

logo

NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1366 NTHI RS01950 WP 011271972.1		0
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NTDB id 1468 ASU2 RS05090 WP 014991734.1		0
NTDB id 64355 AABK06 RS18460 WP 001025210.1		0
NTDB id 1436 BW25113 RS00530 WP 001025145.1		0
NTDB id 1168 A1552VC RS11070 WP 000957200.1		0
NTDB id 1401 DSB67 RS12665 WP 010643259.1		0
NTDB id 1293 VP RS12240 WP 005479695.1		0
NTDB id 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
consensus		

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NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	160
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NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
consensus		

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NTDB id 1028 TT RS08215 WP 011173991.1	EGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL	240
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NTDB id 1251 GC085 RS07725 WP 011946523.1		0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
consensus		

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NTDB id 1028 TT RS08215 WP 011173991.1	KEALEVQKGLPRAKPLGEIIVELGLARPEDVEEALQKQRRGGGRLEDITLVQSGKLRPEALIAQAVATQLGYPYVDPEEDPP	400
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NTDB id 1110 NGFG RS09235 WP 003689817.1	MSVGLLRILVQNQVVTVEQAEHYY.NESQAGKEVLPMFLSDGVISPKSLAALIARVFSYSILDLRHYPR	68
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consensus	*	

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NTDB id 1028 TT RS08215 WP 011173991.1	DPGAPLLLPEDLCRRYGVFPHRLEGNRLVLLMKDPRNILLALDDVRLALKRKGLNVEVAPAVATEAAITKLTIERFYGKAEL	480
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NTDB id 1468 ASU2 RS05090 WP 014991734.1	SLWQKNCEKHILIRYLAIPIQENDDTLWLATDDMNNLSACEIFAFIAHK....TIEPVLISSDELKYLNLALSPEQ	93
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NTDB id 1436 BW25113 RS00530 WP 001025145.1	TALCLRYHGVLLDASEEVVHVAVVDAPSHELLDALHFATTK....RIEITCWTRQQMEGHA.....	63
NTDB id 1168 A1552VC RS11070 WP 000957200.1	ANLCQQLGLRELITRYDALPIAKQGNLILLAVSDPTILQAEEFEFRFATGL....QVELALADHRAALQAATIRRLYGR	144
NTDB id 1401 DSB67 RS12665 WP 010643259.1	ASLCQTLGLRLITIRHNALPLNRTSSTLILAVADPTNLQAEDDFRATGL....QVELVLADFRELTAATIRRLYGR	144
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NTDB id 1251 GC085 RS07725 WP 011946523.1	GTIPVNLVNEKLIKRRHAMVPLFSRGTNLYLATDDPSKQASLKETIQFHTGL....NTHAIVVETDKLSALIDNLTAKES	152
NTDB id 1018 ACIAD RS01685 WP 004920473.1	ALFLKDKIDEKLIQKYRIMPLVHRGHVLYVATSNPTNTEAMDAIRFNSKL....KVEPIIVEHDKLERLLSEHFVEETH	152
NTDB id 1058 ABD1 RS01615 WP 001274986.1	SQIPKDLVDQKLILKHRILPLIQRGQILYVATSNPSNIEAIDAIRFNSKL....LVEPVIVEHHKLEKVLGQHFAEES	152
consensus	*****	

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NTDB id 1058 ABD1 RS01615 WP 001274986.1
consensus

