

Supplementary Material

CHARMM All-Atom Additive Force Field for Sphingomyelin: Elucidation of Hydrogen Bonding and of Positive Curvature

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Table S1. CHelpG charges at the MP2/cc-pVDZ model chemistry for conformations C3 through C6 of the CER6 model for SM. Atom names correspond to Figure 2. Also shown are the charges used in unoptimized CHARMM36 (C36u) and charge optimized C36.

	C3	C4	C5	C6	C36u	C36
C3	+0.601	+0.490	+0.509	+0.421	+0.05	+0.50
H3	-0.090	+0.001	-0.087	-0.035	+0.09	-0.11
O3	-0.744	-0.655	-0.693	-0.646	-0.45	-0.69
HO	+0.429	+0.348	+0.424	+0.348	+0.31	+0.30
C2	+0.220	+0.140	+0.157	+0.284	+0.07	+0.30
H2	-0.002	+0.044	+0.013	+0.048	+0.09	+0.05
N	-0.749	-0.801	-0.776	-0.754	-0.47	-0.70
H	+0.334	+0.443	+0.351	+0.351	+0.31	+0.35
CF	+0.675	+0.743	+0.738	+0.702	+0.51	+0.55
OF	-0.515	-0.592	-0.556	-0.540	-0.51	-0.60

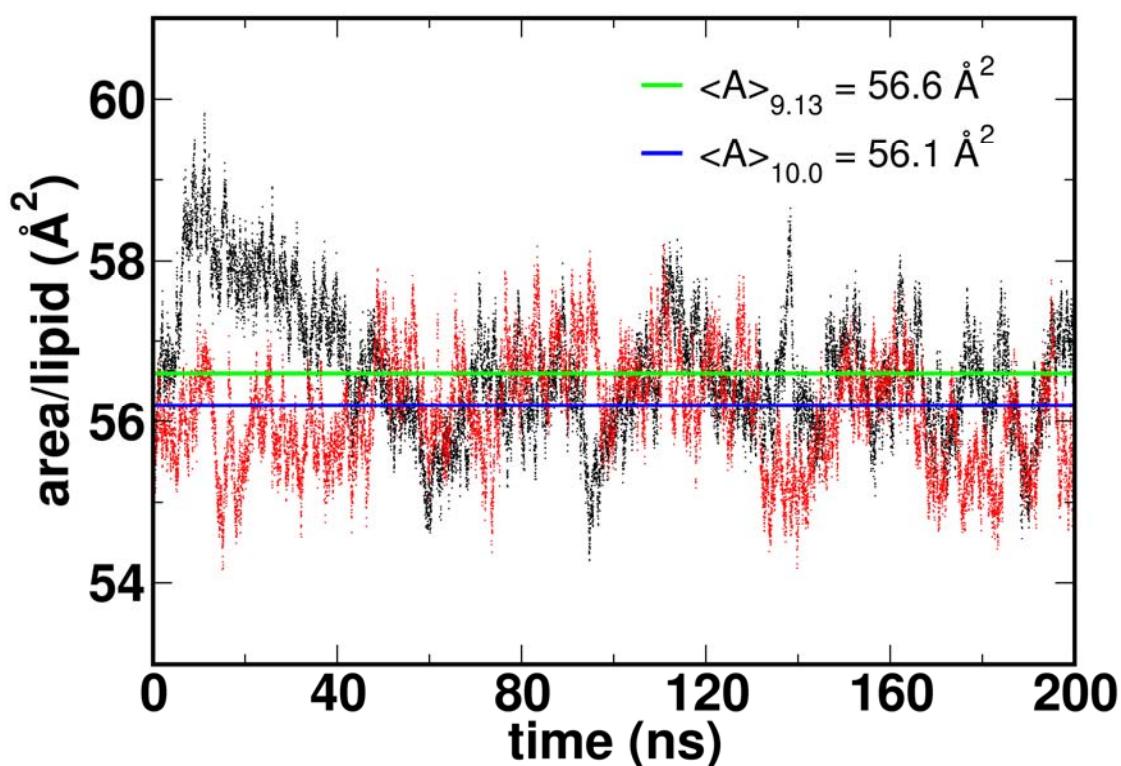


Figure S1. Effect of Anton short range nonbonded interaction cutoff on the area/lipid, A; the blue line is the average for the red time series ($R_{\text{cut}} = 10.0 \text{ \AA}$), and green is the average for the black time series ($R_{\text{cut}} = 9.13 \text{ \AA}$).

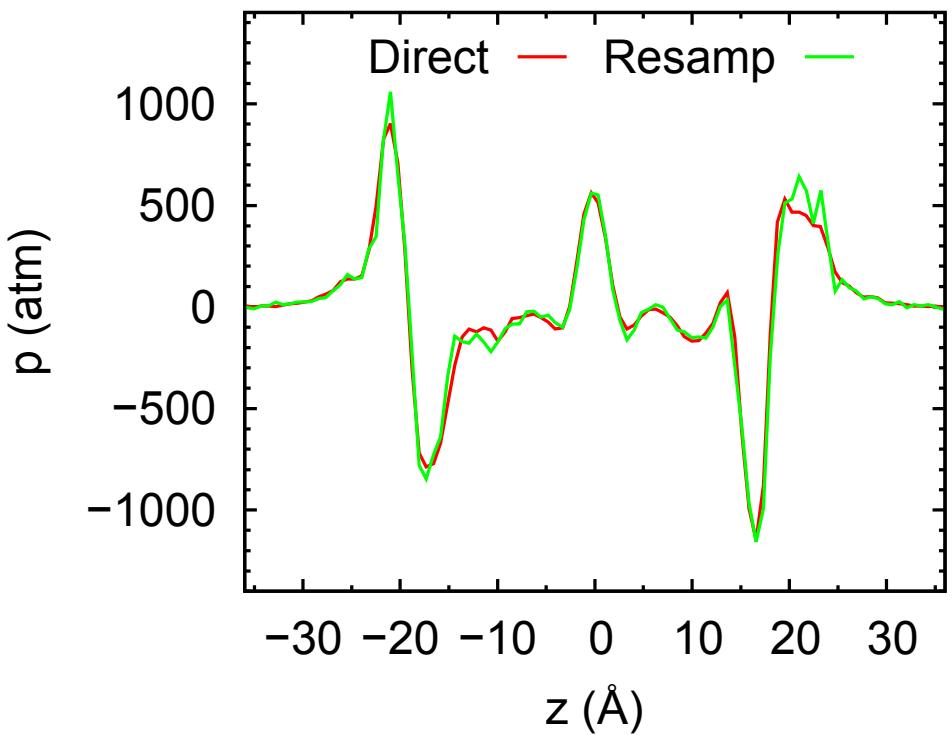


Figure S2. Comparison of pressure profiles from a 100 ns PSM simulation with full sampling (Direct) or via the 10% resampling (Resamp) of 100 ps intervals spaced 1 ns apart, using restart files from the fully sampled simulation. The $\bar{F}'(0)$ values and standard errors obtained are -0.190 ± 0.009 and -0.192 ± 0.004 for the fully sampled and resampled data, respectively.

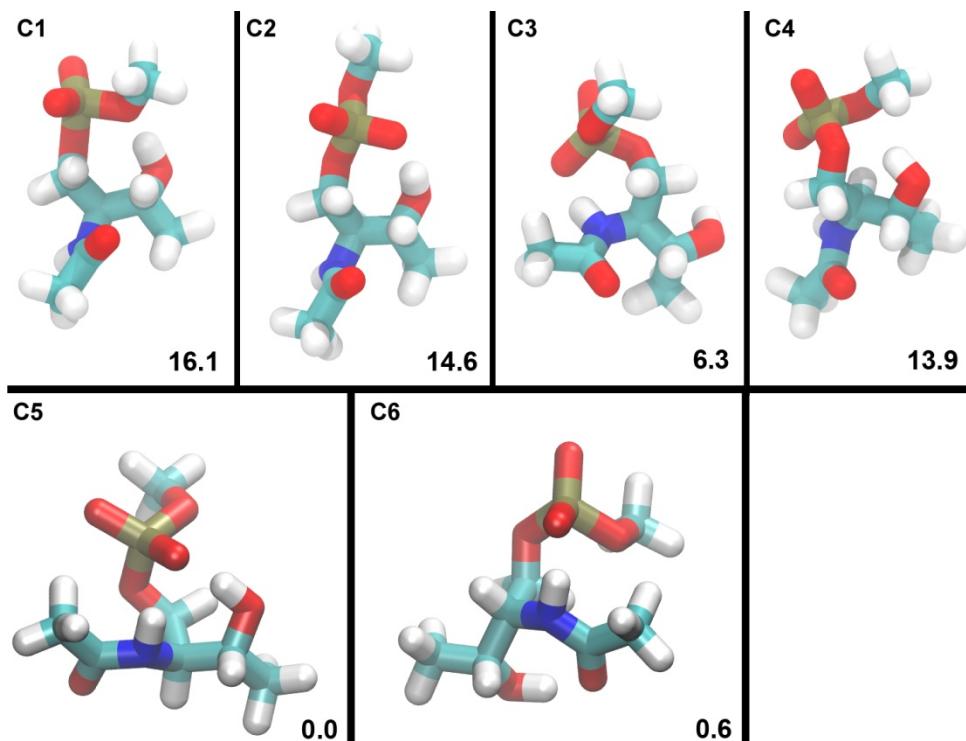


Figure S3. Conformations of local minima calculated at the MP2/cc-pVDZ model chemistry for the CER6 molecule. The energies relative to the global minima are shown in the bottom right corner in kcal/mol.

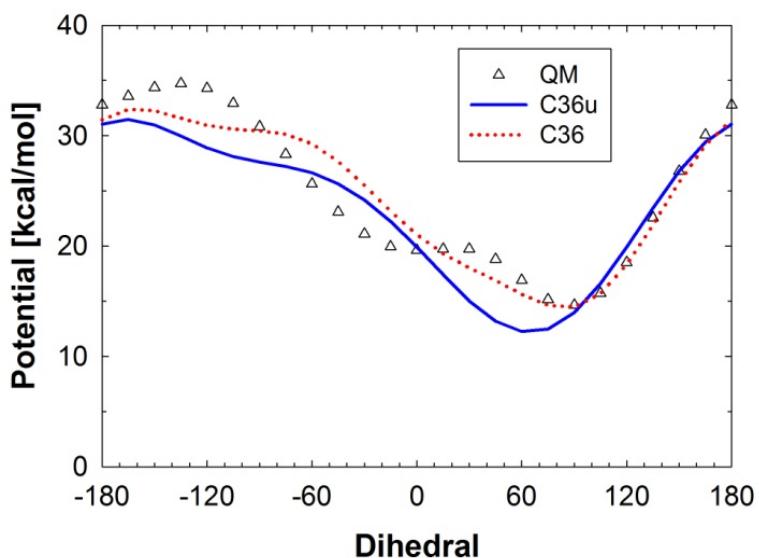


Figure S4. The torsional energy of the tOH (H-O3-C3-C2) torsion of CER6 from QM (MP2/cc-pVDZ) and C36u and C36 relative to the minima of t4. Constraints were included to maintain the C2 conformation in Figure S3.

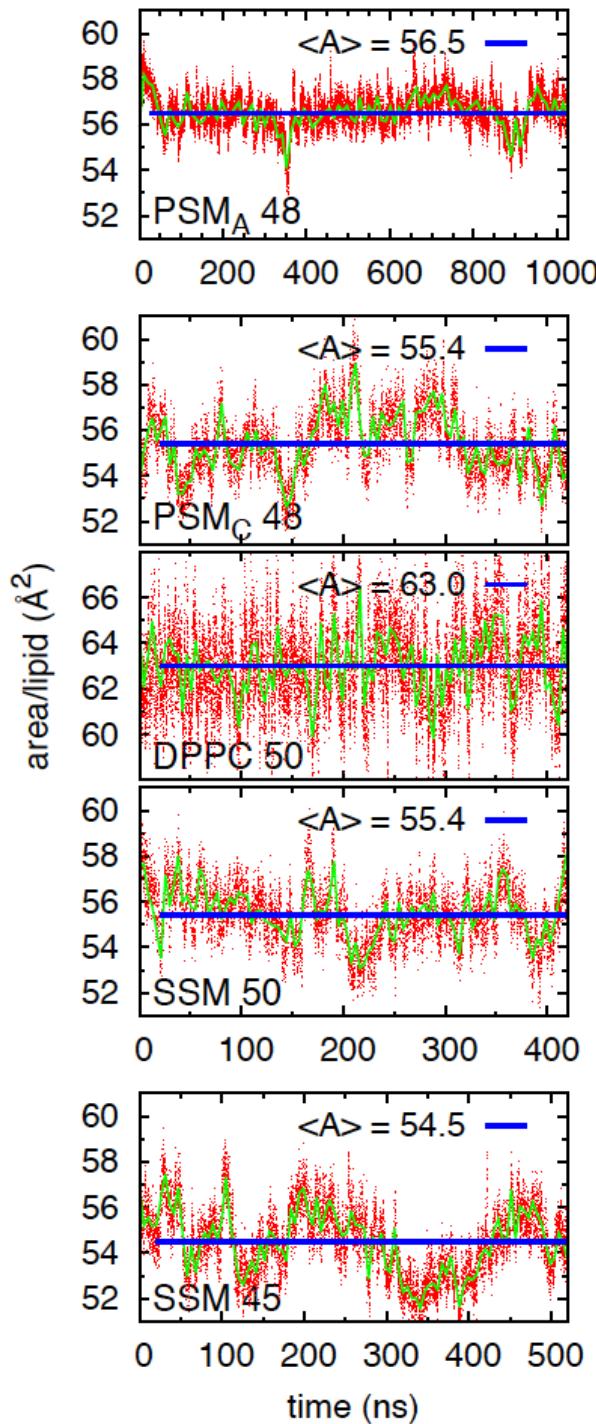


Figure S5. Area/lipid time series from CHARMM and Anton (PSM_A) simulations; red dots are snapshot values, the green line is a smoothed local average, and the heavy blue line is the average over the interval indicated by the line endpoints.

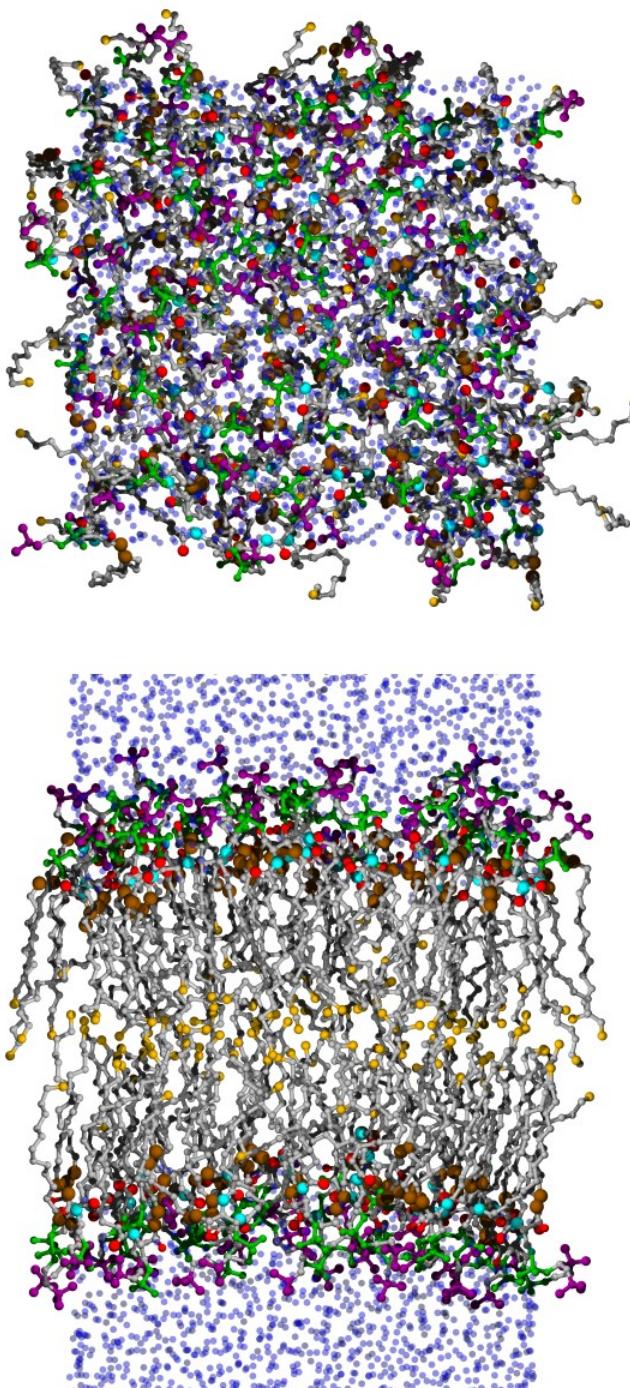


Figure S6. Molecular image snapshot at 200 ns from the PSM simulation run with CHARMM, top and side views. Colors: water is transparent blue, the choline group is purple, the phosphate group is green, amide N is cyan, amide O is orange, hydroxyl O is red, vinyl C=C atoms are brown, chain ends are yellow; all other C atoms are gray.

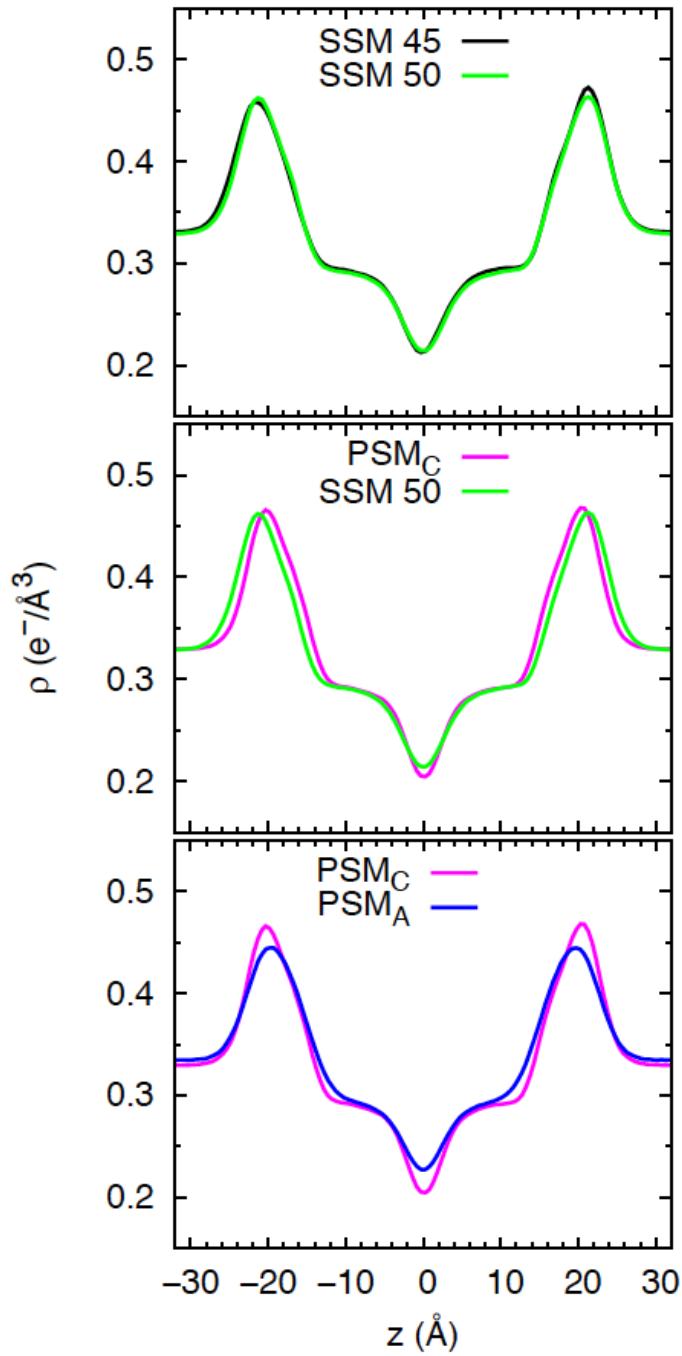


Figure S7. Electron density z profiles for all atoms. Top panel: comparison of SSM at 45 and 50°C; they are nearly identical. Middle panel: CHARMM simulations of SSM at 50°C and PSM at 48°C; the expected smaller $d_{\text{P-P}}$ is observed for PSM. Bottom panel: PSM simulations from CHARMM (PSM_C) and Anton (PSM_A); the difference is most likely due to the different VDW methods and cutoffs.

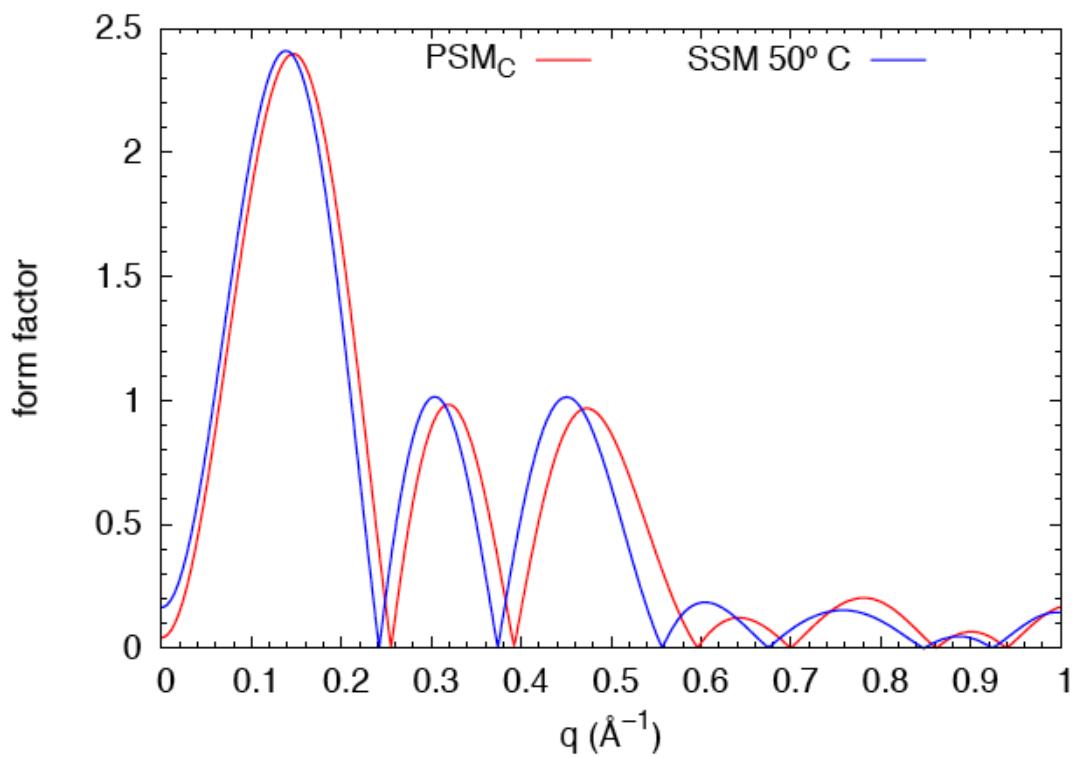


Figure S8. Calculated x-ray scattering form factors from the MD simulations.

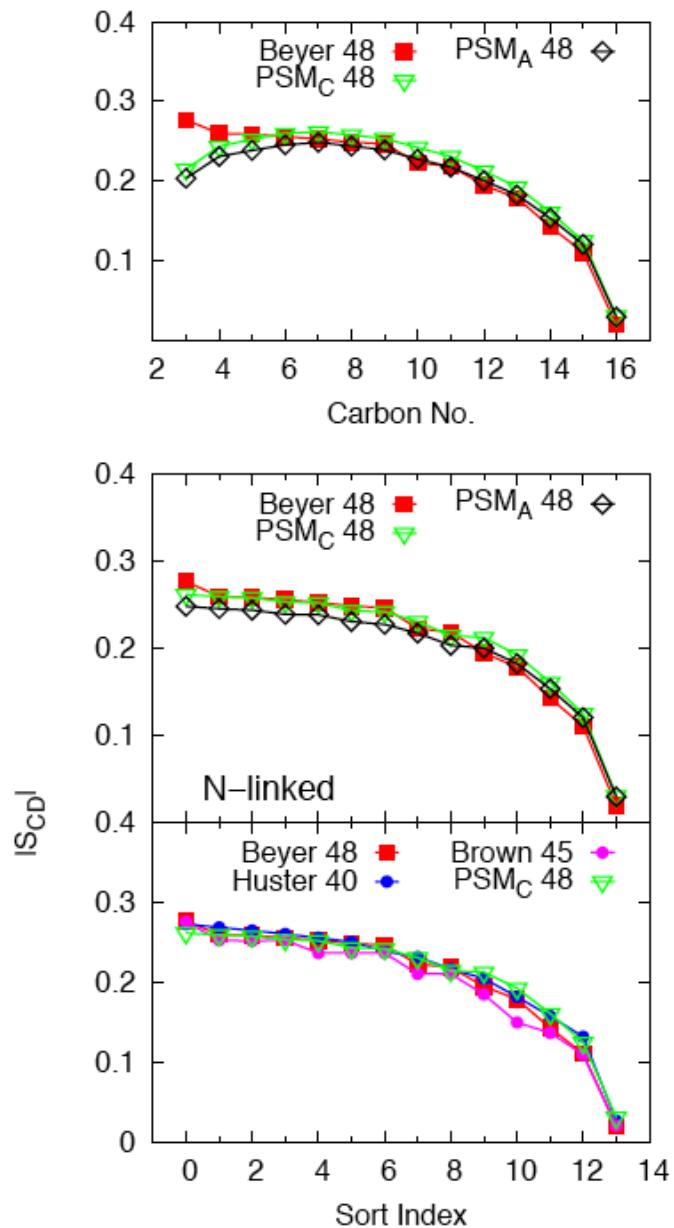


Figure S9. Order parameter profiles based on chain position (top panel) and value-sorted order (lower 2 panels). Labels: PSM_C = PSM simulated with CHARMM; PSM_A = PSM simulated with Anton; Beyer (7), Huster (8), and Brown (9) are the last authors of the respective PSM-d31 NMR papers; the temperature from experiment or simulation is listed in each label. The top panel shows an apparent discrepancy, which is resolved by value sorting the simulation values, as shown in the middle panel. The lower panel compares the sorted S_{CD} profile from the CHARMM simulation to the profiles from all three NMR papers.

PSM lipids with N-H:::O inter-lipid H-bonds

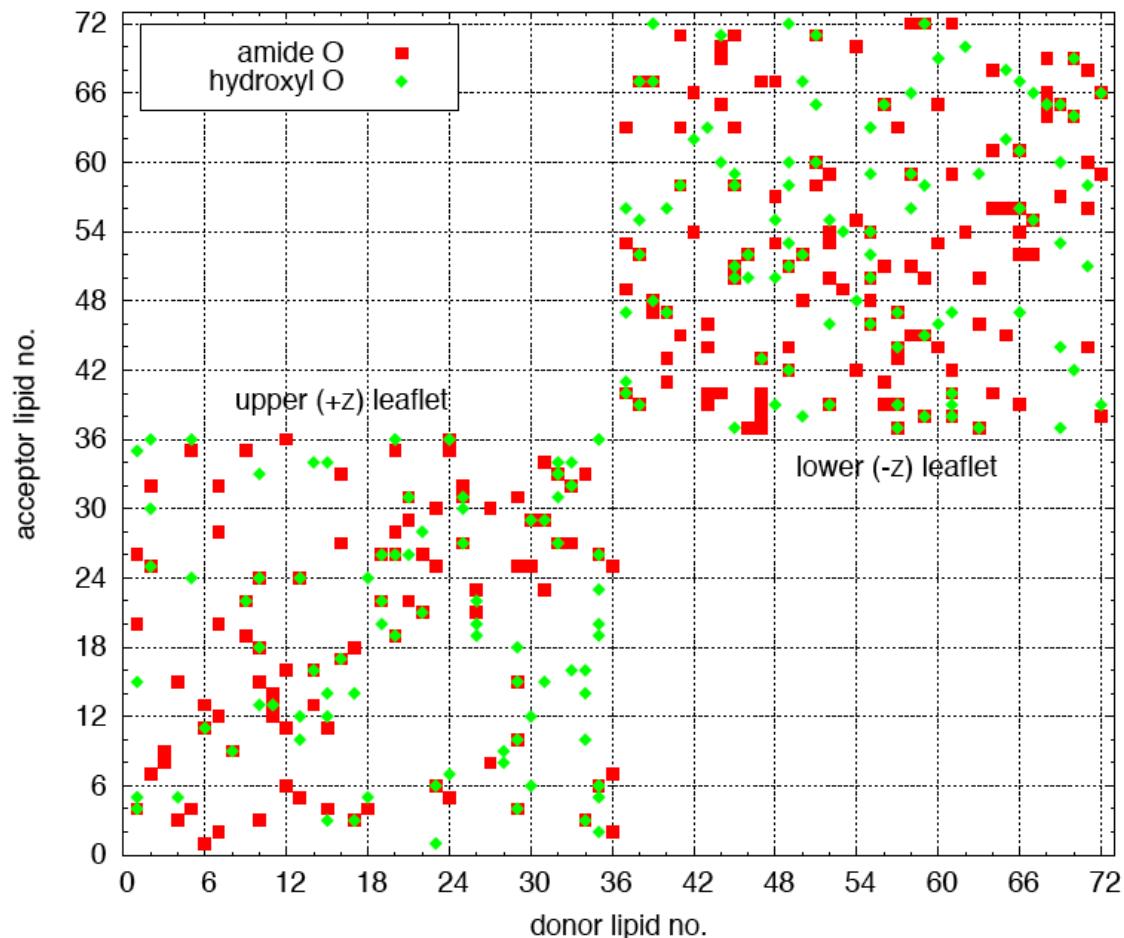


Figure S10. Observed inter-lipid H-bonds with an occupancy of 0.1% or above.

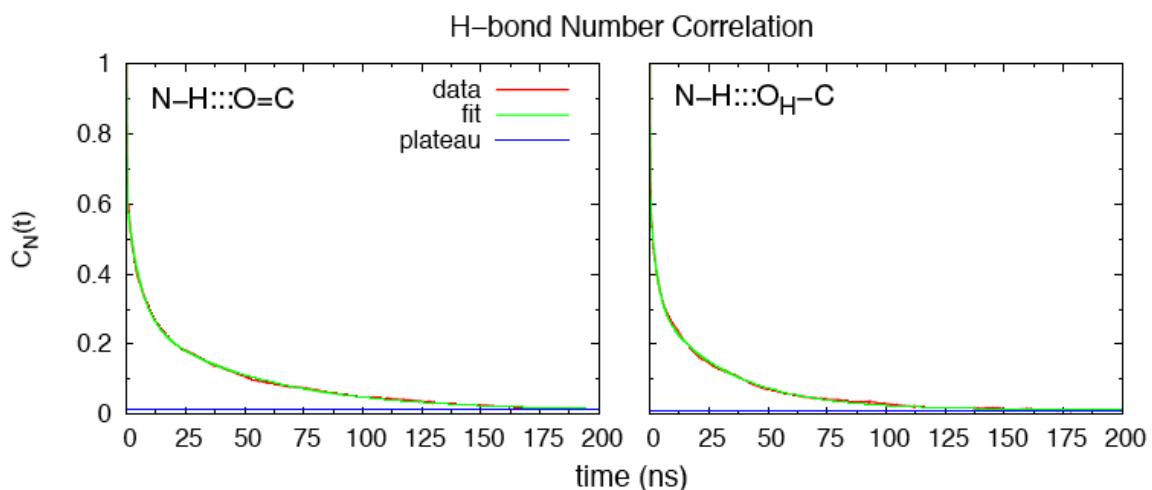


Figure S11. Hydrogen bond number correlation functions and fits for PSM, using the pairs shown in Figure S10; see Table S2 below for fit data.

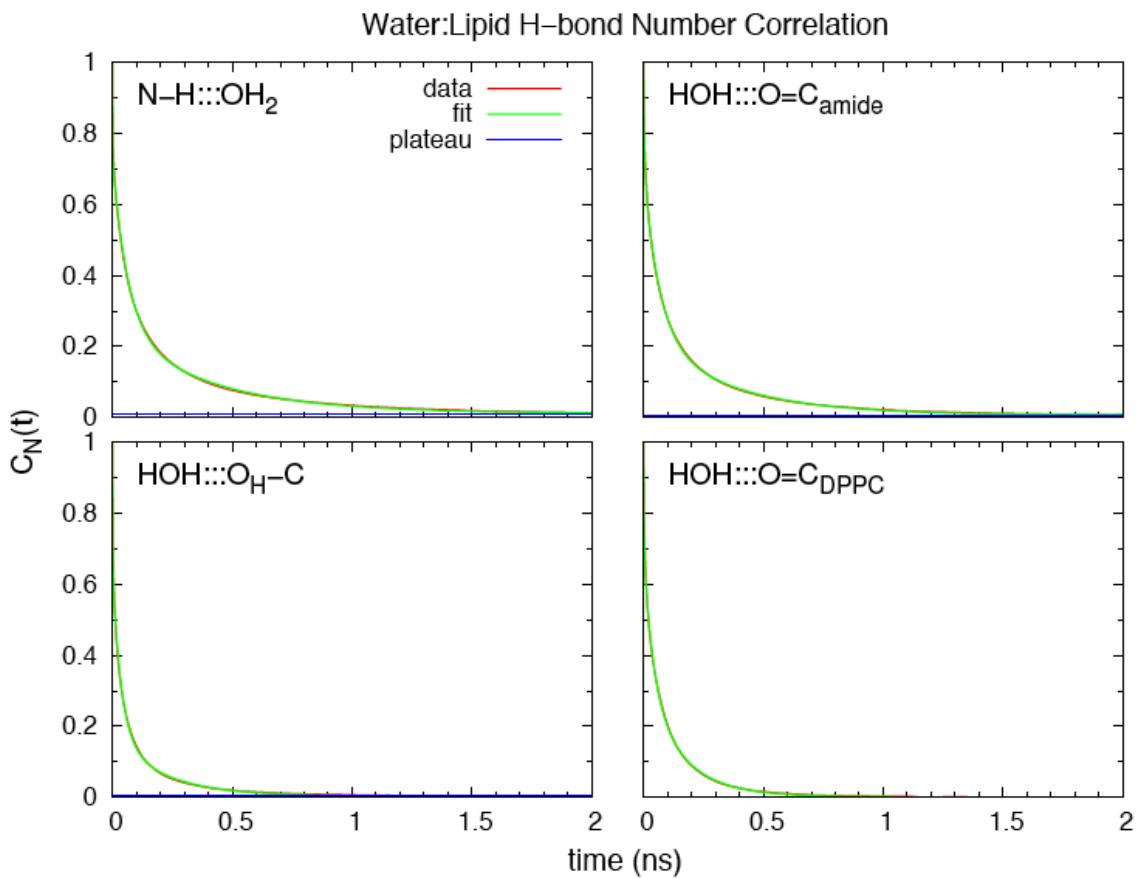


Figure S12. Number correlation functions for water-lipid hydrogen bonds. Correlation functions were evaluated from twenty 5 ns intervals, spaced 20 ns apart; the lower right panel is for the DPPC ester carbonyl O as the acceptor, the other 3 panels represent water:PSM interactions, with the amide H as a donor in the top left panel.

Table S2. Fit coefficients (a) and relaxation times (τ , ns) from 3 exponential fits.

$C_N(t)$	a_1	τ_1	a_2	τ_2	a_3	τ_3	plateau ^a
N-H:::OH ₂	0.268	0.001	0.480	0.064	0.240	0.401	0.011
HOH:::O=C _{amide}	0.291	0.004	0.474	0.065	0.229	0.350	0.006
HOH:::O _H -C	0.363	0.002	0.465	0.037	0.168	0.210	0.003
HOH:::O=C _{DPPC}	0.299	0.003	0.382	0.040	0.318	0.155	0.001

^a The plateau values were free parameters in the three exponential fit, which was constrained by the expression $a_1 + a_2 + a_3 + \text{plateau} = 1.0$.

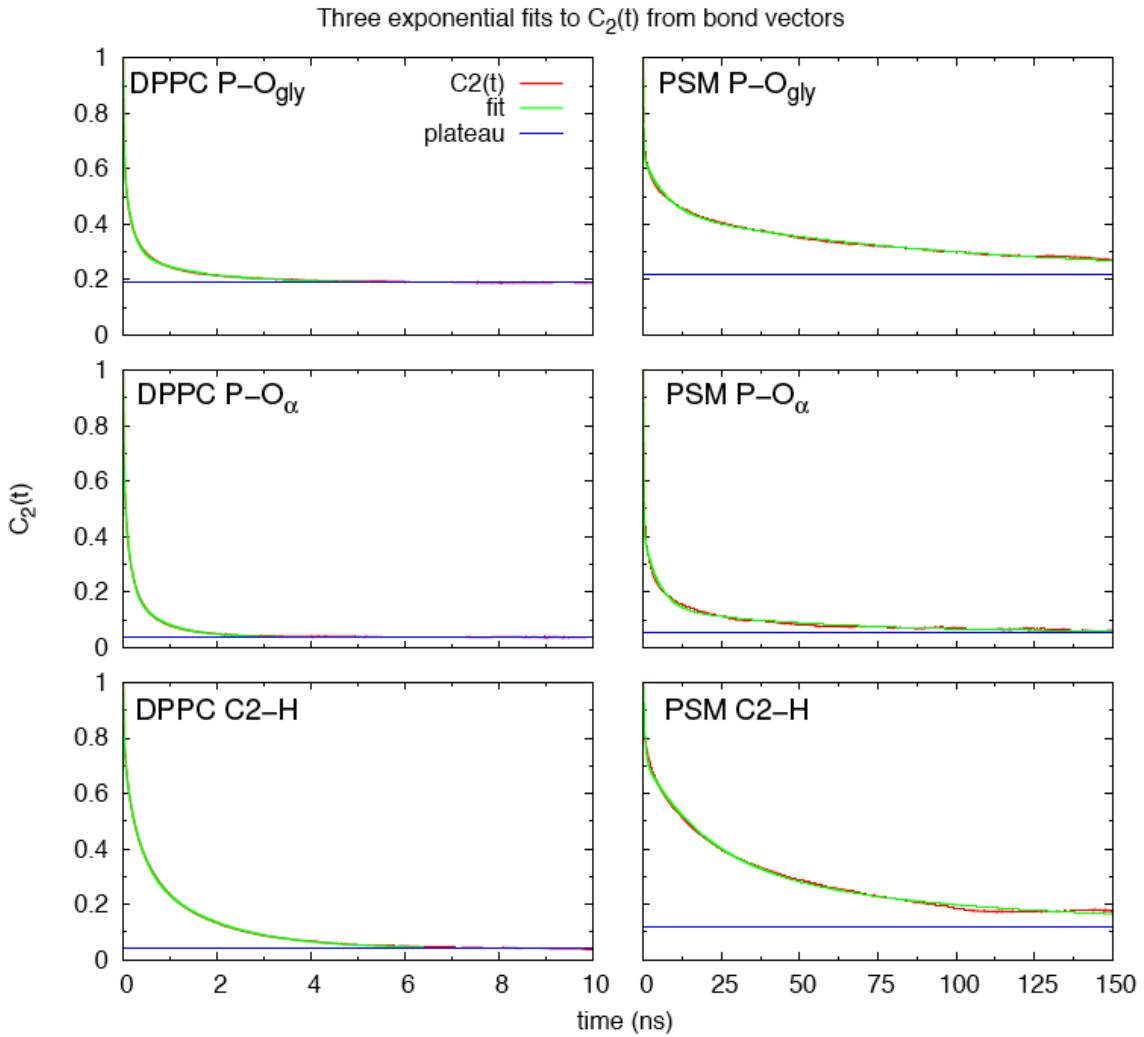


Figure S13. Second rank reorientational correlation functions $C_2(t) = \langle P_2(\hat{\mu}(0) \cdot \hat{\mu}(t)) \rangle$ selected bonds of DPPC (left) and PSM (right).

