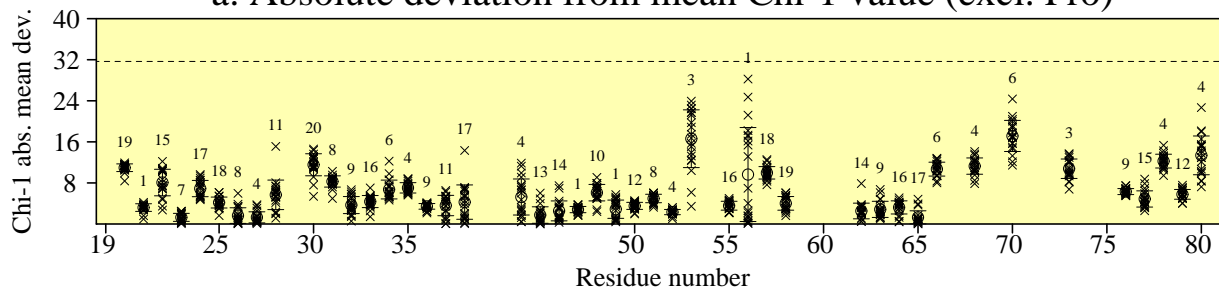
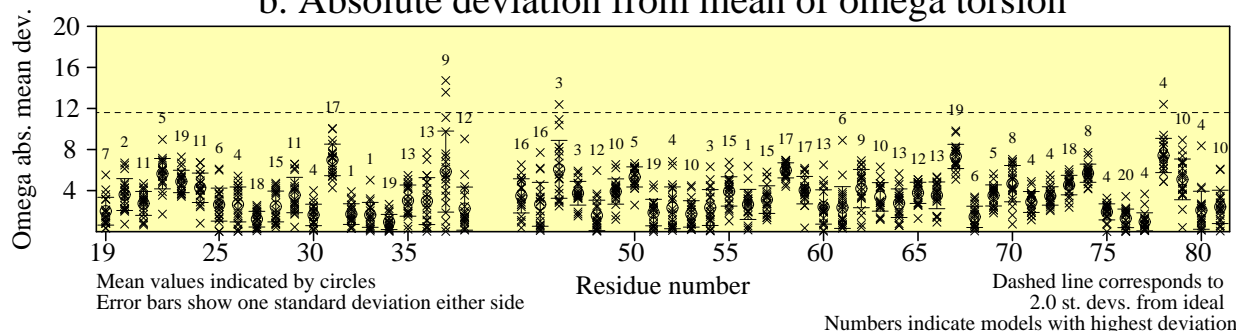


Residue properties HR4527E_R3Cons_em_bcr3 (20 models)**

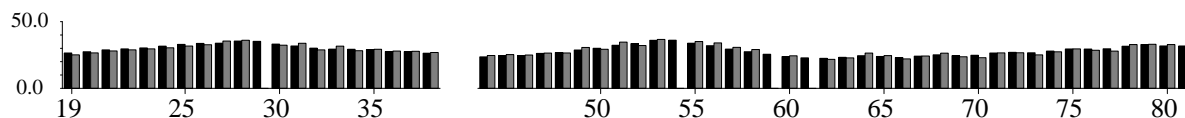
a. Absolute deviation from mean Chi-1 value (excl. Pro)



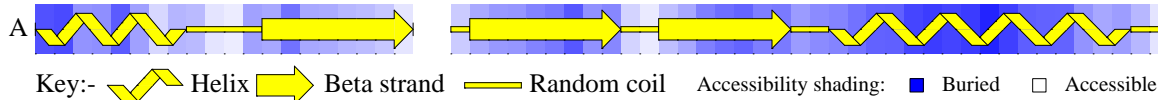
b. Absolute deviation from mean of omega torsion



c. RMS devs from mean coords: main-chain (black) and side-chain (grey)



d. Secondary structure & average estimated accessibility



e. Sequence & average estimated accessibilities



f. Circular variances



g. G-factors

