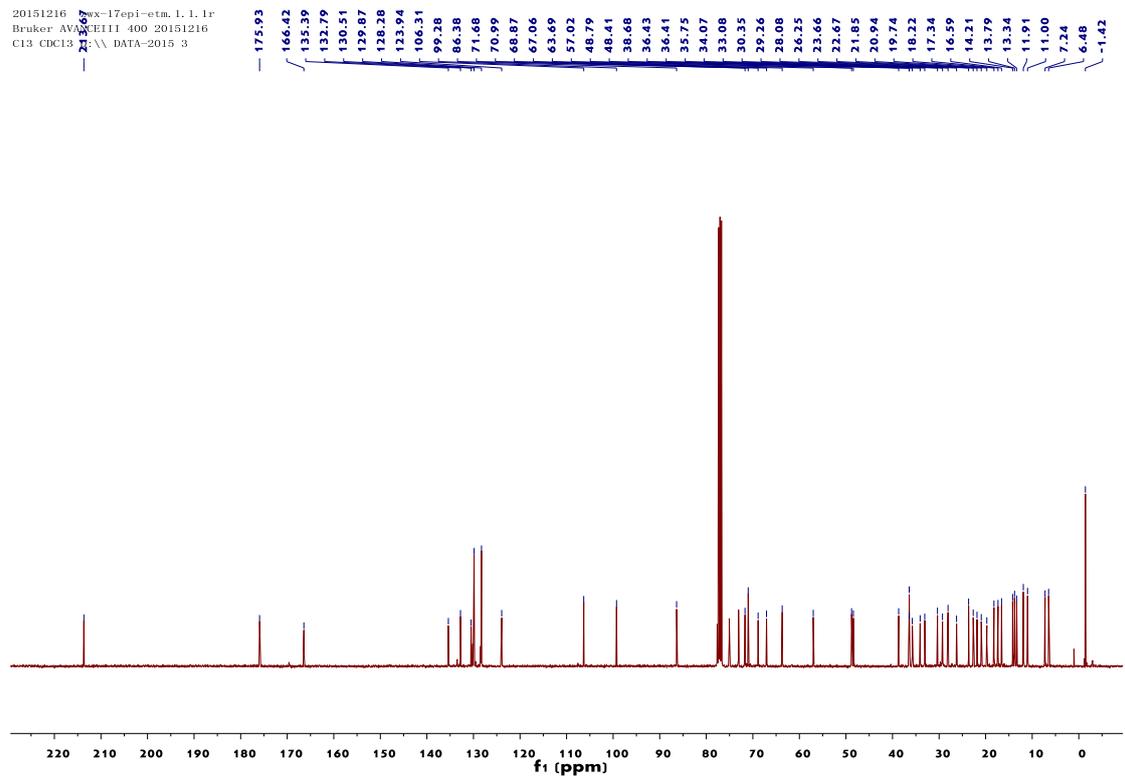
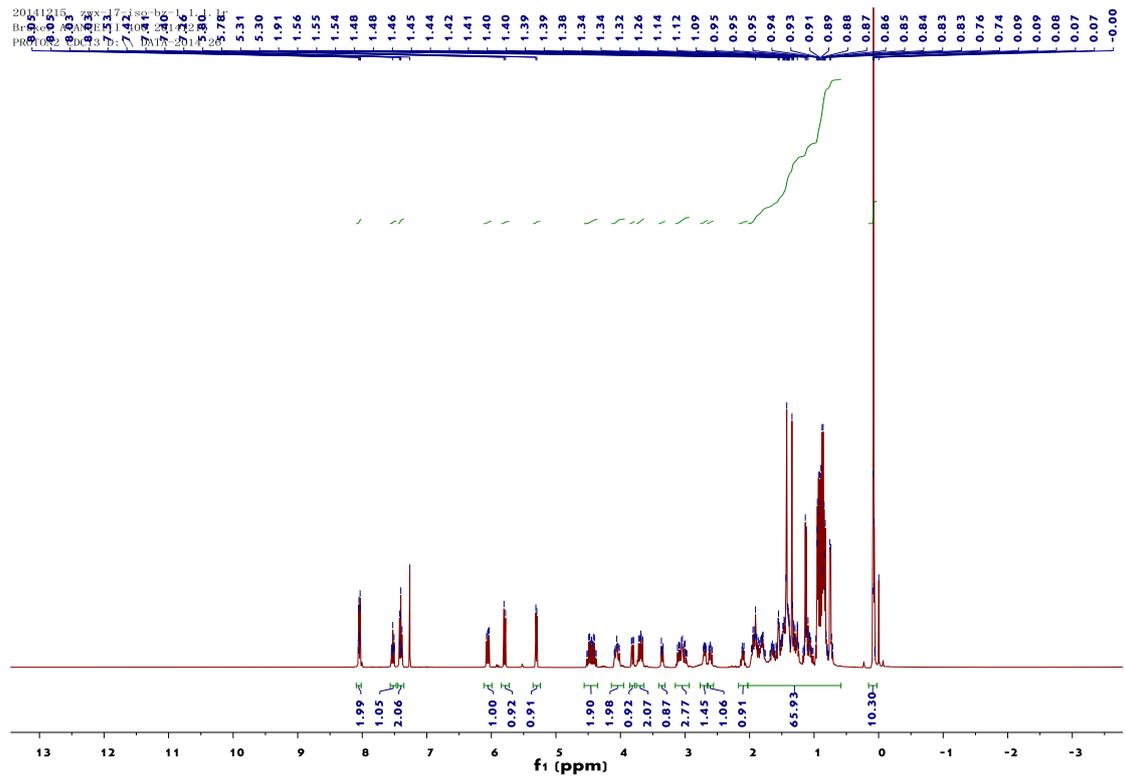


