

Supplementary Information

Sterol methyltransferases in uncultured bacteria complicate eukaryotic biomarker interpretations

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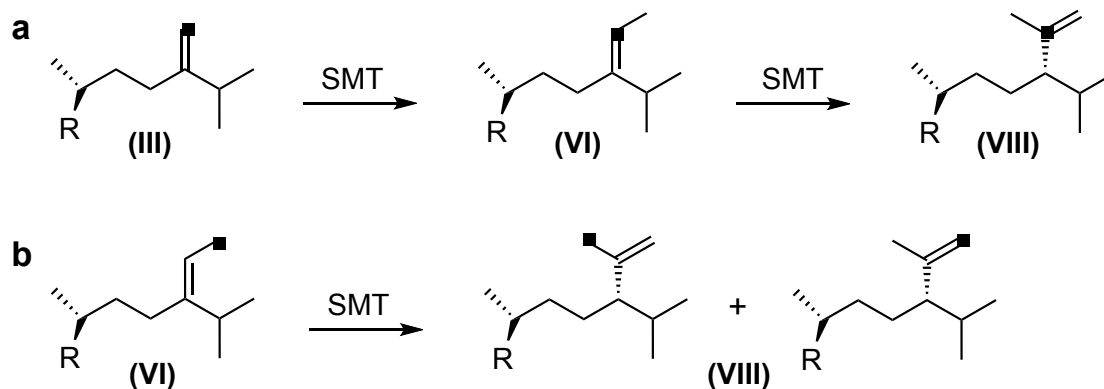
***corresponding author:** welander@stanford.edu

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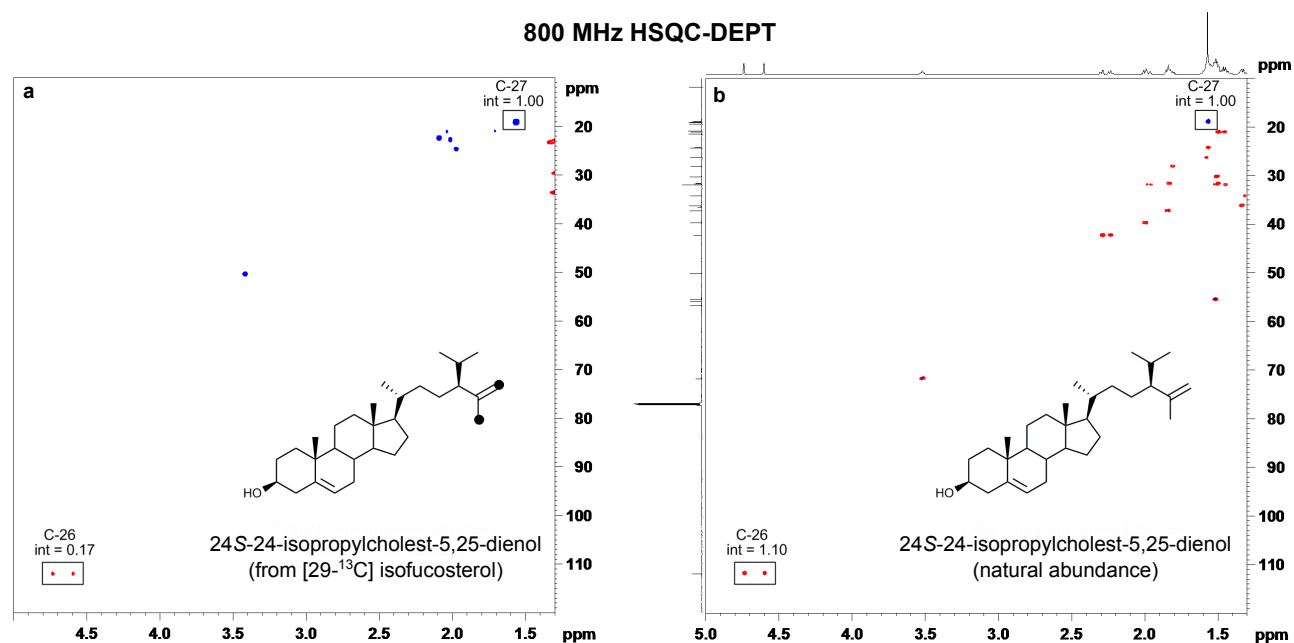
Supplementary Figures 1-7

Supplementary Tables 1-5

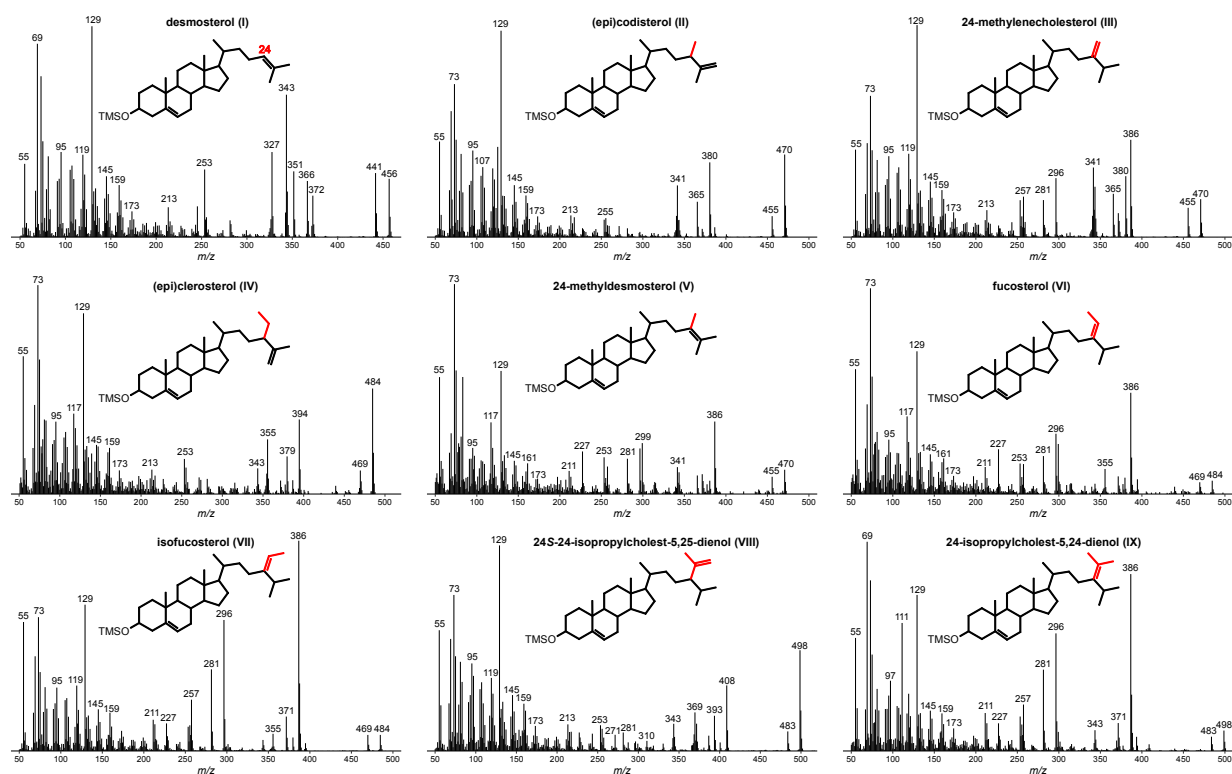
Supplementary References



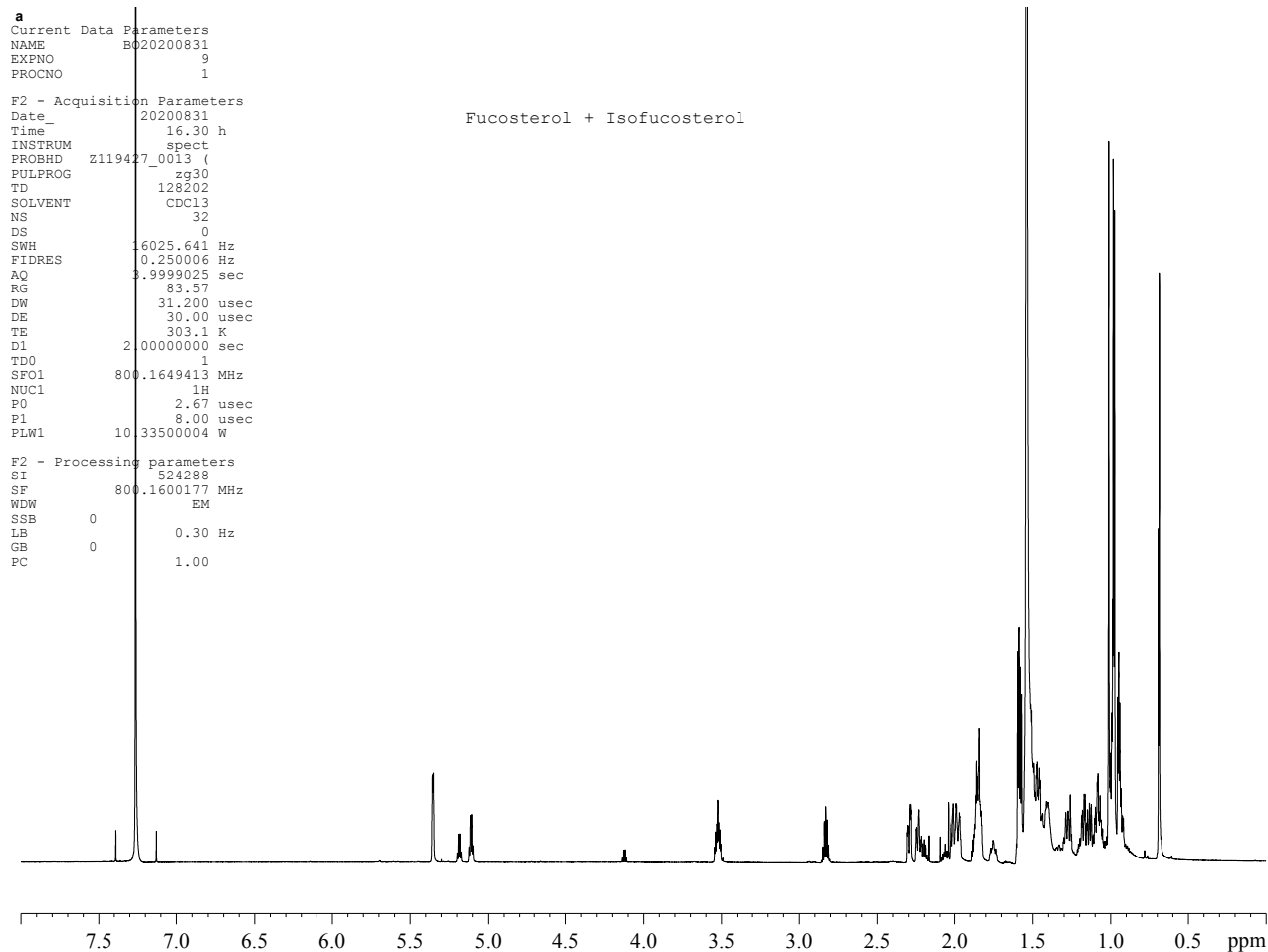
Supplementary Figure 1. ^{13}C -labeling experiments with propylating SMTs from an *Aplysina aerophoba* symbiont and a *Chlamydiae* sp. Positions of the ^{13}C labels are shown for the predominant propylation pathway proceeding from 24-methylenecholesterol (III) to isofucoesterol (VI) to 24S-24-isopropylcholest-5,25,dienol (VIII). **a** Experiments with [28- ^{13}C] 24-methylenecholesterol as the substrate. **b** Experiments with [29- ^{13}C] isofucoesterol as the substrate.



Supplementary Figure 2. **2D NMR (HSQC).** **a** 24S-24-isopropylcholest-5,25,dienol (VIII) from an in vitro reaction performed with [29-¹³C] isofucosterol and a Chlamydiae sp. MAG SMT and **b** naturally abundant 24S-24-isopropylcholest-5,25,dienol (VIII). Additional NMR spectra are shown in Supplementary Figures 4-7 and are summarized in Supplementary Table 3.



Supplementary Figure 3. **Representative mass spectra of sterols identified in the main text.** Data are from in vitro reactions performed with the “*Chlamydiae* sp., drinking water 1 of 2” SMT (**I**, **V**, **VI**, **VII**, **VIII**, **IX**) and the “*Nitrospira* sp., coral reef 1 of 2” SMT (**II**) with desmosterol as the substrate, or the “*Chondrilla nucula*” SMT with 24-methylenecholesterol as the substrate (**III**, **IV**). Total lipid extracts were derivatized to trimethylsilyl ethers. The raw data used to generate this figure are provided in Supplementary Data File 3. Detailed GC-MS methods can be found in the Methods section of the main text.



