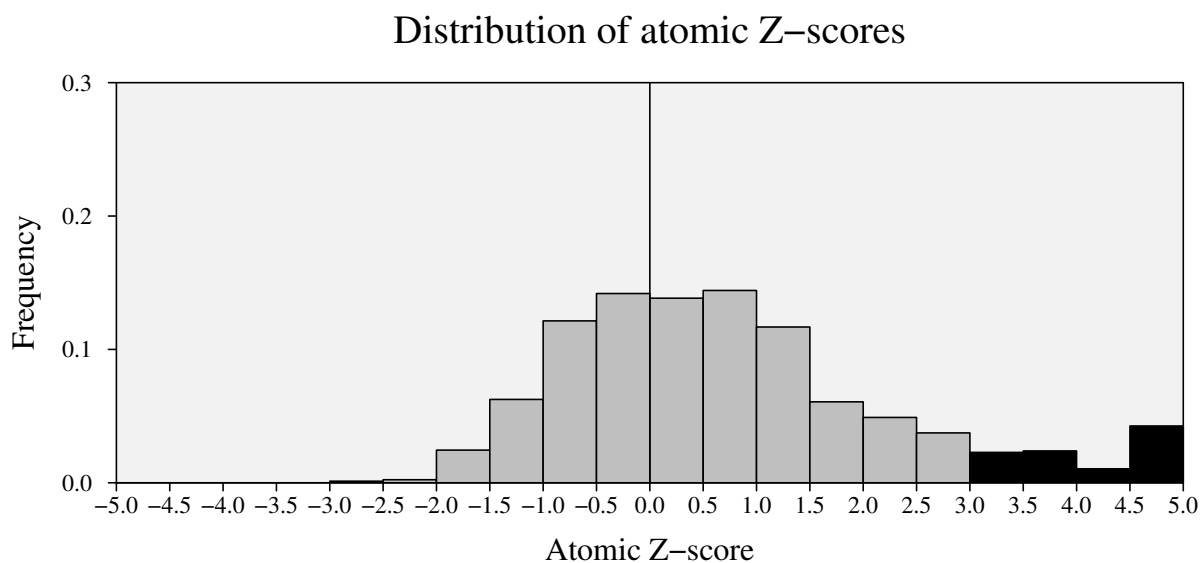
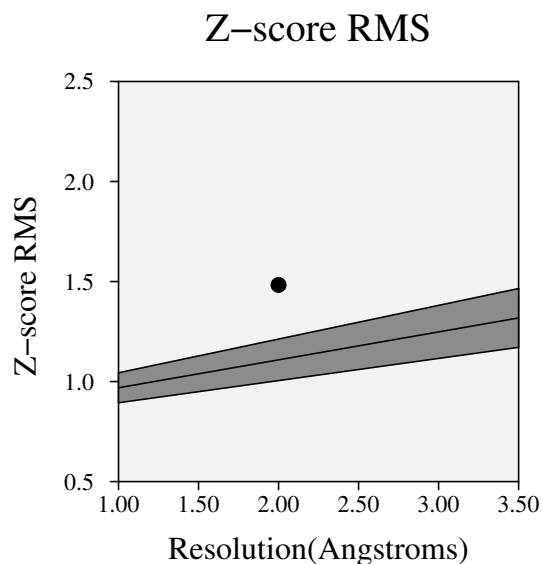
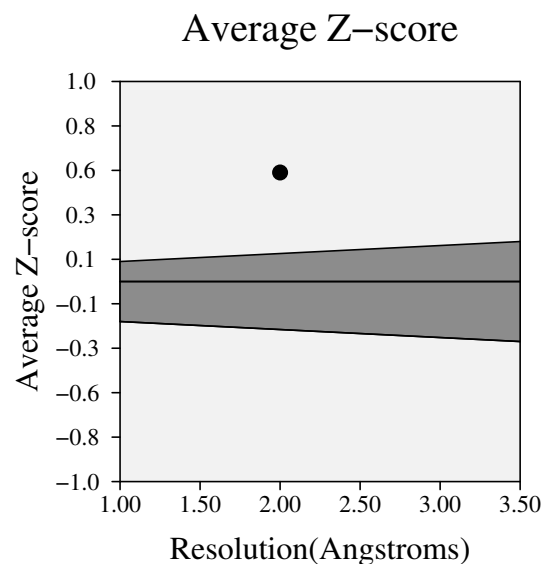


