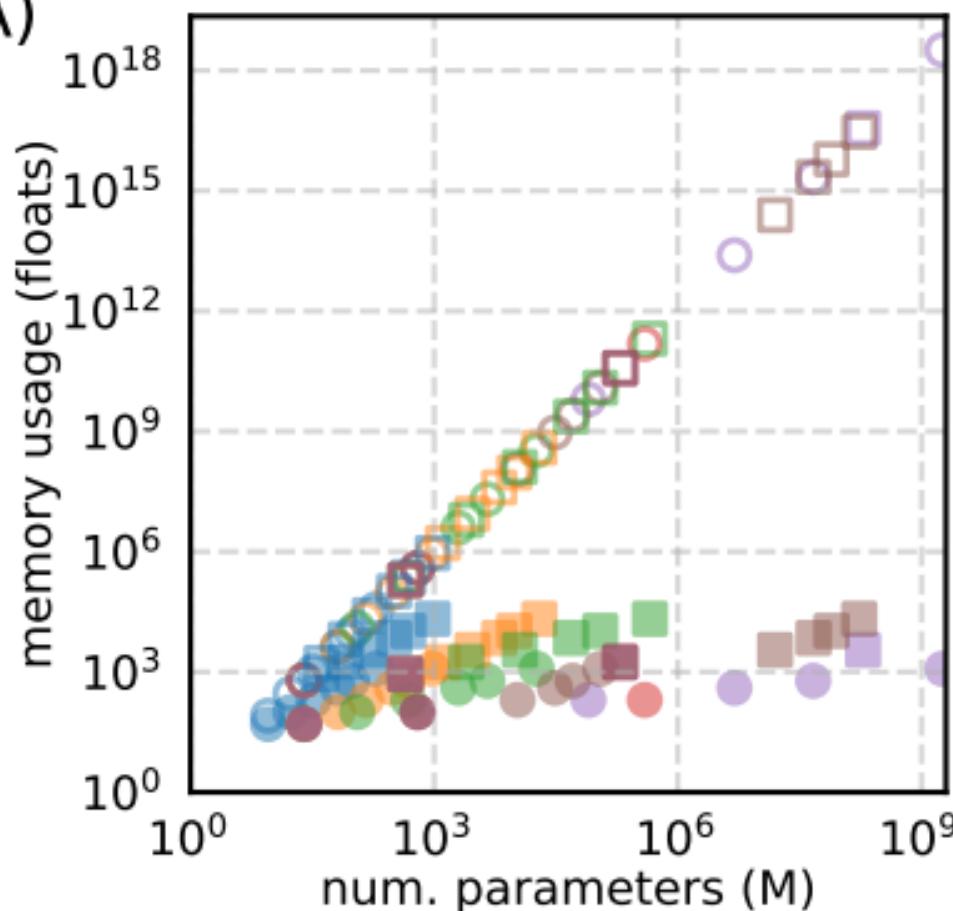
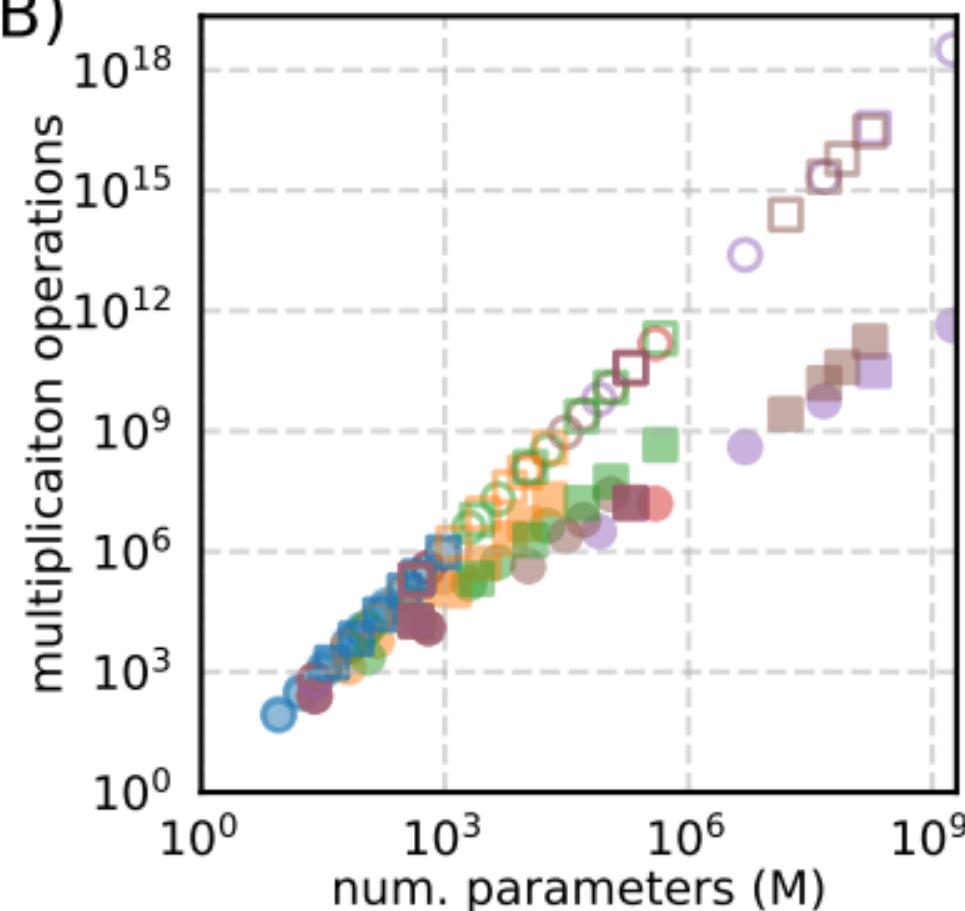


(A)



(B)



model type

additive

neighbor

pairwise

allorder

K-order ($K=5$)K-adjacent ($K=5$)

alphabet size

4 (DNA/RNA)

20 (protein)

algorithm

standard

GaugeFixer