

Supporting Information

Microsecond Molecular Dynamics Studies of Cholesterol-Mediated Myelin Sheath Degeneration in Early Alzheimer's Disease

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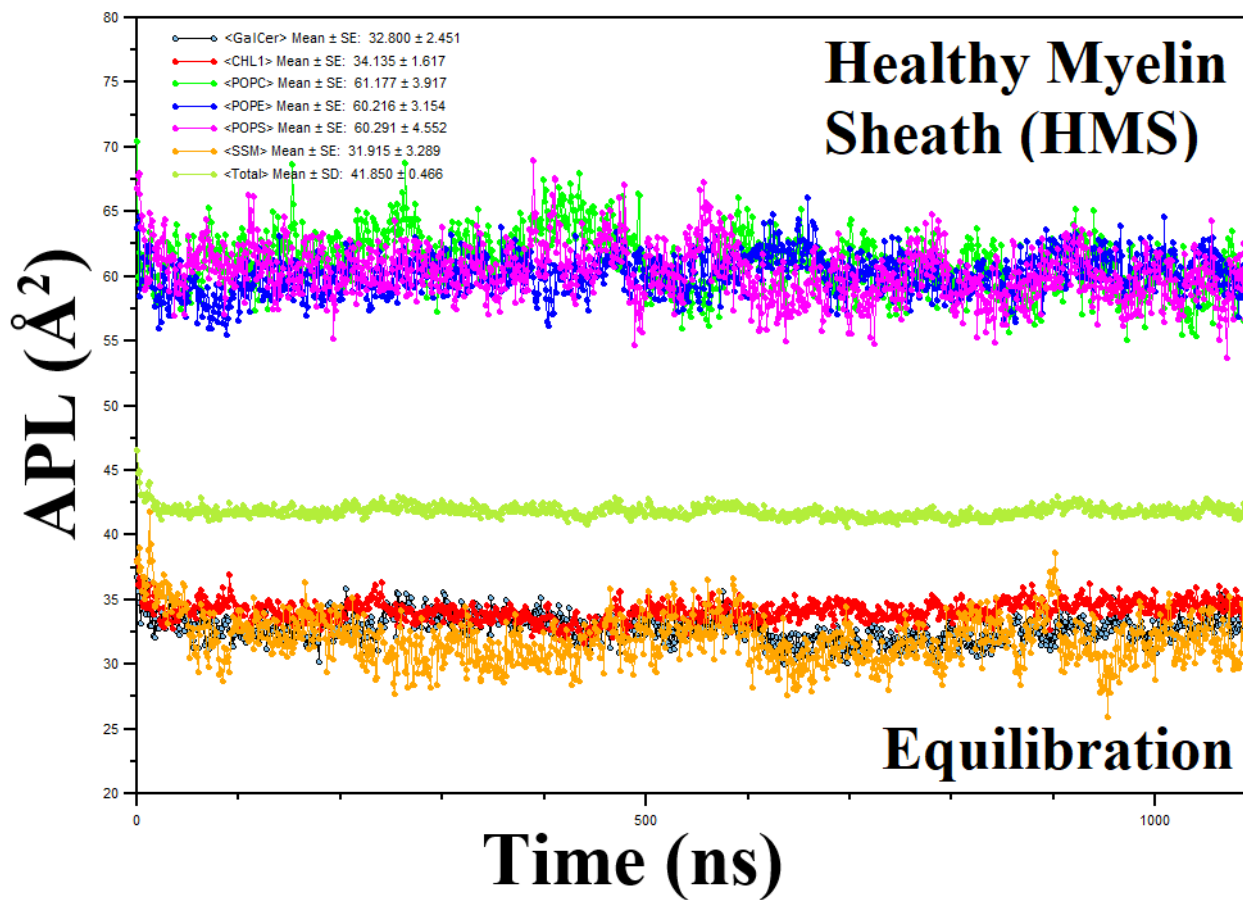


Figure S1: Area per lipid (APL) as a function of equilibration time for healthy myelin sheath membrane (HMS) bilayer system. The component APL (\AA^2) of specific lipid of AP bilayer i.e GalCer, CHL, POPC, POPE, POPS and SSM, is also shown.

