

logo

NTDB id 1436 BW25113 RS00530 WP 001025145.1	MNIPQLTALCLRYHGVLLDASEEVVHVAVVDAPSHELLDALHFATTKRIEITCWTRQQMEGHASRTQQTLPVAVQEKHQP	80
NTDB id 1177 A1552VC RS01165 WP 000422572.1		0
NTDB id 1203 PSJM300 01605 AFN76401.1		0
NTDB id 1019 ACIAD RS04205 WP 004922051.1		0
NTDB id 47407 H7R58 RS04330 WP 182930043.1		0
NTDB id 1448 BW25113 RS15320 WP 001326494.1		0
NTDB id 1307 DR RS10055 WP 010888596.1	MT	2
NTDB id 1086 NMV RS00245 WP 002218487.1		0
NTDB id 1116 NGFG RS10615 WP 003690200.1		0
NTDB id 1185 GTF74 RS11940 WP 000350195.1		0
NTDB id 1176 A1552VC RS01160 WP 000350195.1		0
NTDB id 1283 ACE17W RS02630 WP 010947729.1		0
NTDB id 1259 GC085 RS10305 WP 010947729.1		0
NTDB id 1194 PA0395 NP 249086.1		0
NTDB id 1202 PSJM300 01600 AFN76400.1		0
NTDB id 1020 ACIAD RS04210 WP 011182164.1		0
NTDB id 1081 FDQ49 RS05335 WP 006581387.1		0
NTDB id 1060 ABD1 RS04380 WP 000355489.1		0
NTDB id 1040 HON27 RS13360 WP 000355489.1		0
consensus		

