

logo

NTDB id 1257 GC085 RS04855 WP 163621000.1	MRKTIIA....TLILISASISLAYSKDNSLISVKVLPLPENRVRIDFQFSKPLKQLPASFITQ	58
NTDB id 1284 ACE17W RS12370 WP 010946666.1	MRKTIIA....TLILISASLSMAYSKDNSLISVKVLPLPENRVRIDFQFSKPLKQLPASFITQ	58
NTDB id 1009 ACIAD RS15175 WP 004923730.1	MKQTIIRQFSIG..TV....AYAIMQVASAQVAITNVVPMNIPGQGTEIRVMFNGLPP.QPQAYQLE	59
NTDB id 1039 HON27 RS01460 WP 031993779.1	MNHVFRQFSMG..AV....AIAIMQVASAQVSMNTIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLE	59
NTDB id 1052 ABD1 RS15850 WP 001017033.1	MNHVFRQFSMG..AV....AIAIMQVASAQVSMNTIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLE	59
NTDB id 393440 FIV08 RS03235 WP 152437318.1	MNEKNMHEQNGLRSRLAMFKKLNVCVGI...AIGLWTSLASAVTLEDVSFSSLPGERLEVMTMKFDGPPP.EPTGYTIE	75
NTDB id 1193 PAKAF RS26665 WP 003161249.1	MNSGLSRLGLIALLAAMFAPALLAADLEKLDVAALPGDRVELKLQFDEPVA.APRGYTIE	58
consensus	* * * * *	