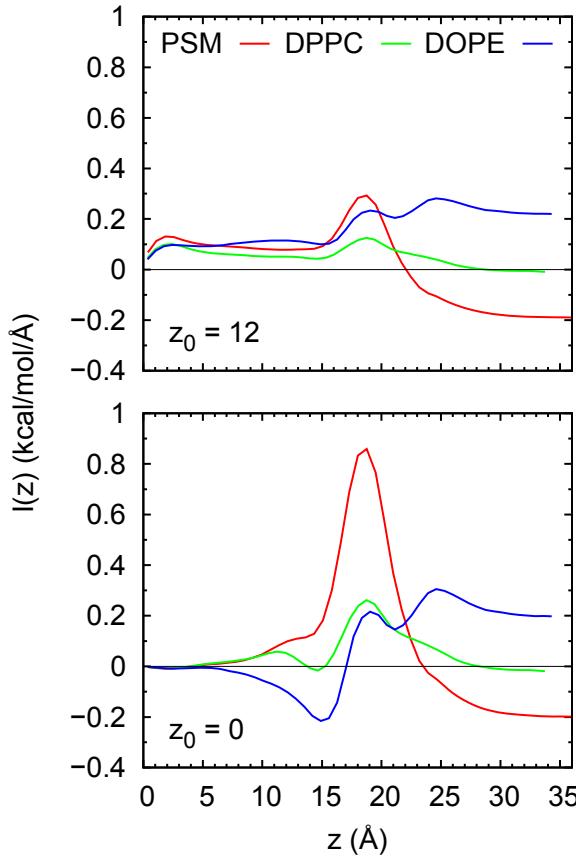


Figure S14. Evaluation of $\bar{F}'(0)$ from integration of the pressure profiles. Equation (2) in the main text can be written more generally as

$$\bar{F}'(0) = - \int_0^{L_z/2} (z - z_0)[p_T(z) - p_N(z)]dz$$

where z_0 is an arbitrary plane in the bilayer. The lower panel of the figure shows the integral

$$I(z) = - \int_0^z (z' - z_0)[p_T(z') - p_N(z')]dz'$$

when $z_0 = 0$ (the bilayer midplane). It is evident that values for $z \approx 35 \text{ \AA}$, which approximate $\bar{F}'(0)$, are quite distinct, lending increased confidence to the result that $\bar{F}'(0)$ for PSM is approximately equal in magnitude to that of DOPE but opposite in sign; i.e., a bilayer of pure PSM has positive curvature. This result does not depend on the value of z_0 : as shown in the top panel for $z_0 = 12 \text{ \AA}$, the intermediate values of $I(z)$ differ for those of $z_0 = 0$, but $\bar{F}'(0)$ is the same.

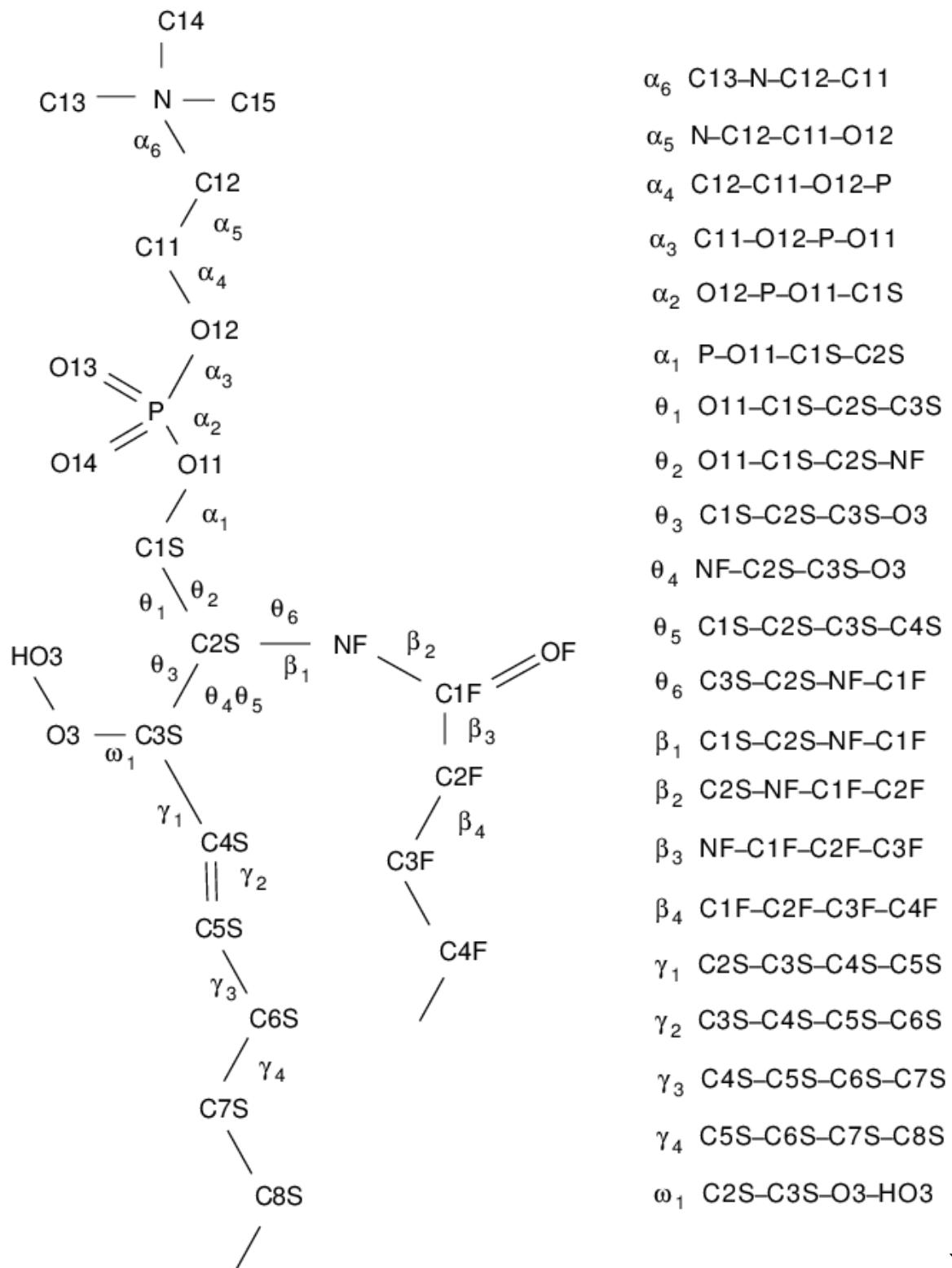


Figure S15. Atom names, and dihedral labels and definitions used for head group conformational analysis, and for the plots in Figure S16.

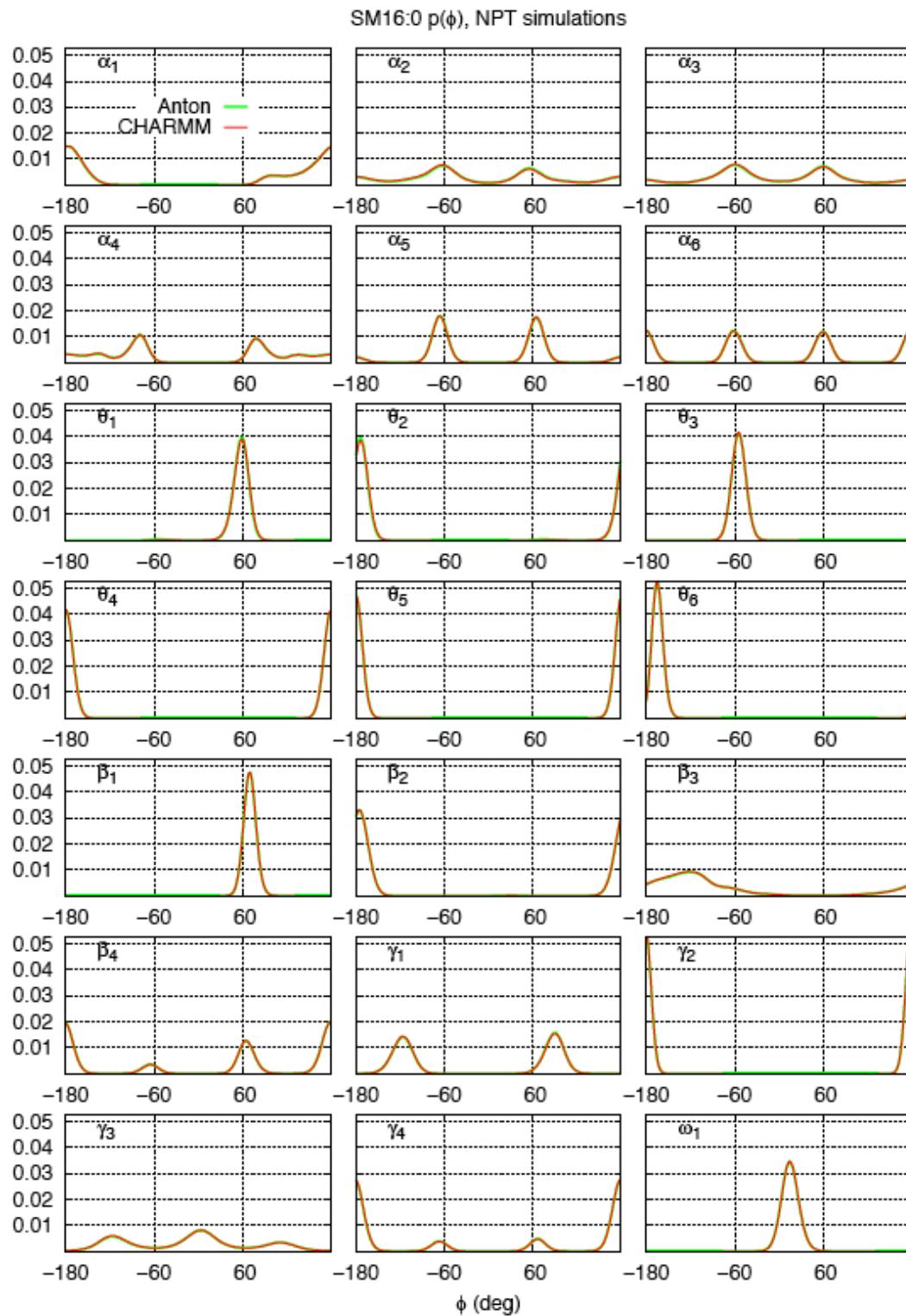


Figure S16. PSM head group dihedral distributions from both CHARMM and ANTON simulations.

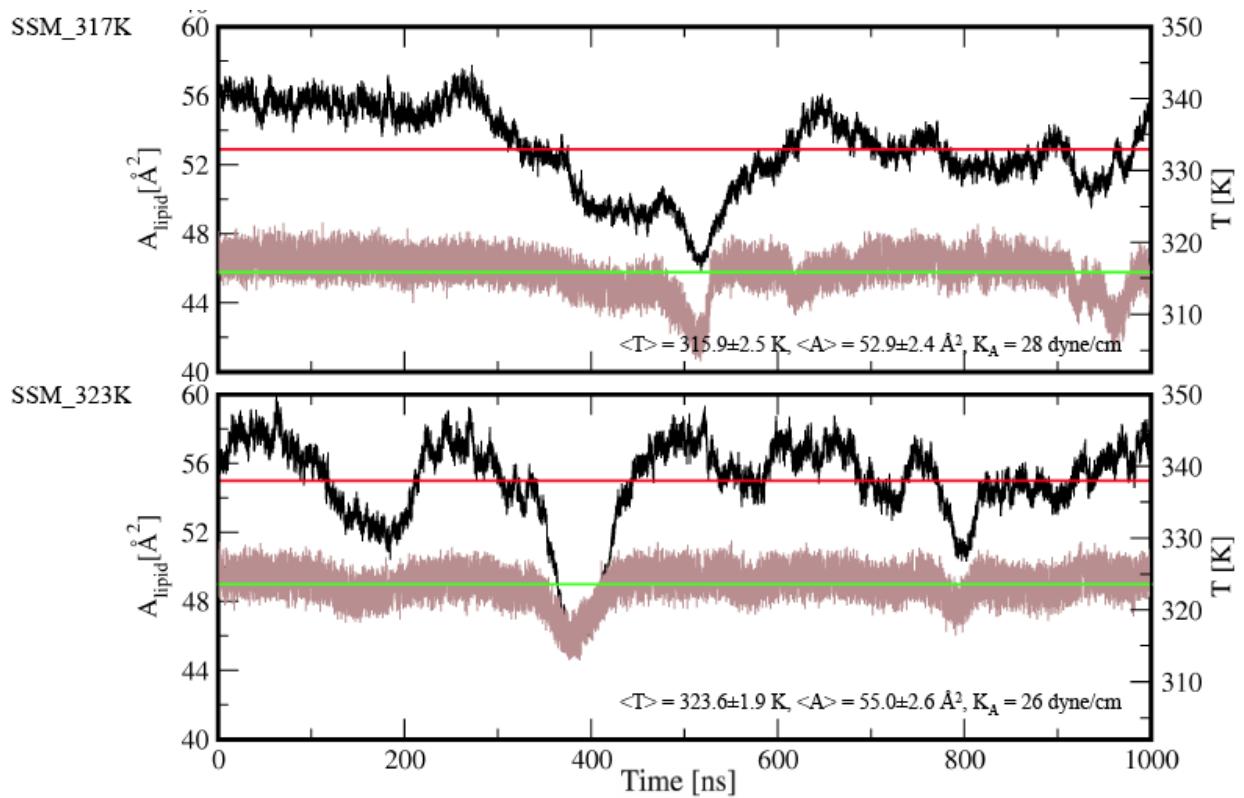


Figure S17. Temperature and area/lipid anomalies observed for SSM on ANTON. The black and red lines are the respective area/lipid time series and averages (left axis); the brown and green lines are time series and averages for the temperature (right axis).

Topology and parameter stream file input for CHARMM.