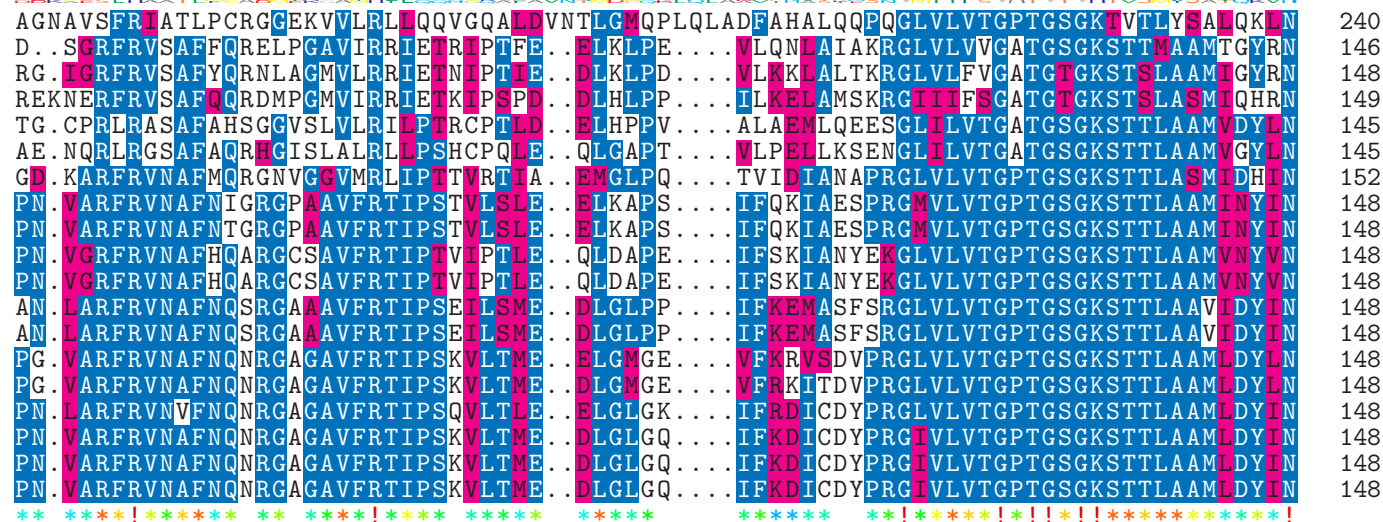
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NTDB id 1436 BW25113 RS00530 WP 001025145.1	KAEELTRTLQSALEQRASDIHIEPADNAYRIRLRIDGVLHPLPDVSPDAGVALTARLKVLGNDIAEHRLPQDGGFTVEL	160
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NTDB id 1203 PSJM300 01605 AFN76401.1	..MEFEKLLRLMVEKGGSDLFITAGV...PPSMKVNGKILPVPNKTPMSPEMTRETVEHGVNMNEQQRREFTENHECNFAISA	75
NTDB id 1019 ACIAD RS04205 WP 004922051.1	..MDFNDLLNLMAKHKASDLFITAGV...EPSLKINGEITPVSKTKLDGDMIAQLIDSITTQKQRQEFIEHTECNFAILN	75
NTDB id 47407 H7R58 RS04330 WP 182930043.1	..MDIQEIVTILSVKHNVSDDLHLCSAG...NARWRRAGKL...ETAPFSAPHLGVLMGWLSESSQQQTWLRQRQLDFSLSL	72
NTDB id 1448 BW25113 RS15320 WP 001326494.1	..MNMEEIVALSVKHNVSDDLHLCSAG...PARWRIRGRM...EAAPFDTPDVEELLREWLDDDQRAILLENGQLDFAVSL	72
NTDB id 1307 DR RS10055 WP 010888596.1	SGIDITDILRLAADKGASDIILAAAGL...PPQFKLSGDYDAQGFEPISASDTRKLMYSMMNERQQRIFFEEKRELDDFSFAI	79
NTDB id 1086 NMV RS00245 WP 002218487.1	..MQITDLLAFGAKNKASDLHLSSGI...SPMIRVHGDMMRRINLPMSAEVGNMVTSMVNDHQKRIYQQNLEVDDFSFEI	75
NTDB id 1116 NGFG RS10615 WP 003690200.1	..MQITDLLAFGAKNKASDLHLSSGI...SPMIRVHGDMMRRINLPMSAEVGNMVTSMVNDHQKRIYQQNLEVDDFSFEI	75
NTDB id 1185 GTF74 RS11940 WP 000350195.1	..MDIAELLEFSVKHNASDLHLSSAGV...PPMVRIDGEVRKLGVPFATHSDVHRLIFEIMNDAQRSEYEEKLEVDDFSFEI	75
NTDB id 1176 A1552VC RS01160 WP 000350195.1	..MDIAELLEFSVKHNASDLHLSSAGV...PPMVRIDGEVRKLGVPFATHSDVHRLIFEIMNDAQRSEYEEKLEVDDFSFEI	75
NTDB id 1283 ACE17W RS02630 WP 010947729.1	..MDIAELLAFSVKNKSSDLHLISAGM...PPMIRVDGDLRKINLPLEHKDVIKIIYDIMNDRQRKEYEEFFETDFSFEI	75
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NTDB id 1202 PSJM300 01600 AFN76400.1	..MDITELLAFSAKQGASDLHLSSAGL...PPMIRVDGDVRRINLPAMDHKQVHALIYDIMNDKQRKDFEEFLETDFSFEV	75
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NTDB id 1081 FDQ49 RS05335 WP 006581387.1	..MDITELLAFSVKNKASDLHLSSAGM...PPMIRVDGEVRRINLPALHKKDVHRLVYDIMNDKQRQDYEEKLETDFSFEV	75
NTDB id 1060 ABD1 RS04380 WP 000355489.1	..MDITELLAFSVKNKASDLHLSSAGM...PPMIRVDGEVRRINLPALHKKDVHRLVYDIMNDKQRQDYEEKLETDFSFEV	75
NTDB id 1040 HON27 RS13360 WP 000355489.1	..MDITELLAFSVKNKASDLHLSSAGM...PPMIRVDGEVRRINLPALHKKDVHRLVYDIMNDKQRQDYEEKLETDFSFEV	75
consensus	*****!*****	

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NTDB id 1436 BW25113 RS00530 WP 001025145.1
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NTDB id 1283 ACE17W RS02630 WP 010947729.1
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NTDB id 1040 HON27 RS13360 WP 000355489.1
consensus

