

logo

NTDB id 1127 NGFG RS10560 WP 003703428.1	MSRGGATPYRFLIRHIVARCGLLFATLKG.KTMKKMFVLF.....CMLFSCAFSLAAVN	54
NTDB id 1126 NGFG RS09375 WP 003703428.1	MSRGGATPYRFLIRHIVARCGLLFATLKG.KTMKKMFVLF.....CMLFSCAFSLAAVN	54
NTDB id 1125 NGFG RS07215 WP 003703428.1	MSRGGATPYRFLIRHIVARCGLLFATLKG.KTMKKMFVLF.....CMLFSCAFSLAAVN	54
NTDB id 1108 NGFG RS06525 WP 003703428.1	MSRGGATPYRFLIRHIVARCGLLFATLKG.KTMKKMFVLF.....CMLFSCAFSLAAVN	54
NTDB id 1014 ACIAD RS13855 WP 004924483.1	MS..IRHFFTFFWA.TLLT LFS...MQIVFAQGFDQQYDAWKTTQKQAHDTLNS....SASHHSLSKPTQQSGTGKVN	69
NTDB id 1440 BW25113 RS02290 WP 000680312.1	...MKHGIKALLI.TLSLACAGMSH.SALAA.....ASVAKPTAVETKAEAPAAQSKAAVPAKASDEEGTRVS	63
NTDB id 1159 ABDM36 RS04765 WP 001166105.1	MQ..IKTKIVT...LFLSLCLPTLPLLANAEETAP.....A.....AQVEEGIVITVN	43
NTDB id 1150 A1552VC RS08580 WP 001166105.1	MQ..IKTKIVT...LFLSLCLPTLPLLANAEETAP.....A.....AQVEEGIVITVN	43
NTDB id 81489 FORC8 RS10580 WP 021449594.1	MK.....WILTLCLVVFAPLSLAA..DT.....K.....ADKYEGIEITVN	34
NTDB id 1294 VP RS04495 WP 021449594.1	MK.....WILTLCLVVFAPLSLAA..DT.....K.....ADKYEGIEITVN	34
NTDB id 1395 DSB67 RS04445 WP 010649155.1	MK.....WMLTLCLLILAPMSWAE.TKT.....K.....ADKYEGIEITVN	35
NTDB id 1362 HI 1008 AAC22669.1	...MKTL...F...TSVVLCGALVVSSSF AEEKAT.....XQTAQSV.VTT.....Q....AEAQVAPAVVSDKLN	52
NTDB id 1459 ASU2 RS10590 WP 015674357.1	...MKTPSKL...GLSL LIG.MLSTGVLAAEKT.....PSVTEPTPVVQNAQATSP....EHSHAEMTNNANLVN	60
NTDB id 1382 A4U84 RS07375 WP 062924234.1	...MKKLSRY...TFSLLFG.LFSATTFAEEKVT.....TST.EHTPVVMQ.....VAATQKVNANVVN	52
consensus	* ** * *	**

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NTDB id 1127 NGFG RS10560 WP 003703428.1	INAASQQELEA.LPGIGPAKAKAIAEYRAQNGAFKSVDDLKIKVKGIGPAVLAKLKDQASVGAPAPKGPAPKPVLPVAVKK	131
NTDB id 1126 NGFG RS09375 WP 003703428.1	INAASQQELEA.LPGIGPAKAKAIAEYRAQNGAFKSVDDLKIKVKGIGPAVLAKLKDQASVGAPAPKGPAPKPVLPVAVKK	131
NTDB id 1125 NGFG RS07215 WP 003703428.1	INAASQQELEA.LPGIGPAKAKAIAEYRAQNGAFKSVDDLKIKVKGIGPAVLAKLKDQASVGAPAPKGPAPKPVLPVAVKK	131
NTDB id 1108 NGFG RS06525 WP 003703428.1	INAASQQELEA.LPGIGPAKAKAIAEYRAQNGAFKSVDDLKIKVKGIGPAVLAKLKDQASVGAPAPKGPAPKPVLPVAVKK	131
NTDB id 1014 ACIAD RS13855 WP 004924483.1	INSATAKELQQ.LHGVGEKKALAIVEYRQQYQKFA SVDDLQKVKGIGPKFIEKNRDLRLAI.....	128
NTDB id 1440 BW25113 RS02290 WP 000680312.1	INNASAEELARAMNGVGLKKAQAIVSYREEYGFPTVEDLQKVPGMGNSLVERNLAVLTL.....	123
NTDB id 1159 ABDM36 RS04765 WP 001166105.1	INTASAEELATLLKGIGLKKQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL.....	103
NTDB id 1150 A1552VC RS08580 WP 001166105.1	INTASAEELATLLKGIGLKKQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL.....	103
NTDB id 81489 FORC8 RS10580 WP 021449594.1	INTASAEELIATMLKGIGEKKQAQSIQDYRTEHGPFKTAADLTNVKGIGEATIKKNEDRILL.....	94
NTDB id 1294 VP RS04495 WP 021449594.1	INTASAEELIATMLKGIGEKKQAQSIQDYRTEHGPFKTAADLTNVKGIGEATIKKNEDRILL.....	94
NTDB id 1395 DSB67 RS04445 WP 010649155.1	VNSATAQEIIATLLIGIGEKKQAQDIVEYRNEHGPFKTAADLTNVKGIGEATVRKNEDRILL.....	95
NTDB id 1362 HI 1008 AAC22669.1	INTATASEIIQKSLTGIGAKKAQAIVQYREKHGNFXNAEQ LLEVQGIGKATLEKNRDRILIF.....	112
NTDB id 1459 ASU2 RS10590 WP 015674357.1	INIATAAEI IQDKLVGIGAKKAQAIVEYRTKNGAFSSLEQLTDVSGIGKATLDKNRDRILLQ.....	121
NTDB id 1382 A4U84 RS07375 WP 062924234.1	INTATASEI IQDKLVGIGAKKAQAIVEYREKNGKFLNVEQLTEVSGIGKATLEKNRDCIVIE.....	113
consensus	*!***! * !*! *!***!***! *! !*****! !*!! *** *****	

- X non conserved
- X similar
- X ≥ 50% conserved