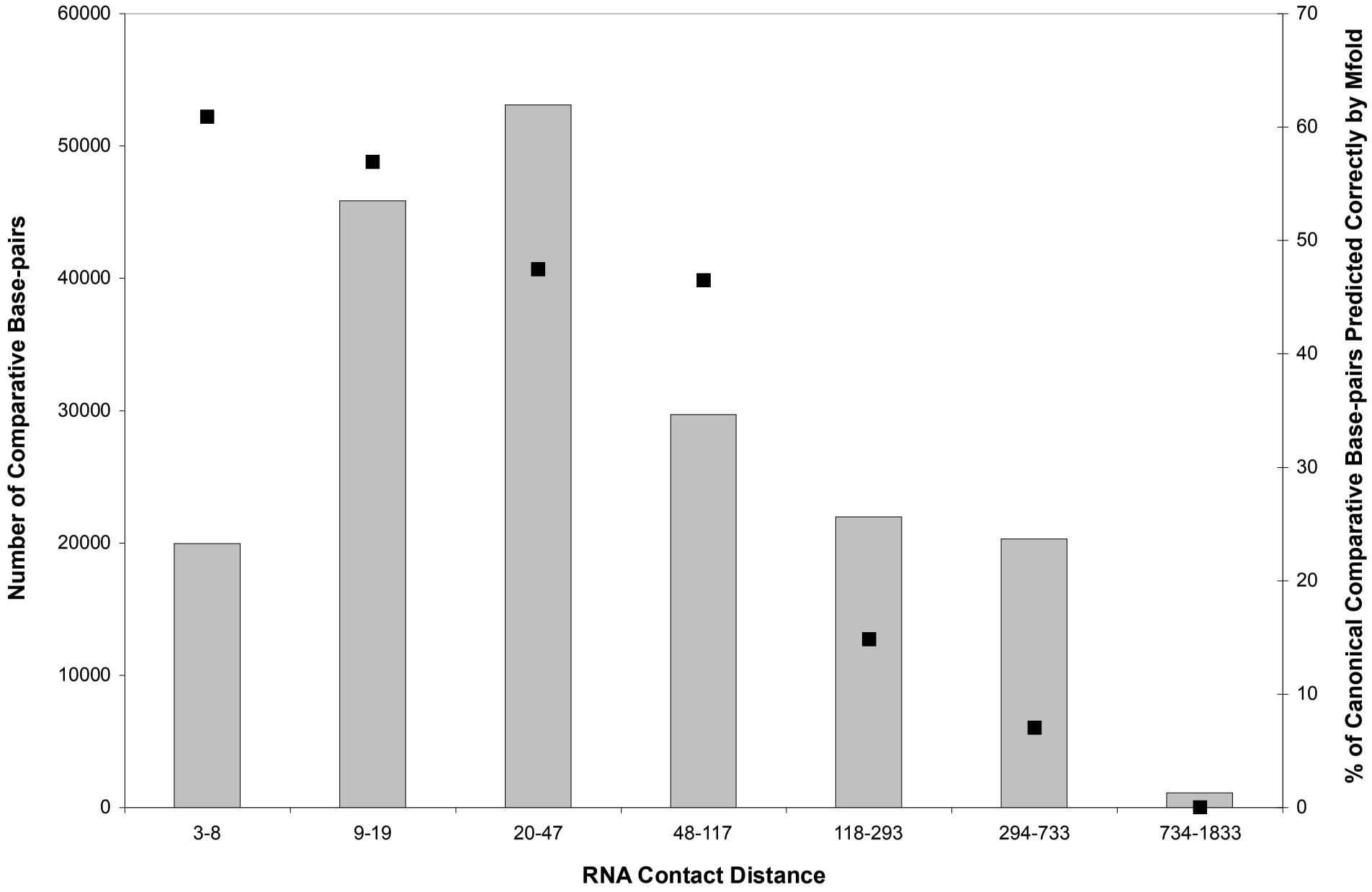
Number of Comparative Base-pairs
60000
50000
40000
30000
20000
10000
0
% of Canonical Comparative Base-pairs Predicted Correctly by Mfold
70
60
50
40
30
20
10
0
3-8
9-19
20-47
48-117
118-293
294-733
734-1833
RNA Contact Distance