QfO benchmarking - QfO:2018-07-07_G_STD_Eukaryota 0.24 0.22 QfO:RF_DISTANCE 0.20 0.18 0.16 better 0.14 1000 2000 3000 5000 6000 4000 QfO:NR_COMPLETED_TREE_SAMPLINGS BBH OMA_GETHOGs_2.0 OrthoFinder_2_BLAST__MSA PANTHER_14.1_LDO_only SonicParanoid_most Ensembl_Compara SonicParanoid_sensitive OMA_Groups_2.0 OrthoFinder_2_DIAMOND__MSA PANTHER_14.1_all Hieranoid2 OMA Pairs 2.0 OrthoFinder 2 Defaults DIAMOND MSA RSD eggNOG

OrthoInspector_3.0

SonicParanoid_default

OrthoFinder_2_BLAST__DendroBLAST_

InParanoid