

NTDB id 85	BSU 00860 NP 387967.1	..MMFGRFTERAQKVLALAQEEALRLGHNNIGTEHILLGLVREGEGIAAKALQALGLGSEKIQKEVESLIGRG.....	71
NTDB id 610	V4T04 RS10165 WP 012897346.1	MKFENIKYIPTDRILEKAEEYAHQYQYGTTESAHLLAAMATTSGSIAYSSLAGMNVDSDDLIDL.....E..DLSSH	72
NTDB id 287	SP RS11210 WP 001109712.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNNDYPYEMDRLEE.....VALELTETDYS	70
NTDB id 290	KZH43 RS10025 WP 001109677.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNNDYPYEMDRLEE.....VALELTETDYS	70
NTDB id 289	SPD RS10700 WP 001109677.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNNDYPYEMDRLEE.....VALELTETDYS	70
NTDB id 329	STU RS10020 WP 011225298.1	MTIYSRKMQAIFHRAQLEAERFESPFLCTWHVLLAMVEVPGSVAYLTFTDFEDRIHSEEIETAAVLAMEKRPKDLS	76
NTDB id 297	STER RS00545 WP 011680614.1	MTIYSRKMQAIFHRAQLEAERFESPFLCTWHVLLAMVEVPGSVAYLTFTDFEDRIHSEEIETAAVLAMEKRPKDLS	76
NTDB id 1136385	DQL06 RS08950 WP 002991294.1	...MTIMYSTKMQDIFRQAQFAARFDSHCLCTWHVLLAMVAVDNLANMI LSEYDAQVAIEEYEAAA ILAMGKTpkEQl	76
NTDB id 377	SMU RS09275 WP 002262344.1	MTDYSLKMGEVFRLAQFEAARFESSYLESWHVLLAMVEIDS SVAGLSFAEFEADVRFDYQAAA ILAIGKKPKSSS	76
consensus		* * * * *	

NTDB id 85	BSU 00860 NP 387967.1	.QEMSQTHYTPRAKKVIELSMDEARKLGHSYVGTEHLLGLIREGEVGAARVLNNLGVSLNKAQQVQLQLGNET...	147
NTDB id 610	V4T04 RS10165 WP 012897346.1	VKVKRSTLRFSPPRAEEVMTAASF LAIHNNSEAVGTEHLLYAILQVEDGFGCLQLIKL.....QKINIVSLRKELEX	142
NTDB id 287	SP RS11210 WP 001109712.1	QDETFTFTELPFSRRLQVLFDEAEYVASVHAKVLGTEHVLVAILHDSNALATRILERAGFSYEDKKDQ.VKIAALRRNLEE	149
NTDB id 290	KZH43 RS10025 WP 001109677.1	QDETFTFTELPFSRRLQILFDEAEYVASVHAKVLGTEHVLVAILHDGNALATRILERAGFSYEDKKDQ.VKIAALRRNLEE	149
NTDB id 289	SPD RS10700 WP 001109677.1	QDETFTFTELPFSRRLQILFDEAEYVASVHAKVLGTEHVLVAILHDGNALATRILERAGFSYEDKKDQ.VKIAALRRNLEE	149
NTDB id 329	STU RS10020 WP 011225298.1	ESDIIDLRAQSPALEAMLQEAQGFIASVTGAVEVGVSEHVLMAFLLHKDLMVCRLLLEVAGFQYKDDSDK.PRIIDLRRSLER	155
NTDB id 297	STER RS00545 WP 011680614.1	ESDIIDLRAQSPALEAMLQEAQEIFIASVTGAVEVGVSEHVLMAFLLHKDLMVCRLLLEVAGFQYKDDSDK.PRIIDLRRSLER	155
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NTDB id 377	SMU RS09275 WP 002262344.1	..NDIMLLEQSHALKRTLAEAAAIISQVTHAKEVGVTEHVLFAILLNPNLLATRIELVGFHAKDDGES.IRLDLLRKVIER	153
consensus		..NDIMLLEQSHALKRTLAEAAAIISQVTHAKEVGVTEHVLFAILLNPNLLATRIELVGFHAKDDGES.IRLDLLRKVIER	

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consensus



logo

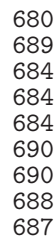
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consensus



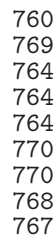
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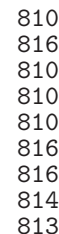
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consensus
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- X non conserved
- X similar
- X >50% conserved