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NTDB id 54 SMU RS08705 WP 002262113.1	MNEALMFLSTMLLTYYTYLYLFLSKYIKVILSKELWYLLSIFLQAYSNFMNPKYIMVWVWLMFKBAEYIFXFTASYSH	74
NTDB id 541 H702 RS10470 WP 236726575.1	MNEALMILSNGLLTLYLTVLFLFLFSKVSNNVTLSSKKELTLFS.....ISNFLIMIAVTMVNVNLFYPAEP.LYFIALSIY	0
NTDB id 543 H702 RS02375 WP 160172984.1		0
NTDB id 1127602 FGL06 RS09355 WP 231997655.1		0
NTDB id 544 H702 RS02380 WP 039696235.1	MIEIVVFIITMLLNFVGVFVIVGILFQQITKVKLTFLEWVLIIFLQAVSFFMNPQKYSMVALVLMKDISVIFCLLSVSFH	80
consensus		

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NTDB id 54 SMU RS08705 WP 002262113.1	LNPNSTSYNIFYGLPYNSYMLREXYLYLHPLFSIQLOSTYMSSTSYTYCMIMTAYLPPIMYYLYSWSLNUDLSN	146
NTDB id 541 H702 RS10470 WP 236726575.1	LNRQNSLSLNIFYGLLPVASSDLFRRRAIFFILDG....TQGIVMGSSIITTY....MIEFAGIALSYFLSVFNVDIGR	0
NTDB id 543 H702 RS02375 WP 160172984.1	MYVIVSRWLNDISN	15
NTDB id 1127602 FGL06 RS09355 WP 231997655.1		0
NTDB id 544 H702 RS02380 WP 039696235.1	HNPKIGIRYHIFYALYPINLYMILHEVYCYIIFAIFSIQLTSLTMTSPSVGKMIGMITAVIIPPIFYVLSKWLNDISN	160
consensus		

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NTDB id 54 SMU RS08705 WP 002262113.1	VSSSLKKLYEKKFLRYMNUMLLYYLMEXLYYKDFENYMNRNLYRAVIFYFILLFELFYLNFRHGGQEQKNFLRN	225
NTDB id 541 H702 RS10470 WP 236726575.1	LKDSITKMKVKRLIPNITMMLLYLLIQVLYVLESNVIP.TLKFRKFVVIYVILLFILLFISLSQYTKQKVQNEIMAQ	16
NTDB id 543 H702 RS02375 WP 160172984.1	MRSYQNRLEKEIASA	94
NTDB id 1127602 FGL06 RS09355 WP 231997655.1	VSSSLKKHIEPKIFNITNLMVSYFLFMDAIAIVKPFHNAD.NDAIRGIVAFIYILLFVFLFYLNFRHGGQEQKNELLRN	39
NTDB id 544 H702 RS02380 WP 039696235.1	VRKTLKNYLSAHALRFVNVLLILFYPGMECFGALAPFEKMSMNRNIYRAVIFYFILLFIFLFYLNFRHGGQEQKNELLRN	240
consensus		

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NTDB id 54 SMU RS08705 WP 002262113.1	QEMELRALENYSKHVESLYQEVRSFRHDYANILMSLKVGLDQCDLDDKKIYDEITADSTKLKVNKFDLTRLANITDSA	305
NTDB id 541 H702 RS10470 WP 236726575.1	KEAQIRNITQYSQQIESLYKDIRSFRHDYLNILTSIRIGIENKDIASIEKIYHQILEKTGHQLQDTRYNIGHLANIQNDA	96
NTDB id 543 H702 RS02375 WP 160172984.1	REQELRYLSAYSKKVEGLYEELRAFRHDYANILVSLKEGIDQDDMGLVRKIYDSVLEDSADFIIRNSKFNIGRLVNIDDDA	174
NTDB id 1127602 FGL06 RS09355 WP 231997655.1	QEMELALANYSKHVESLYQEVRSFRHDYANILMSLKVGLDQCDLDDIKKIYDEITADSTKLKVNKFDLTRLANITDSC	119
NTDB id 544 H702 RS02380 WP 039696235.1	QEMELALANYSKHVESLYQEVRSFRHDYANILMSLKVGLDQCDLDDIKKIYDEITADSTKLKVNKFDLTRLANITDSC	320
consensus		

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NTDB id 54 SMU RS08705 WP 002262113.1	VKSLMSAKFLEAENQGLYSLEVSEKMGTPPLISYIRILSILFDNAIEAALFESNDLVANFYQDGDVFELLEISTK	385
NTDB id 541 H702 RS10470 WP 236726575.1	VKSLMSAKFLEAENQGLYSLEVSEKMGTPPLISYIRILSILFDNAIEAALFESNDLVANFYQDGDVFELLEISTK	176
NTDB id 543 H702 RS02375 WP 160172984.1	VKSLMSAKFLHAENQGIENSLEVSEKMGTPSIPLISYIRILSILFDNAIEAALFESNDLVANFYQDGDVFELLEISTK	254
NTDB id 1127602 FGL06 RS09355 WP 231997655.1	VKSLMSAKFLEAENRGIEISLEVSEKMGTPSIPLISYIRILSILFDNAIEAALFESNDLVANFYQDGDVFELLEISTK	199
NTDB id 544 H702 RS02380 WP 039696235.1	VKSLMSAKFLQAENQGITISLEVSEKMGTPSIPLISYIRILSILFDNAIEAALFESNDLVANFYQDGDVFELLEISTK	400
consensus		

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X non conserved
X similar
X ≥ 50% conserved