

NTDB	id	1127	NGFG	RS10560	WP	003703428.1	MSRGGATPYRFLLRHIVARCGLLFATLKG.KTMKKMFVLF.....CMLFSCAFSLAAVN	54
NTDB	id	1126	NGFG	RS09375	WP	003703428.1	MSRGGATPYRFLLRHIVARCGLLFATLKG.KTMKKMFVLF.....CMLFSCAFSLAAVN	54
NTDB	id	1125	NGFG	RS07215	WP	003703428.1	MSRGGATPYRFLLRHIVARCGLLFATLKG.KTMKKMFVLF.....CMLFSCAFSLAAVN	54
NTDB	id	1108	NGFG	RS06525	WP	003703428.1	MSRGGATPYRFLLRHIVARCGLLFATLKG.KTMKKMFVLF.....CMLFSCAFSLAAVN	54
NTDB	id	1014	ACIAD	RS13855	WP	004924483.1	MS..IRHFFTFFWA.TLLTFS..MQIVFAQQGFDQQYDAWKTTQQAHDTRLNS....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	38607	HK86	RS06430	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....TSVVLCCALVVSSSF.AEEKAT.....XQTAQSV.VTT.....Q....AEAQVAPAVVSDKLN	52	
NTDB	id	1459	ASU2	RS10590	WP	015674357.1	MKTPSKL....GLSLLIIG.MLSTGVLAAEKT.....PSVTEPTPVVQNAQATSP....EHSHAEMTNNANLVN	60
NTDB	id	1382	A4U84	RS07375	WP	062924234.1	MKKLSRY....TFSLLFG.LFSATTF.AEEKVT.....TST.EHTPVVMQ.....VAATQKVNANLVN	52
consensus							* ** * * *	**

- ☐ non conserved
- ☒ similar
- ☒ $\geq 50\%$ conserved