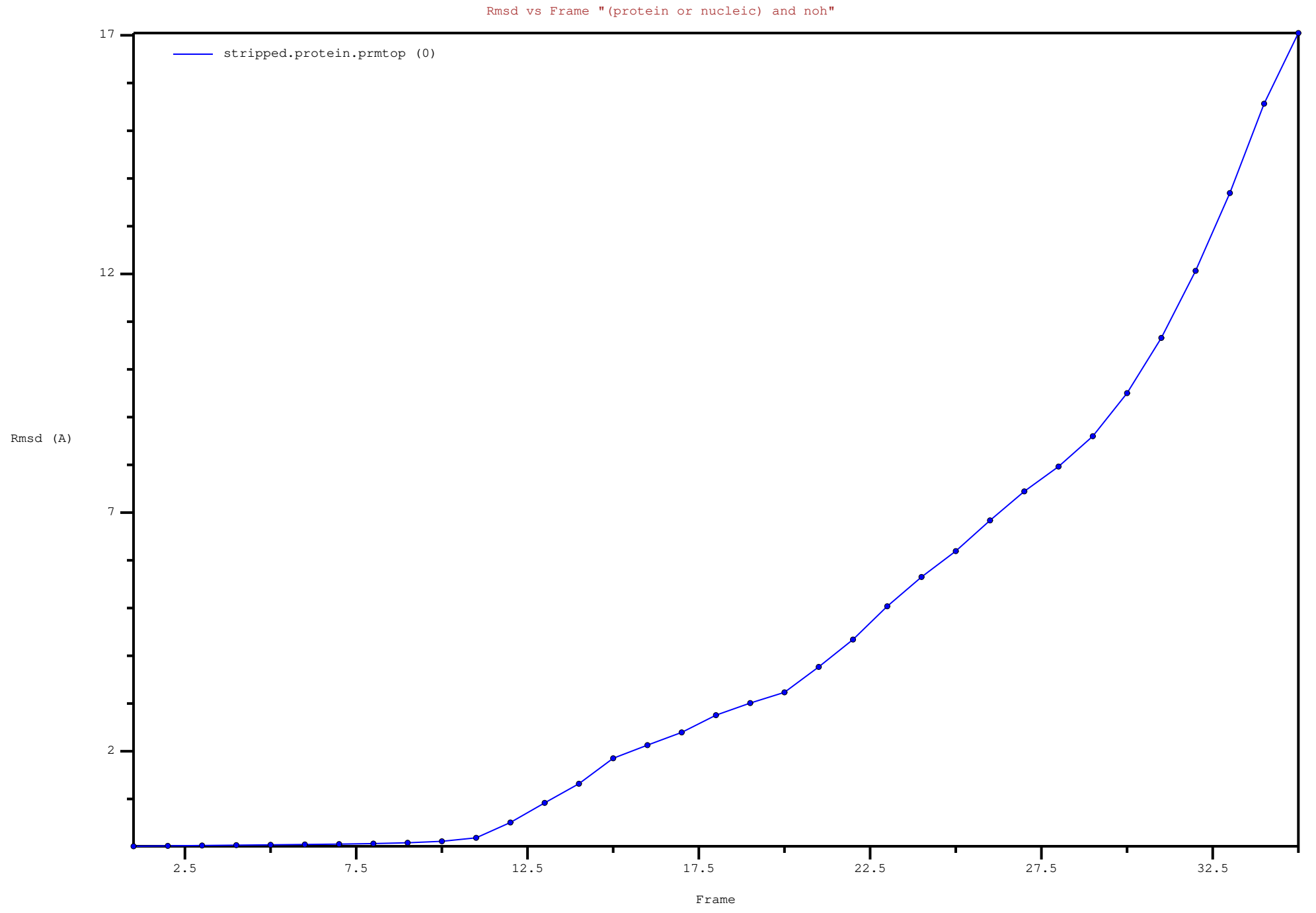
Rmsd vs Frame "(protein or nucleic) and noh"
stripped.protein.prmtop (0)
Rmsd (A)
17
12
7
2
2.5
7.5
12.5
17.5
22.5
27.5
32.5
Frame