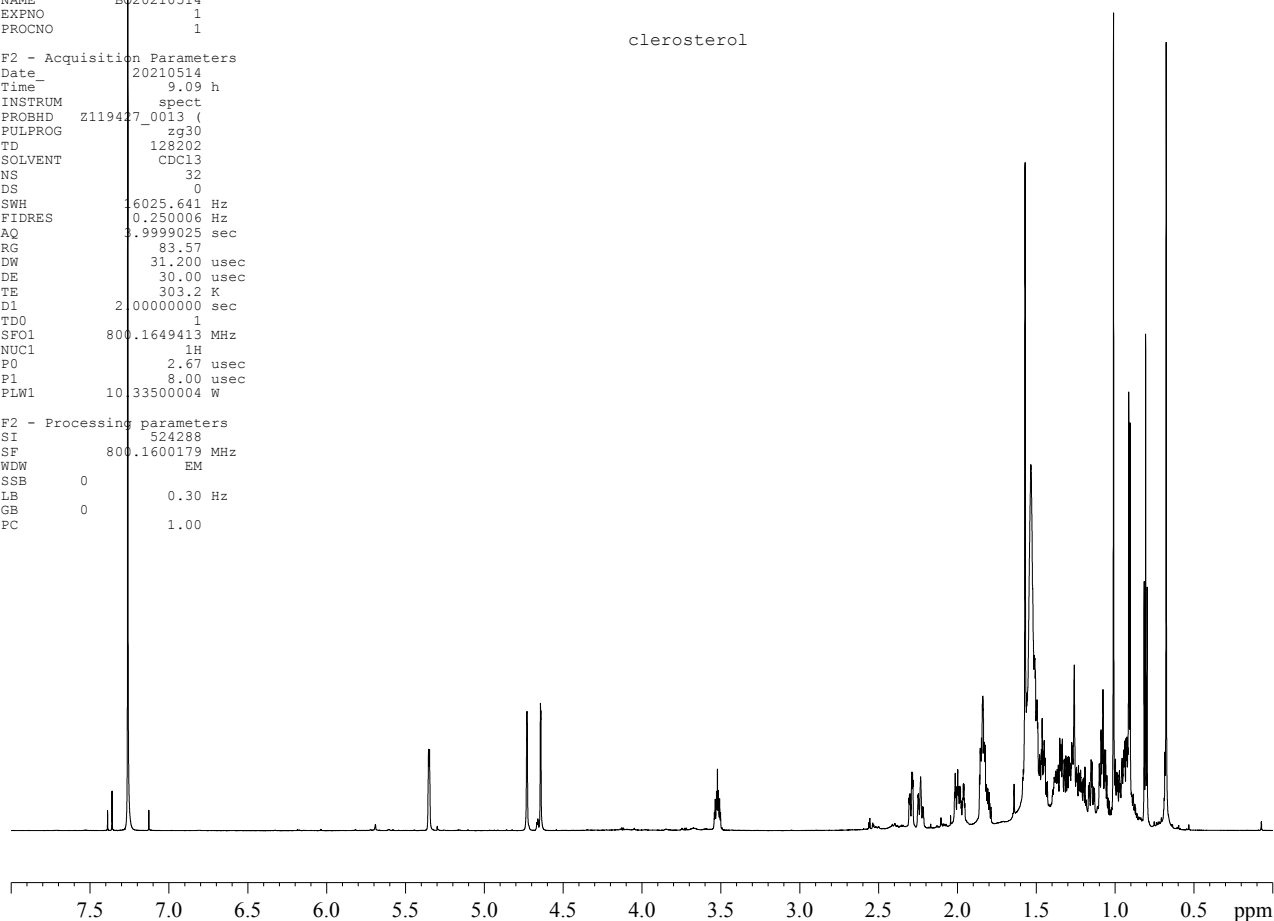
Supplementary Figure 4. **Raw 800 MHz ^1H -NMR spectra of the following sterols: a** fucosterol plus isofucosterol, **b** clerosterol, **c** epiclerosterol, **d** 24*S*-24-isopropylcholesta-5,25-dienol, **e** 24*R*-24-isopropylcholesta-5,25-dienol, **f** [26, 27, 29, 30- ^{13}C] 24-isopropylcholesta-5,24-dienol, and **g** [26, 27, 29, 30- ^{13}C] 24-isopropylcholesta-5,23-dienol plus 24*S*-24-isopropylcholesta-5,25-dienol. Chemical shifts are summarized in Supplementary Table 3.

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PROCNO 1

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FIDRES 0.250006 Hz
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DE 30.00 usec
TE 303.2 K
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P1 8.00 usec
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F2 - Processing parameters
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clerosterol

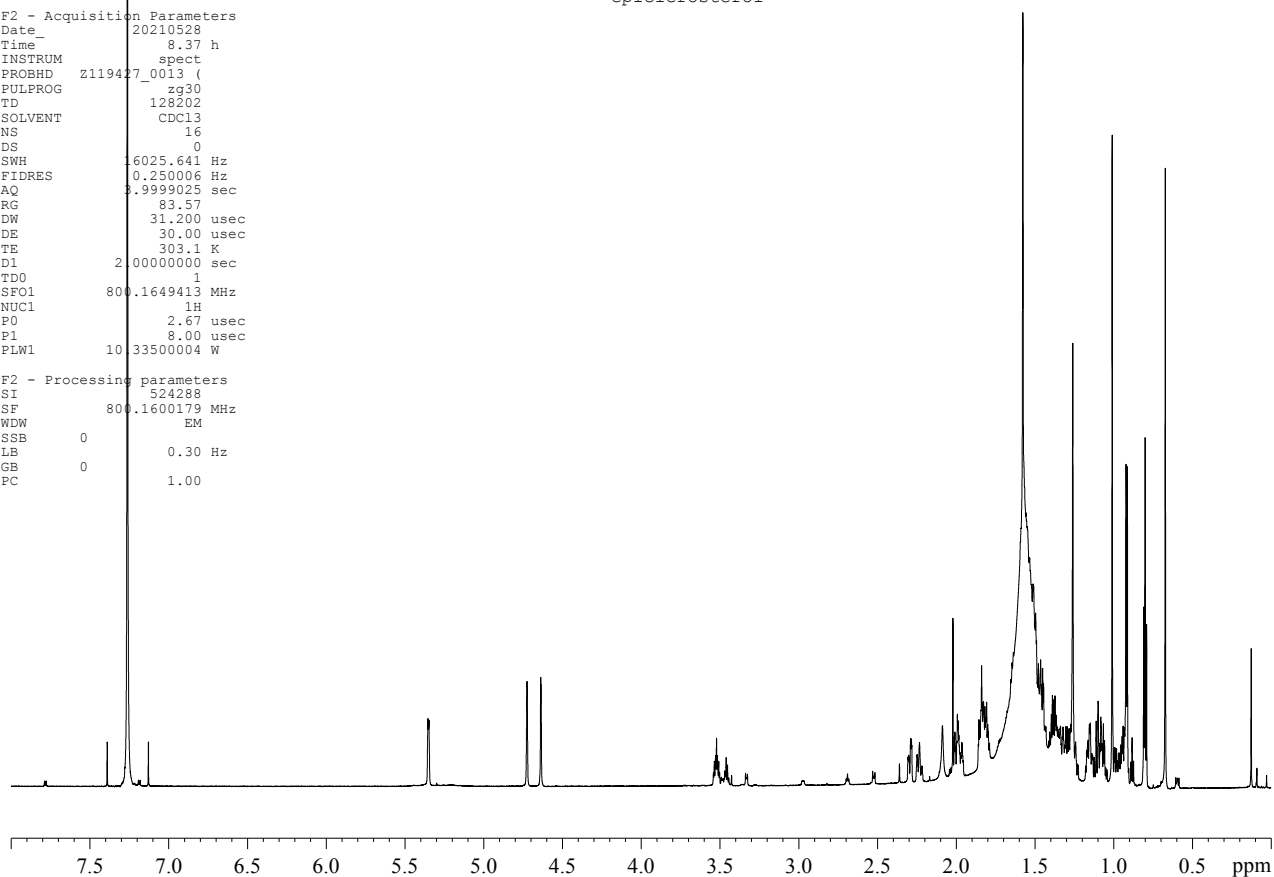


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P0 2.67 usec
P1 8.00 usec
PLW1 10.33500004 W

F2 - Processing parameters
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epiclerosterol



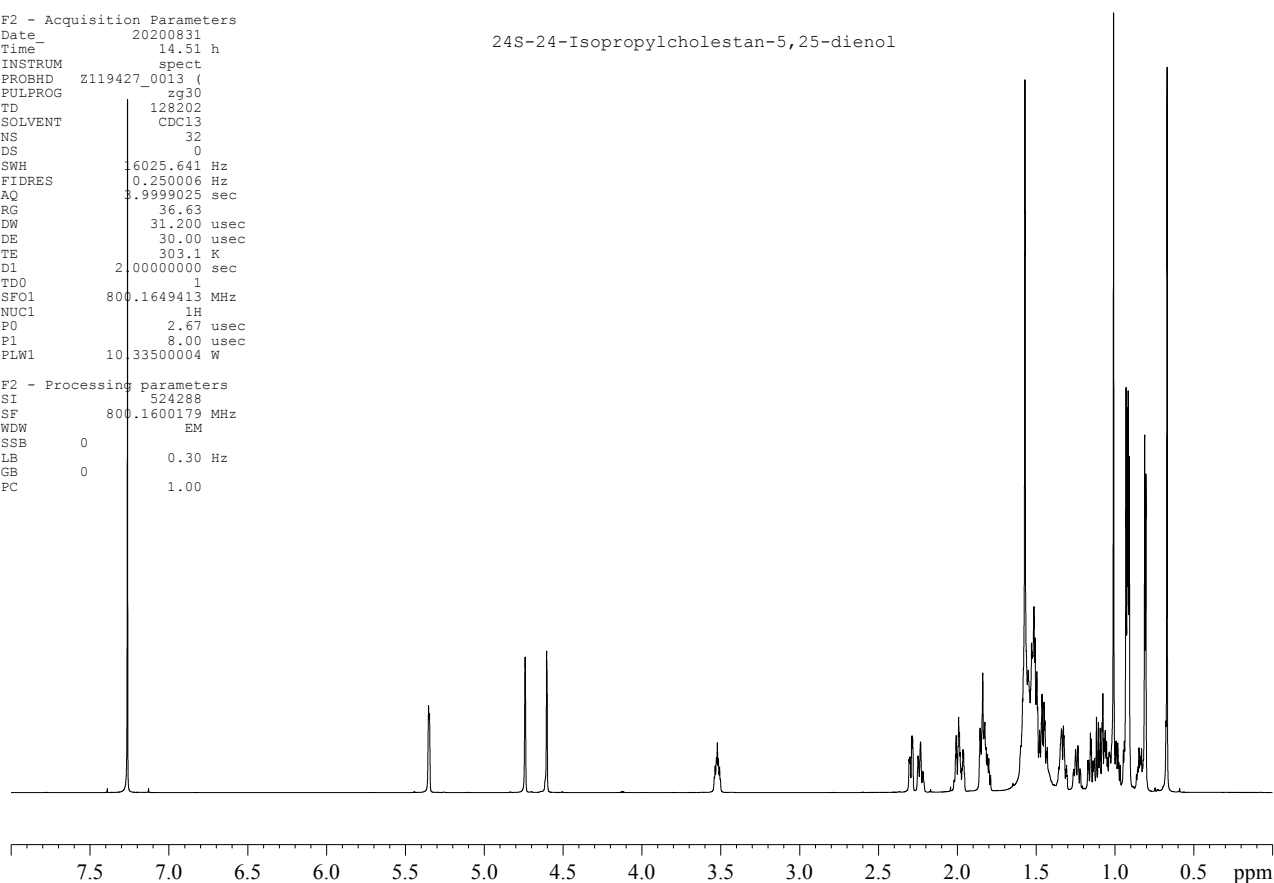
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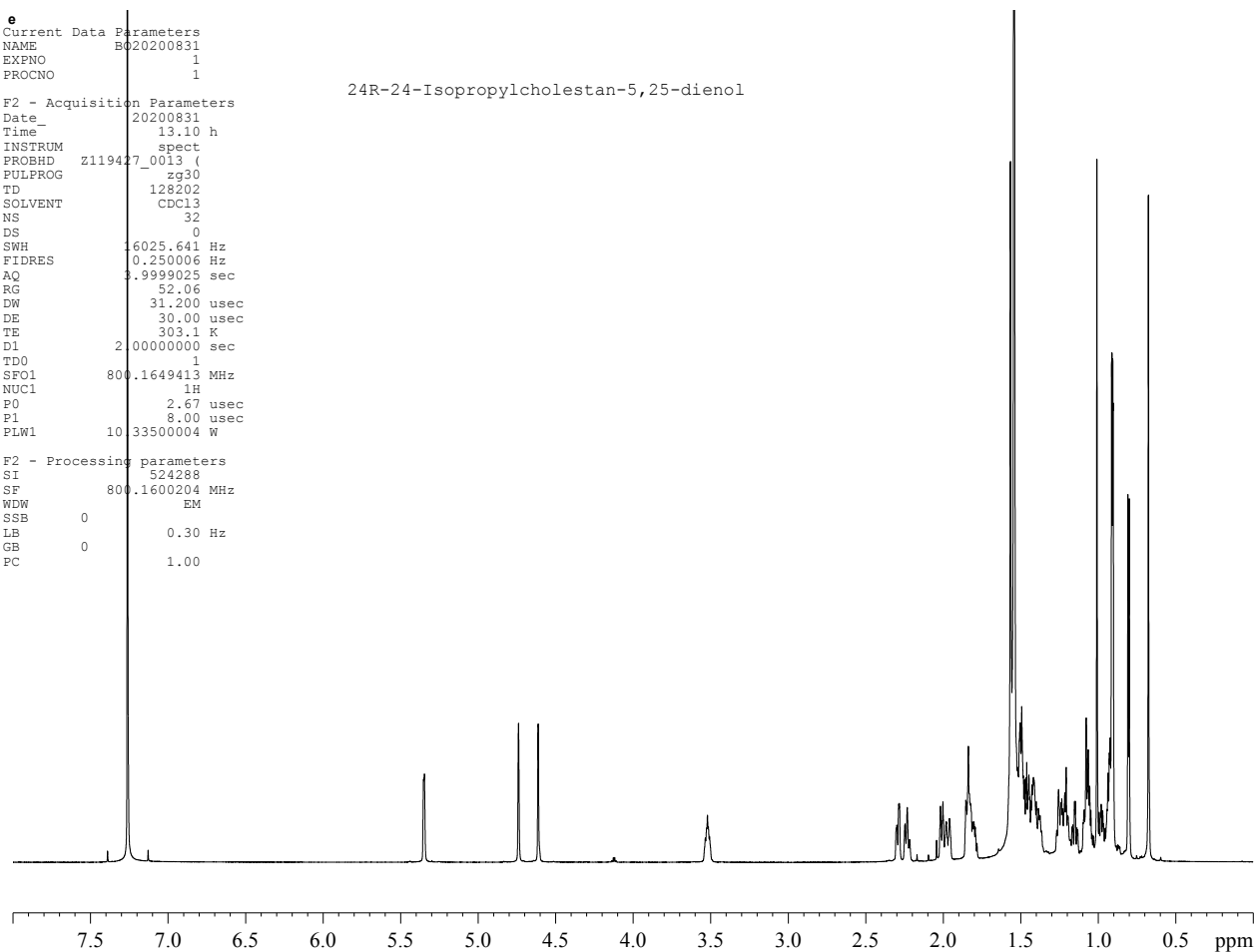


