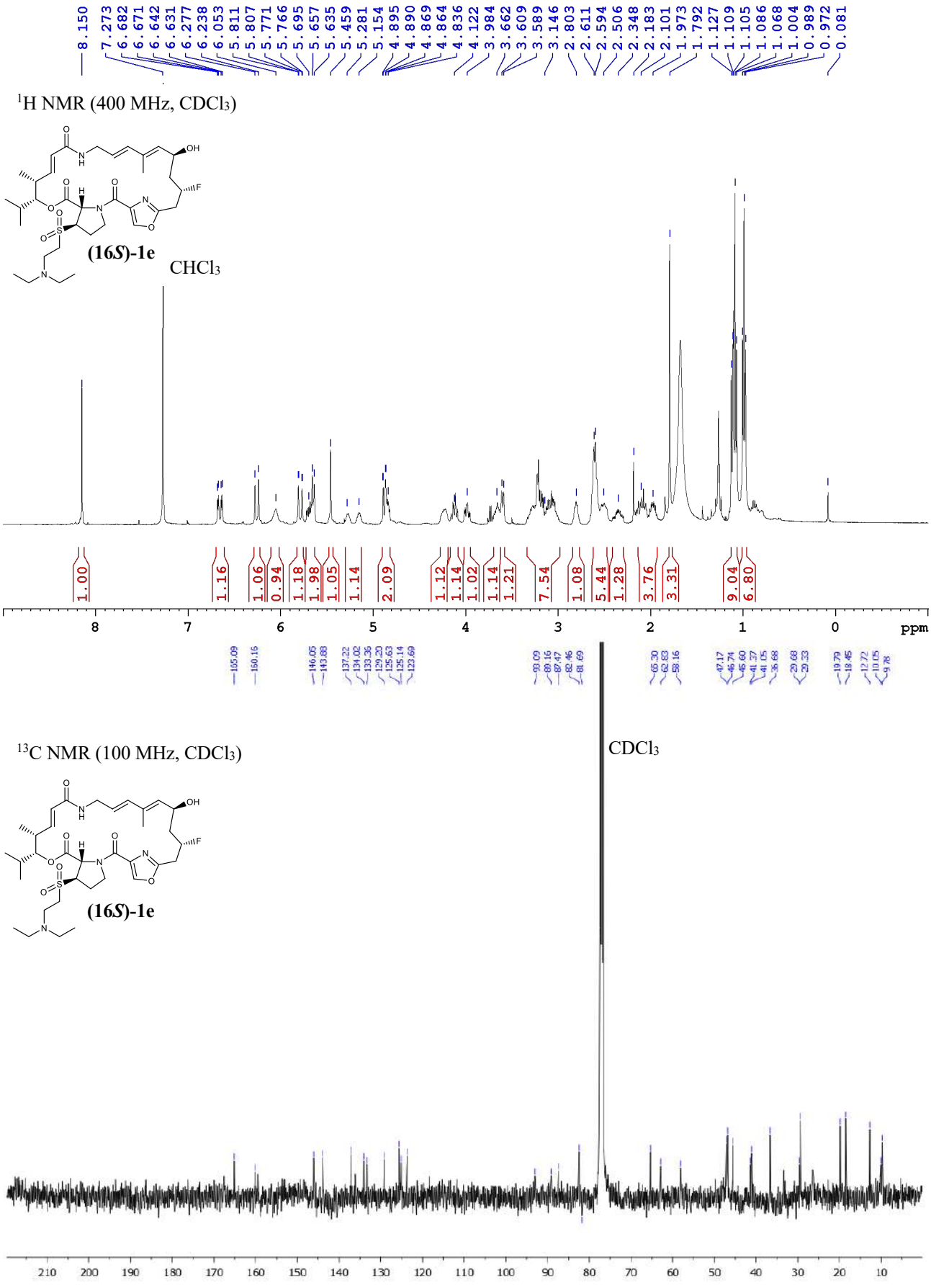


Fig. S54. ^{1}H NMR and ^{13}C NMR spectra of compound **(16*S*)-1e**.

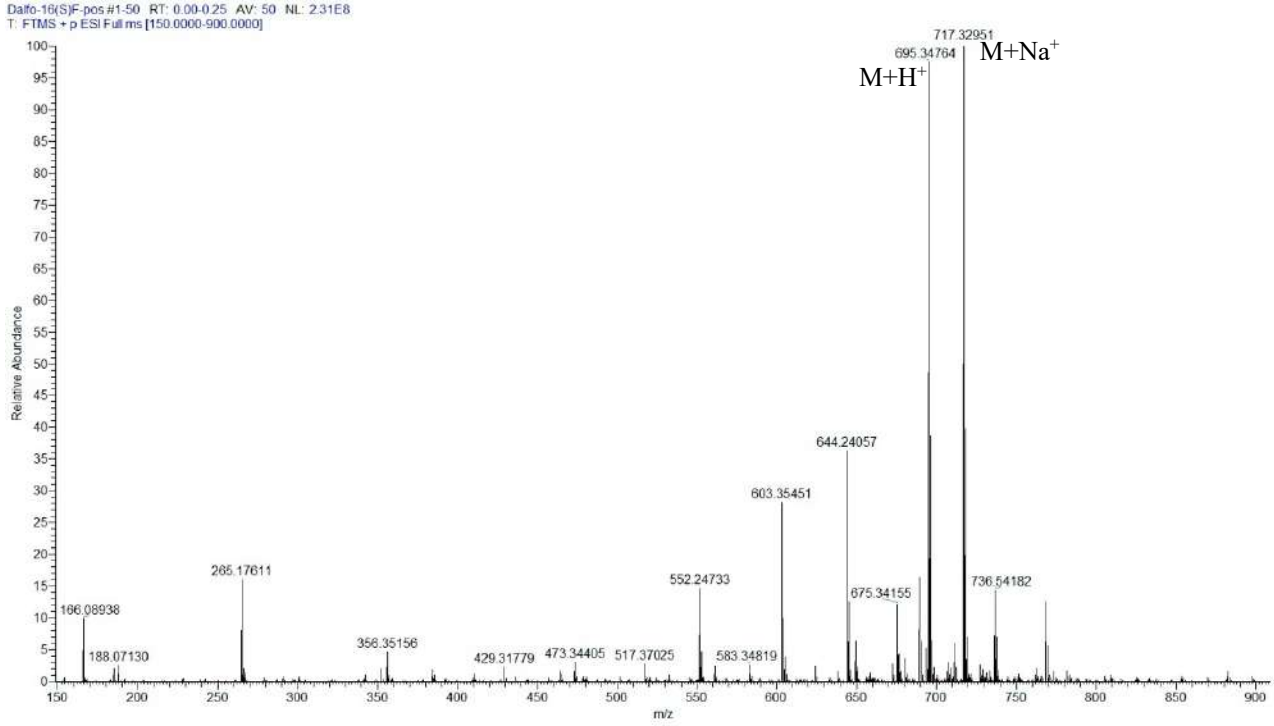
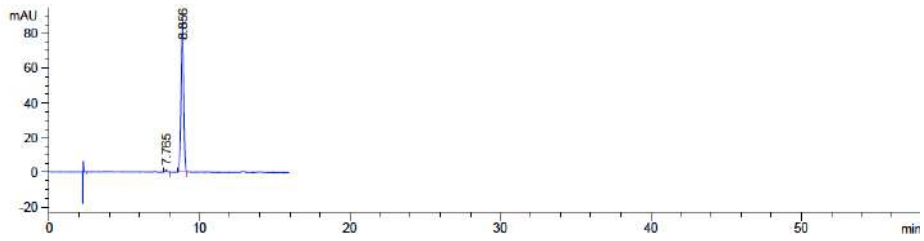


Fig. S55. High resolution ESI(+) MS spectrum of compound **(16*S*)-1e**.



Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	7.765	BB	0.1264	15.27746	1.60035	1.3671
2	8.856	BB	0.1922	1102.20605	89.19329	98.6329
Totals :				1117.48351	90.79364	

Fig. S56. HPLC chromatogram of compound **(16*S*)-1e** (RP18, isocratic water/acetonitrile 30:70 + 1% trifluoroacetic acid).

