

NTDB id	1284	ACE17W	RS12370	WP	010946666.1	MRKIIAILIL..ISASLSMAYS	SKDNSLISVKVLPLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLD	FISTDMQIPSDQ	78
NTDB id	1257	GC085	RS04855	WP	163621000.1	MRKIIAILIL..ISASLSLAY	SKDNSLISVKVLPLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLD	FISTDMQIPSDQ	78
NTDB id	1186	GTF74	RS01320	WP	000788426.1				0
NTDB id	1175	A1552VC	RS12105	WP	000788426.1				0
NTDB id	973118	AABD50	RS01610	WP	005488227.1				0
NTDB id	1408	DSB67	RS13980	WP	010643500.1				0
NTDB id	1193	PAKAF	RS26665	WP	003161249.1	MNSGLSRLGIALIAA.MFAPALLAADLEKLIDVAALPGDRVELKLQFDEPVA.APRGYTIEQPARIALDL	PGVQNKLIG..T	76	
NTDB id	1009	ACTAD	RS15175	WP	004923730.1	MKKIIRQFSIGTVAYAIMQVASAQVAITNVVPMNIPGGTEIRVMFNGLPP.QPQAYQLEQPARLI	LDLFDKAGQKLA..K	77	
NTDB id	1039	HON27	RS01460	WP	031993779.1	MNHVFRQFSMGAVAIATIMQAASAQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRL	LDLFDKAQQGLK..Q	77	
NTDB id	1052	ABD1	RS15850	WP	001017033.1	MNHVFRQFSMGAVAIATIMQAASAQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRL	LDLFDKAQQGLK..Q	77	
consensus						* *	* *	*	

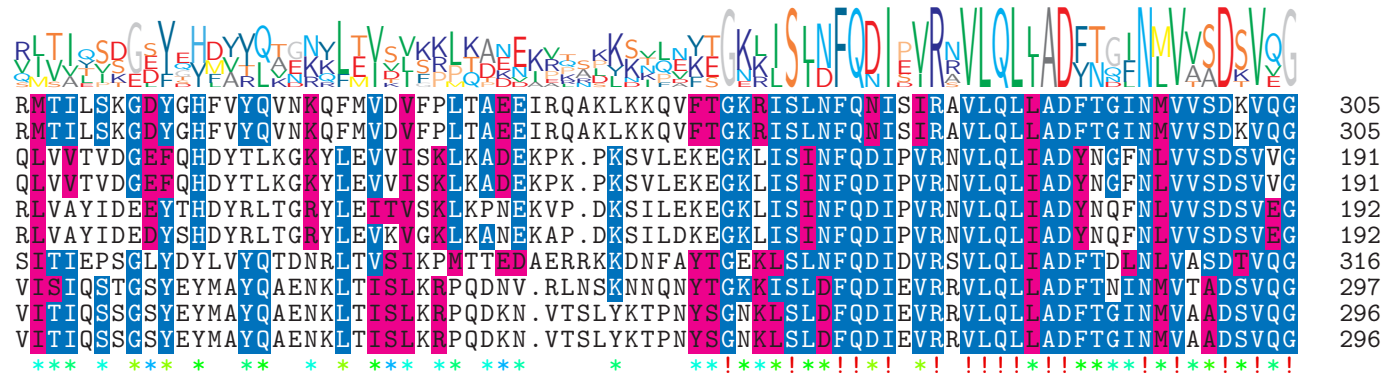
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NTDB id	1257 GC085 RS04855 WP 163621000.1	KTKTITQLGSLHSYNIIVAVGD RVRAILELDRSVSYSGSTAGSVYSVL VLNKGSNDLFENSKE.....VFITNQV	145
NTDB id	1186 GTF74 RS01320 WP 000788426.1	MKNGLKTY..VAQTWLTLWVGL....ALCAS SSM.....VF..SAE	32
NTDB id	1175 A1552VC RS12105 WP 000788426.1	MKNGLKTY..VAQTWLT LWVGL....ALCA SSM.....VF..SAE	32
NTDB id	973118 AABD50 RS01610 WP 005488227.1	MKMRQG LRTPRMLTGFLS WL FL...LLAP.....LA..AAE	33
NTDB id	1408 DSB67 RS13980 WP 010643500.1	M KMRH GL TLRPRKL TG FLS W FS L LL APFV ..AA E	33
NTDB id	1193 PAKAF RS26665 WP 003161249.1	KNREISVG NTRS VTIVE AKDRT RL I IN LT TL SS Y TT RVEGN NFVV GN SP AG AS VA STAPVKAS PAPASYAQPIKP KY	156
NTDB id	1009 ACIAD RS15175 WP 004923730.1	SLIPVTTENAS SIDVSDDKRARVVVN LKDAGAFTTRVEGN VFILKIN NI QSTAT.....PVAT.. SA.....	138
NTDB id	1039 HON27 RS01460 WP 031993779.1	SKIS VATNEASSVDVTSD DQRSRLTVN L KD TA GA FT TR VE GN TF IL KI NS AQ TS NK.....PLP V.. VS.....	138
NTDB id	1052 ABD1 RS15850 WP 001017033.1	SKIS VATNEASSVDVTSD DQRSRLTVN L KD TA GA FT TR VE GN TF IL KI NS AQ TS NK.....PLP V.. VS.....	138
consensus		* * * * ** **** * ***** *	*

