

PQQAEGVLRFELEGGRRARRGRIPQRLRLAWYGGPPVHSGERWRLAVKLKRPAGLVNLLQAFDHEAWLLAQRIGATGTVKDQLIV.
PDRRGDVLRFELEDIHSRHAGTLP SRLRLAWYGGPEIRSGERWLARLKRPSGMVNPSSAFDYEAWLLARRIGATGTIKAGERIAE

[illegible][illegible]

TREHAMLYDAGPRYGDS^DLGERWV^FPSLR^QLG^TERLDL^MLSHAD^SDHAGGALAY^QR^GMPV^GRW^SSG^EPER^LPAY^LPAR^SCR^TGE
 TREHAMLYDAGPRLGDS^DLGARVVLPSLRK^LG^TARLDL^MLSHAD^SADHAGGADAV^QRGL^PVAR^LVSG^DPEGL^PAY^LLP^AEP^CAS^SGE
 TRDHALLYDAGPRYGDF^DIGERVVFPSLR^QLG^TERLDL^MLSHAD^SSDHAGGALA^TQR^AMPV^GAV^LSG^EPER^LLP^AAL^DAR^SCR^TGQ

593
595

logo

NTDB id 66265 PCAU RS08020 WP 096373427.1
NTDB id 1199 PSJM300 12650 AFN78592.1
consensus



678
680

logo

NTDB id 66265 PCAU RS08020 WP 096373427.1
NTDB id 1199 PSJM300 12650 AFN78592.1
consensus



743
741

-  non conserved
-  similar
-  ≥ 50% conserved