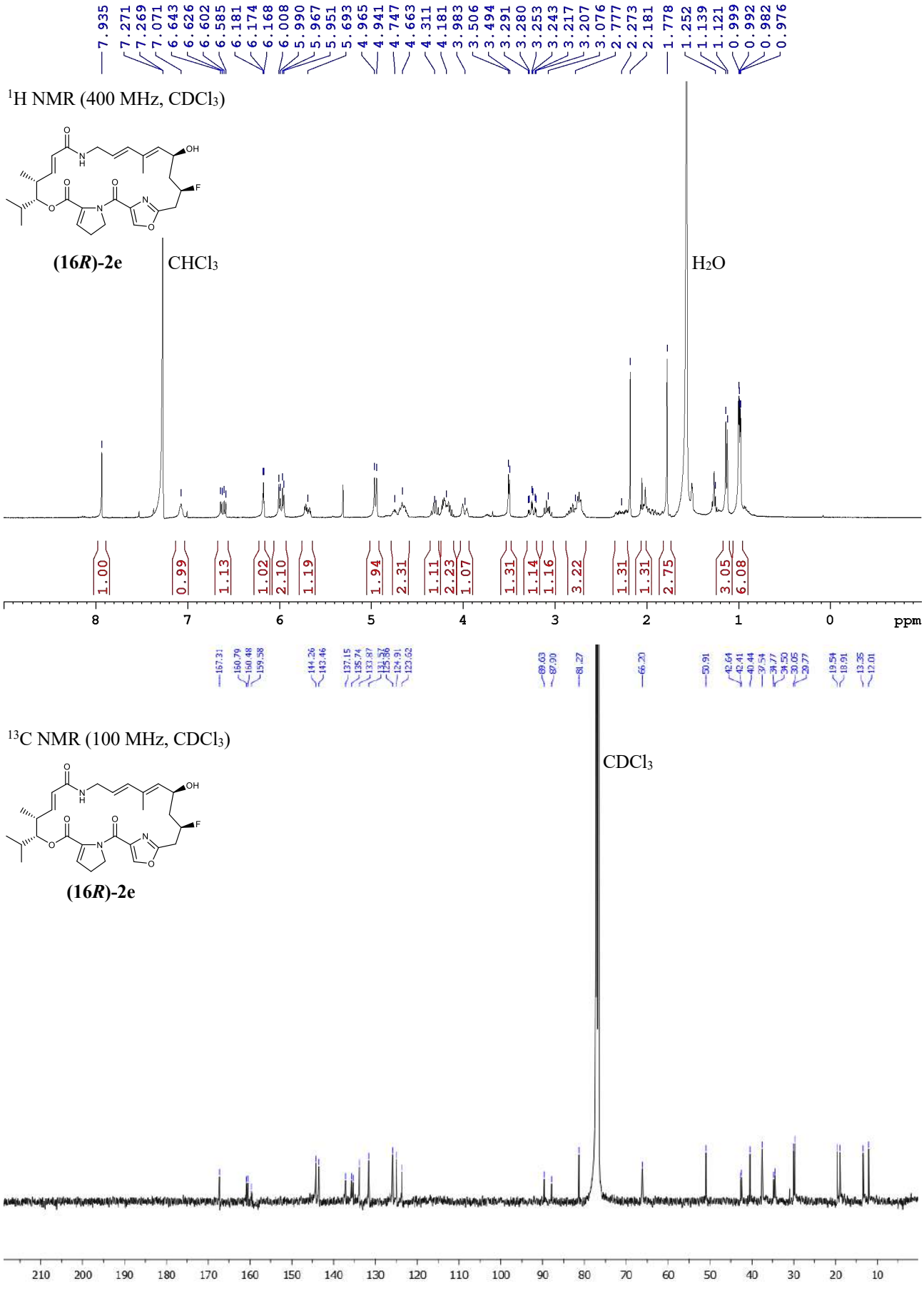


Fig. S57. ^{1}H NMR and ^{13}C NMR spectra of compound **(16*R*)-2e**.

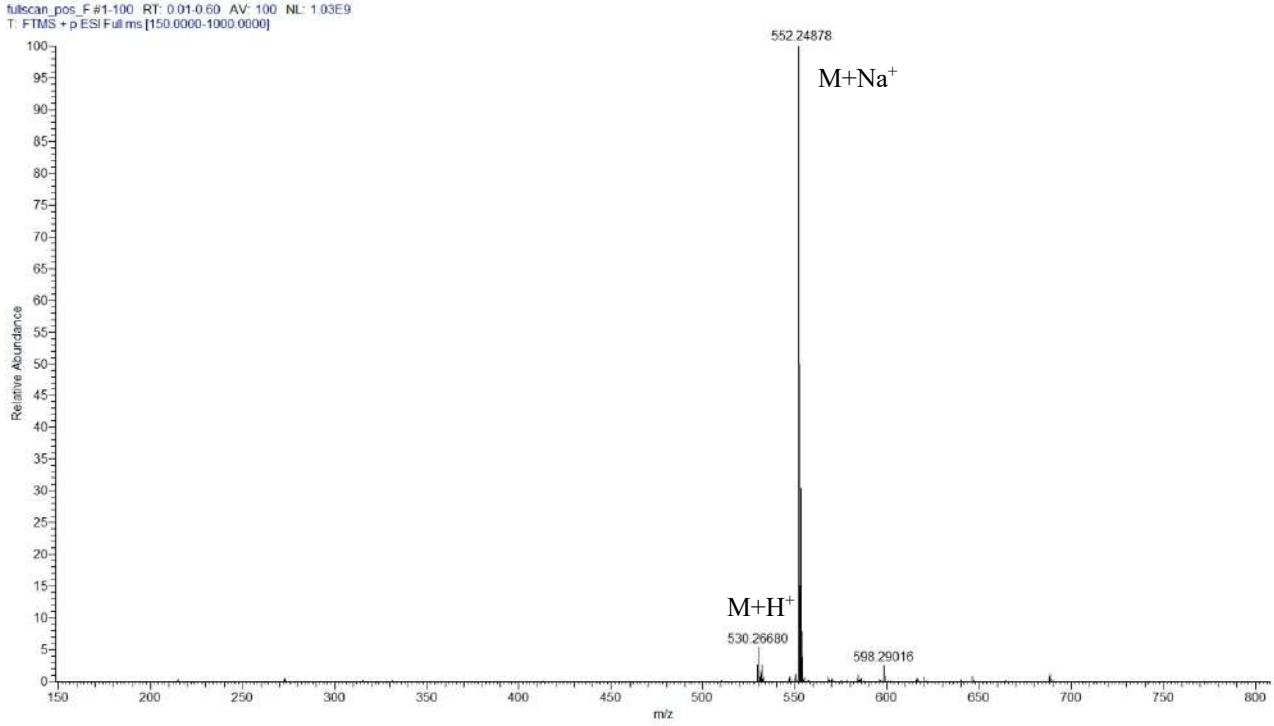
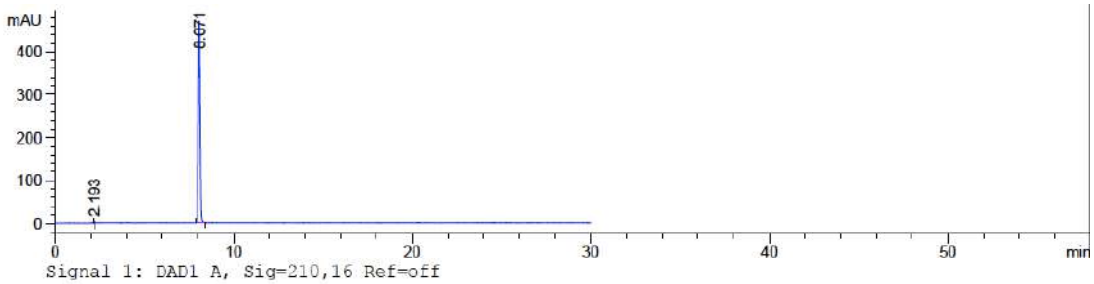


Fig. S58. High resolution ESI(+) MS spectrum of compound **(16*R*)-2e**.



Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	2.193	BB	0.0331	8.81573	3.94144	0.2794
2	8.071	BB	0.1039	3146.79907	471.31345	99.7206
Totals :				3155.61480	475.25489	

Fig. S59. HPLC chromatogram of compound **(16*R*)-2e** (RP18, gradient water/acetonitrile 30:70 to 80:20).

