

logo

NTDB id 1410 SGL RS13950 WP 010873519.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	0
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NTDB id 1168 A1552VC RS11070 WP 000957200.1		0
NTDB id 1293 VP RS12240 WP 005479695.1		0
NTDB id 1401 DSB67 RS12665 WP 010643259.1		0
NTDB id 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 326616 EG365 RS09935 WP 001274990.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	0
NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG	160
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consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	EGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL	0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
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consensus		

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NTDB id 1110	NGFG RS09235 WP 003689817.1		0
NTDB id 1251	GC085 RS07725 WP 011946523.1		0
NTDB id 326616	EG365 RS09935 WP 001274990.1		0
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NTDB id 1018	ACIAD RS01685 WP 004920473.1		0
consensus			

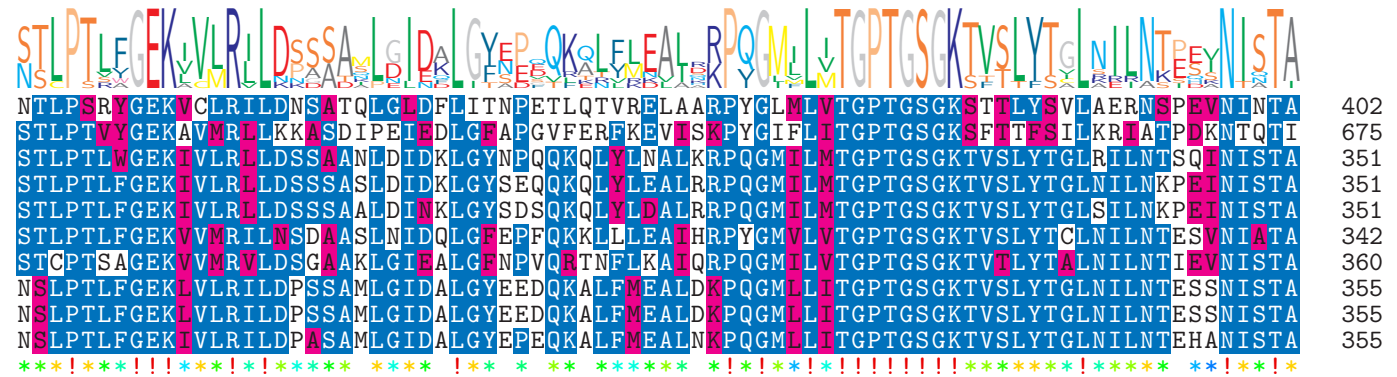
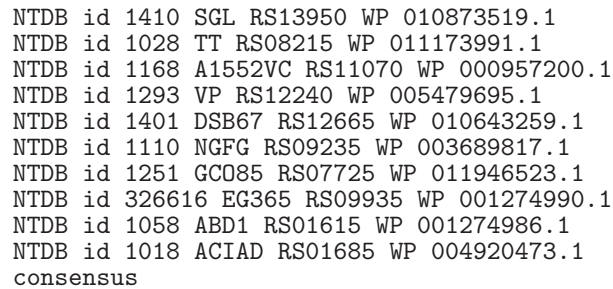
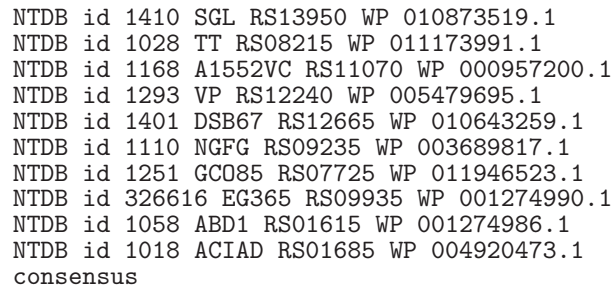
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NTDB id 1028	TT RS08215 WP 011173991.1	KEAIEVQKGGLPFAKPLGETLVLELGLARPEDVEEALQKQRGGGRLEDTLVQ..SGKLRPEALACA.....VATQLGYGP	391
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NTDB id 1110	NGFG RS09235 WP 003689817.1	MSVGLLRLLVQNQVVTVQEAHHYY.NESQAGKEVLPMLFS..DGVISPKS LAAL.....IARVFSYS	59
NTDB id 1251	GCO85 RS07725 WP 011946523.1	...MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLA AA EKI SLLQYIVK..NKILSAEQIALT.....AAQNFQVP	68
NTDB id 326616	EQ365 RS09935 WP 001274990.1	...MSALHTSPKFTGFFRRLVEEKHVSAATMQTALDAAKRAKQDTVAYLIE..EVHLSPSLLAET.....ISAEFAEP	68
NTDB id 1058	ABD1 RS01615 WP 001274986.1	...MSALHTSPKFTGFFRRLVEEKHVSAATMQTALDAAKRAKQDTVAYLIE..EVHLSPSLLAET.....ISAEFAEP	68
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consensus		* * * * *	

