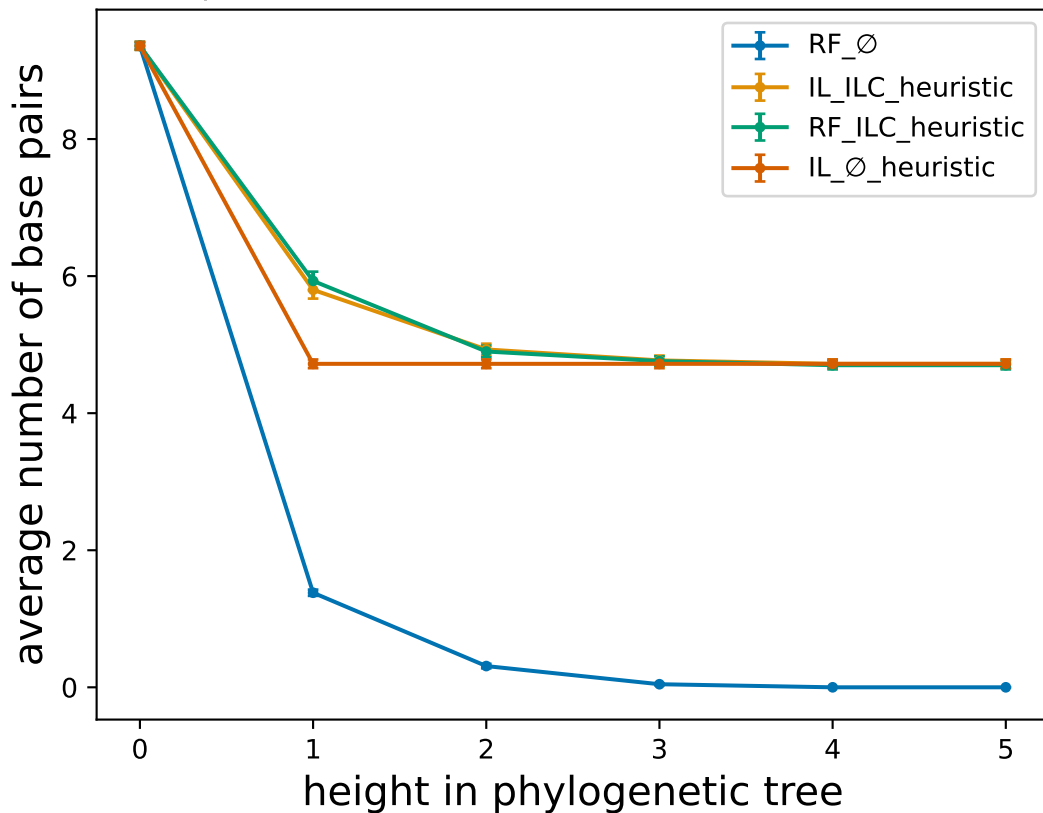
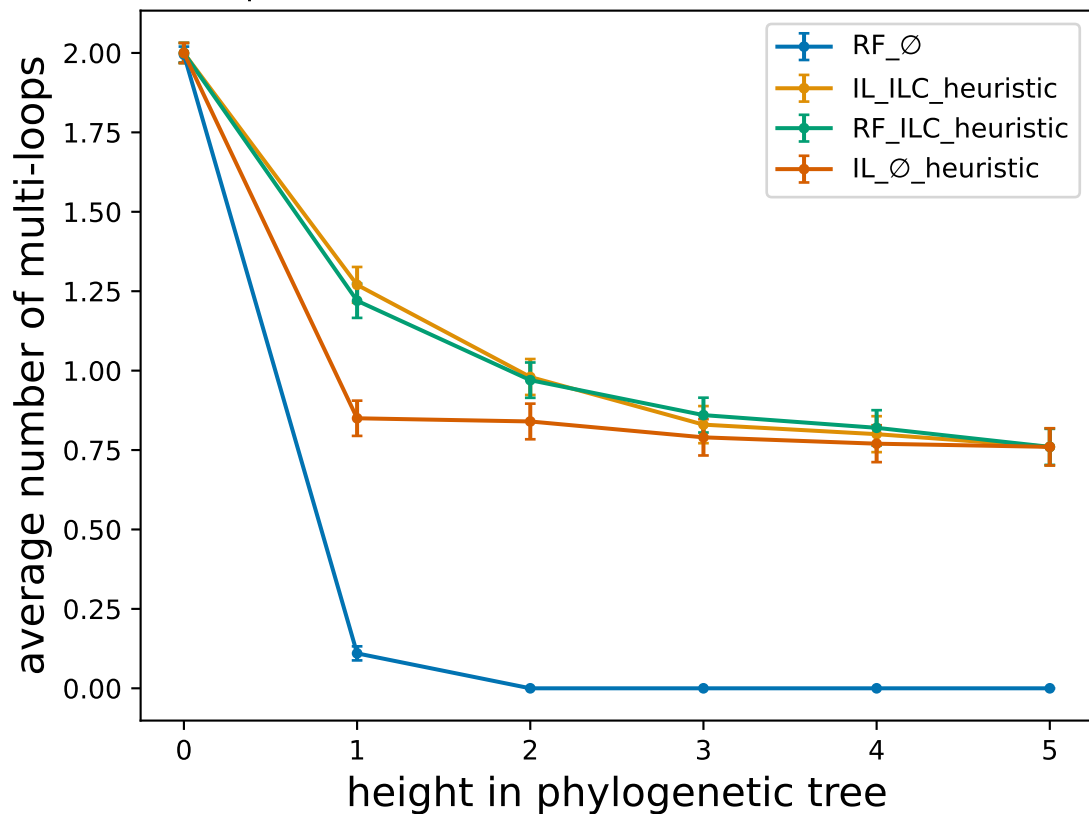
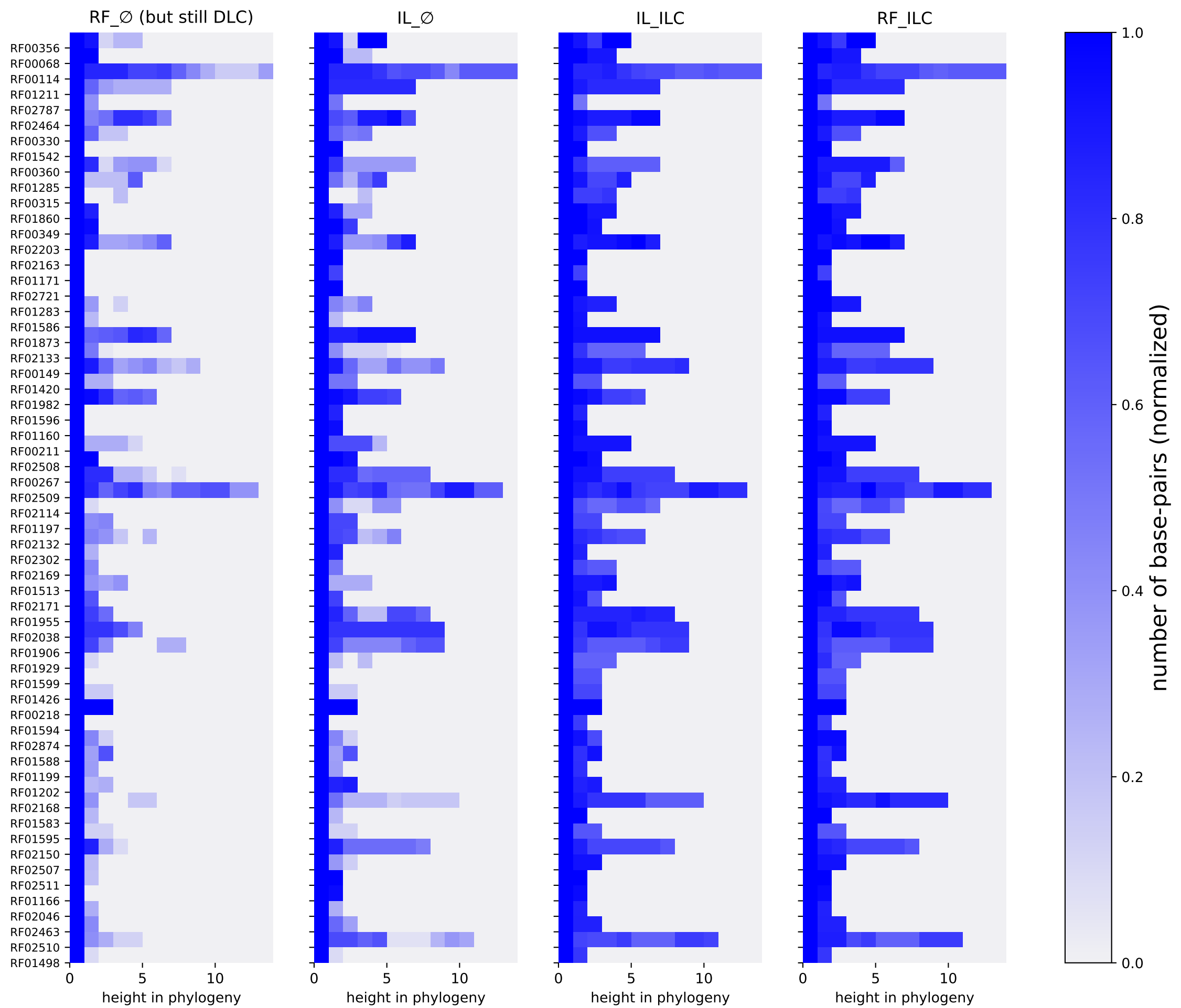


Average maximum number of base-pairs
in predicted ancestral structures (RANDOM dataset)



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





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