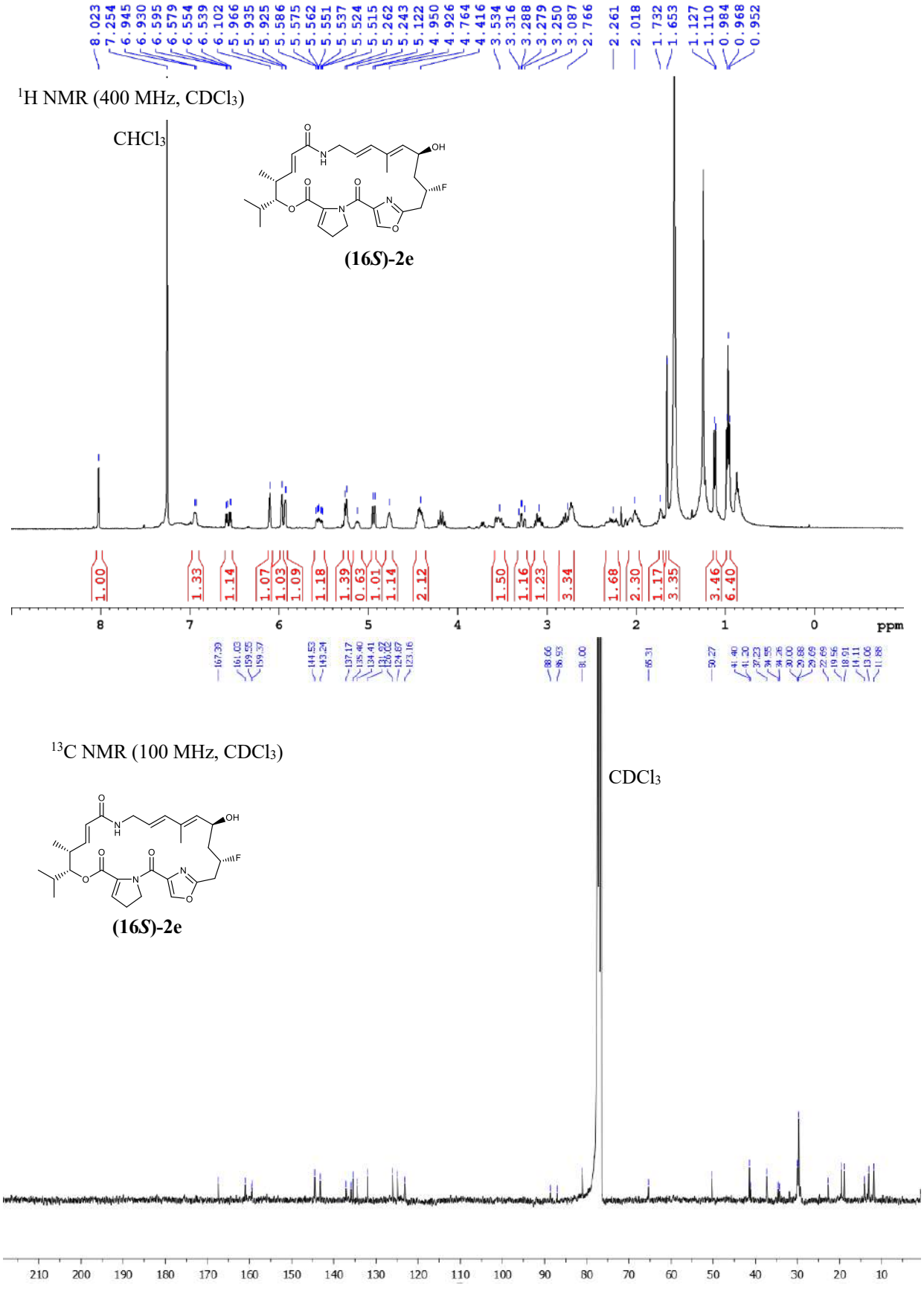


Fig. S60. ^{1}H NMR and ^{13}C NMR spectra of compound **(16*S*)-2e**.

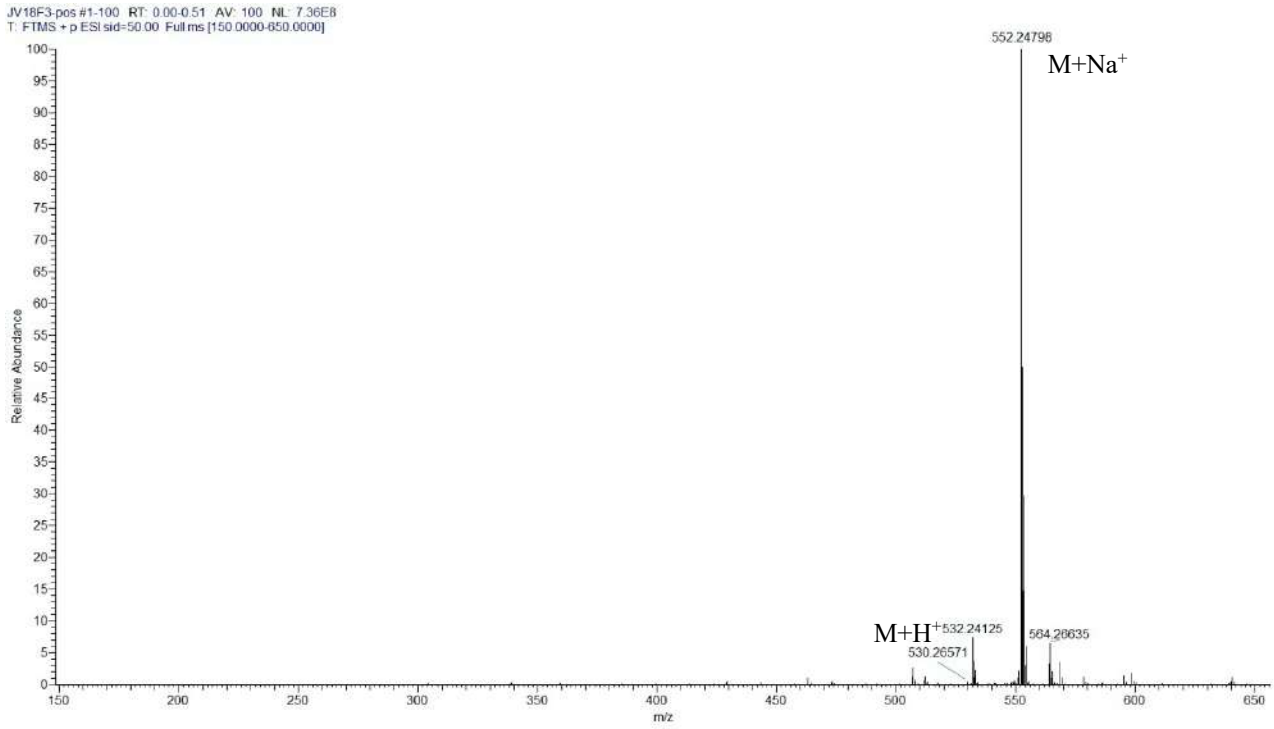
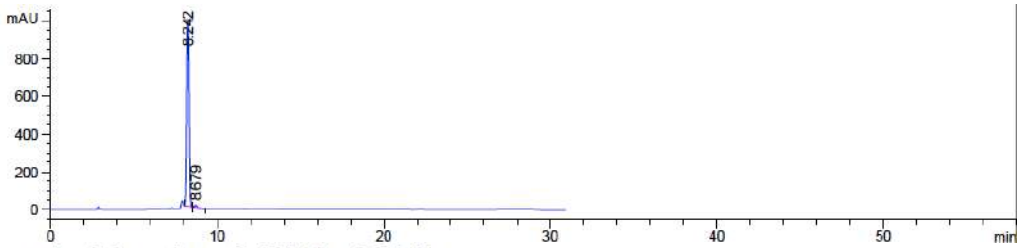


Fig. S61. High resolution ESI(+) MS spectrum of compound **(16*S*)-2e**.



Signal 2: DAD1 B, Sig=254,16 Ref=390,20

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	8.242	BV	0.1502	8991.49316	987.97986	98.9809
2	8.679	VV	0.1284	92.57776	12.79468	1.0191
Totals :				9084.07092	1000.77454	

Fig. S62. HPLC chromatogram of compound **(16*S*)-2e** (RP18, gradient water/acetonitrile 30:70 to 80:20).

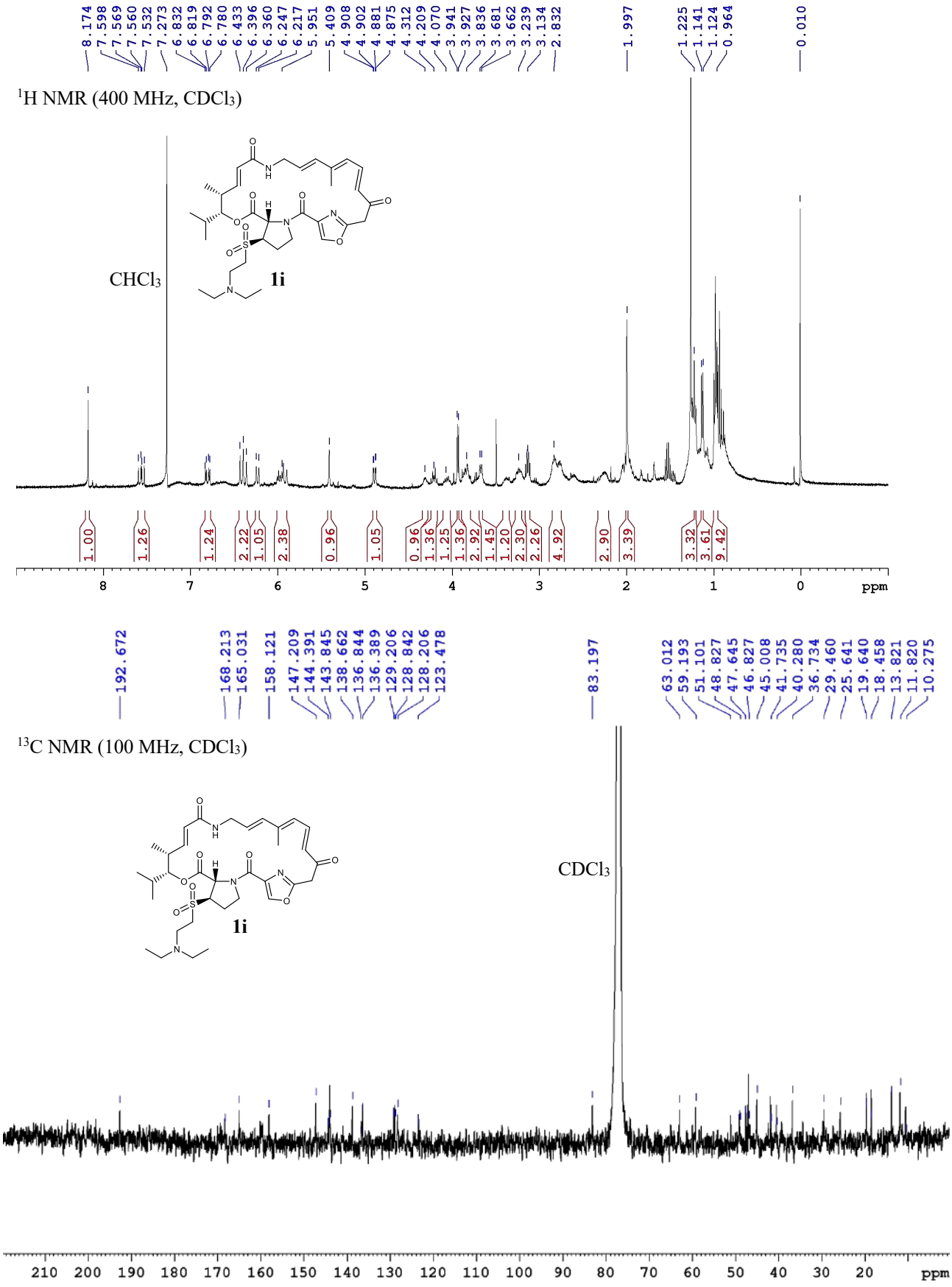


Fig. S63. ^{1}H NMR and ^{13}C NMR spectra of compound **1i**.