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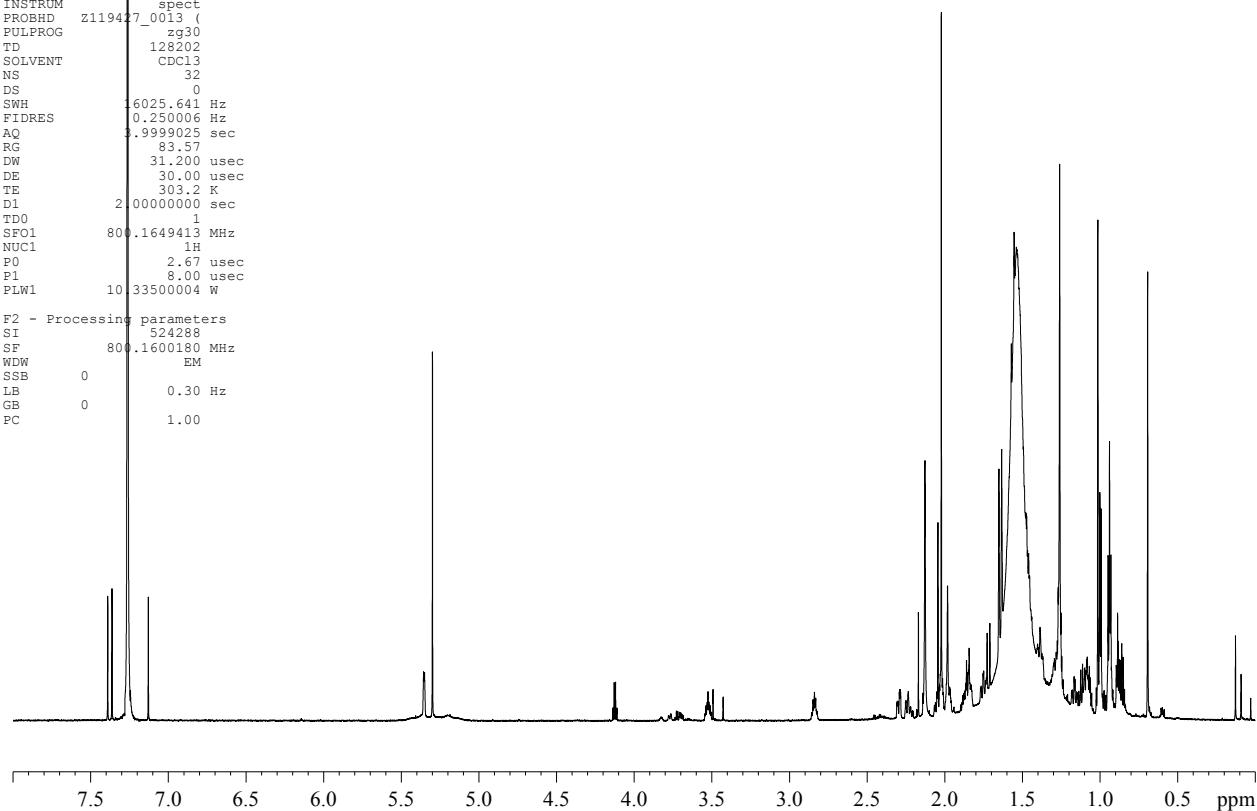
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 P1 8.00 usec
 PLW1 10.33500004 W
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26,27,29,30 - [13C] 24-Isopropylcholestan-5,24-dienol



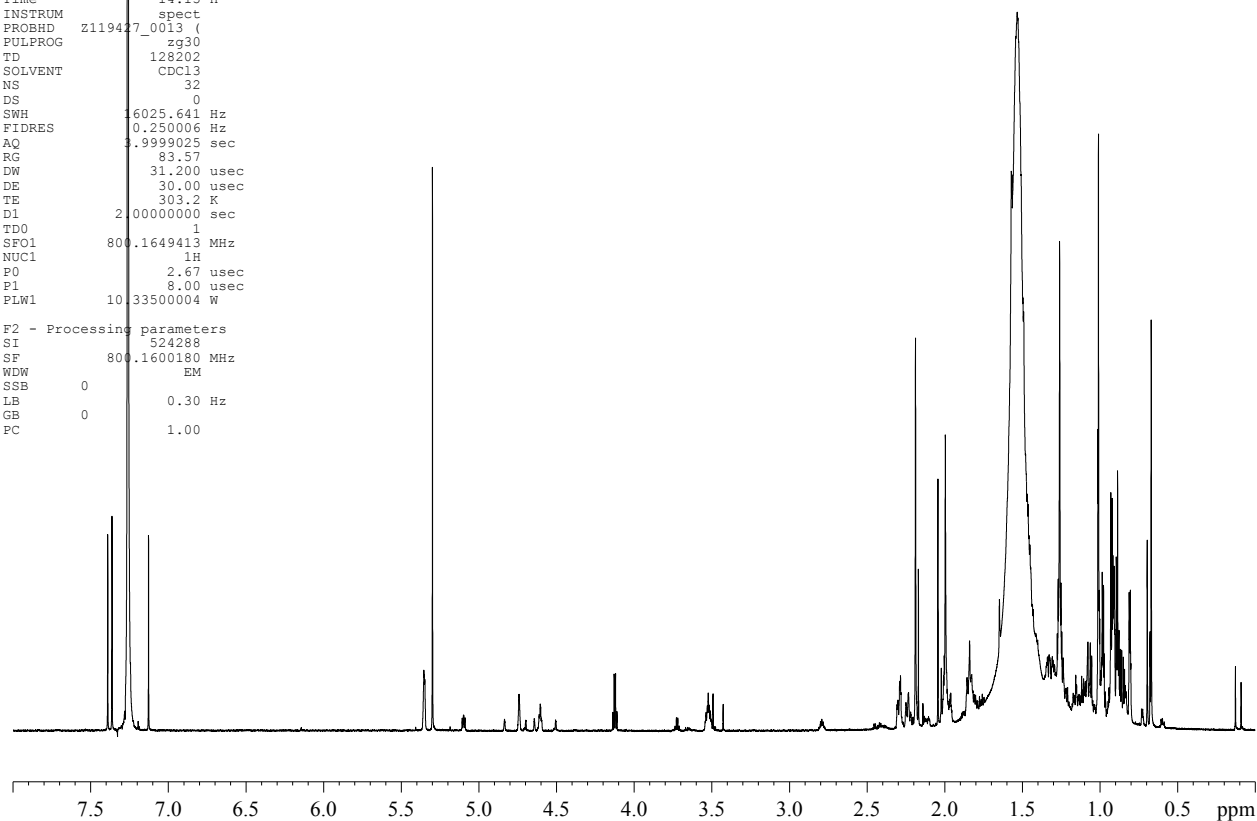
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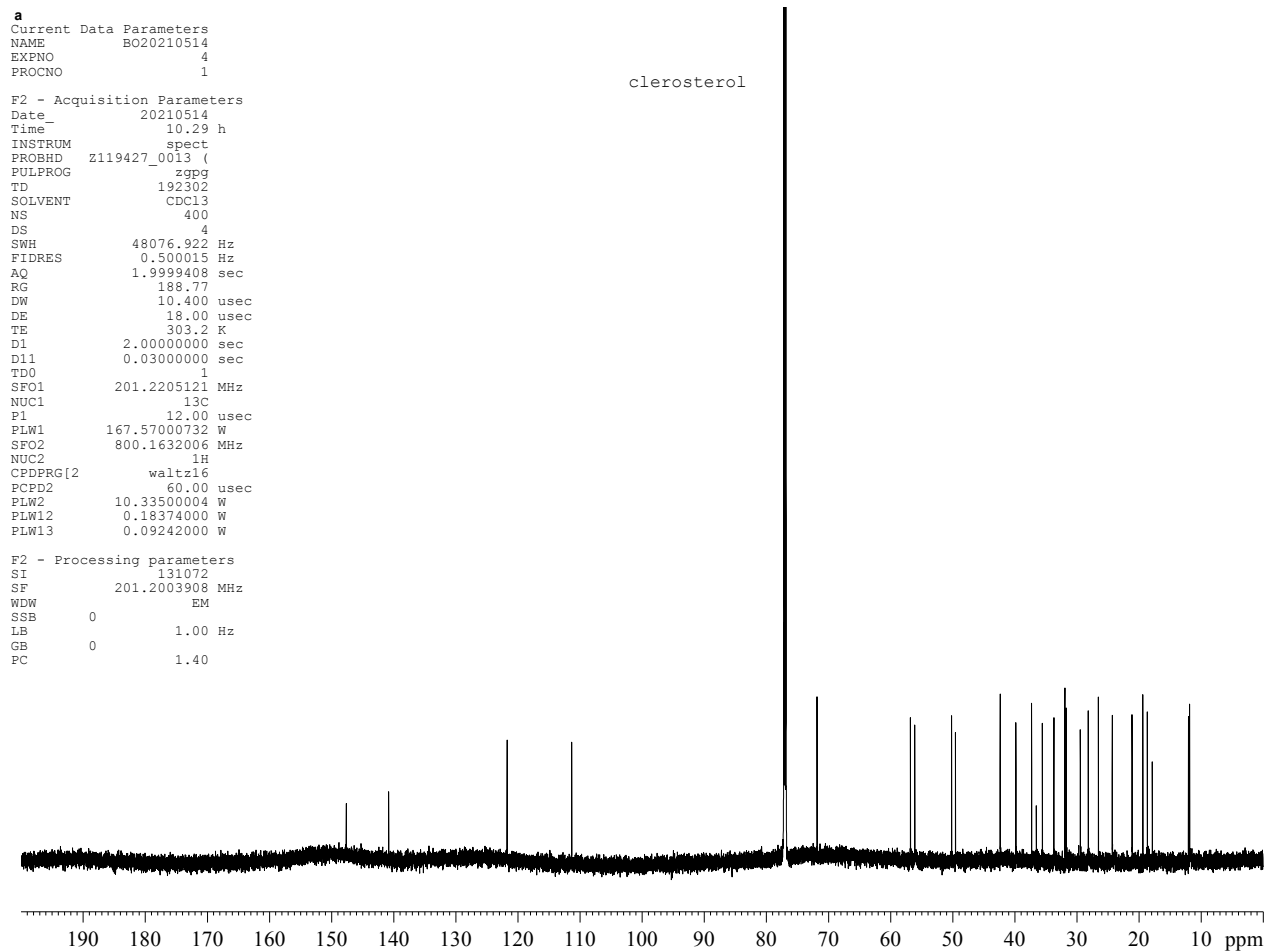
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DE 30.00 usec
TE 303.2 K
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SFO1 800.1649413 MHz
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FO 2.67 usec
P1 8.00 usec
PLW1 10.33500004 W

F2 - Processing parameters
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SF 800.1600180 MHz
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SSB 0
LB 0.30 Hz
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PC 1.00

26,27,29,30 - [13C]
24-Isopropylcholestan-5,23-dienol + 24S-24-Isopropylcholestan-5,25-dienol





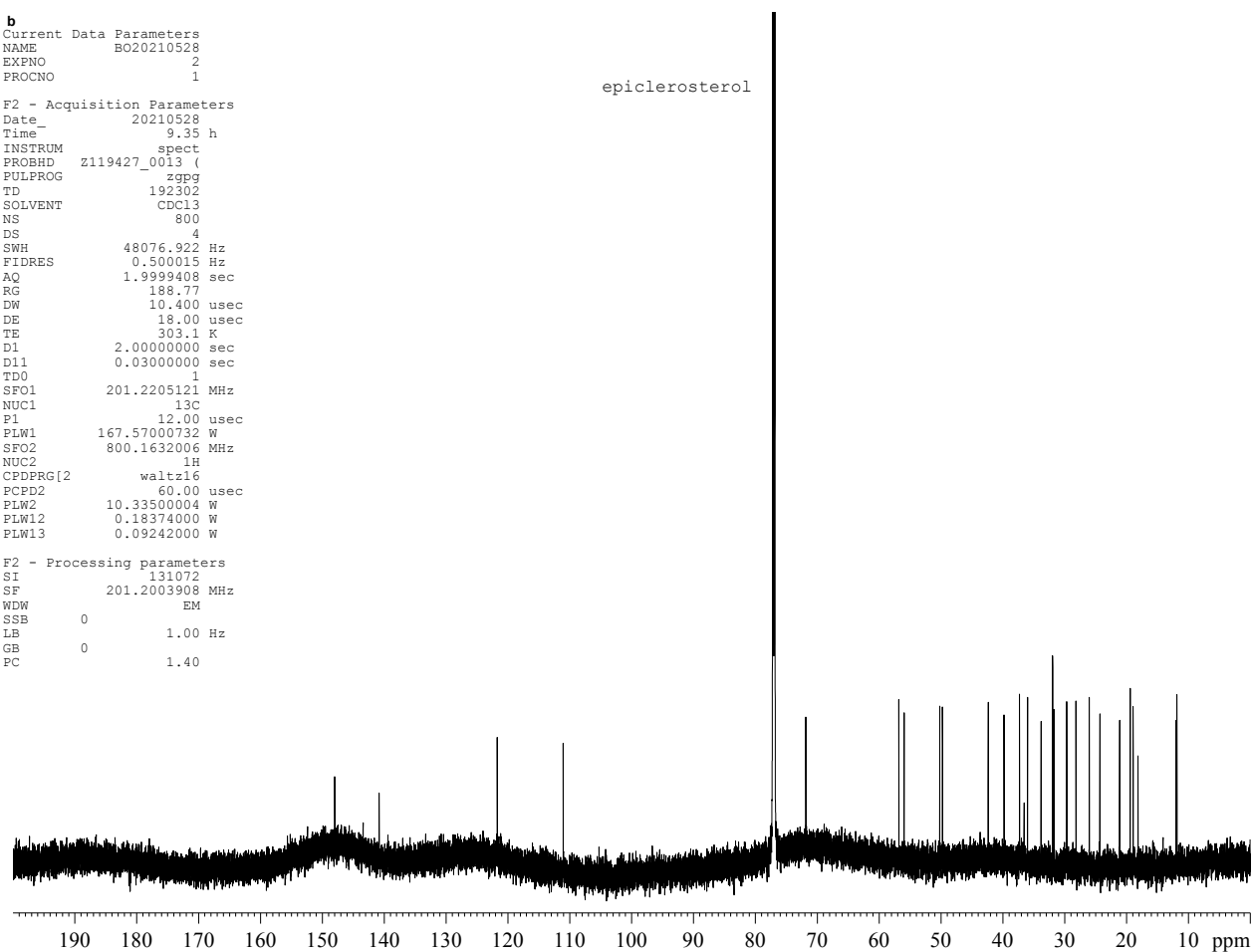
Supplementary Figure 5. Raw 201 MHz ^{13}C -NMR spectra of the following sterols: **a** clerosterol, **b** epiclerosterol, **c** 24*S*-24-isopropylcholesta-5,25-dienol **d** 24*R*-24-isopropylcholesta-5,25-dienol, **e** [26, 27, 29, 30- ^{13}C] 24-isopropylcholesta-5,24-dienol, and **f** [26, 27, 29, 30- ^{13}C] 24-isopropylcholesta-5,23-dienol plus 24*S*-24-isopropylcholesta-5,25-dienol. Chemical shifts are summarized in Supplementary Table 3.