PROVE

saves.pdb

Analysis of entire structure

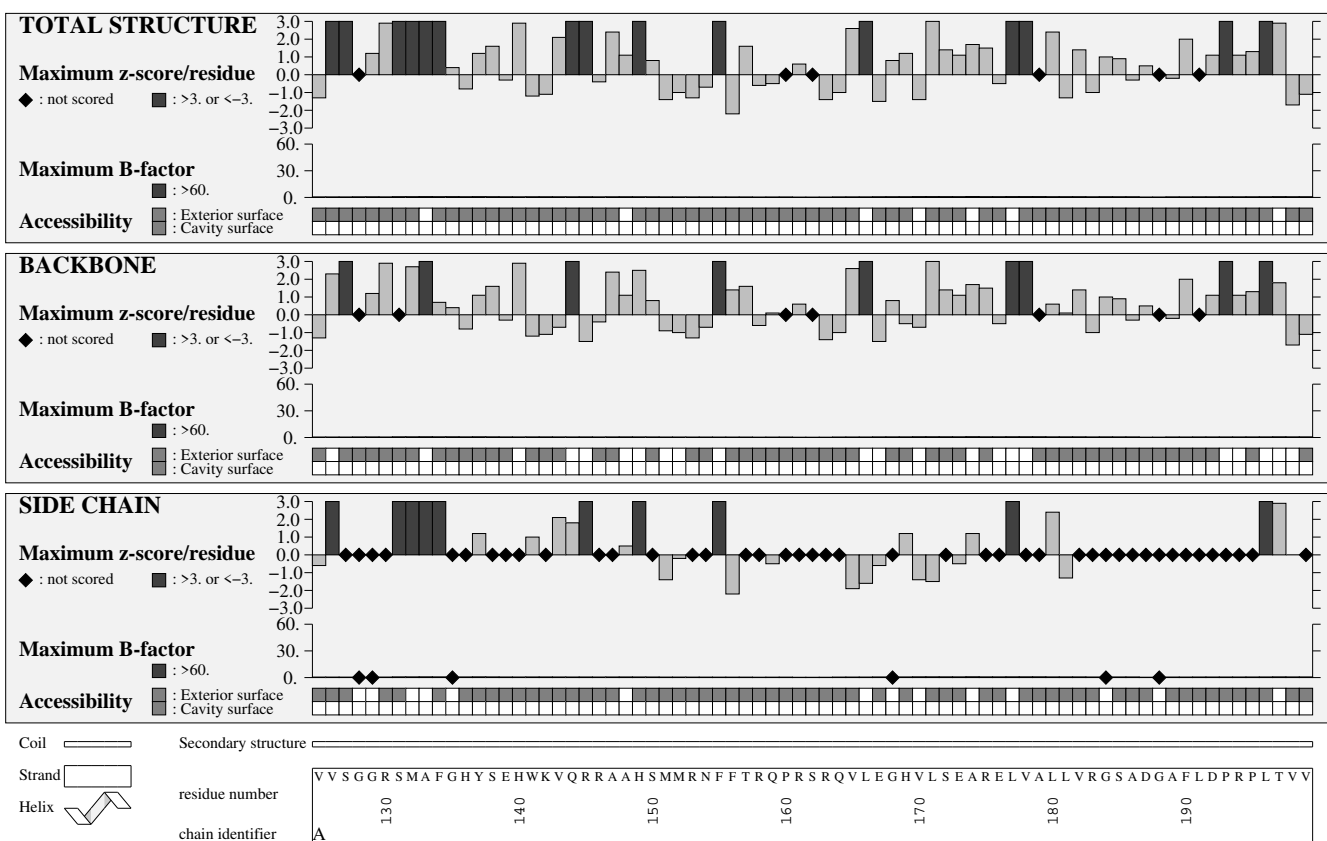
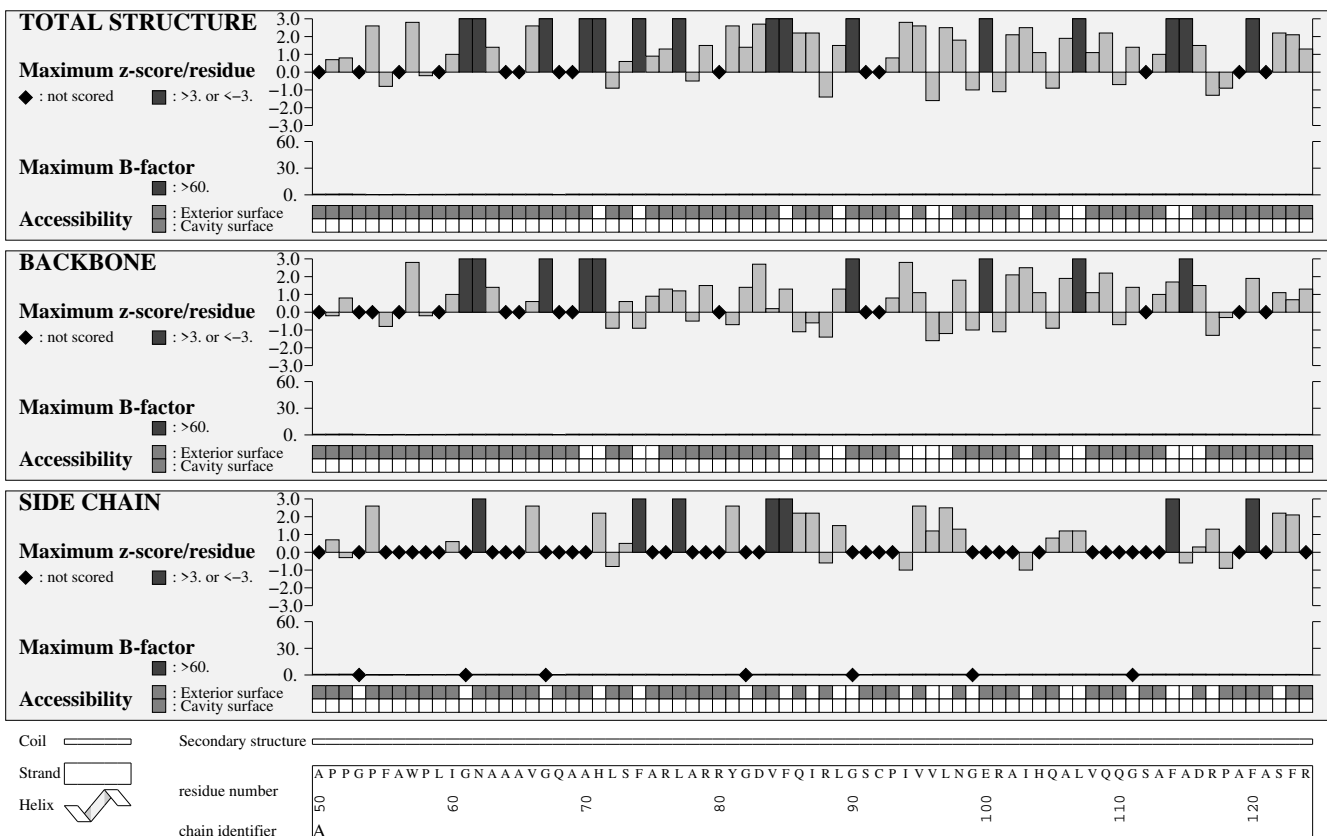


