

logo

NTDB id 318306 D8S77 RS01420 WP 024623426.1

NTDB id 475 HSISS4 RS07005 WP 021143809.1

NTDB id 324 STU RS16140 WP 011226306.1

NTDB id 292 STER RS06940 WP 011681419.1

consensus

KKKSKWLAAGVALLSVGVLAACSSKSNSTSG

KKKSKWLAAGVALLSVGALAACSSKSNSTSG

KKKSKWLAAGVALLSVGALAACSSKSNSTSG

KKKSKWLAAGVALLSVGALAACSSKSNSTSG

!!!!!!!!!!!!!!

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consensus

PSVAEDWSVSKDGLTYTYKIRKGYKMYTSEGEYYAVNTAKDFVTGLKHAADSKAGGLYLVDQSLAGLSDYLSGTTNKDFS

PSVAEDWSVSKDGLTYTYKIRKGYKMYTSEGEYYAVNTAKDFVTGLKHAADSKAGGLYLVDQSLAGLSDYLSGTTNKDFS

PSVAEDWSVSKDGLTYTYKIRKGYKMYTSEGEYYAVNTAKDFVTGLKHAADSKAGGLYLVDQSLAGLSDYLSGTTNKDFS

PSVAEDWSVSKDGLTYTYKIRKGYKMYTSEGEYYAVNTAKDFVTGLKHAADSKAGGLYLVDQSLAGLSDYLSGTTNKDFS

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consensus

VGVKALDDKTVQYTLNKPESYWNKTTYSLFVPVNAFLKSKGKDFGT.TDPTSSILVNGAYFLSAFTSKSSMEFHKNENY

VGVKALDDKTVQYTLNKPESYWNKTTYSLFVPVNAFLKSKGKDFGT.TDPTSSILVNGAYFLSAFTSKSSMEFHKNENY

VGVKALDDKTVQYTLNKPESYWNKTTYSLFVPVNAFLKSKGKDFGT.TDPTSSILVNGAYFLSAFTSKSSMEFHKNENY

VGVKALDDKTVQYTLNKPESYWNKTTYSLFVPVNAFLKSKGKDFGT.TDPTSSILVNGAYFLSAFTSKSSMEFHKNENY

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consensus

WDKKNVHFDKIKESYDGSDDQALVRGFTDGAYSEFARVYPTSSNYAVEKKYKDNLYYTAPGASTSAIGNLDRQSYKYT

WDKKNVHFDKIKESYDGSDDQALVRGFTDGAYSEFARVYPTSSNYAVEKKYKDNLYYTAPGASTSAIGNLDRQSYKYT

WDKKNVHFDKIKESYDGSDDQALVRGFTDGAYSEFARVYPTSSNYAVEKKYKDNLYYTAPGASTSAIGNLDRQSYKYT

WDKKNVHFDKIKESYDGSDDQALVRGFTDGAYSEFARVYPTSSNYAVEKKYKDNLYYTAPGASTSAIGNLDRQSYKYT

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consensus

AKKTDSEKSTKKALLNKDFRQAINFAIDRKAQSQNGKDGLATLAYRNLFVPSDFVSAGDKTFGDLVTEKMSSYGDEWS

AKKTDSEKSTKKALLNKDFRQAINFAIDRKAQSQNGKDGLATLAYRNLFVPSDFVSAGDKTFGDLVTEKMSSYGDEWS

AKKTDSEKSTKKALLNKDFRQAINFAIDRKAQSQNGKDGLATLAYRNLFVPSDFVSAGDKTFGDLVTEKMSSYGDEWS

AKKTDSEKSTKKALLNKDFRQAINFAIDRKAQSQNGKDGLATLAYRNLFVPSDFVSAGDKTFGDLVTEKMSSYGDEWS

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consensus

GVNLFADQDGLYNAEKAKTEFAKAKALQADGVQFPIHLDLPVDQSSKLNVAQAQSLKQTIKSLGSENVVIDYNQLSSD

GVNLFADQDGLYNAEKAKTEFAKAKALQADGVQFPIHLDLPVDQSSKLNVAQAQSLKQTIKSLGSENVVIDYNQLSSD

GVNLFADQDGLYNAEKAKTEFAKAKALQADGVQFPIHLDLPVDQSSKLNVAQAQSLKQTIKSLGSENVVIDYNQLSSD

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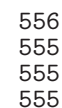
477

476

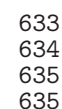
476

476

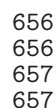
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-  non conserved
-  similar
-  > 50% conserved