

logo

NTDB id 1186	GTF74	RS01320	WP 000788426.1	MKMNGLKLTTPKATWTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKAAVLIVELASPSAVVDVQKVQEGLSI	76
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	..MKNGLKT..YVAQTWLTWVGLALCASSMVFSAESATANQLENIDFRVNKEKAAVLIVELASPSAVVDVQKVQEGLSI	76
NTDB id 1408	DSB67	RS13980	WP 010643500.1	MKMRHGLTLTPRKLTGLFSLWFSLLAP...FVAAEEVQSNLSKSIDFRTNKDKDAVIVVELVSPAAIVDIKRVQEGLSI	77
NTDB id 70994	kpn2146	RS25440	WP 004174030.1		0
NTDB id 1433	BW25113	RS17570	WP 000815987.1		0
NTDB id 1347	HI 0435	AAC22094.1			0
NTDB id 1373	NTHI	RS02645	WP 011272046.1		0
NTDB id 1381	A4U84	RS02010	WP 021113424.1		0
NTDB id 1465	ASU2	RS06170	WP 014991934.1		0
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NTDB id 1186	GTF74	RS01320	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDFGEFQHDYTLKGYLEVVISKKADEKPKPKSVLEKEG	156
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDFGEFQHDYTLKGYLEVVISKKADEKPKPKSVLEKEG	156
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NTDB id 70994	kpn2146	RS25440	WP 004174030.1	MKRMLISLTLPLPLAVAPAAARN	23
NTDB id 1433	BW25113	RS17570	WP 000815987.1	MKQVIAADLLMLIPGVQ...AAKP	21
NTDB id 1347	HI 0435	AAC22094.1		MKKYFLKCGYFLVCFCLPLVFAN...P.KTDN	29
NTDB id 1373	NTHI	RS02645	WP 011272046.1	MKKYFLKCGYFLVCFCLPLVFAN...P.KTDN	29
NTDB id 1381	A4U84	RS02010	WP 021113424.1	MRVLFLLFFATF...P.VLAN	17
NTDB id 1465	ASU2	RS06170	WP 014991934.1	MKRMILLFFIL...P.PVIA	17
consensus					

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NTDB id 1186	GTF74	RS01320	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVVSDSVVGNLTLRLDGVPWQQVLDIILQVKGDKHVDGNVILAPKEFLDLRE	236
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NTDB id 1433	BW25113	RS17570	WP 000815987.1	QKVITLMVDDVPVACVLQALAEQEKLNLVVSPDVSCTVSLHLDTPVPWQALQTVVKSAGLITHQECNLLSVHSIAWQNNNI	101
NTDB id 1347	HI 0435	AAC22094.1		ERFFIRLSQAPLAQTLEQLAFQDDVNLVIGDILENKISLKLNNIDMPRLQITAKSKHILTNKDDGIYYLNGSQSGKGQV	109
NTDB id 1373	NTHI	RS02645	WP 011272046.1	ECFFIRLSQAPLAQTLEQLAFQDDVNLVIGERLEGNISLKLNNIDMPRLKLIKAKSKHILTNKDDGIYYLNGSQSGKGQV	109
NTDB id 1381	A4U84	RS02010	WP 021113424.1	QQISISIRNAPTAETISYLAETGKNITISDEIKDTKNFRVEKSHDEILNSLIKTHQLNLKENGYYIYHQAQEHKQHT	97
NTDB id 1465	ASU2	RS06170	WP 014991934.1	HAFSISLKNAPTAKILSYLAEBHGKNIVLSDNIETNTLRIENSNDVLSKSIHANKITSAYENQIYFIHGKKDDKDPN	97
consensus					