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SFO2 800.1632006 MHz
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PLW13 0.09242000 W

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epiclerosterol

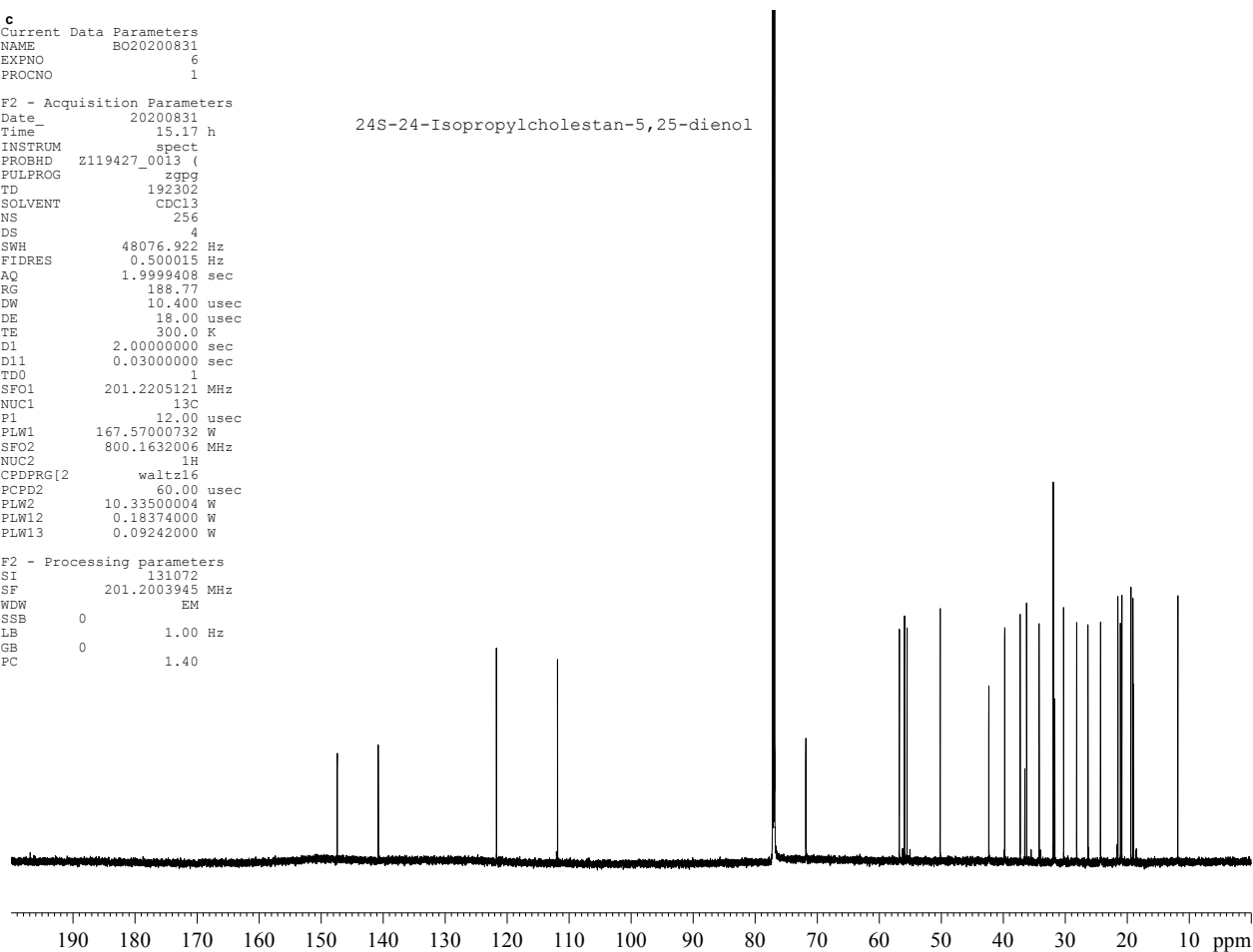


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PLW13 0.09242000 W

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24S-24-Isopropylcholestan-5,25-dienol



d

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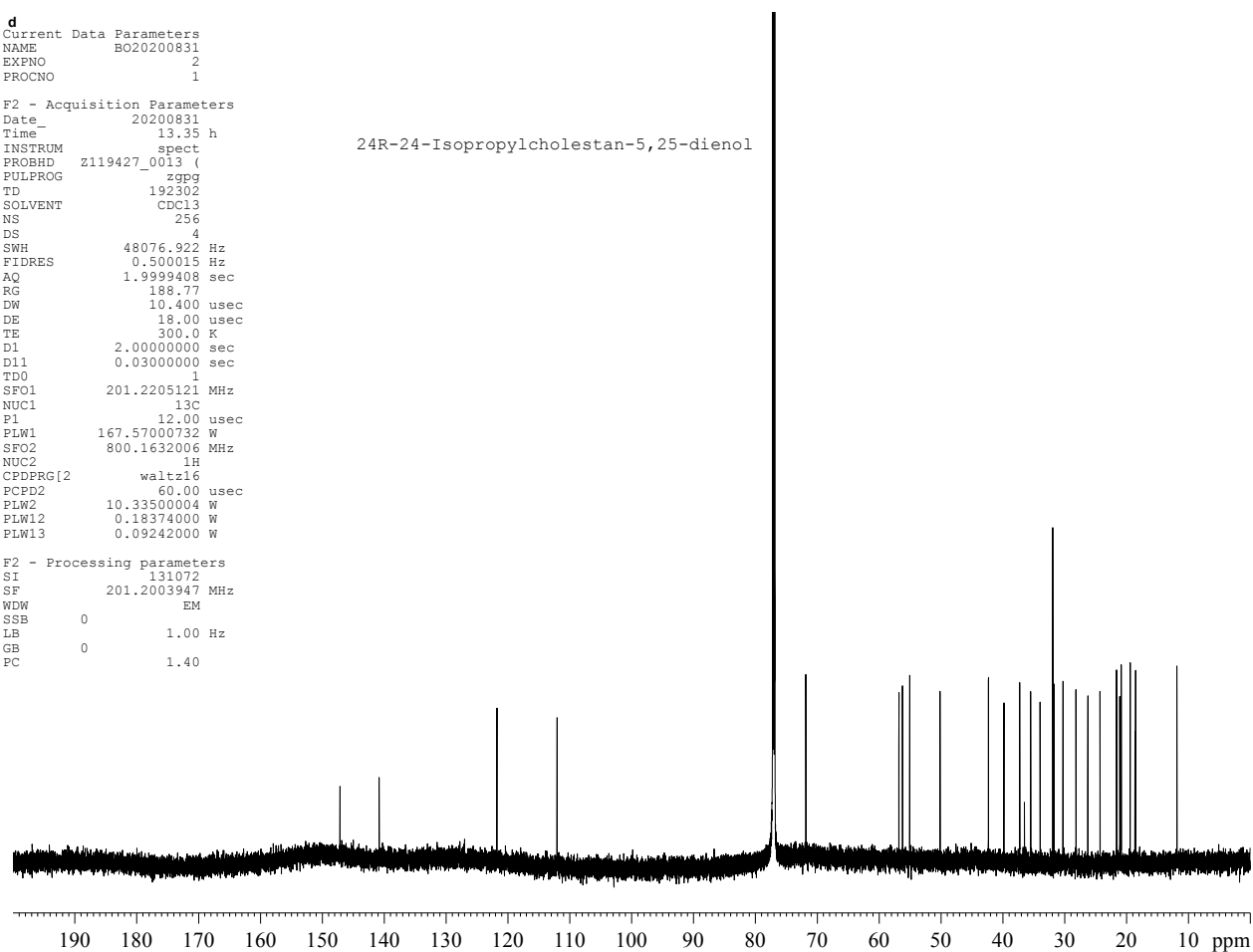
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RG 188.77
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NUC2 1H
CPDPRG[2] waltz16
PCPD2 60.00 usec
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24R-24-Isopropylcholestan-5,25-dienol

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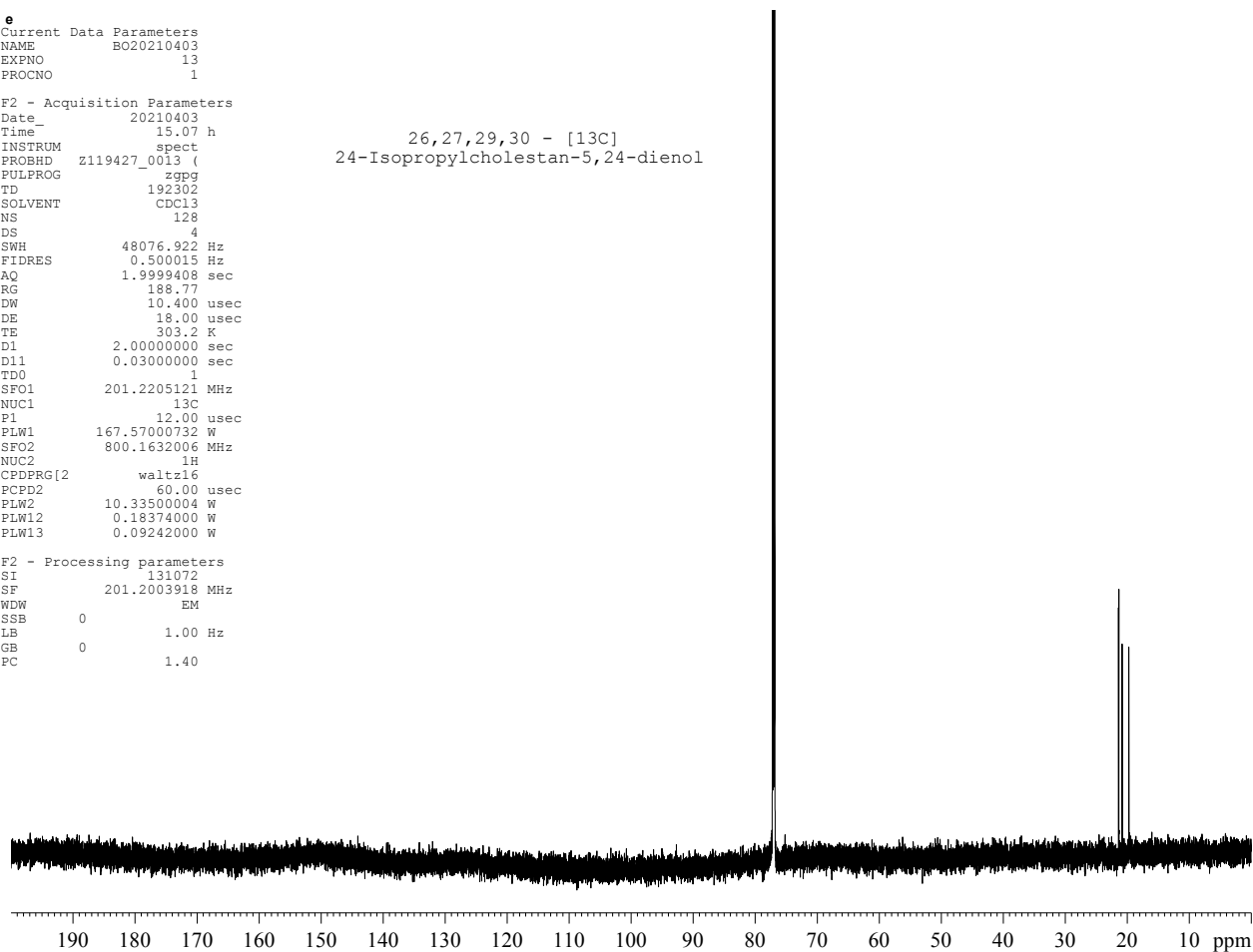


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PLW13 0.09242000 W

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26,27,29,30 - [13C]
24-Isopropylcholestan-5,24-dienol