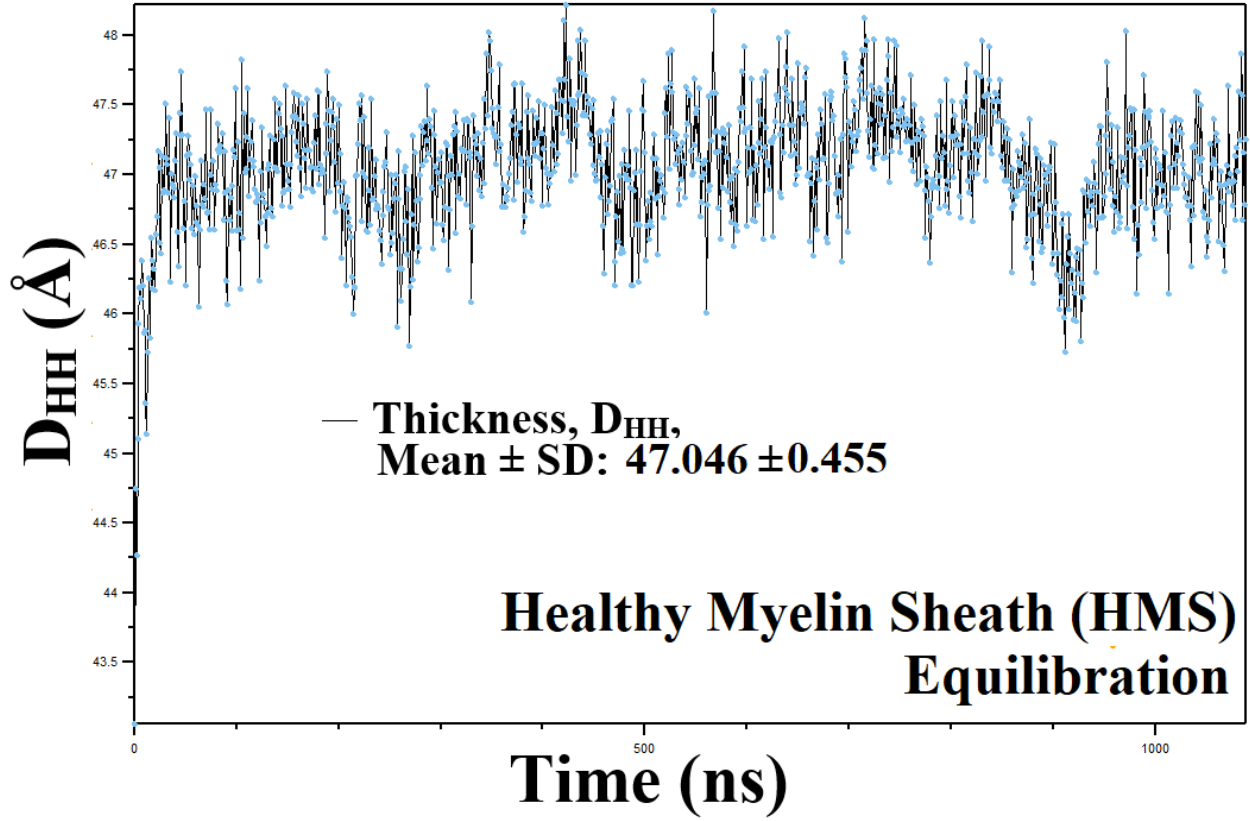


Figure S2: Bilayer thickness (D_{HH}) as a function of equilibration time for healthy myelin sheath membrane (HMS) bilayer system.