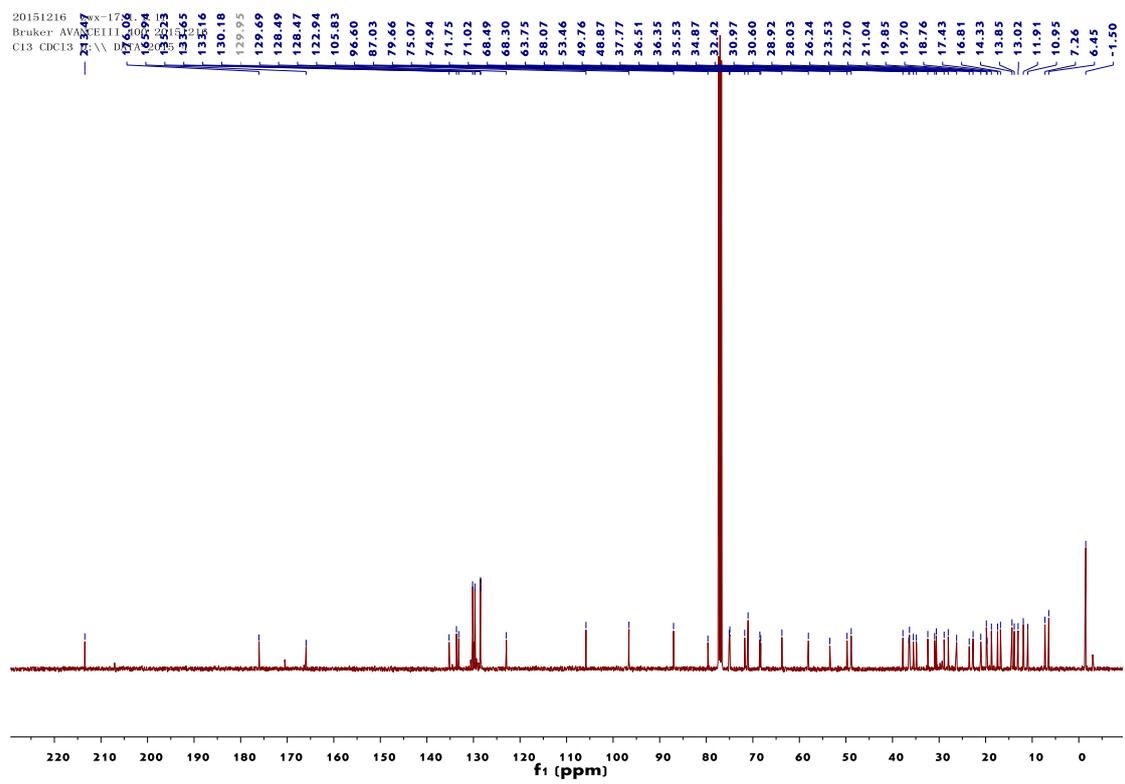
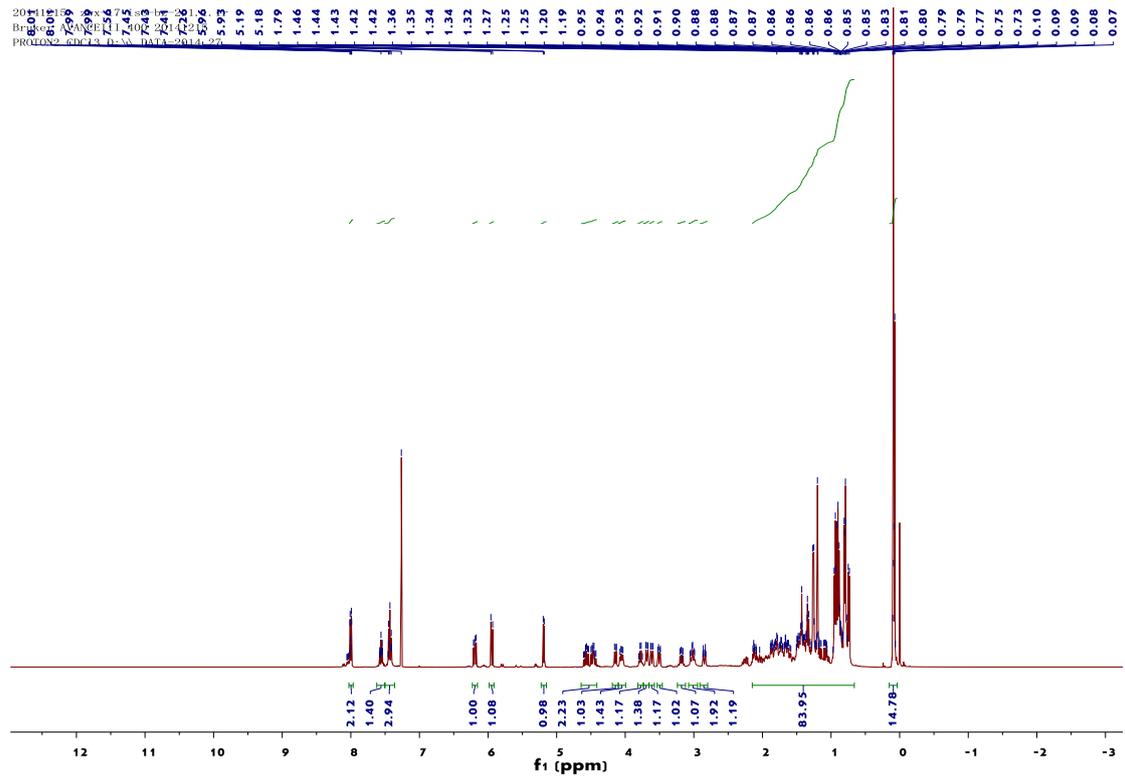
## Supplementary information

1.  $^1\text{H}$  NMR and  $^{13}\text{C}$  NMR of 2,3,4,5,8,9,10
2. Crystal Data of 9
3. Dose-response curves

