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NTDB id 1251 GC085 RS07725 WP 011946523.1		0
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NTDB id 277759 AM467 RS01870 WP 001274992.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	0
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consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	EGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL	0
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consensus		

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consensus		

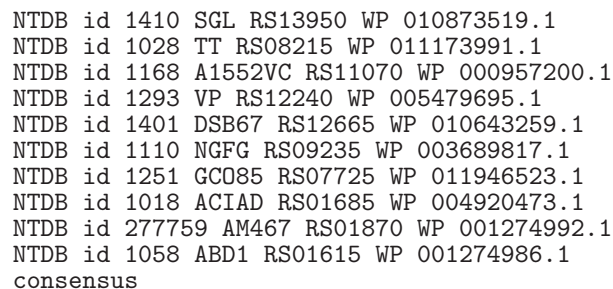
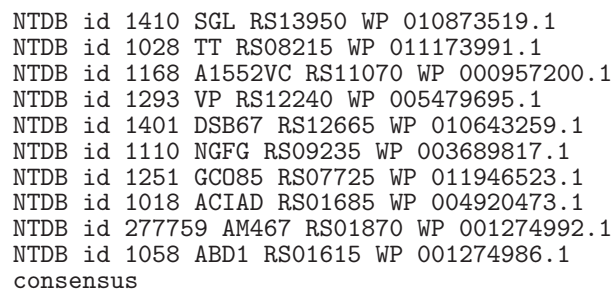
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NTDB id 1251	GCO85 RS07725 WP 011946523.1	...MAL	ATEEYRLQGTGL	LVLEKLLDKTKAIEL	HLKLA AEKISLLQYIV	K..NKILSAEQIALT.....AAQNF	GV	68
NTDB id 1018	ACIAD RS01685 WP 004920473.1	...MSA	FTTPPKFSGFIR	LVEEGYVNAQNM	QQALEKAKKFKQDIV	PYLID..NFSISPLTIAEI.....ISLEF	GE	68
NTDB id 277759	AM467 RS01870 WP 001274992.1	...MSA	LHTSPKFTGFFR	LVEEKHVSAA	TMTALDAAKRAKQDT	VAYLIE..EVHLSPSL	LAET.....ISAEFAEP	68
NTDB id 1058	ABD1 RS01615 WP 001274986.1	...MSA	LHTSPKFTGFFR	LVEEKHVSAA	TMTALDAAKRAKQDT	VAYLIE..EVHLSPSL	LAET.....ISAEFAEP	68
consensus		*	*****	*!*****	*****	*****	*****	

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NTDB	id	1168	A1552VC	RS11070	WP	000957200.1	E	T	D	L	S	R	Y	D	Y	A	N	.	.	.	.	.	.	L	C	Q	Q	L	G	L	R	E	L	I	T	R	Y	D	A	L	P	I	A	K	Q	.	.	G	N	L	L	L	A	V	S	D	P	T	L	L	Q	A	E	E	E	F	R	F	A	T	G	.	.	.	.	.	.	L	Q	V	E	L	A	L	A	D	127	
NTDB	id	1293	VP	RS12240	WP	005479695.1	Q	P	E	L	S	Q	Y	E	Y	A	S	.	.	.	.	.	.	L	C	Q	Q	L	G	L	R	E	L	I	T	R	H	N	A	L	P	L	H	R	T	.	.	P	S	T	L	L	A	V	A	D	P	T	N	Q	Q	A	E	D	D	F	R	F	A	T	G	.	.	.	.	.	.	L	Q	V	E	L	V	L	A	D	127	
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NTDB	id	277759	AM467	RS01870	WP	001274992.1	Y	F	D	L	D	V	Y	D	T	S	Q	.	.	.	.	.	.	I	P	K	D	L	V	D	Q	K	L	I	L	K	H	R	I	L	P	L	I	Q	R	.	.	G	Q	I	L	Y	V	A	T	S	N	P	S																													

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consensus

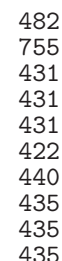
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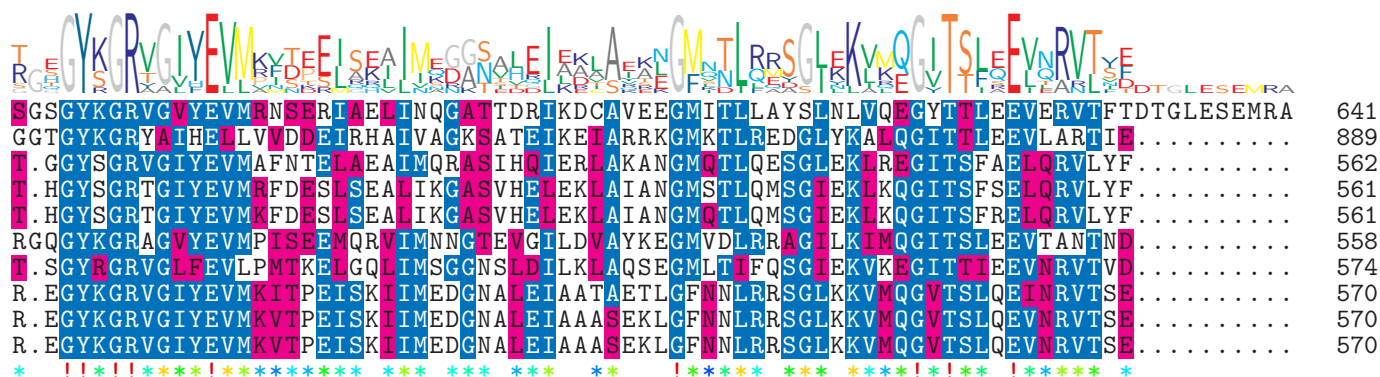
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consensus



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consensus



logo

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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved