

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 1384 A4U84 RS08525 WP 010787006.1		0
NTDB id 1468 ASU2 RS05090 WP 014991734.1		0
NTDB id 30083 SSON53G RS00595 WP 001025185.1		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 GCO85 RS07725 WP 011946523.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
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NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	160
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consensus		

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NTDB id 1028 TT RS08215 WP 011173991.1	EGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLFLESTEGIVPDPSAALLL	240
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consensus		

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NTDB id 1028	TT RS08215 WP 011173991.1	LRSDALRYGAVPIGFQNGEVEVVLSDPRHKEAQAQLLNRPARFYALALPQAWHEELFRRAYPQKNRNLGEVLVQEGKLSREAL	320
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NTDB id 1110	NGFG RS09235 WP 003689817.1		0
NTDB id 1251	GCO85 RS07725 WP 011946523.1		0
NTDB id 1018	ACIAD RS01685 WP 004920473.1		0
NTDB id 1058	ABD1 RS01615 WP 001274986.1		0
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NTDB id 1028	TT RS08215 WP 011173991.1	KEALEVQKGLPRAKPLGEIIVELGLARPEDVEEALQKQRRGGGRVEDTVQSGKLRPEAIAQAVATQLGYPIYVDPEEDPP	400
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NTDB id 1355	HI 0298 AAC21962.1	MTSY....ALLHTQRVTAKNGEVFTI.....SP	24
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NTDB id 1401	DSB67 RS12665 WP 010643259.1	MLTNLSTVLRQAGILTFQSQEESLLEQVKASGISMPAELLSSGFFQSHEITEHLSSVFGLSCTSLNQYEV	69
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NTDB id 1110	NGFG RS09235 WP 003689817.1	MSVGLLRILVQNQVVTVEQAHHY.NESQAGKEVLPMIFSDGVISPKSLAALIARVFSYSILDLRHYPR	68
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NTDB id 1468	ASU2 RS05090 WP 014991734.1	SLWQKNCDEKHILLRYLAIPITQENDDTLWLATDDMNNISACEIFAFIAHK....TIEPVILSSDELKYLLNQLSPEQQS	93
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NTDB id 1168	A1552VC RS11070 WP 000957200.1	ANLCQQLGLRELITRYDALPIAKQGNLLLVAVSDPTLLQAEEFEFRFATGL....QVBLATADHRAQAATRRLYGRSIQ	144
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NTDB id 1251	GCO85 RS07725 WP 011946523.1	GTIPVNLVNEKLIKRRHAMVPLFSRGTNLYLATDDPSKQASLKEIQFHTGL....NTHAIVVETDKLSALIDNLTAKES	152
NTDB id 1018	ACIAD RS01685 WP 004920473.1	ALFLKDKIDEKLIQKYRIMPLVHRGHVLYVATSNPTNTEAMDAIRENSKL....KVEPIIVEHDKLERLSEHFVEETH	152
NTDB id 1058	ABD1 RS01615 WP 001274986.1	SQIPKDLVDQKLLIKHRILPLIQRGQILYVATSNPSNTEAIDAIRENSKL....LVEPVIVEHHKLEKVLGQHFAEES	152
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NTDB	id	1028	TT	RS08215	WP	011173991.1
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NTDB	id	1028	TT	RS08215	WP	011173991.1
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consensus						



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consensus

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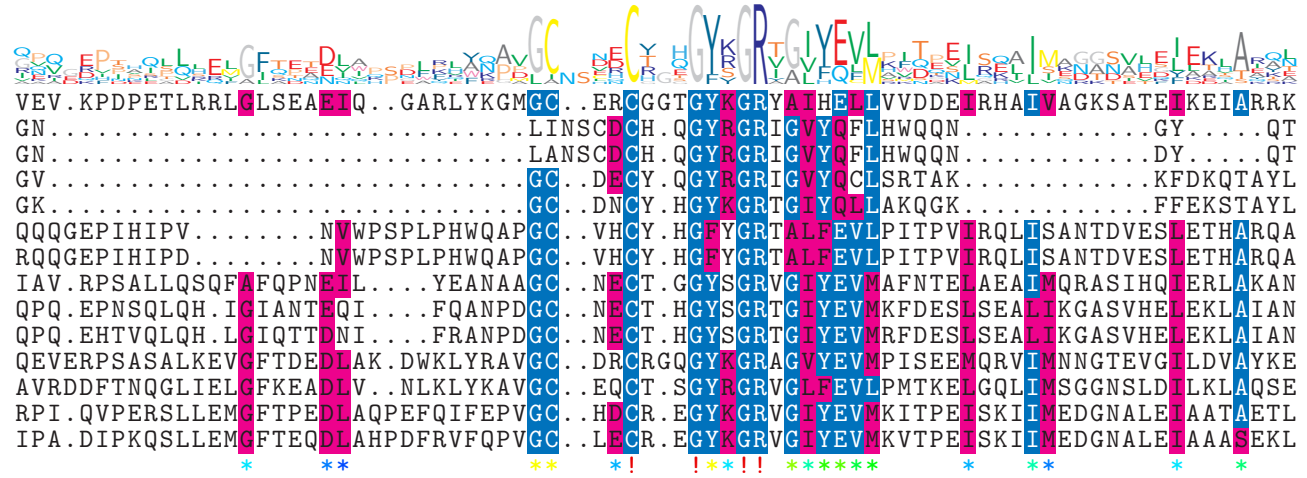


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consensus

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consensus

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464
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461
562
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561
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 non conserved
 similar
 $\geq 50\%$ conserved