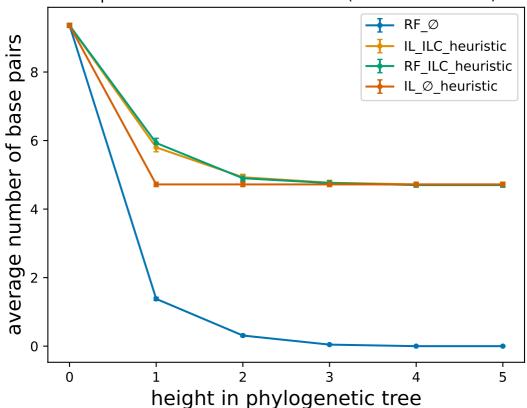
Average maximum number of base-pairs in predicted ancestral structures (RANDOM dataset)



Average maximum number of base-pairs in predicted ancestral structures (RANDOM dataset) RF Ø IL ILC heuristic RF ILC heuristic IL\_Ø\_heuristic height in phylogenetic tree

2.00

1.75

1.50

1.25

1.00

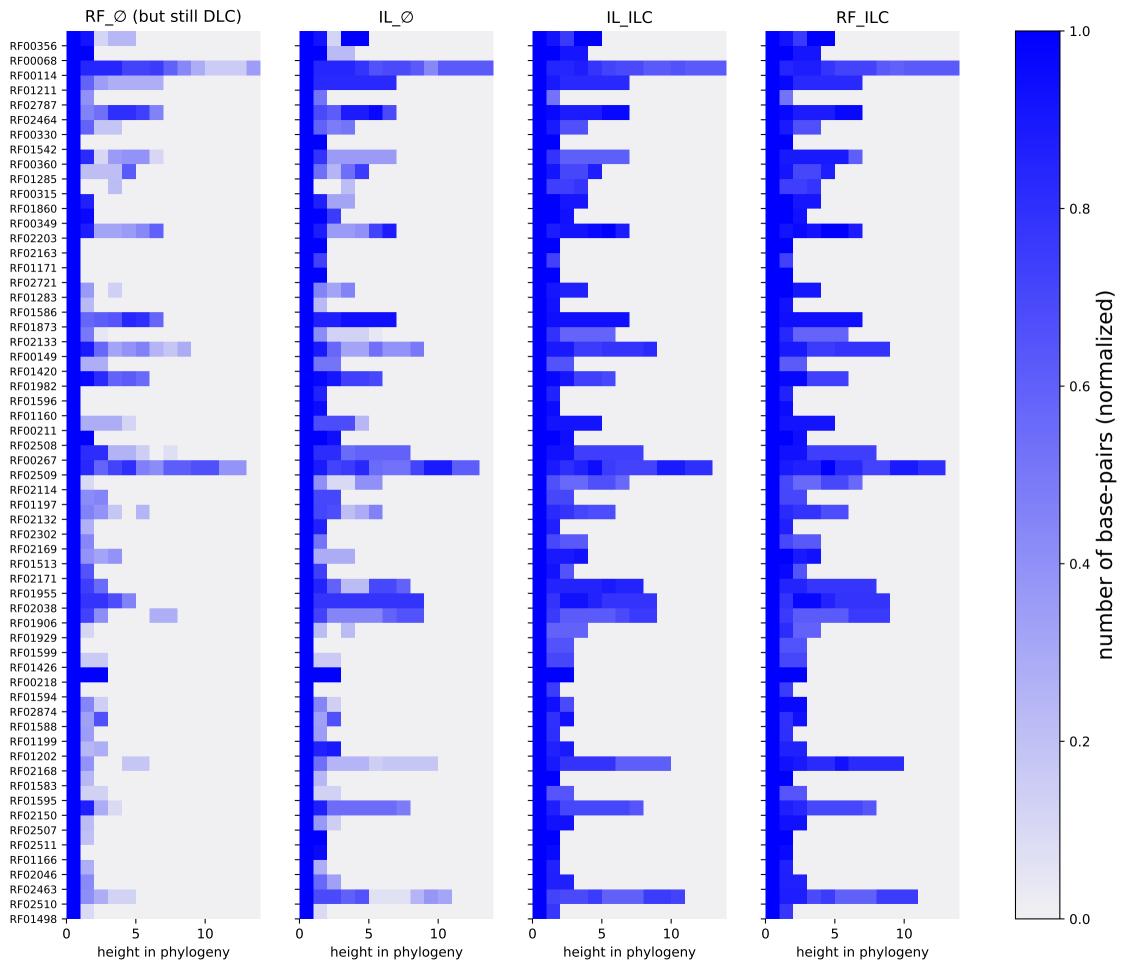
0.75

0.50

0.25

0.00

average number of multi-loops



## Comparison of the number of base-pairs predicted at the root of the phylogeny for each RFAM family

