

logo

NTDB id 342446 ETT45 RS01360 WP 002992224.1	MKKSKWLAAGVALLSVGVLAACSSKSNSTSG	80
NTDB id 475 HSISS4 RS07005 WP 021143809.1	MKKSKWLAAGVALLSVGALAACSSKSNSTSG	76
NTDB id 324 STU RS16140 WP 011226306.1	MKKSKWLAAGVALLGVGVLAACSSKSNSTSG	76
NTDB id 292 STER RS06940 WP 011681419.1	MKKSKWLAAGVALLGVGVLAACSSKSNSTSG	76
consensus	!!!!!!!!!!!!!!	

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NTDB id 342446 ETT45 RS01360 WP 002992224.1	PSVAEDWSVSKDGLTYTYKIRKGYKMYTSEGEYYAVNTAKDFVTGLKHAADSKAGGLYLVDQSLAGLSDYLSGTNKDFS	159
NTDB id 475 HSISS4 RS07005 WP 021143809.1	PSVAEDWSVSKDGLTYTYKIRKGYKMYTSEGEYYAVNTAKDFVTGLKHAADSKAGGLYLVDQSLAGLSDYLSGTNKDFS	156
NTDB id 324 STU RS16140 WP 011226306.1	PSVAEDWSVSKDGLTYTYKIRKGYKMYTSEGEYYAVNTAKDFVTGLKHAADSKAGGLYLVDQSLAGLSDYLSGTNKDFS	156
NTDB id 292 STER RS06940 WP 011681419.1	PSVAEDWSVSKDGLTYTYKIRKGYKMYTSEGEYYAVNTAKDFVTGLKHAADSKAGGLYLVDQSLAGLSDYLSGTNKDFS	156
consensus	!!!!!!!!!!!!!!	

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NTDB id 342446 ETT45 RS01360 WP 002992224.1	VGKKAIDDPYTLQYTLKQPEPYWNSKTTYSLFVFNEDFLKNGKDFGKSTDPSTILYNGPFLKSLTAKSSIELVKNENY	238
NTDB id 475 HSISS4 RS07005 WP 021143809.1	VGKKAIDDPYTLQYTLKQPEPYWNSKTTYSLFVFNEDFLKNGKDFGKSTDPSTILYNGPFLKSLTAKSSIELVKNENY	236
NTDB id 324 STU RS16140 WP 011226306.1	VGKKAIDDPYTLQYTLKQPEPYWNSKTTYSLFVFNEDFLKNGKDFGKSTDPSTILYNGPFLKSLTAKSSIELVKNENY	236
NTDB id 292 STER RS06940 WP 011681419.1	VGKKAIDDPYTLQYTLKQPEPYWNSKTTYSLFVFNEDFLKNGKDFGKSTDPSTILYNGPFLKSLTAKSSIELVKNENY	236
consensus	!!!!!!!!!!!!!!	

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NTDB id 342446 ETT45 RS01360 WP 002992224.1	WDKKNVHFDKIKESYDGSDDQALVRGFTDGAYSEFARVYPTSSNYAVEKKYKDNLYTAPGASTSAIGNLDRQSYKYT	318
NTDB id 475 HSISS4 RS07005 WP 021143809.1	WDKKNVHFDKIKESYDGSDDQALVRGFTDGAYSEFARVYPTSSNYAVEKKYKDNLYTAPGASTSAIGNLDRQSYKYT	316
NTDB id 324 STU RS16140 WP 011226306.1	WDKKNVHFDKIKESYDGSDDQALVRGFTDGAYSEFARVYPTSSNYAVEKKYKDNLYTAPGASTSAIGNLDRQSYKYT	316
NTDB id 292 STER RS06940 WP 011681419.1	WDKKNVHFDKIKESYDGSDDQALVRGFTDGAYSEFARVYPTSSNYAVEKKYKDNLYTAPGASTSAIGNLDRQSYKYT	316
consensus	!!!!!!!!!!!!!!	

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NTDB id 342446 ETT45 RS01360 WP 002992224.1	AKKTDSEKSTKKALLNKDFRQAINFAIDRKAQSQNGKDGLATLAYRNLFVPSDFVSAGDKTFGDLVTEKMSSYGDEWS	397
NTDB id 475 HSISS4 RS07005 WP 021143809.1	AKKTDSEKSTKKALLNKDFRQAINFAIDRKAQSQNGKDGLATLAYRNLFVPSDFVSAGDKTFGDLVTEKMSSYGDEWS	396
NTDB id 324 STU RS16140 WP 011226306.1	AKKTDSEKSTKKALLNKDFRQAINFAIDRKAQSQNGKDGLATLAYRNLFVPSDFVSAGDKTFGDLVTEKMSSYGDEWS	396
NTDB id 292 STER RS06940 WP 011681419.1	AKKTDSEKSTKKALLNKDFRQAINFAIDRKAQSQNGKDGLATLAYRNLFVPSDFVSAGDKTFGDLVTEKMSSYGDEWS	396
consensus	!!!!!!!!!!!!!!	

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NTDB id 342446 ETT45 RS01360 WP 002992224.1	GVNLFADQDGLYNAEKAKTEFAKAKALQADGVQFPIHLDPVDQSSKLNVAQAQSLKQTIKESLGSENVVIDYNQLSSD	477
NTDB id 475 HSISS4 RS07005 WP 021143809.1	GVNLFADQDGLYNAEKAKTEFAKAKALQADGVQFPIHLDPVDQSSKLNVAQAQSLKQTIKESLGSENVVIDYNQLSSD	476
NTDB id 324 STU RS16140 WP 011226306.1	GVNLFADQDGLYNAEKAKTEFAKAKALQADGVQFPIHLDPVDQSSKLNVAQAQSLKQTIKESLGSENVVIDYNQLSSD	476
NTDB id 292 STER RS06940 WP 011681419.1	GVNLFADQDGLYNAEKAKTEFAKAKALQADGVQFPIHLDPVDQSSKLNVAQAQSLKQTIKESLGSENVVIDYNQLSSD	476
consensus	!!!!!!!!!!!!!!	

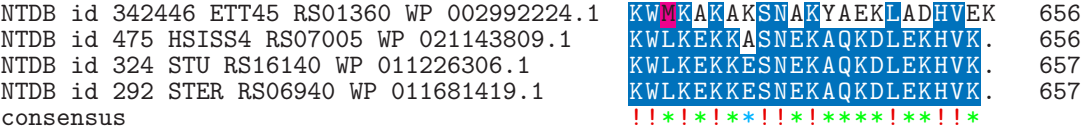
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- non conserved
- similar
- ≥ 50% conserved