Summary of blast-based alignments, Jan 13 2021

PNMA family

Model 0 dN/dS:
ARC: 0.032
CCDC8: 0.145
ZCCHC12/18 12: 0.252
ZCCHC12/18 18: 0.535

Legend:
- ■ Intact
- ■ (red) Intact, used in PAML
- ⊠ Pseud
- ■ (gray) Truncated
- □ Absent
- Not analyzed