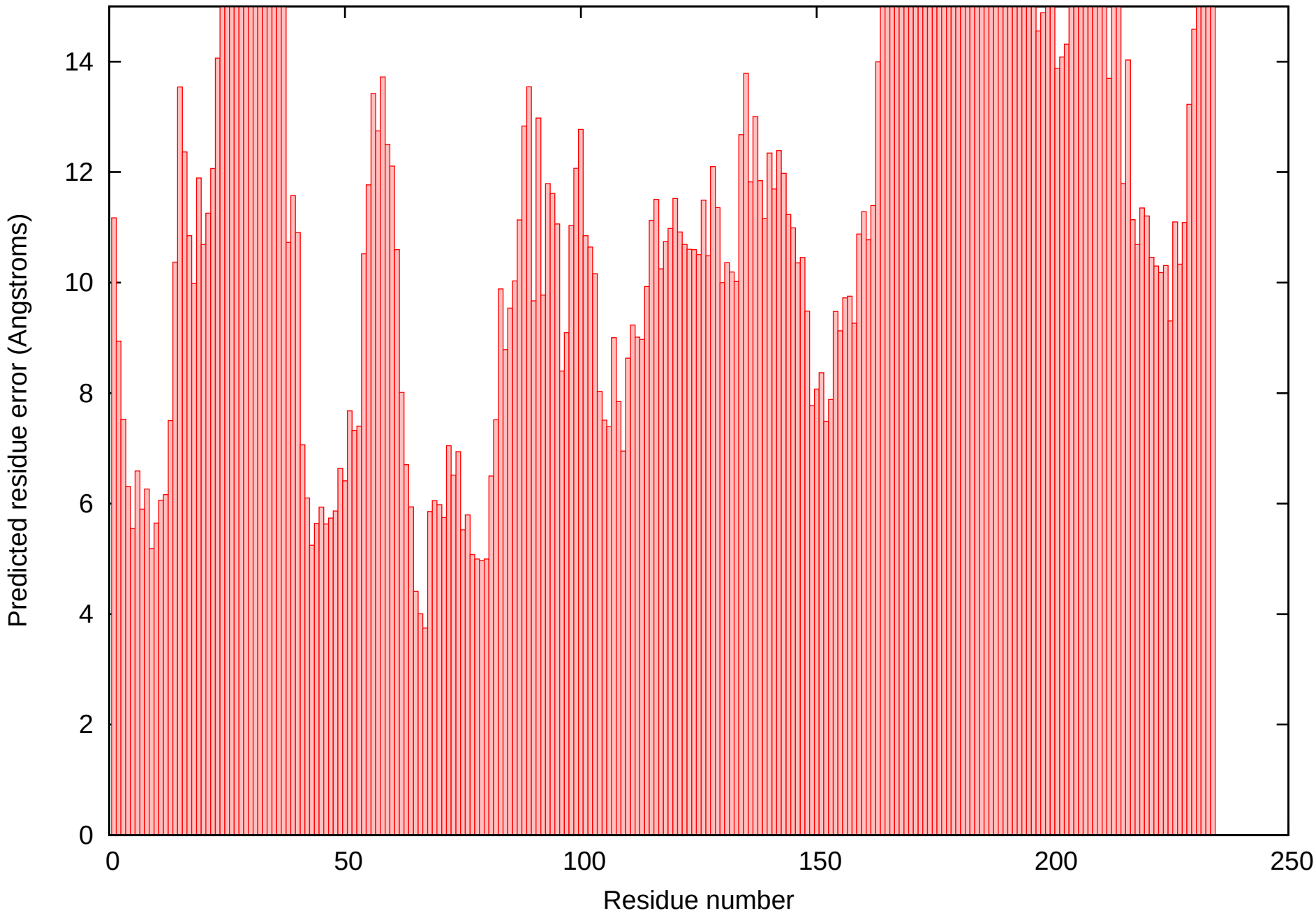
Predicted residue error (Angstroms)
0
2
4
6
8
10
12
14
0
50
100
150
200
250
Residue number