NTDB id	1410	SGL RS13950 WP 010873519.1	SVDPEVAPIATQMAELIGRFFPLDTCRRFKFLPLAQQEGDPSPSVLVAMVPDPNLAQAQDELNRILRVKGF...ELRRLVVT	164
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NTDB id	1293	VP SR12240 WP 005479695.1	QP ELSQYEYAS.....LCQQQLGLRELITRNHALP LHT...PSTLL LAVADPTNQQAEDDFRFATG.....LQVELVLAD	127
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NTDB id	1110	NGFG RS09235 WP 003689817.1	I LDLRHYPRHR.....VLMGVLT EEQMVEFHCVPV FRR...GDKVFFAVSDPT QMPQIQKTVS AAG.....IAVEL VIVE	126
NTDB id	1251	GCO85 RS07725 WP 011946523.1	MLDINCIDVGT.....IPVNLVN EKLIK RHAMV PLFSR...GTNLYLATDDPS KQS LKE IQF HTG.....LNTHAI VE	135
NTDB id	326616	EQ365 RS09935 WP 001274990.1	YFDLDVYDT SQ.....IPKD LV DQKL I LKHRI L P LIQR...GQ IL Y VATSN PSNI EA IDAIR FN SK.....LLVE PVIVE	135
NTDB id	1058	ABD1 RS01615 WP 001274986.1	YFDLDVYDT SQ.....IPKD LV DQKL I LKHRI L P LIQR...GQ IL Y VATSN PSNI EA IDAIR FN SK.....LLVE PVIVE	135
NTDB id	1018	ACIAD RS01685 WP 004920473.1	L LD LGVDEPAL.....FLKDKIDEKLIQKYRI MP LV HR...GHV LY VATSN PTNIE AMD AI RFN SK.....LKVE PIIVE	135
consensus			* * * * * * * * * * ! * * * * * ! * * * * *	

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NTDB id 1058 ABD1 RS01615 WP 001274986.1
NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

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logo

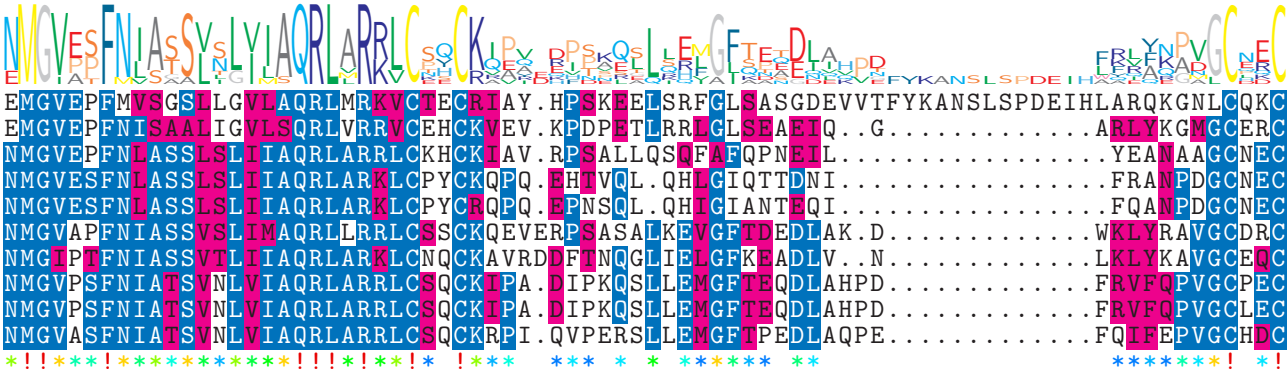
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consensus		



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NTDB id	1410	SGL RS13950 WP 010873519.1
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consensus		



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NTDB id	1410	SGL RS13950 WP 010873519.1
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consensus		



641
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logo

NTDB id 1410 SGL RS13950 WP 010873519.1	KRKSALTCRTCAAELQQEWLDCPYCMTSRFS	672
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NTDB id 1058 ABD1 RS01615 WP 001274986.1		570
NTDB id 1018 ACIAD RS01685 WP 004920473.1		570
consensus		

-  non conserved
-  similar
-  ≥ 50% conserved