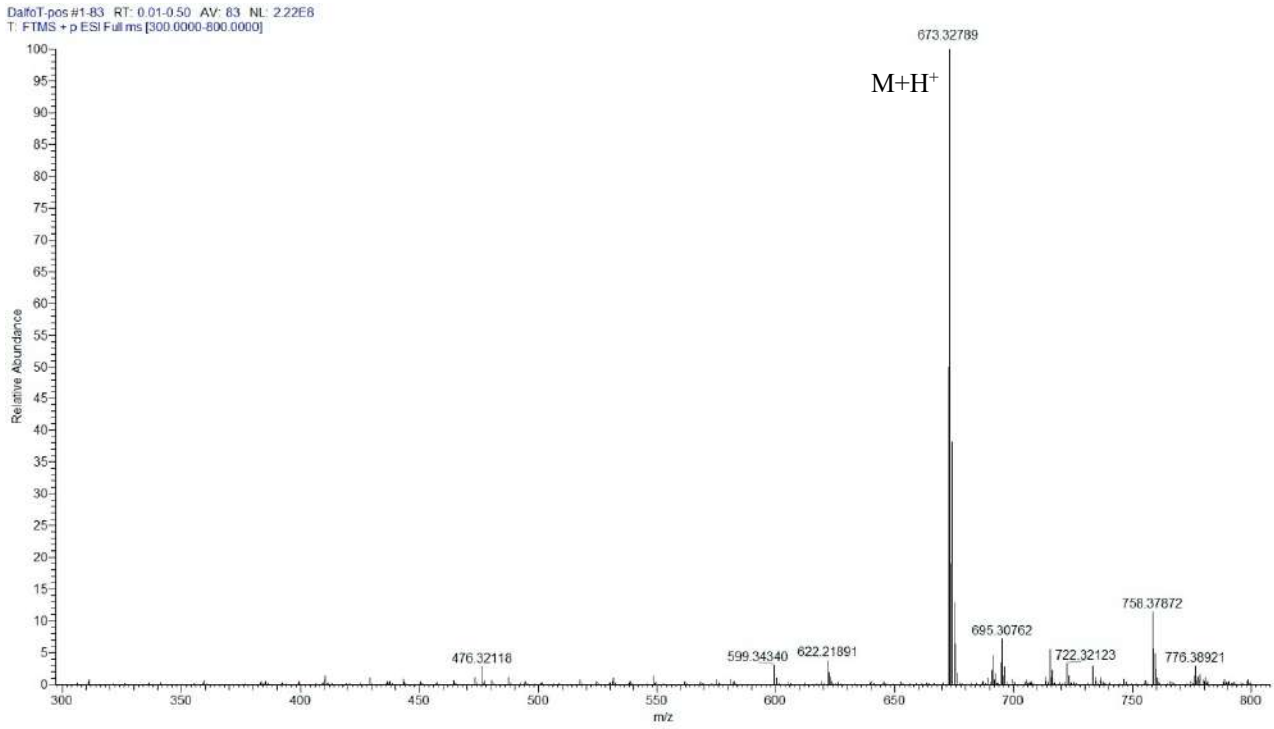
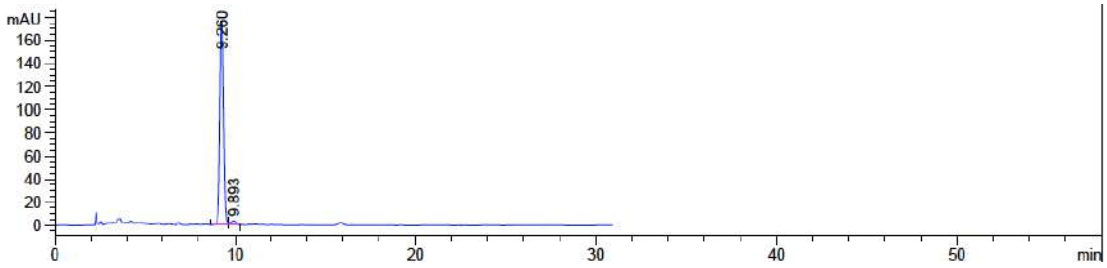


Fig. S64. High resolution ESI(+) MS spectrum of compound **1i**.



Signal 2: DAD1 B, Sig=254,16 Ref=390,20

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	9.260	VV	0.2124	2431.16602	176.79109	97.2352
2	9.893	VV	0.2779	69.12883	3.77092	2.7648
Totals :				2500.29485	180.56201	

Fig. S65. HPLC chromatogram of compound **1i** (RP18, isocratic water/acetonitrile 30:70 + 1% trifluoroacetic acid).

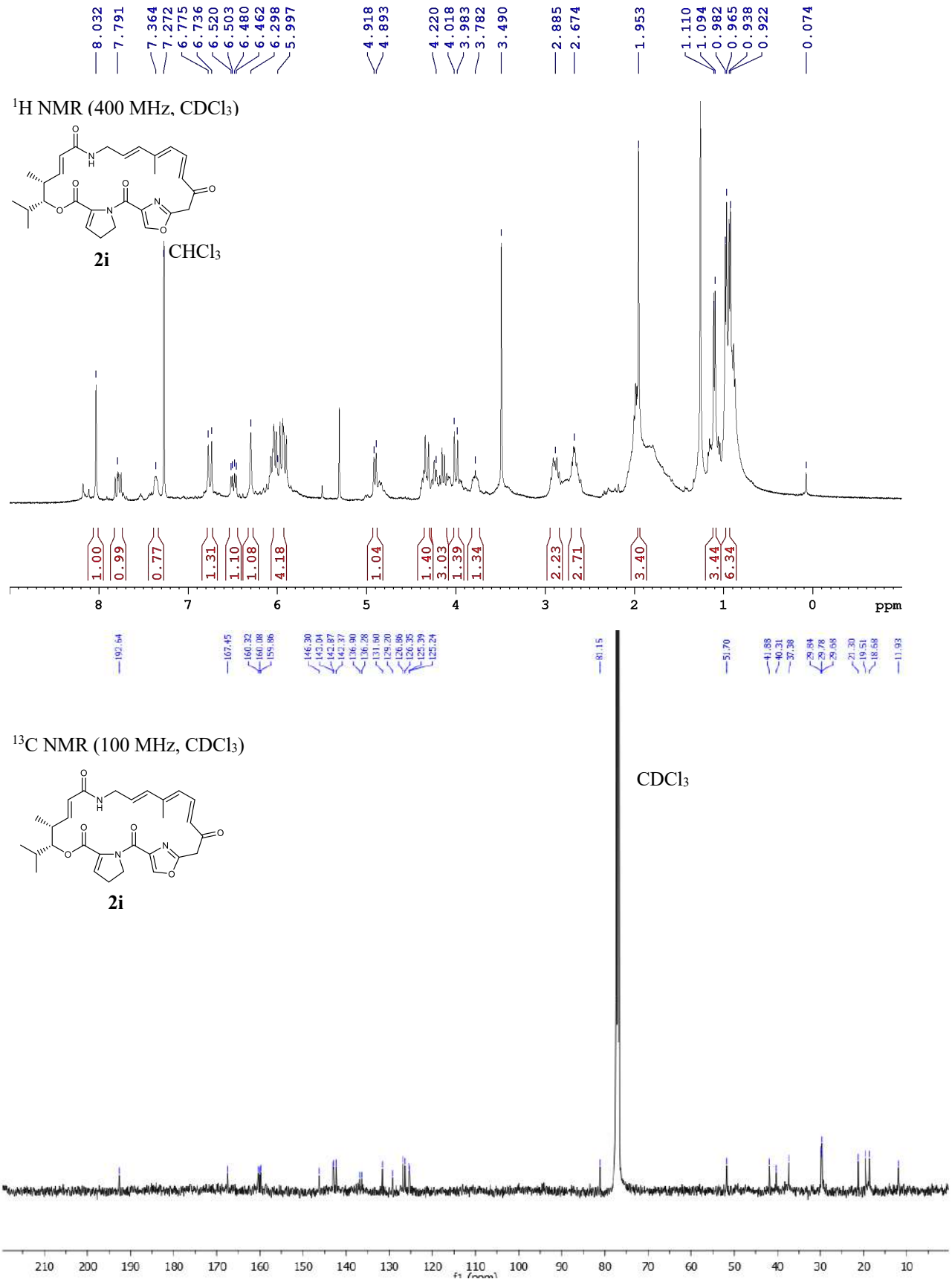


Fig. S66. ^{1}H NMR and ^{13}C NMR spectra of compound **2i**.