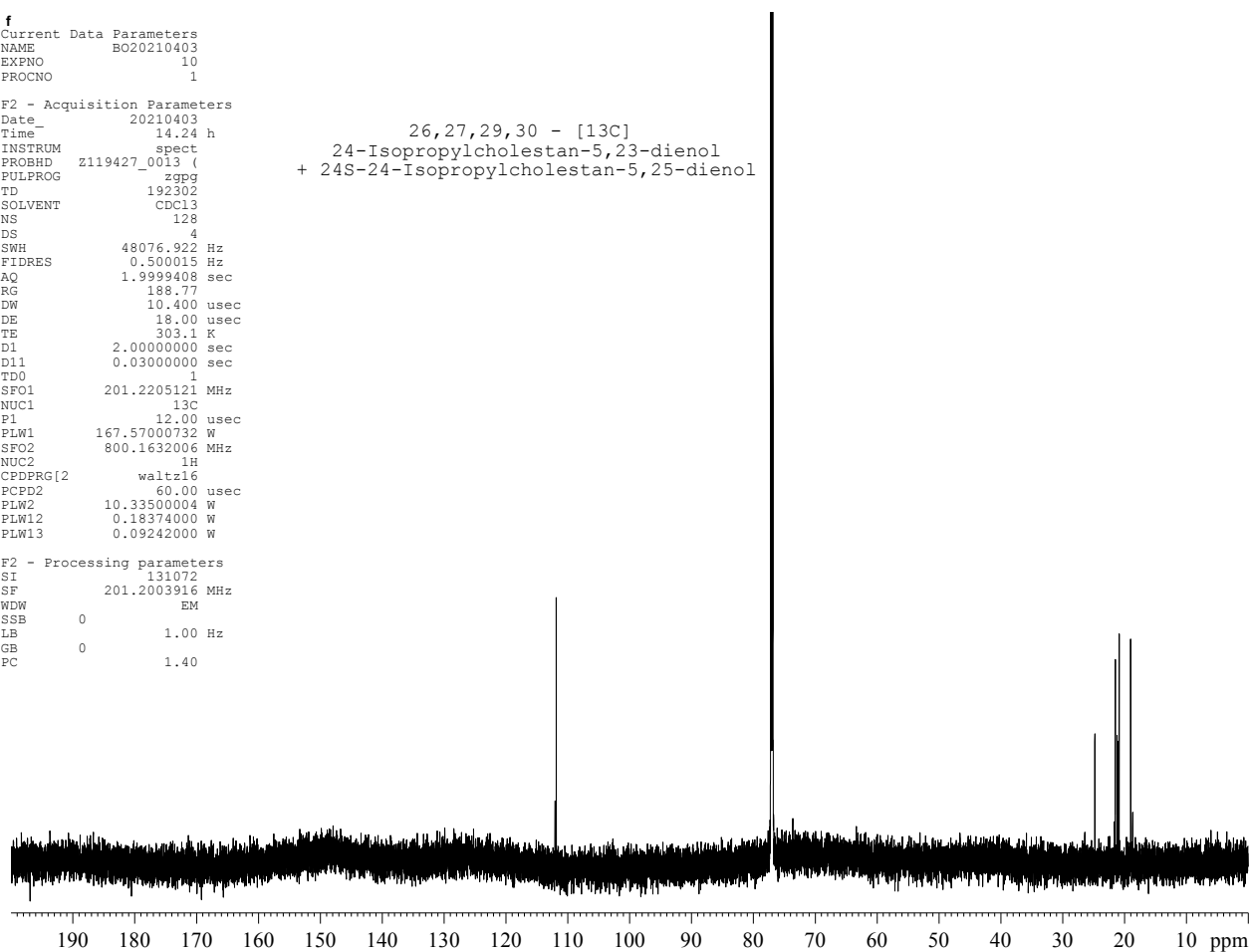


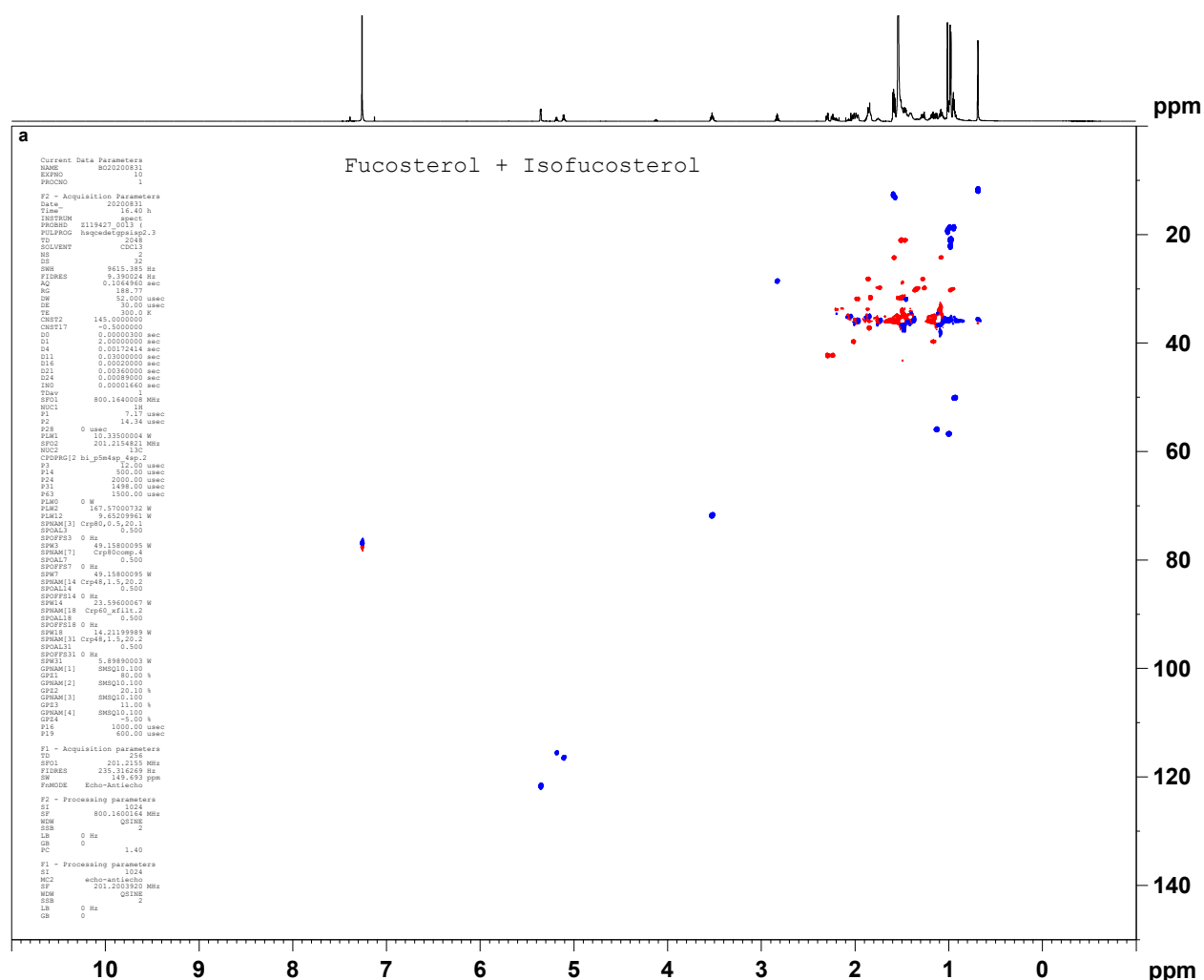
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SOLVENT CDCl3
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FIDRES 0.500015 Hz
AQ 1.9999408 sec
RG 188.77
DW 10.400 usec
DE 18.00 usec
TE 303.1 K
D1 2.00000000 sec
D11 0.03000000 sec
TDO 1
SFO1 201.2205121 MHz
NUC1 13C
P1 12.00 usec
PLW1 167.57000732 W
SFO2 800.1632006 MHz
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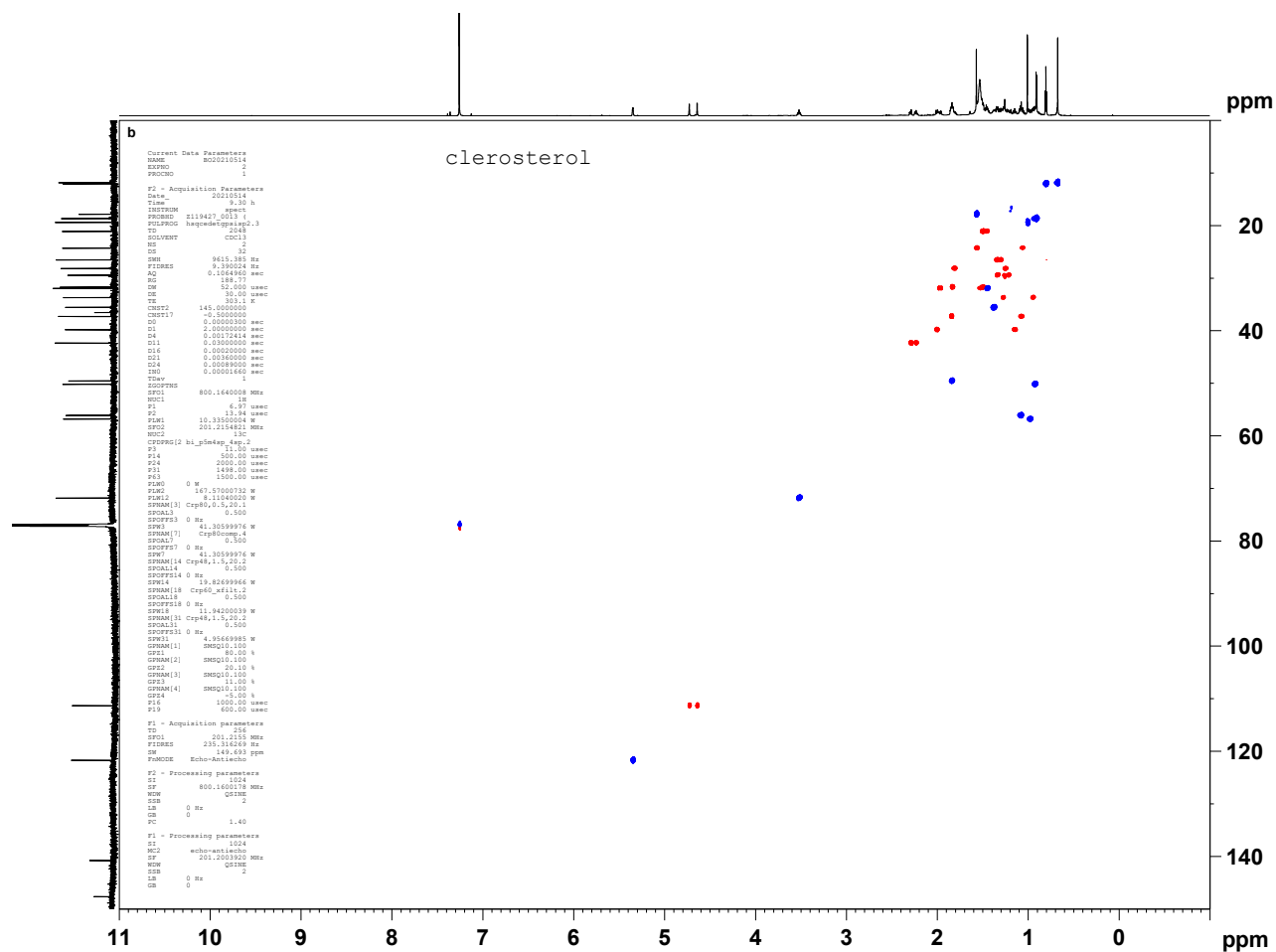
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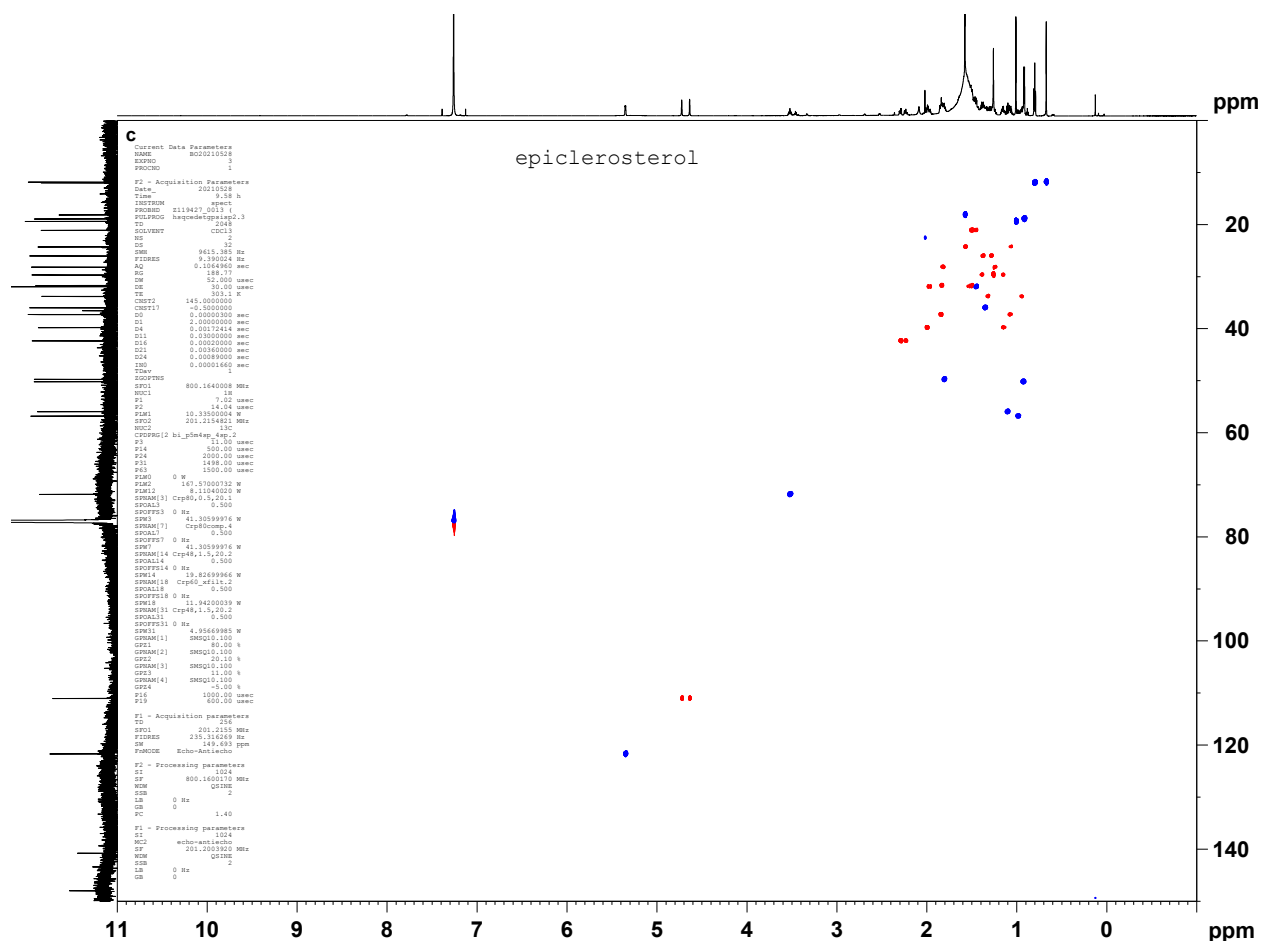
26,27,29,30 - [13C]
24-Isopropylcholestan-5,23-dienol
+ 24S-24-Isopropylcholestan-5,25-dienol

