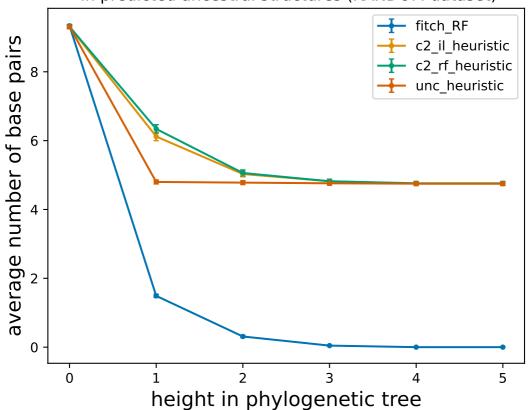
Average maximum number of base-pairs in predicted ancestral structures (RANDOM dataset)



Average maximum number of base-pairs in predicted ancestral structures (RANDOM dataset) fitch RF c2_il_heuristic 🕂 c2 rf heuristic 🕂 unc heuristic height in phylogenetic tree

2.00

1.75

1.50

1.25

1.00

0.75

0.50

0.25

0.00

average number of multi-loops