The following pages are the complete listing of the file with the new molecules and parameters added for sphingomyelins, in CHARMM input format. While complete, using cut-and-paste of these listings to obtain the inputs is inadvisable, due to line wrap issues. The file can be downloaded from <http://terpconnect.umd.edu/~jbklauda/research/download.html>.

```
* Toppar stream for ceramides and sphingomyelins; requires flex format files
* Lipid: top_all36_lipid.rtf and par_all36_lipid.prm
* Protein: top_all36_prot.rtf and par_all36_prot.prm
*
! REFERENCES
!
! PSM and SSM Lipids
!
! Venable, R.M., B. Rogaski, H. Rui, L. Hatcher, A.D. MacKerell, Jr., R.W. Pastor,
! & J.B. Klauda. CHARMM All-Atom Additive Force Field for Sphingomyelin:
! Elucidation of Hydrogen Bonding and of Positive Curvature. In prep. (2014).
!
! Ceramide
!
! Submitted by Sam Tonddast-Navaei of U. Cincinnati. Advised by Russell Devane
!

set nat ?NATC
set app
!We're exploiting what is arguably a bug in the parser. On the left hand side,
!the quotes have priority, so NAT is correctly substituted. On the right hand
!side, the ? has priority and NATC" (sic) is not a valid substitution...
if "@NAT" ne "?NATC" if @nat ne 0 set app append

read rtf card @app
* cer/sm residues
*
36 1

! Ceramides

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MASS    166 CTO1   12.01100 C !
MASS    167 CEL3   12.01100 C !
MASS    168 CTO2   12.01100 C !

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H15B
H15A-C15-H15C
H13C      H14C
H13A-C13---N---C14-H14A (+)
H13B      H14B
H12A-C12-H12B
H11A-C11-H11B
(-) O13 O12
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 |
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 |
 H16F---C16F---H16G
 |
 H16H
 H3S---C3S---O3---HO3
 |
 C4S---H4S
 ||
 C5S---H5S
 |
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 H18T

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!SPHINGOSINE CHAIN

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 !PHOSLPHOCHOLINE
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 BOND C12 N N C13 N C14 N C15
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 BOND C16F H16H
 DOUBLE C4S C5S
 DOUBLE C1F OF
 IMPR C1F C2F NF OF NF C1F C2S HNF
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 IC C17S C16S C15S C14S 1.5337 112.64 180.00 112.58 1.5345
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 IC C13 C12 *N C15 1.4967 109.68 -123.42 110.88 1.4938
 IC C12 N C15 H15A 1.5212 110.88 -177.15 111.31 1.0816
 IC H15A N *C15 H15B 1.0816 111.31 118.98 109.44 1.0875
 IC H15A N *C15 H15C 1.0816 111.31 -122.11 111.46 1.0818
 IC C12 N C14 H14A 1.5212 107.30 -178.89 114.23 1.0739

IC	H14A	N	*C14	H14B	1.0739	114.23	123.44	110.21	1.0968
IC	H14A	N	*C14	H14C	1.0739	114.23	-124.43	111.41	1.0948
IC	C12	N	C13	H13A	1.5212	109.68	176.44	111.15	1.0822
IC	H13A	N	*C13	H13B	1.0822	111.15	121.89	111.75	1.0814
IC	H13A	N	*C13	H13C	1.0822	111.15	-118.50	109.57	1.0892
IC	C13	N	C12	C11	1.4967	109.68	-178.83	116.30	1.5392
IC	C11	N	*C12	H12A	1.5392	116.30	117.04	108.30	1.0985
IC	C11	N	*C12	H12B	1.5392	116.30	-126.61	112.36	1.0787
IC	N	C12	C11	O12	1.5212	116.30	126.79	107.85	1.4239
IC	O12	C12	*C11	H11A	1.4239	107.85	118.10	109.30	1.1131
IC	O12	C12	*C11	H11B	1.4239	107.85	-123.12	113.16	1.1140
IC	C12	C11	O12	P	1.5392	107.85	-68.29	118.09	1.5886
IC	C11	O12	P	O11	1.4239	118.09	-167.58	104.18	1.5725
IC	O11	O12	*P	O13	1.5725	104.18	116.57	107.05	1.4823
IC	O11	O12	*P	O14	1.5725	104.18	-118.73	106.98	1.4805
IC	O12	P	O11	C1S	1.5886	104.18	51.85	121.69	1.4327
IC	P	O11	C1S	C2S	1.5725	121.69	103.66	111.21	1.5586
IC	C2S	O11	*C1S	H1S	1.5586	111.21	118.79	108.35	1.1121
IC	C2S	O11	*C1S	H1T	1.5586	111.21	-123.25	112.85	1.1152
IC	C2S	C3S	O3	H03	1.5343	109.80	46.50	104.54	0.9610
IC	C2S	C3S	C4S	C5S	1.5343	110.14	-84.24	124.28	1.3444
IC	C5S	C3S	*C4S	H4S	1.3444	124.28	178.79	115.93	1.1003
IC	C3S	C4S	C5S	C6S	1.5009	124.28	179.01	124.77	1.5077
IC	C6S	C4S	*C5S	H5S	1.5077	124.77	-178.18	119.00	1.1029
IC	C4S	C5S	C6S	C7S	1.3444	124.77	-117.13	111.46	1.5391
IC	C7S	C5S	*C6S	H6S	1.5391	111.46	120.98	111.41	1.1142
IC	C7S	C5S	*C6S	H6T	1.5391	111.46	-120.60	109.91	1.1125
IC	C5S	C6S	C7S	C8S	1.5077	111.46	-179.61	112.41	1.5354
IC	C8S	C6S	*C7S	H7S	1.5354	112.41	121.26	109.74	1.1138
IC	C8S	C6S	*C7S	H7T	1.5354	112.41	-120.99	109.40	1.1147
IC	C6S	C7S	C8S	C9S	1.5391	112.41	-179.79	112.50	1.5345
IC	C9S	C7S	*C8S	H8S	1.5345	112.50	121.20	108.97	1.1132
IC	C9S	C7S	*C8S	H8T	1.5345	112.50	-121.38	109.07	1.1129
IC	C7S	C8S	C9S	C10S	1.5354	112.50	-179.89	112.66	1.5344
IC	C10S	C8S	*C9S	H9S	1.5344	112.66	121.49	109.05	1.1133
IC	C10S	C8S	*C9S	H9T	1.5344	112.66	-121.38	108.96	1.1136
IC	C8S	C9S	C10S	C11S	1.5345	112.66	-179.87	112.54	1.5345
IC	C11S	C9S	*C10S	H10S	1.5345	112.54	121.27	109.07	1.1133
IC	C11S	C9S	*C10S	H10T	1.5345	112.54	-121.34	109.11	1.1132
IC	C9S	C10S	C11S	C12S	1.5344	112.54	179.97	112.61	1.5344
IC	C12S	C10S	*C11S	H11S	1.5344	112.61	121.36	109.09	1.1133
IC	C12S	C10S	*C11S	H11T	1.5344	112.61	-121.33	109.07	1.1134
IC	C10S	C11S	C12S	C13S	1.5345	112.61	-179.96	112.58	1.5345
IC	C13S	C11S	*C12S	H12S	1.5345	112.58	121.29	109.12	1.1133
IC	C13S	C11S	*C12S	H12T	1.5345	112.58	-121.31	109.15	1.1133
IC	C11S	C12S	C13S	C14S	1.5344	112.58	179.97	112.58	1.5345
IC	C14S	C12S	*C13S	H13S	1.5345	112.58	121.30	109.13	1.1133
IC	C14S	C12S	*C13S	H13T	1.5345	112.58	-121.30	109.13	1.1133
IC	C12S	C13S	C14S	C15S	1.5345	112.58	-179.99	112.58	1.5345
IC	C15S	C13S	*C14S	H14S	1.5345	112.58	121.30	109.12	1.1133
IC	C15S	C13S	*C14S	H14T	1.5345	112.58	-121.31	109.12	1.1133
IC	C13S	C14S	C15S	C16S	1.5345	112.58	180.00	112.58	1.5346
IC	C16S	C14S	*C15S	H15S	1.5346	112.58	121.30	109.11	1.1133
IC	C16S	C14S	*C15S	H15T	1.5346	112.58	-121.30	109.11	1.1133
IC	C14S	C15S	C16S	C17S	1.5345	112.58	180.00	112.64	1.5337
IC	C17S	C15S	*C16S	H16S	1.5337	112.64	121.32	109.14	1.1133
IC	C17S	C15S	*C16S	H16T	1.5337	112.64	-121.32	109.14	1.1133
IC	C15S	C16S	C17S	C18S	1.5346	112.64	180.00	113.23	1.5309
IC	C18S	C16S	*C17S	H17S	1.5309	113.23	121.67	108.77	1.1142
IC	C18S	C16S	*C17S	H17T	1.5309	113.23	-121.67	108.77	1.1142
IC	C16S	C17S	C18S	H18S	1.5337	113.23	180.00	110.63	1.1113
IC	H18S	C17S	*C18S	H18T	1.1113	110.63	120.07	110.47	1.1114
IC	H18S	C17S	*C18S	H18U	1.1113	110.63	-120.07	110.47	1.1114
IC	C2S	NF	C1F	C2F	1.4406	124.85	-156.83	114.97	1.4964
IC	C2F	NF	*C1F	OF	1.4964	114.97	162.54	122.11	1.2268
IC	NF	C1F	C2F	C3F	1.3364	114.97	-146.57	111.92	1.5432
IC	C3F	C1F	*C2F	H2F	1.5432	111.92	-120.83	107.93	1.1115
IC	H2F	C1F	*C2F	H2G	1.1115	107.93	-117.30	109.04	1.1113
IC	C1F	C2F	C3F	C4F	1.4964	111.92	179.54	112.84	1.5310
IC	C4F	C2F	*C3F	H3F	1.5310	112.84	-121.63	109.37	1.1145
IC	H3F	C2F	*C3F	H3G	1.1145	109.37	-117.07	109.08	1.1157

IC	C2F	C3F	C4F	C5F	1.5434	112.23	174.86	112.58	1.5342
IC	C5F	C3F	*C4F	H4F	1.5342	112.58	-120.18	108.65	1.1140
IC	H4F	C3F	*C4F	H4G	1.1140	108.65	-117.52	109.62	1.1127
IC	C3F	C4F	C5F	C6F	1.5340	112.58	-174.06	112.39	1.5340
IC	C6F	C4F	*C5F	H5F	1.5340	112.39	-122.01	109.38	1.1130
IC	H5F	C4F	*C5F	H5G	1.1130	109.38	-117.34	108.92	1.1134
IC	C4F	C5F	C6F	C7F	1.5342	112.39	178.27	112.75	1.5340
IC	C7F	C5F	*C6F	H6F	1.5340	112.75	-120.93	108.96	1.1137
IC	H6F	C5F	*C6F	H6G	1.1137	108.96	-117.41	109.22	1.1131
IC	C5F	C6F	C7F	C8F	1.5340	112.75	-178.89	112.33	1.5342
IC	C8F	C6F	*C7F	H7F	1.5342	112.33	-121.40	109.26	1.1132
IC	H7F	C6F	*C7F	H7G	1.1132	109.26	-117.45	109.12	1.1130
IC	C6F	C7F	C8F	C9F	1.5340	112.33	179.09	112.77	1.5340
IC	C9F	C7F	*C8F	H8F	1.5340	112.77	-121.23	109.16	1.1134
IC	H8F	C7F	*C8F	H8G	1.1134	109.16	-117.38	109.08	1.1133
IC	C7F	C8F	C9F	C10F	1.5342	112.77	179.02	112.34	1.5343
IC	C10F	C8F	*C9F	H9F	1.5343	112.34	-121.27	109.27	1.1132
IC	H9F	C8F	*C9F	H9G	1.1132	109.27	-117.45	109.10	1.1130
IC	C8F	C9F	C10F	C11F	1.5340	112.34	178.16	112.74	1.5340
IC	C11F	C9F	*C10F	H10F	1.5340	112.74	-121.18	109.16	1.1134
IC	H10F	C9F	*C10F	H10G	1.1134	109.16	-117.35	109.10	1.1132
IC	C9F	C10F	C11F	C12F	1.5343	112.74	178.91	112.38	1.5343
IC	C12F	C10F	*C11F	H11F	1.5343	112.38	-121.39	109.34	1.1131
IC	H11F	C10F	*C11F	H11G	1.1131	109.34	-117.43	108.98	1.1132
IC	C10F	C11F	C12F	C13F	1.5340	112.38	176.93	112.68	1.5340
IC	C13F	C11F	*C12F	H12F	1.5340	112.68	-121.02	109.05	1.1134
IC	H12F	C11F	*C12F	H12G	1.1134	109.05	-117.34	109.20	1.1131
IC	C11F	C12F	C13F	C14F	1.5343	112.68	-179.93	112.45	1.5345
IC	C14F	C12F	*C13F	H13F	1.5345	112.45	-121.60	109.42	1.1129
IC	H13F	C12F	*C13F	H13G	1.1129	109.42	-117.40	108.85	1.1134
IC	C12F	C13F	C14F	C15F	1.5340	112.45	175.93	112.68	1.5334
IC	C15F	C13F	*C14F	H14F	1.5334	112.68	-121.00	109.06	1.1131
IC	H14F	C13F	*C14F	H14G	1.1131	109.06	-117.24	109.28	1.1130
IC	C13F	C14F	C15F	C16F	1.5345	112.68	-178.23	113.16	1.5306
IC	C16F	C14F	*C15F	H15F	1.5306	113.16	-121.86	108.84	1.1140
IC	H15F	C14F	*C15F	H15G	1.1140	108.84	-116.70	108.76	1.1144
IC	C14F	C15F	C16F	H16F	1.5334	113.16	-59.91	110.49	1.1113
IC	H16F	C15F	*C16F	H16G	1.1113	110.49	119.86	110.41	1.1115
IC	H16F	C15F	*C16F	H16H	1.1113	110.49	-120.11	110.63	1.1112
IC	C4S	C2S	*C3S	O3	1.5009	110.14	118.36	109.80	1.4235 ! OH chirality (ch
IC	C4S	C2S	*C3S	H3S	1.5009	110.14	-121.14	110.10	1.1140 ! OH chirality (ch
IC	O11	C1S	C2S	C3S	1.4327	111.21	175.67	114.25	1.5343 ! assume ok
IC	C1F	C2S	*NF	HNF	1.3364	124.85	-157.81	113.39	1.0062 ! assume ok
! IC	C1S	C2S	C3S	C4S	1.5586	114.25	-70.50	110.14	1.5009! assume redundant
! SWAPPED									
IC	C4S	C3S	C2S	C1S	1.5009	110.14	160.94	114.25	1.5586
IC	C1S	C2S	NF	C1F	1.5586	111.52	-70.50	124.85	1.3364
! SWAPPED									
IC	C3S	C1S	*C2S	NF	1.5343	114.25	120.00	111.52	1.4406
IC	C3S	C1S	*C2S	H2S	1.5343	114.25	-120.00	105.03	1.1194

PATCH FIRST NONE LAST NONE

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RESI SSM      0.00 ! sphingomyelin d18:1/18:0
GROUP          !
ATOM N      NTL   -0.60 !
ATOM C15    CTL5   -0.35 !
ATOM H15A   HL    0.25 !
ATOM H15B   HL    0.25 !
ATOM H15C   HL    0.25 !
ATOM C14    CTL5   -0.35 !
ATOM H14A   HL    0.25 !
ATOM H14B   HL    0.25 !
ATOM H14C   HL    0.25 !
ATOM C13    CTL5   -0.35 !
ATOM H13A   HL    0.25 !
ATOM H13B   HL    0.25 !
ATOM H13C   HL    0.25 !
ATOM C12    CTL2   -0.10 !
ATOM H12A   HL    0.25 !
ATOM H12B   HL    0.25 !
GROUP          !

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ATOM C11 CTL2 -0.08 !
 ATOM H11A HAL2 0.09 !
 ATOM H11B HAL2 0.09 !
 ATOM P PL 1.50 !
 ATOM O13 O2L -0.78 !
 ATOM O14 O2L -0.78 !
 ATOM O11 OSLP -0.57 !
 ATOM O12 OSLP -0.57 !
 ATOM C1S CTL2 -0.08 !
 ATOM H1S HAL2 0.09 !
 ATOM H1T HAL2 0.09 !
 GROUP !
 ATOM NF NHL -0.70 !
 ATOM HNF H 0.35 !
 ATOM C2S CTL1 0.30 !
 ATOM H2S HAL1 0.05 !
 GROUP !
 ATOM C3S CTO1 0.50 !
 ATOM H3S HAL1 -0.11 !
 ATOM O3 OHL -0.69 !
 ATOM HO3 HOL 0.30 !
 GROUP !
 ATOM C4S CEL3 -0.15 !
 ATOM H4S HEL1 0.15 !
 GROUP !
 ATOM C5S CEL3 -0.15 !
 ATOM H5S HEL1 0.15 !
 GROUP !
 ATOM C6S CTL2 -0.18 !
 ATOM H6S HAL2 0.09 !
 ATOM H6T HAL2 0.09 !
 GROUP !
 ATOM C7S CTL2 -0.18 !
 ATOM H7S HAL2 0.09 !
 ATOM H7T HAL2 0.09 !
 GROUP !
 ATOM C8S CTL2 -0.18 !
 ATOM H8S HAL2 0.09 !
 ATOM H8T HAL2 0.09 !
 GROUP !
 ATOM C9S CTL2 -0.18 !
 ATOM H9S HAL2 0.09 !
 ATOM H9T HAL2 0.09 !
 GROUP !
 ATOM C10S CTL2 -0.18 !
 ATOM H10S HAL2 0.09 !
 ATOM H10T HAL2 0.09 !
 GROUP
 ATOM C11S CTL2 -0.18 !
 ATOM H11S HAL2 0.09 !
 ATOM H11T HAL2 0.09 !
 GROUP
 ATOM C12S CTL2 -0.18 !
 ATOM H12S HAL2 0.09 !
 ATOM H12T HAL2 0.09 !
 GROUP
 ATOM C13S CTL2 -0.18 !
 ATOM H13S HAL2 0.09 !
 ATOM H13T HAL2 0.09 !
 GROUP
 ATOM C14S CTL2 -0.18 !
 ATOM H14S HAL2 0.09 !
 ATOM H14T HAL2 0.09 !
 GROUP
 ATOM C15S CTL2 -0.18 !
 ATOM H15S HAL2 0.09 !
 ATOM H15T HAL2 0.09 !
 GROUP
 ATOM C16S CTL2 -0.18 !
 ATOM H16S HAL2 0.09 !
 ATOM H16T HAL2 0.09 !

GROUP
 ATOM C17S CTL2 -0.18 !
 ATOM H17S HAL2 0.09 !
 ATOM H17T HAL2 0.09 !
 GROUP !
 ATOM C18S CTL3 -0.27 !
 ATOM H18S HAL3 0.09 !
 ATOM H18T HAL3 0.09 !
 ATOM H18U HAL3 0.09 !
 GROUP ! FATTY ACID CHAIN
 ATOM C1F C 0.55 !
 ATOM O1F O -0.60 !
 ATOM C2F CTL2 -0.07 !
 ATOM H2F HAL2 0.06 !
 ATOM H2G HAL2 0.06 !
 GROUP !
 ATOM C3F CTL2 -0.18 !
 ATOM H3F HAL2 0.09 !
 ATOM H3G HAL2 0.09 !
 GROUP !
 ATOM C4F CTL2 -0.18 !
 ATOM H4F HAL2 0.09 !
 ATOM H4G HAL2 0.09 !
 GROUP !
 ATOM C5F CTL2 -0.18 !
 ATOM H5F HAL2 0.09 !
 ATOM H5G HAL2 0.09 !
 GROUP !
 ATOM C6F CTL2 -0.18 !
 ATOM H6F HAL2 0.09 !
 ATOM H6G HAL2 0.09 !
 GROUP !
 ATOM C7F CTL2 -0.18 !
 ATOM H7F HAL2 0.09 !
 ATOM H7G HAL2 0.09 !
 GROUP !
 ATOM C8F CTL2 -0.18 !
 ATOM H8F HAL2 0.09 !
 ATOM H8G HAL2 0.09 !
 GROUP !
 ATOM C9F CTL2 -0.18 !
 ATOM H9F HAL2 0.09 !
 ATOM H9G HAL2 0.09 !
 GROUP !
 ATOM C10F CTL2 -0.18 !
 ATOM H10F HAL2 0.09 !
 ATOM H10G HAL2 0.09 !
 GROUP !
 ATOM C11F CTL2 -0.18 !
 ATOM H11F HAL2 0.09 !
 ATOM H11G HAL2 0.09 !
 GROUP !
 ATOM C12F CTL2 -0.18 !
 ATOM H12F HAL2 0.09 !
 ATOM H12G HAL2 0.09 !
 GROUP !
 ATOM C13F CTL2 -0.18 !
 ATOM H13F HAL2 0.09 !
 ATOM H13G HAL2 0.09 !
 GROUP !
 ATOM C14F CTL2 -0.18 !
 ATOM H14F HAL2 0.09 !
 ATOM H14G HAL2 0.09 !
 GROUP !
 ATOM C15F CTL2 -0.18 !
 ATOM H15F HAL2 0.09 !
 ATOM H15G HAL2 0.09 !
 GROUP !
 ATOM C16F CTL2 -0.18 !
 ATOM H16F HAL2 0.09 !
 ATOM H16G HAL2 0.09 !

GROUP !
 ATOM C17F CTL2 -0.18 !
 ATOM H17F HAL2 0.09 !
 ATOM H17G HAL2 0.09 !
 GROUP !
 ATOM C18F CTL3 -0.27 !
 ATOM H18F HAL3 0.09 !
 ATOM H18G HAL3 0.09 !
 ATOM H18H HAL3 0.09 !

 !SPHINGOSINE CHAIN
 BOND C18S H18S C18S H18T C18S H18U
 BOND C18S C17S C17S H17S C17S H17T
 BOND C17S C16S C16S H16S C16S H16T
 BOND C16S C15S C15S H15S C15S H15T
 BOND C15S C14S C14S H14S C14S H14T
 BOND C14S C13S C13S H13S C13S H13T
 BOND C13S C12S C12S H12S C12S H12T
 BOND C12S C11S C11S H11S C11S H11T
 BOND C11S C10S C10S H10S C10S H10T
 BOND C10S C9S C9S H9S C9S H9T
 BOND C9S C8S C8S H8S C8S H8T
 BOND C8S C7S C7S H7S C7S H7T
 BOND C7S C6S C6S H6S C6S H6T
 BOND C6S C5S C5S H5S C4S H4S
 BOND C4S C3S C3S H3S C3S O3
 BOND O3 HO3
 BOND C3S C2S C2S H2S
 BOND C2S C1S C1S H1S C1S H1T
 BOND C1S O11
 !PHOSLPHOCHOLINE
 BOND O11 P O12 P O13 P O14 P
 BOND O12 C11 C11 H11A C11 H11B
 BOND C11 C12 C12 H12A C12 H12B
 BOND C12 N N C13 N C14 N C15
 BOND C13 H13A C13 H13B C13 H13C
 BOND C14 H14A C14 H14B C14 H14C
 BOND C15 H15A C15 H15B C15 H15C
 !FATTY ACID CHAIN
 BOND C2S NF NF HNF NF C1F
 BOND C1F C2F C2F H2F C2F H2G
 BOND C2F C3F C3F H3F C3F H3G
 BOND C3F C4F C4F H4F C4F H4G
 BOND C4F C5F C5F H5F C5F H5G
 BOND C5F C6F C6F H6F C6F H6G
 BOND C6F C7F C7F H7F C7F H7G
 BOND C7F C8F C8F H8F C8F H8G
 BOND C8F C9F C9F H9F C9F H9G
 BOND C9F C10F C10F H10F C10F H10G
 BOND C10F C11F C11F H11F C11F H11G
 BOND C11F C12F C12F H12F C12F H12G
 BOND C12F C13F C13F H13F C13F H13G
 BOND C13F C14F C14F H14F C14F H14G
 BOND C14F C15F C15F H15F C15F H15G
 BOND C15F C16F C16F H16F C16F H16G
 BOND C16F C17F C17F H17F C17F H17G
 BOND C17F C18F C18F H18F C18F H18G
 BOND C18F H18H

 DOUBLE C4S C5S
 DOUBLE C1F OF
 IMPR C1F C2F NF OF NF C1F C2S HNF

 IC C13 C12 *N C14 1.4967 109.68 116.91 107.30 1.5040
 IC C13 C12 *N C15 1.4967 109.68 -123.42 110.88 1.4938
 IC C12 N C15 H15A 1.5212 110.88 -177.15 111.31 1.0816
 IC H15A N *C15 H15B 1.0816 111.31 118.98 109.44 1.0875
 IC H15A N *C15 H15C 1.0816 111.31 -122.11 111.46 1.0818
 IC C12 N C14 H14A 1.5212 107.30 -178.89 114.23 1.0739
 IC H14A N *C14 H14B 1.0739 114.23 123.44 110.21 1.0968
 IC H14A N *C14 H14C 1.0739 114.23 -124.43 111.41 1.0948

IC	C12	N	C13	H13A	1.5212	109.68	176.44	111.15	1.0822
IC	H13A	N	*C13	H13B	1.0822	111.15	121.89	111.75	1.0814
IC	H13A	N	*C13	H13C	1.0822	111.15	-118.50	109.57	1.0892
IC	C13	N	C12	C11	1.4967	109.68	-178.83	116.30	1.5392
IC	C11	N	*C12	H12A	1.5392	116.30	117.04	108.30	1.0985
IC	C11	N	*C12	H12B	1.5392	116.30	-126.61	112.36	1.0787
IC	N	C12	C11	O12	1.5212	116.30	126.79	107.85	1.4239
IC	O12	C12	*C11	H11A	1.4239	107.85	118.10	109.30	1.1131
IC	O12	C12	*C11	H11B	1.4239	107.85	-123.12	113.16	1.1140
IC	C12	C11	O12	P	1.5392	107.85	-68.29	118.09	1.5886
IC	C11	O12	P	O11	1.4239	118.09	-167.58	104.18	1.5725
IC	O11	O12	*P	O13	1.5725	104.18	116.57	107.05	1.4823
IC	O11	O12	*P	O14	1.5725	104.18	-118.73	106.98	1.4805
IC	O12	P	O11	C1S	1.5886	104.18	51.85	121.69	1.4327
IC	P	O11	C1S	C2S	1.5725	121.69	103.66	111.21	1.5586
IC	C2S	O11	*C1S	H1S	1.5586	111.21	118.79	108.35	1.1121
IC	C2S	O11	*C1S	H1T	1.5586	111.21	-123.25	112.85	1.1152
IC	O11	C1S	C2S	C3S	1.4327	111.21	175.67	114.25	1.5343
IC	C3S	C1S	*C2S	NF	1.5343	114.25	120.00	111.52	1.4406
IC	C3S	C1S	*C2S	H2S	1.5343	114.25	-120.00	105.03	1.1194
IC	C1S	C2S	NF	C1F	1.5586	111.52	160.94	124.85	1.3364
IC	C1F	C2S	*NF	HNF	1.3364	124.85	-157.81	113.39	1.0062
IC	C1S	C2S	C3S	C4S	1.5586	114.25	170.50	110.14	1.5009
IC	C4S	C2S	*C3S	O3	1.5009	110.14	118.36	109.80	1.4235
IC	C4S	C2S	*C3S	H3S	1.5009	110.14	-121.14	110.10	1.1140
IC	C2S	C3S	O3	H03	1.5343	109.80	46.50	104.54	0.9610
IC	C2S	C3S	C4S	C5S	1.5343	110.14	-84.24	124.28	1.3444
IC	C5S	C3S	*C4S	H4S	1.3444	124.28	178.79	115.93	1.1003
IC	C3S	C4S	C5S	C6S	1.5009	124.28	179.01	124.77	1.5077
IC	C6S	C4S	*C5S	H5S	1.5077	124.77	-178.18	119.00	1.1029
IC	C4S	C5S	C6S	C7S	1.3444	124.77	-117.13	111.46	1.5391
IC	C7S	C5S	*C6S	H6S	1.5391	111.46	120.98	111.41	1.1142
IC	C7S	C5S	*C6S	H6T	1.5391	111.46	-120.60	109.91	1.1125
IC	C5S	C6S	C7S	C8S	1.5077	111.46	-179.61	112.41	1.5354
IC	C8S	C6S	*C7S	H7S	1.5354	112.41	121.26	109.74	1.1138
IC	C8S	C6S	*C7S	H7T	1.5354	112.41	-120.99	109.40	1.1147
IC	C6S	C7S	C8S	C9S	1.5391	112.41	-179.79	112.50	1.5345
IC	C9S	C7S	*C8S	H8S	1.5345	112.50	121.20	108.97	1.1132
IC	C9S	C7S	*C8S	H8T	1.5345	112.50	-121.38	109.07	1.1129
IC	C7S	C8S	C9S	C10S	1.5354	112.50	-179.89	112.66	1.5344
IC	C10S	C8S	*C9S	H9S	1.5344	112.66	121.49	109.05	1.1133
IC	C10S	C8S	*C9S	H9T	1.5344	112.66	-121.38	108.96	1.1136
IC	C8S	C9S	C10S	C11S	1.5345	112.66	-179.87	112.54	1.5345
IC	C11S	C9S	*C10S	H10S	1.5345	112.54	121.27	109.07	1.1133
IC	C11S	C9S	*C10S	H10T	1.5345	112.54	-121.34	109.11	1.1132
IC	C9S	C10S	C11S	C12S	1.5344	112.54	179.97	112.61	1.5344
IC	C12S	C10S	*C11S	H11S	1.5344	112.61	121.36	109.09	1.1133
IC	C12S	C10S	*C11S	H11T	1.5344	112.61	-121.33	109.07	1.1134
IC	C10S	C11S	C12S	C13S	1.5345	112.61	-179.96	112.58	1.5345
IC	C13S	C11S	*C12S	H12S	1.5345	112.58	121.29	109.12	1.1133
IC	C13S	C11S	*C12S	H12T	1.5345	112.58	-121.31	109.15	1.1133
IC	C11S	C12S	C13S	C14S	1.5344	112.58	179.97	112.58	1.5345
IC	C14S	C12S	*C13S	H13S	1.5345	112.58	121.30	109.13	1.1133
IC	C14S	C12S	*C13S	H13T	1.5345	112.58	-121.30	109.13	1.1133
IC	C12S	C13S	C14S	C15S	1.5345	112.58	-179.99	112.58	1.5345
IC	C15S	C13S	*C14S	H14S	1.5345	112.58	121.30	109.12	1.1133
IC	C15S	C13S	*C14S	H14T	1.5345	112.58	-121.31	109.12	1.1133
IC	C13S	C14S	C15S	C16S	1.5345	112.58	180.00	112.58	1.5346
IC	C16S	C14S	*C15S	H15S	1.5346	112.58	121.30	109.11	1.1133
IC	C16S	C14S	*C15S	H15T	1.5346	112.58	-121.30	109.11	1.1133
IC	C14S	C15S	C16S	C17S	1.5345	112.58	180.00	112.64	1.5337
IC	C17S	C15S	*C16S	H16S	1.5337	112.64	121.32	109.14	1.1133
IC	C17S	C15S	*C16S	H16T	1.5337	112.64	-121.32	109.14	1.1133
IC	C15S	C16S	C17S	C18S	1.5346	112.64	180.00	113.23	1.5309
IC	C18S	C16S	*C17S	H17S	1.5309	113.23	121.67	108.77	1.1142
IC	C18S	C16S	*C17S	H17T	1.5309	113.23	-121.67	108.77	1.1142
IC	C16S	C17S	C18S	H18S	1.5337	113.23	180.00	110.63	1.1113
IC	H18S	C17S	*C18S	H18T	1.1113	110.63	120.07	110.47	1.1114
IC	H18S	C17S	*C18S	H18U	1.1113	110.63	-120.07	110.47	1.1114
IC	C2S	NF	C1F	C2F	1.4406	124.85	-156.83	114.97	1.4964
IC	C2F	NF	*C1F	OF	1.4964	114.97	162.54	122.11	1.2268

IC	NF	C1F	C2F	C3F	1.3364	114.97	146.57	111.92	1.5432
IC	C3F	C1F	*C2F	H2F	1.5432	111.92	-120.83	107.93	1.1115
IC	H2F	C1F	*C2F	H2G	1.1115	107.93	-117.30	109.04	1.1113
IC	C1F	C2F	C3F	C4F	1.4964	111.92	59.54	112.84	1.5310
IC	C4F	C2F	*C3F	H3F	1.5310	112.84	-121.63	109.37	1.1145
IC	H3F	C2F	*C3F	H3G	1.1145	109.37	-117.07	109.08	1.1157
IC	C2F	C3F	C4F	C5F	1.5434	112.23	174.86	112.58	1.5342
IC	C5F	C3F	*C4F	H4F	1.5342	112.58	-120.18	108.65	1.1140
IC	H4F	C3F	*C4F	H4G	1.1140	108.65	-117.52	109.62	1.1127
IC	C3F	C4F	C5F	C6F	1.5340	112.58	-174.06	112.39	1.5340
IC	C6F	C4F	*C5F	H5F	1.5340	112.39	-122.01	109.38	1.1130
IC	H5F	C4F	*C5F	H5G	1.1130	109.38	-117.34	108.92	1.1134
IC	C4F	C5F	C6F	C7F	1.5342	112.39	178.27	112.75	1.5340
IC	C7F	C5F	*C6F	H6F	1.5340	112.75	-120.93	108.96	1.1137
IC	H6F	C5F	*C6F	H6G	1.1137	108.96	-117.41	109.22	1.1131
IC	C5F	C6F	C7F	C8F	1.5340	112.75	-178.89	112.33	1.5342
IC	C8F	C6F	*C7F	H7F	1.5342	112.33	-121.40	109.26	1.1132
IC	H7F	C6F	*C7F	H7G	1.1132	109.26	-117.45	109.12	1.1130
IC	C6F	C7F	C8F	C9F	1.5340	112.33	179.09	112.77	1.5340
IC	C9F	C7F	*C8F	H8F	1.5340	112.77	-121.23	109.16	1.1134
IC	H8F	C7F	*C8F	H8G	1.1134	109.16	-117.38	109.08	1.1133
IC	C7F	C8F	C9F	C10F	1.5342	112.77	179.02	112.34	1.5343
IC	C10F	C8F	*C9F	H9F	1.5343	112.34	-121.27	109.27	1.1132
IC	H9F	C8F	*C9F	H9G	1.1132	109.27	-117.45	109.10	1.1130
IC	C8F	C9F	C10F	C11F	1.5340	112.34	178.16	112.74	1.5340
IC	C11F	C9F	*C10F	H10F	1.5340	112.74	-121.18	109.16	1.1134
IC	H10F	C9F	*C10F	H10G	1.1134	109.16	-117.35	109.10	1.1132
IC	C9F	C10F	C11F	C12F	1.5343	112.74	178.91	112.38	1.5343
IC	C12F	C10F	*C11F	H11F	1.5343	112.38	-121.39	109.34	1.1131
IC	H11F	C10F	*C11F	H11G	1.1131	109.34	-117.43	108.98	1.1132
IC	C10F	C11F	C12F	C13F	1.5340	112.38	176.93	112.68	1.5340
IC	C13F	C11F	*C12F	H12F	1.5340	112.68	-121.02	109.05	1.1134
IC	H12F	C11F	*C12F	H12G	1.1134	109.05	-117.34	109.20	1.1131
IC	C11F	C12F	C13F	C14F	1.5343	112.68	-179.93	112.45	1.5345
IC	C14F	C12F	*C13F	H13F	1.5345	112.45	-121.60	109.42	1.1129
IC	H13F	C12F	*C13F	H13G	1.1129	109.42	-117.40	108.85	1.1134
IC	C12F	C13F	C14F	C15F	1.5340	112.45	175.93	112.68	1.5334
IC	C15F	C13F	*C14F	H14F	1.5334	112.68	-121.00	109.06	1.1131
IC	H14F	C13F	*C14F	H14G	1.1131	109.06	-117.24	109.28	1.1130
IC	C13F	C14F	C15F	C16F	1.5345	112.68	-178.23	113.16	1.5306
IC	C16F	C14F	*C15F	H15F	1.5306	113.16	-121.86	108.84	1.1140
IC	H15F	C14F	*C15F	H15G	1.1140	108.84	-116.70	108.76	1.1144
IC	C14F	C15F	C16F	C17F	1.5345	112.68	-178.23	113.16	1.5306
IC	C17F	C15F	*C16F	H16F	1.5306	113.16	-121.86	108.84	1.1140
IC	H16F	C15F	*C16F	H16G	1.1140	108.84	-116.70	108.76	1.1144
IC	C15F	C16F	C17F	C18F	1.5345	112.68	-178.23	113.16	1.5306
IC	C18F	C16F	*C17F	H17F	1.5306	113.16	-121.86	108.84	1.1140
IC	H17F	C16F	*C17F	H17G	1.1140	108.84	-116.70	108.76	1.1144
IC	C16F	C17F	C18F	H18F	1.5334	113.16	-59.91	110.49	1.1113
IC	H18F	C17F	*C18F	H18G	1.1113	110.49	119.86	110.41	1.1115
IC	H18F	C17F	*C18F	H18H	1.1113	110.49	-120.11	110.63	1.1112

PATCH FIRST NONE LAST NONE

RESI OSM 0.00 ! sphingomyelin d18:1/18:1 (w/oleic acid)
 GROUP !
 ATOM N NTL -0.60 !
 ATOM C15 CTL5 -0.35 !
 ATOM H15A HL 0.25 ! H15B
 ATOM H15B HL 0.25 ! |
 ATOM H15C HL 0.25 ! H15A-C15-H15C
 ATOM C14 CTL5 -0.35 ! |
 ATOM H14A HL 0.25 ! H13C H14C
 ATOM H14B HL 0.25 ! | |
 ATOM H14C HL 0.25 ! H13A-C13---N--C14-H14A (+)
 ATOM C13 CTL5 -0.35 ! |
 ATOM H13A HL 0.25 ! H13B H14B
 ATOM H13B HL 0.25 ! |
 ATOM H13C HL 0.25 ! H12A-C12-H12B
 ATOM C12 CTL2 -0.10 !
 ATOM H12A HL 0.25 !
 ATOM H12B HL 0.25 ! H11A-C11-H11B

GROUP !
 ATOM C11 CTL2 -0.08 !
 ATOM H11A HAL2 0.09 !
 ATOM H11B HAL2 0.09 !
 ATOM P PL 1.50 !
 ATOM O13 O2L -0.78 !
 ATOM O14 O2L -0.78 !
 ATOM O11 OSLP -0.57 !
 ATOM O12 OSLP -0.57 !
 ATOM C1S CTL2 -0.08 !
 ATOM H1S HAL2 0.09 !
 ATOM H1T HAL2 0.09 !
 GROUP !
 ATOM NF NHL -0.70 !
 ATOM HNF H 0.35 !
 ATOM C2S CTL1 0.30 !
 ATOM H2S HAL1 0.05 !
 GROUP !
 ATOM C3S CTO1 0.50 !
 ATOM H3S HAL1 -0.11 !
 ATOM O3 OHL -0.69 !
 ATOM HO3 HOL 0.30 !
 GROUP !
 ATOM C4S CEL3 -0.15 !
 ATOM H4S HEL1 0.15 !
 GROUP !
 ATOM C5S CEL3 -0.15 !
 ATOM H5S HEL1 0.15 !
 GROUP !
 ATOM C6S CTL2 -0.18 !
 ATOM H6S HAL2 0.09 !
 ATOM H6T HAL2 0.09 !
 GROUP !
 ATOM C7S CTL2 -0.18 !
 ATOM H7S HAL2 0.09 !
 ATOM H7T HAL2 0.09 !
 GROUP !
 ATOM C8S CTL2 -0.18 !
 ATOM H8S HAL2 0.09 !
 ATOM H8T HAL2 0.09 !
 GROUP !
 ATOM C9S CTL2 -0.18 !
 ATOM H9S HAL2 0.09 !
 ATOM H9T HAL2 0.09 !
 GROUP !
 ATOM C10S CTL2 -0.18 !
 ATOM H10S HAL2 0.09 !
 ATOM H10T HAL2 0.09 !
 GROUP !
