

Effects of mutation on the amyloidogenic propensity of apolipoprotein C-II₆₀₋₇₀ peptide

Nevena Todorova¹, Andrew Hung¹, Simon M. Maaser², Michael D.W. Griffin², John Karas³,
Geoffrey J. Howlett² and Irene Yarovsky^{1*}

¹ Health Innovations Research Institute, School of Applied Sciences, RMIT University,
Melbourne, VIC, Australia.

² Biochemistry and Molecular Biology and Bio21 Molecular Science and
Biotechnology Institute, The University of Melbourne, Melbourne, VIC, Australia.

³ Bio21 Molecular Science and Biotechnology Institute, The University of Melbourne,
Melbourne, VIC, Australia.

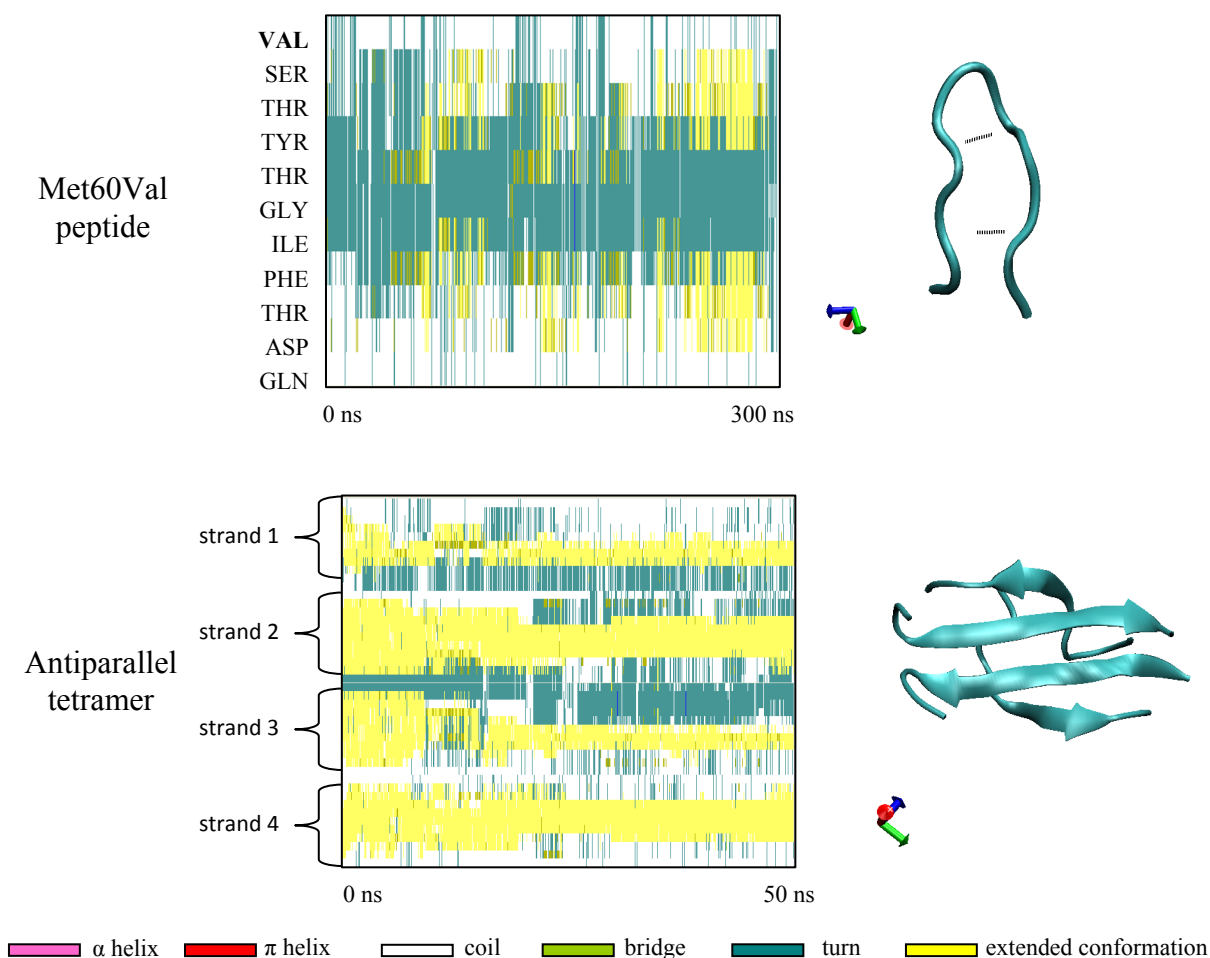


Figure S1. Results of the simulations repeated with reaction field correction for the electrostatic interactions as well as Lennard-Jones cutoff of 1.4 nm. The simulations were conducted for 300ns and 50ns, respectively. Secondary structure evolution plots for valine mutated apoC-II(60-70) peptide and tetramer oligomer in antiparallel arrangement (most stable oligomer), as well as a cartoon representation of their final structure are shown.