NTDB id	1284 ACE17W RS12370 WP 010946666.1	VNAKHEIKRIDFRGIEKKQGGRLVIDVSDTGIPIDVTQVGKEIVVNFLNLTAKIPLNLMKRYDVSDHSPVQIIITMQQEGKKV	225
NTDB id	1257 GC085 RS04855 WP 163621000.1	VNAKHEIKRIDFRGIEKKQGGRLVIDVSDTGIPIDVTQVGKEIVVNFLNLTAKIPLNLMKRYDVSDHSPVQIIITMQQEGKKV	225
NTDB id	1186 GTF74 RS01320 WP 000788426.1	SATANQLENIDFRVNKEKA AVLVELASPSAVVDVVQKVQEGLSIELLKTADVADDKLYLLDVKDFS TPVESVEVFRKEPST	112
NTDB id	1175 A1552VC RS12105 WP 000788426.1	SATANQLENIDFRVNKEKA AVLVELASPSAVVDVVQKVQEGLSIELLKTADVADDKLYLLDVKDFS TPVESVEVFRKEPST	112
NTDB id	973118 AABD50 RS01610 WP 005488227.1	DVQS NALKS IDFRTNKSK DAVIVVELASPAAI VDV KRV QEGLSID L MNTSVKDEQLYLLDVKDFATPVESVEVFRDTSTT	113
NTDB id	1408 DSB67 RS13980 WP 010643500.1	EVQS NSLK SIDFRTNKDKDAVI VVELVS PAAI VDI KR V QEGLSID LL NT SV KDDQLYLLDVKDFS TVVESVEVFRDTSTT	113
NTDB id	1193 PAKAF RS26665 WP 003161249.1	VPAGRAIRNI DFQRGEKG EGNVVIDLSDP TLPDIQEEQQGK I RL DFA KTQLPD ALRVR LD VKDF ATPVQ F VN AS AQSD RT	236
NTDB id	1009 ACTAD RS15175 WP 004923730.1	PAPQGIGIGNIGFQRGAQGEGLVVIDLQGTNT PV DV QQ QS KV V VRFPGIKIPTHLARRLNTTD FATPVAS IDS YNDGSNG	218
NTDB id	1039 HON27 RS01460 WP 031993779.1	.AQPQGVSNIGFQRGSQGEGLVVVDLLG SNT PD VD QQ QS KV VIR TI GT KIPTHLARRLNVNDFATPVSSIDA YNDKGVG	217
NTDB id	1052 ABD1 RS15850 WP 001017033.1	.AQPQGVSNIGFQRGSQGEGLVVVDLLG SNT PD VD QQ QS KV VIR TI GT KIPTHLARRLNVNDFATPVSSIDA YNDKGVG	217
consensus		* *****! * ! * ***** * **!* **** * **** * **********!!**!* ***** *	