 ATOM C11S CTL2 -0.18 !
 ATOM H11S HAL2 0.09 !
 ATOM H11T HAL2 0.09 !
 GROUP !
 ATOM C12S CTL2 -0.18 !
 ATOM H12S HAL2 0.09 !
 ATOM H12T HAL2 0.09 !
 GROUP !
 ATOM C13S CTL2 -0.18 !
 ATOM H13S HAL2 0.09 !
 ATOM H13T HAL2 0.09 !
 GROUP !
 ATOM C14S CTL2 -0.18 !
 ATOM H14S HAL2 0.09 !
 ATOM H14T HAL2 0.09 !
 GROUP !
 ATOM C15S CTL2 -0.18 !
 ATOM H15S HAL2 0.09 !
 ATOM H15T HAL2 0.09 !
 GROUP !
 ATOM C16S CTL2 -0.18 !
 ATOM H16S HAL2 0.09 !

ATOM H16T HAL2 0.09 !
 GROUP
 ATOM C17S CTL2 -0.18 !
 ATOM H17S HAL2 0.09 !
 ATOM H17T HAL2 0.09 !
 GROUP !
 ATOM C18S CTL3 -0.27 !
 ATOM H18S HAL3 0.09 !
 ATOM H18T HAL3 0.09 !
 ATOM H18U HAL3 0.09 !
 GROUP !FATTY ACID CHAIN
 ATOM C1F C 0.55 !
 ATOM O F -0.60 !
 ATOM C2F CTL2 -0.07 !
 ATOM H2F HAL2 0.06 !
 ATOM H2G HAL2 0.06 !
 GROUP !
 ATOM C3F CTL2 -0.18 !
 ATOM H3F HAL2 0.09 !
 ATOM H3G HAL2 0.09 !
 GROUP !
 ATOM C4F CTL2 -0.18 !
 ATOM H4F HAL2 0.09 !
 ATOM H4G HAL2 0.09 !
 GROUP !
 ATOM C5F CTL2 -0.18 !
 ATOM H5F HAL2 0.09 !
 ATOM H5G HAL2 0.09 !
 GROUP !
 ATOM C6F CTL2 -0.18 !
 ATOM H6F HAL2 0.09 !
 ATOM H6G HAL2 0.09 !
 GROUP !
 ATOM C7F CTL2 -0.18 !
 ATOM H7F HAL2 0.09 !
 ATOM H7G HAL2 0.09 !
 GROUP !
 ATOM C8F CTL2 -0.18 !
 ATOM H8F HAL2 0.09 !
 ATOM H8G HAL2 0.09 !
 GROUP !
 ATOM C9F CEL1 -0.15 !
 ATOM H9F HEL1 0.15 !
 GROUP !
 ATOM C10F CEL1 -0.15 !
 ATOM H10F HEL1 0.15 !
 GROUP !
 ATOM C11F CTL2 -0.18 !
 ATOM H11F HAL2 0.09 !
 ATOM H11G HAL2 0.09 !
 GROUP !
 ATOM C12F CTL2 -0.18 !
 ATOM H12F HAL2 0.09 !
 ATOM H12G HAL2 0.09 !
 GROUP !
 ATOM C13F CTL2 -0.18 !
 ATOM H13F HAL2 0.09 !
 ATOM H13G HAL2 0.09 !
 GROUP !
 ATOM C14F CTL2 -0.18 !
 ATOM H14F HAL2 0.09 !
 ATOM H14G HAL2 0.09 !
 GROUP !
 ATOM C15F CTL2 -0.18 !
 ATOM H15F HAL2 0.09 !
 ATOM H15G HAL2 0.09 !
 GROUP !
 ATOM C16F CTL2 -0.18 !
 ATOM H16F HAL2 0.09 !
 ATOM H16G HAL2 0.09 !
 GROUP !

ATOM C17F CTL2 -0.18 !
 ATOM H17F HAL2 0.09 !
 ATOM H17G HAL2 0.09 !
 GROUP !
 ATOM C18F CTL3 -0.27 !
 ATOM H18F HAL3 0.09 !
 ATOM H18G HAL3 0.09 !
 ATOM H18H HAL3 0.09 !

!SPHINGOSINE CHAIN

BOND C18S H18S C18S H18T C18S H18U
 BOND C18S C17S C17S H17S C17S H17T
 BOND C17S C16S C16S H16S C16S H16T
 BOND C16S C15S C15S H15S C15S H15T
 BOND C15S C14S C14S H14S C14S H14T
 BOND C14S C13S C13S H13S C13S H13T
 BOND C13S C12S C12S H12S C12S H12T
 BOND C12S C11S C11S H11S C11S H11T
 BOND C11S C10S C10S H10S C10S H10T
 BOND C10S C9S C9S H9S C9S H9T
 BOND C9S C8S C8S H8S C8S H8T
 BOND C8S C7S C7S H7S C7S H7T
 BOND C7S C6S C6S H6S C6S H6T
 BOND C6S C5S C5S H5S C4S H4S
 BOND C4S C3S C3S H3S C3S O3
 BOND O3 HO3
 BOND C3S C2S C2S H2S
 BOND C2S C1S C1S H1S C1S H1T
 BOND C1S O11

!PHOSLPHOCHOLINE

BOND O11 P O12 P O13 P O14 P
 BOND O12 C11 C11 H11A C11 H11B
 BOND C11 C12 C12 H12A C12 H12B
 BOND C12 N N C13 N C14 N C15
 BOND C13 H13A C13 H13B C13 H13C
 BOND C14 H14A C14 H14B C14 H14C
 BOND C15 H15A C15 H15B C15 H15C

!FATTY ACID CHAIN

BOND C2S NF NF HNF NF C1F
 BOND C1F C2F C2F H2F C2F H2G
 BOND C2F C3F C3F H3F C3F H3G
 BOND C3F C4F C4F H4F C4F H4G
 BOND C4F C5F C5F H5F C5F H5G
 BOND C5F C6F C6F H6F C6F H6G
 BOND C6F C7F C7F H7F C7F H7G
 BOND C7F C8F C8F H8F C8F H8G
 BOND C8F C9F C9F H9F
 BOND C10F H10F
 BOND C10F C11F C11F H11F C11F H11G
 BOND C11F C12F C12F H12F C12F H12G
 BOND C12F C13F C13F H13F C13F H13G
 BOND C13F C14F C14F H14F C14F H14G
 BOND C14F C15F C15F H15F C15F H15G
 BOND C15F C16F C16F H16F C16F H16G
 BOND C16F C17F C17F H17F C17F H17G
 BOND C17F C18F C18F H18F C18F H18G
 BOND C18F H18H

DOUBLE C4S C5S

DOUBLE C9F C10F

DOUBLE C1F OF

IMPR C1F C2F NF OF NF C1F C2S HNF

IC	C18S	C17S	C16S	C15S	1.5309	113.23	180.00	112.64	1.5346
IC	C17S	C16S	C15S	C14S	1.5337	112.64	180.00	112.58	1.5345
IC	C16S	C15S	C14S	C13S	1.5346	112.58	180.00	112.58	1.5345
IC	C15S	C14S	C13S	C12S	1.5345	112.58	-179.99	112.58	1.5345
IC	C14S	C13S	C12S	C11S	1.5345	112.58	179.97	112.58	1.5344
IC	C13S	C12S	C11S	C10S	1.5345	112.58	-179.96	112.61	1.5345
IC	C12S	C11S	C10S	C9S	1.5344	112.61	179.97	112.54	1.5344
IC	C11S	C10S	C9S	C8S	1.5345	112.54	-179.87	112.66	1.5345

IC	C10S	C9S	C8S	C7S	1.5344	112.66	-179.89	112.50	1.5354
IC	C9S	C8S	C7S	C6S	1.5345	112.50	-179.79	112.41	1.5391
IC	C8S	C7S	C6S	C5S	1.5354	112.41	-179.61	111.46	1.5077
IC	C7S	C6S	C5S	C4S	1.5391	111.46	-117.13	124.77	1.3444
IC	C6S	C5S	C4S	C3S	1.5077	124.77	179.01	124.28	1.5009
IC	C5S	C4S	C3S	C2S	1.3444	124.28	-84.24	110.14	1.5343
IC	C5S	C4S	C3S	O3	1.3444	124.28	156.07	107.64	1.4235
IC	C4S	C3S	C2S	C1S	1.5009	110.14	-70.50	114.25	1.5586
IC	C13	C12	*N	C14	1.4967	109.68	116.91	107.30	1.5040
IC	C13	C12	*N	C15	1.4967	109.68	-123.42	110.88	1.4938
IC	C12	N	C15	H15A	1.5212	110.88	-177.15	111.31	1.0816
IC	H15A	N	*C15	H15B	1.0816	111.31	118.98	109.44	1.0875
IC	H15A	N	*C15	H15C	1.0816	111.31	-122.11	111.46	1.0818
IC	C12	N	C14	H14A	1.5212	107.30	-178.89	114.23	1.0739
IC	H14A	N	*C14	H14B	1.0739	114.23	123.44	110.21	1.0968
IC	H14A	N	*C14	H14C	1.0739	114.23	-124.43	111.41	1.0948
IC	C12	N	C13	H13A	1.5212	109.68	176.44	111.15	1.0822
IC	H13A	N	*C13	H13B	1.0822	111.15	121.89	111.75	1.0814
IC	H13A	N	*C13	H13C	1.0822	111.15	-118.50	109.57	1.0892
IC	C13	N	C12	C11	1.4967	109.68	-178.83	116.30	1.5392
IC	C11	N	*C12	H12A	1.5392	116.30	117.04	108.30	1.0985
IC	C11	N	*C12	H12B	1.5392	116.30	-126.61	112.36	1.0787
IC	N	C12	C11	O12	1.5212	116.30	126.79	107.85	1.4239
IC	O12	C12	*C11	H11A	1.4239	107.85	118.10	109.30	1.1131
IC	O12	C12	*C11	H11B	1.4239	107.85	-123.12	113.16	1.1140
IC	C12	C11	O12	P	1.5392	107.85	-68.29	118.09	1.5886
IC	C11	O12	P	O11	1.4239	118.09	-167.58	104.18	1.5725
IC	O11	O12	*P	O13	1.5725	104.18	116.57	107.05	1.4823
IC	O11	O12	*P	O14	1.5725	104.18	-118.73	106.98	1.4805
IC	O12	P	O11	C1S	1.5886	104.18	51.85	121.69	1.4327
IC	P	O11	C1S	C2S	1.5725	121.69	103.66	111.21	1.5586
IC	C2S	O11	*C1S	H1S	1.5586	111.21	118.79	108.35	1.1121
IC	C2S	O11	*C1S	H1T	1.5586	111.21	-123.25	112.85	1.1152
IC	O11	C1S	C2S	C3S	1.4327	111.21	175.67	114.25	1.5343
IC	C3S	C1S	*C2S	NF	1.5343	114.25	120.00	111.52	1.4406
IC	C3S	C1S	*C2S	H2S	1.5343	114.25	-120.00	105.03	1.1194
IC	C1S	C2S	NF	C1F	1.5586	111.52	160.94	124.85	1.3364
IC	C1F	C2S	*NF	HNF	1.3364	124.85	-157.81	113.39	1.0062
IC	C1S	C2S	C3S	C4S	1.5586	114.25	-70.50	110.14	1.5009
IC	C4S	C2S	*C3S	O3	1.5009	110.14	118.36	109.80	1.4235
IC	C4S	C2S	*C3S	H3S	1.5009	110.14	-121.14	110.10	1.1140
IC	C2S	C3S	O3	H03	1.5343	109.80	46.50	104.54	0.9610
IC	C2S	C3S	C4S	C5S	1.5343	110.14	-84.24	124.28	1.3444
IC	C5S	C3S	*C4S	H4S	1.3444	124.28	178.79	115.93	1.1003
IC	C3S	C4S	C5S	C6S	1.5009	124.28	179.01	124.77	1.5077
IC	C6S	C4S	*C5S	H5S	1.5077	124.77	-178.18	119.00	1.1029
IC	C4S	C5S	C6S	C7S	1.3444	124.77	-117.13	111.46	1.5391
IC	C7S	C5S	*C6S	H6S	1.5391	111.46	120.98	111.41	1.1142
IC	C7S	C5S	*C6S	H6T	1.5391	111.46	-120.60	109.91	1.1125
IC	C5S	C6S	C7S	C8S	1.5077	111.46	-179.61	112.41	1.5354
IC	C8S	C6S	*C7S	H7S	1.5354	112.41	121.26	109.74	1.1138
IC	C8S	C6S	*C7S	H7T	1.5354	112.41	-120.99	109.40	1.1147
IC	C6S	C7S	C8S	C9S	1.5391	112.41	-179.79	112.50	1.5345
IC	C9S	C7S	*C8S	H8S	1.5345	112.50	121.20	108.97	1.1132
IC	C9S	C7S	*C8S	H8T	1.5345	112.50	-121.38	109.07	1.1129
IC	C7S	C8S	C9S	C10S	1.5354	112.50	-179.89	112.66	1.5344
IC	C10S	C8S	*C9S	H9S	1.5344	112.66	121.49	109.05	1.1133
IC	C10S	C8S	*C9S	H9T	1.5344	112.66	-121.38	108.96	1.1136
IC	C8S	C9S	C10S	C11S	1.5345	112.66	-179.87	112.54	1.5345
IC	C11S	C9S	*C10S	H10S	1.5345	112.54	121.27	109.07	1.1133
IC	C11S	C9S	*C10S	H10T	1.5345	112.54	-121.34	109.11	1.1132
IC	C9S	C10S	C11S	C12S	1.5344	112.54	179.97	112.61	1.5344
IC	C12S	C10S	*C11S	H11S	1.5344	112.61	121.36	109.09	1.1133
IC	C12S	C10S	*C11S	H11T	1.5344	112.61	-121.33	109.07	1.1134
IC	C10S	C11S	C12S	C13S	1.5345	112.61	-179.96	112.58	1.5345
IC	C13S	C11S	*C12S	H12S	1.5345	112.58	121.29	109.12	1.1133
IC	C13S	C11S	*C12S	H12T	1.5345	112.58	-121.31	109.15	1.1133
IC	C11S	C12S	C13S	C14S	1.5344	112.58	179.97	112.58	1.5345
IC	C14S	C12S	*C13S	H13S	1.5345	112.58	121.30	109.13	1.1133
IC	C14S	C12S	*C13S	H13T	1.5345	112.58	-121.30	109.13	1.1133
IC	C12S	C13S	C14S	C15S	1.5345	112.58	-179.99	112.58	1.5345

IC	C15S	C13S	*C14S	H14S	1.5345	112.58	121.30	109.12	1.1133
IC	C15S	C13S	*C14S	H14T	1.5345	112.58	-121.31	109.12	1.1133
IC	C13S	C14S	C15S	C16S	1.5345	112.58	180.00	112.58	1.5346
IC	C16S	C14S	*C15S	H15S	1.5346	112.58	121.30	109.11	1.1133
IC	C16S	C14S	*C15S	H15T	1.5346	112.58	-121.30	109.11	1.1133
IC	C14S	C15S	C16S	C17S	1.5345	112.58	180.00	112.64	1.5337
IC	C17S	C15S	*C16S	H16S	1.5337	112.64	121.32	109.14	1.1133
IC	C17S	C15S	*C16S	H16T	1.5337	112.64	-121.32	109.14	1.1133
IC	C15S	C16S	C17S	C18S	1.5346	112.64	180.00	113.23	1.5309
IC	C18S	C16S	*C17S	H17S	1.5309	113.23	121.67	108.77	1.1142
IC	C18S	C16S	*C17S	H17T	1.5309	113.23	-121.67	108.77	1.1142
IC	C16S	C17S	C18S	H18S	1.5337	113.23	180.00	110.63	1.1113
IC	H18S	C17S	*C18S	H18T	1.1113	110.63	120.07	110.47	1.1114
IC	H18S	C17S	*C18S	H18U	1.1113	110.63	-120.07	110.47	1.1114
IC	C2S	NF	C1F	C2F	1.4406	124.85	-156.83	114.97	1.4964
IC	C2F	NF	*C1F	OF	1.4964	114.97	162.54	122.11	1.2268
IC	NF	C1F	C2F	C3F	1.3364	114.97	-146.57	111.92	1.5432
IC	C3F	C1F	*C2F	H2F	1.5432	111.92	-120.83	107.93	1.1115
IC	H2F	C1F	*C2F	H2G	1.1115	107.93	-117.30	109.04	1.1113
IC	C1F	C2F	C3F	C4F	1.4964	111.92	179.54	112.84	1.5310
IC	C4F	C2F	*C3F	H3F	1.5310	112.84	-121.63	109.37	1.1145
IC	H3F	C2F	*C3F	H3G	1.1145	109.37	-117.07	109.08	1.1157
IC	C2F	C3F	C4F	C5F	1.5434	112.23	174.86	112.58	1.5342
IC	C5F	C3F	*C4F	H4F	1.5342	112.58	-120.18	108.65	1.1140
IC	H4F	C3F	*C4F	H4G	1.1140	108.65	-117.52	109.62	1.1127
IC	C3F	C4F	C5F	C6F	1.5340	112.58	-174.06	112.39	1.5340
IC	C6F	C4F	*C5F	H5F	1.5340	112.39	-122.01	109.38	1.1130
IC	H5F	C4F	*C5F	H5G	1.1130	109.38	-117.34	108.92	1.1134
IC	C4F	C5F	C6F	C7F	1.5342	112.39	178.27	112.75	1.5340
IC	C7F	C5F	*C6F	H6F	1.5340	112.75	-120.93	108.96	1.1137
IC	H6F	C5F	*C6F	H6G	1.1137	108.96	-117.41	109.22	1.1131
IC	C5F	C6F	C7F	C8F	1.5340	112.75	-178.89	112.33	1.5342
IC	C8F	C6F	*C7F	H7F	1.5342	112.33	-121.40	109.26	1.1132
IC	H7F	C6F	*C7F	H7G	1.1132	109.26	-117.45	109.12	1.1130
IC	C6F	C7F	C8F	C9F	1.5340	112.33	179.09	112.77	1.5340
IC	C9F	C7F	*C8F	H8F	1.5340	112.77	-121.23	109.16	1.1134
IC	H8F	C7F	*C8F	H8G	1.1134	109.16	-117.38	109.08	1.1133
IC	C7F	C8F	C9F	C10F	1.5342	112.77	179.02	112.34	1.5343
IC	C10F	C8F	*C9F	H9F	1.5343	112.34	-121.27	109.27	1.1132
IC	C8F	C9F	C10F	C11F	1.5340	112.34	0.00	112.74	1.5340
IC	C11F	C9F	*C10F	H10F	1.5340	112.74	-121.18	109.16	1.1134
IC	C9F	C10F	C11F	C12F	1.5343	112.74	180.00	112.38	1.5343
IC	C12F	C10F	*C11F	H11F	1.5343	112.38	-121.39	109.34	1.1131
IC	H11F	C10F	*C11F	H11G	1.1131	109.34	-117.43	108.98	1.1132
IC	C10F	C11F	C12F	C13F	1.5340	112.38	0.00	112.68	1.5340
IC	C13F	C11F	*C12F	H12F	1.5340	112.68	-121.02	109.05	1.1134
IC	H12F	C11F	*C12F	H12G	1.1134	109.05	-117.34	109.20	1.1131
IC	C11F	C12F	C13F	C14F	1.5343	112.68	-179.93	112.45	1.5345
IC	C14F	C12F	*C13F	H13F	1.5345	112.45	-121.60	109.42	1.1129
IC	H13F	C12F	*C13F	H13G	1.1129	109.42	-117.40	108.85	1.1134
IC	C12F	C13F	C14F	C15F	1.5340	112.45	175.93	112.68	1.5334
IC	C15F	C13F	*C14F	H14F	1.5334	112.68	-121.00	109.06	1.1131
IC	H14F	C13F	*C14F	H14G	1.1131	109.06	-117.24	109.28	1.1130
IC	C13F	C14F	C15F	C16F	1.5345	112.68	-178.23	113.16	1.5306
IC	C16F	C14F	*C15F	H15F	1.5306	113.16	-121.86	108.84	1.1140
IC	H15F	C14F	*C15F	H15G	1.1140	108.84	-116.70	108.76	1.1144
IC	C14F	C15F	C16F	C17F	1.5345	112.68	-178.23	113.16	1.5306
IC	C17F	C15F	*C16F	H16F	1.5306	113.16	-121.86	108.84	1.1140
IC	H16F	C15F	*C16F	H16G	1.1140	108.84	-116.70	108.76	1.1144
IC	C15F	C16F	C17F	C18F	1.5345	112.68	-178.23	113.16	1.5306
IC	C18F	C16F	*C17F	H17F	1.5306	113.16	-121.86	108.84	1.1140
IC	H17F	C16F	*C17F	H17G	1.1140	108.84	-116.70	108.76	1.1144
IC	C16F	C17F	C18F	H18F	1.5334	113.16	-59.91	110.49	1.1113
IC	H18F	C17F	*C18F	H18G	1.1113	110.49	119.86	110.41	1.1115
IC	H18F	C17F	*C18F	H18H	1.1113	110.49	-120.11	110.63	1.1112

PATCH FIRST NONE LAST NONE

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RESI ASM      0.00 ! sphingomyelin d18:1/20:0
GROUP        !
ATOM N      NTL   -0.60 !
ATOM C15    CTL5  -0.35 !

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ATOM H15A HL 0.25 ! H15A-C15-H15C
 ATOM H15B HL 0.25 ! H13C | H14C
 ATOM H15C HL 0.25 ! H13A-C13---N--C14-H14A (+)
 ATOM C14 CTL5 -0.35 ! H13B | H14B
 ATOM H14A HL 0.25 !
 ATOM H14B HL 0.25 !
 ATOM H14C HL 0.25 !
 ATOM C13 CTL5 -0.35 !
 ATOM H13A HL 0.25 !
 ATOM H13B HL 0.25 !
 ATOM H13C HL 0.25 ! H12A-C12-H12B
 ATOM C12 CTL2 -0.10 !
 ATOM H12A HL 0.25 !
 ATOM H12B HL 0.25 ! H11A-C11-H11B
 GROUP !
 ATOM C11 CTL2 -0.08 !
 ATOM H11A HAL2 0.09 !
 ATOM H11B HAL2 0.09 ! (-) O13 O12
 ATOM P PL 1.50 ! \ /
 ATOM O13 O2L -0.78 ! P (+)
 ATOM O14 O2L -0.78 ! / \
 ATOM O11 OSLP -0.57 ! (-) O14 O11
 ATOM O12 OSLP -0.57 !
 ATOM C1S CTL2 -0.08 !
 ATOM H1S HAL2 0.09 ! H1S---C1S---H1T
 ATOM H1T HAL2 0.09 ! H2S---C2S-----NF--HNF
 GROUP !
 ATOM NF NHL -0.70 ! C1F=OF (FATTY ACID CHAIN)
 ATOM HNF H 0.35 !
 ATOM C2S CTL1 0.30 !
 ATOM H2S HAL1 0.05 !
 GROUP !
 ATOM C3S CTO1 0.50 !
 ATOM H3S HAL1 -0.11 !
 ATOM O3 OHL -0.69 !
 ATOM HO3 HOL 0.30 !
 GROUP !
 ATOM C4S CEL3 -0.15 !
 ATOM H4S HEL1 0.15 !
 GROUP !
 ATOM C5S CEL3 -0.15 !
 ATOM H5S HEL1 0.15 !
 GROUP !
 ATOM C6S CTL2 -0.18 !
 ATOM H6S HAL2 0.09 !
 ATOM H6T HAL2 0.09 !
 GROUP !
 ATOM C7S CTL2 -0.18 !
 ATOM H7S HAL2 0.09 !
 ATOM H7T HAL2 0.09 !
 GROUP !
 ATOM C8S CTL2 -0.18 !
 ATOM H8S HAL2 0.09 !
 ATOM H8T HAL2 0.09 !
 GROUP !
 ATOM C9S CTL2 -0.18 !
 ATOM H9S HAL2 0.09 !
 ATOM H9T HAL2 0.09 !
 GROUP !
 ATOM C10S CTL2 -0.18 !
 ATOM H10S HAL2 0.09 !
 ATOM H10T HAL2 0.09 !
 GROUP !
 ATOM C11S CTL2 -0.18 !
 ATOM H11S HAL2 0.09 !
 ATOM H11T HAL2 0.09 !
 GROUP !
 ATOM C12S CTL2 -0.18 !
 ATOM H12S HAL2 0.09 !
 ATOM H12T HAL2 0.09 !
 GROUP !

ATOM C13S CTL2 -0.18 !
 ATOM H13S HAL2 0.09 !
 ATOM H13T HAL2 0.09 !
 GROUP
 ATOM C14S CTL2 -0.18 !
 ATOM H14S HAL2 0.09 !
 ATOM H14T HAL2 0.09 !
 GROUP
 ATOM C15S CTL2 -0.18 !
 ATOM H15S HAL2 0.09 !
 ATOM H15T HAL2 0.09 !
 GROUP
 ATOM C16S CTL2 -0.18 !
 ATOM H16S HAL2 0.09 !
 ATOM H16T HAL2 0.09 !
 GROUP
 ATOM C17S CTL2 -0.18 !
 ATOM H17S HAL2 0.09 !
 ATOM H17T HAL2 0.09 !
 GROUP
 ATOM C18S CTL3 -0.27 !
 ATOM H18S HAL3 0.09 !
 ATOM H18T HAL3 0.09 !
 ATOM H18U HAL3 0.09 !
 GROUP !FATTY ACID CHAIN
 ATOM C1F C 0.55 !
 ATOM O F -0.60 !
 ATOM C2F CTL2 -0.07 !
 ATOM H2F HAL2 0.06 !
 ATOM H2G HAL2 0.06 !
 GROUP !
 ATOM C3F CTL2 -0.18 !
 ATOM H3F HAL2 0.09 !
 ATOM H3G HAL2 0.09 !
 GROUP !
 ATOM C4F CTL2 -0.18 !
 ATOM H4F HAL2 0.09 !
 ATOM H4G HAL2 0.09 !
 GROUP !
 ATOM C5F CTL2 -0.18 !
 ATOM H5F HAL2 0.09 !
 ATOM H5G HAL2 0.09 !
 GROUP !
 ATOM C6F CTL2 -0.18 !
 ATOM H6F HAL2 0.09 !
 ATOM H6G HAL2 0.09 !
 GROUP !
 ATOM C7F CTL2 -0.18 !
 ATOM H7F HAL2 0.09 !
 ATOM H7G HAL2 0.09 !
 GROUP !
 ATOM C8F CTL2 -0.18 !
 ATOM H8F HAL2 0.09 !
 ATOM H8G HAL2 0.09 !
 GROUP !
 ATOM C9F CTL2 -0.18 !
 ATOM H9F HAL2 0.09 !
 ATOM H9G HAL2 0.09 !
 GROUP !
 ATOM C10F CTL2 -0.18 !
 ATOM H10F HAL2 0.09 !
 ATOM H10G HAL2 0.09 !
 GROUP !
 ATOM C11F CTL2 -0.18 !
 ATOM H11F HAL2 0.09 !
 ATOM H11G HAL2 0.09 !
 GROUP !
 ATOM C12F CTL2 -0.18 !
 ATOM H12F HAL2 0.09 !
 ATOM H12G HAL2 0.09 !
 GROUP !

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ATOM C13F CTL2   -0.18 !
ATOM H13F HAL2   0.09 !
ATOM H13G HAL2   0.09 !
GROUP
ATOM C14F CTL2   -0.18 !
ATOM H14F HAL2   0.09 !
ATOM H14G HAL2   0.09 !
GROUP
ATOM C15F CTL2   -0.18 !
ATOM H15F HAL2   0.09 !
ATOM H15G HAL2   0.09 !
GROUP
ATOM C16F CTL2   -0.18 !
ATOM H16F HAL2   0.09 !
ATOM H16G HAL2   0.09 !
GROUP
ATOM C17F CTL2   -0.18 !
ATOM H17F HAL2   0.09 !
ATOM H17G HAL2   0.09 !
GROUP
ATOM C18F CTL2   -0.18 !
ATOM H18F HAL2   0.09 !
ATOM H18G HAL2   0.09 !
GROUP
ATOM C19F CTL2   -0.18 !
ATOM H19F HAL2   0.09 !
ATOM H19G HAL2   0.09 !
GROUP
ATOM C20F CTL3   -0.27 !
ATOM H20F HAL3   0.09 !
ATOM H20G HAL3   0.09 !
ATOM H20H HAL3   0.09 !

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!SPHINGOSINE CHAIN

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BOND C18S H18S C18S H18T C18S H18U
BOND C18S C17S C17S H17S C17S H17T
BOND C17S C16S C16S H16S C16S H16T
BOND C16S C15S C15S H15S C15S H15T
BOND C15S C14S C14S H14S C14S H14T
BOND C14S C13S C13S H13S C13S H13T
BOND C13S C12S C12S H12S C12S H12T
BOND C12S C11S C11S H11S C11S H11T
BOND C11S C10S C10S H10S C10S H10T
BOND C10S C9S C9S H9S C9S H9T
BOND C9S C8S C8S H8S C8S H8T
BOND C8S C7S C7S H7S C7S H7T
BOND C7S C6S C6S H6S C6S H6T
BOND C6S C5S C5S H5S C4S H4S
BOND C4S C3S C3S H3S C3S O3
BOND O3 HO3
BOND C3S C2S C2S H2S
BOND C2S C1S C1S H1S C1S H1T
BOND C1S O11

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!PHOSLPHOCHOLINE

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BOND O11 P O12 P O13 P O14 P
BOND O12 C11 C11 H11A C11 H11B
BOND C11 C12 C12 H12A C12 H12B
BOND C12 N N C13 N C14 N C15
BOND C13 H13A C13 H13B C13 H13C
BOND C14 H14A C14 H14B C14 H14C
BOND C15 H15A C15 H15B C15 H15C

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!FATTY ACID CHAIN

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BOND C2S NF NF HNF NF C1F
BOND C1F C2F C2F H2F C2F H2G
BOND C2F C3F C3F H3F C3F H3G
BOND C3F C4F C4F H4F C4F H4G
BOND C4F C5F C5F H5F C5F H5G
BOND C5F C6F C6F H6F C6F H6G
BOND C6F C7F C7F H7F C7F H7G
BOND C7F C8F C8F H8F C8F H8G
BOND C8F C9F C9F H9F C9F H9G

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BOND C9F C10F C10F H10F C10F H10G
BOND C10F C11F C11F H11F C11F H11G
BOND C11F C12F C12F H12F C12F H12G
BOND C12F C13F C13F H13F C13F H13G
BOND C13F C14F C14F H14F C14F H14G
BOND C14F C15F C15F H15F C15F H15G
BOND C15F C16F C16F H16F C16F H16G
BOND C16F C17F C17F H17F C17F H17G
BOND C17F C18F C18F H18F C18F H18G
BOND C18F C19F C19F H19F C19F H19G
BOND C19F C20F C20F H20F C20F H20G
BOND C20F H20H

DOUBLE C4S C5S

DOUBLE C1F OF

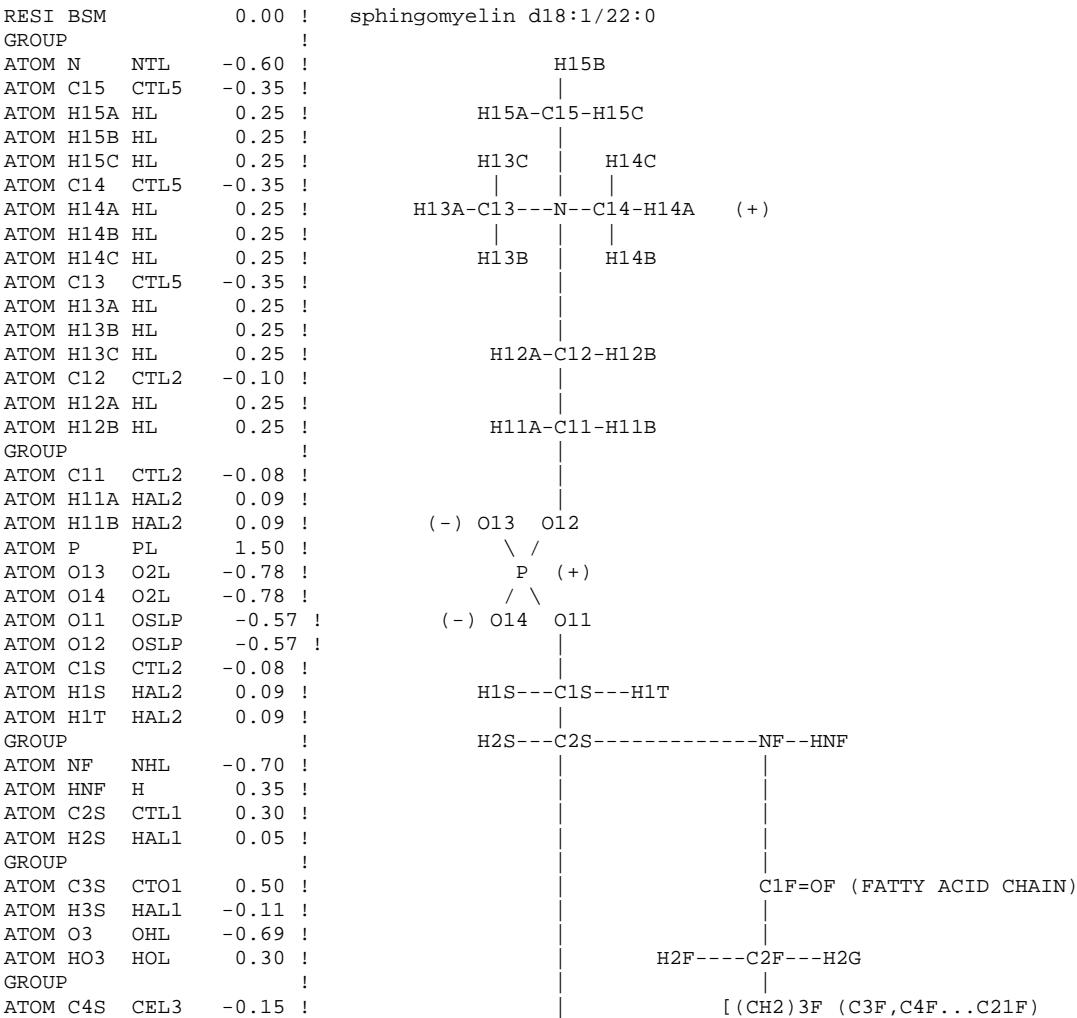
IMPR C1F C2F NF OF NF C1F C2S HNF

IC	C18S	C17S	C16S	C15S	1.5309	113.23	180.00	112.64	1.5346
IC	C17S	C16S	C15S	C14S	1.5337	112.64	180.00	112.58	1.5345
IC	C16S	C15S	C14S	C13S	1.5346	112.58	180.00	112.58	1.5345
IC	C15S	C14S	C13S	C12S	1.5345	112.58	-179.99	112.58	1.5345
IC	C14S	C13S	C12S	C11S	1.5345	112.58	179.97	112.58	1.5344
IC	C13S	C12S	C11S	C10S	1.5345	112.58	-179.96	112.61	1.5345
IC	C12S	C11S	C10S	C9S	1.5344	112.61	179.97	112.54	1.5344
IC	C11S	C10S	C9S	C8S	1.5345	112.54	-179.87	112.66	1.5345
IC	C10S	C9S	C8S	C7S	1.5344	112.66	-179.89	112.50	1.5354
IC	C9S	C8S	C7S	C6S	1.5345	112.50	-179.79	112.41	1.5391
IC	C8S	C7S	C6S	C5S	1.5354	112.41	-179.61	111.46	1.5077
IC	C7S	C6S	C5S	C4S	1.5391	111.46	-117.13	124.77	1.3444
IC	C6S	C5S	C4S	C3S	1.5077	124.77	179.01	124.28	1.5009
IC	C5S	C4S	C3S	C2S	1.3444	124.28	-84.24	110.14	1.5343
IC	C5S	C4S	C3S	O3	1.3444	124.28	156.07	107.64	1.4235
IC	C4S	C3S	C2S	C1S	1.5009	110.14	-70.50	114.25	1.5586
IC	C13	C12	*N	C14	1.4967	109.68	116.91	107.30	1.5040
IC	C13	C12	*N	C15	1.4967	109.68	-123.42	110.88	1.4938
IC	C12	N	C15	H15A	1.5212	110.88	-177.15	111.31	1.0816
IC	H15A	N	*C15	H15B	1.0816	111.31	118.98	109.44	1.0875
IC	H15A	N	*C15	H15C	1.0816	111.31	-122.11	111.46	1.0818
IC	C12	N	C14	H14A	1.5212	107.30	-178.89	114.23	1.0739
IC	H14A	N	*C14	H14B	1.0739	114.23	123.44	110.21	1.0968
IC	H14A	N	*C14	H14C	1.0739	114.23	-124.43	111.41	1.0948
IC	C12	N	C13	H13A	1.5212	109.68	176.44	111.15	1.0822
IC	H13A	N	*C13	H13B	1.0822	111.15	121.89	111.75	1.0814
IC	H13A	N	*C13	H13C	1.0822	111.15	-118.50	109.57	1.0892
IC	C13	N	C12	C11	1.4967	109.68	-178.83	116.30	1.5392
IC	C11	N	*C12	H12A	1.5392	116.30	117.04	108.30	1.0985
IC	C11	N	*C12	H12B	1.5392	116.30	-126.61	112.36	1.0787
IC	N	C12	C11	O12	1.5212	116.30	126.79	107.85	1.4239
IC	O12	C12	*C11	H11A	1.4239	107.85	118.10	109.30	1.1131
IC	O12	C12	*C11	H11B	1.4239	107.85	-123.12	113.16	1.1140
IC	C12	C11	O12	P	1.5392	107.85	-68.29	118.09	1.5886
IC	C11	O12	P	O11	1.4239	118.09	-167.58	104.18	1.5725
IC	O11	O12	*P	O13	1.5725	104.18	116.57	107.05	1.4823
IC	O11	O12	*P	O14	1.5725	104.18	-118.73	106.98	1.4805
IC	O12	P	O11	C1S	1.5886	104.18	51.85	121.69	1.4327
IC	P	O11	C1S	C2S	1.5725	121.69	103.66	111.21	1.5586
IC	C2S	O11	*C1S	H1S	1.5586	111.21	118.79	108.35	1.1121
IC	C2S	O11	*C1S	H1T	1.5586	111.21	-123.25	112.85	1.1152
IC	O11	C1S	C2S	C3S	1.4327	111.21	175.67	114.25	1.5343
IC	C3S	C1S	*C2S	NF	1.5343	114.25	120.00	111.52	1.4406
IC	C3S	C1S	*C2S	H2S	1.5343	114.25	-120.00	105.03	1.1194
IC	C1S	C2S	NF	C1F	1.5586	111.52	160.94	124.85	1.3364
IC	C1F	C2S	*NF	HNF	1.3364	124.85	-157.81	113.39	1.0062
IC	C1S	C2S	C3S	C4S	1.5586	114.25	-70.50	110.14	1.5009
IC	C4S	C2S	*C3S	O3	1.5009	110.14	118.36	109.80	1.4235
IC	C4S	C2S	*C3S	H3S	1.5009	110.14	-121.14	110.10	1.1140
IC	C2S	C3S	O3	H03	1.5343	109.80	46.50	104.54	0.9610
IC	C2S	C3S	C4S	C5S	1.5343	110.14	-84.24	124.28	1.3444
IC	C5S	C3S	*C4S	H4S	1.3444	124.28	178.79	115.93	1.1003
IC	C3S	C4S	C5S	C6S	1.5009	124.28	179.01	124.77	1.5077
IC	C6S	C4S	*C5S	H5S	1.5077	124.77	-178.18	119.00	1.1029

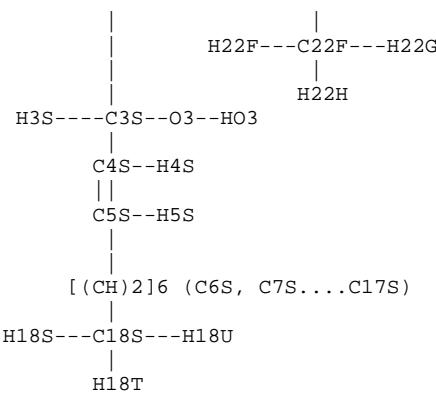
IC	C4S	C5S	C6S	C7S	1.3444	124.77	-117.13	111.46	1.5391
IC	C7S	C5S	*C6S	H6S	1.5391	111.46	120.98	111.41	1.1142
IC	C7S	C5S	*C6S	H6T	1.5391	111.46	-120.60	109.91	1.1125
IC	C5S	C6S	C7S	C8S	1.5077	111.46	-179.61	112.41	1.5354
IC	C8S	C6S	*C7S	H7S	1.5354	112.41	121.26	109.74	1.1138
IC	C8S	C6S	*C7S	H7T	1.5354	112.41	-120.99	109.40	1.1147
IC	C6S	C7S	C8S	C9S	1.5391	112.41	-179.79	112.50	1.5345
IC	C9S	C7S	*C8S	H8S	1.5345	112.50	121.20	108.97	1.1132
IC	C9S	C7S	*C8S	H8T	1.5345	112.50	-121.38	109.07	1.1129
IC	C7S	C8S	C9S	C10S	1.5354	112.50	-179.89	112.66	1.5344
IC	C10S	C8S	*C9S	H9S	1.5344	112.66	121.49	109.05	1.1133
IC	C10S	C8S	*C9S	H9T	1.5344	112.66	-121.38	108.96	1.1136
IC	C8S	C9S	C10S	C11S	1.5345	112.66	-179.87	112.54	1.5345
IC	C11S	C9S	*C10S	H10S	1.5345	112.54	121.27	109.07	1.1133
IC	C11S	C9S	*C10S	H10T	1.5345	112.54	-121.34	109.11	1.1132
IC	C9S	C10S	C11S	C12S	1.5344	112.54	179.97	112.61	1.5344
IC	C12S	C10S	*C11S	H11S	1.5344	112.61	121.36	109.09	1.1133
IC	C12S	C10S	*C11S	H11T	1.5344	112.61	-121.33	109.07	1.1134
IC	C10S	C11S	C12S	C13S	1.5345	112.61	-179.96	112.58	1.5345
IC	C13S	C11S	*C12S	H12S	1.5345	112.58	121.29	109.12	1.1133
IC	C13S	C11S	*C12S	H12T	1.5345	112.58	-121.31	109.15	1.1133
IC	C11S	C12S	C13S	C14S	1.5344	112.58	179.97	112.58	1.5345
IC	C14S	C12S	*C13S	H13S	1.5345	112.58	121.30	109.13	1.1133
IC	C14S	C12S	*C13S	H13T	1.5345	112.58	-121.30	109.13	1.1133
IC	C12S	C13S	C14S	C15S	1.5345	112.58	-179.99	112.58	1.5345
IC	C15S	C13S	*C14S	H14S	1.5345	112.58	121.30	109.12	1.1133
IC	C15S	C13S	*C14S	H14T	1.5345	112.58	-121.31	109.12	1.1133
IC	C13S	C14S	C15S	C16S	1.5345	112.58	180.00	112.58	1.5346
IC	C16S	C14S	*C15S	H15S	1.5346	112.58	121.30	109.11	1.1133
IC	C16S	C14S	*C15S	H15T	1.5346	112.58	-121.30	109.11	1.1133
IC	C14S	C15S	C16S	C17S	1.5345	112.58	180.00	112.64	1.5337
IC	C17S	C15S	*C16S	H16S	1.5337	112.64	121.32	109.14	1.1133
IC	C17S	C15S	*C16S	H16T	1.5337	112.64	-121.32	109.14	1.1133
IC	C15S	C16S	C17S	C18S	1.5346	112.64	180.00	113.23	1.5309
IC	C18S	C16S	*C17S	H17S	1.5309	113.23	121.67	108.77	1.1142
IC	C18S	C16S	*C17S	H17T	1.5309	113.23	-121.67	108.77	1.1142
IC	C16S	C17S	C18S	H18S	1.5337	113.23	180.00	110.63	1.1113
IC	H18S	C17S	*C18S	H18T	1.1113	110.63	120.07	110.47	1.1114
IC	H18S	C17S	*C18S	H18U	1.1113	110.63	-120.07	110.47	1.1114
IC	C2S	NF	C1F	C2F	1.4406	124.85	-156.83	114.97	1.4964
IC	C2F	NF	*C1F	OF	1.4964	114.97	162.54	122.11	1.2268
IC	NF	C1F	C2F	C3F	1.3364	114.97	-146.57	111.92	1.5432
IC	C3F	C1F	*C2F	H2F	1.5432	111.92	-120.83	107.93	1.1115
IC	H2F	C1F	*C2F	H2G	1.1115	107.93	-117.30	109.04	1.1113
IC	C1F	C2F	C3F	C4F	1.4964	111.92	179.54	112.84	1.5310
IC	C4F	C2F	*C3F	H3F	1.5310	112.84	-121.63	109.37	1.1145
IC	H3F	C2F	*C3F	H3G	1.1145	109.37	-117.07	109.08	1.1157
IC	C2F	C3F	C4F	C5F	1.5434	112.23	174.86	112.58	1.5342
IC	C5F	C3F	*C4F	H4F	1.5342	112.58	-120.18	108.65	1.1140
IC	H4F	C3F	*C4F	H4G	1.1140	108.65	-117.52	109.62	1.1127
IC	C3F	C4F	C5F	C6F	1.5340	112.58	-174.06	112.39	1.5340