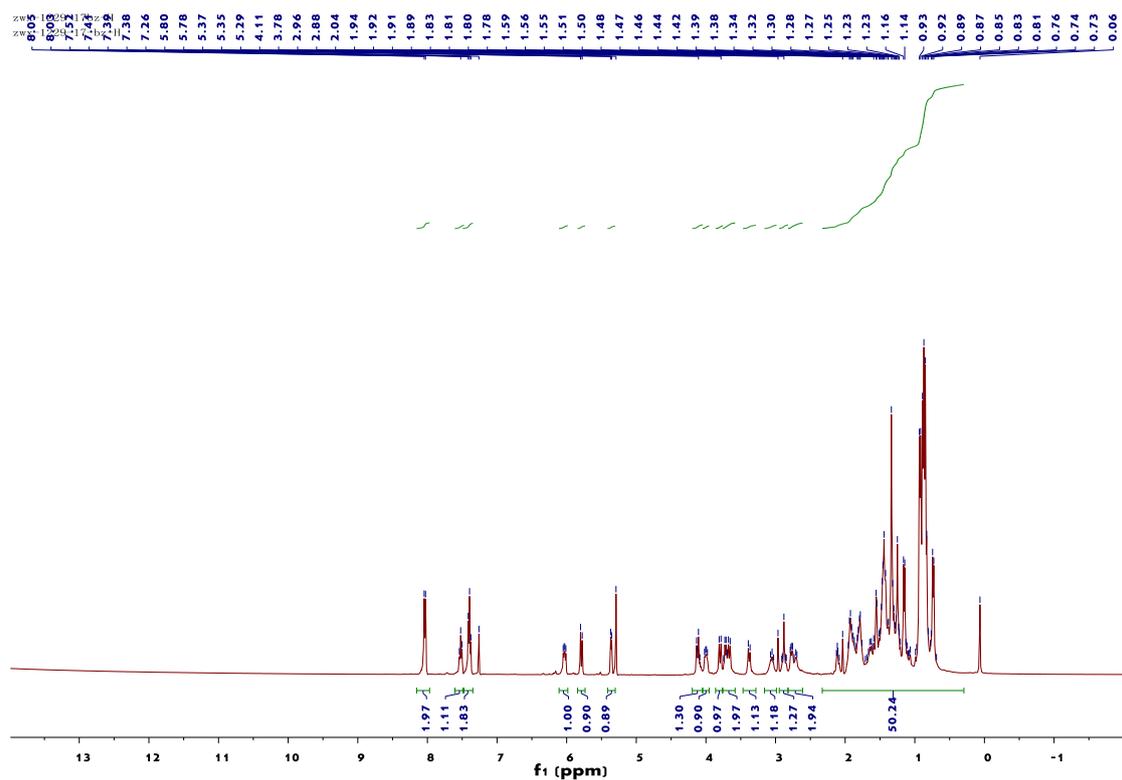
intermediate 8



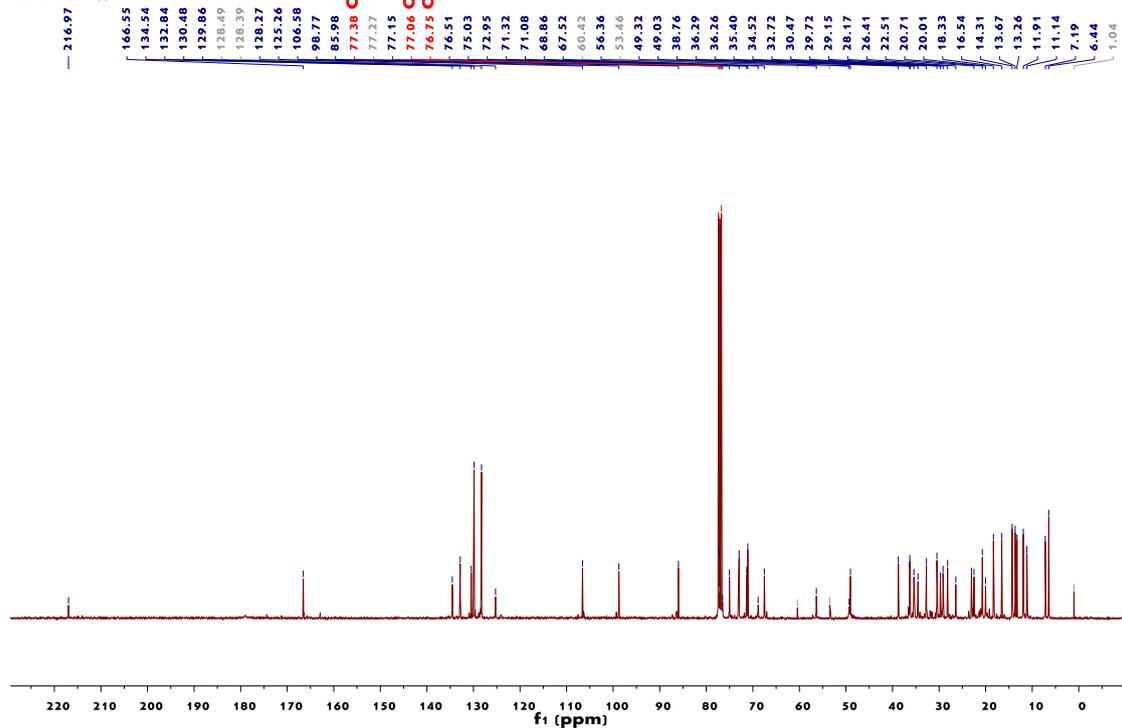
intermediate 9