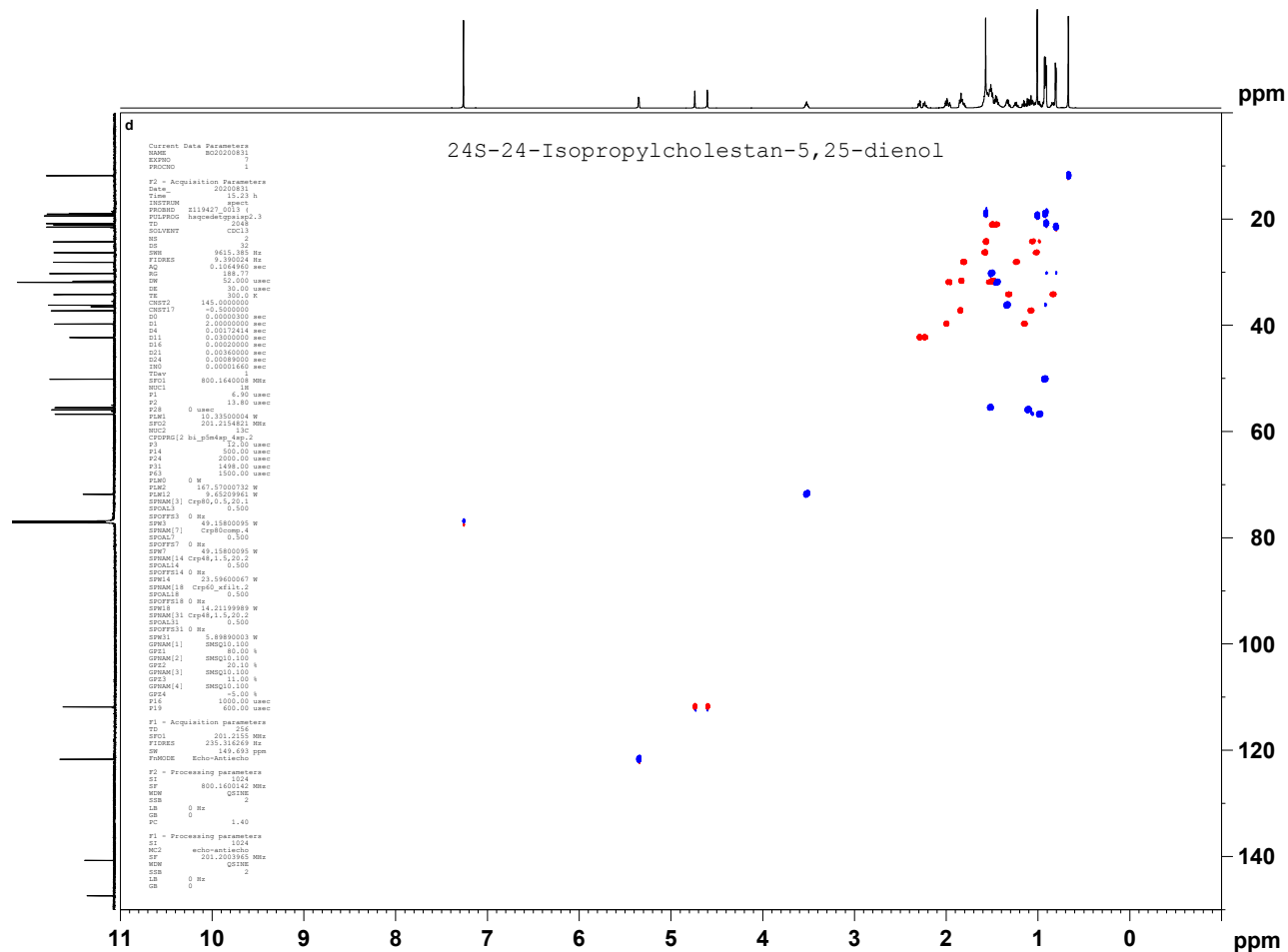


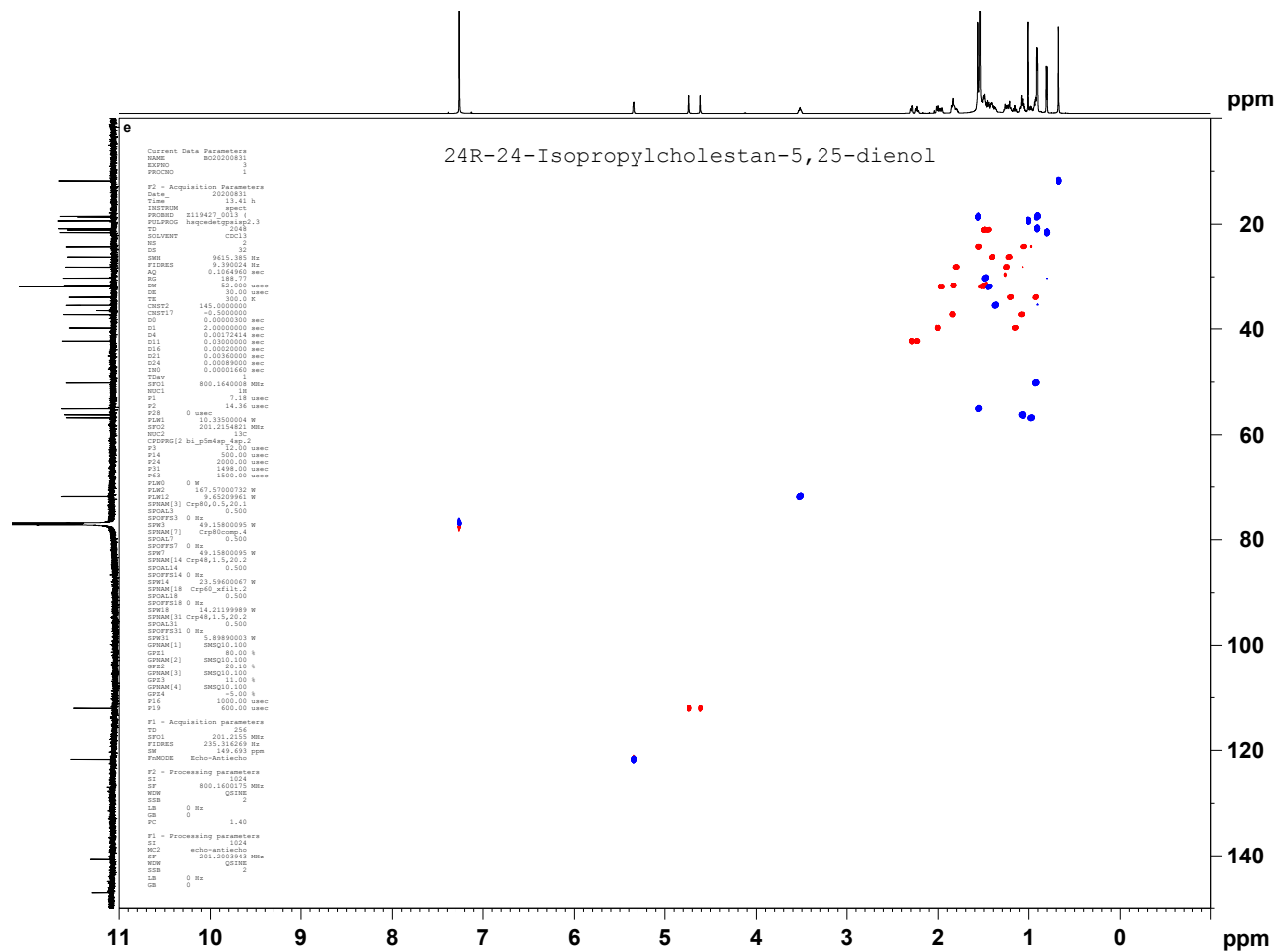


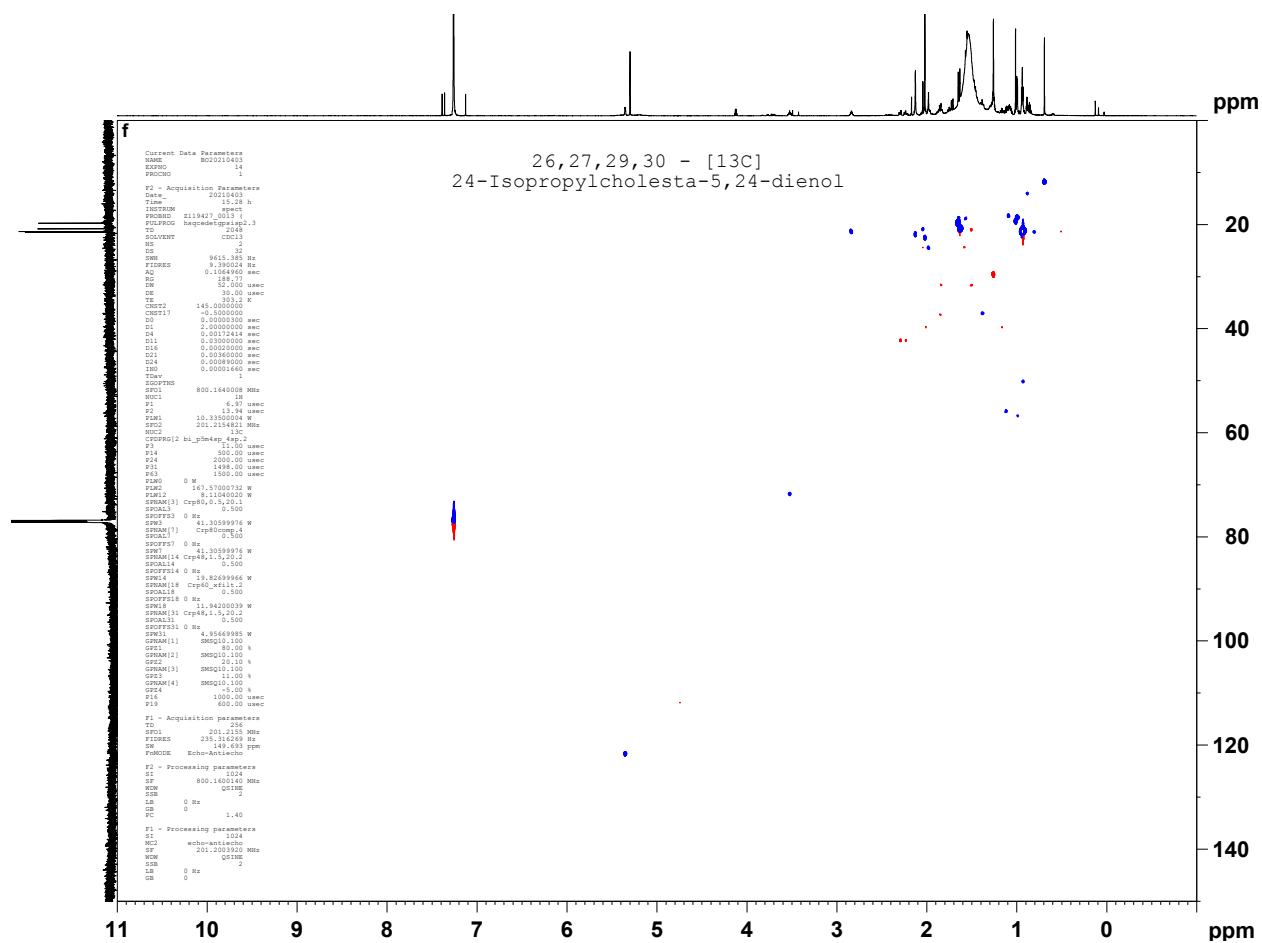
Supplementary Figure 6. Raw 2D NMR (HSQC) spectra of the following sterols: **a** fucosterol plus isofucosterol, **b** clerosterol, **c** epiclerosterol, **d** 24*S*-24-isopropylcholesta-5,25-dienol, **e** 24*R*-24-isopropylcholesta-5,25-dienol, **f** [26, 27, 29, 30-¹³C] 24-isopropylcholesta-5,24-dienol, and **g** [26, 27, 29, 30-¹³C] 24-isopropylcholesta-5,23-dienol plus 24*S*-24-isopropylcholesta-5,25-dienol. Chemical shifts are summarized in Supplementary Table 3.