IC	C6F	C4F	*C5F	H5F	1.5340	112.39	-122.01	109.38	1.1130
IC	H5F	C4F	*C5F	H5G	1.1130	109.38	-117.34	108.92	1.1134
IC	C4F	C5F	C6F	C7F	1.5342	112.39	178.27	112.75	1.5340
IC	C7F	C5F	*C6F	H6F	1.5340	112.75	-120.93	108.96	1.1137
IC	H6F	C5F	*C6F	H6G	1.1137	108.96	-117.41	109.22	1.1131
IC	C5F	C6F	C7F	C8F	1.5340	112.75	-178.89	112.33	1.5342
IC	C8F	C6F	*C7F	H7F	1.5342	112.33	-121.40	109.26	1.1132
IC	H7F	C6F	*C7F	H7G	1.1132	109.26	-117.45	109.12	1.1130
IC	C6F	C7F	C8F	C9F	1.5340	112.33	179.09	112.77	1.5340
IC	C9F	C7F	*C8F	H8F	1.5340	112.77	-121.23	109.16	1.1134
IC	H8F	C7F	*C8F	H8G	1.1134	109.16	-117.38	109.08	1.1133
IC	C7F	C8F	C9F	C10F	1.5342	112.77	179.02	112.34	1.5343
IC	C10F	C8F	*C9F	H9F	1.5343	112.34	-121.27	109.27	1.1132
IC	H9F	C8F	*C9F	H9G	1.1132	109.27	-117.45	109.10	1.1130
IC	C8F	C9F	C10F	C11F	1.5340	112.34	178.16	112.74	1.5340
IC	C11F	C9F	*C10F	H10F	1.5340	112.74	-121.18	109.16	1.1134
IC	H10F	C9F	*C10F	H10G	1.1134	109.16	-117.35	109.10	1.1132
IC	C9F	C10F	C11F	C12F	1.5343	112.74	178.91	112.38	1.5343
IC	C12F	C10F	*C11F	H11F	1.5343	112.38	-121.39	109.34	1.1131
IC	H11F	C10F	*C11F	H11G	1.1131	109.34	-117.43	108.98	1.1132

IC	C10F	C11F	C12F	C13F	1.5340	112.38	176.93	112.68	1.5340
IC	C13F	C11F	*C12F	H12F	1.5340	112.68	-121.02	109.05	1.1134
IC	H12F	C11F	*C12F	H12G	1.1134	109.05	-117.34	109.20	1.1131
IC	C11F	C12F	C13F	C14F	1.5343	112.68	-179.93	112.45	1.5345
IC	C14F	C12F	*C13F	H13F	1.5345	112.45	-121.60	109.42	1.1129
IC	H13F	C12F	*C13F	H13G	1.1129	109.42	-117.40	108.85	1.1134
IC	C12F	C13F	C14F	C15F	1.5340	112.45	175.93	112.68	1.5334
IC	C15F	C13F	*C14F	H14F	1.5334	112.68	-121.00	109.06	1.1131
IC	H14F	C13F	*C14F	H14G	1.1131	109.06	-117.24	109.28	1.1130
IC	C13F	C14F	C15F	C16F	1.5345	112.68	-178.23	113.16	1.5306
IC	C16F	C14F	*C15F	H15F	1.5306	113.16	-121.86	108.84	1.1140
IC	H15F	C14F	*C15F	H15G	1.1140	108.84	-116.70	108.76	1.1144
IC	C14F	C15F	C16F	C17F	1.5345	112.68	-178.23	113.16	1.5306
IC	C17F	C15F	*C16F	H16F	1.5306	113.16	-121.86	108.84	1.1140
IC	H16F	C15F	*C16F	H16G	1.1140	108.84	-116.70	108.76	1.1144
IC	C15F	C16F	C17F	C18F	1.5345	112.68	-178.23	113.16	1.5306
IC	C18F	C16F	*C17F	H17F	1.5306	113.16	-121.86	108.84	1.1140
IC	H17F	C16F	*C17F	H17G	1.1140	108.84	-116.70	108.76	1.1144
IC	C16F	C17F	C18F	C19F	1.5345	112.68	-178.23	113.16	1.5306
IC	C19F	C17F	*C18F	H18F	1.5306	113.16	-121.86	108.84	1.1140
IC	H18F	C17F	*C18F	H18G	1.1140	108.84	-116.70	108.76	1.1144
IC	C17F	C18F	C19F	C20F	1.5345	112.68	-178.23	113.16	1.5306
IC	C20F	C18F	*C19F	H19F	1.5306	113.16	-121.86	108.84	1.1140
IC	H19F	C18F	*C19F	H19G	1.1140	108.84	-116.70	108.76	1.1144
IC	C18F	C19F	C20F	H20F	1.5334	113.16	-59.91	110.49	1.1113
IC	H20F	C19F	*C20F	H20G	1.1113	110.49	119.86	110.41	1.1115
IC	H20F	C19F	*C20F	H20H	1.1113	110.49	-120.11	110.63	1.1112

PATCH FIRST NONE LAST NONE



ATOM H4S HEL1 0.15 !
 GROUP !
 ATOM C5S CEL3 -0.15 !
 ATOM H5S HEL1 0.15 !
 GROUP !
 ATOM C6S CTL2 -0.18 !
 ATOM H6S HAL2 0.09 !
 ATOM H6T HAL2 0.09 !
 GROUP !
 ATOM C7S CTL2 -0.18 !
 ATOM H7S HAL2 0.09 !
 ATOM H7T HAL2 0.09 !
 GROUP !
 ATOM C8S CTL2 -0.18 !
 ATOM H8S HAL2 0.09 !
 ATOM H8T HAL2 0.09 !
 GROUP !
 ATOM C9S CTL2 -0.18 !
 ATOM H9S HAL2 0.09 !
 ATOM H9T HAL2 0.09 !
 GROUP !
 ATOM C10S CTL2 -0.18 !
 ATOM H10S HAL2 0.09 !
 ATOM H10T HAL2 0.09 !
 GROUP
 ATOM C11S CTL2 -0.18 !
 ATOM H11S HAL2 0.09 !
 ATOM H11T HAL2 0.09 !
 GROUP
 ATOM C12S CTL2 -0.18 !
 ATOM H12S HAL2 0.09 !
 ATOM H12T HAL2 0.09 !
 GROUP
 ATOM C13S CTL2 -0.18 !
 ATOM H13S HAL2 0.09 !
 ATOM H13T HAL2 0.09 !
 GROUP
 ATOM C14S CTL2 -0.18 !
 ATOM H14S HAL2 0.09 !
 ATOM H14T HAL2 0.09 !
 GROUP
 ATOM C15S CTL2 -0.18 !
 ATOM H15S HAL2 0.09 !
 ATOM H15T HAL2 0.09 !
 GROUP
 ATOM C16S CTL2 -0.18 !
 ATOM H16S HAL2 0.09 !
 ATOM H16T HAL2 0.09 !
 GROUP
 ATOM C17S CTL2 -0.18 !
 ATOM H17S HAL2 0.09 !
 ATOM H17T HAL2 0.09 !
 GROUP !
 ATOM C18S CTL3 -0.27 !
 ATOM H18S HAL3 0.09 !
 ATOM H18T HAL3 0.09 !
 ATOM H18U HAL3 0.09 !
 GROUP !FATTY ACID CHAIN
 ATOM C1F C 0.55 !
 ATOM O F -0.60 !
 ATOM C2F CTL2 -0.07 !
 ATOM H2F HAL2 0.06 !
 ATOM H2G HAL2 0.06 !
 GROUP !
 ATOM C3F CTL2 -0.18 !
 ATOM H3F HAL2 0.09 !
 ATOM H3G HAL2 0.09 !
 GROUP !
 ATOM C4F CTL2 -0.18 !
 ATOM H4F HAL2 0.09 !
 ATOM H4G HAL2 0.09 !



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GROUP          !
ATOM C5F  CTL2   -0.18 !
ATOM H5F  HAL2    0.09 !
ATOM H5G  HAL2    0.09 !
GROUP          !
ATOM C6F  CTL2   -0.18 !
ATOM H6F  HAL2    0.09 !
ATOM H6G  HAL2    0.09 !
GROUP          !
ATOM C7F  CTL2   -0.18 !
ATOM H7F  HAL2    0.09 !
ATOM H7G  HAL2    0.09 !
GROUP          !
ATOM C8F  CTL2   -0.18 !
ATOM H8F  HAL2    0.09 !
ATOM H8G  HAL2    0.09 !
GROUP          !
ATOM C9F  CTL2   -0.18 !
ATOM H9F  HAL2    0.09 !
ATOM H9G  HAL2    0.09 !
GROUP          !
ATOM C10F CTL2   -0.18 !
ATOM H10F HAL2    0.09 !
ATOM H10G HAL2    0.09 !
GROUP          !
ATOM C11F CTL2   -0.18 !
ATOM H11F HAL2    0.09 !
ATOM H11G HAL2    0.09 !
GROUP          !
ATOM C12F CTL2   -0.18 !
ATOM H12F HAL2    0.09 !
ATOM H12G HAL2    0.09 !
GROUP          !
ATOM C13F CTL2   -0.18 !
ATOM H13F HAL2    0.09 !
ATOM H13G HAL2    0.09 !
GROUP          !
ATOM C14F CTL2   -0.18 !
ATOM H14F HAL2    0.09 !
ATOM H14G HAL2    0.09 !
GROUP          !
ATOM C15F CTL2   -0.18 !
ATOM H15F HAL2    0.09 !
ATOM H15G HAL2    0.09 !
GROUP          !
ATOM C16F CTL2   -0.18 !
ATOM H16F HAL2    0.09 !
ATOM H16G HAL2    0.09 !
GROUP          !
ATOM C17F CTL2   -0.18 !
ATOM H17F HAL2    0.09 !
ATOM H17G HAL2    0.09 !
GROUP          !
ATOM C18F CTL2   -0.18 !
ATOM H18F HAL2    0.09 !
ATOM H18G HAL2    0.09 !
GROUP          !
ATOM C19F CTL2   -0.18 !
ATOM H19F HAL2    0.09 !
ATOM H19G HAL2    0.09 !
GROUP          !
ATOM C20F CTL2   -0.18 !
ATOM H20F HAL2    0.09 !
ATOM H20G HAL2    0.09 !
GROUP          !
ATOM C21F CTL2   -0.18 !
ATOM H21F HAL2    0.09 !
ATOM H21G HAL2    0.09 !
GROUP          !
ATOM C22F CTL3   -0.27 !
ATOM H22F HAL3    0.09 !

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ATOM H22G HAL3 0.09 !
 ATOM H22H HAL3 0.09 !

!SPHINGOSINE CHAIN

BOND C18S H18S C18S H18T C18S H18U
 BOND C18S C17S C17S H17S C17S H17T
 BOND C17S C16S C16S H16S C16S H16T
 BOND C16S C15S C15S H15S C15S H15T
 BOND C15S C14S C14S H14S C14S H14T
 BOND C14S C13S C13S H13S C13S H13T
 BOND C13S C12S C12S H12S C12S H12T
 BOND C12S C11S C11S H11S C11S H11T
 BOND C11S C10S C10S H10S C10S H10T
 BOND C10S C9S C9S H9S C9S H9T
 BOND C9S C8S C8S H8S C8S H8T
 BOND C8S C7S C7S H7S C7S H7T
 BOND C7S C6S C6S H6S C6S H6T
 BOND C6S C5S C5S H5S C4S H4S
 BOND C4S C3S C3S H3S C3S O3
 BOND O3 HO3
 BOND C3S C2S C2S H2S
 BOND C2S C1S C1S H1S C1S H1T
 BOND C1S O11

!PHOSLPHOCHOLINE

BOND O11 P O12 P O13 P O14 P
 BOND O12 C11 C11 H11A C11 H11B
 BOND C11 C12 C12 H12A C12 H12B
 BOND C12 N N C13 N C14 N C15
 BOND C13 H13A C13 H13B C13 H13C
 BOND C14 H14A C14 H14B C14 H14C
 BOND C15 H15A C15 H15B C15 H15C

!FATTY ACID CHAIN

BOND C2S NF NF HNF NF C1F
 BOND C1F C2F C2F H2F C2F H2G
 BOND C2F C3F C3F H3F C3F H3G
 BOND C3F C4F C4F H4F C4F H4G
 BOND C4F C5F C5F H5F C5F H5G
 BOND C5F C6F C6F H6F C6F H6G
 BOND C6F C7F C7F H7F C7F H7G
 BOND C7F C8F C8F H8F C8F H8G
 BOND C8F C9F C9F H9F C9F H9G
 BOND C9F C10F C10F H10F C10F H10G
 BOND C10F C11F C11F H11F C11F H11G
 BOND C11F C12F C12F H12F C12F H12G
 BOND C12F C13F C13F H13F C13F H13G
 BOND C13F C14F C14F H14F C14F H14G
 BOND C14F C15F C15F H15F C15F H15G
 BOND C15F C16F C16F H16F C16F H16G
 BOND C16F C17F C17F H17F C17F H17G
 BOND C17F C18F C18F H18F C18F H18G
 BOND C18F C19F C19F H19F C19F H19G
 BOND C19F C20F C20F H20F C20F H20G
 BOND C20F C21F C21F H21F C21F H21G
 BOND C21F C22F C22F H22F C22F H22G
 BOND C22F H22H

DOUBLE C4S C5S

DOUBLE C1F OF

IMPR C1F C2F NF OF NF C1F C2S HNF

IC	C18S	C17S	C16S	C15S	1.5309	113.23	180.00	112.64	1.5346
IC	C17S	C16S	C15S	C14S	1.5337	112.64	180.00	112.58	1.5345
IC	C16S	C15S	C14S	C13S	1.5346	112.58	180.00	112.58	1.5345
IC	C15S	C14S	C13S	C12S	1.5345	112.58	-179.99	112.58	1.5345
IC	C14S	C13S	C12S	C11S	1.5345	112.58	179.97	112.58	1.5344
IC	C13S	C12S	C11S	C10S	1.5345	112.58	-179.96	112.61	1.5345
IC	C12S	C11S	C10S	C9S	1.5344	112.61	179.97	112.54	1.5344
IC	C11S	C10S	C9S	C8S	1.5345	112.54	-179.87	112.66	1.5345
IC	C10S	C9S	C8S	C7S	1.5344	112.66	-179.89	112.50	1.5354
IC	C9S	C8S	C7S	C6S	1.5345	112.50	-179.79	112.41	1.5391
IC	C8S	C7S	C6S	C5S	1.5354	112.41	-179.61	111.46	1.5077

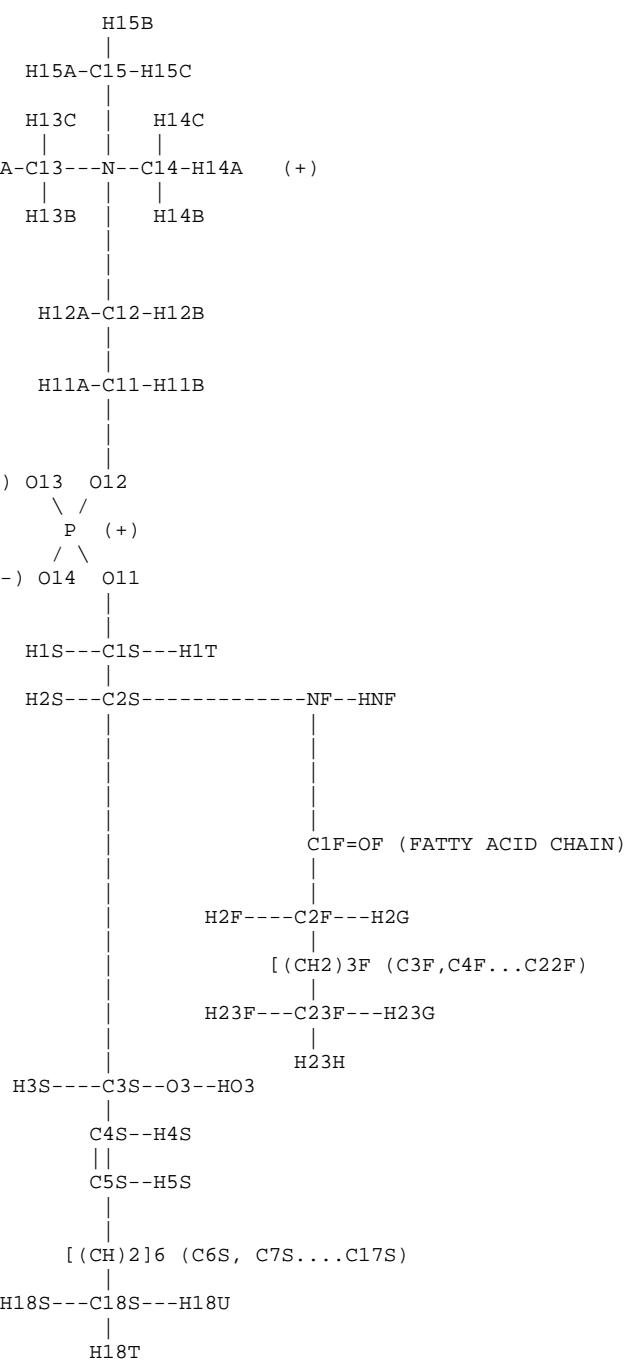
IC	C7S	C6S	C5S	C4S	1.5391	111.46	-117.13	124.77	1.3444
IC	C6S	C5S	C4S	C3S	1.5077	124.77	179.01	124.28	1.5009
IC	C5S	C4S	C3S	C2S	1.3444	124.28	-84.24	110.14	1.5343
IC	C5S	C4S	C3S	O3	1.3444	124.28	156.07	107.64	1.4235
IC	C4S	C3S	C2S	C1S	1.5009	110.14	-70.50	114.25	1.5586
IC	C13	C12	*N	C14	1.4967	109.68	116.91	107.30	1.5040
IC	C13	C12	*N	C15	1.4967	109.68	-123.42	110.88	1.4938
IC	C12	N	C15	H15A	1.5212	110.88	-177.15	111.31	1.0816
IC	H15A	N	*C15	H15B	1.0816	111.31	118.98	109.44	1.0875
IC	H15A	N	*C15	H15C	1.0816	111.31	-122.11	111.46	1.0818
IC	C12	N	C14	H14A	1.5212	107.30	-178.89	114.23	1.0739
IC	H14A	N	*C14	H14B	1.0739	114.23	123.44	110.21	1.0968
IC	H14A	N	*C14	H14C	1.0739	114.23	-124.43	111.41	1.0948
IC	C12	N	C13	H13A	1.5212	109.68	176.44	111.15	1.0822
IC	H13A	N	*C13	H13B	1.0822	111.15	121.89	111.75	1.0814
IC	H13A	N	*C13	H13C	1.0822	111.15	-118.50	109.57	1.0892
IC	C13	N	C12	C11	1.4967	109.68	-178.83	116.30	1.5392
IC	C11	N	*C12	H12A	1.5392	116.30	117.04	108.30	1.0985
IC	C11	N	*C12	H12B	1.5392	116.30	-126.61	112.36	1.0787
IC	N	C12	C11	O12	1.5212	116.30	126.79	107.85	1.4239
IC	O12	C12	*C11	H11A	1.4239	107.85	118.10	109.30	1.1131
IC	O12	C12	*C11	H11B	1.4239	107.85	-123.12	113.16	1.1140
IC	C12	C11	O12	P	1.5392	107.85	-68.29	118.09	1.5886
IC	C11	O12	P	O11	1.4239	118.09	-167.58	104.18	1.5725
IC	O11	O12	*P	O13	1.5725	104.18	116.57	107.05	1.4823
IC	O11	O12	*P	O14	1.5725	104.18	-118.73	106.98	1.4805
IC	O12	P	O11	C1S	1.5886	104.18	51.85	121.69	1.4327
IC	P	O11	C1S	C2S	1.5725	121.69	103.66	111.21	1.5586
IC	C2S	O11	*C1S	H1S	1.5586	111.21	118.79	108.35	1.1121
IC	C2S	O11	*C1S	H1T	1.5586	111.21	-123.25	112.85	1.1152
IC	O11	C1S	C2S	C3S	1.4327	111.21	175.67	114.25	1.5343
IC	C3S	C1S	*C2S	NF	1.5343	114.25	120.00	111.52	1.4406
IC	C3S	C1S	*C2S	H2S	1.5343	114.25	-120.00	105.03	1.1194
IC	C1S	C2S	NF	C1F	1.5586	111.52	160.94	124.85	1.3364
IC	C1F	C2S	*NF	HNF	1.3364	124.85	-157.81	113.39	1.0062
IC	C1S	C2S	C3S	C4S	1.5586	114.25	-70.50	110.14	1.5009
IC	C4S	C2S	*C3S	O3	1.5009	110.14	118.36	109.80	1.4235
IC	C4S	C2S	*C3S	H3S	1.5009	110.14	-121.14	110.10	1.1140
IC	C2S	C3S	O3	H03	1.5343	109.80	46.50	104.54	0.9610
IC	C2S	C3S	C4S	C5S	1.5343	110.14	-84.24	124.28	1.3444
IC	C5S	C3S	*C4S	H4S	1.3444	124.28	178.79	115.93	1.1003
IC	C3S	C4S	C5S	C6S	1.5009	124.28	179.01	124.77	1.5077
IC	C6S	C4S	*C5S	H5S	1.5077	124.77	-178.18	119.00	1.1029
IC	C4S	C5S	C6S	C7S	1.3444	124.77	-117.13	111.46	1.5391
IC	C7S	C5S	*C6S	H6S	1.5391	111.46	120.98	111.41	1.1142
IC	C7S	C5S	*C6S	H6T	1.5391	111.46	-120.60	109.91	1.1125
IC	C5S	C6S	C7S	C8S	1.5077	111.46	-179.61	112.41	1.5354
IC	C8S	C6S	*C7S	H7S	1.5354	112.41	121.26	109.74	1.1138
IC	C8S	C6S	*C7S	H7T	1.5354	112.41	-120.99	109.40	1.1147
IC	C6S	C7S	C8S	C9S	1.5391	112.41	-179.79	112.50	1.5345
IC	C9S	C7S	*C8S	H8S	1.5345	112.50	121.20	108.97	1.1132
IC	C9S	C7S	*C8S	H8T	1.5345	112.50	-121.38	109.07	1.1129
IC	C7S	C8S	C9S	C10S	1.5354	112.50	-179.89	112.66	1.5344
IC	C10S	C8S	*C9S	H9S	1.5344	112.66	121.49	109.05	1.1133
IC	C10S	C8S	*C9S	H9T	1.5344	112.66	-121.38	108.96	1.1136
IC	C8S	C9S	C10S	C11S	1.5345	112.66	-179.87	112.54	1.5345
IC	C11S	C9S	*C10S	H10S	1.5345	112.54	121.27	109.07	1.1133
IC	C11S	C9S	*C10S	H10T	1.5345	112.54	-121.34	109.11	1.1132
IC	C9S	C10S	C11S	C12S	1.5344	112.54	179.97	112.61	1.5344
IC	C12S	C10S	*C11S	H11S	1.5344	112.61	121.36	109.09	1.1133
IC	C12S	C10S	*C11S	H11T	1.5344	112.61	-121.33	109.07	1.1134
IC	C10S	C11S	C12S	C13S	1.5345	112.61	-179.96	112.58	1.5345
IC	C13S	C11S	*C12S	H12S	1.5345	112.58	121.29	109.12	1.1133
IC	C13S	C11S	*C12S	H12T	1.5345	112.58	-121.31	109.15	1.1133
IC	C11S	C12S	C13S	C14S	1.5344	112.58	179.97	112.58	1.5345
IC	C14S	C12S	*C13S	H13S	1.5345	112.58	121.30	109.13	1.1133
IC	C14S	C12S	*C13S	H13T	1.5345	112.58	-121.30	109.13	1.1133
IC	C12S	C13S	C14S	C15S	1.5345	112.58	-179.99	112.58	1.5345
IC	C15S	C13S	*C14S	H14S	1.5345	112.58	121.30	109.12	1.1133
IC	C15S	C13S	*C14S	H14T	1.5345	112.58	-121.31	109.12	1.1133
IC	C13S	C14S	C15S	C16S	1.5345	112.58	180.00	112.58	1.5346

IC	C16S	C14S	*C15S	H15S	1.5346	112.58	121.30	109.11	1.1133
IC	C16S	C14S	*C15S	H15T	1.5346	112.58	-121.30	109.11	1.1133
IC	C14S	C15S	C16S	C17S	1.5345	112.58	180.00	112.64	1.5337
IC	C17S	C15S	*C16S	H16S	1.5337	112.64	121.32	109.14	1.1133
IC	C17S	C15S	*C16S	H16T	1.5337	112.64	-121.32	109.14	1.1133
IC	C15S	C16S	C17S	C18S	1.5346	112.64	180.00	113.23	1.5309
IC	C18S	C16S	*C17S	H17S	1.5309	113.23	121.67	108.77	1.1142
IC	C18S	C16S	*C17S	H17T	1.5309	113.23	-121.67	108.77	1.1142
IC	C16S	C17S	C18S	H18S	1.5337	113.23	180.00	110.63	1.1113
IC	H18S	C17S	*C18S	H18T	1.1113	110.63	120.07	110.47	1.1114
IC	H18S	C17S	*C18S	H18U	1.1113	110.63	-120.07	110.47	1.1114
IC	C2S	NF	C1F	C2F	1.4406	124.85	-156.83	114.97	1.4964
IC	C2F	NF	*C1F	OF	1.4964	114.97	162.54	122.11	1.2268
IC	NF	C1F	C2F	C3F	1.3364	114.97	-146.57	111.92	1.5432
IC	C3F	C1F	*C2F	H2F	1.5432	111.92	-120.83	107.93	1.1115
IC	H2F	C1F	*C2F	H2G	1.1115	107.93	-117.30	109.04	1.1113
IC	C1F	C2F	C3F	C4F	1.4964	111.92	179.54	112.84	1.5310
IC	C4F	C2F	*C3F	H3F	1.5310	112.84	-121.63	109.37	1.1145
IC	H3F	C2F	*C3F	H3G	1.1145	109.37	-117.07	109.08	1.1157
IC	C2F	C3F	C4F	C5F	1.5434	112.23	174.86	112.58	1.5342
IC	C5F	C3F	*C4F	H4F	1.5342	112.58	-120.18	108.65	1.1140
IC	H4F	C3F	*C4F	H4G	1.1140	108.65	-117.52	109.62	1.1127
IC	C3F	C4F	C5F	C6F	1.5340	112.58	-174.06	112.39	1.5340
IC	C6F	C4F	*C5F	H5F	1.5340	112.39	-122.01	109.38	1.1130
IC	H5F	C4F	*C5F	H5G	1.1130	109.38	-117.34	108.92	1.1134
IC	C4F	C5F	C6F	C7F	1.5342	112.39	178.27	112.75	1.5340
IC	C7F	C5F	*C6F	H6F	1.5340	112.75	-120.93	108.96	1.1137
IC	H6F	C5F	*C6F	H6G	1.1137	108.96	-117.41	109.22	1.1131
IC	C5F	C6F	C7F	C8F	1.5340	112.75	-178.89	112.33	1.5342
IC	C8F	C6F	*C7F	H7F	1.5342	112.33	-121.40	109.26	1.1132
IC	H7F	C6F	*C7F	H7G	1.1132	109.26	-117.45	109.12	1.1130
IC	C6F	C7F	C8F	C9F	1.5340	112.33	179.09	112.77	1.5340
IC	C9F	C7F	*C8F	H8F	1.5340	112.77	-121.23	109.16	1.1134
IC	H8F	C7F	*C8F	H8G	1.1134	109.16	-117.38	109.08	1.1133
IC	C7F	C8F	C9F	C10F	1.5342	112.77	179.02	112.34	1.5343
IC	C10F	C8F	*C9F	H9F	1.5343	112.34	-121.27	109.27	1.1132
IC	H9F	C8F	*C9F	H9G	1.1132	109.27	-117.45	109.10	1.1130
IC	C8F	C9F	C10F	C11F	1.5340	112.34	178.16	112.74	1.5340
IC	C11F	C9F	*C10F	H10F	1.5340	112.74	-121.18	109.16	1.1134
IC	H10F	C9F	*C10F	H10G	1.1134	109.16	-117.35	109.10	1.1132
IC	C9F	C10F	C11F	C12F	1.5343	112.74	178.91	112.38	1.5343
IC	C12F	C10F	*C11F	H11F	1.5343	112.38	-121.39	109.34	1.1131
IC	H11F	C10F	*C11F	H11G	1.1131	109.34	-117.43	108.98	1.1132
IC	C10F	C11F	C12F	C13F	1.5340	112.38	176.93	112.68	1.5340
IC	C13F	C11F	*C12F	H12F	1.5340	112.68	-121.02	109.05	1.1134
IC	H12F	C11F	*C12F	H12G	1.1134	109.05	-117.34	109.20	1.1131
IC	C11F	C12F	C13F	C14F	1.5343	112.68	-179.93	112.45	1.5345
IC	C14F	C12F	*C13F	H13F	1.5345	112.45	-121.60	109.42	1.1129
IC	H13F	C12F	*C13F	H13G	1.1129	109.42	-117.40	108.85	1.1134
IC	C12F	C13F	C14F	C15F	1.5340	112.45	175.93	112.68	1.5334
IC	C15F	C13F	*C14F	H14F	1.5334	112.68	-121.00	109.06	1.1131
IC	H14F	C13F	*C14F	H14G	1.1131	109.06	-117.24	109.28	1.1130
IC	C13F	C14F	C15F	C16F	1.5345	112.68	-178.23	113.16	1.5306
IC	C16F	C14F	*C15F	H15F	1.5306	113.16	-121.86	108.84	1.1140
IC	H15F	C14F	*C15F	H15G	1.1140	108.84	-116.70	108.76	1.1144
IC	C14F	C15F	C16F	C17F	1.5345	112.68	-178.23	113.16	1.5306
IC	C17F	C15F	*C16F	H16F	1.5306	113.16	-121.86	108.84	1.1140
IC	H16F	C15F	*C16F	H16G	1.1140	108.84	-116.70	108.76	1.1144
IC	C15F	C16F	C17F	C18F	1.5345	112.68	-178.23	113.16	1.5306
IC	C18F	C16F	*C17F	H17F	1.5306	113.16	-121.86	108.84	1.1140
IC	H17F	C16F	*C17F	H17G	1.1140	108.84	-116.70	108.76	1.1144
IC	C16F	C17F	C18F	C19F	1.5345	112.68	-178.23	113.16	1.5306
IC	C19F	C17F	*C18F	H18F	1.5306	113.16	-121.86	108.84	1.1140
IC	H18F	C17F	*C18F	H18G	1.1140	108.84	-116.70	108.76	1.1144
IC	C17F	C18F	C19F	C20F	1.5344	112.57	179.80	112.59	1.5346
IC	C20F	C18F	*C19F	H19F	1.5346	112.59	-121.28	109.10	1.1133
IC	H19F	C18F	*C19F	H19G	1.1133	109.10	-117.38	109.12	1.1133
IC	C18F	C19F	C20F	C21F	1.5345	112.59	-179.89	112.63	1.5337
IC	C21F	C19F	*C20F	H20F	1.5337	112.63	-121.33	109.16	1.1133
IC	H20F	C19F	*C20F	H20G	1.1133	109.16	-117.38	109.14	1.1133
IC	C19F	C20F	C21F	C22F	1.5346	112.63	179.93	113.24	1.5309

IC	C22F	C20F	*C21F	H21F	1.5309	113.24	-121.66	108.77	1.1142
IC	H21F	C20F	*C21F	H21G	1.1142	108.77	-116.65	108.78	1.1142
IC	C20F	C21F	*C22F	H22F	1.5337	113.24	-59.91	110.47	1.1114
IC	H22F	C21F	*C22F	H22G	1.1114	110.47	119.87	110.48	1.1113
IC	H22F	C21F	*C22F	H22H	1.1114	110.47	-120.06	110.63	1.1113

PATCH FIRST NONE LAST NONE

RESI 23SM 0.00 ! sphingomyelin d18:1/23:0
 GROUP !
 ATOM N NTL -0.60 !
 ATOM C15 CTL5 -0.35 !
 ATOM H15A HL 0.25 !
 ATOM H15B HL 0.25 !
 ATOM H15C HL 0.25 !
 ATOM C14 CTL5 -0.35 !
 ATOM H14A HL 0.25 !
 ATOM H14B HL 0.25 !
 ATOM H14C HL 0.25 !
 ATOM C13 CTL5 -0.35 !
 ATOM H13A HL 0.25 !
 ATOM H13B HL 0.25 !
 ATOM H13C HL 0.25 !
 ATOM C12 CTL2 -0.10 !
 ATOM H12A HL 0.25 !
 ATOM H12B HL 0.25 !
 GROUP !
 ATOM C11 CTL2 -0.08 !
 ATOM H11A HAL2 0.09 !
 ATOM H11B HAL2 0.09 !
 ATOM P PL 1.50 !
 ATOM O13 O2L -0.78 !
 ATOM O14 O2L -0.78 !
 ATOM O11 OSLP -0.57 !
 ATOM O12 OSLP -0.57 !
 ATOM C1S CTL2 -0.08 !
 ATOM H1S HAL2 0.09 !
 ATOM H1T HAL2 0.09 !
 GROUP !
 ATOM NF NHL -0.70 !
 ATOM HNF H 0.35 !
 ATOM C2S CTL1 0.30 !
 ATOM H2S HAL1 0.05 !
 GROUP !
 ATOM C3S CTO1 0.50 !
 ATOM H3S HAL1 -0.11 !
 ATOM O3 OHL -0.69 !
 ATOM HO3 HOL 0.30 !
 GROUP !
 ATOM C4S CEL3 -0.15 !
 ATOM H4S HEL1 0.15 !
 GROUP !
 ATOM C5S CEL3 -0.15 !
 ATOM H5S HEL1 0.15 !
 GROUP !
 ATOM C6S CTL2 -0.18 !
 ATOM H6S HAL2 0.09 !
 ATOM H6T HAL2 0.09 !
 GROUP !
 ATOM C7S CTL2 -0.18 !
 ATOM H7S HAL2 0.09 !
 ATOM H7T HAL2 0.09 !
 GROUP !
 ATOM C8S CTL2 -0.18 !
 ATOM H8S HAL2 0.09 !
 ATOM H8T HAL2 0.09 !
 GROUP !
 ATOM C9S CTL2 -0.18 !
 ATOM H9S HAL2 0.09 !
 ATOM H9T HAL2 0.09 !
 GROUP !
 ATOM C10S CTL2 -0.18 !



ATOM H10S HAL2 0.09 !
 ATOM H10T HAL2 0.09 !
 GROUP
 ATOM C11S CTL2 -0.18 !
 ATOM H11S HAL2 0.09 !
 ATOM H11T HAL2 0.09 !
 GROUP
 ATOM C12S CTL2 -0.18 !
 ATOM H12S HAL2 0.09 !
 ATOM H12T HAL2 0.09 !
 GROUP
 ATOM C13S CTL2 -0.18 !
 ATOM H13S HAL2 0.09 !
 ATOM H13T HAL2 0.09 !
 GROUP
 ATOM C14S CTL2 -0.18 !
 ATOM H14S HAL2 0.09 !
 ATOM H14T HAL2 0.09 !
 GROUP
 ATOM C15S CTL2 -0.18 !
 ATOM H15S HAL2 0.09 !
 ATOM H15T HAL2 0.09 !
 GROUP
 ATOM C16S CTL2 -0.18 !
 ATOM H16S HAL2 0.09 !
 ATOM H16T HAL2 0.09 !
 GROUP
 ATOM C17S CTL2 -0.18 !
 ATOM H17S HAL2 0.09 !
 ATOM H17T HAL2 0.09 !
 GROUP
 ATOM C18S CTL3 -0.27 !
 ATOM H18S HAL3 0.09 !
 ATOM H18T HAL3 0.09 !
 ATOM H18U HAL3 0.09 !
 GROUP !FATTY ACID CHAIN
 ATOM C1F C 0.55 !
 ATOM O F O -0.60 !
 ATOM C2F CTL2 -0.07 !
 ATOM H2F HAL2 0.06 !
 ATOM H2G HAL2 0.06 !
 GROUP !
 ATOM C3F CTL2 -0.18 !
 ATOM H3F HAL2 0.09 !
 ATOM H3G HAL2 0.09 !
 GROUP !
 ATOM C4F CTL2 -0.18 !
 ATOM H4F HAL2 0.09 !
 ATOM H4G HAL2 0.09 !
 GROUP !
 ATOM C5F CTL2 -0.18 !
 ATOM H5F HAL2 0.09 !
 ATOM H5G HAL2 0.09 !
 GROUP !
 ATOM C6F CTL2 -0.18 !
 ATOM H6F HAL2 0.09 !
 ATOM H6G HAL2 0.09 !
 GROUP !
 ATOM C7F CTL2 -0.18 !
 ATOM H7F HAL2 0.09 !
 ATOM H7G HAL2 0.09 !
 GROUP !
 ATOM C8F CTL2 -0.18 !
 ATOM H8F HAL2 0.09 !
 ATOM H8G HAL2 0.09 !
 GROUP !
 ATOM C9F CTL2 -0.18 !
 ATOM H9F HAL2 0.09 !
 ATOM H9G HAL2 0.09 !
 GROUP !
 ATOM C10F CTL2 -0.18 !

ATOM H10F HAL2 0.09 !
 ATOM H10G HAL2 0.09 !
 GROUP
 ATOM C11F CTL2 -0.18 !
 ATOM H11F HAL2 0.09 !
 ATOM H11G HAL2 0.09 !
 GROUP
 ATOM C12F CTL2 -0.18 !
 ATOM H12F HAL2 0.09 !
 ATOM H12G HAL2 0.09 !
 GROUP
 ATOM C13F CTL2 -0.18 !
 ATOM H13F HAL2 0.09 !
 ATOM H13G HAL2 0.09 !
 GROUP
 ATOM C14F CTL2 -0.18 !
 ATOM H14F HAL2 0.09 !
 ATOM H14G HAL2 0.09 !
 GROUP
 ATOM C15F CTL2 -0.18 !
 ATOM H15F HAL2 0.09 !
 ATOM H15G HAL2 0.09 !
 GROUP
 ATOM C16F CTL2 -0.18 !
 ATOM H16F HAL2 0.09 !
 ATOM H16G HAL2 0.09 !
 GROUP
 ATOM C17F CTL2 -0.18 !
 ATOM H17F HAL2 0.09 !
 ATOM H17G HAL2 0.09 !
 GROUP
 ATOM C18F CTL2 -0.18 !
 ATOM H18F HAL2 0.09 !
 ATOM H18G HAL2 0.09 !
 GROUP
 ATOM C19F CTL2 -0.18 !
 ATOM H19F HAL2 0.09 !
 ATOM H19G HAL2 0.09 !
 GROUP
 ATOM C20F CTL2 -0.18 !
 ATOM H20F HAL2 0.09 !
 ATOM H20G HAL2 0.09 !
 GROUP
 ATOM C21F CTL2 -0.18 !
 ATOM H21F HAL2 0.09 !
 ATOM H21G HAL2 0.09 !
 GROUP
 ATOM C22F CTL2 -0.18 !
 ATOM H22F HAL2 0.09 !
 ATOM H22G HAL2 0.09 !
 GROUP
 ATOM C23F CTL3 -0.27 !
 ATOM H23F HAL3 0.09 !
 ATOM H23G HAL3 0.09 !
 ATOM H23H HAL3 0.09 !

!SPHINGOSINE CHAIN
 BOND C18S H18S C18S H18T C18S H18U
 BOND C18S C17S C17S H17S C17S H17T
 BOND C17S C16S C16S H16S C16S H16T
 BOND C16S C15S C15S H15S C15S H15T
 BOND C15S C14S C14S H14S C14S H14T
 BOND C14S C13S C13S H13S C13S H13T
 BOND C13S C12S C12S H12S C12S H12T
 BOND C12S C11S C11S H11S C11S H11T
 BOND C11S C10S C10S H10S C10S H10T
 BOND C10S C9S C9S H9S C9S H9T
 BOND C9S C8S C8S H8S C8S H8T
 BOND C8S C7S C7S H7S C7S H7T
 BOND C7S C6S C6S H6S C6S H6T
 BOND C6S C5S C5S H5S C4S H4S

BOND C4S C3S C3S H3S C3S O3
 BOND O3 HO3
 BOND C3S C2S C2S H2S
 BOND C2S C1S C1S H1S C1S H1T
 BOND C1S O11
 !PHOSLPHOCHOLINE
 BOND O11 P O12 P O13 P O14 P
 BOND O12 C11 C11 H11A C11 H11B
 BOND C11 C12 C12 H12A C12 H12B
 BOND C12 N N C13 N C14 N C15
 BOND C13 H13A C13 H13B C13 H13C
 BOND C14 H14A C14 H14B C14 H14C
 BOND C15 H15A C15 H15B C15 H15C
 !FATTY ACID CHAIN
 BOND C2S NF NF HNF NF C1F
 BOND C1F C2F C2F H2F C2F H2G
 BOND C2F C3F C3F H3F C3F H3G
 BOND C3F C4F C4F H4F C4F H4G
 BOND C4F C5F C5F H5F C5F H5G
 BOND C5F C6F C6F H6F C6F H6G
 BOND C6F C7F C7F H7F C7F H7G
 BOND C7F C8F C8F H8F C8F H8G
 BOND C8F C9F C9F H9F C9F H9G
 BOND C9F C10F C10F H10F C10F H10G
 BOND C10F C11F C11F H11F C11F H11G
 BOND C11F C12F C12F H12F C12F H12G
 BOND C12F C13F C13F H13F C13F H13G
 BOND C13F C14F C14F H14F C14F H14G
 BOND C14F C15F C15F H15F C15F H15G
 BOND C15F C16F C16F H16F C16F H16G
 BOND C16F C17F C17F H17F C17F H17G
 BOND C17F C18F C18F H18F C18F H18G
 BOND C18F C19F C19F H19F C19F H19G
 BOND C19F C20F C20F H20F C20F H20G
 BOND C20F C21F C21F H21F C21F H21G
 BOND C21F C22F C22F H22F C22F H22G
 BOND C22F C23F C23F H23F C23F H23G
 BOND C23F H23H

DOUBLE C4S C5S
 DOUBLE C1F OF
 IMPR C1F C2F NF OF NF C1F C2S HNF

IC	C18S	C17S	C16S	C15S	1.5309	113.23	180.00	112.64	1.5346
IC	C17S	C16S	C15S	C14S	1.5337	112.64	180.00	112.58	1.5345
IC	C16S	C15S	C14S	C13S	1.5346	112.58	180.00	112.58	1.5345
IC	C15S	C14S	C13S	C12S	1.5345	112.58	-179.99	112.58	1.5345
IC	C14S	C13S	C12S	C11S	1.5345	112.58	179.97	112.58	1.5344
IC	C13S	C12S	C11S	C10S	1.5345	112.58	-179.96	112.61	1.5345
IC	C12S	C11S	C10S	C9S	1.5344	112.61	179.97	112.54	1.5344
IC	C11S	C10S	C9S	C8S	1.5345	112.54	-179.87	112.66	1.5345
IC	C10S	C9S	C8S	C7S	1.5344	112.66	-179.89	112.50	1.5354
IC	C9S	C8S	C7S	C6S	1.5345	112.50	-179.79	112.41	1.5391
IC	C8S	C7S	C6S	C5S	1.5354	112.41	-179.61	111.46	1.5077
IC	C7S	C6S	C5S	C4S	1.5391	111.46	-117.13	124.77	1.3444
IC	C6S	C5S	C4S	C3S	1.5077	124.77	179.01	124.28	1.5009
IC	C5S	C4S	C3S	C2S	1.3444	124.28	-84.24	110.14	1.5343
IC	C5S	C4S	C3S	O3	1.3444	124.28	156.07	107.64	1.4235
IC	C4S	C3S	C2S	C1S	1.5009	110.14	-70.50	114.25	1.5586
IC	C13	C12	*N	C14	1.4967	109.68	116.91	107.30	1.5040
IC	C13	C12	*N	C15	1.4967	109.68	-123.42	110.88	1.4938
IC	C12	N	C15	H15A	1.5212	110.88	-177.15	111.31	1.0816
IC	H15A	N	*C15	H15B	1.0816	111.31	118.98	109.44	1.0875
IC	H15A	N	*C15	H15C	1.0816	111.31	-122.11	111.46	1.0818
IC	C12	N	C14	H14A	1.5212	107.30	-178.89	114.23	1.0739
IC	H14A	N	*C14	H14B	1.0739	114.23	123.44	110.21	1.0968
IC	H14A	N	*C14	H14C	1.0739	114.23	-124.43	111.41	1.0948
IC	C12	N	C13	H13A	1.5212	109.68	176.44	111.15	1.0822
