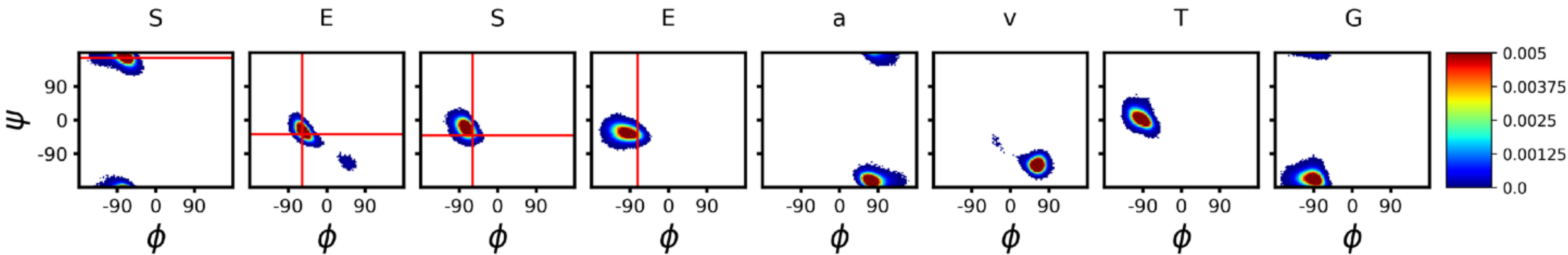
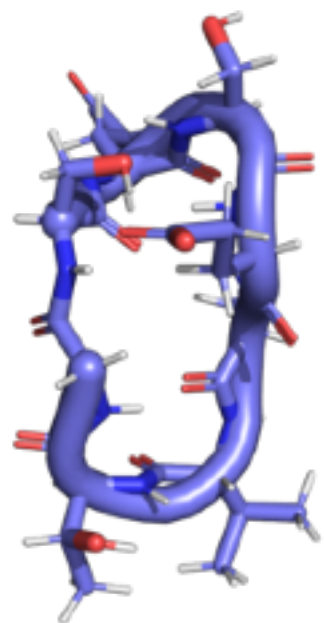
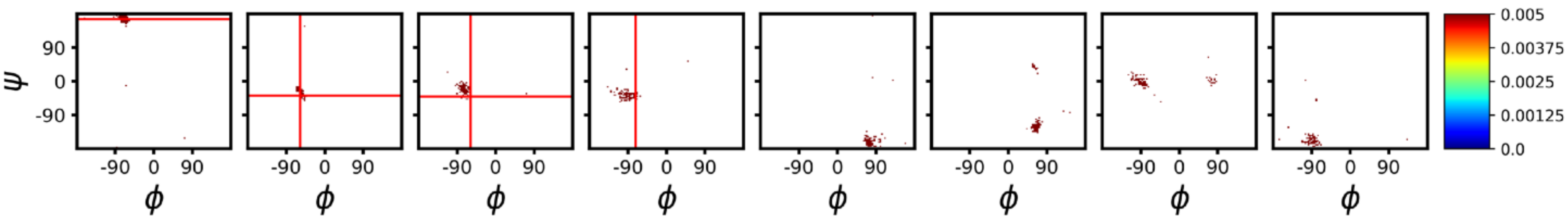
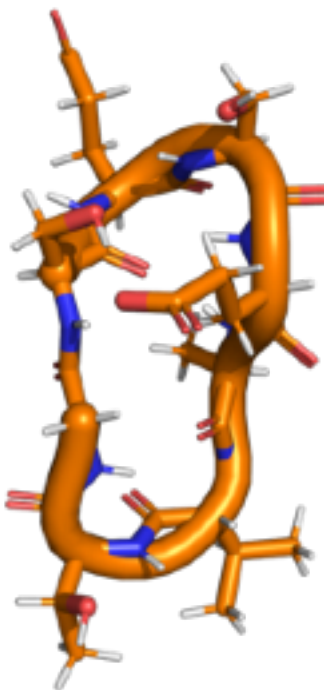