Z-score mean	0.545
Z-score stddev	1.380
Z-score RMS	1.483
# scored atoms	1718
# outliers	92
% outliers	5.400

PROVE

saves.pdb

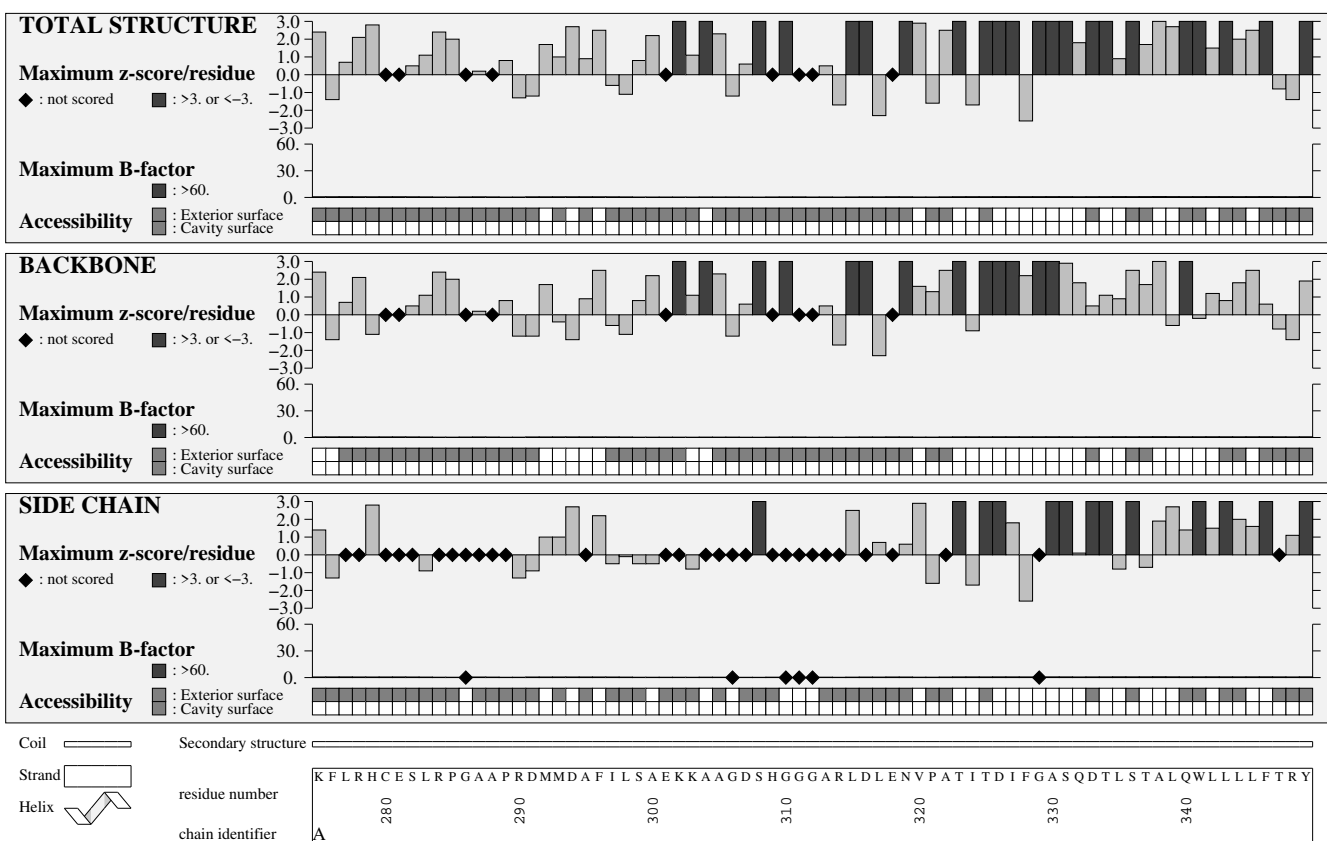
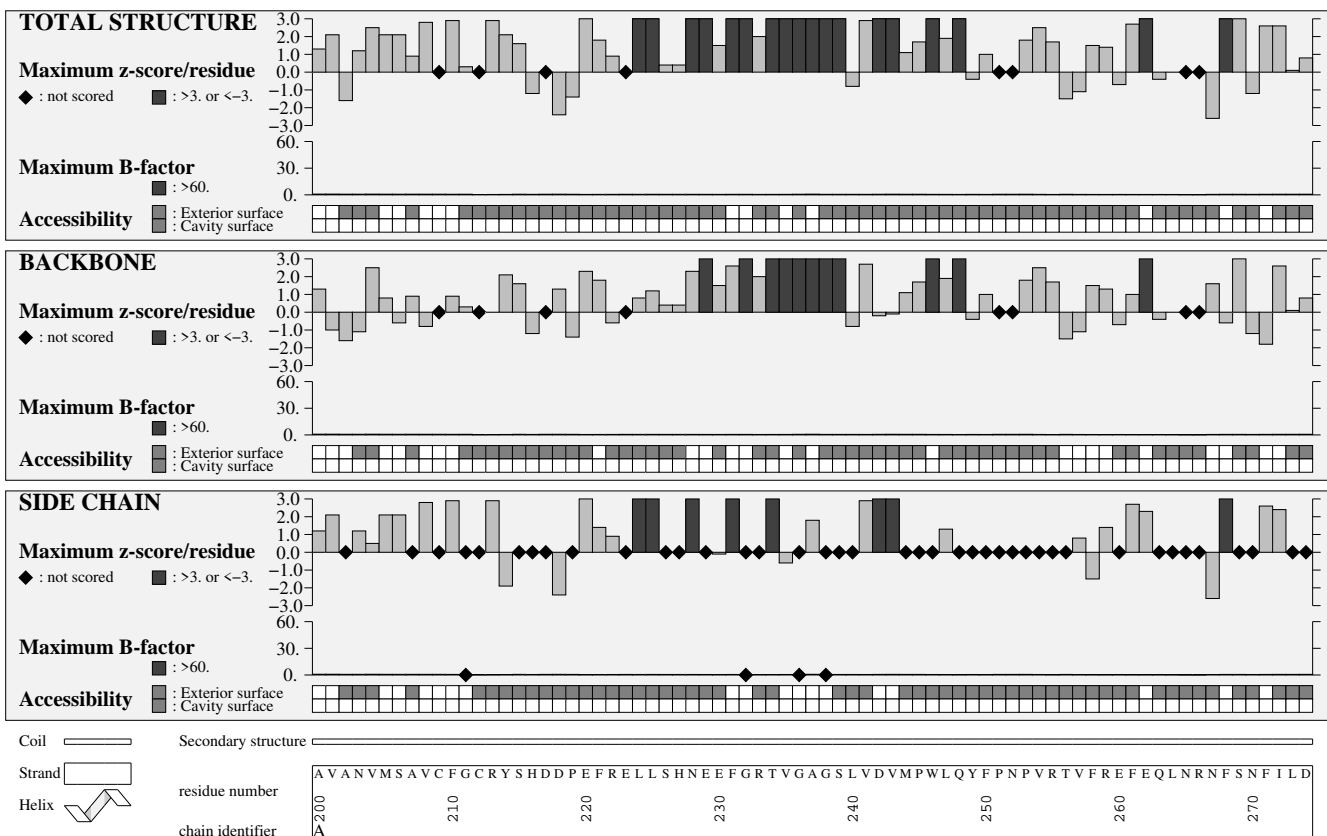
Analysis of residues



