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NTDB id 324126 ECB94 RS13650 WP 124940765.1	MSRGGATPYRFLLRHIVARCGLLFATLKGGKMKKLELFCM	23
NTDB id 1159 ABDM36 RS04765 WP 001166105.1	MKYL.KNVVLIV..LLVAPVSFSY..A.....	29
NTDB id 1150 A1552VC RS08580 WP 001166105.1	MQIKTK.IVTLFLSLCLPTLPLLANAEETA.....	29
NTDB id 1294 VP RS04495 WP 021449594.1	MQIKTK.IVTLFLSLCLPTLPLLANAEETA.....	20
NTDB id 1395 DSB67 RS04445 WP 010649155.1	MKWILTLCVLVVFAPLSLAA..D.....	21
NTDB id 1041 HON27 RS14740 WP 005124713.1	MKWMLTLCVLILAPMSWAE.TK.....	58
NTDB id 1014 ACIAD RS13855 WP 004924483.1	MQYSMRFWKN.....KYFLLIMLCGLVISPSVCAQSFDQNFQEWKAKQMYDQKLNITKSSHS	54
NTDB id 1127 NGFG RS10560 WP 003703428.1	MSIRHFFT...FFWAT.....LLTLFSM....QIVFAQGFDQYDAWKTTQKKAHTRLNSSASHHS	44
NTDB id 1126 NGFG RS09375 WP 003703428.1	MSRGGATPYRFLLRHIVARCGLLFATLKGGTMMKMFVLFM....LF.....	44
NTDB id 1125 NGFG RS07215 WP 003703428.1	MSRGGATPYRFLLRHIVARCGLLFATLKGGTMMKMFVLFM....LF.....	44
NTDB id 1108 NGFG RS06525 WP 003703428.1	MSRGGATPYRFLLRHIVARCGLLFATLKGGTMMKMFVLFM....LF.....	44
NTDB id 1242 Cj0011c YP 002343483.1	MKKLLFLF.....FA.....	10
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NTDB id 324126 ECB94 RS13650 WP 124940765.1	SEAKKLEGIEITVNINQAGPEELAEELKGVGEEKAKDIVEYRDQHGSFKSADDLTQVKGIGEATVEKNRSRITL.	97
NTDB id 1159 ABDM36 RS04765 WP 001166105.1	PAAQVEEGIVITVNINTASAEELATLLKGIGLKKAAQIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL.	103
NTDB id 1150 A1552VC RS08580 WP 001166105.1	PAAQVEEGIVITVNINTASAEELATLLKGIGLKKAAQIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL.	103
NTDB id 1294 VP RS04495 WP 021449594.1	TKADKYEGIEITVNINTASAEETIATMLKGIGEKKAAQIVDYRTEHGPFKTAADLTNVKGIGEATIKKNEDRILL.	94
NTDB id 1395 DSB67 RS04445 WP 010649155.1	TKADKYEGIEITVNINSATAQEIATLLIGIGEEKKAQDIVEYRNEHGPFKTAADLTQVKGIGEATVRKNEDRILL.	95
NTDB id 1041 HON27 RS14740 WP 005124713.1	NGSKISQTKNFNDSTGQIHLNQANVNEFQQ.LKGVGEEKKAQIVVEYRQKNGSFKNIDEIKNVKGIGPAIFEKKNKSRLLAL.	136
NTDB id 1014 ACIAD RS13855 WP 004924483.1	LSKP....TQSGTGKVNINSATAKELQQLHGVGEEKKALAIVEYRQQYQKGFASVDDLQKVKGIGPKFIEKNRDLRLAI.	128
NTDB id 1127 NGFG RS10560 WP 003703428.1	SCAFSLAAVNINAASQQELEA.LPGIGPAKAKAIAEYRAQNGAFKSVDLLIKVKGIGPAVLAKLKDQASVG	114
NTDB id 1126 NGFG RS09375 WP 003703428.1	SCAFSLAAVNINAASQQELEA.LPGIGPAKAKAIAEYRAQNGAFKSVDLLIKVKGIGPAVLAKLKDQASVG	114
NTDB id 1125 NGFG RS07215 WP 003703428.1	SCAFSLAAVNINAASQQELEA.LPGIGPAKAKAIAEYRAQNGAFKSVDLLIKVKGIGPAVLAKLKDQASVG	114
NTDB id 1108 NGFG RS06525 WP 003703428.1	SCAFSLAAVNINAASQQELEA.LPGIGPAKAKAIAEYRAQNGAFKSVDLLIKVKGIGPAVLAKLKDQASVG	114
NTDB id 1242 Cj0011c YP 002343483.1	LTAFLFGAVNINTATLKELES.LNGIGEAKAKAILEYRKE.ANFTSIDDLKVKGIGDKLFEKIKNDITIE	79
consensus	*****	

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NTDB id 324126 ECB94 RS13650 WP 124940765.1	APAPKGPAKPVLPVKK	97
NTDB id 1159 ABDM36 RS04765 WP 001166105.1		103
NTDB id 1150 A1552VC RS08580 WP 001166105.1		103
NTDB id 1294 VP RS04495 WP 021449594.1		94
NTDB id 1395 DSB67 RS04445 WP 010649155.1		95
NTDB id 1041 HON27 RS14740 WP 005124713.1		136
NTDB id 1014 ACIAD RS13855 WP 004924483.1		128
NTDB id 1127 NGFG RS10560 WP 003703428.1	APAPKGPAKPVLPVKK	131
NTDB id 1126 NGFG RS09375 WP 003703428.1	APAPKGPAKPVLPVKK	131
NTDB id 1125 NGFG RS07215 WP 003703428.1	APAPKGPAKPVLPVKK	131
NTDB id 1108 NGFG RS06525 WP 003703428.1	APAPKGPAKPVLPVKK	131
NTDB id 1242 Cj0011c YP 002343483.1		79
consensus		

- non conserved
- similar
- ≥ 50% conserved