IC	H13A	N	*C13	H13B	1.0822	111.15	121.89	111.75	1.0814
IC	H13A	N	*C13	H13C	1.0822	111.15	-118.50	109.57	1.0892
IC	C13	N	C12	C11	1.4967	109.68	-178.83	116.30	1.5392

IC	C11	N	*C12	H12A	1.5392	116.30	117.04	108.30	1.0985
IC	C11	N	*C12	H12B	1.5392	116.30	-126.61	112.36	1.0787
IC	N	C12	C11	O12	1.5212	116.30	126.79	107.85	1.4239
IC	O12	C12	*C11	H11A	1.4239	107.85	118.10	109.30	1.1131
IC	O12	C12	*C11	H11B	1.4239	107.85	-123.12	113.16	1.1140
IC	C12	C11	O12	P	1.5392	107.85	-68.29	118.09	1.5886
IC	C11	O12	P	O11	1.4239	118.09	-167.58	104.18	1.5725
IC	O11	O12	*P	O13	1.5725	104.18	116.57	107.05	1.4823
IC	O11	O12	*P	O14	1.5725	104.18	-118.73	106.98	1.4805
IC	O12	P	O11	C1S	1.5886	104.18	51.85	121.69	1.4327
IC	P	O11	C1S	C2S	1.5725	121.69	103.66	111.21	1.5586
IC	C2S	O11	*C1S	H1S	1.5586	111.21	118.79	108.35	1.1121
IC	C2S	O11	*C1S	H1T	1.5586	111.21	-123.25	112.85	1.1152
IC	O11	C1S	C2S	C3S	1.4327	111.21	175.67	114.25	1.5343
IC	C3S	C1S	*C2S	NF	1.5343	114.25	120.00	111.52	1.4406
IC	C3S	C1S	*C2S	H2S	1.5343	114.25	-120.00	105.03	1.1194
IC	C1S	C2S	NF	C1F	1.5586	111.52	160.94	124.85	1.3364
IC	C1F	C2S	*NF	HNF	1.3364	124.85	-157.81	113.39	1.0062
IC	C1S	C2S	C3S	C4S	1.5586	114.25	-70.50	110.14	1.5009
IC	C4S	C2S	*C3S	O3	1.5009	110.14	118.36	109.80	1.4235
IC	C4S	C2S	*C3S	H3S	1.5009	110.14	-121.14	110.10	1.1140
IC	C2S	C3S	O3	H03	1.5343	109.80	46.50	104.54	0.9610
IC	C2S	C3S	C4S	C5S	1.5343	110.14	-84.24	124.28	1.3444
IC	C5S	C3S	*C4S	H4S	1.3444	124.28	178.79	115.93	1.1003
IC	C3S	C4S	C5S	C6S	1.5009	124.28	179.01	124.77	1.5077
IC	C6S	C4S	*C5S	H5S	1.5077	124.77	-178.18	119.00	1.1029
IC	C4S	C5S	C6S	C7S	1.3444	124.77	-117.13	111.46	1.5391
IC	C7S	C5S	*C6S	H6S	1.5391	111.46	120.98	111.41	1.1142
IC	C7S	C5S	*C6S	H6T	1.5391	111.46	-120.60	109.91	1.1125
IC	C5S	C6S	C7S	C8S	1.5077	111.46	-179.61	112.41	1.5354
IC	C8S	C6S	*C7S	H7S	1.5354	112.41	121.26	109.74	1.1138
IC	C8S	C6S	*C7S	H7T	1.5354	112.41	-120.99	109.40	1.1147
IC	C6S	C7S	C8S	C9S	1.5391	112.41	-179.79	112.50	1.5345
IC	C9S	C7S	*C8S	H8S	1.5345	112.50	121.20	108.97	1.1132
IC	C9S	C7S	*C8S	H8T	1.5345	112.50	-121.38	109.07	1.1129
IC	C7S	C8S	C9S	C10S	1.5354	112.50	-179.89	112.66	1.5344
IC	C10S	C8S	*C9S	H9S	1.5344	112.66	121.49	109.05	1.1133
IC	C10S	C8S	*C9S	H9T	1.5344	112.66	-121.38	108.96	1.1136
IC	C8S	C9S	C10S	C11S	1.5345	112.66	-179.87	112.54	1.5345
IC	C11S	C9S	*C10S	H10S	1.5345	112.54	121.27	109.07	1.1133
IC	C11S	C9S	*C10S	H10T	1.5345	112.54	-121.34	109.11	1.1132
IC	C9S	C10S	C11S	C12S	1.5344	112.54	179.97	112.61	1.5344
IC	C12S	C10S	*C11S	H11S	1.5344	112.61	121.36	109.09	1.1133
IC	C12S	C10S	*C11S	H11T	1.5344	112.61	-121.33	109.07	1.1134
IC	C10S	C11S	C12S	C13S	1.5345	112.61	-179.96	112.58	1.5345
IC	C13S	C11S	*C12S	H12S	1.5345	112.58	121.29	109.12	1.1133
IC	C13S	C11S	*C12S	H12T	1.5345	112.58	-121.31	109.15	1.1133
IC	C11S	C12S	C13S	C14S	1.5344	112.58	179.97	112.58	1.5345
IC	C14S	C12S	*C13S	H13S	1.5345	112.58	121.30	109.13	1.1133
IC	C14S	C12S	*C13S	H13T	1.5345	112.58	-121.30	109.13	1.1133
IC	C12S	C13S	C14S	C15S	1.5345	112.58	-179.99	112.58	1.5345
IC	C15S	C13S	*C14S	H14S	1.5345	112.58	121.30	109.12	1.1133
IC	C15S	C13S	*C14S	H14T	1.5345	112.58	-121.31	109.12	1.1133
IC	C13S	C14S	C15S	C16S	1.5345	112.58	180.00	112.58	1.5346
IC	C16S	C14S	*C15S	H15S	1.5346	112.58	121.30	109.11	1.1133
IC	C16S	C14S	*C15S	H15T	1.5346	112.58	-121.30	109.11	1.1133
IC	C14S	C15S	C16S	C17S	1.5345	112.58	180.00	112.64	1.5337
IC	C17S	C15S	*C16S	H16S	1.5337	112.64	121.32	109.14	1.1133
IC	C17S	C15S	*C16S	H16T	1.5337	112.64	-121.32	109.14	1.1133
IC	C15S	C16S	C17S	C18S	1.5346	112.64	180.00	113.23	1.5309
IC	C18S	C16S	*C17S	H17S	1.5309	113.23	121.67	108.77	1.1142
IC	C18S	C16S	*C17S	H17T	1.5309	113.23	-121.67	108.77	1.1142
IC	C16S	C17S	C18S	H18S	1.5337	113.23	180.00	110.63	1.1113
IC	H18S	C17S	*C18S	H18T	1.1113	110.63	120.07	110.47	1.1114
IC	H18S	C17S	*C18S	H18U	1.1113	110.63	-120.07	110.47	1.1114
IC	C2S	NF	C1F	C2F	1.4406	124.85	-156.83	114.97	1.4964
IC	C2F	NF	*C1F	OF	1.4964	114.97	162.54	122.11	1.2268
IC	NF	C1F	C2F	C3F	1.3364	114.97	-146.57	111.92	1.5432
IC	C3F	C1F	*C2F	H2F	1.5432	111.92	-120.83	107.93	1.1115
IC	H2F	C1F	*C2F	H2G	1.1115	107.93	-117.30	109.04	1.1113
IC	C1F	C2F	C3F	C4F	1.4964	111.92	179.54	112.84	1.5310

IC	C4F	C2F	*C3F	H3F	1.5310	112.84	-121.63	109.37	1.1145
IC	H3F	C2F	*C3F	H3G	1.1145	109.37	-117.07	109.08	1.1157
IC	C2F	C3F	C4F	C5F	1.5434	112.23	174.86	112.58	1.5342
IC	C5F	C3F	*C4F	H4F	1.5342	112.58	-120.18	108.65	1.1140
IC	H4F	C3F	*C4F	H4G	1.1140	108.65	-117.52	109.62	1.1127
IC	C3F	C4F	C5F	C6F	1.5340	112.58	-174.06	112.39	1.5340
IC	C6F	C4F	*C5F	H5F	1.5340	112.39	-122.01	109.38	1.1130
IC	H5F	C4F	*C5F	H5G	1.1130	109.38	-117.34	108.92	1.1134
IC	C4F	C5F	C6F	C7F	1.5342	112.39	178.27	112.75	1.5340
IC	C7F	C5F	*C6F	H6F	1.5340	112.75	-120.93	108.96	1.1137
IC	H6F	C5F	*C6F	H6G	1.1137	108.96	-117.41	109.22	1.1131
IC	C5F	C6F	C7F	C8F	1.5340	112.75	-178.89	112.33	1.5342
IC	C8F	C6F	*C7F	H7F	1.5342	112.33	-121.40	109.26	1.1132
IC	H7F	C6F	*C7F	H7G	1.1132	109.26	-117.45	109.12	1.1130
IC	C6F	C7F	C8F	C9F	1.5340	112.33	179.09	112.77	1.5340
IC	C9F	C7F	*C8F	H8F	1.5340	112.77	-121.23	109.16	1.1134
IC	H8F	C7F	*C8F	H8G	1.1134	109.16	-117.38	109.08	1.1133
IC	C7F	C8F	C9F	C10F	1.5342	112.77	179.02	112.34	1.5343
IC	C10F	C8F	*C9F	H9F	1.5343	112.34	-121.27	109.27	1.1132
IC	H9F	C8F	*C9F	H9G	1.1132	109.27	-117.45	109.10	1.1130
IC	C8F	C9F	C10F	C11F	1.5340	112.34	178.16	112.74	1.5340
IC	C11F	C9F	*C10F	H10F	1.5340	112.74	-121.18	109.16	1.1134
IC	H10F	C9F	*C10F	H10G	1.1134	109.16	-117.35	109.10	1.1132
IC	C9F	C10F	C11F	C12F	1.5343	112.74	178.91	112.38	1.5343
IC	C12F	C10F	*C11F	H11F	1.5343	112.38	-121.39	109.34	1.1131
IC	H11F	C10F	*C11F	H11G	1.1131	109.34	-117.43	108.98	1.1132
IC	C10F	C11F	C12F	C13F	1.5340	112.38	176.93	112.68	1.5340
IC	C13F	C11F	*C12F	H12F	1.5340	112.68	-121.02	109.05	1.1134
IC	H12F	C11F	*C12F	H12G	1.1134	109.05	-117.34	109.20	1.1131
IC	C11F	C12F	C13F	C14F	1.5343	112.68	-179.93	112.45	1.5345
IC	C14F	C12F	*C13F	H13F	1.5345	112.45	-121.60	109.42	1.1129
IC	H13F	C12F	*C13F	H13G	1.1129	109.42	-117.40	108.85	1.1134
IC	C12F	C13F	C14F	C15F	1.5340	112.45	175.93	112.68	1.5334
IC	C15F	C13F	*C14F	H14F	1.5334	112.68	-121.00	109.06	1.1131
IC	H14F	C13F	*C14F	H14G	1.1131	109.06	-117.24	109.28	1.1130
IC	C13F	C14F	C15F	C16F	1.5345	112.68	-178.23	113.16	1.5306
IC	C16F	C14F	*C15F	H15F	1.5306	113.16	-121.86	108.84	1.1140
IC	H15F	C14F	*C15F	H15G	1.1140	108.84	-116.70	108.76	1.1144
IC	C14F	C15F	C16F	C17F	1.5345	112.68	-178.23	113.16	1.5306
IC	C17F	C15F	*C16F	H16F	1.5306	113.16	-121.86	108.84	1.1140
IC	H16F	C15F	*C16F	H16G	1.1140	108.84	-116.70	108.76	1.1144
IC	C15F	C16F	C17F	C18F	1.5345	112.68	-178.23	113.16	1.5306
IC	C18F	C16F	*C17F	H17F	1.5306	113.16	-121.86	108.84	1.1140
IC	H17F	C16F	*C17F	H17G	1.1140	108.84	-116.70	108.76	1.1144
IC	C16F	C17F	C18F	C19F	1.5345	112.68	-178.23	113.16	1.5306
IC	C19F	C17F	*C18F	H18F	1.5306	113.16	-121.86	108.84	1.1140
IC	H18F	C17F	*C18F	H18G	1.1140	108.84	-116.70	108.76	1.1144
IC	C17F	C18F	C19F	C20F	1.5344	112.57	179.80	112.59	1.5346
IC	C20F	C18F	*C19F	H19F	1.5346	112.59	-121.28	109.10	1.1133
IC	H19F	C18F	*C19F	H19G	1.1133	109.10	-117.38	109.12	1.1133
IC	C18F	C19F	C20F	C21F	1.5345	112.59	-179.89	112.63	1.5337
IC	C21F	C19F	*C20F	H20F	1.5337	112.63	-121.33	109.16	1.1133
IC	H20F	C19F	*C20F	H20G	1.1133	109.16	-117.38	109.14	1.1133
IC	C19F	C20F	C21F	C22F	1.5346	112.63	179.93	113.24	1.5309
IC	C22F	C20F	*C21F	H21F	1.5309	113.24	-121.66	108.77	1.1142
IC	H21F	C20F	*C21F	H21G	1.1142	108.77	-116.65	108.78	1.1142
IC	C20F	C21F	C22F	C23F	1.5346	112.63	179.93	113.24	1.5309
IC	C23F	C21F	*C22F	H22F	1.5309	113.24	-121.66	108.77	1.1142
IC	H22F	C21F	*C22F	H22G	1.1142	108.77	-116.65	108.78	1.1142
IC	C21F	C22F	C23F	H23F	1.5337	113.24	-59.91	110.47	1.1114
IC	H23F	C22F	*C23F	H23G	1.1114	110.47	119.87	110.48	1.1113
IC	H23F	C22F	*C23F	H23H	1.1114	110.47	-120.06	110.63	1.1113

PATCH FIRST NONE LAST NONE

RESI LSM	0.00 ! sphingomyelin d18:1/24:0
GROUP	!
ATOM N NTL	-0.60 ! H15B
ATOM C15 CTL5	-0.35 !
ATOM H15A HL	0.25 ! H15A-C15-H15C
ATOM H15B HL	0.25 !
ATOM H15C HL	0.25 ! H13C H14C

ATOM C14 CTL5 -0.35 !
 ATOM H14A HL 0.25 !
 ATOM H14B HL 0.25 !
 ATOM H14C HL 0.25 !
 ATOM C13 CTL5 -0.35 !
 ATOM H13A HL 0.25 !
 ATOM H13B HL 0.25 !
 ATOM H13C HL 0.25 !
 ATOM C12 CTL2 -0.10 !
 ATOM H12A HL 0.25 !
 ATOM H12B HL 0.25 !
 GROUP !
 ATOM C11 CTL2 -0.08 !
 ATOM H11A HAL2 0.09 !
 ATOM H11B HAL2 0.09 !
 ATOM P PL 1.50 !
 ATOM O13 O2L -0.78 !
 ATOM O14 O2L -0.78 !
 ATOM O11 OSLP -0.57 !
 ATOM O12 OSLP -0.57 !
 ATOM C1S CTL2 -0.08 !
 ATOM H1S HAL2 0.09 !
 ATOM H1T HAL2 0.09 !
 GROUP !
 ATOM NF NHL -0.70 !
 ATOM HNF H 0.35 !
 ATOM C2S CTL1 0.30 !
 ATOM H2S HAL1 0.05 !
 GROUP !
 ATOM C3S CTO1 0.50 !
 ATOM H3S HAL1 -0.11 !
 ATOM O3 OHL -0.69 !
 ATOM HO3 HOL 0.30 !
 GROUP !
 ATOM C4S CEL3 -0.15 !
 ATOM H4S HEL1 0.15 !
 GROUP !
 ATOM C5S CEL3 -0.15 !
 ATOM H5S HEL1 0.15 !
 GROUP !
 ATOM C6S CTL2 -0.18 !
 ATOM H6S HAL2 0.09 !
 ATOM H6T HAL2 0.09 !
 GROUP !
 ATOM C7S CTL2 -0.18 !
 ATOM H7S HAL2 0.09 !
 ATOM H7T HAL2 0.09 !
 GROUP !
 ATOM C8S CTL2 -0.18 !
 ATOM H8S HAL2 0.09 !
 ATOM H8T HAL2 0.09 !
 GROUP !
 ATOM C9S CTL2 -0.18 !
 ATOM H9S HAL2 0.09 !
 ATOM H9T HAL2 0.09 !
 GROUP !
 ATOM C10S CTL2 -0.18 !
 ATOM H10S HAL2 0.09 !
 ATOM H10T HAL2 0.09 !
 GROUP !
 ATOM C11S CTL2 -0.18 !
 ATOM H11S HAL2 0.09 !
 ATOM H11T HAL2 0.09 !
 GROUP !
 ATOM C12S CTL2 -0.18 !
 ATOM H12S HAL2 0.09 !
 ATOM H12T HAL2 0.09 !
 GROUP !
 ATOM C13S CTL2 -0.18 !
 ATOM H13S HAL2 0.09 !
 ATOM H13T HAL2 0.09 !

GROUP
 ATOM C14S CTL2 -0.18 !
 ATOM H14S HAL2 0.09 !
 ATOM H14T HAL2 0.09 !
 GROUP
 ATOM C15S CTL2 -0.18 !
 ATOM H15S HAL2 0.09 !
 ATOM H15T HAL2 0.09 !
 GROUP
 ATOM C16S CTL2 -0.18 !
 ATOM H16S HAL2 0.09 !
 ATOM H16T HAL2 0.09 !
 GROUP
 ATOM C17S CTL2 -0.18 !
 ATOM H17S HAL2 0.09 !
 ATOM H17T HAL2 0.09 !
 GROUP
 ATOM C18S CTL3 -0.27 !
 ATOM H18S HAL3 0.09 !
 ATOM H18T HAL3 0.09 !
 ATOM H18U HAL3 0.09 !
 GROUP !FATTY ACID CHAIN
 ATOM C1F C 0.55 !
 ATOM O F -0.60 !
 ATOM C2F CTL2 -0.07 !
 ATOM H2F HAL2 0.06 !
 ATOM H2G HAL2 0.06 !
 GROUP !
 ATOM C3F CTL2 -0.18 !
 ATOM H3F HAL2 0.09 !
 ATOM H3G HAL2 0.09 !
 GROUP !
 ATOM C4F CTL2 -0.18 !
 ATOM H4F HAL2 0.09 !
 ATOM H4G HAL2 0.09 !
 GROUP !
 ATOM C5F CTL2 -0.18 !
 ATOM H5F HAL2 0.09 !
 ATOM H5G HAL2 0.09 !
 GROUP !
 ATOM C6F CTL2 -0.18 !
 ATOM H6F HAL2 0.09 !
 ATOM H6G HAL2 0.09 !
 GROUP !
 ATOM C7F CTL2 -0.18 !
 ATOM H7F HAL2 0.09 !
 ATOM H7G HAL2 0.09 !
 GROUP !
 ATOM C8F CTL2 -0.18 !
 ATOM H8F HAL2 0.09 !
 ATOM H8G HAL2 0.09 !
 GROUP !
 ATOM C9F CTL2 -0.18 !
 ATOM H9F HAL2 0.09 !
 ATOM H9G HAL2 0.09 !
 GROUP !
 ATOM C10F CTL2 -0.18 !
 ATOM H10F HAL2 0.09 !
 ATOM H10G HAL2 0.09 !
 GROUP !
 ATOM C11F CTL2 -0.18 !
 ATOM H11F HAL2 0.09 !
 ATOM H11G HAL2 0.09 !
 GROUP !
 ATOM C12F CTL2 -0.18 !
 ATOM H12F HAL2 0.09 !
 ATOM H12G HAL2 0.09 !
 GROUP !
 ATOM C13F CTL2 -0.18 !
 ATOM H13F HAL2 0.09 !
 ATOM H13G HAL2 0.09 !

GROUP !
 ATOM C14F CTL2 -0.18 !
 ATOM H14F HAL2 0.09 !
 ATOM H14G HAL2 0.09 !
 GROUP !
 ATOM C15F CTL2 -0.18 !
 ATOM H15F HAL2 0.09 !
 ATOM H15G HAL2 0.09 !
 GROUP !
 ATOM C16F CTL2 -0.18 !
 ATOM H16F HAL2 0.09 !
 ATOM H16G HAL2 0.09 !
 GROUP !
 ATOM C17F CTL2 -0.18 !
 ATOM H17F HAL2 0.09 !
 ATOM H17G HAL2 0.09 !
 GROUP !
 ATOM C18F CTL2 -0.18 !
 ATOM H18F HAL2 0.09 !
 ATOM H18G HAL2 0.09 !
 GROUP !
 ATOM C19F CTL2 -0.18 !
 ATOM H19F HAL2 0.09 !
 ATOM H19G HAL2 0.09 !
 GROUP !
 ATOM C20F CTL2 -0.18 !
 ATOM H20F HAL2 0.09 !
 ATOM H20G HAL2 0.09 !
 GROUP !
 ATOM C21F CTL2 -0.18 !
 ATOM H21F HAL2 0.09 !
 ATOM H21G HAL2 0.09 !
 GROUP !
 ATOM C22F CTL2 -0.18 !
 ATOM H22F HAL2 0.09 !
 ATOM H22G HAL2 0.09 !
 GROUP !
 ATOM C23F CTL2 -0.18 !
 ATOM H23F HAL2 0.09 !
 ATOM H23G HAL2 0.09 !
 GROUP !
 ATOM C24F CTL3 -0.27 !
 ATOM H24F HAL3 0.09 !
 ATOM H24G HAL3 0.09 !
 ATOM H24H HAL3 0.09 !

!SPHINGOSINE CHAIN
 BOND C18S H18S C18S H18T C18S H18U
 BOND C18S C17S C17S H17S C17S H17T
 BOND C17S C16S C16S H16S C16S H16T
 BOND C16S C15S C15S H15S C15S H15T
 BOND C15S C14S C14S H14S C14S H14T
 BOND C14S C13S C13S H13S C13S H13T
 BOND C13S C12S C12S H12S C12S H12T
 BOND C12S C11S C11S H11S C11S H11T
 BOND C11S C10S C10S H10S C10S H10T
 BOND C10S C9S C9S H9S C9S H9T
 BOND C9S C8S C8S H8S C8S H8T
 BOND C8S C7S C7S H7S C7S H7T
 BOND C7S C6S C6S H6S C6S H6T
 BOND C6S C5S C5S H5S C4S H4S
 BOND C4S C3S C3S H3S C3S O3
 BOND O3 HO3
 BOND C3S C2S C2S H2S
 BOND C2S C1S C1S H1S C1S H1T
 BOND C1S O11

!PHOSPHOCHOLINE
 BOND O11 P O12 P O13 P O14 P
 BOND O12 C11 C11 H11A C11 H11B
 BOND C11 C12 C12 H12A C12 H12B
 BOND C12 N N C13 N C14 N C15

BOND C13 H13A C13 H13B C13 H13C
 BOND C14 H14A C14 H14B C14 H14C
 BOND C15 H15A C15 H15B C15 H15C
 !FATTY ACID CHAIN
 BOND C2S NF NF HNF NF C1F
 BOND C1F C2F C2F H2F C2F H2G
 BOND C2F C3F C3F H3F C3F H3G
 BOND C3F C4F C4F H4F C4F H4G
 BOND C4F C5F C5F H5F C5F H5G
 BOND C5F C6F C6F H6F C6F H6G
 BOND C6F C7F C7F H7F C7F H7G
 BOND C7F C8F C8F H8F C8F H8G
 BOND C8F C9F C9F H9F C9F H9G
 BOND C9F C10F C10F H10F C10F H10G
 BOND C10F C11F C11F H11F C11F H11G
 BOND C11F C12F C12F H12F C12F H12G
 BOND C12F C13F C13F H13F C13F H13G
 BOND C13F C14F C14F H14F C14F H14G
 BOND C14F C15F C15F H15F C15F H15G
 BOND C15F C16F C16F H16F C16F H16G
 BOND C16F C17F C17F H17F C17F H17G
 BOND C17F C18F C18F H18F C18F H18G
 BOND C18F C19F C19F H19F C19F H19G
 BOND C19F C20F C20F H20F C20F H20G
 BOND C20F C21F C21F H21F C21F H21G
 BOND C21F C22F C22F H22F C22F H22G
 BOND C22F C23F C23F H23F C23F H23G
 BOND C23F C24F C24F H24F C24F H24G
 BOND C24F H24H

DOUBLE C4S C5S
 DOUBLE C1F OF
 IMPR C1F C2F NF OF NF C1F C2S HNF

IC	C18S	C17S	C16S	C15S	1.5309	113.23	180.00	112.64	1.5346
IC	C17S	C16S	C15S	C14S	1.5337	112.64	180.00	112.58	1.5345
IC	C16S	C15S	C14S	C13S	1.5346	112.58	180.00	112.58	1.5345
IC	C15S	C14S	C13S	C12S	1.5345	112.58	-179.99	112.58	1.5345
IC	C14S	C13S	C12S	C11S	1.5345	112.58	179.97	112.58	1.5344
IC	C13S	C12S	C11S	C10S	1.5345	112.58	-179.96	112.61	1.5345
IC	C12S	C11S	C10S	C9S	1.5344	112.61	179.97	112.54	1.5344
IC	C11S	C10S	C9S	C8S	1.5345	112.54	-179.87	112.66	1.5345
IC	C10S	C9S	C8S	C7S	1.5344	112.66	-179.89	112.50	1.5345
IC	C9S	C8S	C7S	C6S	1.5345	112.50	-179.79	112.41	1.5391
IC	C8S	C7S	C6S	C5S	1.5354	112.41	-179.61	111.46	1.5077
IC	C7S	C6S	C5S	C4S	1.5391	111.46	-117.13	124.77	1.3444
IC	C6S	C5S	C4S	C3S	1.5077	124.77	179.01	124.28	1.5009
IC	C5S	C4S	C3S	C2S	1.3444	124.28	-84.24	110.14	1.5343
IC	C5S	C4S	C3S	O3	1.3444	124.28	156.07	107.64	1.4235
IC	C4S	C3S	C2S	C1S	1.5009	110.14	-70.50	114.25	1.5586
IC	C13	C12	*N	C14	1.4967	109.68	116.91	107.30	1.5040
IC	C13	C12	*N	C15	1.4967	109.68	-123.42	110.88	1.4938
IC	C12	N	C15	H15A	1.5212	110.88	-177.15	111.31	1.0816
IC	H15A	N	*C15	H15B	1.0816	111.31	118.98	109.44	1.0875
IC	H15A	N	*C15	H15C	1.0816	111.31	-122.11	111.46	1.0818
IC	C12	N	C14	H14A	1.5212	107.30	-178.89	114.23	1.0739
IC	H14A	N	*C14	H14B	1.0739	114.23	123.44	110.21	1.0968
IC	H14A	N	*C14	H14C	1.0739	114.23	-124.43	111.41	1.0948
IC	C12	N	C13	H13A	1.5212	109.68	176.44	111.15	1.0822
IC	H13A	N	*C13	H13B	1.0822	111.15	121.89	111.75	1.0814
IC	H13A	N	*C13	H13C	1.0822	111.15	-118.50	109.57	1.0892
IC	C13	N	C12	C11	1.4967	109.68	-178.83	116.30	1.5392
IC	C11	N	*C12	H12A	1.5392	116.30	117.04	108.30	1.0985
IC	C11	N	*C12	H12B	1.5392	116.30	-126.61	112.36	1.0787
IC	N	C12	C11	O12	1.5212	116.30	126.79	107.85	1.4239
IC	O12	C12	*C11	H11A	1.4239	107.85	118.10	109.30	1.1131
IC	O12	C12	*C11	H11B	1.4239	107.85	-123.12	113.16	1.1140
IC	C12	C11	O12	P	1.5392	107.85	-68.29	118.09	1.5886
IC	C11	O12	P	O11	1.4239	118.09	-167.58	104.18	1.5725
IC	O11	O12	*P	O13	1.5725	104.18	116.57	107.05	1.4823
IC	O11	O12	*P	O14	1.5725	104.18	-118.73	106.98	1.4805

IC	O12	P	O11	C1S	1.5886	104.18	51.85	121.69	1.4327
IC	P	O11	C1S	C2S	1.5725	121.69	103.66	111.21	1.5586
IC	C2S	O11	*C1S	H1S	1.5586	111.21	118.79	108.35	1.1121
IC	C2S	O11	*C1S	H1T	1.5586	111.21	-123.25	112.85	1.1152
IC	O11	C1S	C2S	C3S	1.4327	111.21	175.67	114.25	1.5343
IC	C3S	C1S	*C2S	NF	1.5343	114.25	120.00	111.52	1.4406
IC	C3S	C1S	*C2S	H2S	1.5343	114.25	-120.00	105.03	1.1194
IC	C1S	C2S	NF	C1F	1.5586	111.52	160.94	124.85	1.3364
IC	C1F	C2S	*NF	HNF	1.3364	124.85	-157.81	113.39	1.0062
IC	C1S	C2S	C3S	C4S	1.5586	114.25	-70.50	110.14	1.5009
IC	C4S	C2S	*C3S	O3	1.5009	110.14	118.36	109.80	1.4235
IC	C4S	C2S	*C3S	H3S	1.5009	110.14	-121.14	110.10	1.1140
IC	C2S	C3S	O3	HO3	1.5343	109.80	46.50	104.54	0.9610
IC	C2S	C3S	C4S	C5S	1.5343	110.14	-84.24	124.28	1.3444
IC	C5S	C3S	*C4S	H4S	1.3444	124.28	178.79	115.93	1.1003
IC	C3S	C4S	C5S	C6S	1.5009	124.28	179.01	124.77	1.5077
IC	C6S	C4S	*C5S	H5S	1.5077	124.77	-178.18	119.00	1.1029
IC	C4S	C5S	C6S	C7S	1.3444	124.77	-117.13	111.46	1.5391
IC	C7S	C5S	*C6S	H6S	1.5391	111.46	120.98	111.41	1.1142
IC	C7S	C5S	*C6S	H6T	1.5391	111.46	-120.60	109.91	1.1125
IC	C5S	C6S	C7S	C8S	1.5077	111.46	-179.61	112.41	1.5354
IC	C8S	C6S	*C7S	H7S	1.5354	112.41	121.26	109.74	1.1138
IC	C8S	C6S	*C7S	H7T	1.5354	112.41	-120.99	109.40	1.1147
IC	C6S	C7S	C8S	C9S	1.5391	112.41	-179.79	112.50	1.5345
IC	C9S	C7S	*C8S	H8S	1.5345	112.50	121.20	108.97	1.1132
IC	C9S	C7S	*C8S	H8T	1.5345	112.50	-121.38	109.07	1.1129
IC	C7S	C8S	C9S	C10S	1.5354	112.50	-179.89	112.66	1.5344
IC	C10S	C8S	*C9S	H9S	1.5344	112.66	121.49	109.05	1.1133
IC	C10S	C8S	*C9S	H9T	1.5344	112.66	-121.38	108.96	1.1136
IC	C8S	C9S	C10S	C11S	1.5345	112.66	-179.87	112.54	1.5345
IC	C11S	C9S	*C10S	H10S	1.5345	112.54	121.27	109.07	1.1133
IC	C11S	C9S	*C10S	H10T	1.5345	112.54	-121.34	109.11	1.1132
IC	C9S	C10S	C11S	C12S	1.5344	112.54	179.97	112.61	1.5344
IC	C12S	C10S	*C11S	H11S	1.5344	112.61	121.36	109.09	1.1133
IC	C12S	C10S	*C11S	H11T	1.5344	112.61	-121.33	109.07	1.1134
IC	C10S	C11S	C12S	C13S	1.5345	112.61	-179.96	112.58	1.5345
IC	C13S	C11S	*C12S	H12S	1.5345	112.58	121.29	109.12	1.1133
IC	C13S	C11S	*C12S	H12T	1.5345	112.58	-121.31	109.15	1.1133
IC	C11S	C12S	C13S	C14S	1.5344	112.58	179.97	112.58	1.5345
IC	C14S	C12S	*C13S	H13S	1.5345	112.58	121.30	109.13	1.1133
IC	C14S	C12S	*C13S	H13T	1.5345	112.58	-121.30	109.13	1.1133
IC	C12S	C13S	C14S	C15S	1.5345	112.58	-179.99	112.58	1.5345
IC	C15S	C13S	*C14S	H14S	1.5345	112.58	121.30	109.12	1.1133
IC	C15S	C13S	*C14S	H14T	1.5345	112.58	-121.31	109.12	1.1133
IC	C13S	C14S	C15S	C16S	1.5345	112.58	180.00	112.58	1.5346
IC	C16S	C14S	*C15S	H15S	1.5346	112.58	121.30	109.11	1.1133
IC	C16S	C14S	*C15S	H15T	1.5346	112.58	-121.30	109.11	1.1133
IC	C14S	C15S	C16S	C17S	1.5345	112.58	180.00	112.64	1.5337
IC	C17S	C15S	*C16S	H16S	1.5337	112.64	121.32	109.14	1.1133
IC	C17S	C15S	*C16S	H16T	1.5337	112.64	-121.32	109.14	1.1133
IC	C15S	C16S	C17S	C18S	1.5346	112.64	180.00	113.23	1.5309
IC	C18S	C16S	*C17S	H17S	1.5309	113.23	121.67	108.77	1.1142
IC	C18S	C16S	*C17S	H17T	1.5309	113.23	-121.67	108.77	1.1142
IC	C16S	C17S	C18S	H18S	1.5337	113.23	180.00	110.63	1.1113
IC	H18S	C17S	*C18S	H18T	1.1113	110.63	120.07	110.47	1.1114
IC	H18S	C17S	*C18S	H18U	1.1113	110.63	-120.07	110.47	1.1114
IC	C2S	NF	C1F	C2F	1.4406	124.85	-156.83	114.97	1.4964
IC	C2F	NF	*C1F	OF	1.4964	114.97	162.54	122.11	1.2268
IC	NF	C1F	C2F	C3F	1.3364	114.97	-146.57	111.92	1.5432
IC	C3F	C1F	*C2F	H2F	1.5432	111.92	-120.83	107.93	1.1115
IC	H2F	C1F	*C2F	H2G	1.1115	107.93	-117.30	109.04	1.1113
IC	C1F	C2F	C3F	C4F	1.4964	111.92	179.54	112.84	1.5310
IC	C4F	C2F	*C3F	H3F	1.5310	112.84	-121.63	109.37	1.1145
IC	H3F	C2F	*C3F	H3G	1.1145	109.37	-117.07	109.08	1.1157
IC	C2F	C3F	C4F	C5F	1.5434	112.23	174.86	112.58	1.5342
IC	C5F	C3F	*C4F	H4F	1.5342	112.58	-120.18	108.65	1.1140
IC	H4F	C3F	*C4F	H4G	1.1140	108.65	-117.52	109.62	1.1127
IC	C3F	C4F	C5F	C6F	1.5340	112.58	-174.06	112.39	1.5340
IC	C6F	C4F	*C5F	H5F	1.5340	112.39	-122.01	109.38	1.1130
IC	H5F	C4F	*C5F	H5G	1.1130	109.38	-117.34	108.92	1.1134
IC	C4F	C5F	C6F	C7F	1.5342	112.39	178.27	112.75	1.5340

IC	C7F	C5F	*C6F	H6F	1.5340	112.75	-120.93	108.96	1.1137
IC	H6F	C5F	*C6F	H6G	1.1137	108.96	-117.41	109.22	1.1131
IC	C5F	C6F	C7F	C8F	1.5340	112.75	-178.89	112.33	1.5342
IC	C8F	C6F	*C7F	H7F	1.5342	112.33	-121.40	109.26	1.1132
IC	H7F	C6F	*C7F	H7G	1.1132	109.26	-117.45	109.12	1.1130
IC	C6F	C7F	C8F	C9F	1.5340	112.33	179.09	112.77	1.5340
IC	C9F	C7F	*C8F	H8F	1.5340	112.77	-121.23	109.16	1.1134
IC	H8F	C7F	*C8F	H8G	1.1134	109.16	-117.38	109.08	1.1133
IC	C7F	C8F	C9F	C10F	1.5342	112.77	179.02	112.34	1.5343
IC	C10F	C8F	*C9F	H9F	1.5343	112.34	-121.27	109.27	1.1132
IC	H9F	C8F	*C9F	H9G	1.1132	109.27	-117.45	109.10	1.1130
IC	C8F	C9F	C10F	C11F	1.5340	112.34	178.16	112.74	1.5340
IC	C11F	C9F	*C10F	H10F	1.5340	112.74	-121.18	109.16	1.1134
IC	H10F	C9F	*C10F	H10G	1.1134	109.16	-117.35	109.10	1.1132
IC	C9F	C10F	C11F	C12F	1.5343	112.74	178.91	112.38	1.5343
IC	C12F	C10F	*C11F	H11F	1.5343	112.38	-121.39	109.34	1.1131
IC	H11F	C10F	*C11F	H11G	1.1131	109.34	-117.43	108.98	1.1132
IC	C10F	C11F	C12F	C13F	1.5340	112.38	176.93	112.68	1.5340
IC	C13F	C11F	*C12F	H12F	1.5340	112.68	-121.02	109.05	1.1134
IC	H12F	C11F	*C12F	H12G	1.1134	109.05	-117.34	109.20	1.1131
IC	C11F	C12F	C13F	C14F	1.5343	112.68	-179.93	112.45	1.5345
IC	C14F	C12F	*C13F	H13F	1.5345	112.45	-121.60	109.42	1.1129
IC	H13F	C12F	*C13F	H13G	1.1129	109.42	-117.40	108.85	1.1134
IC	C12F	C13F	C14F	C15F	1.5340	112.45	175.93	112.68	1.5334
IC	C15F	C13F	*C14F	H14F	1.5334	112.68	-121.00	109.06	1.1131
IC	H14F	C13F	*C14F	H14G	1.1131	109.06	-117.24	109.28	1.1130
IC	C13F	C14F	C15F	C16F	1.5345	112.68	-178.23	113.16	1.5306
IC	C16F	C14F	*C15F	H15F	1.5306	113.16	-121.86	108.84	1.1140
IC	H15F	C14F	*C15F	H15G	1.1140	108.84	-116.70	108.76	1.1144
IC	C14F	C15F	C16F	C17F	1.5345	112.68	-178.23	113.16	1.5306
IC	C17F	C15F	*C16F	H16F	1.5306	113.16	-121.86	108.84	1.1140
IC	H16F	C15F	*C16F	H16G	1.1140	108.84	-116.70	108.76	1.1144
IC	C15F	C16F	C17F	C18F	1.5345	112.68	-178.23	113.16	1.5306
IC	C18F	C16F	*C17F	H17F	1.5306	113.16	-121.86	108.84	1.1140
IC	H17F	C16F	*C17F	H17G	1.1140	108.84	-116.70	108.76	1.1144
IC	C16F	C17F	C18F	C19F	1.5345	112.68	-178.23	113.16	1.5306
IC	C19F	C17F	*C18F	H18F	1.5306	113.16	-121.86	108.84	1.1140
IC	H18F	C17F	*C18F	H18G	1.1140	108.84	-116.70	108.76	1.1144
IC	C17F	C18F	C19F	C20F	1.5344	112.57	179.80	112.59	1.5346
IC	C20F	C18F	*C19F	H19F	1.5346	112.59	-121.28	109.10	1.1133
IC	H19F	C18F	*C19F	H19G	1.1133	109.10	-117.38	109.12	1.1133
IC	C18F	C19F	C20F	C21F	1.5345	112.59	-179.89	112.63	1.5337
IC	C21F	C19F	*C20F	H20F	1.5337	112.63	-121.33	109.16	1.1133
IC	H20F	C19F	*C20F	H20G	1.1133	109.16	-117.38	109.14	1.1133
IC	C19F	C20F	C21F	C22F	1.5346	112.63	179.93	113.24	1.5309
IC	C22F	C20F	*C21F	H21F	1.5309	113.24	-121.66	108.77	1.1142
IC	H21F	C20F	*C21F	H21G	1.1142	108.77	-116.65	108.78	1.1142
IC	C20F	C21F	C22F	C23F	1.5346	112.63	179.93	113.24	1.5309
IC	C23F	C21F	*C22F	H22F	1.5309	113.24	-121.66	108.77	1.1142
IC	H22F	C21F	*C22F	H22G	1.1142	108.77	-116.65	108.78	1.1142
IC	C21F	C22F	C23F	C24F	1.5346	112.63	179.93	113.24	1.5309
IC	C24F	C22F	*C23F	H23F	1.5309	113.24	-121.66	108.77	1.1142
IC	H23F	C22F	*C23F	H23G	1.1142	108.77	-116.65	108.78	1.1142
IC	C22F	C23F	C24F	H24F	1.5337	113.24	-59.91	110.47	1.1114
IC	H24F	C23F	*C24F	H24G	1.1114	110.47	119.87	110.48	1.1113
IC	H24F	C23F	*C24F	H24H	1.1114	110.47	-120.06	110.63	1.1113

PATCH FIRST NONE LAST NONE

RESI NSM 0.00 ! sphingomyelin d18:1/24:1 (w/nervonic acid)

GROUP !

ATOM N	NTL	-0.60 !	H15B			
ATOM C15	CTL5	-0.35 !	H15A-C15-H15C			
ATOM H15A	HL	0.25 !	H13C		H14C	
ATOM H15B	HL	0.25 !				
ATOM H15C	HL	0.25 !				
ATOM C14	CTL5	-0.35 !	H13A-C13---N--C14-H14A	(+)		
ATOM H14A	HL	0.25 !				
ATOM H14B	HL	0.25 !				
ATOM H14C	HL	0.25 !	H13B		H14B	
ATOM C13	CTL5	-0.35 !				
ATOM H13A	HL	0.25 !				

ATOM H13B HL 0.25 !
 ATOM H13C HL 0.25 !
 ATOM C12 CTL2 -0.10 !
 ATOM H12A HL 0.25 !
 ATOM H12B HL 0.25 !
 GROUP !
 ATOM C11 CTL2 -0.08 !
 ATOM H11A HAL2 0.09 !
 ATOM H11B HAL2 0.09 !
 ATOM P PL 1.50 !
 ATOM O13 O2L -0.78 !
 ATOM O14 O2L -0.78 !
 ATOM O11 OSLP -0.57 !
 ATOM O12 OSLP -0.57 !
 ATOM C1S CTL2 -0.08 !