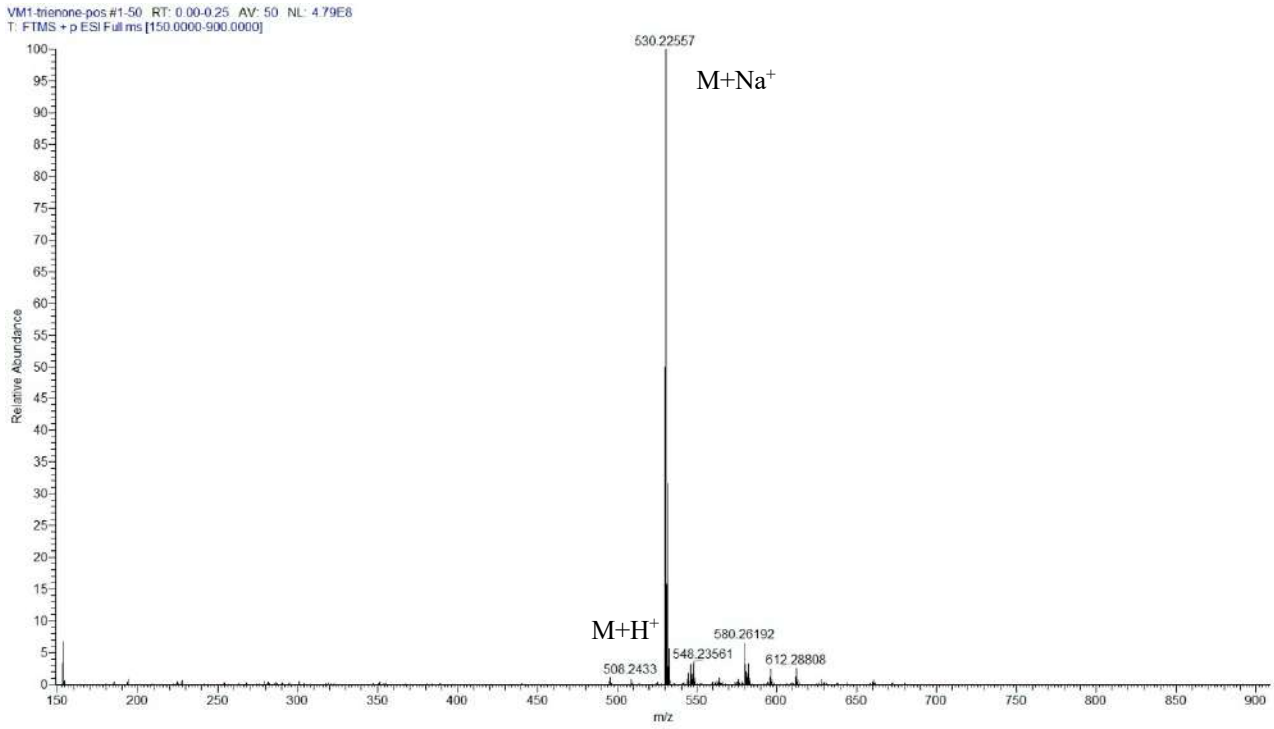
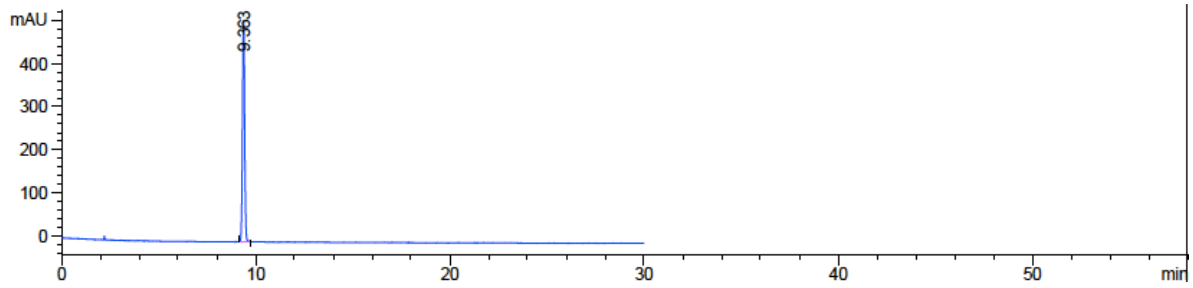


Fig. S67. High resolution ESI(+) MS spectra of compound **2i**.




```
Signal 1: DAD1 A, Sig=210,16 Ref=off

Peak RetTime Type  Width     Area      Height     Area
  #   [min]        [min]   [mAU*s]     [mAU]       %
----|-------|----|-------|----------|----------|--------|
   1   9.363 BB    0.1211 3994.25659  511.33139 100.0000

Totals :                  3994.25659  511.33139
```

Fig. S68. HPLC chromatogram of compound **2i** (RP18, gradient water/acetonitrile 30:70 to 80:20).

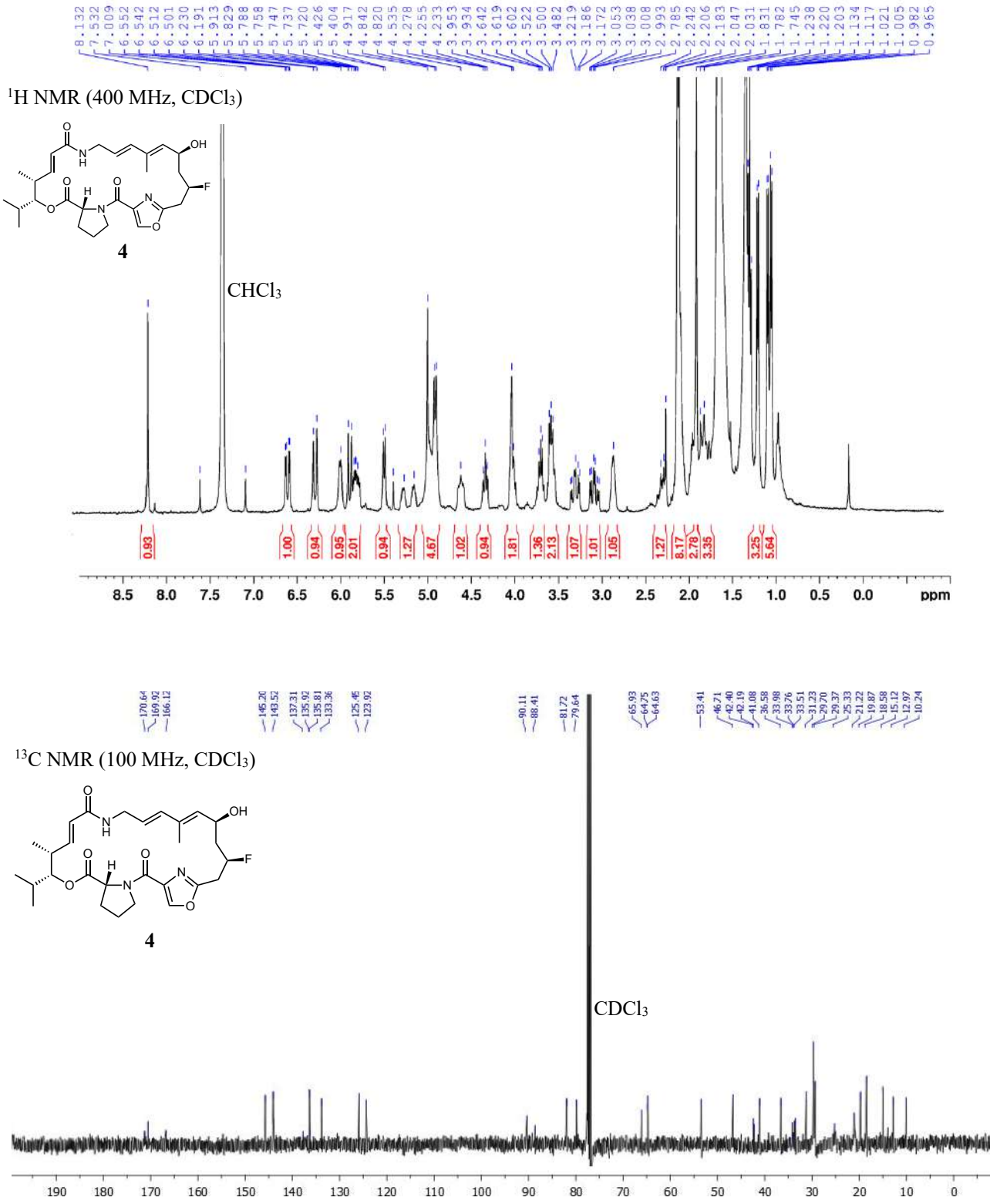


Fig. S69. ^{1}H NMR and ^{13}C NMR spectra of compound **4**.