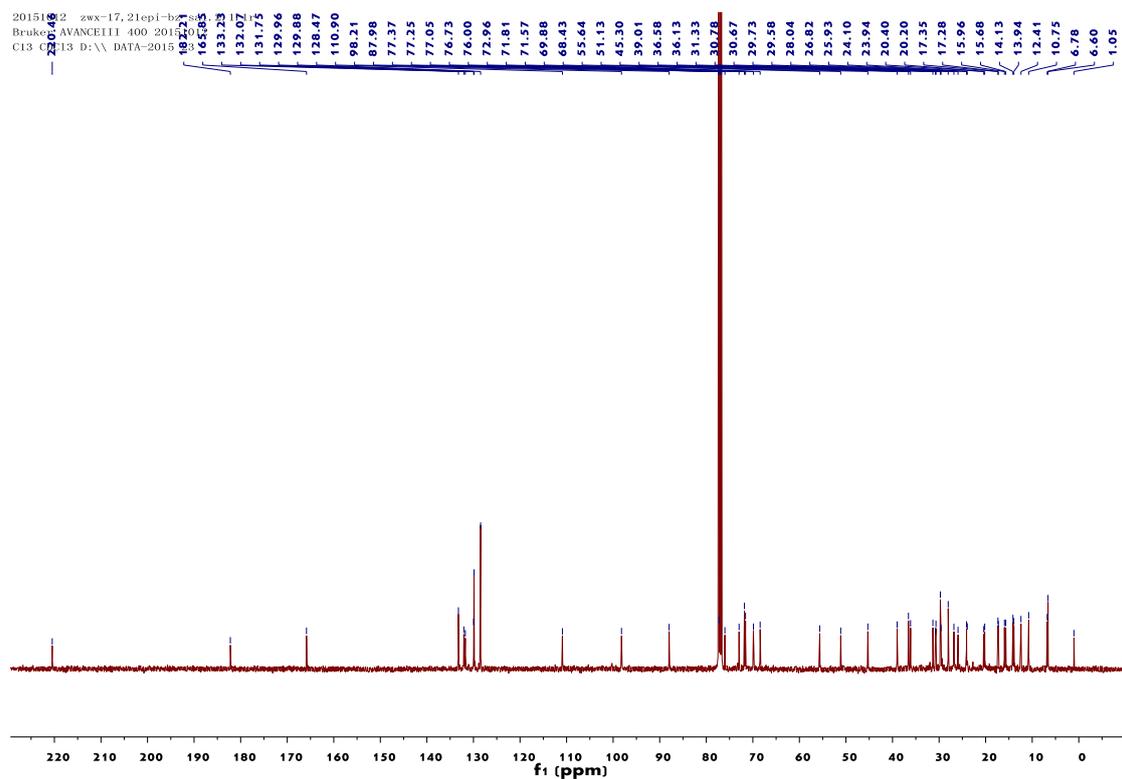
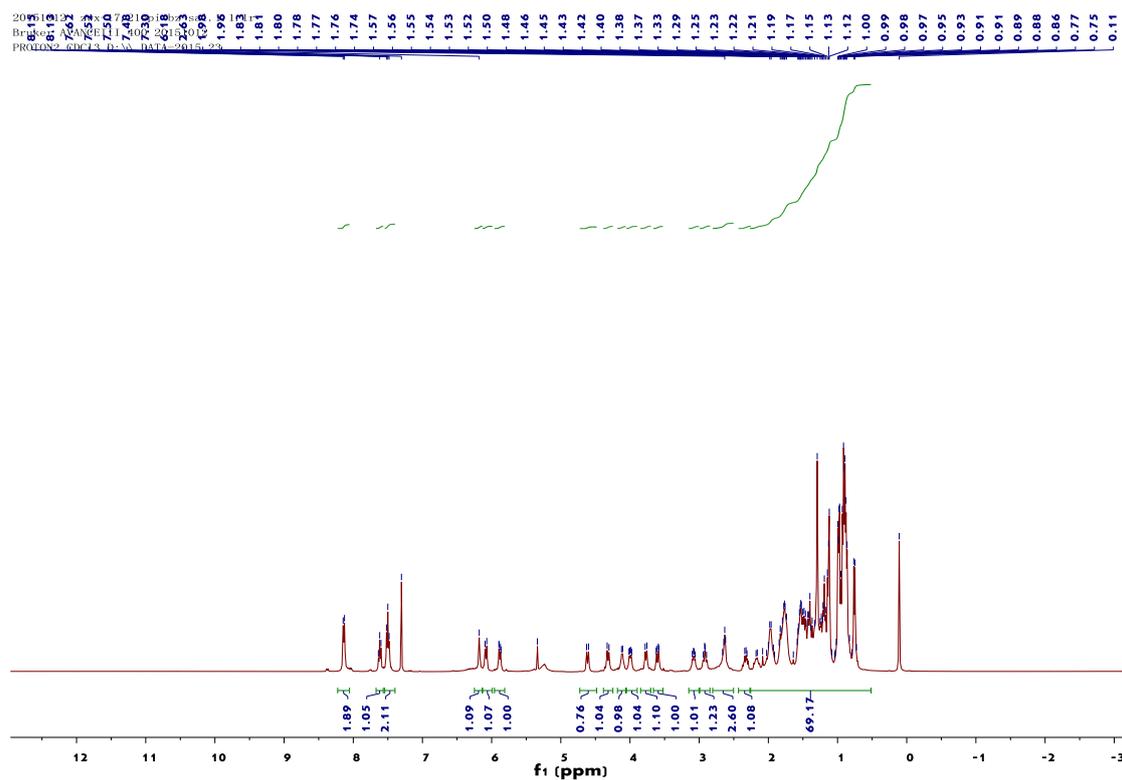
17-*epi*-20-*O*-Bz-salinomycin sodium salt (4)