 ATOM H1S HAL2 0.09 !
 ATOM H1T HAL2 0.09 !
 GROUP !
 ATOM NF NHL -0.70 !
 ATOM HNF H 0.35 !
 ATOM C2S CTL1 0.30 !
 ATOM H2S HAL1 0.05 !
 GROUP !
 ATOM C3S CTO1 0.50 !
 ATOM H3S HAL1 -0.11 !
 ATOM O3 OHL -0.69 !
 ATOM HO3 HOL 0.30 !
 GROUP !
 ATOM C4S CEL3 -0.15 !
 ATOM H4S HEL1 0.15 !
 GROUP !
 ATOM C5S CEL3 -0.15 !
 ATOM H5S HEL1 0.15 !
 GROUP !
 ATOM C6S CTL2 -0.18 !
 ATOM H6S HAL2 0.09 !
 ATOM H6T HAL2 0.09 !
 GROUP !
 ATOM C7S CTL2 -0.18 !
 ATOM H7S HAL2 0.09 !
 ATOM H7T HAL2 0.09 !
 GROUP !
 ATOM C8S CTL2 -0.18 !
 ATOM H8S HAL2 0.09 !
 ATOM H8T HAL2 0.09 !
 GROUP !
 ATOM C9S CTL2 -0.18 !
 ATOM H9S HAL2 0.09 !
 ATOM H9T HAL2 0.09 !
 GROUP !
 ATOM C10S CTL2 -0.18 !
 ATOM H10S HAL2 0.09 !
 ATOM H10T HAL2 0.09 !
 GROUP
 ATOM C11S CTL2 -0.18 !
 ATOM H11S HAL2 0.09 !
 ATOM H11T HAL2 0.09 !
 GROUP
 ATOM C12S CTL2 -0.18 !
 ATOM H12S HAL2 0.09 !
 ATOM H12T HAL2 0.09 !
 GROUP
 ATOM C13S CTL2 -0.18 !
 ATOM H13S HAL2 0.09 !
 ATOM H13T HAL2 0.09 !
 GROUP
 ATOM C14S CTL2 -0.18 !
 ATOM H14S HAL2 0.09 !
 ATOM H14T HAL2 0.09 !
 GROUP
 ATOM C15S CTL2 -0.18 !

ATOM H15S HAL2 0.09 !
 ATOM H15T HAL2 0.09 !
 GROUP
 ATOM C16S CTL2 -0.18 !
 ATOM H16S HAL2 0.09 !
 ATOM H16T HAL2 0.09 !
 GROUP
 ATOM C17S CTL2 -0.18 !
 ATOM H17S HAL2 0.09 !
 ATOM H17T HAL2 0.09 !
 GROUP
 ATOM C18S CTL3 -0.27 !
 ATOM H18S HAL3 0.09 !
 ATOM H18T HAL3 0.09 !
 ATOM H18U HAL3 0.09 !
 GROUP !FATTY ACID CHAIN
 ATOM C1F C 0.55 !
 ATOM O F -0.60 !
 ATOM C2F CTL2 -0.07 !
 ATOM H2F HAL2 0.06 !
 ATOM H2G HAL2 0.06 !
 GROUP !
 ATOM C3F CTL2 -0.18 !
 ATOM H3F HAL2 0.09 !
 ATOM H3G HAL2 0.09 !
 GROUP !
 ATOM C4F CTL2 -0.18 !
 ATOM H4F HAL2 0.09 !
 ATOM H4G HAL2 0.09 !
 GROUP !
 ATOM C5F CTL2 -0.18 !
 ATOM H5F HAL2 0.09 !
 ATOM H5G HAL2 0.09 !
 GROUP !
 ATOM C6F CTL2 -0.18 !
 ATOM H6F HAL2 0.09 !
 ATOM H6G HAL2 0.09 !
 GROUP !
 ATOM C7F CTL2 -0.18 !
 ATOM H7F HAL2 0.09 !
 ATOM H7G HAL2 0.09 !
 GROUP !
 ATOM C8F CTL2 -0.18 !
 ATOM H8F HAL2 0.09 !
 ATOM H8G HAL2 0.09 !
 GROUP !
 ATOM C9F CTL2 -0.18 !
 ATOM H9F HAL2 0.09 !
 ATOM H9G HAL2 0.09 !
 GROUP !
 ATOM C10F CTL2 -0.18 !
 ATOM H10F HAL2 0.09 !
 ATOM H10G HAL2 0.09 !
 GROUP !
 ATOM C11F CTL2 -0.18 !
 ATOM H11F HAL2 0.09 !
 ATOM H11G HAL2 0.09 !
 GROUP !
 ATOM C12F CTL2 -0.18 !
 ATOM H12F HAL2 0.09 !
 ATOM H12G HAL2 0.09 !
 GROUP !
 ATOM C13F CTL2 -0.18 !
 ATOM H13F HAL2 0.09 !
 ATOM H13G HAL2 0.09 !
 GROUP !
 ATOM C14F CTL2 -0.18 !
 ATOM H14F HAL2 0.09 !
 ATOM H14G HAL2 0.09 !
 GROUP !
 ATOM C15F CEL1 -0.15 !

ATOM H15F HEL1 0.15 !
 GROUP !
 ATOM C16F CEL1 -0.15 !
 ATOM H16F HEL1 0.15 !
 GROUP !
 ATOM C17F CTL2 -0.18 !
 ATOM H17F HAL2 0.09 !
 ATOM H17G HAL2 0.09 !
 GROUP !
 ATOM C18F CTL2 -0.18 !
 ATOM H18F HAL2 0.09 !
 ATOM H18G HAL2 0.09 !
 GROUP !
 ATOM C19F CTL2 -0.18 !
 ATOM H19F HAL2 0.09 !
 ATOM H19G HAL2 0.09 !
 GROUP !
 ATOM C20F CTL2 -0.18 !
 ATOM H20F HAL2 0.09 !
 ATOM H20G HAL2 0.09 !
 GROUP !
 ATOM C21F CTL2 -0.18 !
 ATOM H21F HAL2 0.09 !
 ATOM H21G HAL2 0.09 !
 GROUP !
 ATOM C22F CTL2 -0.18 !
 ATOM H22F HAL2 0.09 !
 ATOM H22G HAL2 0.09 !
 GROUP !
 ATOM C23F CTL2 -0.18 !
 ATOM H23F HAL2 0.09 !
 ATOM H23G HAL2 0.09 !
 GROUP !
 ATOM C24F CTL3 -0.27 !
 ATOM H24F HAL3 0.09 !
 ATOM H24G HAL3 0.09 !
 ATOM H24H HAL3 0.09 !

!SPHINGOSINE CHAIN

BOND C18S H18S C18S H18T C18S H18U
 BOND C18S C17S C17S H17S C17S H17T
 BOND C17S C16S C16S H16S C16S H16T
 BOND C16S C15S C15S H15S C15S H15T
 BOND C15S C14S C14S H14S C14S H14T
 BOND C14S C13S C13S H13S C13S H13T
 BOND C13S C12S C12S H12S C12S H12T
 BOND C12S C11S C11S H11S C11S H11T
 BOND C11S C10S C10S H10S C10S H10T
 BOND C10S C9S C9S H9S C9S H9T
 BOND C9S C8S C8S H8S C8S H8T
 BOND C8S C7S C7S H7S C7S H7T
 BOND C7S C6S C6S H6S C6S H6T
 BOND C6S C5S C5S H5S C4S H4S
 BOND C4S C3S C3S H3S C3S O3
 BOND O3 HO3
 BOND C3S C2S C2S H2S
 BOND C2S C1S C1S H1S C1S H1T
 BOND C1S O11

!PHOSPHOCHOLINE

BOND O11 P O12 P O13 P O14 P
 BOND O12 C11 C11 H11A C11 H11B
 BOND C11 C12 C12 H12A C12 H12B
 BOND C12 N N C13 N C14 N C15
 BOND C13 H13A C13 H13B C13 H13C
 BOND C14 H14A C14 H14B C14 H14C
 BOND C15 H15A C15 H15B C15 H15C

FATTY ACID CHAIN

BOND C2S NF NF HNF NF C1F
 BOND C1F C2F C2F H2F C2F H2G
 BOND C2F C3F C3F H3F C3F H3G
 BOND C3F C4F C4F H4F C4F H4G

BOND C4F C5F C5F H5F C5F H5G
 BOND C5F C6F C6F H6F C6F H6G
 BOND C6F C7F C7F H7F C7F H7G
 BOND C7F C8F C8F H8F C8F H8G
 BOND C8F C9F C9F H9F C9F H9G
 BOND C9F C10F C10F H10F C10F H10G
 BOND C10F C11F C11F H11F C11F H11G
 BOND C11F C12F C12F H12F C12F H12G
 BOND C12F C13F C13F H13F C13F H13G
 BOND C13F C14F C14F H14F C14F H14G
 BOND C14F C15F C15F H15F
 BOND C16F H16F
 BOND C16F C17F C17F H17F C17F H17G
 BOND C17F C18F C18F H18F C18F H18G
 BOND C18F C19F C19F H19F C19F H19G
 BOND C19F C20F C20F H20F C20F H20G
 BOND C20F C21F C21F H21F C21F H21G
 BOND C21F C22F C22F H22F C22F H22G
 BOND C22F C23F C23F H23F C23F H23G
 BOND C23F C24F C24F H24F C24F H24G
 BOND C24F H24H

DOUBLE C4S C5S
 DOUBLE C15F C16F
 DOUBLE C1F OF
 IMPR C1F C2F NF OF NF C1F C2S HNF

IC	C18S	C17S	C16S	C15S	1.5309	113.23	180.00	112.64	1.5346
IC	C17S	C16S	C15S	C14S	1.5337	112.64	180.00	112.58	1.5345
IC	C16S	C15S	C14S	C13S	1.5346	112.58	180.00	112.58	1.5345
IC	C15S	C14S	C13S	C12S	1.5345	112.58	-179.99	112.58	1.5345
IC	C14S	C13S	C12S	C11S	1.5345	112.58	179.97	112.58	1.5344
IC	C13S	C12S	C11S	C10S	1.5345	112.58	-179.96	112.61	1.5345
IC	C12S	C11S	C10S	C9S	1.5344	112.61	179.97	112.54	1.5344
IC	C11S	C10S	C9S	C8S	1.5345	112.54	-179.87	112.66	1.5345
IC	C10S	C9S	C8S	C7S	1.5344	112.66	-179.89	112.50	1.5354
IC	C9S	C8S	C7S	C6S	1.5345	112.50	-179.79	112.41	1.5391
IC	C8S	C7S	C6S	C5S	1.5354	112.41	-179.61	111.46	1.5077
IC	C7S	C6S	C5S	C4S	1.5391	111.46	-117.13	124.77	1.3444
IC	C6S	C5S	C4S	C3S	1.5077	124.77	179.01	124.28	1.5009
IC	C5S	C4S	C3S	C2S	1.3444	124.28	-84.24	110.14	1.5343
IC	C5S	C4S	C3S	O3	1.3444	124.28	156.07	107.64	1.4235
IC	C4S	C3S	C2S	C1S	1.5009	110.14	-70.50	114.25	1.5586
IC	C13	C12	*N	C14	1.4967	109.68	116.91	107.30	1.5040
IC	C13	C12	*N	C15	1.4967	109.68	-123.42	110.88	1.4938
IC	C12	N	C15	H15A	1.5212	110.88	-177.15	111.31	1.0816
IC	H15A	N	*C15	H15B	1.0816	111.31	118.98	109.44	1.0875
IC	H15A	N	*C15	H15C	1.0816	111.31	-122.11	111.46	1.0818
IC	C12	N	C14	H14A	1.5212	107.30	-178.89	114.23	1.0739
IC	H14A	N	*C14	H14B	1.0739	114.23	123.44	110.21	1.0968
IC	H14A	N	*C14	H14C	1.0739	114.23	-124.43	111.41	1.0948
IC	C12	N	C13	H13A	1.5212	109.68	176.44	111.15	1.0822
IC	H13A	N	*C13	H13B	1.0822	111.15	121.89	111.75	1.0814
IC	H13A	N	*C13	H13C	1.0822	111.15	-118.50	109.57	1.0892
IC	C13	N	C12	C11	1.4967	109.68	-178.83	116.30	1.5392
IC	C11	N	*C12	H12A	1.5392	116.30	117.04	108.30	1.0985
IC	C11	N	*C12	H12B	1.5392	116.30	-126.61	112.36	1.0787
IC	N	C12	C11	O12	1.5212	116.30	126.79	107.85	1.4239
IC	O12	C12	*C11	H11A	1.4239	107.85	118.10	109.30	1.1131
IC	O12	C12	*C11	H11B	1.4239	107.85	-123.12	113.16	1.1140
IC	C12	C11	O12	P	1.5392	107.85	-68.29	118.09	1.5886
IC	C11	O12	P	O11	1.4239	118.09	-167.58	104.18	1.5725
IC	O11	O12	*P	O13	1.5725	104.18	116.57	107.05	1.4823
IC	O11	O12	*P	O14	1.5725	104.18	-118.73	106.98	1.4805
IC	O12	P	O11	C1S	1.5886	104.18	51.85	121.69	1.4327
IC	P	O11	C1S	C2S	1.5725	121.69	103.66	111.21	1.5586
IC	C2S	O11	*C1S	H1S	1.5586	111.21	118.79	108.35	1.1121
IC	C2S	O11	*C1S	H1T	1.5586	111.21	-123.25	112.85	1.1152
IC	O11	C1S	C2S	C3S	1.4327	111.21	175.67	114.25	1.5343
IC	C3S	C1S	*C2S	NF	1.5343	114.25	120.00	111.52	1.4406
IC	C3S	C1S	*C2S	H2S	1.5343	114.25	-120.00	105.03	1.1194

IC	C1S	C2S	NF	C1F	1.5586	111.52	160.94	124.85	1.3364
IC	C1F	C2S	*NF	HNF	1.3364	124.85	-157.81	113.39	1.0062
IC	C1S	C2S	C3S	C4S	1.5586	114.25	-70.50	110.14	1.5009
IC	C4S	C2S	*C3S	O3	1.5009	110.14	118.36	109.80	1.4235
IC	C4S	C2S	*C3S	H3S	1.5009	110.14	-121.14	110.10	1.1140
IC	C2S	C3S	O3	HO3	1.5343	109.80	46.50	104.54	0.9610
IC	C2S	C3S	C4S	C5S	1.5343	110.14	-84.24	124.28	1.3444
IC	C5S	C3S	*C4S	H4S	1.3444	124.28	178.79	115.93	1.1003
IC	C3S	C4S	C5S	C6S	1.5009	124.28	179.01	124.77	1.5077
IC	C6S	C4S	*C5S	H5S	1.5077	124.77	-178.18	119.00	1.1029
IC	C4S	C5S	C6S	C7S	1.3444	124.77	-117.13	111.46	1.5391
IC	C7S	C5S	*C6S	H6S	1.5391	111.46	120.98	111.41	1.1142
IC	C7S	C5S	*C6S	H6T	1.5391	111.46	-120.60	109.91	1.1125
IC	C5S	C6S	C7S	C8S	1.5077	111.46	-179.61	112.41	1.5354
IC	C8S	C6S	*C7S	H7S	1.5354	112.41	121.26	109.74	1.1138
IC	C8S	C6S	*C7S	H7T	1.5354	112.41	-120.99	109.40	1.1147
IC	C6S	C7S	C8S	C9S	1.5391	112.41	-179.79	112.50	1.5345
IC	C9S	C7S	*C8S	H8S	1.5345	112.50	121.20	108.97	1.1132
IC	C9S	C7S	*C8S	H8T	1.5345	112.50	-121.38	109.07	1.1129
IC	C7S	C8S	C9S	C10S	1.5354	112.50	-179.89	112.66	1.5344
IC	C10S	C8S	*C9S	H9S	1.5344	112.66	121.49	109.05	1.1133
IC	C10S	C8S	*C9S	H9T	1.5344	112.66	-121.38	108.96	1.1136
IC	C8S	C9S	C10S	C11S	1.5345	112.66	-179.87	112.54	1.5345
IC	C11S	C9S	*C10S	H10S	1.5345	112.54	121.27	109.07	1.1133
IC	C11S	C9S	*C10S	H10T	1.5345	112.54	-121.34	109.11	1.1132
IC	C9S	C10S	C11S	C12S	1.5344	112.54	179.97	112.61	1.5344
IC	C12S	C10S	*C11S	H11S	1.5344	112.61	121.36	109.09	1.1133
IC	C12S	C10S	*C11S	H11T	1.5344	112.61	-121.33	109.07	1.1134
IC	C10S	C11S	C12S	C13S	1.5345	112.61	-179.96	112.58	1.5345
IC	C13S	C11S	*C12S	H12S	1.5345	112.58	121.29	109.12	1.1133
IC	C13S	C11S	*C12S	H12T	1.5345	112.58	-121.31	109.15	1.1133
IC	C11S	C12S	C13S	C14S	1.5344	112.58	179.97	112.58	1.5345
IC	C14S	C12S	*C13S	H13S	1.5345	112.58	121.30	109.13	1.1133
IC	C14S	C12S	*C13S	H13T	1.5345	112.58	-121.30	109.13	1.1133
IC	C12S	C13S	C14S	C15S	1.5345	112.58	-179.99	112.58	1.5345
IC	C15S	C13S	*C14S	H14S	1.5345	112.58	121.30	109.12	1.1133
IC	C15S	C13S	*C14S	H14T	1.5345	112.58	-121.31	109.12	1.1133
IC	C13S	C14S	C15S	C16S	1.5345	112.58	180.00	112.58	1.5346
IC	C16S	C14S	*C15S	H15S	1.5346	112.58	121.30	109.11	1.1133
IC	C16S	C14S	*C15S	H15T	1.5346	112.58	-121.30	109.11	1.1133
IC	C14S	C15S	C16S	C17S	1.5345	112.58	180.00	112.64	1.5337
IC	C17S	C15S	*C16S	H16S	1.5337	112.64	121.32	109.14	1.1133
IC	C17S	C15S	*C16S	H16T	1.5337	112.64	-121.32	109.14	1.1133
IC	C15S	C16S	C17S	C18S	1.5346	112.64	180.00	113.23	1.5309
IC	C18S	C16S	*C17S	H17S	1.5309	113.23	121.67	108.77	1.1142
IC	C18S	C16S	*C17S	H17T	1.5309	113.23	-121.67	108.77	1.1142
IC	C16S	C17S	C18S	H18S	1.5337	113.23	180.00	110.63	1.1113
IC	H18S	C17S	*C18S	H18T	1.1113	110.63	120.07	110.47	1.1114
IC	H18S	C17S	*C18S	H18U	1.1113	110.63	-120.07	110.47	1.1114
IC	C2S	NF	C1F	C2F	1.4406	124.85	-156.83	114.97	1.4964
IC	C2F	NF	*C1F	OF	1.4964	114.97	162.54	122.11	1.2268
IC	NF	C1F	C2F	C3F	1.3364	114.97	-146.57	111.92	1.5432
IC	C3F	C1F	*C2F	H2F	1.5432	111.92	-120.83	107.93	1.1115
IC	H2F	C1F	*C2F	H2G	1.1115	107.93	-117.30	109.04	1.1113
IC	C1F	C2F	C3F	C4F	1.4964	111.92	179.54	112.84	1.5310
IC	C4F	C2F	*C3F	H3F	1.5310	112.84	-121.63	109.37	1.1145
IC	H3F	C2F	*C3F	H3G	1.1145	109.37	-117.07	109.08	1.1157
IC	C2F	C3F	C4F	C5F	1.5434	112.23	174.86	112.58	1.5342
IC	C5F	C3F	*C4F	H4F	1.5342	112.58	-120.18	108.65	1.1140
IC	H4F	C3F	*C4F	H4G	1.1140	108.65	-117.52	109.62	1.1127
IC	C3F	C4F	C5F	C6F	1.5340	112.58	-174.06	112.39	1.5340
IC	C6F	C4F	*C5F	H5F	1.5340	112.39	-122.01	109.38	1.1130
IC	H5F	C4F	*C5F	H5G	1.1130	109.38	-117.34	108.92	1.1134
IC	C4F	C5F	C6F	C7F	1.5342	112.39	178.27	112.75	1.5340
IC	C7F	C5F	*C6F	H6F	1.5340	112.75	-120.93	108.96	1.1137
IC	H6F	C5F	*C6F	H6G	1.1137	108.96	-117.41	109.22	1.1131
IC	C5F	C6F	C7F	C8F	1.5340	112.75	-178.89	112.33	1.5342
IC	C8F	C6F	*C7F	H7F	1.5342	112.33	-121.40	109.26	1.1132
IC	H7F	C6F	*C7F	H7G	1.1132	109.26	-117.45	109.12	1.1130
IC	C6F	C7F	C8F	C9F	1.5340	112.33	179.09	112.77	1.5340
IC	C9F	C7F	*C8F	H8F	1.5340	112.77	-121.23	109.16	1.1134

IC	H8F	C7F	*C8F	H8G	1.1134	109.16	-117.38	109.08	1.1133
IC	C7F	C8F	C9F	C10F	1.5342	112.77	179.02	112.34	1.5343
IC	C10F	C8F	*C9F	H9F	1.5343	112.34	-121.27	109.27	1.1132
IC	H9F	C8F	*C9F	H9G	1.1132	109.27	-117.45	109.10	1.1130
IC	C8F	C9F	C10F	C11F	1.5340	112.34	178.16	112.74	1.5340
IC	C11F	C9F	*C10F	H10F	1.5340	112.74	-121.18	109.16	1.1134
IC	H10F	C9F	*C10F	H10G	1.1134	109.16	-117.35	109.10	1.1132
IC	C9F	C10F	C11F	C12F	1.5343	112.74	178.91	112.38	1.5343
IC	C12F	C10F	*C11F	H11F	1.5343	112.38	-121.39	109.34	1.1131
IC	H11F	C10F	*C11F	H11G	1.1131	109.34	-117.43	108.98	1.1132
IC	C10F	C11F	C12F	C13F	1.5340	112.38	176.93	112.68	1.5340
IC	C13F	C11F	*C12F	H12F	1.5340	112.68	-121.02	109.05	1.1134
IC	H12F	C11F	*C12F	H12G	1.1134	109.05	-117.34	109.20	1.1131
IC	C11F	C12F	C13F	C14F	1.5343	112.68	-179.93	112.45	1.5345
IC	C14F	C12F	*C13F	H13F	1.5345	112.45	-121.60	109.42	1.1129
IC	H13F	C12F	*C13F	H13G	1.1129	109.42	-117.40	108.85	1.1134
IC	C12F	C13F	C14F	C15F	1.5340	112.45	175.93	112.68	1.5334
IC	C15F	C13F	*C14F	H14F	1.5334	112.68	-121.00	109.06	1.1131
IC	H14F	C13F	*C14F	H14G	1.1131	109.06	-117.24	109.28	1.1130
IC	C13F	C14F	C15F	C16F	1.5345	112.68	-178.23	113.16	1.5306
IC	C16F	C14F	*C15F	H15F	1.5306	113.16	-121.86	108.84	1.1140
IC	C14F	C15F	C16F	C17F	1.5345	112.68	0.00	113.16	1.5306
IC	C17F	C15F	*C16F	H16F	1.5306	113.16	-121.86	108.84	1.1140
IC	C15F	C16F	C17F	C18F	1.5345	112.68	-178.23	113.16	1.5306
IC	C18F	C16F	*C17F	H17F	1.5306	113.16	-121.86	108.84	1.1140
IC	H17F	C16F	*C17F	H17G	1.1140	108.84	-116.70	108.76	1.1144
IC	C16F	C17F	C18F	C19F	1.5345	112.68	0.00	113.16	1.5306
IC	C19F	C17F	*C18F	H18F	1.5306	113.16	-121.86	108.84	1.1140
IC	H18F	C17F	*C18F	H18G	1.1140	108.84	-116.70	108.76	1.1144
IC	C17F	C18F	C19F	C20F	1.5344	112.57	180.00	112.59	1.5346
IC	C20F	C18F	*C19F	H19F	1.5346	112.59	-121.28	109.10	1.1133
IC	H19F	C18F	*C19F	H19G	1.1133	109.10	-117.38	109.12	1.1133
IC	C18F	C19F	C20F	C21F	1.5345	112.59	-179.89	112.63	1.5337
IC	C21F	C19F	*C20F	H20F	1.5337	112.63	-121.33	109.16	1.1133
IC	H20F	C19F	*C20F	H20G	1.1133	109.16	-117.38	109.14	1.1133
IC	C19F	C20F	C21F	C22F	1.5346	112.63	179.93	113.24	1.5309
IC	C22F	C20F	*C21F	H21F	1.5309	113.24	-121.66	108.77	1.1142
IC	H21F	C20F	*C21F	H21G	1.1142	108.77	-116.65	108.78	1.1142
IC	C20F	C21F	C22F	C23F	1.5346	112.63	179.93	113.24	1.5309
IC	C23F	C21F	*C22F	H22F	1.5309	113.24	-121.66	108.77	1.1142
IC	H22F	C21F	*C22F	H22G	1.1142	108.77	-116.65	108.78	1.1142
IC	C21F	C22F	C23F	C24F	1.5346	112.63	179.93	113.24	1.5309
IC	C24F	C22F	*C23F	H23F	1.5309	113.24	-121.66	108.77	1.1142
IC	H23F	C22F	*C23F	H23G	1.1142	108.77	-116.65	108.78	1.1142
IC	C22F	C23F	C24F	H24F	1.5337	113.24	-59.91	110.47	1.1114
IC	H24F	C23F	*C24F	H24G	1.1114	110.47	119.87	110.48	1.1113
IC	H24F	C23F	*C24F	H24H	1.1114	110.47	-120.06	110.63	1.1113

PATCH FIRST NONE LAST NONE

RESI CER2 0.00 ! CERAMIDE by by Sam Tonddast-Navaei (U. Cin/Russel Lab)

GROUP ! HO1

ATOM O1 OHL -0.69 ! \

ATOM HO1 HOL 0.30 ! O1

ATOM C1S CTO2 0.49 ! H1S---C1S---H1T

ATOM H1S HAL2 -0.05 !

ATOM H1T HAL2 -0.05 !

GROUP ! H2S---C2S-----NF--HNF

ATOM NF NHL -0.70 !

ATOM HNF H 0.35 !

ATOM C2S CTL1 0.30 !

ATOM H2S HAL1 0.05 !

GROUP !

ATOM C3S CTO1 0.50 ! C1F=OF (FATTY ACID CHAIN)

ATOM H3S HAL1 -0.11 !

ATOM O3 OHL -0.69 !

ATOM HO3 HOL 0.30 !

GROUP !

ATOM C4S CEL3 -0.15 ! [(CH2)3F (C3F,C4F...C23F)

ATOM H4S HEL1 0.15 !

GROUP ! H24F---C24F---H24G

ATOM C5S CEL3 -0.15 !
 ATOM H5S HEL1 0.15 !
 GROUP !
 ATOM C6S CTL2 -0.18 !
 ATOM H6S HAL2 0.09 !
 ATOM H6T HAL2 0.09 !
 GROUP !
 ATOM C7S CTL2 -0.18 !
 ATOM H7S HAL2 0.09 !
 ATOM H7T HAL2 0.09 !
 GROUP !
 ATOM C8S CTL2 -0.18 !
 ATOM H8S HAL2 0.09 !
 ATOM H8T HAL2 0.09 !
 GROUP !
 ATOM C9S CTL2 -0.18 !
 ATOM H9S HAL2 0.09 !
 ATOM H9T HAL2 0.09 !
 GROUP !
 ATOM C10S CTL2 -0.18 !
 ATOM H10S HAL2 0.09 !
 ATOM H10T HAL2 0.09 !
 GROUP
 ATOM C11S CTL2 -0.18 !
 ATOM H11S HAL2 0.09 !
 ATOM H11T HAL2 0.09 !
 GROUP
 ATOM C12S CTL2 -0.18 !
 ATOM H12S HAL2 0.09 !
 ATOM H12T HAL2 0.09 !
 GROUP
 ATOM C13S CTL2 -0.18 !
 ATOM H13S HAL2 0.09 !
 ATOM H13T HAL2 0.09 !
 GROUP
 ATOM C14S CTL2 -0.18 !
 ATOM H14S HAL2 0.09 !
 ATOM H14T HAL2 0.09 !
 GROUP
 ATOM C15S CTL2 -0.18 !
 ATOM H15S HAL2 0.09 !
 ATOM H15T HAL2 0.09 !
 GROUP
 ATOM C16S CTL2 -0.18 !
 ATOM H16S HAL2 0.09 !
 ATOM H16T HAL2 0.09 !
 GROUP
 ATOM C17S CTL2 -0.18 !
 ATOM H17S HAL2 0.09 !
 ATOM H17T HAL2 0.09 !
 GROUP !
 ATOM C18S CTL3 -0.27 !
 ATOM H18S HAL3 0.09 !
 ATOM H18T HAL3 0.09 !
 ATOM H18U HAL3 0.09 !
 GROUP !FATTY ACID CHAIN
 ATOM C1F C 0.55 !
 ATOM OF O -0.60 !
 ATOM C2F CTL2 -0.07 !
 ATOM H2F HAL2 0.06 !
 ATOM H2G HAL2 0.06 !
 GROUP !
 ATOM C3F CTL2 -0.18 !
 ATOM H3F HAL2 0.09 !
 ATOM H3G HAL2 0.09 !
 GROUP !
 ATOM C4F CTL2 -0.18 !
 ATOM H4F HAL2 0.09 !
 ATOM H4G HAL2 0.09 !
 GROUP !
 ATOM C5F CTL2 -0.18 !

ATOM H5F HAL2 0.09 !
 ATOM H5G HAL2 0.09 !
 GROUP !
 ATOM C6F CTL2 -0.18 !
 ATOM H6F HAL2 0.09 !
 ATOM H6G HAL2 0.09 !
 GROUP !
 ATOM C7F CTL2 -0.18 !
 ATOM H7F HAL2 0.09 !
 ATOM H7G HAL2 0.09 !
 GROUP !
 ATOM C8F CTL2 -0.18 !
 ATOM H8F HAL2 0.09 !
 ATOM H8G HAL2 0.09 !
 GROUP !
 ATOM C9F CTL2 -0.18 !
 ATOM H9F HAL2 0.09 !
 ATOM H9G HAL2 0.09 !
 GROUP !
 ATOM C10F CTL2 -0.18 !
 ATOM H10F HAL2 0.09 !
 ATOM H10G HAL2 0.09 !
 GROUP !
 ATOM C11F CTL2 -0.18 !
 ATOM H11F HAL2 0.09 !
 ATOM H11G HAL2 0.09 !
 GROUP !
 ATOM C12F CTL2 -0.18 !
 ATOM H12F HAL2 0.09 !
 ATOM H12G HAL2 0.09 !
 GROUP !
 ATOM C13F CTL2 -0.18 !
 ATOM H13F HAL2 0.09 !
 ATOM H13G HAL2 0.09 !
 GROUP !
 ATOM C14F CTL2 -0.18 !
 ATOM H14F HAL2 0.09 !
 ATOM H14G HAL2 0.09 !
 GROUP !
 ATOM C15F CTL2 -0.18 !
 ATOM H15F HAL2 0.09 !
 ATOM H15G HAL2 0.09 !
 GROUP !
 ATOM C16F CTL2 -0.18 !
 ATOM H16F HAL2 0.09 !
 ATOM H16G HAL2 0.09 !
 GROUP !
 ATOM C17F CTL2 -0.18 !
 ATOM H17F HAL2 0.09 !
 ATOM H17G HAL2 0.09 !
 GROUP !
 ATOM C18F CTL2 -0.18 !
 ATOM H18F HAL2 0.09 !
 ATOM H18G HAL2 0.09 !
 GROUP !
 ATOM C19F CTL2 -0.18 !
 ATOM H19F HAL2 0.09 !
 ATOM H19G HAL2 0.09 !
 GROUP !
 ATOM C20F CTL2 -0.18 !
 ATOM H20F HAL2 0.09 !
 ATOM H20G HAL2 0.09 !
 GROUP !
 ATOM C21F CTL2 -0.18 !
 ATOM H21F HAL2 0.09 !
 ATOM H21G HAL2 0.09 !
 GROUP !
 ATOM C22F CTL2 -0.18 !
 ATOM H22F HAL2 0.09 !
 ATOM H22G HAL2 0.09 !
 GROUP !

ATOM C23F CTL2 -0.18 !
 ATOM H23F HAL2 0.09 !
 ATOM H23G HAL2 0.09 !
 GROUP !
 ATOM C24F CTL3 -0.27 !
 ATOM H24F HAL3 0.09 !
 ATOM H24G HAL3 0.09 !
 ATOM H24H HAL3 0.09 !

!SPHINGOSINE CHAIN

BOND C18S H18S C18S H18T C18S H18U
 BOND C18S C17S C17S H17S C17S H17T
 BOND C17S C16S C16S H16S C16S H16T
 BOND C16S C15S C15S H15S C15S H15T
 BOND C15S C14S C14S H14S C14S H14T
 BOND C14S C13S C13S H13S C13S H13T
 BOND C13S C12S C12S H12S C12S H12T
 BOND C12S C11S C11S H11S C11S H11T
 BOND C11S C10S C10S H10S C10S H10T
 BOND C10S C9S C9S H9S C9S H9T
 BOND C9S C8S C8S H8S C8S H8T
 BOND C8S C7S C7S H7S C7S H7T
 BOND C7S C6S C6S H6S C6S H6T
 BOND C6S C5S C5S H5S C4S H4S
 BOND C4S C3S C3S H3S C3S O3
 BOND O3 HO3
 BOND C3S C2S C2S H2S
 BOND C2S C1S C1S H1S C1S H1T
 BOND C1S O1 O1 HO1

!FATTY ACID CHAIN

BOND C2S NF NF HNF NF C1F
 BOND C1F C2F C2F H2F C2F H2G
 BOND C2F C3F C3F H3F C3F H3G
 BOND C3F C4F C4F H4F C4F H4G
 BOND C4F C5F C5F H5F C5F H5G
 BOND C5F C6F C6F H6F C6F H6G
 BOND C6F C7F C7F H7F C7F H7G
 BOND C7F C8F C8F H8F C8F H8G
 BOND C8F C9F C9F H9F C9F H9G
 BOND C9F C10F C10F H10F C10F H10G
 BOND C10F C11F C11F H11F C11F H11G
 BOND C11F C12F C12F H12F C12F H12G
 BOND C12F C13F C13F H13F C13F H13G
 BOND C13F C14F C14F H14F C14F H14G
 BOND C14F C15F C15F H15F C15F H15G
 BOND C15F C16F C16F H16F C16F H16G
 BOND C16F C17F C17F H17F C17F H17G
 BOND C17F C18F C18F H18F C18F H18G
 BOND C18F C19F C19F H19F C19F H19G
 BOND C19F C20F C20F H20F C20F H20G
 BOND C20F C21F C21F H21F C21F H21G
 BOND C21F C22F C22F H22F C22F H22G
 BOND C22F C23F C23F H23F C23F H23G
 BOND C23F C24F C24F H24F C24F H24G
 BOND C24F H24H

DOUBLE C4S C5S

DOUBLE C1F OF

IMPR C1F C2F NF OF NF C1F C2S HNF

IC	C18S	C17S	C16S	C15S	1.5309	113.23	180.00	112.64	1.5346
IC	C17S	C16S	C15S	C14S	1.5337	112.64	180.00	112.58	1.5345
IC	C16S	C15S	C14S	C13S	1.5346	112.58	180.00	112.58	1.5345
IC	C15S	C14S	C13S	C12S	1.5345	112.58	-179.99	112.58	1.5345
IC	C14S	C13S	C12S	C11S	1.5345	112.58	179.97	112.58	1.5344
IC	C13S	C12S	C11S	C10S	1.5345	112.58	-179.96	112.61	1.5345
IC	C12S	C11S	C10S	C9S	1.5344	112.61	179.97	112.54	1.5344
IC	C11S	C10S	C9S	C8S	1.5345	112.54	-179.87	112.66	1.5345
IC	C10S	C9S	C8S	C7S	1.5344	112.66	-179.89	112.50	1.5345
IC	C9S	C8S	C7S	C6S	1.5345	112.50	-179.79	112.41	1.5391

IC	C8S	C7S	C6S	C5S	1.5354	112.41	-179.61	111.46	1.5077
IC	C7S	C6S	C5S	C4S	1.5391	111.46	-117.13	124.77	1.3444
IC	C6S	C5S	C4S	C3S	1.5077	124.77	179.01	124.28	1.5009
IC	C5S	C4S	C3S	C2S	1.3444	124.28	-84.24	110.14	1.5343
IC	C5S	C4S	C3S	O3	1.3444	124.28	156.07	107.64	1.4235
IC	C4S	C3S	C2S	C1S	1.5009	110.14	-70.50	114.25	1.5586
IC	H01	O1	C1S	C2S	1.5725	121.69	103.66	111.21	1.5586
IC	C2S	O1	*C1S	H1S	1.5586	111.21	118.79	108.35	1.1121
IC	C2S	O1	*C1S	H1T	1.5586	111.21	-123.25	112.85	1.1152
IC	O1	C1S	C2S	C3S	1.4327	111.21	175.67	114.25	1.5343
IC	C3S	C1S	*C2S	NF	1.5343	114.25	120.00	111.52	1.4406
IC	C3S	C1S	*C2S	H2S	1.5343	114.25	-120.00	105.03	1.1194
IC	C1S	C2S	NF	C1F	1.5586	111.52	160.94	124.85	1.3364
IC	C1F	C2S	*NF	HNF	1.3364	124.85	-157.81	113.39	1.0062
IC	C1S	C2S	C3S	C4S	1.5586	114.25	-70.50	110.14	1.5009
IC	C4S	C2S	*C3S	O3	1.5009	110.14	118.36	109.80	1.4235
IC	C4S	C2S	*C3S	H3S	1.5009	110.14	-121.14	110.10	1.1140
IC	C2S	C3S	O3	H03	1.5343	109.80	46.50	104.54	0.9610
IC	C2S	C3S	C4S	C5S	1.5343	110.14	-84.24	124.28	1.3444
IC	C5S	C3S	*C4S	H4S	1.3444	124.28	178.79	115.93	1.1003
IC	C3S	C4S	C5S	C6S	1.5009	124.28	179.01	124.77	1.5077
IC	C6S	C4S	*C5S	H5S	1.5077	124.77	-178.18	119.00	1.1029
IC	C4S	C5S	C6S	C7S	1.3444	124.77	-117.13	111.46	1.5391
IC	C7S	C5S	*C6S	H6S	1.5391	111.46	120.98	111.41	1.1142
IC	C7S	C5S	*C6S	H6T	1.5391	111.46	-120.60	109.91	1.1125
IC	C5S	C6S	C7S	C8S	1.5077	111.46	-179.61	112.41	1.5354
IC	C8S	C6S	*C7S	H7S	1.5354	112.41	121.26	109.74	1.1138
IC	C8S	C6S	*C7S	H7T	1.5354	112.41	-120.99	109.40	1.1147
IC	C6S	C7S	C8S	C9S	1.5391	112.41	-179.79	112.50	1.5345
IC	C9S	C7S	*C8S	H8S	1.5345	112.50	121.20	108.97	1.1132
IC	C9S	C7S	*C8S	H8T	1.5345	112.50	-121.38	109.07	1.1129
IC	C7S	C8S	C9S	C10S	1.5354	112.50	-179.89	112.66	1.5344
IC	C10S	C8S	*C9S	H9S	1.5344	112.66	121.49	109.05	1.1133
IC	C10S	C8S	*C9S	H9T	1.5344	112.66	-121.38	108.96	1.1136
IC	C8S	C9S	C10S	C11S	1.5345	112.66	-179.87	112.54	1.5345
IC	C11S	C9S	*C10S	H10S	1.5345	112.54	121.27	109.07	1.1133
IC	C11S	C9S	*C10S	H10T	1.5345	112.54	-121.34	109.11	1.1132
IC	C9S	C10S	C11S	C12S	1.5344	112.54	179.97	112.61	1.5344
IC	C12S	C10S	*C11S	H11S	1.5344	112.61	121.36	109.09	1.1133
IC	C12S	C10S	*C11S	H11T	1.5344	112.61	-121.33	109.07	1.1134
IC	C10S	C11S	C12S	C13S	1.5345	112.61	-179.96	112.58	1.5345
IC	C13S	C11S	*C12S	H12S	1.5345	112.58	121.29	109.12	1.1133
IC	C13S	C11S	*C12S	H12T	1.5345	112.58	-121.31	109.15	1.1133
IC	C11S	C12S	C13S	C14S	1.5344	112.58	179.97	112.58	1.5345
IC	C14S	C12S	*C13S	H13S	1.5345	112.58	121.30	109.13	1.1133
IC	C14S	C12S	*C13S	H13T	1.5345	112.58	-121.30	109.13	1.1133
IC	C12S	C13S	C14S	C15S	1.5345	112.58	-179.99	112.58	1.5345
IC	C15S	C13S	*C14S	H14S	1.5345	112.58	121.30	109.12	1.1133
IC	C15S	C13S	*C14S	H14T	1.5345	112.58	-121.31	109.12	1.1133
IC	C13S	C14S	C15S	C16S	1.5345	112.58	180.00	112.58	1.5346
IC	C16S	C14S	*C15S	H15S	1.5346	112.58	121.30	109.11	1.1133
IC	C16S	C14S	*C15S	H15T	1.5346	112.58	-121.30	109.11	1.1133
IC	C14S	C15S	C16S	C17S	1.5345	112.58	180.00	112.64	1.5337
IC	C17S	C15S	*C16S	H16S	1.5337	112.64	121.32	109.14	1.1133
IC	C17S	C15S	*C16S	H16T	1.5337	112.64	-121.32	109.14	1.1133
IC	C15S	C16S	C17S	C18S	1.5346	112.64	180.00	113.23	1.5309
IC	C18S	C16S	*C17S	H17S	1.5309	113.23	121.67	108.77	1.1142
IC	C18S	C16S	*C17S	H17T	1.5309	113.23	-121.67	108.77	1.1142
IC	C16S	C17S	C18S	H18S	1.5337	113.23	180.00	110.63	1.1113
IC	H18S	C17S	*C18S	H18T	1.1113	110.63	120.07	110.47	1.1114
IC	H18S	C17S	*C18S	H18U	1.1113	110.63	-120.07	110.47	1.1114
IC	C2S	NF	C1F	C2F	1.4406	124.85	-156.83	114.97	1.4964
IC	C2F	NF	*C1F	OF	1.4964	114.97	162.54	122.11	1.2268
IC	NF	C1F	C2F	C3F	1.3364	114.97	-146.57	111.92	1.5432
IC	C3F	C1F	*C2F	H2F	1.5432	111.92	-120.83	107.93	1.1115
IC	H2F	C1F	*C2F	H2G	1.1115	107.93	-117.30	109.04	1.1113