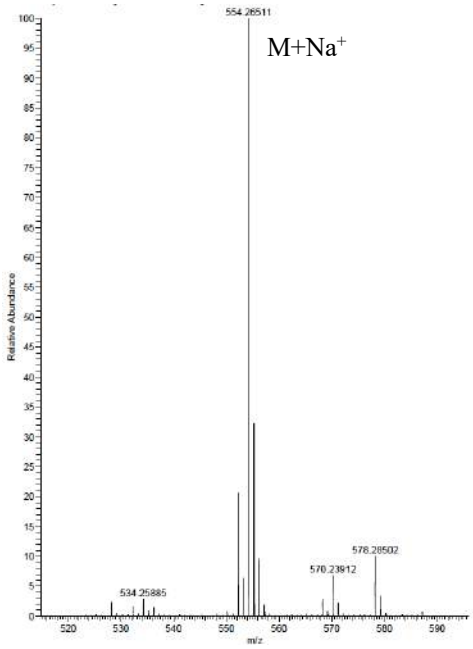
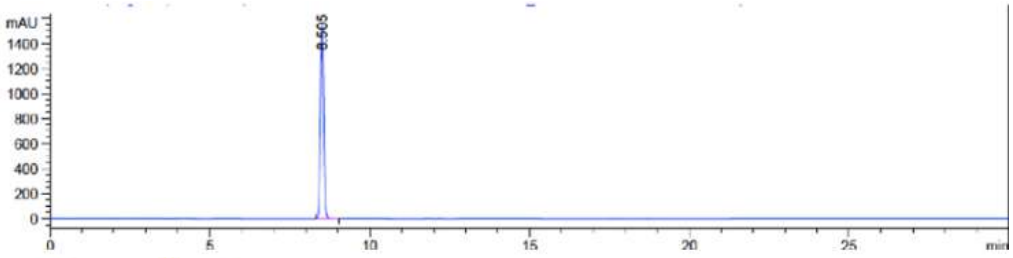


Fig. S70. High resolution ESI(+) MS spectrum of compound **4**.



Signal 1: DAD1 A, Sig=210,16 Ref=off

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	8.505	BB	0.1174	1.16428e4	1553.29126	100.0000
Totals :				1.16428e4	1553.29126	

Fig. S71. HPLC chromatogram of compound **4** (RP18, gradient water/acetonitrile 30:70 to 80:20).

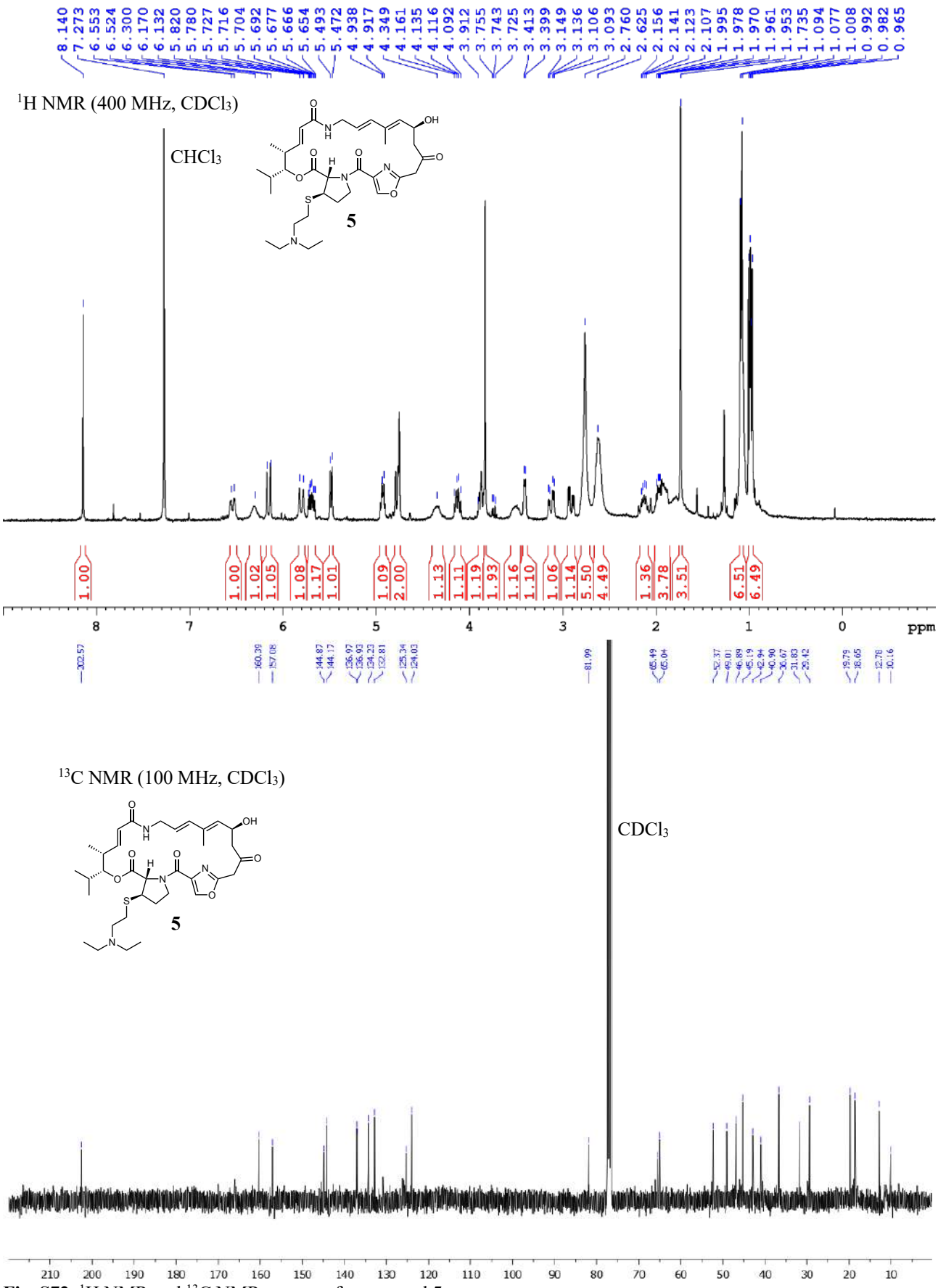


Fig. S72. ^{1}H NMR and ^{13}C NMR spectra of compound **5**.

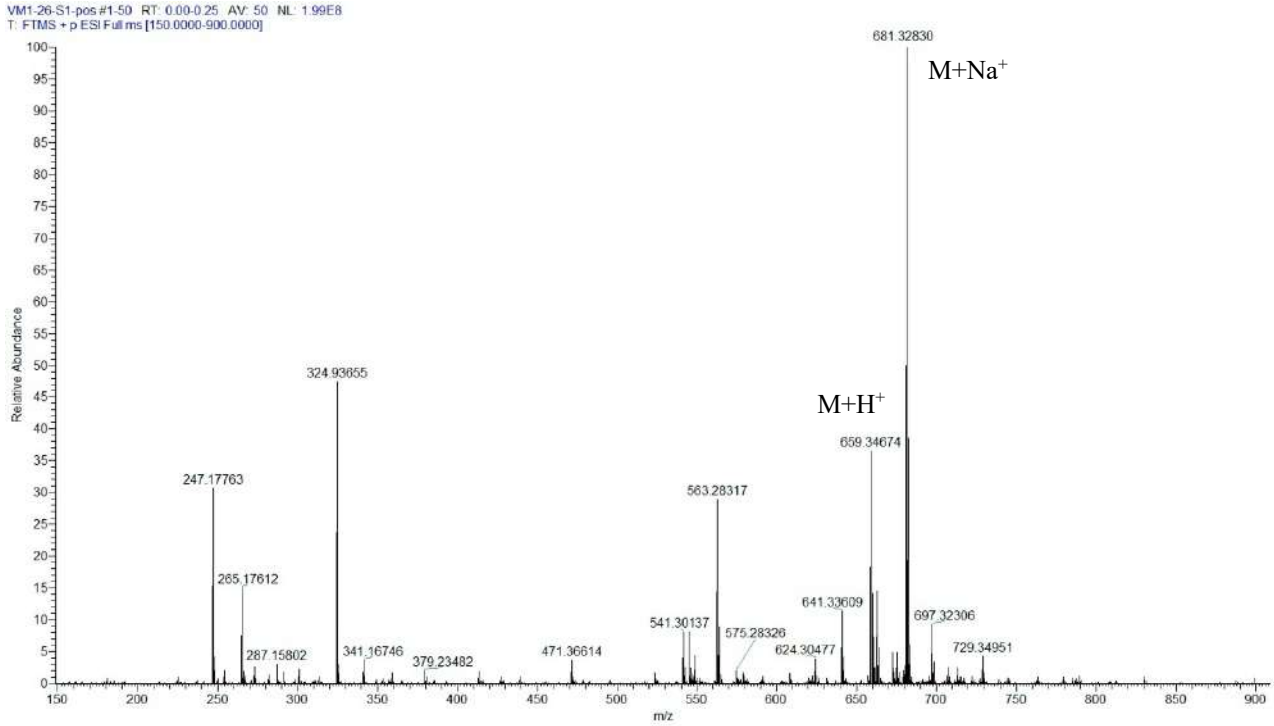
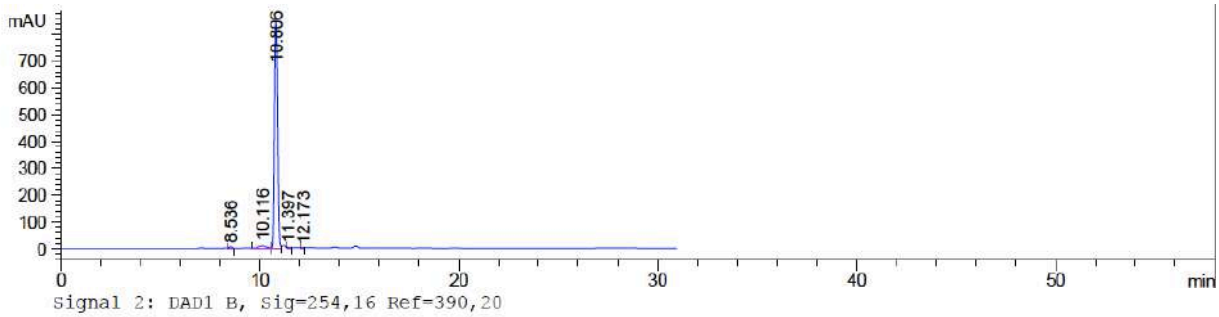


Fig. S73. High resolution ESI(+) MS spectrum of compound **5**.



