

Quick Notes on Phylip usage

Phylip tools used

Seqboot
Protpars
Consense

Files used with Seqboot

phylip_alignment_for_Phylip

Commands used

% seqboot

> phylip_alignment_for_Phylip

*The output of Seqboot is always a file called: outfile. To simplify this folder, the name of this file was altered.

Results

result_from_Seqboot_step1

Files used with Protpars

result_from_Seqboot_step1

Commands used

% protpars

> result_from_Seqboot_step1

*The output of Protpars is always two files: outfile & outtree. To simplify this folder, the name of both these files was altered.

Results

file_result_from_Protpars_step2
tree_result_from_Protpars_step2

Files used with Consense

tree_result_from_Protpars_step2

Commands used

% consense

> tree_result_from_Protpars_step2

*The output of Consense is always two files: outfile & outtree. To simplify this folder, the name of both these files was altered.

Results

consense_result_file_step3
consense_result_tree_step3