Boxplot sorting: Comparison of MSA sorting per tree phylip_file: ./data/10c/10.phy.uniq sample_root: midpoint sample trees: 100

sample_root. Midpoint
sample_trees: 100
program_runtime: 55 minutes and 53 seconds
data_file: ./output_sorting1/2015–11–14 20–28–39 data.csv
graph_file_name: graphs/data=10c 2015–11–14 20–28–39

