

logo

NTDB id 1247 GC085 RS11680 WP 015961431.1	MKQQLGSCCQYCAYPELSDDTYLVCQGQVCR	28
NTDB id 1270 LPP RS11540 WP 015961431.1	MLQKFLSITQNLRLPSICTLGNQFHKS.....QLAVCSNCEMFMKQLGSCCQYCAYPELSDDTYLVCQGQVCR	66
NTDB id 616 LCA RS09740 WP 148664867.1		0
NTDB id 109 BSU 35450 NP 391425.2	MSNEAGGGIHKICLLCDSQFSQDVTWRAFLFLKPD.EKVCYSRCSKLKKITGHCPLCGRPQSV..HAVCRDCEV	73
NTDB id 594 KW2 RS12255 WP 244878749.1		0
NTDB id 1089286 ACLSIV RS02325 WP 163604800.1		0
NTDB id 278 KZH43 RS10085 WP 000649974.1	MKCLLCGGQTMKTVLTFSLLLRNDSDCLCSDCDSTFERIGEENCPNCKMTE.L..STKCQDCQL	62
NTDB id 237 SPD RS10760 WP 000649974.1	MKCLLCGGQTMKTVLTFSLLLRNDSDCLCSDCDSTFERIGEENCPNCKMTE.L..STKCQDCQL	62
NTDB id 203 SPR RS10245 WP 000649974.1	MKCLLCGGQTMKTVLTFSLLLRNDSDCLCSDCDSTFERIGEENCPNCKMTE.L..STKCQDCQL	62
NTDB id 168 SP RS11270 WP 000649971.1	MKCLLCGGQTMKTVLTFSLLLRNDSDCLCSDCDSTFERIGEENCPNCKMTE.L..STKCQDCQL	62
NTDB id 508 SM12261 RS09235 WP 000649953.1	MKCLLCGGQTMKAVLTFSLLLRNDASCLCLDCDSTFERIGEENCPNCKMTD.L..SMKCQDCQL	62
NTDB id 536 SMSK321 RS0102865 WP 000649958.1	MKCLLCGGQTMKAVLTFSLLLRNDVSCCLCLDCDSTFERIGEENCPNCKMTA.L..STQCQDCQF	62
consensus	***** * **** * * * * * * * * * *	

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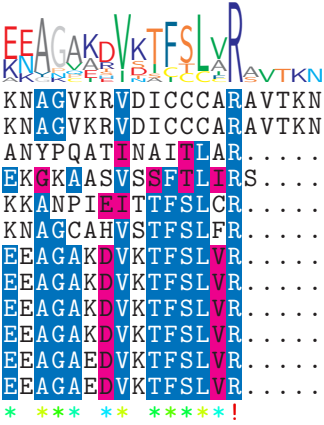
[illegible]

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NTDB	id	1247	GC085	RS11680	WP	O15961431.1	AVLTRLRLARQLNIPYDLYYCQKIINTASQANLDGEQRKRNLRHAFYVP...PVTYEHVMIIVDDLLTTGSTANEIAHT.L	176
NTDB	id	1270	LPP	RS11540	WP	O15961431.1	AVLTRLRLARQLNIPYDLYYCQKIINTASQANLDGEQRKRNLRHAFYVP...PVTYEHVMIIVDDLLTTGSTANEIAHT.L	214
NTDB	id	616	LCA	RS09740	WP	I48664867.1	QGIFETCF....PLTSLKKRPTEKEGQAQKNRAERL.ASPQFELIPQELPKPQSIIILDDIYYTGRTVWHAQQCLR	127
NTDB	id	109	BSU	35450	NP	391425.2	HLLAECLDR....PSHHPLIRLNNEKQSKKKTERLLSECFIDTKNN.SAEGMNIILLIDDLYTTGATHFAARCCLL	224
NTDB	id	594	KW2	RS12255	WP	244878749.1	TAFIKD..R....TYLDLLEKEDTVRQSSLNREGRLSTINPFKLKEG.EKIPQKIILLIDDYYTTGTTFYHAIKT.L	119
NTDB	id	1089286	ACLSIV	RS02325	WP	I63604800.1	TGFLSSAKL....PYLELTKEEETHQHSHKNRAERLASENPFHLKKN.VTIPEHVVLVDLDFITGTTLKNGASIL.L	94
NTDB	id	278	KZH43	RS10085	WP	O00649974.1	EGLVEAAGF....EYLDLLEKREERASSSKNRSERLGTELPPFIKSG.VTIPKKILLIDDYYTTGATINRVKKLL.L	205
NTDB	id	237	SPD	RS10760	WP	O00649974.1	EGLVEAAGF....EYLDLLEKREERASSSKNRSERLGTELPPFIKSG.VTIPKKILLIDDYYTTGATINRVKKLL.L	205
NTDB	id	203	SPR	RS10245	WP	O00649974.1	EGLVEAAGF....EYLDLLEKREERASSSKNRSERLGTELPPFIKSG.VTIPKKILLIDDYYTTGATINRVKKLL.L	205
NTDB	id	168	SP	RS11270	WP	O00649971.1	EGLVEAAGF....EYLDLLEKREERASSSKNRSERLGTELPPFIKSG.VTIPKKILLIDDYYTTGATINRVKKLL.L	205
NTDB	id	508	SM12261	RS09235	WP	O00649953.1	EGLVEAAGF....AYLDLLEKREERASSKSRSERLETFFPFIKSG.VTIPKKILLIDDYYTTGATINRVKKLL.L	205
NTDB	id	536	SMSK321	RS0102865	WP	O00649958.1	EGLVEAAGF....SFQELLEKREERASSSKNRSERLGTELPPFIKRG.VTIPKKILLIDDYYTTGSTINRVKKLL.L	205
consensus							***** ** * ***** * **** * **** * ! * **** * **** * **** * !! * * !! * ! * * * *	

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NTDB id 1247 GC085 RS11680 WP 015961431.1	KNAGVKKRV	DICCCARAVTKN	196
NTDB id 1270 LPP RS11540 WP 015961431.1	KNAGVKKRV	DICCCARAVTKN	234
NTDB id 616 LCA RS09740 WP 148664867.1	ANYPQATINAITLAR.....		142
NTDB id 109 BSU 35450 NP 391425.2	EKCKAASVSSFILIRS.....		240
NTDB id 594 KW2 RS12255 WP 244878749.1	KKANPIEITTFSLCR.....		134
NTDB id 1089286 ACLSIV RS02325 WP 163604800.1	KNAGCAHVSTFSLFR.....		109
NTDB id 278 KZH43 RS10085 WP 000649974.1	EEAGAKDVKTFSLV.....		220
NTDB id 237 SPD RS10760 WP 000649974.1	EEAGAKDVKTFSLV.....		220
NTDB id 203 SPR RS10245 WP 000649974.1	EEAGAKDVKTFSLV.....		220
NTDB id 168 SP RS11270 WP 000649971.1	EEAGAKDVKTFSLV.....		220
NTDB id 508 SM12261 RS09235 WP 000649953.1	EEAGAEDVKTFSLV.....		220
NTDB id 536 SMSK321 RS0102865 WP 000649958.1	EEAGAEDVKTFSLV.....		220
consensus	* ** * * * !		



X non conserved
X similar
X ≥50% conserved