

logo

NTDB id	1284	ACE17W	RS12370	WP	010946666.1	MRKIIAILIL..ISASLSMAYSKDNSLISVKVLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDFISTDMQIPSDQ	78
NTDB id	1257	GC085	RS04855	WP	163621000.1	MRKIIAILIL..ISASLSLAYSKDNSLISVKVLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDFISTDMQIPSDQ	78
NTDB id	1186	GTF74	RS01320	WP	000788426.1		0
NTDB id	1175	A1552VC	RS12105	WP	000788426.1		0
NTDB id	284244	C7Y67	RS01200	WP	005488227.1		0
NTDB id	1408	DSB67	RS13980	WP	010643500.1		0
NTDB id	1193	PAKAF	RS26665	WP	003161249.1	MNSGLSLRGIALIAA.MFAPALLAADLEKLDVAALPGDRVELKLQFDEPVA.APRGYTIEQPARIALDLPGVQNKG..T	76
NTDB id	1009	ACIAD	RS15175	WP	004923730.1	MKQIRQFSIGTVAYAIMQVASAQVAITNVVPMNIPGGTEIRVMFNGLPP.QPQAYQLEQPARLILDFDKAGQKLA..K	77
NTDB id	1039	HON27	RS01460	WP	031993779.1	MNHVFRQFSMGAVAIAIMQAASAQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRLILDFDKAQGGLK..Q	77
NTDB id	1052	ABD1	RS15850	WP	001017033.1	MNHVFRQFSMGAVAIAIMQAASAQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRLILDFDKAQGGLK..Q	77
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[illegible]

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NTDB	id	1284	ACE17W	RS12370	WP	O10946666.1	VNAKHEIKRIDFRGIEKQGGRVLIDVSDTGIPIDVTQVGKEIVVNFLNNTKIPLNLMKRYDVSDFHSPVQIITMQEGKKV	225
NTDB	id	1257	GCO85	RS04855	WP	163621000.1	VNAKHEIKRIDFRGIEKQGGRVLIDVSDTGIPIDVTQVGKEIVVNFLNNTKIPLNLMKRYDVSDFHSPVQIITMQEGKKV	225
NTDB	id	1186	GTFT4	RS01320	WP	000788426.1	SATANQLENIDFRVNKEKA AVLIVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFS TPVESVEVFRKEPST	112
NTDB	id	1175	A1552VC	RS12105	WP	000788426.1	SATANQLENIDFRVNKEKA AVLIVELASPSAVVDVQKVQEGLSIELLKTDVADDKLYLLDVKDFS TPVESVEVFRKEPST	112
NTDB	id	284244	CY767	RS01200	WP	005488227.1	DVQSNALESIDFRTNKSKDAVVVELLASPA AIVDKVRVQEGLSIDLMTNSVKDEQLYLLDVKDFATV VESVEVFRDTSTT	113
NTDB	id	1408	DSB67	RS13980	WP	010643500.1	EVQSN SLKSIDFR TNKD KDAIVVVELVSPA AIVDKRVQEGLSIDLMTNSVKDDQLYLLDVKDFS TVVESVEVFRDTSTT	113
NTDB	id	1193	PAKAF	RS26665	WP	003161249.1	VPAGRAIRNIDFRQRGEKGECNVV TDLSDPTLSPIEQEGGKIRLDFAKTQLPDALRVRLDVKDFATPV QFVNASAQSDRT	236
NTDB	id	1009	ACIAD	RS15175	WP	004923730.1	PAPQQIGIGNIFGRGAQGECLVV IDLQGTNT PVDVQQQGSKV VVRFPGIKIPTHLAR RLNTTDFATPVA SIDS YNDGSGN	218
NTDB	id	1039	HON27	RS01460	WP	031993779.1	.AQPQGVSNIGFQRGSQGEGLVVVDLLGSNT PV D V Q Q Q G S K V V I R T I G T K I P T H L A R R L N V N D F A T P V S I D A Y N D K G V G	217
NTDB	id	1052	ABD1	RS15850	WP	001017033.1	.AQPQGVSNIGFQRGSQGEGLVVVDLLGSNT PV D V Q Q Q G S K V V I R T I G T K I P T H L A R R L N V N D F A T P V S I D A Y N D K G V G	217
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NTDB id 284244 CTY67 RS01200 WP 005488227.1
NTDB id 1408 DSB67 RS13980 WP 010643500.1
NTDB id 1193 PAKAF RS26665 WP 003161249.1
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NTDB id 1052 ABD1 RS15850 WP 001017033.1
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consensus

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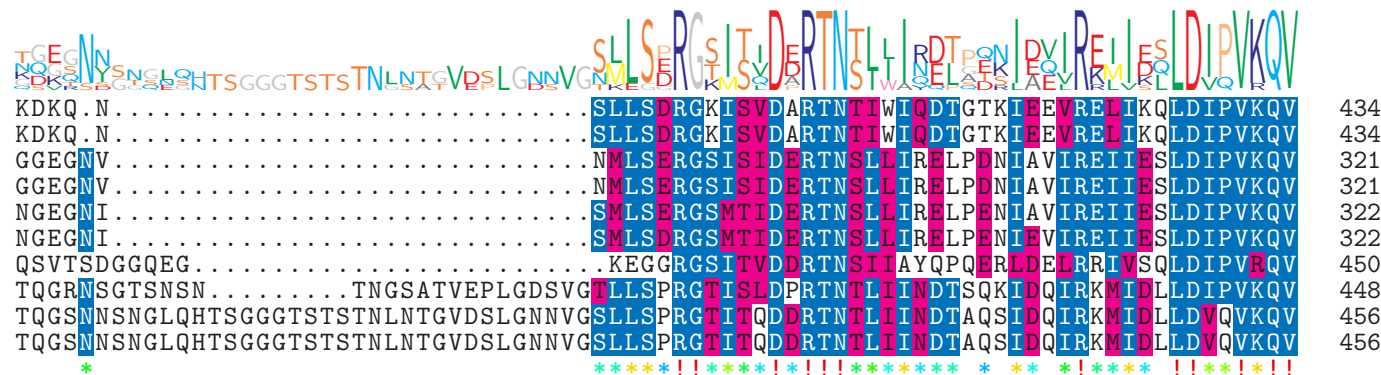


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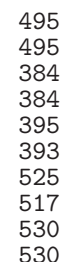
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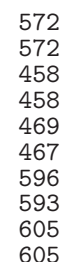
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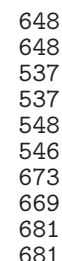
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X non conserved
X similar
X ≥ 50% conserved