

Mean alignment error

0.75
0.50
0.25
0.00

Acetyltransf

aldosered

biotin

blmb

ghf13

hla

hom

myb

PDZ

Rhodanese

rrm

rvp

sdr

tRNA-synt

zf-CCHH

Average

SEPP(Default) UPP(Default)

