

logo

NTDB id 1037792 ACB362 RS01415 WP 024623426.1	MKKSKWLAAGVALLSVGLAACSSKSNSTSG	80
NTDB id 475 HSISS4 RS07005 WP 021143809.1	MKKSKWLAAGVALLSVGLAACSSKSNSTSG	76
NTDB id 324 STU RS16140 WP 011226306.1	MKKSKWLAAGVALLSVGLAACSSKSNSTSG	76
NTDB id 292 STER RS06940 WP 011681419.1	MKKSKWLAAGVALLSVGLAACSSKSNSTSG	76
consensus	!!!!!!!!!!!!!!	

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NTDB id 1037792 ACB362 RS01415 WP 024623426.1	PSVAEDWSVSKDGLTYTYKIRKGYKMYTSEGEEYAVTAKDFVTGLKHAADSKAGGLYLVDQSIAGLSDYLSGYNKDFSN	159
NTDB id 475 HSISS4 RS07005 WP 021143809.1	PSVAEDWSVSKDGLTYTYKIRKGYKMYTSEGEEYAVTAKDFVTGLKHAADSKAGGLYLVDQSIAGLSDYLSGYNKDFSN	156
NTDB id 324 STU RS16140 WP 011226306.1	PSVAEDWSVSKDGLTYTYKIRKGYKMYTSEGEEYAVTAKDFVTGLKHAADSKAGGLYLVDQSIAGLSDYLSGYNKDFSN	156
NTDB id 292 STER RS06940 WP 011681419.1	PSVAEDWSVSKDGLTYTYKIRKGYKMYTSEGEEYAVTAKDFVTGLKHAADSKAGGLYLVDQSIAGLSDYLSGYNKDFSN	156
consensus	!!!!!!!!!!!!!!	

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NTDB id 1037792 ACB362 RS01415 WP 024623426.1	VGKKAIDDDYTLQYTLKQPEPYWNSKTTYSLFPVNEFLKNGKDFGKSTDPSTSLYNGPFLLKSLTAKSSIELVKNENY	238
NTDB id 475 HSISS4 RS07005 WP 021143809.1	VGKKAIDDDYTLQYTLKQPEPYWNSKTTYSLFPVNEFLKNGKDFGKSTDPSTSLYNGPFLLKSLTAKSSIELVKNENY	236
NTDB id 324 STU RS16140 WP 011226306.1	VGKKAIDDDYTLQYTLKQPEPYWNSKTTYSLFPVNEFLKNGKDFGKSTDPSTSLYNGPFLLKSLTAKSSIELVKNENY	236
NTDB id 292 STER RS06940 WP 011681419.1	VGKKAIDDDYTLQYTLKQPEPYWNSKTTYSLFPVNEFLKNGKDFGKSTDPSTSLYNGPFLLKSLTAKSSIELVKNENY	236
consensus	!!!!!!!!!!!!!!	

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NTDB id 1037792 ACB362 RS01415 WP 024623426.1	WDKKNVHFDIAIKSYDGSQDALVRGFTDGAYSEARVYPTSSNYSVSEKKYKDNLYYTAPGASTSAIGINLDRQSYKYT	318
NTDB id 475 HSISS4 RS07005 WP 021143809.1	WDKKNVHFDIAIKSYDGSQDALVRGFTDGAYSEARVYPTSSNYSVSEKKYKDNLYYTAPGASTSAIGINLDRQSYKYT	316
NTDB id 324 STU RS16140 WP 011226306.1	WDKKNVHFDIAIKSYDGSQDALVRGFTDGAYSEARVYPTSSNYSVSEKKYKDNLYYTAPGASTSAIGINLDRQSYKYT	316
NTDB id 292 STER RS06940 WP 011681419.1	WDKKNVHFDIAIKSYDGSQDALVRGFTDGAYSEARVYPTSSNYSVSEKKYKDNLYYTAPGASTSAIGINLDRQSYKYT	316
consensus	!!!!!!!!!!!!!!	

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NTDB id 1037792 ACB362 RS01415 WP 024623426.1	AKKTDSEKSTKKALLNKDFRQAINFAIDRKAQSQINGKDGATLAYRNLFVPSDFVSAGDKTFGDLVTEKMSSYGDEWS	397
NTDB id 475 HSISS4 RS07005 WP 021143809.1	AKKTDSEKSTKKALLNKDFRQAINFAIDRKAQSQINGKDGATLAYRNLFVPSDFVSAGDKTFGDLVTEKMSSYGDEWS	396
NTDB id 324 STU RS16140 WP 011226306.1	AKKTDSEKSTKKALLNKDFRQAINFAIDRKAQSQINGKDGATLAYRNLFVPSDFVSAGDKTFGDLVTEKMSSYGDEWS	396
NTDB id 292 STER RS06940 WP 011681419.1	AKKTDSEKSTKKALLNKDFRQAINFAIDRKAQSQINGKDGATLAYRNLFVPSDFVSAGDKTFGDLVTEKMSSYGDEWS	396
consensus	!!!!!!!!!!!!!!	

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NTDB id 1037792 ACB362 RS01415 WP 024623426.1	GVNLAADQDGLYNAEKAKTEFAKAKEALQADGVQFPIHLDPVDQSSKLNVAQAQSLKQTIIEKSLGSENVVIDYNQLSSD	477
NTDB id 475 HSISS4 RS07005 WP 021143809.1	GVNLAADQDGLYNAEKAKTEFAKAKEALQADGVQFPIHLDPVDQSSKLNVAQAQSLKQTIIEKSLGSENVVIDYNQLSSD	476
NTDB id 324 STU RS16140 WP 011226306.1	GVNLAADQDGLYNAEKAKTEFAKAKEALQADGVQFPIHLDPVDQSSKLNVAQAQSLKQTIIEKSLGSENVVIDYNQLSSD	476
NTDB id 292 STER RS06940 WP 011681419.1	GVNLAADQDGLYNAEKAKTEFAKAKEALQADGVQFPIHLDPVDQSSKLNVAQAQSLKQTIIEKSLGSENVVIDYNQLSSD	476
consensus	!!!!!!!!!!!!!!	

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NTDB id 292 STER RS06940 WP 011681419.1

consensus



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consensus



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NTDB id 1037792 ACB362 RS01415 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



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656

657

657

-  non conserved
-  similar
-  ≥ 50% conserved