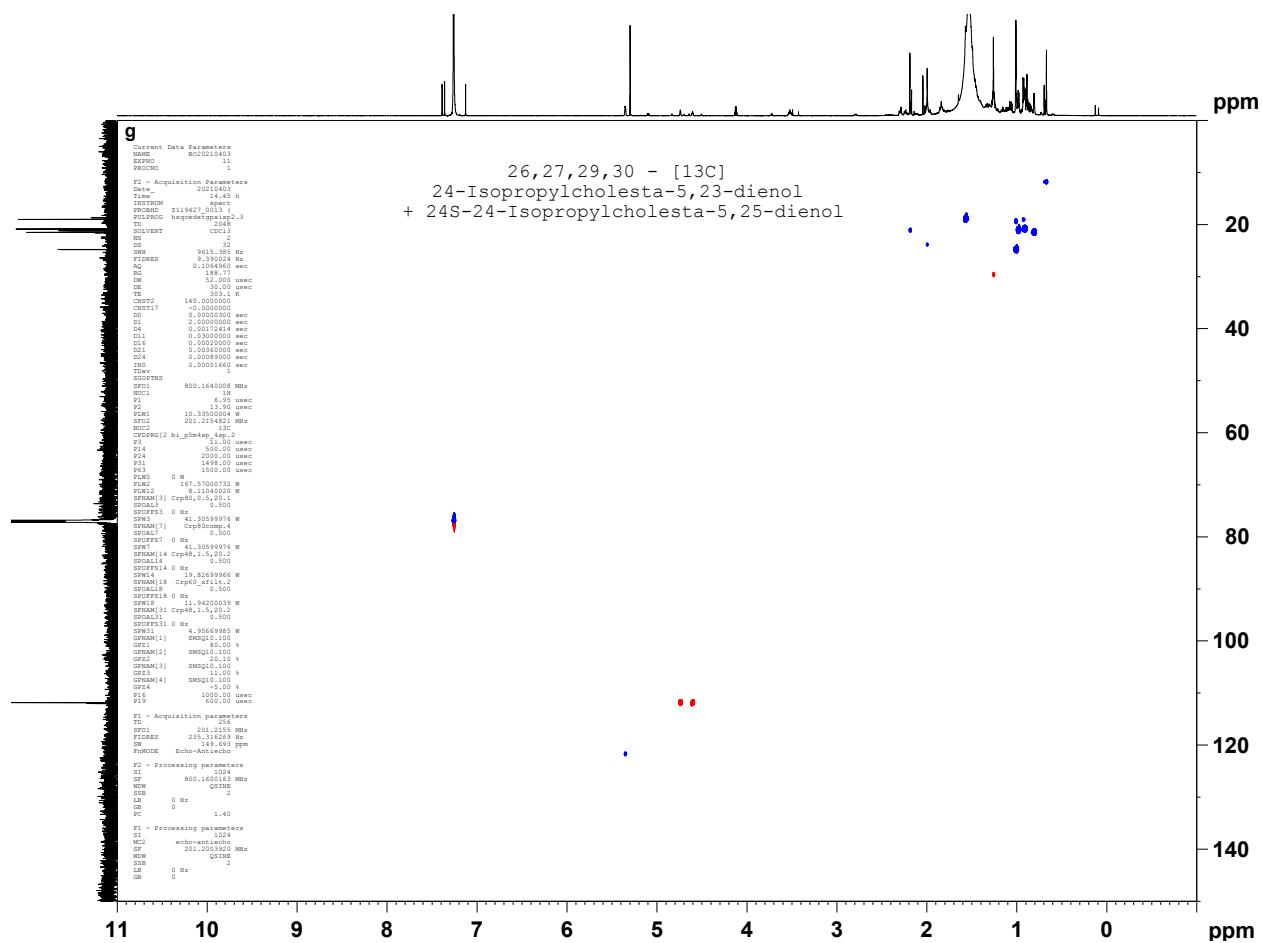


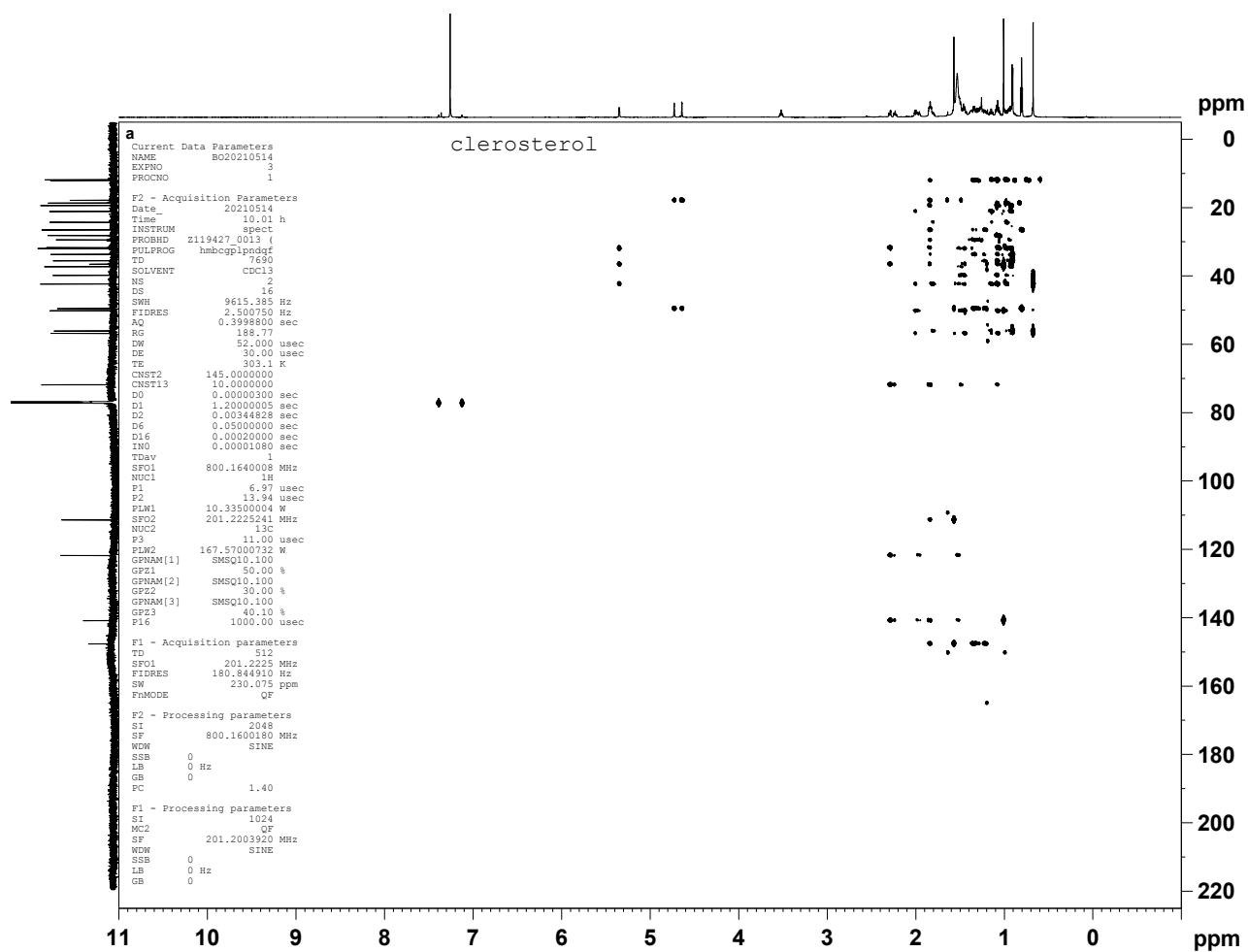




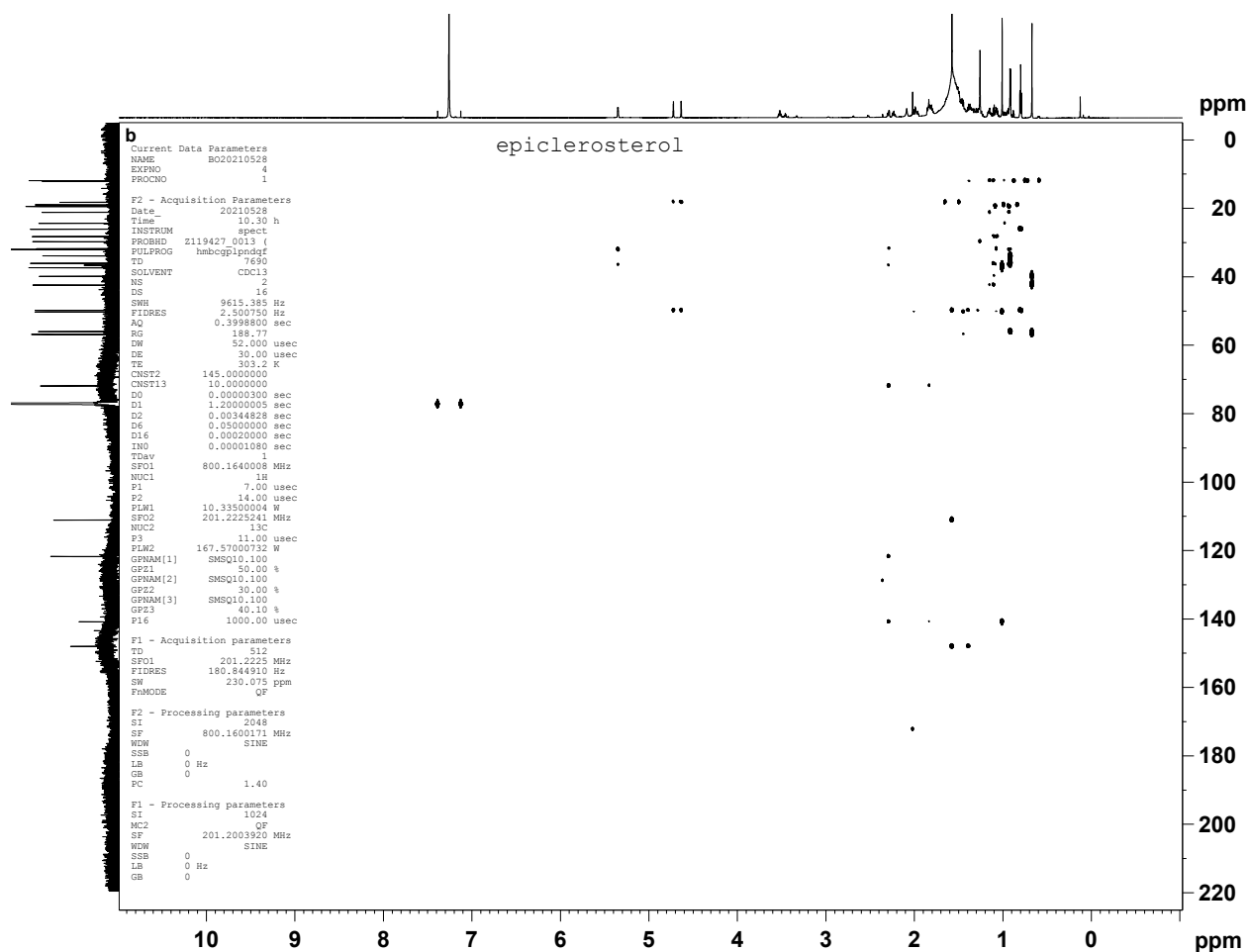


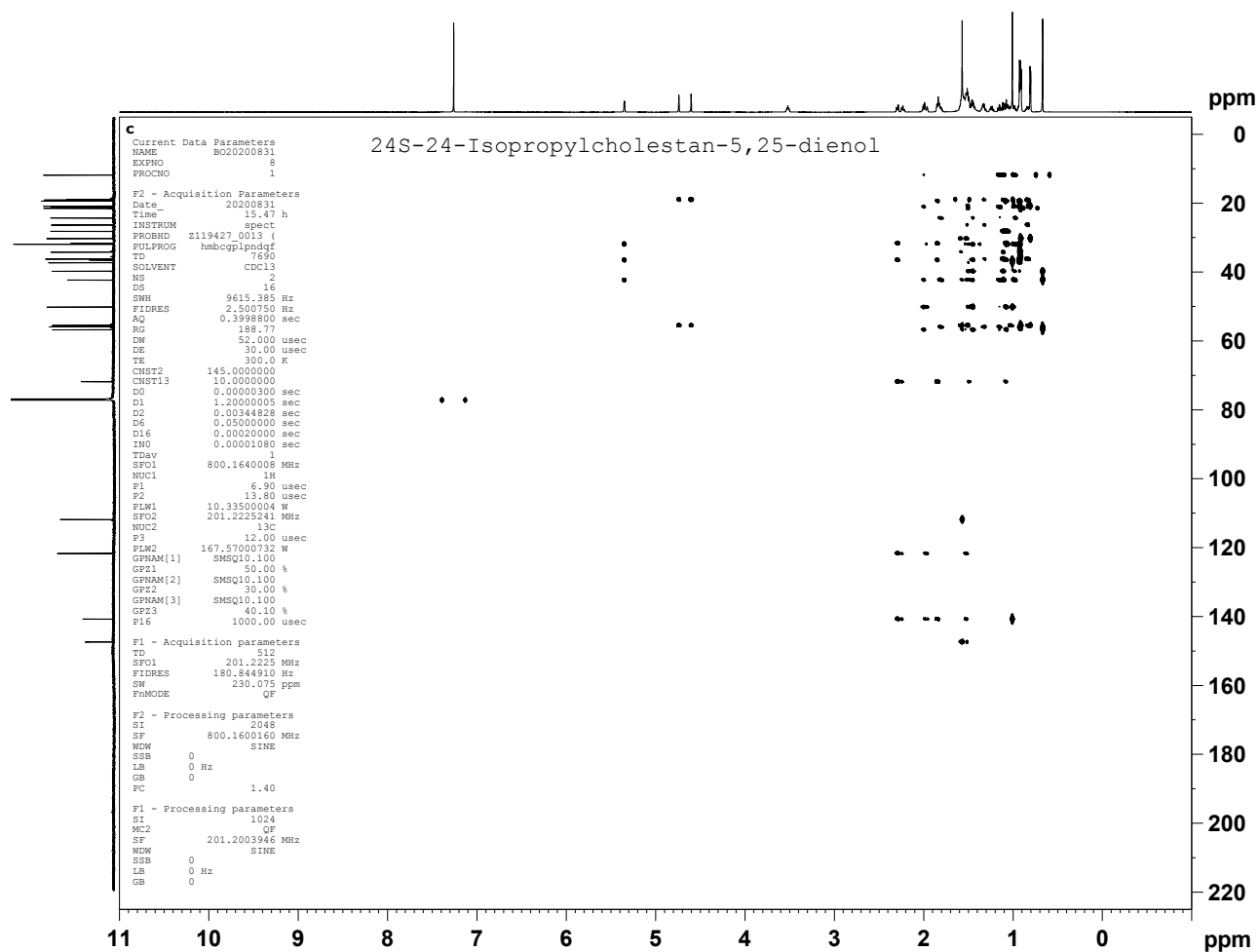


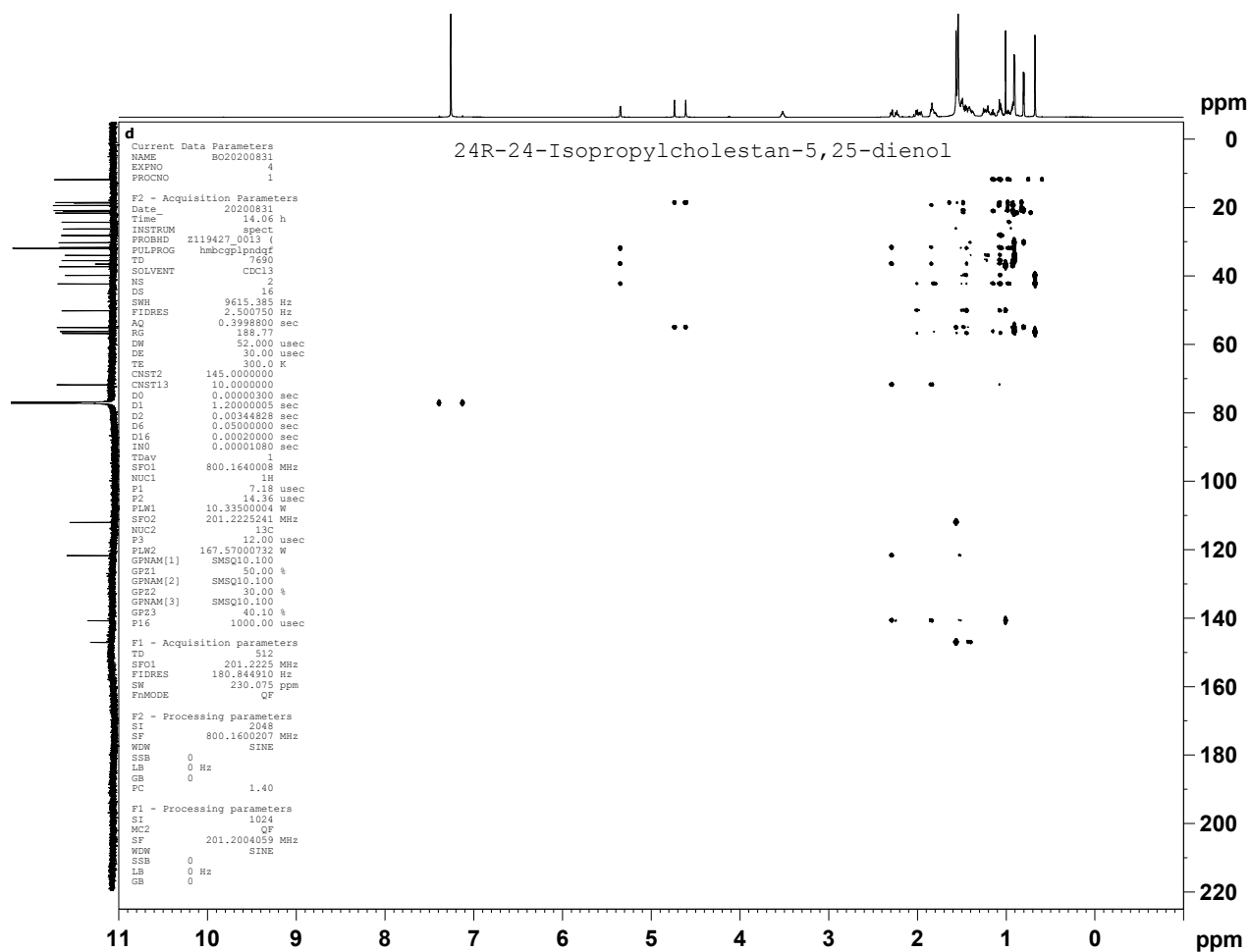


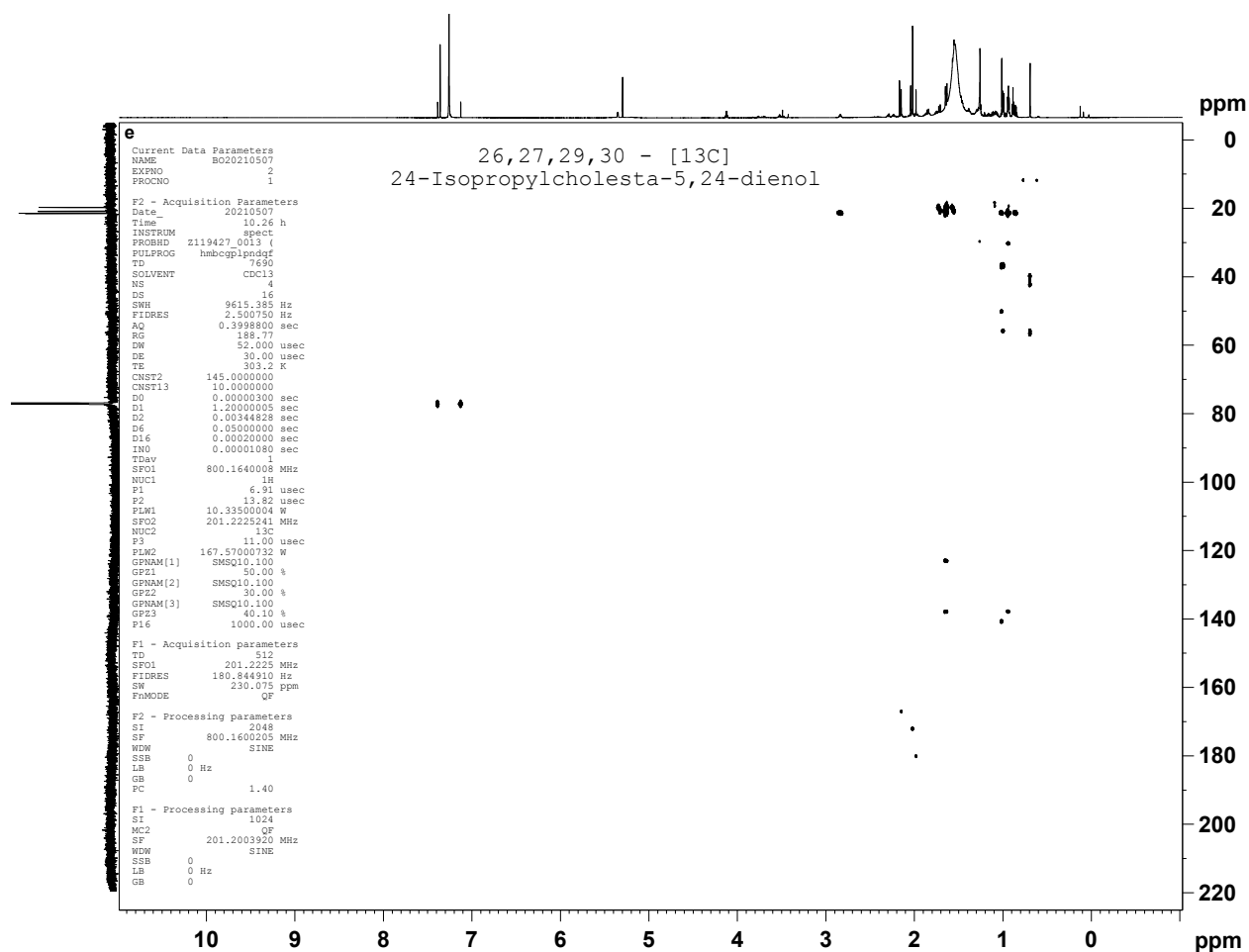


Supplementary Figure 7. **Raw 2D NMR (HMBC) spectra of the following sterols: a clerosterol, b epiclerosterol, c 24*S*-24-isopropylcholesta-5,25-dienol, d 24*R*-24-isopropylcholesta-5,25-dienol, and e [26, 27, 29, 30-¹³C] 24-isopropylcholesta-5,24-dienol.** Chemical shifts are summarized in Supplementary Table 3.









Supplementary Table 1. **Sterols previously identified in sponges relevant to this study.** The major sterol and the presence and/or percentages of C₂₇-C₃₀ sterols for each species are given when available.

Sponge species	Major sterol	C ₂₇	C ₂₈	C ₂₉	C ₃₀	Reference
DEMOSPONGIAE						
<i>Amphimedon queenslandica</i>	cholesterol	40	24	36		Gold et al., 2017 ¹
<i>Chondrilla nucula</i>	cholestanol	yes	yes	yes		Bergmann and McTigue, 1948 ²
<i>Sarcotragus fasciculatus</i>			yes	yes		Venkateswarlu et al., 1996 ³
<i>Haliclona amboinensis</i>	fucosterol	19	24	56		Fromont et al., 1994 ⁴
<i>Aplysina aerophoba</i>	aplysterol	13	9	77	1*	Nechev et al., 2002 ⁵ Zumberge et al., 2018 ⁶
<i>Petrosia ficiformis</i>	petrosterol	>2	>6	>60	yes	Khalil and Djerassi, 1980 ⁷
<i>Theonella swinhoei</i>	theonellasterol	yes	yes	yes		Kho 1980 ⁸
HOMOSCLEROMORPHA						
<i>Oscarella lobularis</i>		yes	yes	yes		Aiello et al., 1991 ⁹
CALCAREA						
<i>Sycon</i> sp. 1	cholesterol	33	28	34		Bergquist et al., 1986 ¹⁰
<i>Sycon</i> sp. 2	cholesterol	54	25	21		Bergquist et al., 1986 ¹⁰

*Nechev et al., 2002⁵ reports 1.2% 24-propyl-cholest-5,24(28)-dien-3 β -ol in *A. aerophoba*. Zumberge et al., 2018⁶ reports 0.5% 24-npc and 0.4% 24-ipc after catalytic hydropyrolysis of *A. aerophoba* biomass. 24-isopropyl sterols have not been reported as components of *A. aerophoba*'s native sterols, to our knowledge.

Supplementary Table 2. Whokaryote predicts all metagenomic contigs containing sterol methyltransferases analyzed in this study as prokaryotic.

		contig	nr	ratio same	ID general	ID general	gene	gene			rbs	tiara		
SMT		contig	length	genes	orientation	avg	std	density	length	ID Q1	ID Q3	ratio	pred	predicted
<i>Aplysina aerophoba</i>	JGIcombinedJ30088_10000835	JGIcombinedJ30088_10000835	32339	23	0.68	128.14	183.70	0.91	1285.09	3.75	181.00	0.96	1.0	prokaryote
<i>Aplysina aerophoba</i>	Ga0209021_1000146	Ga0209021_1000146	97163	103	0.68	115.34	117.55	0.88	829.83	31.00	173.75	0.90	1.0	prokaryote
<i>Petrosia ficiformis</i>	LXNJ01014139	LXNJ01014139.1	3313	4	0.00	100.67	62.53	0.91	752.00	56.50	124.50	0.25	1.0	prokaryote
<i>Petrosia ficiformis</i>	LXNJ01003202	LXNJ01003202.1	8597	8	0.57	115.57	167.57	0.91	975.50	0.00	131.00	0.88	1.0	prokaryote
<i>Petrosia ficiformis</i>	LXNJ01000389	LXNJ01000389.1	31336	22	0.67	79.90	78.14	0.94	1341.91	4.00	147.00	0.77	1.0	prokaryote
<i>Petrosia ficiformis</i>	LXNJ01004423	LXNJ01004423.1	7072	6	0.60	102.20	58.00	0.93	1093.00	63.00	126.00	0.67	1.0	prokaryote
<i>Sarcotragus foetidus</i>		LXNIO1004480.1	7225	5	0.50	105.25	156.88	0.94	1359.20	6.75	121.00	0.80	1.0	prokaryote
<i>Theonella swinhoei</i> associated Chromatiales sp. 1 of 2		2690316284	444506	427	0.74	74.66	122.32	0.93	968.13	1.00	104.25	0.68	1.0	prokaryote
<i>Theonella swinhoei</i> associated Chromatiales sp. 2 of 2		2690316289	117771	97	0.83	74.25	116.07	0.94	1142.96	0.00	97.50	0.65	1.0	prokaryote
Marine sediment, Gulf of Thailand		Ga0118733_100012602	21160	19	0.89	72.06	133.71	0.94	1042.53	0.00	65.50	0.84	1.0	prokaryote
Marine sediment, Helgoland, North Sea		Ga0055584_100171735	2192	3	0.50	80.50	75.50	0.93	676.00	42.75	118.25	0.00	1.0	prokaryote
Western Arctic Ocean		Ga0133547_10420286	2727	3	0.00	147.00	126.00	0.89	810.00	84.00	210.00	0.33	1.0	prokaryote
Freshwater, Lake Fryxell, Antarctica		Ga0105048_10034691	7428	9	0.63	86.00	80.31	0.88	722.33	13.50	104.75	0.44	1.0	prokaryote
Lake sediment, Walker Lake, Nevada		Ga0114946_10008497	6500	5	0.50	77.25	71.98	0.93	1212.80	22.50	103.75	0.40	1.0	prokaryote
Hotspring, Beatty, Nevada		Ga0114945_10021081	3480	3	0.00	63.00	7.00	0.96	1108.00	59.50	66.50	0.33	1.0	prokaryote
Hotspring sediment, Dewar Creek, BC		Ga0073932_1004097	18020	12	0.82	164.73	132.22	0.89	1342.75	67.00	269.00	0.92	1.0	prokaryote
Chlamydiae sp., drinking water 1 and 2		2619626156	1146800	1105	0.68	85.18	128.28	0.92	956.10	0.00	118.00	0.42	1.0	prokaryote
<i>Nitrospira</i> sp., coral reef 1 and 2		2681831735	70077	63	0.74	147.08	145.66	0.87	962.71	27.00	196.00	0.89	1.0	prokaryote
