

FN tree error

1.00
0.75
0.50
0.25
0.00

aat

Acetyltransf

adh

aldosered

biotin

blmb

ghf13

gluts

hla

hom

myb

p450

PDZ

Rhodanese

rrm

rvp

sdr

tRNA-synt

zf-CCHH

Average

Clustal-Omega Muscle MAFFT UPP(Default)

