

logo

NTDB id 1186	GTF74	RS01320	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKAAVLVVELASPSAVVDVQKVQEGLSI	76
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKAAVLVVELASPSAVVDVQKVQEGLSI	76
NTDB id 1408	DSB67	RS13980	WP 010643500.1	MKMRHGLTLTPRKLTGLFSLWFSLLAP...FVAAEEVQSNLSKSIDFRNTNKDKDAVIVVELVSPA AIVDIKRVQEGLSI	77
NTDB id 50323	SMKP03	RS01815	WP 121593180.1		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	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDYTLKGKYLEVVISKLKADEKPKPKSVLEKEG	156
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDYTLKGKYLEVVISKLKADEKPKPKSVLEKEG	156
NTDB id 1408	DSB67	RS13980	WP 010643500.1	DLLNTSVKDDQLYLLDVKDFSTVSVESVEVFRDSTTTRLVAYIDEDYSHDYRLTGRYLEVKVGK LKAN EKAPDKSILDKEG	157
NTDB id 50323	SMKP03	RS01815	WP 121593180.1	MMRWISLLLLLLPLAVAPAARNND	23
NTDB id 1433	BW25113	RS17570	WP 000815987.1	MKQWIAALLLMLIPGVQ...AAKP	21
NTDB id 1347	HI 0435	AAC22094.1		MKKYFLKCGYFLVCFCLPLIVFAN....P.KTDN	29
NTDB id 1373	NTHI	RS02645	WP 011272046.1	MKKYFLKCGYFLVCFCLPLIVFAN....P.KTDN	29
NTDB id 1381	A4U84	RS02010	WP 021113424.1	MRYLFLFFATF....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	KLISINFQDIPVRNVLQLIADYNGFNLVVSDSVVGNTLRLDGVPPWQVLDIILQVKGLDKRVDGNVILIAPKEELDRLRE	236
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NTDB id 50323	SMKP03	RS01815	WP 121593180.1	KPVSLVIDDAPVAQVLQALAEENHKNLVVAPDVSGTSLRLQKVPWSQALRAVADSAGLSLQQGTVIYAHTQVWVKANQ	103
NTDB id 1433	BW25113	RS17570	WP 000815987.1	QKVTLMVDDVPVAQVLQALAEQEKLNLVVSPDVSGTVSLHLTDVPWKQALQTVVKSAGLITRQEGNLSVHSIAWQNNNI	101
NTDB id 1347	HI 0435	AAC22094.1		ERFFIRLSQAPLAQTLEQLAFQQDVNLVIGDILENKISLKLNNIDMPRLQLTIAKSKHLTLNKDDGIYYLNGSQSGKGQV	109
NTDB id 1373	NTHI	RS02645	WP 011272046.1	ECFFIRLSQAPLAQTLEQLAFQQDVNLVIGERLEGNISLKLNNIDMPRLKLTIAKSKHLTLNKDDGVYYLNGSQSGKGQV	109
NTDB id 1381	A4U84	RS02010	WP 021113424.1	QQISLSIRNAPTAETISYLAEEETGKNITISDEIKDTKNFRVEKSHFDEILNSLIKTHQLNLKKENGIIYIYHQAQEHKQHT	97
NTDB id 1465	ASU2	RS06170	WP 014991934.1	HAFSISLKNAPTAKILSYLAEEHGKNIVLSDNIETNTTLRIENSNFDSVLKSIITRANKLTSAYENQIYFIGHKDDKDPN	97
consensus					

