

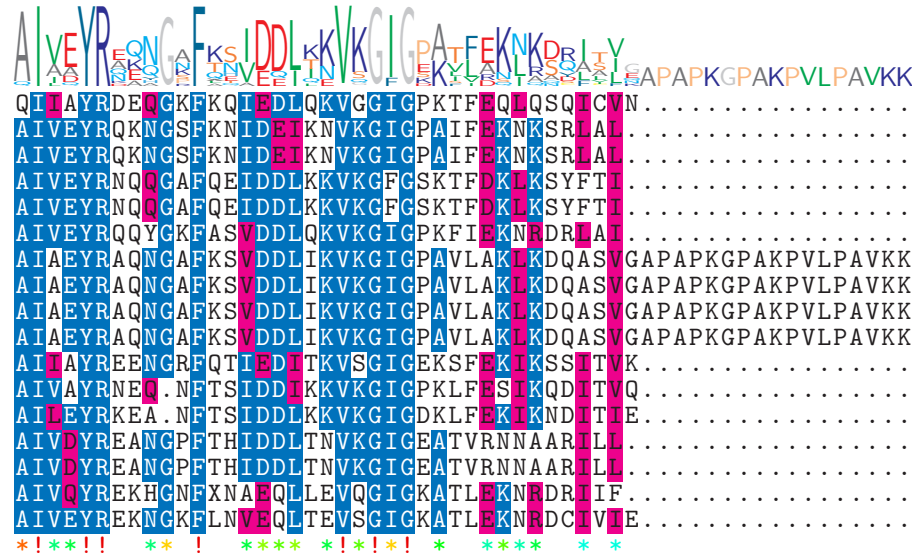
logo

NTDB id 613 LCA RS05340 WP 231952379.1	MMWQDYWQRYRRFWVPCLVLLGIIAGGFYWHQSQQKQKVTQNEQL.LAT.....SASKTKPATKKTH	62
NTDB id 1069 ABD1 RS03060 WP 043960841.1		0
NTDB id 1041 HON27 RS14740 WP 005124713.1		0
NTDB id 20 SA RS08005 WP 001802022.1	MVLLYQFLLRYKDFLTQWKLYI....ISAVVLIMVLIGFIFWRQDDYTSRNFENKDTALKQSTSENNSLSKLEDVQVKDG	76
NTDB id 36 MW RS08230 WP 001793467.1	MVLLYQFLLRYKDFLTQWKLYI....ISAVVLIMVLIGFIFWRQDDYTSRNFENKDNALKQSTSENNSSLSKVEDVQVKDG	76
NTDB id 1014 ACIAD RS13855 WP 004924483.1		0
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 105 BSU 25590 NP 390437.1	MNWLNQHKKAI....ILAASAAVFTAIMIFLATG....KNKE....PVKQAVPT....ETENTVVVKQEA	53
NTDB id 238567 CAV RS00865 WP 094324639.1		0
NTDB id 1242 Cj0011c YP 002343483.1		0
NTDB id 1159 ABDM36 RS04765 WP 001166105.1		0
NTDB id 1150 A1552VC RS08580 WP 001166105.1		0
NTDB id 1362 HI 1008 AAC22669.1		0
NTDB id 1382 A4U84 RS07375 WP 062924234.1		0
consensus		

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NTDB id 613 LCA RS05340 WP 231952379.1	QAATKKKGGFVDIKGAVVYPGIYPITVSETRLFVLLKNAGGPTDAADTKRLNLAQKLTIDQLI.....I	125
NTDB id 1069 ABD1 RS03060 WP 043960841.1	MQYSMRFWKNKYFVLIMLC LGVISPSV	27
NTDB id 1041 HON27 RS14740 WP 005124713.1	MQYSMRFWKNKYFLLIMLC LGVISPSV	27
NTDB id 20 SA RS08005 WP 001802022.1	DNSKNKGPVYVDVKGAVKHPNVYKMTSK.DRVVDLLDKAQLLDDA.DVSRINLSEKLTIDQK.....MI	137
NTDB id 36 MW RS08230 WP 001793467.1	DNSKNKGPVYVDVKGAVKHPNVYKMTSK.DRVVDLLDKAQLLEDA.DVSQINLSEKLTIDQK.....MI	137
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NTDB id 1127 NGFG RS10560 WP 003703428.1	MSRGGATPY....RFLLRHIVARCGL.....L	24
NTDB id 1126 NGFG RS09375 WP 003703428.1	MSRGGATPY....RFLLRHIVARCGL.....L	24
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NTDB id 1108 NGFG RS06525 WP 003703428.1	MSRGGATPY....RFLLRHIVARCGL.....L	24
NTDB id 105 BSU 25590 NP 390437.1	NNDESNETIVIDIKGAVQHPGVYEMRTG.DRVSQLAIEKAGGTSEQADEAQVNLAELQDGTV.....V	115
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NTDB id 1150 A1552VC RS08580 WP 001166105.1	MQIKTKIVTLFSLCLPTLP LLA	23
NTDB id 1362 HI 1008 AAC22669.1	MKTLL..FTSVVLCGALVVSSS	19
NTDB id 1382 A4U84 RS07375 WP 062924234.1	MKKLSRYTFSLLFG.LFSATT	20
consensus	*	*

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consensus

- X non conserved
- X similar
- X $\geq 50\%$ conserved