

logo

NTDB id 1186	GTF74	RS01320	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKAAVLVIVELASPSAVVDVQKVQEGLSI	76
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	..MKNGLKT..YVAQTWLTLWVGLALCASSMVFSAESATANQLENIDFRVNKEKAAVLVIVELASPSAVVDVQKVQEGLSI	76
NTDB id 1408	DSB67	RS13980	WP 010643500.1	MKMRHGLTLTPRKLTGLFSLWFSLLAP...FVAAEEVQSNLSKSIDFRNTKDKDAVIVVELVSPAIVDIKRVQEGLSI	77
NTDB id 1347	HI 0435	AAC22094.1			0
NTDB id 1373	NTHI	RS02645	WP 011272046.1		0
NTDB id 253412	CRH16	RS07475	WP 010785953.1		0
NTDB id 1381	A4U84	RS02010	WP 021113424.1		0
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NTDB id 1186	GTF74	RS01320	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDYTLKGYKYLEVVISKLKADEKPKPKSVLEKEG	156
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	ELLKTDVADDKLYLLDVKDFSTPVESVEVFRKEPSTQLVVTVDGEFQHDYTLKGYKYLEVVISKLKADEKPKPKSVLEKEG	156
NTDB id 1408	DSB67	RS13980	WP 010643500.1	DLNNTSVKDDQLYLLDVKDFSTVSVESVEVFRDSTTTRLVAYIDEDYSHDYRLTGRYLEVKGKLGKANEKAPDKSILDKEG	157
NTDB id 1347	HI 0435	AAC22094.1		MKKYFLKCGYFLVCFCLPLIVF...A..NPKTDN	29
NTDB id 1373	NTHI	RS02645	WP 011272046.1	MKKYFLKCGYFLVCFCLPLIVF...A..NPKTDN	29
NTDB id 253412	CRH16	RS07475	WP 010785953.1	MRYLFLFFAT.....FPVLAN	17
NTDB id 1381	A4U84	RS02010	WP 021113424.1	MRYLFLFFAT.....FPVLAN	17
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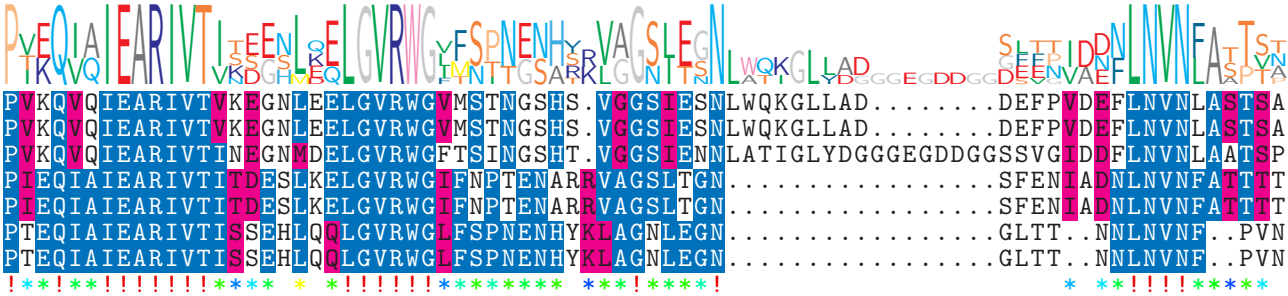
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NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLTLRLDGVPWQQVLDIILQVKGLDKRVDGNVILVAPKEELDRLRE	236
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLTLRLDGVPWQQVLDIILQVKGLDKRVDGNVILVAPKDELRLRE	237
NTDB id 1347	HI 0435	AAC22094.1		ERFFIRLSQAPLAQTLEQLAFQDDVNLVIGDILENKISLKLNNIDMPRLQIIAKSKHLTLNKDDGYIYLNQSQS.....	104
NTDB id 1373	NTHI	RS02645	WP 011272046.1	ECFFIRLSQAPLAQTLEQLAFQDDVNLVIGERLEGNISLKLNNIDMPRLQIIAKSKHLTLNKDDGVYIYLNQSQS.....	104
NTDB id 253412	CRH16	RS07475	WP 010785953.1	QQISLSIRNAPTAETIIGYLAETGKNITISDEIKDTKNFRVEKSHFDEILNSLIKTHQLNLKKENGITYYIHQAQE.....	92
NTDB id 1381	A4U84	RS02010	WP 021113424.1	QQISLSIRNAPTAETIISYLAETGKNITISDEIKDTKNFRVEKSHFDEILNSLIKTHQLNLKKENGITYYIHQAQE.....	92
consensus					

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NTDB id 1186	GTF74	RS01320	WP 000788426.1	KQALEKARLAEEELGDLKSEIIKINFAKASDIAAMIGGEGNVNMLSERGSISIDERTNSLLIRELPDNIIVIREIIESLDI	316
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KQALEKARLAEEELGDLKSEIIKINFAKASDIAAMIGGEGNVNMLSERGSISIDERTNSLLIRELPDNIIVIREIIESLDI	316
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KQALEKQKLEKEIGELSSSEIKVNFASDIAEMINGEGNISMLSDRGSMTIDERTNSLLIRELPENIEVIREIIESLDI	317
NTDB id 1347	HI 0435	AAC22094.1		GKQQVAGNLTNNEPHLVSHTVKLHFAKASELMKSLT.TGSGSLLSPAGSITFDDRSNLLVIQDEPRSVQNIKKLIAEMDK	183
NTDB id 1373	NTHI	RS02645	WP 011272046.1	GKQQVAGNLTNNEPHLVSHTVKLHFAKASELMKSLT.TGSGSLLSAGSITFDDRSNLLVIQDEPRFVQNIKKLIAEMDK	183
NTDB id 253412	CRH16	RS07475	WP 010785953.1	HKQHTTAQLVNALPKLITKTIKLYHASKASEVIESLT.KGQGNLLSENGYLFHDDRSNSIIVKDSAAVKNFTQLIETLDK	171
NTDB id 1381	A4U84	RS02010	WP 021113424.1	HKQHTTAQLVNALPKLITKTIKLYHASKASEVIESLT.KGQGNLLSENGYLFHDDRSNSIIVKDSAAVKNFTQLIESLDK	171
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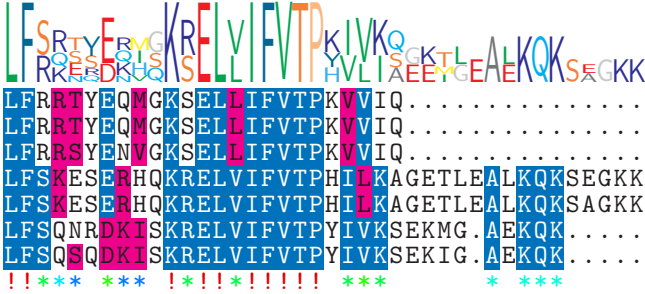
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consensus



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NTDB id 253412 CRH16 RS07475 WP 010785953.1
NTDB id 1381 A4U84 RS02010 WP 021113424.1
consensus



571
571
580
445
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☒ non conserved
☒ similar
☒ ≥ 50% conserved