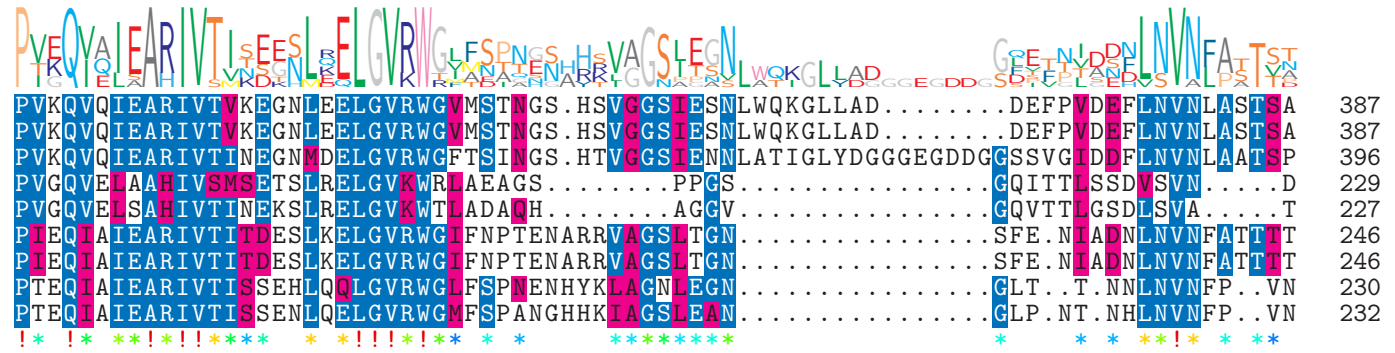
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NTDB id 1433	BW25113	RS17570	WP 000815987.1	ARQEAQARAQANLPLENRSITLQYADAGEL...A.KAGEKLLSAKGSMTVDKRTNRLLRDKNKTALSALEQWVAQMDL	176
NTDB id 1347	HI 0435	AAC22094.1		.A....GNLTTNEPHLVSHTVKHLHFAKASELMKSLT.TGSGSLLSPAGSITFDDRSNLLVIQDEPRSVQNIKKLIAEMDK	183
NTDB id 1373	NTHI	RS02645	WP 011272046.1	.A....GNLTTNEPHLVSHTVKHLHFAKASELMKSLT.TGSGSLLSPAGSITFDDRSNLLVIQDEPRFVQNIKKLIAEMDK	183
NTDB id 1381	A4U84	RS02010	WP 021113424.1	.T....AQLVNALPKLITKTIKHLHYSKASEVIESLT.KGQGNLLSESGYLHFDPRSNSIIVKDSAASVKNFQTQLIESLDK	171
NTDB id 1465	ASU2	RS06170	WP 014991934.1	RG....LNLELLKPKLITKTIKLDYAKAAEVIESLT.KGSGNFLSENGYLHFDPRSNSLIKIDSPESMKNTVKLIRNLDR	172
consensus					

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NTDB id 1186 GTF74 RS01320 WP 000788426.1
NTDB id 1175 A1552VC RS12105 WP 000788426.1
NTDB id 1408 DSB67 RS13980 WP 010643500.1
NTDB id 50323 SMKPO3 RS01815 WP 121593180.1
NTDB id 1433 BW25113 RS17570 WP 000815987.1
NTDB id 1347 HI 0435 AAC22094.1
NTDB id 1373 NTHI RS02645 WP 011272046.1
NTDB id 1381 A4U84 RS02010 WP 021113424.1
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consensus

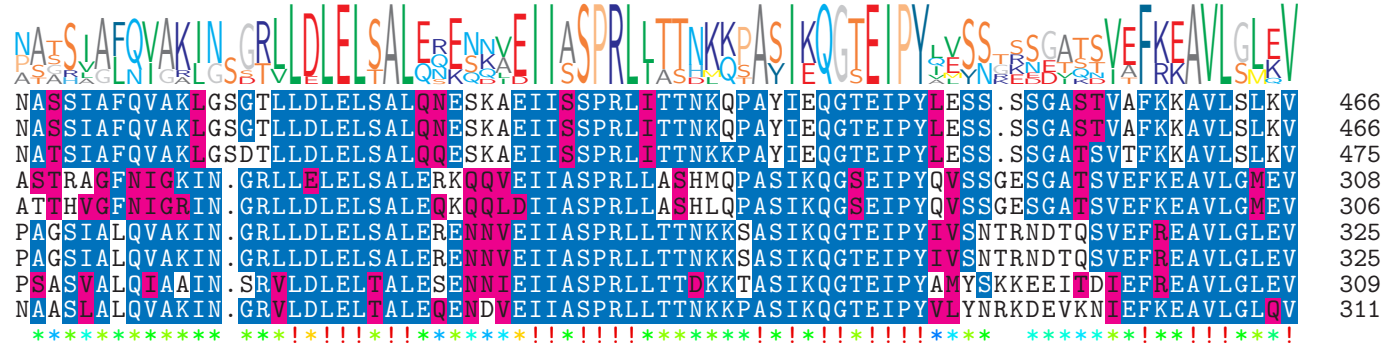
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consensus

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 non conserved
 similar
 $\geq 50\%$ conserved