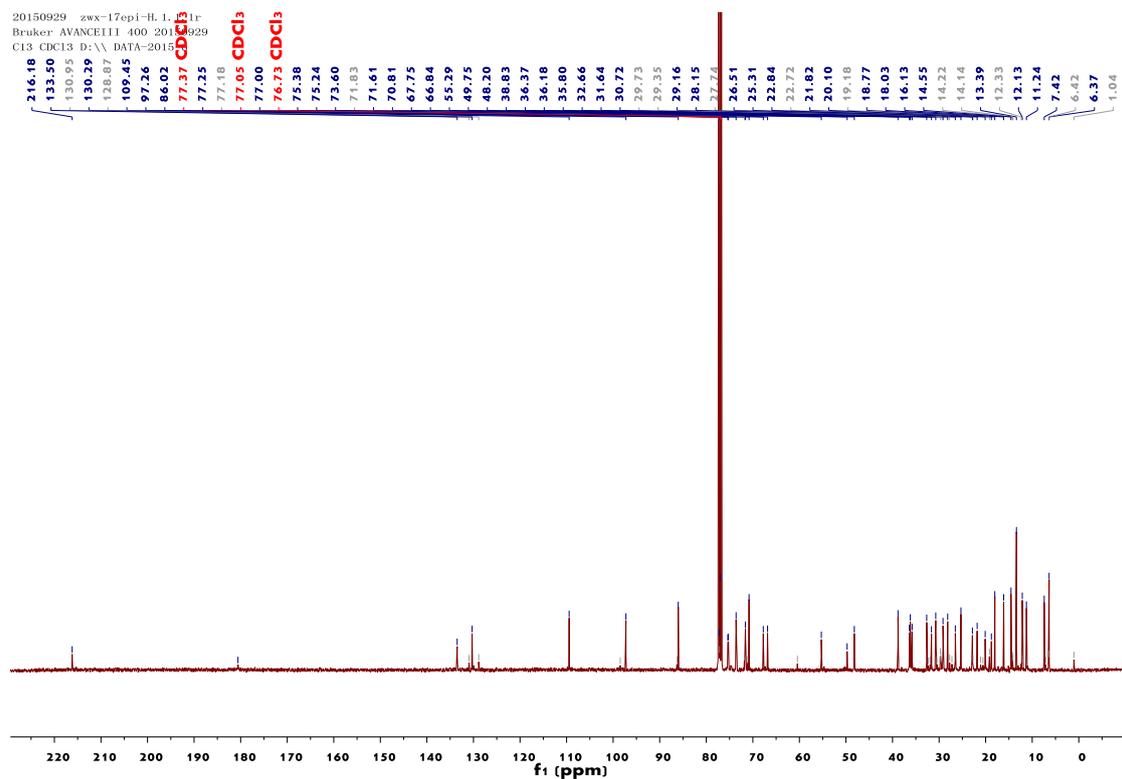
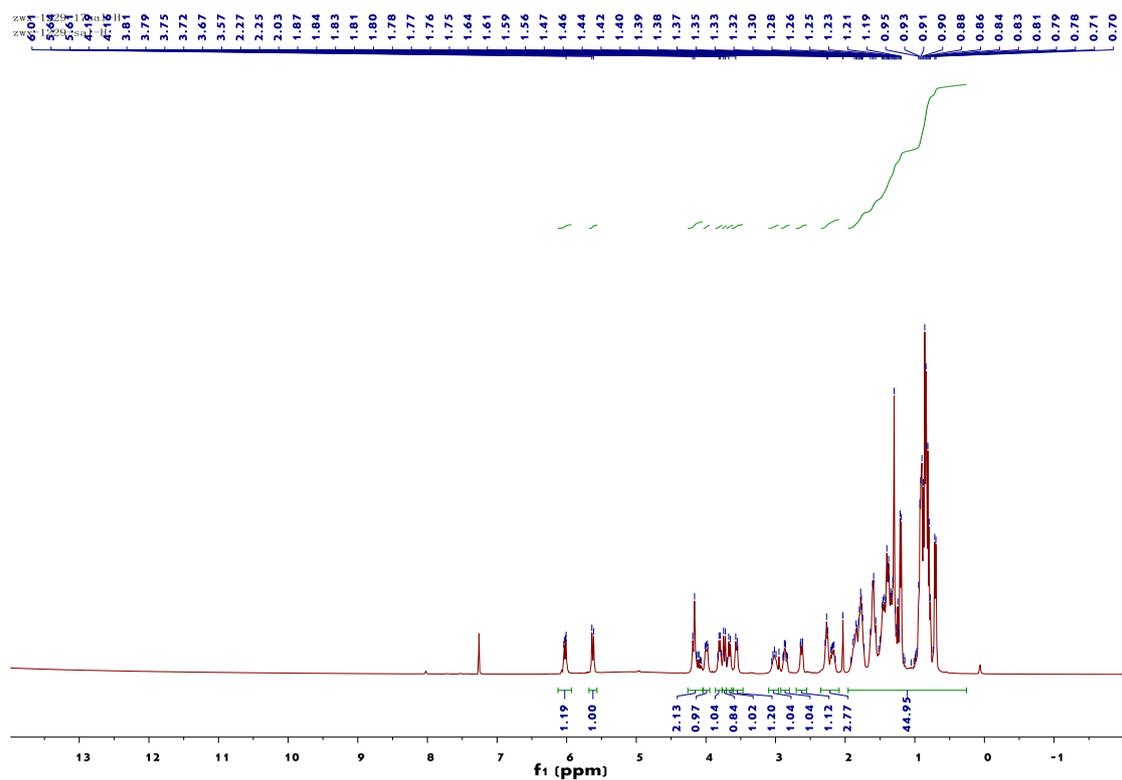
20150104 zwx-17epi-Bz.1.1.1r  
 Bruker AVANCEIII 400 20150104  
 C13-CDCl3 D:\DATA-2014 45