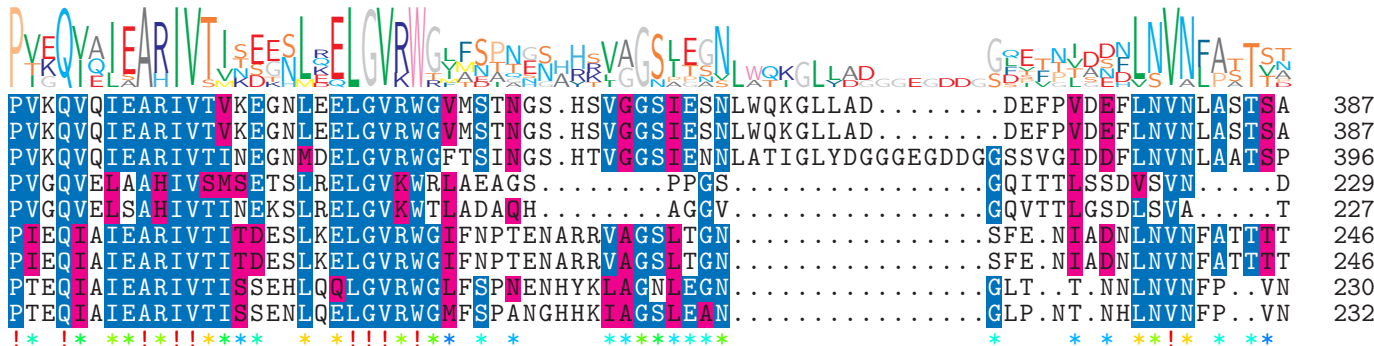
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NTDB id 1433	BW25113	RS17570	WP 000815987.1	ARQEAQARAQANLPLENRSITLQYADAGEL...A.KAGEKLLSAKGSMTVDKRTNRLLRDNKTAISALEQWVAQMDI	176
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NTDB id 1465	ASU2	RS06170	WP 014991934.1	RG....LNLLELLKPKLITIKTLHYKASEVIESI.T.KCQCNLLSESCYLFHFDPRSNSIIVKDSAASVKNFTQLIESLDK	172
consensus					

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NTDB id 70994 kpn2146 RS25440 WP 004174030.1
NTDB id 1433 BW25113 RS17570 WP 000815987.1
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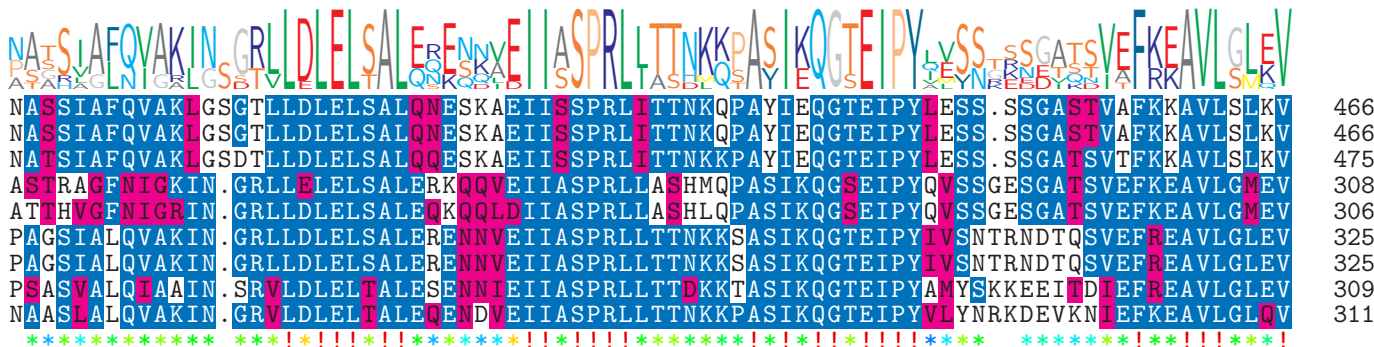
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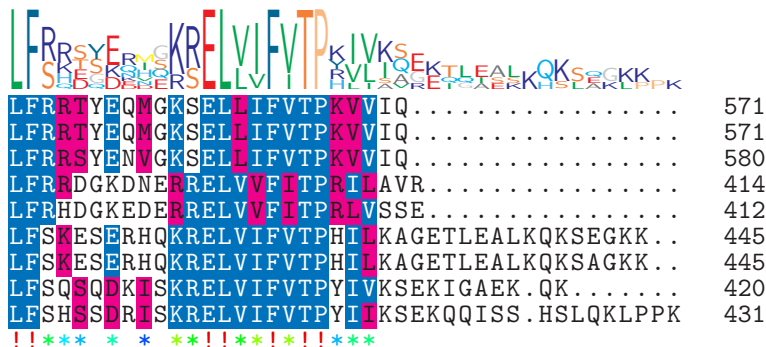
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NTDB id 1465 ASU2 RS06170 WP 014991934.1
consensus

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 non conserved
 similar
 $\geq 50\%$ conserved