IC	C1F	C2F	C3F	C4F	1.4964	111.92	179.54	112.84	1.5310
IC	C4F	C2F	*C3F	H3F	1.5310	112.84	-121.63	109.37	1.1145
IC	H3F	C2F	*C3F	H3G	1.1145	109.37	-117.07	109.08	1.1157
IC	C2F	C3F	C4F	C5F	1.5434	112.23	174.86	112.58	1.5342
IC	C5F	C3F	*C4F	H4F	1.5342	112.58	-120.18	108.65	1.1140

IC	H4F	C3F	*C4F	H4G	1.1140	108.65	-117.52	109.62	1.1127
IC	C3F	C4F	C5F	C6F	1.5340	112.58	-174.06	112.39	1.5340
IC	C6F	C4F	*C5F	H5F	1.5340	112.39	-122.01	109.38	1.1130
IC	H5F	C4F	*C5F	H5G	1.1130	109.38	-117.34	108.92	1.1134
IC	C4F	C5F	C6F	C7F	1.5342	112.39	178.27	112.75	1.5340
IC	C7F	C5F	*C6F	H6F	1.5340	112.75	-120.93	108.96	1.1137
IC	H6F	C5F	*C6F	H6G	1.1137	108.96	-117.41	109.22	1.1131
IC	C5F	C6F	C7F	C8F	1.5340	112.75	-178.89	112.33	1.5342
IC	C8F	C6F	*C7F	H7F	1.5342	112.33	-121.40	109.26	1.1132
IC	H7F	C6F	*C7F	H7G	1.1132	109.26	-117.45	109.12	1.1130
IC	C6F	C7F	C8F	C9F	1.5340	112.33	179.09	112.77	1.5340
IC	C9F	C7F	*C8F	H8F	1.5340	112.77	-121.23	109.16	1.1134
IC	H8F	C7F	*C8F	H8G	1.1134	109.16	-117.38	109.08	1.1133
IC	C7F	C8F	C9F	C10F	1.5342	112.77	179.02	112.34	1.5343
IC	C10F	C8F	*C9F	H9F	1.5343	112.34	-121.27	109.27	1.1132
IC	H9F	C8F	*C9F	H9G	1.1132	109.27	-117.45	109.10	1.1130
IC	C8F	C9F	C10F	C11F	1.5340	112.34	178.16	112.74	1.5340
IC	C11F	C9F	*C10F	H10F	1.5340	112.74	-121.18	109.16	1.1134
IC	H10F	C9F	*C10F	H10G	1.1134	109.16	-117.35	109.10	1.1132
IC	C9F	C10F	C11F	C12F	1.5343	112.74	178.91	112.38	1.5343
IC	C12F	C10F	*C11F	H11F	1.5343	112.38	-121.39	109.34	1.1131
IC	H11F	C10F	*C11F	H11G	1.1131	109.34	-117.43	108.98	1.1132
IC	C10F	C11F	C12F	C13F	1.5340	112.38	176.93	112.68	1.5340
IC	C13F	C11F	*C12F	H12F	1.5340	112.68	-121.02	109.05	1.1134
IC	H12F	C11F	*C12F	H12G	1.1134	109.05	-117.34	109.20	1.1131
IC	C11F	C12F	C13F	C14F	1.5343	112.68	-179.93	112.45	1.5345
IC	C14F	C12F	*C13F	H13F	1.5345	112.45	-121.60	109.42	1.1129
IC	H13F	C12F	*C13F	H13G	1.1129	109.42	-117.40	108.85	1.1134
IC	C12F	C13F	C14F	C15F	1.5340	112.45	175.93	112.68	1.5334
IC	C15F	C13F	*C14F	H14F	1.5334	112.68	-121.00	109.06	1.1131
IC	H14F	C13F	*C14F	H14G	1.1131	109.06	-117.24	109.28	1.1130
IC	C13F	C14F	C15F	C16F	1.5345	112.68	-178.23	113.16	1.5306
IC	C16F	C14F	*C15F	H15F	1.5306	113.16	-121.86	108.84	1.1140
IC	H15F	C14F	*C15F	H15G	1.1140	108.84	-116.70	108.76	1.1144
IC	C14F	C15F	C16F	C17F	1.5345	112.68	-178.23	113.16	1.5306
IC	C17F	C15F	*C16F	H16F	1.5306	113.16	-121.86	108.84	1.1140
IC	H16F	C15F	*C16F	H16G	1.1140	108.84	-116.70	108.76	1.1144
IC	C15F	C16F	C17F	C18F	1.5345	112.68	-178.23	113.16	1.5306
IC	C18F	C16F	*C17F	H17F	1.5306	113.16	-121.86	108.84	1.1140
IC	H17F	C16F	*C17F	H17G	1.1140	108.84	-116.70	108.76	1.1144
IC	C16F	C17F	C18F	C19F	1.5345	112.68	-178.23	113.16	1.5306
IC	C19F	C17F	*C18F	H18F	1.5306	113.16	-121.86	108.84	1.1140
IC	H18F	C17F	*C18F	H18G	1.1140	108.84	-116.70	108.76	1.1144
IC	C17F	C18F	C19F	C20F	1.5344	112.57	179.80	112.59	1.5346
IC	C20F	C18F	*C19F	H19F	1.5346	112.59	-121.28	109.10	1.1133
IC	H19F	C18F	*C19F	H19G	1.1133	109.10	-117.38	109.12	1.1133
IC	C18F	C19F	C20F	C21F	1.5345	112.59	-179.89	112.63	1.5337
IC	C21F	C19F	*C20F	H20F	1.5337	112.63	-121.33	109.16	1.1133
IC	H20F	C19F	*C20F	H20G	1.1133	109.16	-117.38	109.14	1.1133
IC	C19F	C20F	C21F	C22F	1.5346	112.63	179.93	113.24	1.5309
IC	C22F	C20F	*C21F	H21F	1.5309	113.24	-121.66	108.77	1.1142
IC	H21F	C20F	*C21F	H21G	1.1142	108.77	-116.65	108.78	1.1142
IC	C20F	C21F	C22F	C23F	1.5346	112.63	179.93	113.24	1.5309
IC	C23F	C21F	*C22F	H22F	1.5309	113.24	-121.66	108.77	1.1142
IC	H22F	C21F	*C22F	H22G	1.1142	108.77	-116.65	108.78	1.1142
IC	C21F	C22F	C23F	C24F	1.5346	112.63	179.93	113.24	1.5309
IC	C24F	C22F	*C23F	H23F	1.5309	113.24	-121.66	108.77	1.1142
IC	H23F	C22F	*C23F	H23G	1.1142	108.77	-116.65	108.78	1.1142
IC	C22F	C23F	C24F	H24F	1.5337	113.24	-59.91	110.47	1.1114
IC	H24F	C23F	*C24F	H24G	1.1114	110.47	119.87	110.48	1.1113
IC	H24F	C23F	*C24F	H24H	1.1114	110.47	-120.06	110.63	1.1113

PATCH FIRST NONE LAST NONE

end

read para card flex @app
 * new parameters

*

ATOMS

MASS 165 NHL 14.00700 !N !

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MASS    166 CTO1  12.01100 !C !
MASS    167 CEL3   12.01100 !C !
MASS    168 CTO2   12.01100 !C !

BONDS
!
!V(bond) = Kb(b - b0)**2
!
!Kb: kcal/mole/A**2
!b0: A
!
!atom type Kb          b0
! Cyclopropane Moeity Addition
!

! Ceramides
! CC
CTL1  CEL3  250.000  1.490 ! prot22; Ala Dipeptide ab initio calc's (LK) fixed from 10/90 (5/91); CT1-C
CTO1  CEL3  250.000  1.490 ! prot22; Ala Dipeptide ab initio calc's (LK) fixed from 10/90 (5/91); CT1-C
CTO1  CTL1  222.500  1.500 ! lip36; alkanes, 3/92; CTL1-CTL1
CTO1  CTL3  222.500  1.538 ! lip36; alkanes, 3/92; CTL1-CTL3
CEL3   CTL2  365.000  1.502 ! lip36; butene; from propene, yin,adm jr., 12/95; CEL1-CTL2
CEL3   CEL3  440.000  1.340 ! lip36; butene, yin,adm jr., 12/95; CEL1-CEL1
CTL2   C     250.000  1.490 ! prot22; Ala Dipeptide ab initio calc's (LK) fixed from 10/90 (5/91); CT2-C
CTL3   C     250.000  1.490 ! prot22; Ala Dipeptide ab initio calc's (LK) fixed from 10/90 (5/91); CT3-C
CTO1  OHL   428.000  1.420 ! lip36; glycerol; OHL-CTL1
CTO1  OSL   340.000  1.430 ! lipid ester; rmv nov 2009
CTO1  HAL1  309.000  1.111 ! lip36; alkanes, 3/92; CTL1-HAL1
HEL1  CEL3  360.500  1.100 ! lip36; propene, yin,adm jr., 12/95; HEL1-CEL1
NHL   H     440.000  0.997 ! prot22; Alanine Dipeptide ab initio calc's (LK); NH1-H
NHL   CTL3  320.000  1.430 ! prot22; NMA Gas & Liquid Phase IR Spectra (LK); NH1-CT3
NHL   C     370.000  1.345 ! prot22; Alanine Dipeptide ab initio calc's (LK); NH1-C
NHL   CTL1  320.000  1.430 ! prot22; NMA Gas & Liquid Phase IR Spectra (LK); NH1-CT1
CTO2  HAL2  309.000  1.111 ! lip36; alkanes, 4/98
CTO2  CTL1  222.500  1.538 ! lip36; alkanes, 3/92
CTO2  OHL   428.000  1.420 ! glycerol

ANGLES
!
!V(angle) = Ktheta(Theta - Theta0)**2
!
!V(Urey-Bradley) = Kub(S - S0)**2
!
!Ktheta: kcal/mole/rad**2
!Theta0: degrees
!Kub: kcal/mole/A**2 (Urey-Bradley)
!S0: A
!
!atom types      Ktheta    Theta0    Kub      S0
! Ceramides
!CCC
CTL3   CTL1  CEL3   52.000   108.00 ! prot22; Alanine Dipeptide ab initio calc's (LK); CT3-CT1-C
CTL3   CTO1  CEL3   52.000   108.00 ! prot22; Alanine Dipeptide ab initio calc's (LK); CT3-CT1-C
CTL1   CEL3   CEL3   48.00    123.50 ! lip36; 2-butene,from 2-butene, yin,adm jr., 12/95; CEL1-CEL1-CTL2
CTO1   CEL3   CEL3   48.00    123.50 ! lip36; 2-butene,from 2-butene, yin,adm jr., 12/95; CEL1-CEL1-CTL2
CTL1   CTL1   CEL3   52.000   108.00 ! prot22; Alanine Dipeptide ab initio calc's (LK); CT1 CT1 C
CTL1   CTO1   CEL3   52.000   108.00 ! prot22; Alanine Dipeptide ab initio calc's (LK); CT1 CT1 C
C     CTL2   CTL2   52.000   108.00 ! prot22; Alanine Dipeptide ab initio calc's (LK); CT3-CT1-C
CTL2   CTL1   CTO1   58.350   113.50 11.16   2.561 ! lip36; glycerol; CTL1-CTL1-CTL2
CEL3   CEL3   CTL2   48.00    123.50 ! lip36; 2-butene,from 2-butene, yin,adm jr., 12/95; CEL1-CEL1-CTL2
CEL3   CTL2   CTL3   32.00    112.20 ! lip36; 1-butene; from propene, yin,adm jr., 12/95; CEL1-CTL2-CTL3
CEL3   CTL2   CTL2   32.00    112.20 ! lip36; 1-butene; from propene, yin,adm jr., 12/95; CEL1-CTL2-CTL2
CTL3   CTL2   C     52.000   108.00 ! prot22; Alanine Dipeptide ab initio calc's (LK); CT3-CT1-C
OHL   CTL1   CEL3   75.700   110.10 ! prot22; MeOH, EMB, 10/10/89; OH1-CT1-CT1
OHL   CTO1   CEL3   75.700   110.10 ! prot22; MeOH, EMB, 10/10/89; OH1-CT1-CT1
OHL   CTL1   CTL3   75.700   110.10 ! prot22; MeOH, EMB, 10/10/89; OH1 CT1 CT3
OHL   CTO1   CTL3   75.700   110.10 ! prot22; MeOH, EMB, 10/10/89; OH1 CT1 CT3
OHL   CTO1   CTL1   75.700   110.10 ! prot22; MeOH, EMB, 10/10/89; OH1 CT1 CT1
O     C     CTL2   80.000   121.00 ! prot22; Alanine Dipeptide ab initio calc's (LK); O-C-CT2

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O C CTL3 80.000 121.00 ! prot22; Alanine Dipeptide ab initio calc's (LK); O C CT3
 OSL CTO1 CTL1 75.700 110.10 ! analog for acetyl at sphingosine O3; rmv nov 2009
 OSL CTO1 CEL3 75.700 110.10 ! analog for acetyl at sphingosine O3; rmv nov 2009
 CL OSL CTO1 40.00 109.60 30.0 2.2651 ! analog for acetyl at sphingosine O3; rmv nov 2009
 CTL1 CEL3 HEL1 40.00 116.00 ! lip36; 1-butene, from propene,yin.adm jr.,12/95; HEL1-CEL1-CTL2
 CTO1 CEL3 HEL1 40.00 116.00 ! lip36; 1-butene, from propene,yin.adm jr.,12/95; HEL1-CEL1-CTL2
 HAL1 CTL1 CEL3 33.000 109.50 30.00 2.16300 ! prot22; alanine dipeptide, LK, replaced, adm jr.,
 5/09/91; HA-CT1-C (Protein)
 HAL1 CTO1 CEL3 33.000 109.50 30.00 2.16300 ! prot22; alanine dipeptide, LK, replaced, adm jr.,
 5/09/91; HA-CT1-C (Protein)
 HAL1 CTO1 CTL3 34.500 110.10 22.53 2.179 ! lip36; alkane, 3/92; HAL1-CTL1-CTL3
 HAL1 CTO1 CTL1 34.500 110.10 22.53 2.179 ! lip36; alkane, 3/92; HAL1-CTL1-CTL1
 HAL1 CTL1 CTO1 34.500 110.10 22.53 2.179 ! lip36; alkane, 3/92; HAL1-CTL1-CTL1
 HAL3 CTL3 CTO1 33.430 110.10 22.53 2.179 ! lip36; alkane, 4/98; HAL3-CTL3-CTL
 HEL1 CEL3 CEL3 52.00 119.50 ! lip36; 2-butene, yin,adm jr., 12/95; HEL1-CEL1-CEL1
 HEL1 CEL3 CTL2 40.00 116.00 ! lip36; 1-butene; from propene, yin,adm jr., 12/95; HEL1-CEL1-CTL2
 CEL3 CTL2 HAL2 45.00 111.50 ! lip36; 1-butene; from propene, yin,adm jr., 12/95; HAL2-CTL2-CEL1
 HAL2 CTL2 C 33.000 109.50 30.00 2.16300 ! prot22; alanine dipeptide, LK, replaced, adm jr.,
 5/09/91; HA-CT2-C
 HAL3 CTL3 C 33.000 109.50 30.00 2.16300 ! prot22; alanine dipeptide, LK, replaced, adm jr.,
 5/09/91; C CT3 HA
 NHL CTL1 OSL 80.000 122.50 ! prot22; NMA Vib Modes (LK); O-C-NH1
 O C NHL 80.000 122.50 ! prot22; NMA Vib Modes (LK); O-C-NH1
 HOL OHL CTO1 57.500 106.00 ! lip36; glycerol; HOL-OHL-CTL1
 OHL CTO1 HAL1 45.900 108.89 ! lip36; glycerol; OHL-CTL1-HAL1
 OSL CTO1 HAL1 45.900 108.89 ! lip36; glycerol; OHL-CTL1-HAL1; rmv nov 2009
 NHL CTL1 CTO1 70.000 113.50 ! prot22; Alanine Dipeptide ab initio calc's (LK); NH1-CT1-CT1
 NHL C CTL2 80.000 116.50 ! prot22; NMA Vib Modes (LK); NH1-C-CT2
 NHL CTL1 CTL2 70.000 113.50 ! prot22; Alanine Dipeptide ab initio calc's (LK); NH1-CT1-CT2
 NHL CTL1 CTL3 70.000 113.50 ! prot22;
 NHL CTL1 CTL1 70.000 113.50 ! prot22; Alanine Dipeptide ab initio calc's (LK); NH1-CT1-CT1
 NHL C CTL3 80.000 116.50 ! prot22; NMA Vib Modes (LK); NH1-C-CT3
 H NHL CTL3 35.000 117.00 ! prot22; NMA Vibrational Modes (LK); H-NH1-CT3
 H NHL C 34.000 123.00 ! prot22; NMA Vib Modes (LK); H-NH1-C
 H NHL CTL1 35.000 117.00 ! prot22; NMA Vibrational Modes (LK); H-NH1-CT1
 CTL3 NHL C 50.000 120.00 ! prot22; NMA Vib Modes (LK); CT3-NH1-C
 CTL1 NHL C 50.000 120.00 ! prot22; NMA Vib Modes (LK); CT1 NH1 C
 NHL CTL3 HAL3 51.500 109.50 ! prot22; NMA crystal (JCS); NH1-CT3-HA
 NHL CTL1 HAL1 48.000 108.00 ! prot22; Alanine Dipeptide ab initio calc's (LK); NH1-CT1-HB
 HOL OHL CTO2 57.500 106.00 ! lip36; glycerol; HOL-OHL-CTL2
 OHL CTO2 HAL2 45.900 108.89 ! lip36; glycerol; OHL-CTL2-HAL1
 HAL2 CTO2 CTL1 26.500 110.10 22.53 2.179 ! lip36; alkane, 4/98; HAL2-CTL2-CTL1
 HAL1 CTL1 CTO2 34.500 110.10 22.53 2.179 ! lip36; alkane, 3/92; HAL1-CTL1-CTL2
 NHL CTL1 CTO2 70.000 113.50 ! prot22; Alanine Dipeptide ab initio calc's(LK);NH1-CT1-CTO2
 CTO2 CTL1 CTO1 58.350 113.50 11.16 2.561 ! lip36; glycerol; CTL1-CTL1-CTL2
 HAL2 CTO2 HAL2 35.500 109.00 5.40 1.80200 ! alkane, 3/92 HAL2-CLT2-HAL2
 OHL CTO2 CTL1 75.700 110.10 ! prot22; MeOH, EMB, 10/10/89; OH1 CT2 CT2

!
 DIHEDRALS
 !
 !V(dihedral) = Kchi(1 + cos(n(chi) - delta))
 !
 !Kchi: kcal/mole
 !n: multiplicity
 !delta: degrees
 !
 !atom types Kchi n delta
 !
 ! Ceramides
 !CCCC
 CTL3 CTL1 CEL3 CEL3 0.9000 1 180.00 ! lip36; 2-pentene and 3-heptene; CEL1 CEL1 CTL2 CTL3
 CEL3 CEL3 CTL2 CTL3 0.9100 1 180.00 ! lip36; 2-hexene, adm jr., 11/09
 CEL3 CEL3 CTL2 CTL3 0.1800 2 180.00 !
 CEL3 CEL3 CTL2 CTL3 0.1700 3 180.00 ! ""
 CEL3 CEL3 CTL2 CTL2 0.9100 1 180.00 ! lip36; 2-hexene, adm jr., 11/09
 CEL3 CEL3 CTL2 CTL2 0.1800 2 180.00 !
 CEL3 CEL3 CTL2 CTL2 0.1700 3 180.00 ! ""
 CTL1 CEL3 CEL3 CTL2 0.4500 1 180.00 ! lip36; 2-butene, adm jr., 4/04; X CEL1 CEL1 X
 CTL1 CEL3 CEL3 CTL2 8.5000 2 180.00 ! lip36; 2-butene, adm jr., 4/04; X CEL1 CEL1 X

CTO1 CEL3 CEL3 CTL2 0.1500 1 0.00 ! prot27; 2-butene, adm jr., 2/00 update; X CEL1 CEL1 X
 CTO1 CEL3 CEL3 CTL2 8.5000 2 180.00 ! prot27; 2-butene, adm jr., 2/00 update; X CEL1 CEL1 X
 CTL1 CTL1 CEL3 CEL3 0.6000 1 180.00 ! lip36; 2-hexene; CEL1 CEL1 CTL2 CTL2
 !CCCO
 O C CTL2 CTL3 1.400 1 0.00 ! prot22; lactams, adm jr; O C CT2 CT2/O C CT1 CT3
 O C CTL2 CTL2 1.400 1 0.00 ! prot22; lactams, adm jr; O C CT2 CT2/O C CT1 CT3
 OHL CTL1 CEL3 CEL3 0.600 1 180.00 ! lip36; 2-hexene; CEL3 CEL3 CTL2 CTL2
 OSL CTO1 CEL3 CEL3 1.280 1 180.0 ! torsfit MP2/ccpVTZ, cerl model, erh 8/09
 OSL CTO1 CEL3 CEL3 0.560 2 180.0 ! "" analogy fct TACSPH, rmv nov 2009
 OSL CTO1 CEL3 CEL3 0.860 3 0.0 ! ""
 CTL2 CTL1 CTO1 OHL 0.200 3 0.00 ! lip36; alkane, 3/92, X CTL1 CTL1 X
 CTO1 CTL1 CTL2 OSL 0.390 1 180.00 ! torsfit MP2/ccpVTZ, sph model, erh 8/09; transferred from cer4
 CTO1 CTL1 CTL2 OSL 0.510 2 180.00 ! ""
 CTO1 CTL1 CTL2 OSL 0.870 3 180.00 ! ""
 CTL2 CTL1 CTO1 OSL 0.390 1 180.00 ! torsfit MP2/ccpVTZ, sph model, erh 8/09; transferred from cer4
 CTL2 CTL1 CTO1 OSL 0.510 2 180.00 ! "" analogy for TACSPH, rmv nov 2009
 CTL2 CTL1 CTO1 OSL 0.870 3 180.00 ! ""
 !HCOH
 HAL1 CTO1 OHL HOL 0.140 3 0.00 ! lip36; glycerol, X CTL1 OHL X
 !COCC
 CL OSL CTO1 CTL1 0.000 4 0.00 ! glycerol, betal 6/07; from lip36 for TACSPH, rmv nov 2009
 CL OSL CTO1 CTL1 0.150 3 180.00 ! glycerol, betal 6/07
 CL OSL CTO1 CTL1 1.453 2 180.00 ! glycerol, betal 6/07
 CL OSL CTO1 CTL1 0.837 1 180.00 ! glycerol, betal 6/07
 CEL3 CTO1 OSL CL 0.220 1 0.00 ! og OMeTHP compounds 2 and 3; rmv nov 2009
 CEL3 CTO1 OSL CL 0.280 2 180.00 ! " MCSA fit
 CEL3 CTO1 OSL CL 0.890 3 0.00 ! " MP2/cc-pVTZ//MP2/6-31G*
 !COCH
 HAL1 CTO1 OSL CL 0.00 3 0.00 ! lip36, X CT1 OSL X; TACCSPH, rmv nov 2009
 !CCCH
 HEL1 CEL3 CTL1 CTL3 0.1200 3 0.00 ! prot22; for butene, yin/adm jr., 12/95; HE1 CE1 CT2 CT3
 HEL1 CEL3 CTO1 CTL3 0.1200 3 0.00 ! prot22; for butene, yin/adm jr., 12/95; HE1 CE1 CT2 CT3
 HEL1 CEL3 CTL2 CTL2 0.1200 3 0.00 ! prot22; for butene, yin/adm jr., 12/95; HE1 CE1 CT2 CT3
 HAL1 CTL1 CEL3 CEL3 0.0300 3 0.00 ! prot22; for butene, yin/adm jr., 12/95; CE1 CE1 CT3 HA
 HAL1 CTO1 CEL3 CEL3 0.0300 3 0.00 ! prot22; for butene, yin/adm jr., 12/95; CE1 CE1 CT3 HA
 CTL1 CTL1 CEL3 HEL1 0.1200 3 0.00 ! lip36; butene, yin, adm. jr; CTL2 CTL2 CEL1 HEL1
 HAL1 CTL3 CTO1 CEL3 0.2000 3 0.00 ! lip36; alkane, 3/92; X CTL1 CTL3 X
 CTL2 CTL1 CTO1 HAL1 0.200 3 0.00 ! lip36; alkane, 3/92; X CTL1 CTL3 X
 CTL1 CTO1 CEL3 HEL1 0.1200 3 0.00 ! lip36 butene, yin, adm. jr; CTL2 CTL2 CEL1 HEL1
 HAL1 CTL1 CTO1 CEL3 0.200 3 0.00 ! lip36; alkane, 3/92; X CTL1 CTL3 X
 CTO1 CEL3 CEL3 HEL1 0.1500 1 180.00 ! lip36; 2-butene, adm jr., 2/00 update; X CEL1 CEL1 X
 CTO1 CEL3 CEL3 HEL1 8.5000 2 180.00 ! lip36; 2-butene, adm jr., 2/00 update; X CEL1 CEL1 X
 CEL3 CEL3 CTL2 HAL2 0.3000 3 180.00 ! lip36; 2-butene, adm jr., 4/04; CEL1 CEL1 CTL2 HAL2
 HEL1 CEL3 CEL3 CTL2 0.1500 1 180.00 ! lip36; 2-butene, adm jr., 2/00 update; X CEL1 CEL1 X
 HEL1 CEL3 CEL3 CTL2 8.5000 2 180.00 ! lip36; 2-butene, adm jr., 2/00 update; X CEL1 CEL1 X
 HEL1 CEL3 CTL2 CTL3 0.1200 3 0.00 ! lip36; butene, yin, adm jr., 12/95; HEL1 CEL1 CTL2 CTL3
 !HCCH
 HEL1 CEL3 CTL1 HAL1 0.000 3 0.00 ! prot22; butene, adm jr., 2/00 update; HE1 CE1 CT2 HA
 HEL1 CEL3 CTO1 HAL1 0.000 3 0.00 ! prot22; butene, adm jr., 2/00 update; HE1 CE1 CT2 HA
 HAL3 CTL3 CTO1 HAL1 0.200 3 0.00 ! lip36; alkane, 3/92; X CTL1 CTL3 X
 HAL1 CTL1 CTO1 HAL1 0.200 3 0.00 ! lip36; alkane, 3/92; X CTL1 CTL1 X
 HEL1 CEL3 CEL3 HEL1 1.000 2 180.00 ! lip36; 2-butene, adm jr., 8/98 update; HEL1 CEL1 CEL1 HEL1
 HEL1 CEL3 CTL2 HAL2 0.000 3 0.00 ! lip36; butene, adm jr., 2/00 update; HEL1 CEL1 CTL2 HAL2
 !OCCH
 OHL CTL1 CEL3 HEL1 0.120 3 180.00 ! lip36; butene, yin, adm jr., 12/95; HEL1 CEL1 CTL1 CTL3
 OHL CTO1 CEL3 HEL1 0.120 3 180.00 ! lip36; butene, yin, adm jr., 12/95; HEL1 CEL1 CTL1 CTL3
 OSL CTO1 CEL3 HEL1 0.120 3 180.00 ! lip36; butene, yin, adm jr., 12/95; HEL1 CEL1 CTL1 CTL3 rmv nov
 2009
 O C CTL2 HAL2 0.000 3 180.00 ! prot22; adm jr., 8/13/90 acetamide geometry and vibrations; O C
 CT2 HA
 O C CTL3 HAL3 0.000 3 180.00 ! prot22; adm jr., 8/13/90 acetamide geometry and vibrations; O C
 CT3 HA
 HAL3 CTL3 CTO1 OHL 0.200 3 0.00 ! lip36; alkane, 3/92; X CTL1 CTL3 X
 HAL1 CTL1 CTO1 OHL 0.200 3 0.00 ! lip36; alkane, 3/92; X CTL1 CTL1 X
 HAL1 CTL1 CTO1 OSL 0.200 3 0.00 ! lip36; alkane, 3/92; X CTL1 CTL1 X; rmv nov 2009
 !NCCH
 NHL CTL1 CTL1 CTL3 0.260 1 180.00 ! torsfit MP2/ccpVTZ, cer3 model, erh 8/09
 NHL CTL1 CTL1 CTL3 0.390 2 180.00 ! ""
 NHL CTL1 CTL1 CTL3 0.120 3 180.00 ! ""

NHL C CTL2 HAL2 0.000 3 0.00 ! prot22; LK for autogenerate dihe, sp2-methyl, no dihedral potential; NHL C CT2 HA
 NHL C CTL3 HAL3 0.000 3 0.00 ! prot22; LK for autogenerate dihe, sp2-methyl, no dihedral potential; NHL C CT3 HA
 NHL CTL1 CTO1 HAL1 0.200 3 0.00 ! lip36; alkane, 3/92; X CTL1 CTL1 X
 !HNCH
 HAL3 CTL3 NHL H 0.000 3 0.00 ! prot22; LK for autogenerate dihe, sp2-methyl, no dihedral potential; HA CT3 NH1 H
 HAL1 CTL1 NHL H 0.000 1 0.00 ! prot22; Alanine Dipeptide ab initio calc's (LK); HB CT1 NH1 H
 !OCNH HNCC for NMA/CERN cis/trans barrier
 H NHL C O 0.530 1 0.00 ! fit to cern QM profile, jbk 10/12
 H NHL C O 1.270 2 180.00 ! fit to cern QM profile, jbk 10/12
 H NHL C O 0.140 3 0.00 ! fit to cern QM profile, jbk 10/12
 H NHL C O 0.780 4 0.00 ! fit to cern QM profile, jbk 10/12
 H NHL C CTL2 2.500 2 180.00 ! prot22; Gives appropriate NMA cis/trans barrier. (LK); H NH1 C
 CT2
 H NHL CTL1 CTL2 0.000 1 0.00 ! prot22; Alanine Dipeptide ab initio calc's (LK); H NH1 CT1 CT2
 H NHL CTL1 CTL1 0.000 1 0.00 ! prot22; Alanine Dipeptide ab initio calc's (LK); H NH1 CT1 CT1
 H NHL C CTL3 2.500 2 180.00 ! prot22; Gives appropriate NMA cis/trans barrier. (LK); H NH1 C
 CT3
 H NHL CTL1 CTL3 2.500 2 180.00 ! prot22;
 H NHL CTL1 CTO1 0.000 1 0.00 ! prot22; Alanine Dipeptide ab initio calc's (LK); H NH1 CT1 CT1
 !OCNC
 O C NHL CTL3 2.500 2 180.00 ! prot22; Gives appropriate NMA cis/trans barrier. (LK); O C NH1
 CT3
 O C NHL CTL1 2.500 2 180.00 ! prot22; Gives appropriate NMA cis/trans barrier. (LK); O C NH1
 CT1
 !CCNC
 CTL1 NHL C CTL2 0.000 1 0.00 ! set to zero in fits
 CTL1 NHL C CTL3 0.000 1 0.00 ! set to zero in fits
 ! torsion 4 and OH torsion
 CTL2 C NHL CTL3 1.600 1 0.00 ! prot22; from CT2 C NH1 CT2, adm jr. 10/21/96; CT2 C NH1 CT3
 CTL3 C NHL CTL3 1.600 1 0.00 ! prot22; from CT2 C NH1 CT2, adm jr. 10/21/96; CT2 C NH1 CT3
 CTL1 CTL1 NHL C 0.000 1 0.00 ! set to zero in fit
 CTL2 CTL1 NHL C 0.000 1 0.00 ! set to zero in fit
 CTL3 CTL1 NHL C 0.000 1 0.00 ! set to zero in fit
 C NHL CTL1 CTO1 4.080 1 0.00 ! torsion fit to cer6 qm, jbk 10/12
 C NHL CTL1 CTO1 1.560 2 180.00 ! torsion fit to cer6 qm, jbk 10/12
 C NHL CTL1 CTO1 1.800 3 0.00 ! torsion fit to cer6 qm, jbk 10/12
 CTL1 CTO1 OHL HOL 5.000 1 180.00 ! torsion fit to cer6 qm, jbk 10/12
 CTL1 CTO1 OHL HOL 0.220 2 0.00 ! torsion fit to cer6 qm, jbk 10/12
 CTL1 CTO1 OHL HOL 1.110 3 180.00 ! torsion fit to cer6 qm, jbk 10/12
 HOL OHL CTL1 CEL3 1.330 1 0.00 ! prot22; 2-propanol OH hf/6-31g* torsional surface, adm jr.,
 3/2/93; H OHL CT1 CT1
 HOL OHL CTO1 CEL3 1.330 1 0.00 ! prot22; 2-propanol OH hf/6-31g* torsional surface, adm jr.,
 3/2/93; H OHL CT1 CT1
 HOL OHL CTO1 CTL3 0.000 3 0.00 ! torsion fit to cer6 qm, jbk 9/12
 !HCNC
 HAL3 CTL3 NHL C 0.000 3 0.00 ! prot22; LK for autogenerate dihe, sp2-methyl, no dihedral potential; HA CT3 NH1 C
 HAL1 CTL1 NHL C 0.000 1 0.00 ! prot22; Alanine Dipeptide ab initio calc's (LK); HB CT1 NH1 C
 !NCCO
 NHL CTL1 CTL2 OSL 1.630 1 180.00 ! torsfit MP2/ccpVTZ, cer3 model, erh 8/09; transferred
 NHL CTL1 CTL2 OSL 0.490 2 0.00 ! ""
 NHL CTL1 CTL2 OSL 0.570 3 0.00 ! ""
 NHL CTL1 CTO1 OSL 1.630 1 180.00 ! torsfit MP2/ccpVTZ, cer3 model, erh 8/09; transferred
 NHL CTL1 CTO1 OSL 0.490 2 0.00 ! "" analogy for TACSPH, rmv nov 2009
 NHL CTL1 CTO1 OSL 0.570 3 0.00 ! ""
 !OPOC
 OSL PL OSL CTL2 1.200 1 180.00 ! lip36; phosphate, new NA, 4/98, adm jr.
 OSL PL OSL CTL2 0.100 2 180.00 ! lip36; phosphate, new NA, 4/98, adm jr.
 OSL PL OSL CTL2 0.100 3 180.00 ! lip36; phosphate, new NA, 4/98, adm jr.
 !torsion1
 CEL3 CTO1 CTL1 NHL 0.940 1 0.00 ! torfit to cer4 qm, jbk 5/13
 CEL3 CTO1 CTL1 NHL 0.880 2 180.00 ! torfit to cer4 qm, jbk 5/13
 CEL3 CTO1 CTL1 NHL 2.090 3 0.00 ! torfit to cer4 qm, jbk 5/13
 NHL CTL1 CTO1 OHL 0.420 1 0.00 ! torfit to cer4 qm, jbk 5/13
 NHL CTL1 CTO1 OHL 1.530 2 180.00 ! torfit to cer4 qm, jbk 5/13
 NHL CTL1 CTO1 OHL 1.580 3 180.00 ! torfit to cer4 qm, jbk 5/13
 CEL3 CTO1 CTL1 CTL2 0.000 1 0.00 ! torfit to cer4 qm, jbk 10/12
 !torsion2

CEL3 CEL3 CTO1 CTL1 0.860 1 0.00 ! torfit to cer4 qm w/adj, jbk 5/13
 CEL3 CEL3 CTO1 CTL1 1.910 2 0.00 ! torfit to cer4 qm w/adj, jbk 5/13
 CEL3 CEL3 CTO1 CTL1 1.520 3 180.00 ! torfit to cer4 qm w/adj, jbk 5/13
 CEL3 CEL3 CTO1 OHL 1.380 1 180.00 ! torfit to cer4 qm w/adj, jbk 5/13
 CEL3 CEL3 CTO1 OHL 0.580 2 0.00 ! torfit to cer4 qm w/adj, jbk 5/13
 CEL3 CEL3 CTO1 OHL 1.030 3 0.00 ! torfit to cer4 qm w/adj, jbk 5/13
 !torsion3
 NHL CTL1 CTL2 OSLP 1.370 1 0.00 ! torfit to cer3 qm, jbk 5/13
 NHL CTL1 CTL2 OSLP 0.340 2 0.00 ! torfit to cer3 qm, jbk 5/13
 NHL CTL1 CTL2 OSLP 0.200 3 180.00 ! torfit to cer3 qm, jbk 5/13
 NHL CTL1 CTL2 OSLP 0.020 4 180.00 ! torfit to cer3 qm, jbk 5/13
 NHL CTL1 CTL2 OSLP 0.140 5 0.00 ! torfit to cer3 qm, jbk 5/13
 NHL CTL1 CTL2 OSLP 0.060 6 0.00 ! torfit to cer3 qm, jbk 5/13
 CTL1 CTL1 CTL2 OSLP 0.000 1 0.00 ! torfit to cer3 qm, jbk 5/13
 !torsion5
 CTL3 CTL2 C NHL 1.990 1 0.00 ! torfit to cer2 qm, jbk 10/12
 CTL3 CTL2 C NHL 0.640 2 180.00 ! torfit to cer2 qm, jbk 10/12
 CTL3 CTL2 C NHL 0.280 3 180.00 ! torfit to cer2 qm, jbk 10/12
 CTL2 CTL2 C NHL 1.990 1 0.00 ! torfit to cer2 qm, jbk 10/12
 CTL2 CTL2 C NHL 0.640 2 180.00 ! torfit to cer2 qm, jbk 10/12
 CTL2 CTL2 C NHL 0.280 3 180.00 ! torfit to cer2 qm, jbk 10/12
 X CTO1 X 0.200 3 0.00 ! alkane, 3/92
 X CTO1 CTL3 X 0.200 3 0.00 ! alkane, 3/92
 !Added by STN for CER24, 3/13
 NHL CTO2 OHL 0.550 1 0.00 ! jbk; NHL CTL1 CTO1 OHL
 NHL CTO1 CTO2 OHL 1.630 2 180.00 ! jbk; NHL CTL1 CTO1 OHL
 NHL CTO1 CTO2 OHL 1.430 3 180.00 ! jbk; NHL CTL1 CTO1 OHL
 C NHL CTL1 CTO2 4.080 1 0.00 ! jbk; C NHL CTL1 CTO1
 C NHL CTL1 CTO2 1.560 2 180.00 ! jbk; C NHL CTL1 CTO1
 C NHL CTL1 CTO2 1.800 3 0.00 ! jbk; C NHL CTL1 CTO1
 X CTO1 CTO2 X 0.200 3 0.00 ! alkane, 3/92; X CTL1 CTL2 X
 X CTO2 OHL X 0.140 3 0.00 ! glycerol; X CTL2 OHL X
 CTO2 CTL1 CTO1 OHL 0.200 3 0.00 ! lip36; alkane, 3/92, X CTL1 CTL1 X
 H NHL CTL1 CTO2 0.000 1 0.00 ! prot22; Alanine Dipeptide ab initio calc's (LK); H NH1 CT1 CT2
 CEL3 CTO1 CTL1 CTO2 0.000 1 0.00 ! torfit to cer4 qm, jbk 10/12

!

IMPROPER

!

!V(improper) = Kpsi(psi - psi0)**2

!

!Kpsi: kcal/mole/rad**2

!psi0: degrees

!note that the second column of numbers (0) is ignored

!

!atom types Kpsi psi0

!

! Ceramides

NHL X X H 20. 0 0.00 ! amide, from peptide; rmv nov 2009

NONBONDED nbxmod 5 atom cdiel shift vatom vdistance vswitch -
 cutnb 14.0 ctofnb 12.0 ctonnb 10.0 eps 1.0 el14fac 1.0 wmin 1.5
 !adm jr., 5/08/91, suggested cutoff scheme

!

!V(Lennard-Jones) = Eps,i,j[(Rmin,i,j/ri,j)**12 - 2(Rmin,i,j/ri,j)**6]

!

!epsilon: kcal/mole, Eps,i,j = sqrt(eps,i * eps,j)

!Rmin/2: A, Rmin,i,j = Rmin/2,i + Rmin/2,j

!

!atom ignored epsilon Rmin/2 ignored eps,1-4 Rmin/2,1-4

! Ceramides

NHL 0.0 -0.2000 1.850 0.0 -0.20 1.55 ! ALLOW PEP POL ARO
 ! This 1,4 vdW allows the C5 dipeptide minimum to exist.(LK) - TAKEN FROM NH1

CTO1 0.0 -0.0200 2.275 0.0 -0.01 1.9 ! alkane, 3/92
 ! Transferred from CTL1; lip36

CEL3 0.0 -0.0680 2.090 ! alkene, yin,adm jr., 12/95
 ! Transferred from CEL1; lip36! Added my STN for CER24

CTO2 0.0 -0.0560 2.010 0.0 -0.01 1.9 ! STN, 3/13

end

return