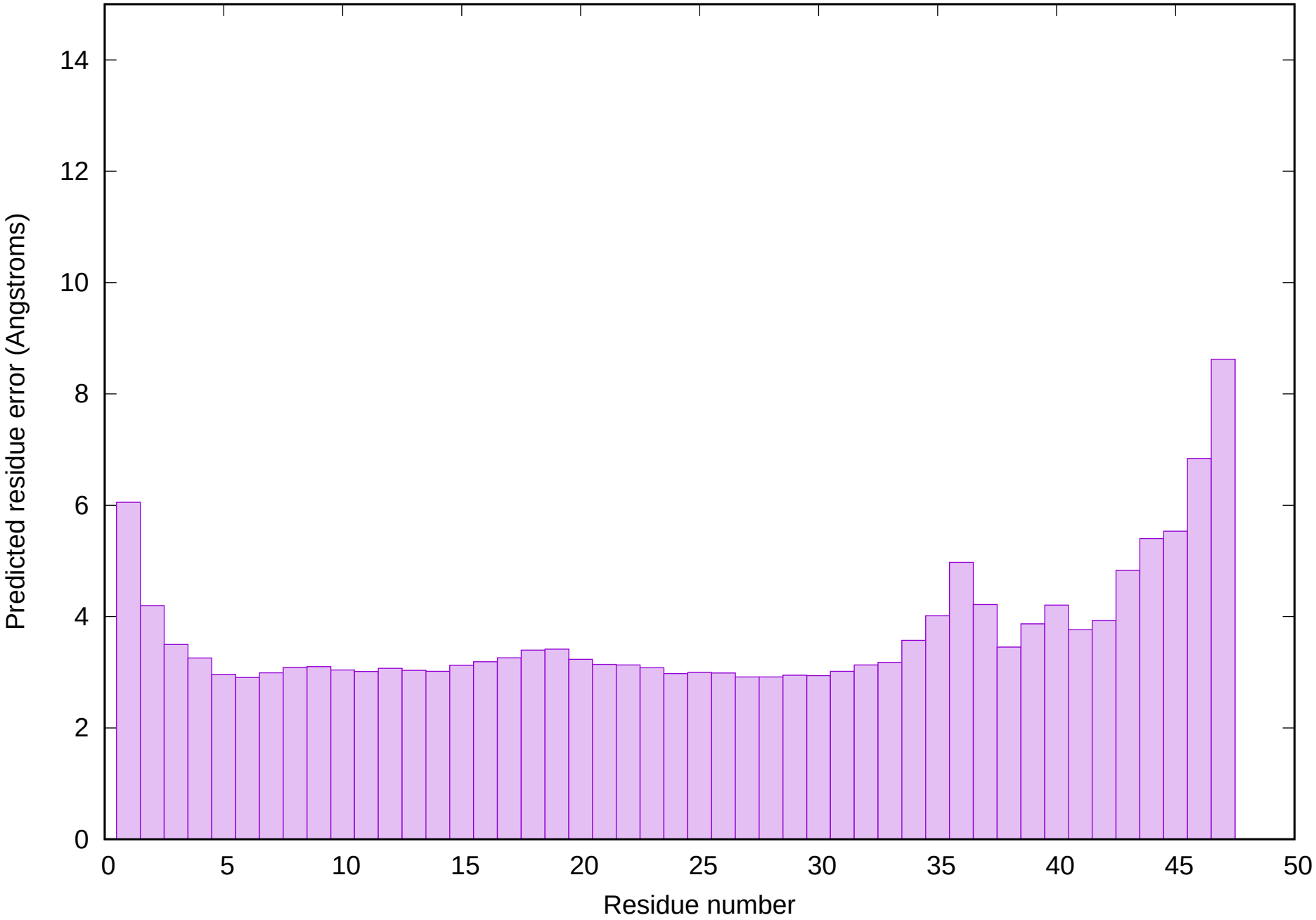
Predicted residue error (Angstroms)
0
2
4
6
8
10
12
14
0
5
10
15
20
25
30
35
40
45
50
Residue number