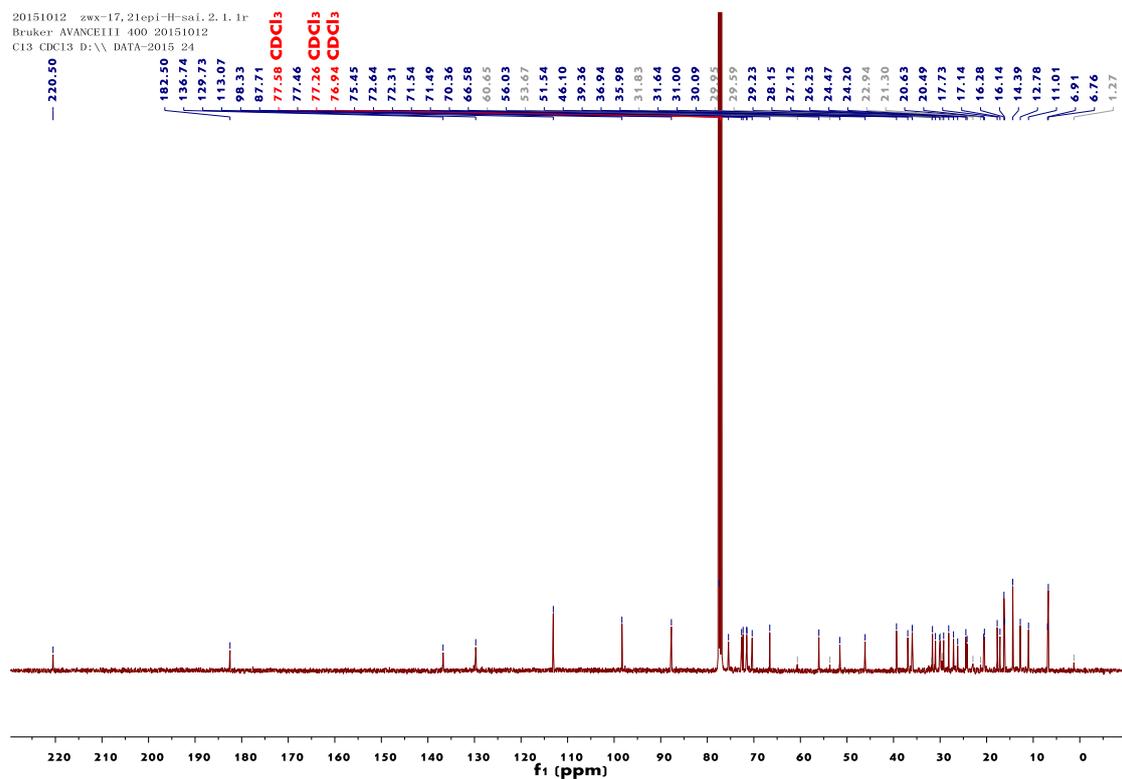
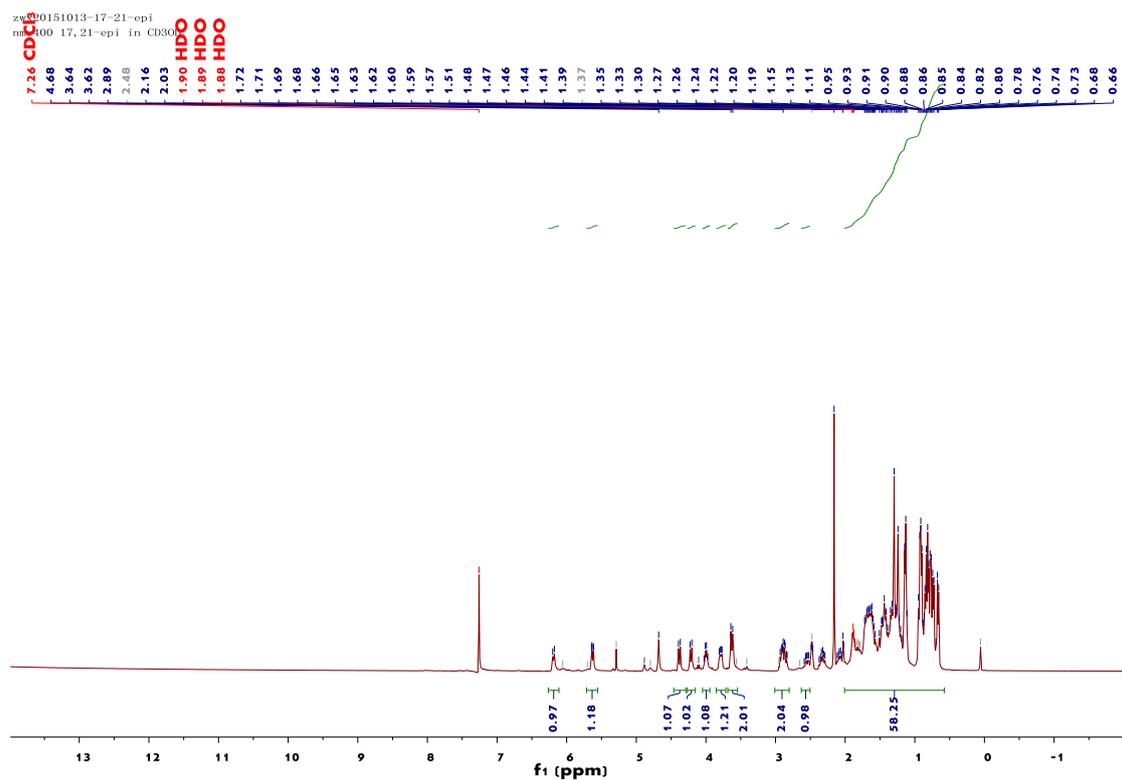
17,21-di-epi-20-O-Bz-salinomycin sodium salt (5)



17-epi-salinomycin sodium salt (2)



17, 21-di-epi-salinomycin sodium salt (3)



**Crystal data and structure refinement details of compound 9:**

A suitable crystal was selected and analyzed on a Xcalibur, Atlas, Gemini ultra diffractometer. The crystal was kept at 293(2) K during data collection. Using Olex2, the structure was solved with the Superflip structure solution program using Charge Flipping and refined with the ShelXL refinement package using Least Squares minimisation.