<i>Sandaracinus</i> sp., marine 1 and 2		2775515747	24518	17	0.88	112.25	106.29	0.89	1282.47	22.50	163.50	0.76	1.0	prokaryote
<i>Spirochaetes</i> sp., soil		2708772188	70818	55	0.54	151.69	167.86	0.89	1142.60	6.25	242.50	0.24	1.0	prokaryote

Supplementary Table 3. **800 MHz ¹H and 201 MHz ¹³C NMR assignments of selected sterol side-chain positions at 30 °C.** ¹³C data are in boldface. The assignments were determined by HSQC-DEPT, HMBC, and COSY 2D-NMR experiments. ¹H and ¹³C chemical shifts are given to three and two decimal places, respectively, if directly observed, two and one, respectively, if determined from 2D spectra. Diastereotopic signals are not assigned. The raw spectra used to generate this table are provided in Supplementary Fig. 4-7.

	C-24	C-25	C-26	C-27	C-28	C-29	C-30
Fucosterol	145.6	2.199 34.7			5.183 115.58	1.574 13.16	
Isofucosterol	147.1	2.829 28.6			5.108 116.48	1.591 12.76	
Clerosterol*	1.838 49.54	147.61	4.730, 4.642 111.33	1.570 17.85	1.305, 1.341 26.53	0.805 12.04	
Epiclerosterol	1.81 49.75	147.98	4.725, 4.636 111.06	1.576 18.13	1.290, 1.380 26.01	0.800 11.96	
24 <i>R</i> -24-Isopropylcholest-5,25-dienol*	1.56 55.03	147.29	4.613, 4.738 112.02	1.567 18.63	1.49 30.25	0.911 20.85	0.803 21.60
24 <i>S</i> -24-Isopropylcholest-5,25-dienol	1.52 55.50	147.38	4.602 4.740 111.85	1.570 18.95	1.51 30.24	0.913 20.85	0.807 21.49
24-Isopropylcholest-5,23-dienol			1.013 24.77	1.013 24.79		0.980 20.96	0.980 21.18
24-Isopropylcholest-5,24-dienol	137.0	123.1	1.633 20.81	1.651 19.73		0.938 21.34	0.938 21.41

*Clerosterol and 24*R*-24-isopropylcholest-5,25-dienol were not detected as products of the *A. aerophoba* symbiont and Chlamydiae SMTs but are shown for comparative purposes.

Supplementary Table 4. Sources of sterol methyltransferases analyzed in this study.

SMT	Source	Genome ID/ BioProject	Gene ID/Locus
DEMOSPONGIAE			
<i>Amphimedon queenslandica</i>	JGI IMG	2507525020	2508051218
<i>Chondrilla nucula</i>	(¹¹)		
<i>Sarcotragus fasciculatus</i>	(¹¹)		
<i>Haliclona amboinensis</i>	Compagen		
<i>Haliclona tubifera</i>	Compagen		
HOMOSCLEROMORPHA			
<i>Oscarella pearsei</i>	Compagen		
<i>Oscarella carmela</i>	Compagen		
CALCAREA			
<i>Sycon ciliatum</i>	Compagen		
DEMOSPONGE METAGENOME			
<i>Aplysina aerophoba</i> JGIcombinedJ30088_10000835 1 of 2	JGI IMG	3300002448	JGIcombinedJ30088_1000083511
<i>Aplysina aerophoba</i> JGIcombinedJ30088_10000835 2 of 2	JGI IMG	3300002448	JGIcombinedJ30088_100008359
<i>Aplysina aerophoba</i> Ga0209021_1000146	JGI IMG	3300027386	Ga0209021_100014628
<i>Petrosia ficiformis</i> LXNJ01014139	GenBank	PRJNA318959	LXNJ01014139.1
<i>Petrosia ficiformis</i> LXNJ01003202	GenBank	PRJNA318959	LXNJ01003202.1
<i>Petrosia ficiformis</i> LXNJ01000389	GenBank	PRJNA318959	LXNJ01000389.1
<i>Petrosia ficiformis</i> LXNJ01004423	GenBank	PRJNA318959	LXNJ01004423.1
<i>Sarcotragus foetidus</i>	GenBank	PRJNA318959	LXNI01004480.1
DEMOSPONGE METAGENOME-ASSEMBLED GENOME			
<i>Theonella swinhoei</i> associated Chromatiales sp. 1 of 2	JGI IMG	2690315631	2690455922
<i>Theonella swinhoei</i> associated Chromatiales sp. 2 of 2	JGI IMG	2690315631	2690456335
OTHER METAGENOME			
Marine sediment, Gulf of Thailand	JGI IMG	3300010430	Ga0118733_1000126028
Marine sediment, Helgoland, North Sea	JGI IMG	3300004097	Ga0055584_1001717352
Western Arctic Ocean	JGI IMG	3300010883	Ga0133547_104202862
Freshwater, Lake Fryxell, Antarctica	JGI IMG	3300009032	Ga0105048_100346915
Lake sediment, Walker Lake, Nevada	JGI IMG	3300009504	Ga0114946_100084975
Hot spring, Beatty, Nevada	JGI IMG	3300009444	Ga0114945_100210811
Hot spring sediment, Dewer Creek, BC	JGI IMG	3300007072	Ga0073932_10040971
OTHER METAGENOME-ASSEMBLED GENOME			
<i>Chlamydiae</i> sp., drinking water 1 of 2	JGI IMG	2619618861	2619789262
<i>Chlamydiae</i> sp., drinking water 2 of 2	JGI IMG	2619618861	2619789261
<i>Nitrospira</i> sp., coral reef 1 of 2	JGI IMG	2681812944	2682229274
<i>Nitrospira</i> sp., coral reef 2 of 2	JGI IMG	2681812944	2682229270
<i>Sandaracinus</i> sp., marine 1 of 2	JGI IMG	2775506781	2775843484
<i>Sandaracinus</i> sp., marine 2 of 2	JGI IMG	2775506781	2775843485
<i>Spirochaetes</i> sp., soil	JGI IMG	2708742746	2709604172

Supplementary Table 5. Best amino acid substitution models, gamma shape parameters, and invariable sites proportions under the Bayesian Information Criterion and Akaike Information Criterion using ModelTest-NG on XSEDE for generation of sterol methyltransferase and oxidosqualene cyclase maximum-likelihood trees.

Protein	Substitution Model	Gamma Shape Parameter	Invariable Sites Proportion
sterol methyltransferase	LG+I+G	1.260	0.020
oxidosqualene cyclase	LG+I+G	1.212	0.022

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