

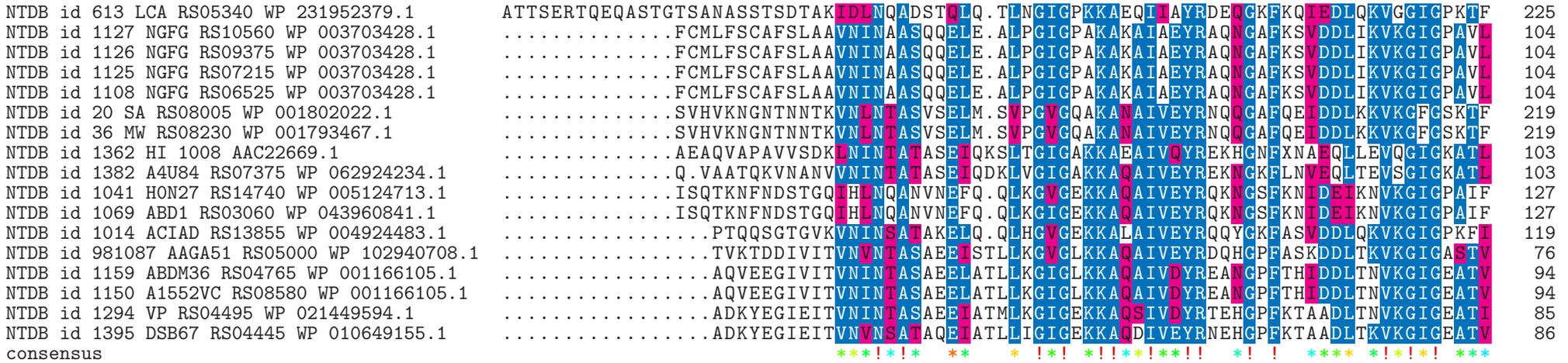
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NTDB id 613 LCA RS05340 WP 231952379.1		.MMWQDYWQR...YRRFW...WVPCLVILGIIAGGFYWHQSQQKQKVTQNEQL.LATSA.....SKTKPATKKTTHQAAT	66
NTDB id 1127 NGFG RS10560 WP 003703428.1		.....	0
NTDB id 1126 NGFG RS09375 WP 003703428.1		.....	0
NTDB id 1125 NGFG RS07215 WP 003703428.1		.....	0
NTDB id 1108 NGFG RS06525 WP 003703428.1		.....	0
NTDB id 20 SA RS08005 WP 001802022.1		MVLLYQFLLRYKDFLTQWKLYIISAVVILIMVLIGFIFWRQDDYTSRNFENKDTALKQSTSENNSLSKLEDVQVKDGDNSK	80
NTDB id 36 MW RS08230 WP 001793467.1		MVLLYQFLLRYKDFLTQWKLYIISAVVILIMVLIGFIFWRQDDYTSRNFENKDNALKQSTSENSSLSKVEDVQVKDGDNSK	80
NTDB id 1362 HI 1008 AAC22669.1		....MKT.....FTSV.....VLCGALV.....	15
NTDB id 1382 A4U84 RS07375 WP 062924234.1		....MKKLS.....RYTFS.....LLFG.LF.....	16
NTDB id 1041 HON27 RS14740 WP 005124713.1		....MQYSMRFWKNKYFLLI.....MLCLGVI.....	23
NTDB id 1069 ABD1 RS03060 WP 043960841.1		....MQYSMRFWKNKYFVLI.....MLCLGVI.....	23
NTDB id 1014 ACIAD RS13855 WP 004924483.1		....MSIRHF...FTFFWAT.....LLTLFS.....	19
NTDB id 981087 AAGA51 RS05000 WP 102940708.1		.....	0
NTDB id 1159 ABDM36 RS04765 WP 001166105.1		....MQIKTK...IV.TLFL...SLCLPT.....	18
NTDB id 1150 A1552VC RS08580 WP 001166105.1		....MQIKTK...IV.TLFL...SLCLPT.....	18
NTDB id 1294 VP RS04495 WP 021449594.1		.....M.KWIL...TLCLVV.....	11
NTDB id 1395 DSB67 RS04445 WP 010649155.1		.....M.KWML...TLCLLI.....	11
consensus		.....*	

