

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...FVAAEEVQSNLSKSIDFRNTKDKDAVIVVELVSPA AIVDIKRVQEGLSI	77
NTDB id 1347	HI 0435	AAC22094.1			0
NTDB id 1373	NTHI	RS02645	WP 011272046.1		0
NTDB id 78796	JL26	RS05960	WP 026917118.1		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	DLLNTSVKDDQLYLLDVKDFSTVSVESVEVFRDSTTTRLVAYIDEDYSHDYRLTGRYLEVKGKLGKANEKAPDKSILDKEG	157
NTDB id 1347	HI 0435	AAC22094.1		MKKYFLKCGYFLVCFCLPLIVFA.....NPKTDN	29
NTDB id 1373	NTHI	RS02645	WP 011272046.1	MKKYFLKCGYFLVCFCLPLIVFA.....NPKTDN	29
NTDB id 78796	JL26	RS05960	WP 026917118.1	MRYLFLFFAT.....FPVLAN	17
NTDB id 1381	A4U84	RS02010	WP 021113424.1	MRYLFLFFAT.....FPVLAN	17
NTDB id 1465	ASU2	RS06170	WP 014991934.1	MKRMILLFFI.....LPPVIA	17
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NTDB id 1186	GTF74	RS01320	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLTLRLDGVPWQQVLDIILQVKGLDKRVDGNVILTAPKEELDLRE	236
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLTLRLDGVPWQQVLDIILQVKGLDKRVDGNVILTAPKEELDLRE	236
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KLISINFQDIPVRNVLQLIADYNGFNLVSDSVVGNLTLRLDGVPWQQVLDIILQVKGLDKRVDGNVILVAPKDELDLRE	237
NTDB id 1347	HI 0435	AAC22094.1		ERFFIRISQAPLAQTLEQLAFQDDVNLVIGDILENKISLKLNNIDMPRLQIIAKSKHLLTNKDDGIYYLNGSQS....G	105
NTDB id 1373	NTHI	RS02645	WP 011272046.1	ECFFIRISQAPLAQTLEQLAFQDDVNLVIGDILENKISLKLNNIDMPRLQIIAKSKHLLTNKDDGIYYLNGSQS....G	105
NTDB id 78796	JL26	RS05960	WP 026917118.1	QQISLSIRNAPTAETIISYLAEEETGKNITISDEIKDTKNFRVEKSHFDEILNSLIKTHQLNLKKENGIIYIHQAQE....H	93
NTDB id 1381	A4U84	RS02010	WP 021113424.1	QQISLSIRNAPTAETIISYLAEEETGKNITISDEIKDTKNFRVEKSHFDEILNSLIKTHQLNLKKENGIIYIHQAQE....H	93
NTDB id 1465	ASU2	RS06170	WP 014991934.1	HAFSISLKNAPTAKILSYLAEEHGKNIVLSDNIETNTLRINIENSNFDSVLKSIITRANKLTSAYENQIYFIGHKKD....D	93
consensus					

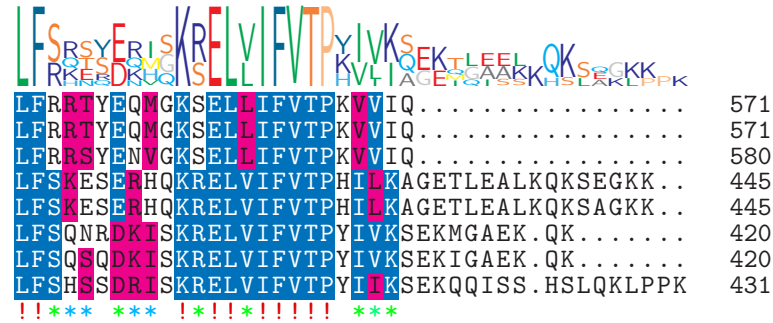
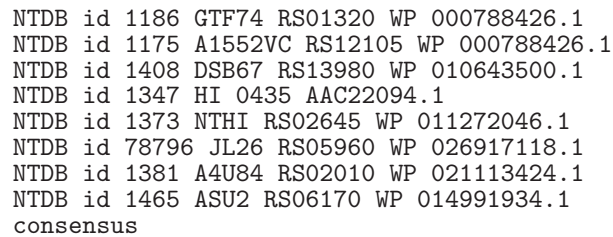
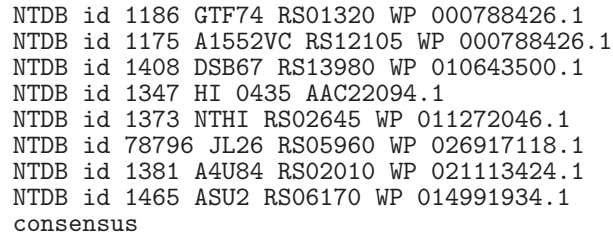
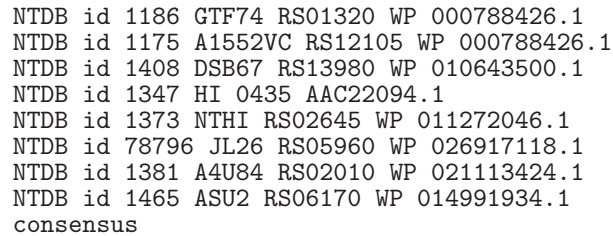
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NTDB id 1186	GTF74	RS01320	WP 000788426.1	KQALEKARLAEEELGDLKSEIIKINFAKASDIAAMIGGEGNVNMLSERGSISIDERTNSLLIRELPDNIIVIRETIESLDI	316
NTDB id 1175	A1552VC	RS12105	WP 000788426.1	KQALEKARLAEEELGDLKSEIIKINFAKASDIAAMIGGEGNVNMLSERGSISIDERTNSLLIRELPDNIIVIRETIESLDI	316
NTDB id 1408	DSB67	RS13980	WP 010643500.1	KQALEKQKLEKEIGELSSSEIKVNFAKASDIAEMINEGNIISMLSDRGSMITIDERTNSLLIRELPENIEVIRETIESLDI	317
NTDB id 1347	HI 0435	AAC22094.1		KGQV..AGNLTITNEPHLVSHTVKLHFAKASELMKSLT..TGSGSLLSPAGSITFDDRSNLLVIQDEPRSVQNIKKLIAEMDK	183
NTDB id 1373	NTHI	RS02645	WP 011272046.1	KGQV..AGNLTITNEPHLVSHTVKLHFAKASELMKSLT..TGSGSLLSSAGSITFDDRSNLLVIQDEPRFVQNIKKLIAEMDK	183
NTDB id 78796	JL26	RS05960	WP 026917118.1	KQHT..TAQLVNALPKLITKTIKLYSKASEVIESLT..KGQGNLLSENGYLHFDDRSNSIIVKDSAAASVKNFQTQLIESLDK	171
NTDB id 1381	A4U84	RS02010	WP 021113424.1	KQHT..TAQLVNALPKLITKTIKLYSKASEVIESLT..KGQGNLLSENGYLHFDDRSNSIIVKDSAAASVKNFQTQLIESLDK	171
NTDB id 1465	ASU2	RS06170	WP 014991934.1	KDPNRGLNLELLKPKLITKTIKLYAKAAEVIESLT..KGSNGFLSENGYLHFDDRSNSLIKDSPESMKNITVQLIRNLDR	172
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NTDB id 1465 ASU2 RS06170 WP 014991934.1
consensus

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 non conserved
 similar
 $\geq 50\%$ conserved