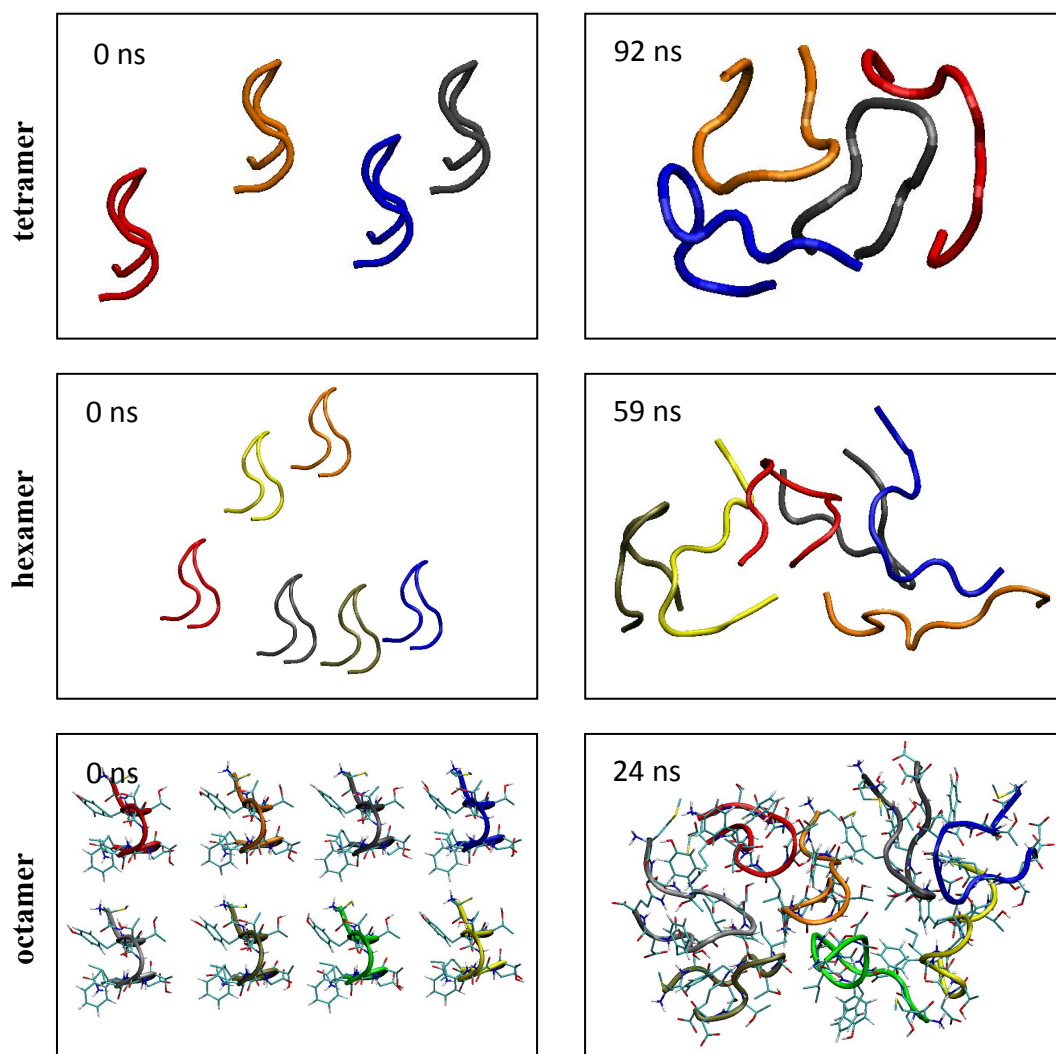


Figure S2. Screen shots showing the starting and final structures of several β -hairpin oligomers of apoC-II(60-70).