**Crystal Data** for  $C_{55}H_{88}O_{11}Si$  ( $M = 953.34$  g/mol): monoclinic, space group  $P2_1$  (no. 4),  $a=12.1784(3)$  Å  $b=10.4550(2)$  Å  $c=22.4818(4)$  Å  $\alpha=90.0000^\circ$   $\beta=99.8798(18)^\circ$   $\gamma=90.0000^\circ$   $V=2820.05(10)$  Å<sup>3</sup>  $Z=2$ ,  $\mu(\text{CuK}\alpha) = 0.803$  mm<sup>-1</sup>,  $D_{\text{calc}} = 1.123$  g/cm<sup>3</sup>, 41478 reflections measured ( $7.368^\circ \leq 2^\theta \leq 134.002^\circ$ ), 10003 unique ( $R_{\text{int}} = 0.0390$ ,  $R_{\text{sigma}} = 0.0221$ ) which were used in all calculations. The final  $R_1$  was 0.0447 ( $I > 2\sigma(I)$ ) and  $wR_2$  was 0.1313 (all data).

Atomic parameters:

Atom	x/a	y/b	z/c	U [Å <sup>2</sup> ]
O1	-0.7429(2)	-0.4023(2)	-0.58095(9)	
O7	-0.8458(2)	-0.5181(3)	-0.66126(9)	
O41	-0.5484(2)	-0.5023(2)	-0.97813(9)	
O38	-0.7989(2)	-0.3382(3)	-0.46195(9)	
O18	-0.6722(2)	-0.6807(2)	-0.86992(10)	
H18	-0.60440	-0.68960	-0.86480	0.1080
O28	-0.7669(2)	0.0078(2)	-0.71355(10)	
O24	-0.8571(2)	-0.2301(2)	-0.61603(10)	
O20	-0.6904(3)	-0.4000(3)	-0.79542(11)	
O49	-0.3543(3)	-0.4861(3)	-0.86019(11)	
C2	-0.8111(4)	-0.5119(4)	-0.59784(14)	
O51	-0.4361(2)	-0.6776(3)	-0.86476(12)	
O40	-0.9389(3)	-0.2849(3)	-0.41476(12)	
C6	-0.7921(3)	-0.2879(3)	-0.56473(14)	
C5	-0.8704(3)	-0.3112(4)	-0.51975(15)	
H5	-0.91470	-0.23420	-0.51620	0.0920
C11	-0.7611(3)	-0.5298(3)	-0.69843(13)	
H11	-0.71740	-0.45060	-0.69510	0.0780
C46	-0.6173(3)	-0.6130(3)	-0.99444(12)	
H46	-0.58030	-0.68710	-0.97300	0.0730
C39	-0.8452(4)	-0.3221(3)	-0.41287(15)	
C13	-0.7640(3)	-0.4780(3)	-0.80978(14)	
C12	-0.8234(3)	-0.5429(3)	-0.76432(13)	
H12	-0.82600	-0.63440	-0.77400	0.0780
C15	-0.7025(3)	-0.5641(3)	-0.90184(12)	
H15	-0.64000	-0.50420	-0.89260	0.0750
C16	-0.7248(3)	-0.5887(3)	-0.97073(13)	
H16	-0.75830	-0.51090	-0.99030	0.0760
C42	-0.4391(3)	-0.5073(4)	-0.99249(15)	
H42	-0.40670	-0.42190	-0.98470	0.0880
C32	-0.8695(3)	0.0122(3)	-0.81580(17)	

C29	-0.8191(3)	-0.1086(3)	-0.69893(15)	
H29	-0.77610	-0.18080	-0.71070	0.0830
C60	-0.7678(4)	-0.3542(3)	-0.35677(15)	
C35	-0.9293(3)	0.1234(3)	-0.80229(19)	
H35A	-0.91430	0.13890	-0.75960	0.1160
H35B	-1.00780	0.10980	-0.81510	0.1160
H35C	-0.90590	0.19590	-0.82310	0.1160
C45	-0.6303(3)	-0.6389(3)	-1.06252(13)	
H45	-0.66820	-0.72140	-1.07070	0.0840
C47	-0.3630(3)	-0.6011(4)	-0.95101(16)	
H47	-0.37950	-0.68810	-0.96620	0.0940
C48	-0.3888(3)	-0.5934(4)	-0.88814(16)	
C14	-0.8014(3)	-0.5089(4)	-0.87696(13)	
H14	-0.86090	-0.57300	-0.88100	0.0820
C33	-0.7542(3)	0.0163(4)	-0.77588(15)	
H33	-0.72330	0.10120	-0.78130	0.0900
C27	-0.8092(4)	-0.1098(4)	-0.63073(16)	
C3	-0.9172(4)	-0.5094(5)	-0.57390(17)	
H3	-0.96760	-0.57550	-0.58510	0.1210
C30	-0.9361(3)	-0.1171(4)	-0.73425(17)	
H30A	-0.98130	-0.04880	-0.72210	0.0940
H30B	-0.96940	-0.19810	-0.72600	0.0940
C4	-0.9452(4)	-0.4209(5)	-0.53812(17)	
H4	-1.01260	-0.42700	-0.52420	0.1130
C21	-0.9456(3)	-0.4951(4)	-0.77461(16)	
H21A	-0.94960	-0.41790	-0.75110	0.0970
H21B	-0.96790	-0.47290	-0.81690	0.0970
C31	-0.9329(3)	-0.1064(4)	-0.80153(16)	
H31A	-0.89740	-0.18190	-0.81460	0.0900
H31B	-1.00850	-0.10270	-0.82380	0.0900
C25	-0.6960(4)	-0.1961(4)	-0.54315(16)	
H25A	-0.71210	-0.14250	-0.51050	0.1000
H25B	-0.62720	-0.24230	-0.52970	0.1000
C17	-0.8083(3)	-0.6970(4)	-0.98692(15)	
H17A	-0.82640	-0.70490	-1.03000	0.1140
H17B	-0.87480	-0.67860	-0.97090	0.1140
H17C	-0.77630	-0.77560	-0.97000	0.1140
C10	-0.6825(4)	-0.6413(4)	-0.67906(16)	
H10	-0.72490	-0.72060	-0.68810	0.1050
C26	-0.6888(4)	-0.1174(4)	-0.59879(17)	
H26A	-0.64240	-0.15910	-0.62400	0.1050
H26B	-0.65910	-0.03290	-0.58790	0.1050