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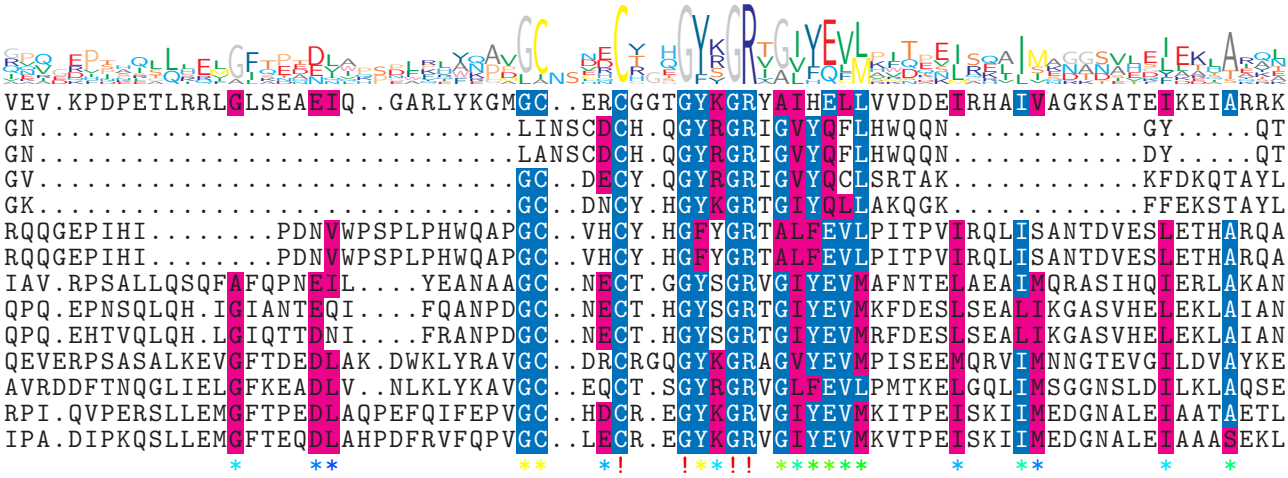
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NTDB	id	1028	TT	RS08215	WP	011173991.1
NTDB	id	1366	NTHI	RS01950	WP	011271972.1
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consensus						



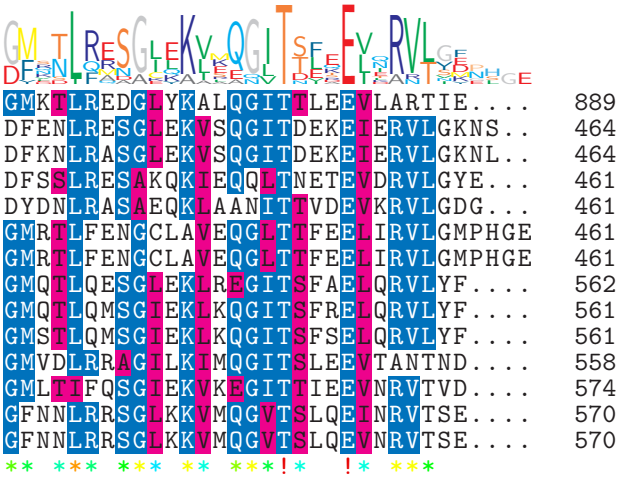
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NTDB	id	1028	TT	RS08215	WP	011173991.1
NTDB	id	1366	NTHI	RS01950	WP	011271972.1
NTDB	id	1355	HI	0298	AAC21962.1	
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NTDB	id	1468	ASU2	RS05090	WP	014991734.1
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NTDB	id	1436	BW25113	RS00530	WP	001025145.1
NTDB	id	1168	A1552VC	RS11070	WP	000957200.1
NTDB	id	1401	DSB67	RS12665	WP	010643259.1
NTDB	id	1293	VP	RS12240	WP	005479695.1
NTDB	id	1110	NGFG	RS09235	WP	003689817.1
NTDB	id	1251	GC085	RS07725	WP	011946523.1
NTDB	id	1018	ACIAD	RS01685	WP	004920473.1
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consensus						



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NTDB	id	1028	TT	RS08215	WP	011173991.1
NTDB	id	1366	NTHI	RS01950	WP	011271972.1
NTDB	id	1355	HI	0298	AAC21962.1	
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NTDB	id	1251	GC085	RS07725	WP	011946523.1
NTDB	id	1018	ACIAD	RS01685	WP	004920473.1
NTDB	id	1058	ABD1	RS01615	WP	001274986.1
consensus						



 non conserved
 similar
 $\geq 50\%$ conserved