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NTDB id 1257 GC085 RS04855 WP 163621000.1	KPPRLVLDLFISTDMQIPSDQTKTKTIQLGSLHSYNIVAVGDRVRAILELDRSVSYSGSTAGSVYSLVLNGKSNDLFENSKE	138
NTDB id 1284 ACE17W RS12370 WP 010946666.1	KPPRLVLDLFISTDMQIPSDQTKTKTIQLGSLHSYNIVAVGDRVRAILELDRSVSYSGSTAGSIYSLVLNGKSNDLFENSKE	138
NTDB id 1009 ACIAD RS15175 WP 004923730.1	QPARLILDFDKAGQKL..AKSLIPVTTNEASSIDVSSDDKRARVVVNKLDAAGFTTRVEGNVFILKINNIQSTATPVAT.	136
NTDB id 1039 HON27 RS01460 WP 031993779.1	NPSRLILDFDKAQGGL..KQSKISVATNEASSVDVTSDDQRSRLTVNLKDTGAFTTRVEGNTFILKINSAQTSNKPLPV.	136
NTDB id 1052 ABD1 RS15850 WP 001017033.1	NPSRLILDFDKAQGGL..KQSKISVATNEASSVDVTSDDQRSRLTVNLKDTGAFTTRVEGNTFILKINSAQTSNKPLPV.	136
NTDB id 393440 FIV08 RS03235 WP 152437318.1	RPARIAVDLKDTTSGL..NQRSIPLGSGNAQSVTVVETKDRTRLIFNLVELVPHDTPVRQNNSLVMTIGGASGAPTVTAGA	153
NTDB id 1193 PAKAF RS26665 WP 003161249.1	QPARIALDLPGVQNKLL..GTKNRELSVGNTRSVTVVEAKDRTRLIINLTTLSSYTTRVEGNLNFVVVGNSPAGASVASTA	136
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NTDB id 1257 GC085 RS04855 WP 163621000.1	VFITNQVVNAKHEIKRIDFRGIEKQGGRVLIDVSDTGIPIDVTQVGKEIVVNFLNTKTIPLNLMKRYD	205
NTDB id 1284 ACE17W RS12370 WP 010946666.1	VFITNQVVNAKHEIKRIDFRGIEKQGGRVLIDVSDTGIPIDVTQVGKEIVVNFLNTKTIPLNLMKRYD	205
NTDB id 1009 ACIAD RS15175 WP 004923730.1	SAPAPQCGIGNIGFQRGAQGEGLVVIDLQGTINTPVDVQQQGSKVVRFPFGIKIPHTHLARRLN	198
NTDB id 1039 HON27 RS01460 WP 031993779.1	VS.AQPQGVSNIGFQRGSQGEGLVVVDLLGSNTPVDVQQQGSKVVIRTIGTKIPHTHLARRLN	197
NTDB id 1052 ABD1 RS15850 WP 001017033.1	VS.AQPQGVSNIGFQRGSQGEGLVVVDLLGSNTPVDVQQQGSKVVIRTIGTKIPHTHLARRLN	197
NTDB id 393440 FIV08 RS03235 WP 152437318.1	QQSRTGTTNASAAANALAGVDFRRSKEGEGRVIVDLGSASAPVNLSELAGRIRLTMPGLEVPVVDLRRRLD	222
NTDB id 1193 PAKAF RS26665 WP 003161249.1	PVKASPAPASYAQPIKPKPYVPAGRAIRNIDFQRGEKGEGNVVIDLSDPTLSPDIQEQQGKIRLDFAKTQLPDALRVRLD	216
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NTDB id 1257 GC085 RS04855 WP 163621000.1	VSDFHSPVQIITMQQEGKKVRMTILSKGDYGHFVYQVQNKQFMDVFPDLTAAEIRQAKLKKQVFTGKRISLNFQNISIRAV	285
NTDB id 1284 ACE17W RS12370 WP 010946666.1	VSDFHSPVQIITMQQEGKKVRMTILSKGDYGHFVYQVQNKQFMDVFPDLTAAEIRQAKLKKQVFTGKRISLNFQNISIRAV	285
NTDB id 1009 ACIAD RS15175 WP 004923730.1	TIDFATPVASIDSYNDGSNGVISIQSTGSYEYMAQAEENKLTISLKRQDQNV.RLNSKNNQNYTGKKISLDFQDIEVRRV	277
NTDB id 1039 HON27 RS01460 WP 031993779.1	VNDFATPVSSIDAYNDKGVGVITIQSSGSYEYMAQAEENKLTISLKRQDQKN.VTSLYKTPNYSGNKLSLDFQDIEVRRV	276
NTDB id 1052 ABD1 RS15850 WP 001017033.1	VNDFATPVSSIDAYNDKGVGVITIQSSGSYEYMAQAEENKLTISLKRQDQKN.VTSLYKTPNYSGNKLSLDFQDIEVRRV	276
NTDB id 393440 FIV08 RS03235 WP 152437318.1	VIDFATPVTRIDTFIEDGNAVVEIRPEGNVDYIAYQSGREFTVSVVERISEEAEATREEEKFPYTGDKLSLNFQDIEVRSV	302
NTDB id 1193 PAKAF RS26665 WP 003161249.1	VKDFATPVQFVNASDAQSDRTSITIEPSSGLDYDLVYQTDNRLTVSIKPMTTEDAEERRKKNDFAYTGEKLSLNFQDIDVRSV	296
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NTDB id 1193 PAKAF RS26665 WP 003161249.1
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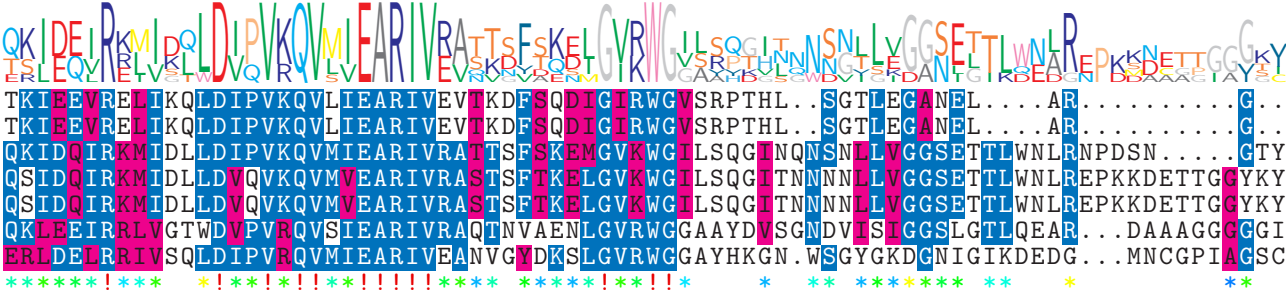
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consensus



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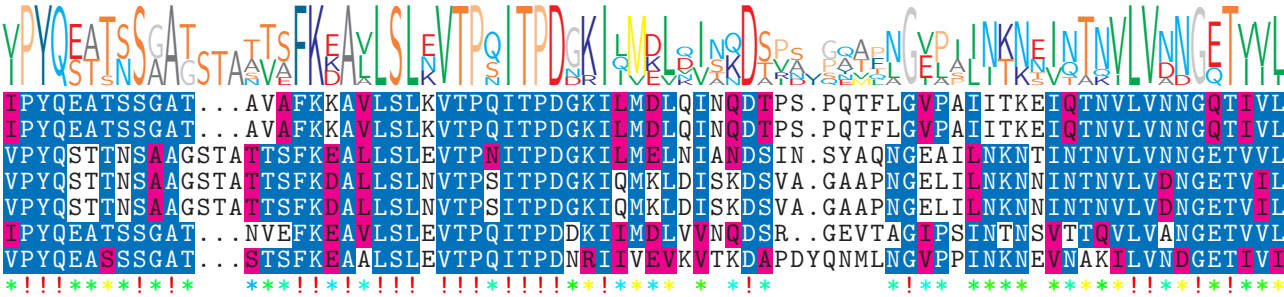
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NTDB id 1193 PAKAF RS26665 WP 003161249.1
consensus



X non conserved
X similar
X ≥ 50% conserved