Signal 2: DAD1 B, Sig=254,16 Ref=390,20

Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	8.536	VV	0.1631	57.80320	5.66381	0.6365
2	10.116	VV	0.2975	236.64586	10.72316	2.6060
3	10.806	VV	0.1681	8728.30664	847.67499	96.1192
4	11.397	VV	0.1841	33.97823	2.82913	0.3742
5	12.173	VV	0.1968	23.97721	1.73948	0.2640
Totals :				9080.71114	868.63057	

Fig. S74. HPLC chromatogram of compound **5** (RP18, isocratic water/acetonitrile 30:70 + 1% trifluoroacetic acid).

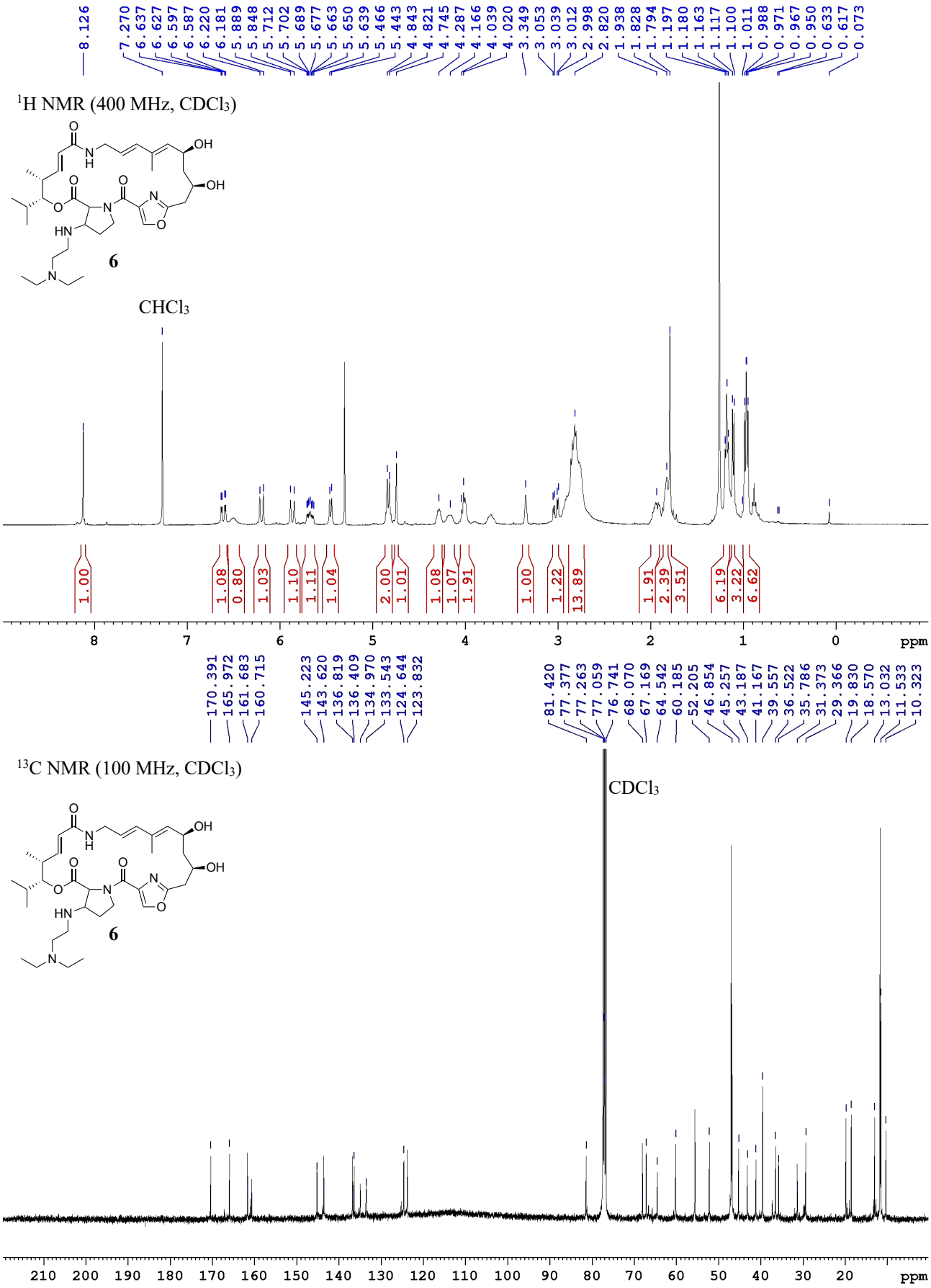


Fig. S75. ^{1}H NMR and ^{13}C NMR spectra of compound **6**.

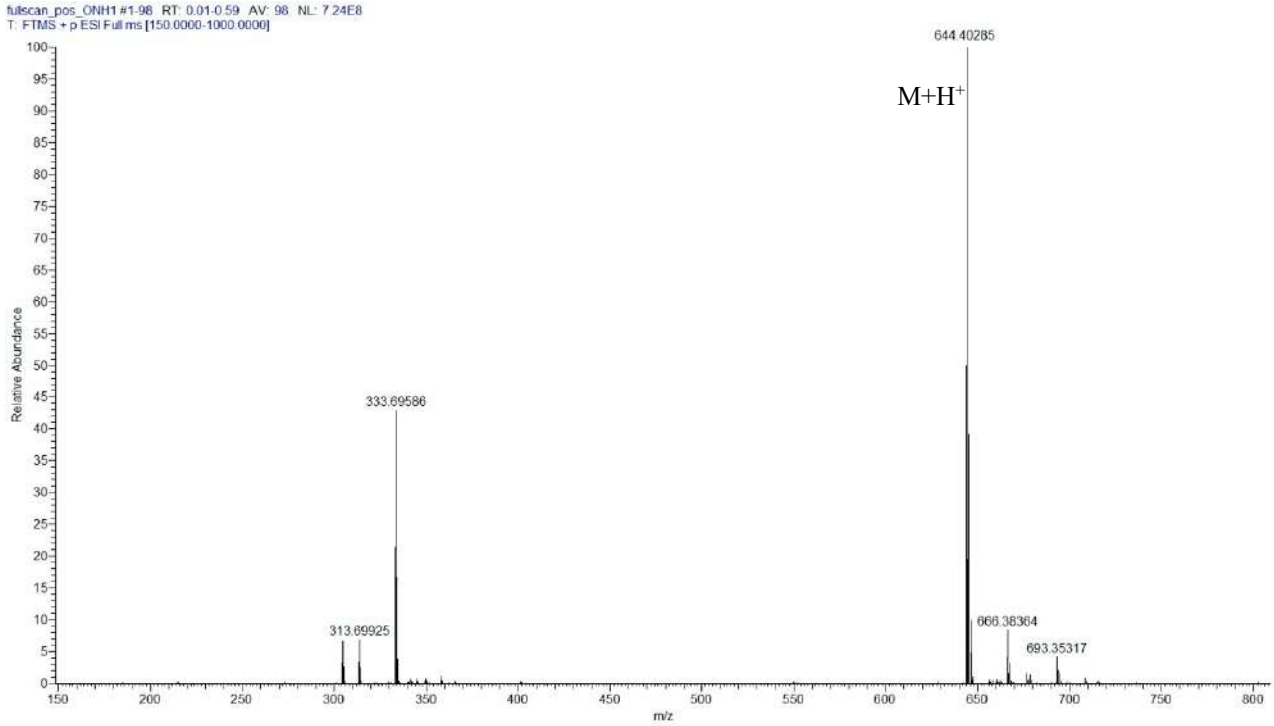
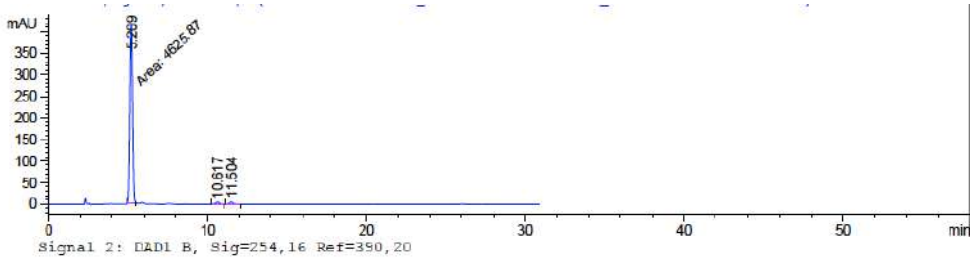


Fig. S76. High resolution ESI(+) MS spectrum of compound **6**.



Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	5.209	MM	0.1862	4625.86572	414.00119	97.1521
2	10.617	BB	0.2526	69.83024	4.31944	1.4666
3	11.504	BB	0.2140	65.77187	4.73754	1.3813
Totals :				4761.46783	423.05817	

Fig. S77. HPLC chromatogram of compound **6** (RP18, isocratic water/acetonitrile 30:70 + 1% trifluoroacetic acid).

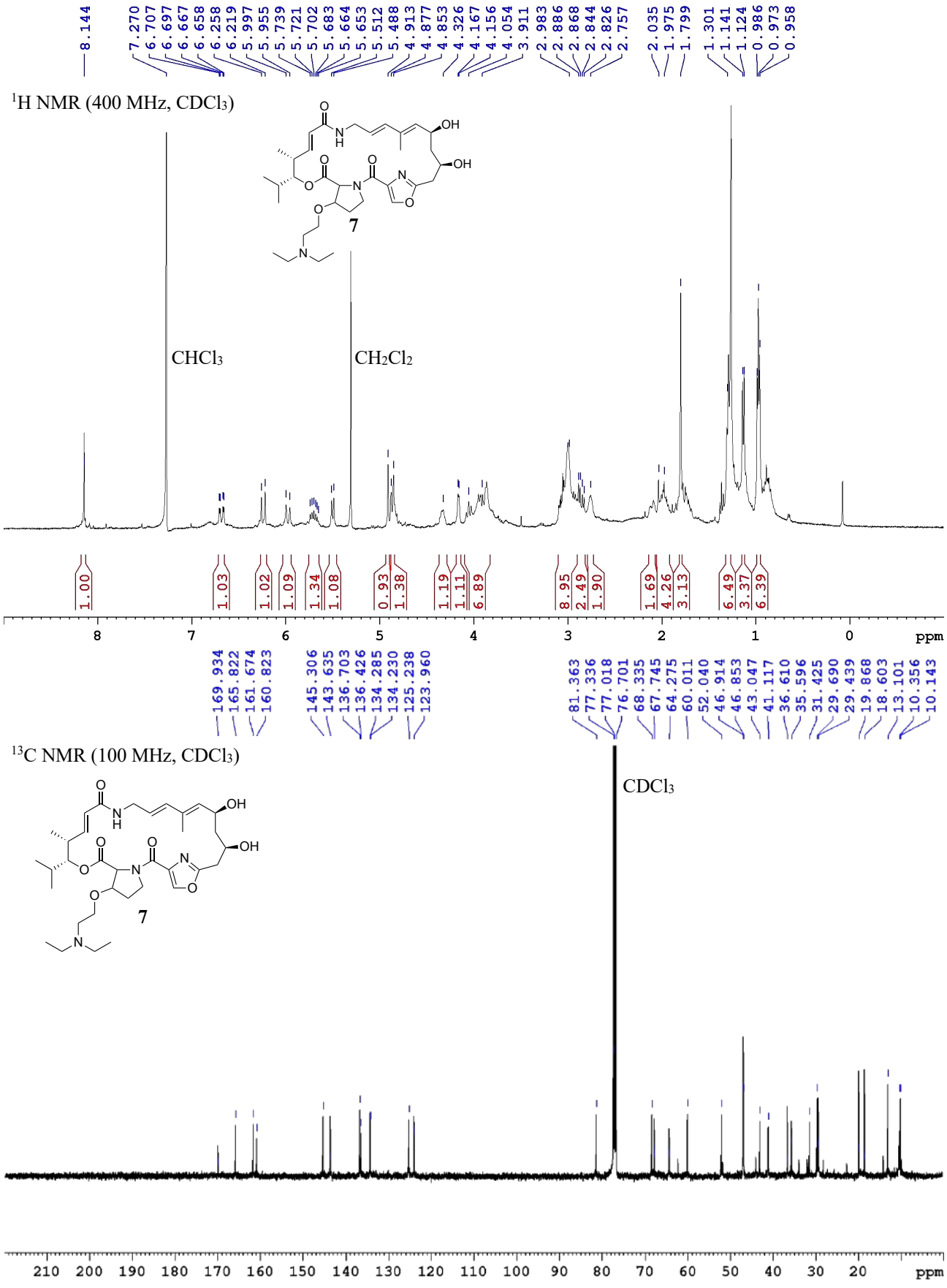


Fig. S78. ^{1}H NMR and ^{13}C NMR spectra of compound **7**.

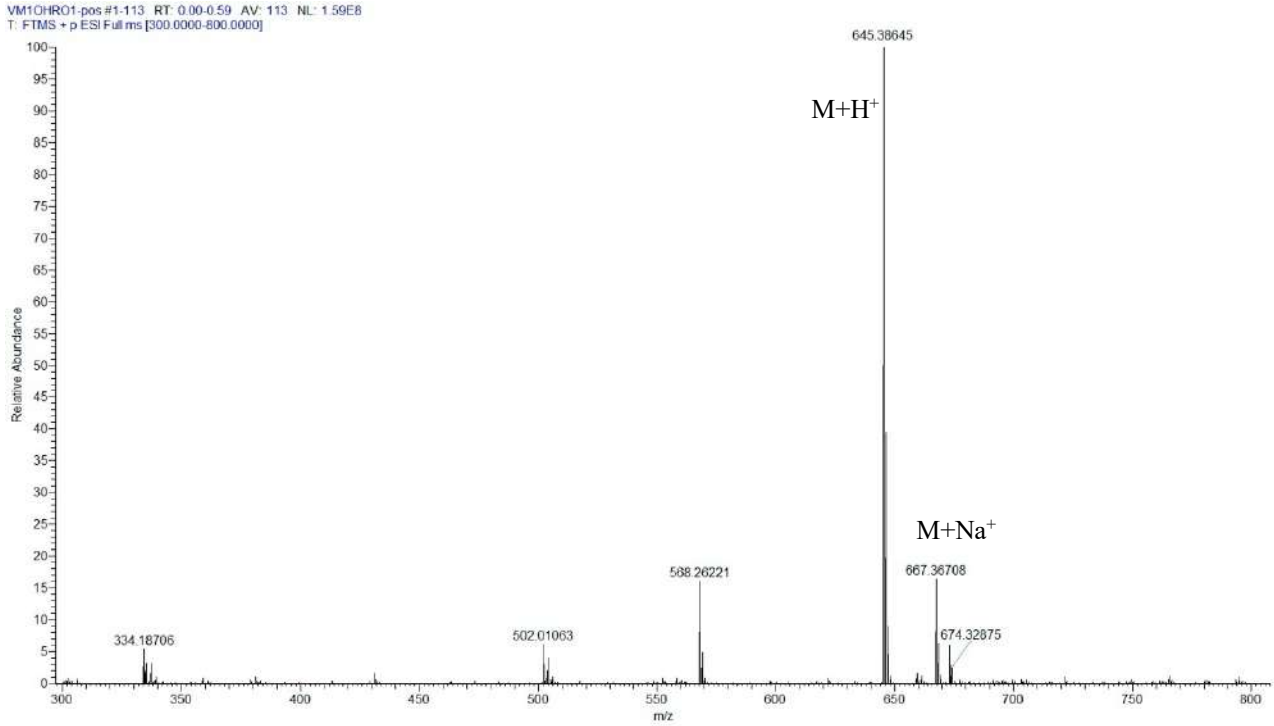
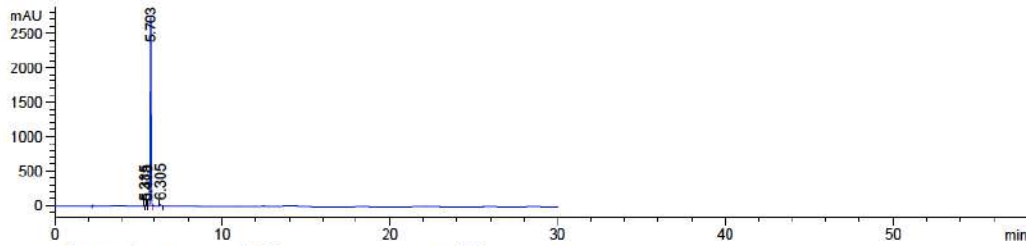


Fig. S79. High resolution ESI(+) MS spectrum of compound **7**.



Peak #	RetTime [min]	Type	Width [min]	Area [mAU*s]	Height [mAU]	Area %
1	5.385	VV	0.0473	29.69088	9.27708	0.2961
2	5.478	VV	0.0652	46.01634	10.91695	0.4589
3	5.703	VV	0.0569	9871.58789	2747.98120	98.4549
4	6.305	VV	0.0507	79.21222	22.68969	0.7900
Totals :				1.00265e4	2790.86492	

Fig. S80. HPLC chromatogram of compound **7** (RP18, isocratic water/acetonitrile 30:70 + 1% trifluoroacetic acid).