

Scatterplot: Distribution of site dependencies for all trees

phylip_file: ./data/128/128.phy.uniq

sample_root: midpoint

sample_trees: 1

number_of_partitions: 34

number_of_taxa: 128

number_of_sites: 19866

program_runtime: 1 minutes and 33 seconds

data_file: ./output_site_dependencies_count/2015-12-18 11-24-38 data.csv

graph_file_name: graphs/data-128 2015-12-18 11-24-38

./data/128/parsimony_trees/RAxML_parsimonyTree.T8.RUN.0

./data/128/parsimony_trees/RAxML_result.T8.RUN.0

./data/128/random_trees/RAxML_result.S7.RUN.0

count

count

count

sorted sites

partition

- 12S_rRNA
- 16S_rRNA
- ADORA3
- ADRB2
- APOB
- APP
- ATP6
- ATP7A
- ATP8
- BDNF
- BMI
- BRCA1
- CNR1
- COX1
- COX2
- COX3
- CREM
- Cytb
- EDG1
- IRBP
- ND1
- ND2
- ND3
- ND4
- ND4L
- ND5
- ND6
- PLCB4
- PNOC
- RAG1
- RAG2
- tRNAs
- TYR
- ZFX