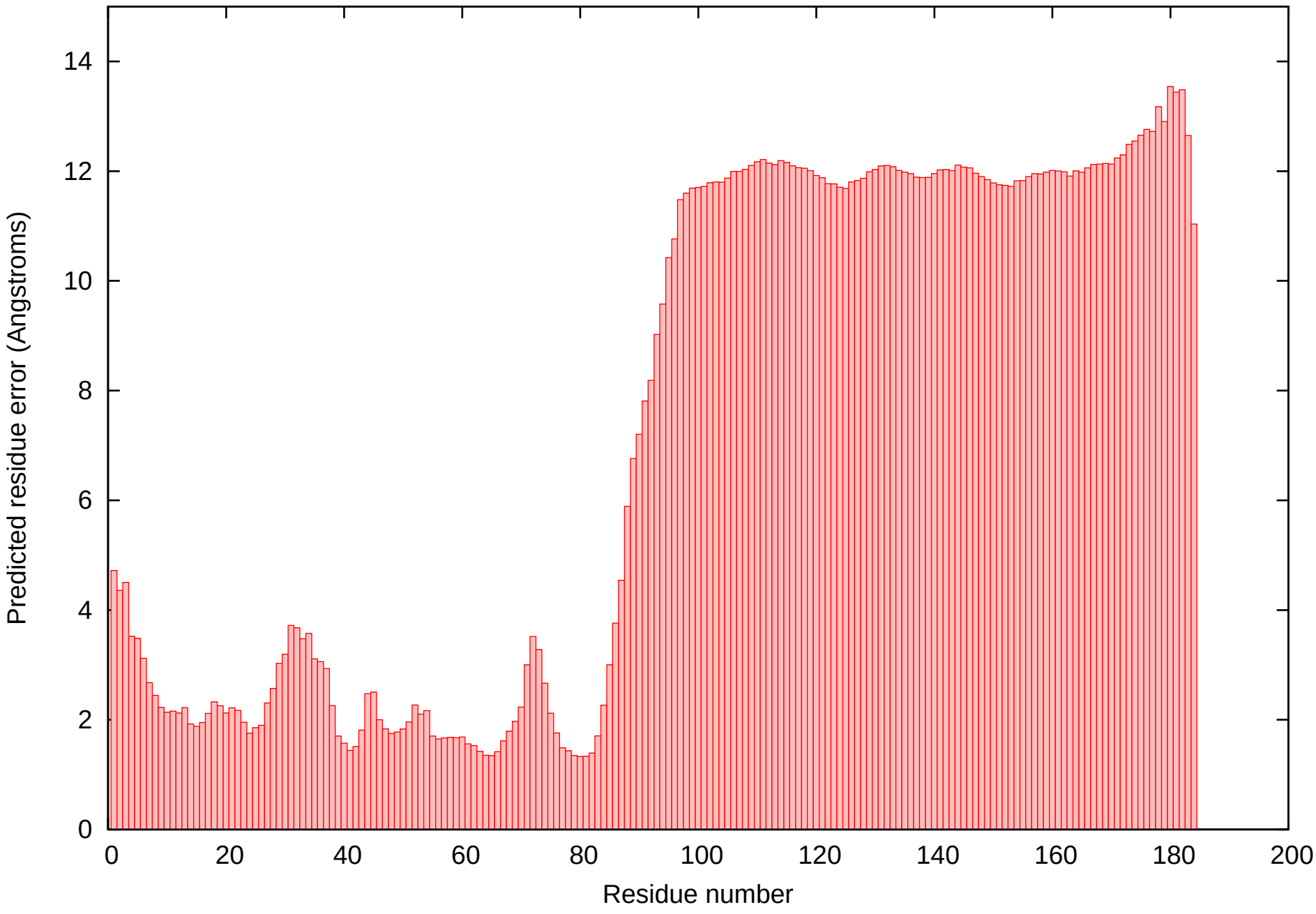
Predicted residue error (Angstroms)
Residue number
0
2
4
6
8
10
12
14
0
20
40
60
80
100
120
140
160
180
200