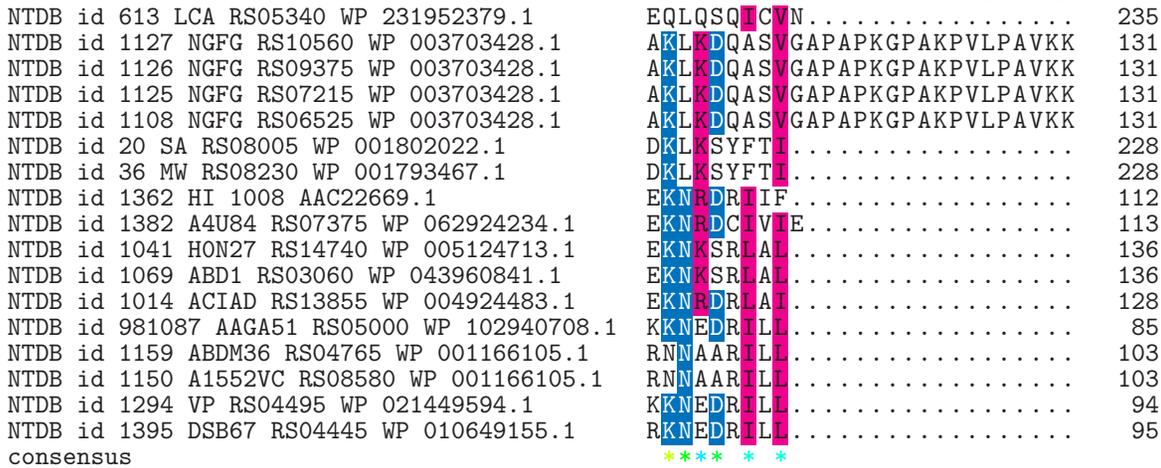
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NTDB id 613 LCA RS05340 WP 231952379.1		KKKGGFVDIKGAVVYPGIYPITVSETRLFEVLKKNAGGPTDAADTKRLNLAQKLTQQLIYIPKKGEVVDQELIQAPQQI	146
NTDB id 1127 NGFG RS10560 WP 003703428.1		.....MSRGA..TPYRFL.....LIRHIVARCGLLFATLKQKTMKKM..FVL....	39
NTDB id 1126 NGFG RS09375 WP 003703428.1		.....MSRGA..TPYRFL.....LIRHIVARCGLLFATLKQKTMKKM..FVL....	39
NTDB id 1125 NGFG RS07215 WP 003703428.1		.....MSRGA..TPYRFL.....LIRHIVARCGLLFATLKQKTMKKM..FVL....	39
NTDB id 1108 NGFG RS06525 WP 003703428.1		.....MSRGA..TPYRFL.....LIRHIVARCGLLFATLKQKTMKKM..FVL....	39
NTDB id 20 SA RS08005 WP 001802022.1		NKGPVYVDVKGAVKHPNVYKMTSKD.RVVDLLDK.AQLLEDADVSRINLSEKLTQKMFIFPHKQKKNVEPQIEVN...	154
NTDB id 36 MW RS08230 WP 001793467.1		NKGPVYVDVKGAVKHPNVYKMTSKD.RVVDLLDK.AQLLEDADVSRINLSEKLTQKMFIFPHKQKKNVEPQIEVN...	154
NTDB id 1362 HI 1008 AAC22669.1		.....VSSSFA.....E.....EKATXQT.....AQSIVTTQ.....	37
NTDB id 1382 A4U84 RS07375 WP 062924234.1		.....SATTEA.....E.....EKVTTST.....EHTPVVMQ.....	38
NTDB id 1041 HON27 RS14740 WP 005124713.1		.....SPSVCA.....QSFDQ.NFQEWKAKQQMYDQKLNIT..KSSHSNGSK.....	62
NTDB id 1069 ABD1 RS03060 WP 043960841.1		.....SPSVCA.....QSFDQ.NFQEWKVKQQMYDQKLNIS..KSSHSNSSK.....	62
NTDB id 1014 ACIAD RS13855 WP 004924483.1		.....MQIVFA.....QGFDQ.QYDAWKTTQQKAHDTRINSS..ASHHSLSK.....	57
NTDB id 981087 AAGA51 RS05000 WP 102940708.1		.....MT.....SVS..AADNSTNQ.....	13
NTDB id 1159 ABDM36 RS04765 WP 001166105.1		.....LP.....LLA..NAEETAPA.....	31
NTDB id 1150 A1552VC RS08580 WP 001166105.1		.....LP.....LLA..NAEETAPA.....	31
NTDB id 1294 VP RS04495 WP 021449594.1		.....FA.....PLS..LAA..DTK.....	22
NTDB id 1395 DSB67 RS04445 WP 010649155.1		.....LA.....PMS..WAE..TKTK.....	23
consensus		.....* * * *	

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- non conserved
- similar
- ≥ 50% conserved