C64	-0.7314(8)	-0.3668(6)	-0.24852(19)	
H64	-0.75480	-0.35630	-0.21160	0.1560
C43	-0.4436(4)	-0.5337(5)	-1.05920(16)	
H43A	-0.36870	-0.54730	-1.06710	0.1050
H43B	-0.47470	-0.46000	-1.08240	0.1050
C65	-0.8045(5)	-0.3403(4)	-0.30196(16)	
H65	-0.87700	-0.31380	-0.30090	0.1240
C50	-0.3734(4)	-0.4725(5)	-0.79751(18)	
H50A	-0.35490	-0.55170	-0.77570	0.1170
H50B	-0.45130	-0.45350	-0.79720	0.1170
C44	-0.5146(4)	-0.6511(4)	-1.07912(15)	
H44A	-0.52150	-0.66140	-1.12250	0.1010
H44B	-0.47820	-0.72660	-1.06000	0.1010
C54	-0.3023(4)	-0.3669(5)	-0.76795(18)	
H54A	-0.32720	-0.28750	-0.78820	0.1130
H54B	-0.22640	-0.38170	-0.77410	0.1130
C8	-0.7377(6)	-0.6269(5)	-0.57594(18)	
H8	-0.78340	-0.70390	-0.58510	0.1330
C34	-0.6715(3)	-0.0790(5)	-0.79158(17)	
H34A	-0.60920	-0.08410	-0.75910	0.1300
H34B	-0.64620	-0.05240	-0.82780	0.1300
H34C	-0.70630	-0.16140	-0.79780	0.1300
C61	-0.6602(4)	-0.3964(4)	-0.35704(18)	
H61	-0.63480	-0.40600	-0.39350	0.1080
C59	-0.7002(4)	-0.5376(5)	-1.10077(15)	
H59A	-0.77550	-0.54120	-1.09350	0.1370
H59B	-0.69920	-0.55350	-1.14270	0.1370
H59C	-0.66970	-0.45440	-1.09010	0.1370
C62	-0.5907(5)	-0.4242(5)	-0.3038(2)	
H62	-0.51910	-0.45450	-0.30440	0.1290
C36	-0.8583(4)	0.0208(5)	-0.88194(18)	
H36A	-0.81190	0.09390	-0.88710	0.1070
H36B	-0.81980	-0.05510	-0.89230	0.1070
C23	-0.5855(5)	-0.6443(6)	-0.7134(2)	
H23A	-0.61280	-0.66280	-0.75520	0.1700
H23B	-0.53360	-0.70930	-0.69670	0.1700
H23C	-0.54890	-0.56270	-0.71010	0.1700
C57	-0.2656(5)	-0.4965(5)	-0.6443(2)	
H57A	-0.32460	-0.55710	-0.65610	0.1610
H57B	-0.25680	-0.48190	-0.60160	0.1610
H57C	-0.19740	-0.52960	-0.65410	0.1610
C19	-0.8465(5)	-0.3870(5)	-0.91012(17)	

H19A	-0.90250	-0.34990	-0.89020	0.1570
H19B	-0.87860	-0.40760	-0.95100	0.1570
H19C	-0.78670	-0.32710	-0.91000	0.1570
C37	-0.8735(6)	-0.0005(5)	-0.6075(2)	
H37A	-0.95190	-0.01240	-0.62110	0.1730
H37B	-0.85060	0.07940	-0.62250	0.1730
H37C	-0.85840	0.00030	-0.56410	0.1730
C9	-0.6420(5)	-0.6359(5)	-0.61069(18)	
H9A	-0.59870	-0.71200	-0.59810	0.1330
H9B	-0.59370	-0.56230	-0.60120	0.1330
C56	-0.4383(4)	-0.2849(6)	-0.6718(3)	
H56A	-0.46040	-0.21350	-0.69800	0.1650
H56B	-0.43290	-0.25840	-0.63050	0.1650
H56C	-0.49270	-0.35190	-0.68030	0.1650
C58	-0.1914(5)	-0.2257(5)	-0.6576(2)	
H58A	-0.11940	-0.26190	-0.65900	0.1620
H58B	-0.19590	-0.20190	-0.61690	0.1620
H58C	-0.20230	-0.15140	-0.68310	0.1620
C52	-0.2397(4)	-0.5729(6)	-0.9525(2)	
H52A	-0.22920	-0.56650	-0.99420	0.1240
H52B	-0.22010	-0.49100	-0.93340	0.1240
C63	-0.6261(7)	-0.4078(6)	-0.2503(2)	
H63	-0.57780	-0.42480	-0.21450	0.1560
C22	-1.0271(4)	-0.5922(6)	-0.7575(2)	
H22A	-1.02460	-0.66840	-0.78110	0.1580
H22B	-1.10100	-0.55720	-0.76530	0.1580
H22C	-1.00720	-0.61250	-0.71540	0.1580
Si55	-0.30047(9)	-0.34503(10)	-0.68460(4)	
C1	-0.9659(4)	0.0330(7)	-0.9257(2)	
H1A	-0.95020	0.04520	-0.96570	0.1930
H1B	-1.00700	0.10500	-0.91470	0.1930
H1C	-1.00920	-0.04340	-0.92460	0.1930
C1AA	-0.1630(5)	-0.6733(8)	-0.9214(3)	
H1AA	-0.17880	-0.75370	-0.94170	0.2130
H1AB	-0.17390	-0.68120	-0.88020	0.2130
H1AC	-0.08720	-0.64940	-0.92220	0.2130
C0AA	-0.6957(8)	-0.6252(6)	-0.5069(2)	
H0AA	-0.65460	-0.54790	-0.49600	0.2360
H0AB	-0.64810	-0.69760	-0.49570	0.2360
H0AC	-0.75820	-0.62900	-0.48610	0.2360

