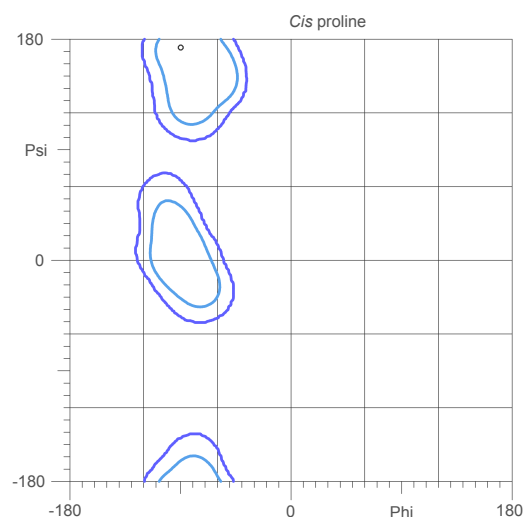
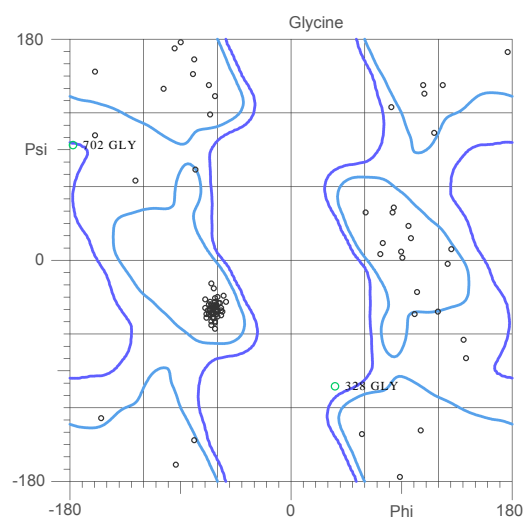
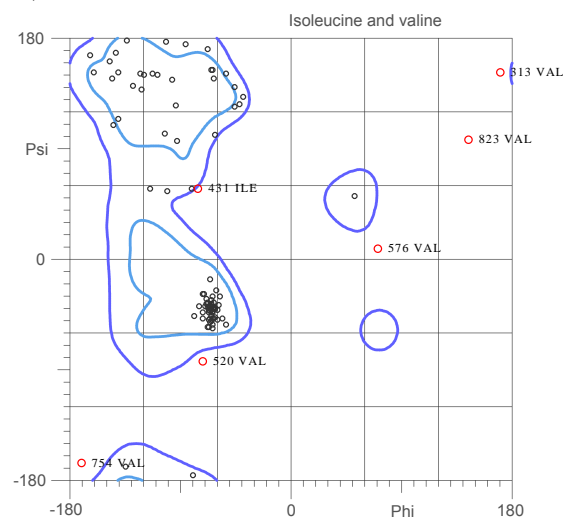


1yq2.B99990039.pdb, model 1



87.3% (894/1018) of all residues were in favoured (98%) regions.

These were 28 outliers (phi, psi):

216 ALA (170.5, 131.1)
216 ALA (190.8, 129.7)
311 ARG (178.1, 103.0)
311 ARG (171.8, 103.0)
328 GLY (166.8, 104.9)
328 GLY (166.8, 104.9)
370 SER (167.7, 170.8)
371 SER (168.9, 168.3)
412 THR (174.8, 108.8)
412 THR (163.7, 172.1)
436 SER (163.7, 172.1)
529 VAL (173.3, 103.8)
529 VAL (169.6, 172.3)
716 VAL (171.8, 94.8)
991 ASP (169.4, 168.9)
991 GLN (162.7, 110.8)
991 SER (176.8, 169.9)
992 GLY (178.2, 94.9)
993 SER (176.2, 161.8)
994 ARG (176.8, 103.0)
994 VAL (171.8, 94.8)
995 ALA (166.4, 99.3)
996 ARG (172.8, 108.7)
998 ASN (172.8, 141.2)
998 ASP (126.7, 74.3)
916 ALA (164.2, 77.9)
916 VAL (167.7, 98.2)
917 THR (167.4, 77.4)
940 ASP (132.2, 124.9)
940 ASP (170.7, 123.7)