PROVE

saves.pdb

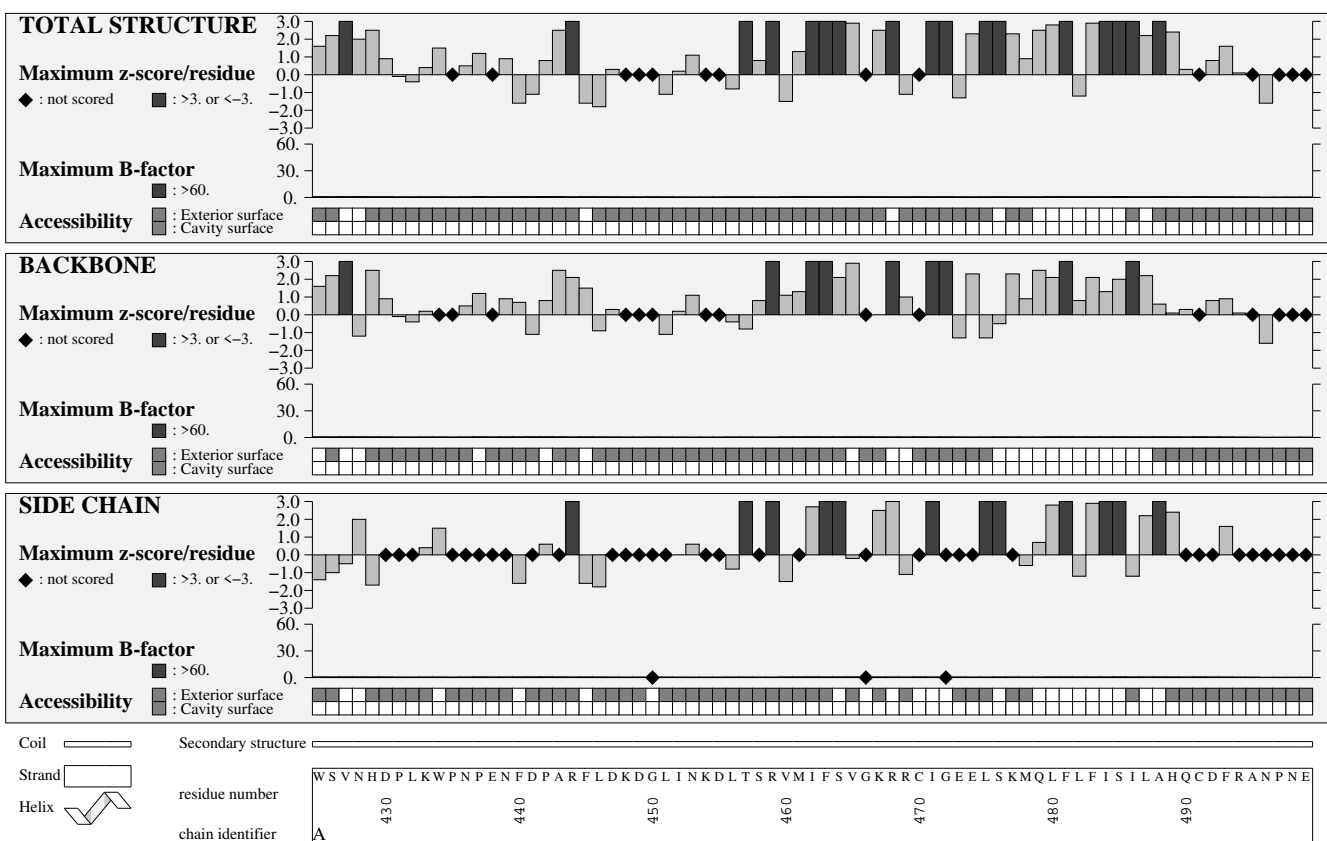
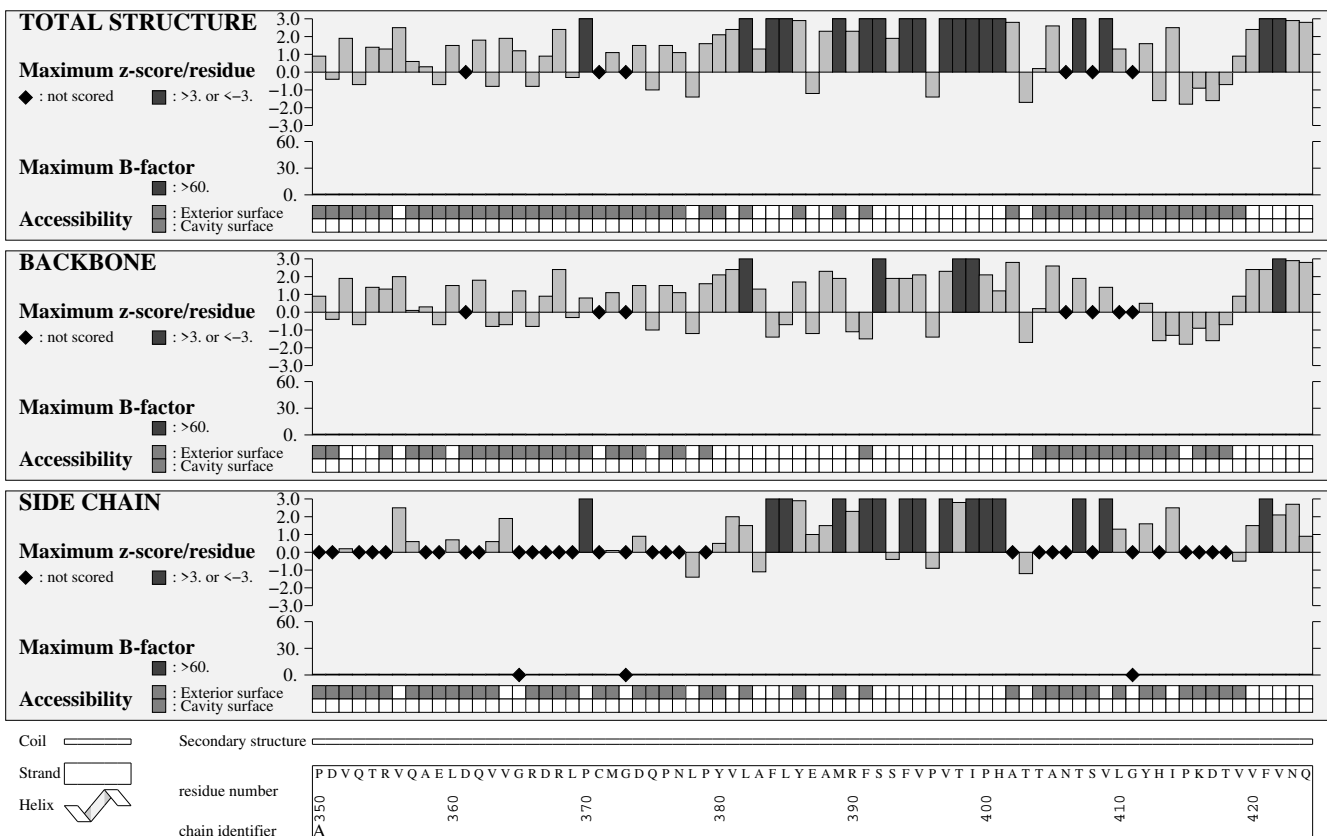
Analysis of residues (2)



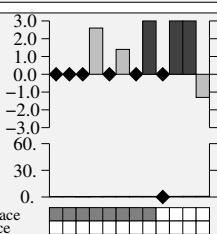
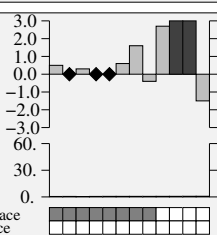
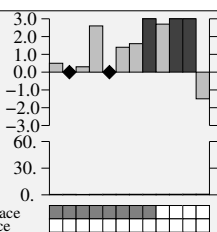
PROVE

saves.pdb

Analysis of residues (3)



Analysis of residues (4)



chain identifier

50A