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consensus

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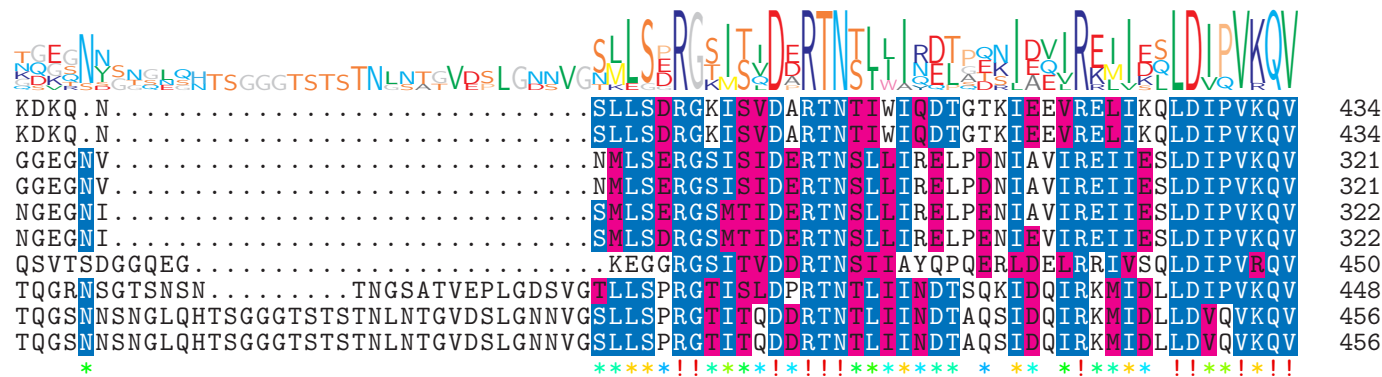
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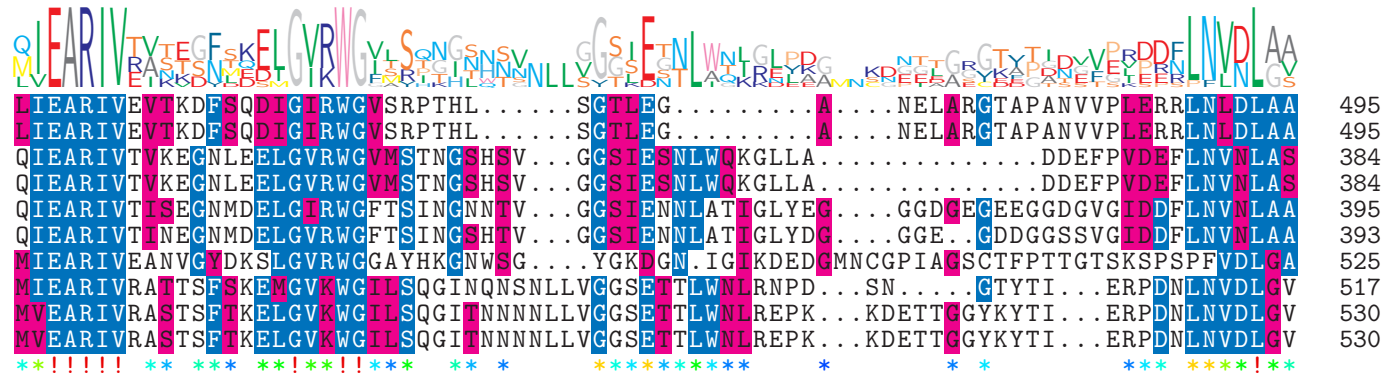
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consensus

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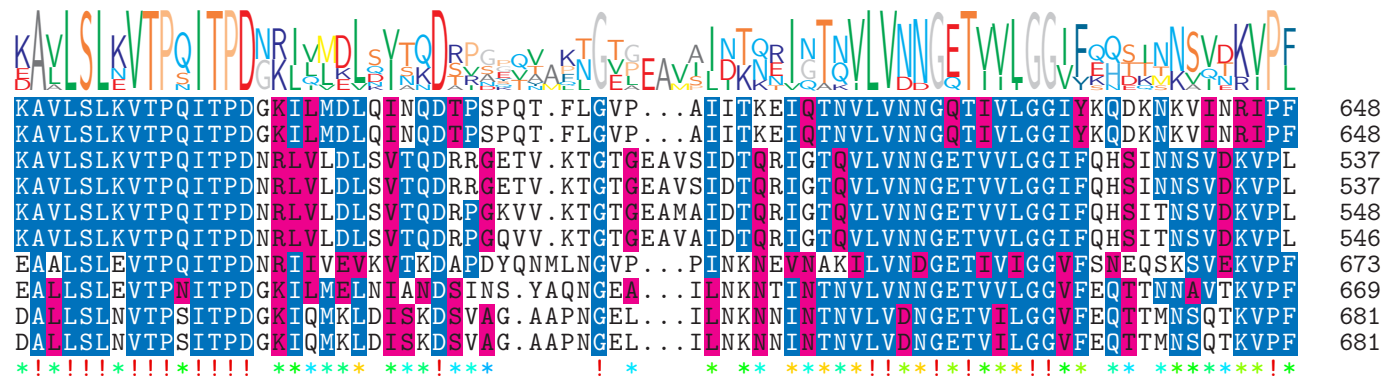
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consensus
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consensus

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logo



non conserved
 similar
 ≥ 50% conserved