# NMR confirms structural prediction for cyclo-(SESEavTG)

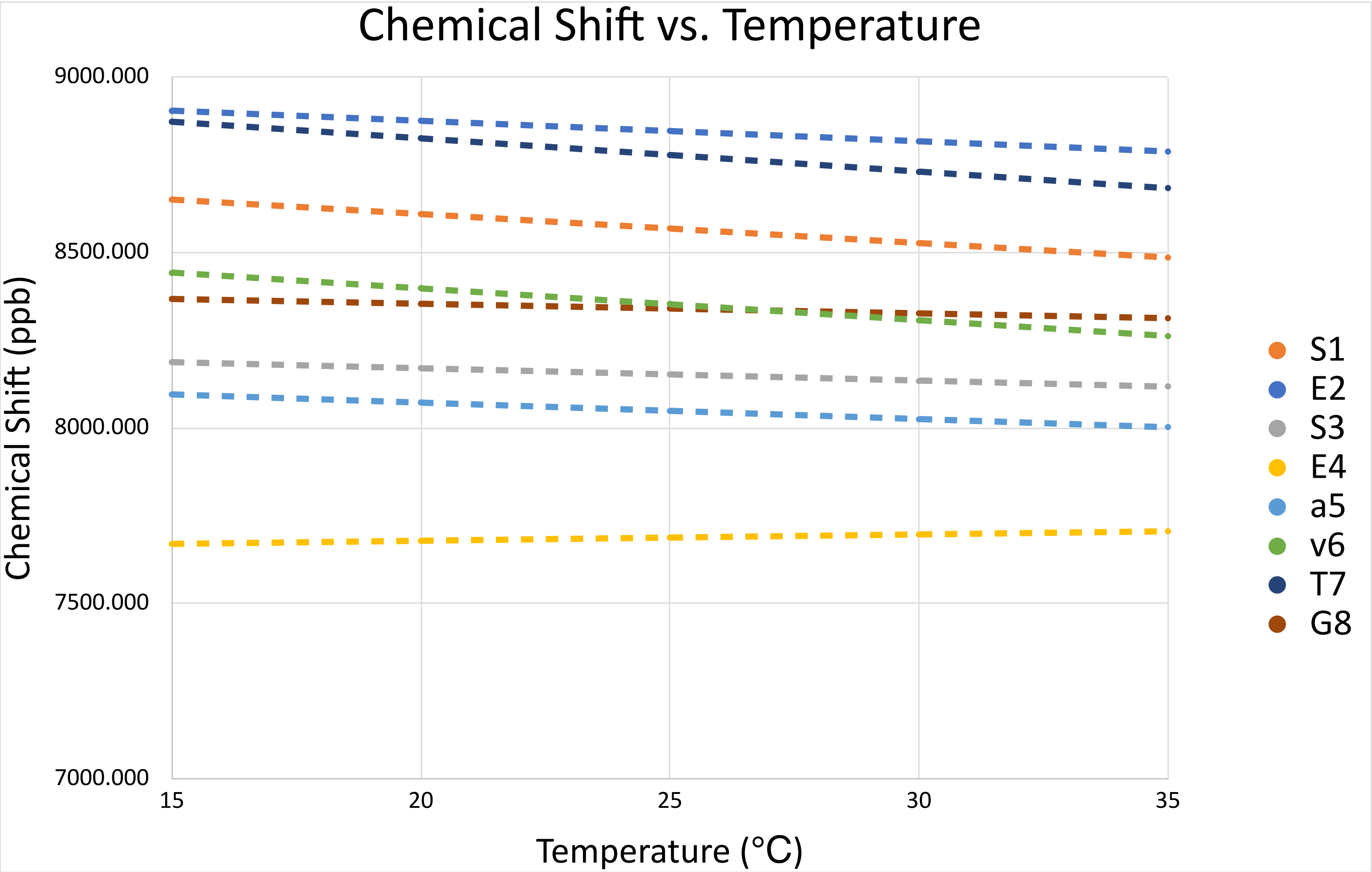
Predicted  
Structure



Constructed  
Structure



# NMR reveals hydrogen bond network in cyclo-(SESEavTG)



Residue	Slope (ppb/°C)
Ser1	-8.3
Glu2	-5.8
Ser3	-3.5
Glu4	1.8
D-Ala5	-4.7
D-Val6	-9
Thr7	-9.5
Gly8	-2.7

\*using -4.5 ppb/°C as cutoff

Baxter, N. J. & Williamson, M. P., *J. Biomol. NMR* **9**, 359-369, (1997)