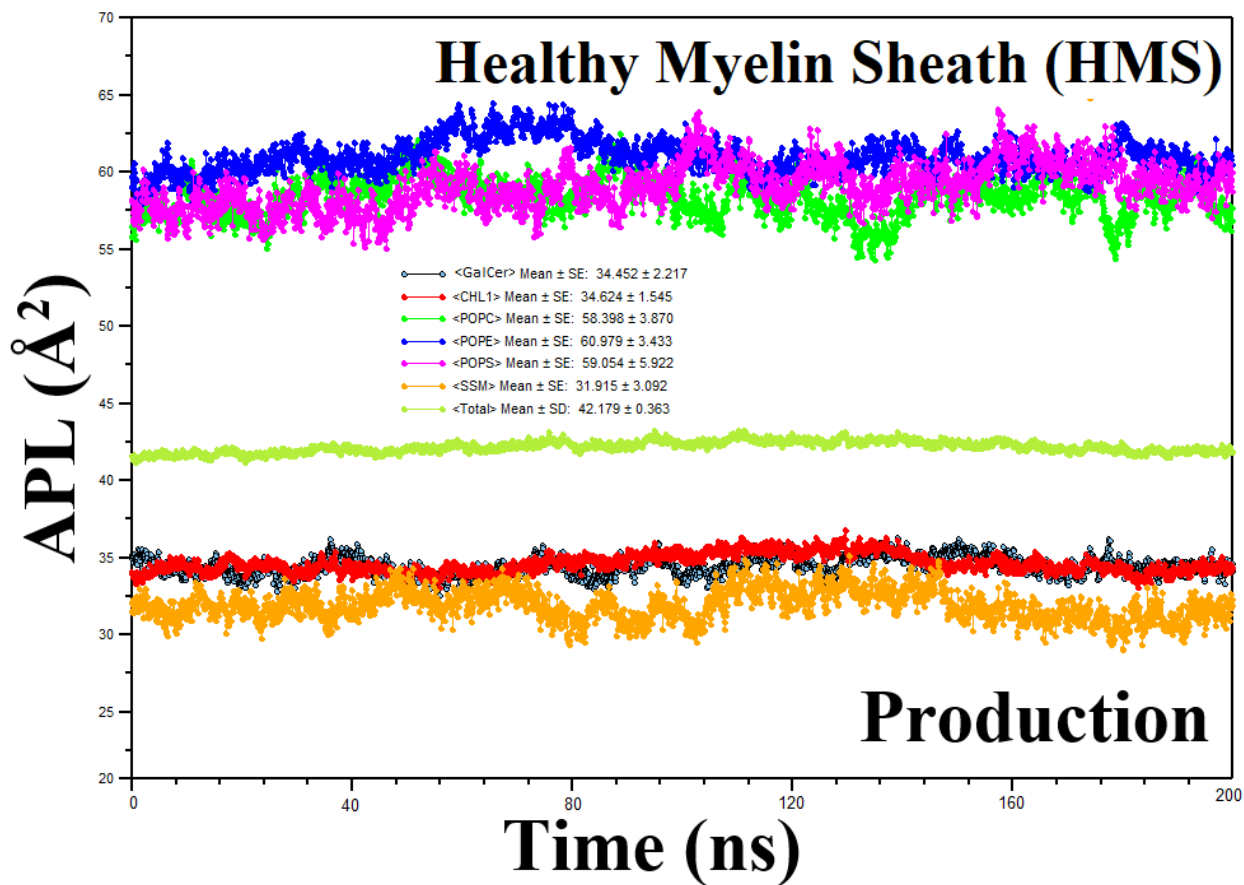


Figure S3: Area per lipid (APL) as a function of time for healthy myelin sheath membrane (HMS) bilayer system at production stage (after equilibration). The component APL (Å²) of specific lipid of AP bilayer i.e GalCer, CHL, POPC, POPE, POPS and SSM, is also shown.