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NTDB id 1436 BW25113 RS00530 WP 001025145.1	LHTNSTCETIIVRIQQ.....MGVARWMLSSALTLVIAQRLVRKLCPHCRRQQGGEPIHIPDNVWPSPLPHWQAPGCVHCY	389
NTDB id 1177 A1552VC RS01165 WP 000422572.1	LHANNANQALERILHLVLPKEQREQFLDLSLNLKGVIQAQQLLRDK.....	270
NTDB id 1203 PSJM300 01605 AFN76401.1	LHANNANQALDRIINFFPADRQNVWMDLSLNLKAIVAQQQLVPTP.....	272
NTDB id 1019 ACIAD RS04205 WP 004922051.1	LHANNANQTLERIIHFFDADRHHQQTIFMDLSLNLKAVISQQLIPTH.....	273
NTDB id 47407 H7R58 RS04330 WP 182930043.1	LHTRGAAQAIERLVDVFPAGEKEPVNRQLASSLCAVLAAQKLQPD.....	268
NTDB id 1448 BW25113 RS15320 WP 001326494.1	LHTRGAAQAVERLVDSFPAQEKDPVRNQLAGSLRAVLSSQKLEVD.....	268
NTDB id 1307 DR RS10055 WP 010888596.1	LHTNSAPESIDRIVDVFPPEEQEQEIRVQLANNLVAVMTQQLLPRA.....	276
NTDB id 1086 NMV RS00245 WP 002218487.1	LHTTGAAKTVDRIVDVFPAGEKEMVRSMLSESLTAVISQNLKTH.....	272
NTDB id 1116 NGFG RS10615 WP 003690200.1	LHTTGAAKTVDRIVDVFPAGEKEMVRSMLSESLTAVISQNLKTH.....	272
NTDB id 1185 GTF74 RS11940 WP 000350195.1	LHTSSAAKTIDRIIDVFPGSDKDMVRSMLSESLRAVIAAQKLLKR.....	271
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NTDB id 1283 ACE17W RS02630 WP 010947729.1	LHTNSATKTIINRIIDVFPAGEKSMVRSMLSESLQAVVAQSLLKK.....	271
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NTDB id 1060 ABD1 RS04380 WP 000355489.1	LHTTSAAKTIDRVIDVFPAGEKDMVRAMLSESLQAVISQTLKK.....	271
NTDB id 1040 HON27 RS13360 WP 000355489.1	LHTTSAAKTIDRVIDVFPAGEKDMVRAMLSESLQAVISQTLKK.....	271
consensus	!!*****!!********** *!*!!* *!!*!! *!!*!! *!!*	

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NTDB id 1436 BW25113 RS00530 WP 001025145.1	HGFYGRVALFEVLPIPTVIRQLISANTDVE..SLETHARQAGMRTLFEENGCLAVEQGLTTFEELIRVLGMPHGE.....	461
NTDB id 1177 A1552VC RS01165 WP 000422572.1	.NGKGRRHGVFEVLINSPRIADLIRRGELHELKATMARSQEVGMQTFDQALYQLVVDKISEQDALHSADSANDLRLMLKT	349
NTDB id 1203 PSJM300 01605 AFN76401.1	.DGKGRRRAVIEVLINTPLAADLIRKGEVHELKGLMKRSTDLGMQTFDQALYNLYVQGEITYEDALLHADSSNDLRLMIKL	351
NTDB id 1019 ACIAD RS04205 WP 004922051.1	.DGQSRRAAVEVLINTPLLSDIYIRQGEIYKIKELMARSQELGMQTFDQALFDLYDHQHISYADALKHADSPNNLRLQIKL	352
NTDB id 47407 H7R58 RS04330 WP 182930043.1	.NRGGRVALFEMLINTPAVGNLIREGKTHQIAGVMQTGLQSGMQTFSQSYQQRVKAGMLS.....	327
NTDB id 1448 BW25113 RS15320 WP 001326494.1	.KQEGRVALFELLINTPAVGNLIREGKTHQLPHVVIQTGQQVGMITTFQSSYQHRVGEGR.....	326
NTDB id 1307 DR RS10055 WP 010888596.1	.DGQGRVLAYEILLANPAVRRLIREGKTFQITSVMTGTAREGMVTMDAYLAGLFRHLITYDKGLERAID.PKEFARLA.	353
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NTDB id 1194 PA0395 NP 249086.1	.IGGGRVAAHEIMIGTPAIRNLIREDKVAQMYSIIQTGSLGMQTLDMCLKGLVAKGLISRENAREKAKI.PENF.....	344
NTDB id 1202 PSJM300 01600 AFN76400.1	.VGGGRVAAHEIMIGTPAIRNLIREDKVAQMYSIIQTGSLGMQTLDSCLKGLIAKGLISRDSAKEKAKQ.PENF.....	344
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consensus	***!*****!**********!*** ********** *!!*!!* * * * * * * *	

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NTDB id 1436 BW25113 RS00530 WP 001025145.1		461
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NTDB id 1203 PSJM300 01605 AFN76401.1	GSETDGEHLTSVS.QGLTL.....EMSDDDPGRSFR.....	381
NTDB id 1019 ACIAD RS04205 WP 004922051.1	SEEANKPLDNRERMSEMER.....KMTFDGQRRNLK.....	383
NTDB id 47407 H7R58 RS04330 WP 182930043.1		327
NTDB id 1448 BW25113 RS15320 WP 001326494.1		326
NTDB id 1307 DR RS10055 WP 010888596.1	.NDASAGL.GGASMTPAAPPVPAYNSSSSSTGRGGDFGRGAAAPATPAAPAAGRSEPGRAGGFGRR	418
NTDB id 1086 NMV RS00245 WP 002218487.1		347
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NTDB id 1185 GTF74 RS11940 WP 000350195.1		345
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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved