

logo

NTDB id 1272 LPP RS04360 WP 010946546.1	MNWLNQHKKAII LAASA AVFTAIMIFLATGKNKEPVKQAVPTETENTVVKQEANNDNESNETIVIDIKGAVQHPGVYEMRT	0
NTDB id 1249 GC085 RS04230 WP 010946546.1		0
NTDB id 1159 ABDM36 RS04765 WP 001166105.1		0
NTDB id 1150 A1552VC RS08580 WP 001166105.1		0
NTDB id 1294 VP RS04495 WP 021449594.1		0
NTDB id 105 BSU 25590 NP 390437.1	MNWLNQHKKAII LAASA AVFTAIMIFLATGKNKEPVKQAVPTETENTVVKQEANNDNESNETIVIDIKGAVQHPGVYEMRT	80
NTDB id 1127 NGFG RS10560 WP 003703428.1		0
NTDB id 1126 NGFG RS09375 WP 003703428.1		0
NTDB id 1125 NGFG RS07215 WP 003703428.1		0
NTDB id 1108 NGFG RS06525 WP 003703428.1		0
NTDB id 152138 CUREO RS07735 WP 018713931.1		0
NTDB id 1242 Cj0011c YP 002343483.1		0
consensus		

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NTDB id 1272 LPP RS04360 WP 010946546.1	MSRGGATPY...RFLLRHIVARCGLLFATLKGKTMKKMFVLF...TPAKQVLVKGKINLNTADVSLTGSFKG	55
NTDB id 1249 GC085 RS04230 WP 010946546.1	MKARLIIAILLSFIVIPFSSQAENEPQK.....TPAKQVLVKGKINLNTADVSLTGSFKG	55
NTDB id 1159 ABDM36 RS04765 WP 001166105.1	MQIKTKIVTLFSLCLPTLPLLANAEETA.....PAAQVEEGIVITVNINTASAEELATLLKG	58
NTDB id 1150 A1552VC RS08580 WP 001166105.1	MQIKTKIVTLFSLCLPTLPLLANAEETA.....PAAQVEEGIVITVNINTASAEELATLLKG	58
NTDB id 1294 VP RS04495 WP 021449594.1	MKWILTLCVVFPAPLSLAAD.....TKADKYEGIEITVNINTASAEETATMLKG	49
NTDB id 105 BSU 25590 NP 390437.1	GDRVSAIEKAGGTSEQADEAQVNLAELIQDGTVVYIPKKGEETAVQQGGGGSVQSDGGKGALVNINTATLEELQG. ISG	159
NTDB id 1127 NGFG RS10560 WP 003703428.1	MSRGGATPY...RFLLRHIVARCGLLFATLKGKTMKKMFVLF...TPAKQVLVKGKINLNTADVSLTGSFKG	68
NTDB id 1126 NGFG RS09375 WP 003703428.1	MSRGGATPY...RFLLRHIVARCGLLFATLKGKTMKKMFVLF...TPAKQVLVKGKINLNTADVSLTGSFKG	68
NTDB id 1125 NGFG RS07215 WP 003703428.1	MSRGGATPY...RFLLRHIVARCGLLFATLKGKTMKKMFVLF...TPAKQVLVKGKINLNTADVSLTGSFKG	68
NTDB id 1108 NGFG RS06525 WP 003703428.1	MSRGGATPY...RFLLRHIVARCGLLFATLKGKTMKKMFVLF...TPAKQVLVKGKINLNTADVSLTGSFKG	68
NTDB id 152138 CUREO RS07735 WP 018713931.1	MKRLLLLFV..FSFSLIFGAVNINTASKEELMT.LKG	34
NTDB id 1242 Cj0011c YP 002343483.1	MKKLLFLFF..ALTAFLFGAVNINTATLKEELKS.LNG	34
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NTDB id 1272 LPP RS04360 WP 010946546.1	IGKKRAEAI IAYRENHQGFKSLEELA EVKGIGQRFVDKNREKLKEIFVIN.....	105
NTDB id 1249 GC085 RS04230 WP 010946546.1	IGKKRAEAI IAYRENHQGFKSLEELA EVKGIGQRFVDKNREKLKEIFVIN.....	105
NTDB id 1159 ABDM36 RS04765 WP 001166105.1	IGLKKQAIVDYREANGPFTHIDDLTNVKGIGEAIVRNNAARILL.....	103
NTDB id 1150 A1552VC RS08580 WP 001166105.1	IGLKKQAIVDYREANGPFTHIDDLTNVKGIGEAIVRNNAARILL.....	103
NTDB id 1294 VP RS04495 WP 021449594.1	IGEKKAQSI VDYRTEHGPFKTAADLTNVKGIGEAITKKNEDRILL.....	94
NTDB id 105 BSU 25590 NP 390437.1	VGPSKAEAI IAYREENGRFQTIEDITKVSIGEKSFEEKSSITVK.....	205
NTDB id 1127 NGFG RS10560 WP 003703428.1	IGPAKAKAIAEYRAQNGAFKSVDDLKVKGIGPAVLAKLKDQASVGAPAPKGPAPKPVLPVAVKK	131
NTDB id 1126 NGFG RS09375 WP 003703428.1	IGPAKAKAIAEYRAQNGAFKSVDDLKVKGIGPAVLAKLKDQASVGAPAPKGPAPKPVLPVAVKK	131
NTDB id 1125 NGFG RS07215 WP 003703428.1	IGPAKAKAIAEYRAQNGAFKSVDDLKVKGIGPAVLAKLKDQASVGAPAPKGPAPKPVLPVAVKK	131
NTDB id 1108 NGFG RS06525 WP 003703428.1	IGPAKAKAIAEYRAQNGAFKSVDDLKVKGIGPAVLAKLKDQASVGAPAPKGPAPKPVLPVAVKK	131
NTDB id 152138 CUREO RS07735 WP 018713931.1	IGEATAEAI IEYRKEN. KFTKIEDIKNVKGIGDKKFESIKEDI EVKDNKK.....	83
NTDB id 1242 Cj0011c YP 002343483.1	IGEA KAKAILEYRKEA. NFTSIDDLLKKVKGIGDKLFEKIKNDITIE.....	79
consensus	*! *! *!***! ** !***** !!!!! * * ***** * *	

☒ non conserved
☒ similar
☒ ≥ 50% conserved