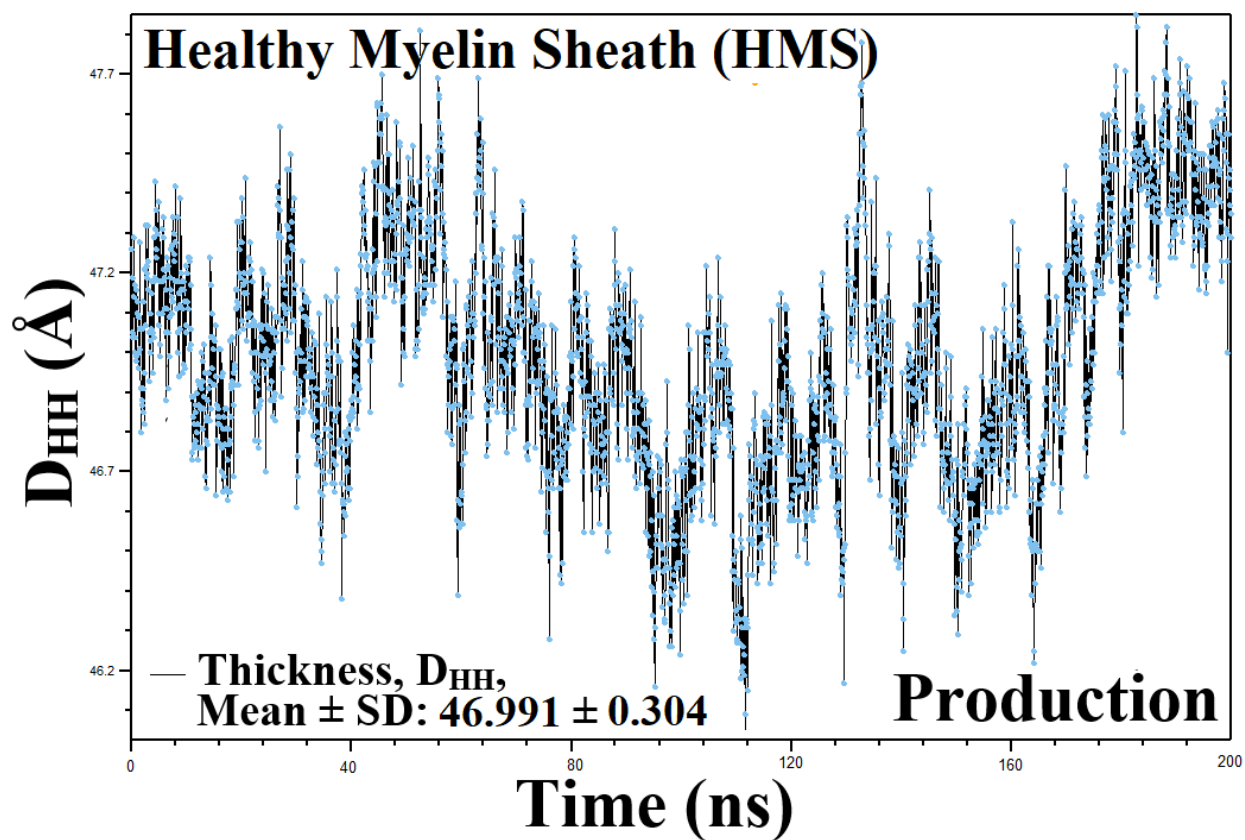


Figure S4: Bilayer thickness (D_{HH}) as a function of time for healthy myelin sheath membrane (HMS) bilayer system at production stage (after equilibration).

Table S1: The average P-N angle for POPC, POPE, POPS, SSM lipids and average cholesterol tilt angle (C13-C10) of reference membrane systems at 310 K. The standard deviation in the results over the 0.2 μ S simulation trajectory is also given.

S. No.	Bilayer	χ_{CHL}	P-N angle (deg)				C13-C10 angle (deg)
			POPC	POPE	POPS	SSM	
1	PC	0	67.953+1.92	--	--	--	--
2	PC:PE	0	62.951+2.57	82.084+2.66	--	--	--
3	PC:PE:CHL	0.34	64.206+3.21	79.068+3.34	--	--	14.349+0.96
4	PC:PE:CHL:SSM	0.25	64.467+3.58	81.687+3.54	--	67.541+3.37	13.427+1.12
5	PC:PE:CHL:GalCer	0.25	63.320+3.72	77.892+3.79	--	--	11.967+0.92

Table S2: The average fraction of contacts and mass overlap, interdigitation within lipid bilayers systems of reference membrane systems at 310 K. The standard deviation in the results over the 0.2 μ S simulation trajectory is also given.

S.No.	Bilayer	χ_{CHL}	Interdigitation of bilayer leaflets i.e. width of overlapping regions $W\rho$ (nm)	Fraction of mass overlap ($I\rho$)	Fraction of contacts (I_c)
1	PC	0	0.520+0.050	0.288+0.019	0.053+0.004
2	PC:PE	0	0.468+0.049	0.252+0.020	0.047+0.004
3	PC:PE:CHL	0.34	0.323+0.044	0.164+0.019	0.033+0.003
4	PC:PE:CHL:SSM	0.25	0.314+0.049	0.153+0.016	0.031+0.003
5	PC:PE:CHL:GalCer	0.25	0.288+0.045	0.153+0.018	0.026+0.003

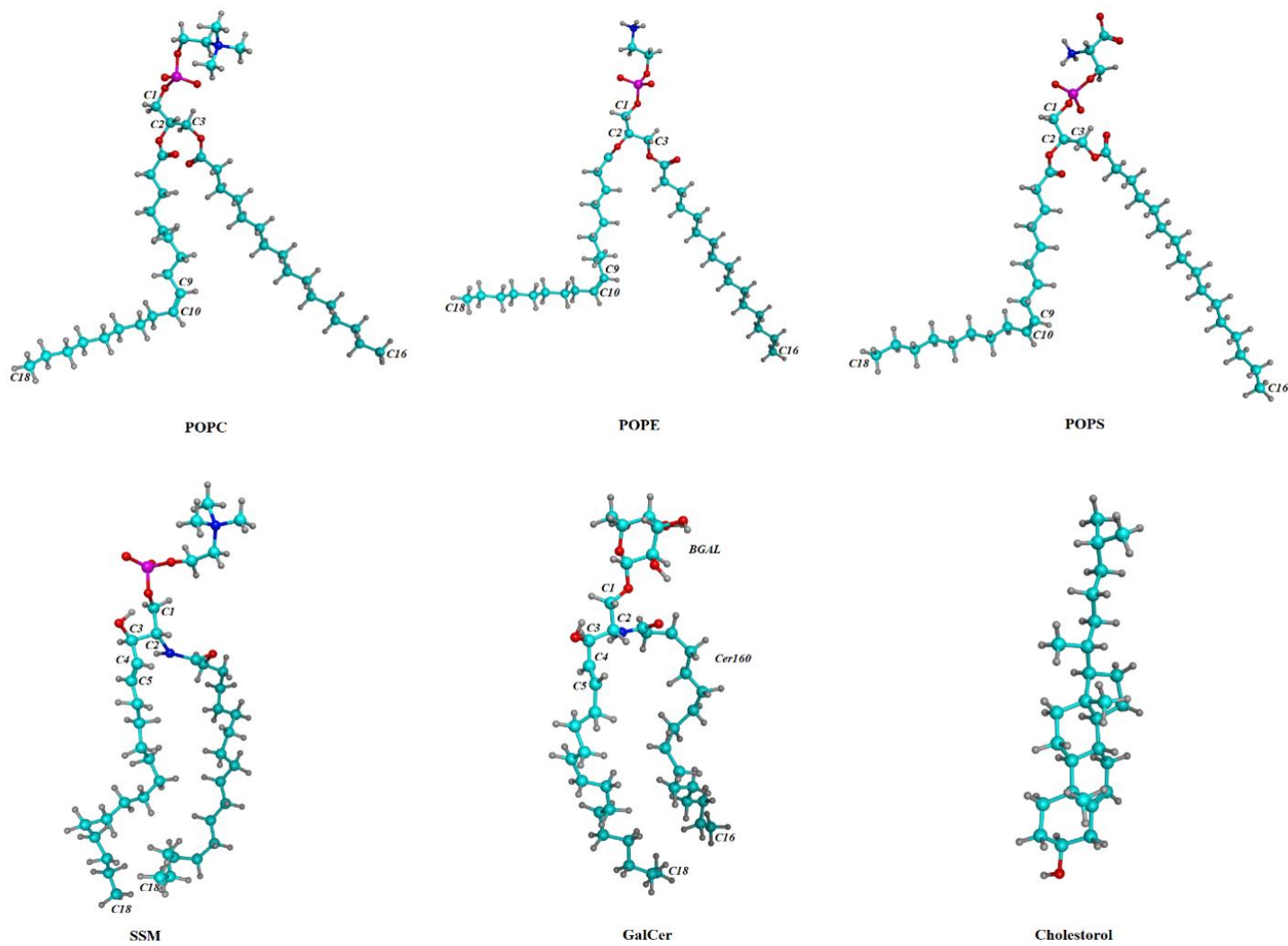


Figure S5: Structure of various lipids and cholesterol molecules used in bilayer systems.