rmsd before Boxplot using best by rmsd Methods classic-abinitio sade mc ffi9 02 sade mc final sade remc sade remc ffi9 02 sade remc final 112y 2mr9 1acw 1ail 1crn 1enh 1rop 1utg 1zdd 1wqc Protein

12

10

8

6

RMSD (Lower is better)