

logo

NTDB id 1306 DR RS10060 WP 027479822.1	MSXLIIGDRRLGAALLDQGYLDLQLRALERHSEVGGRLAELVLYDSGGEERRIASLIETSGIPLVLAHVETSPKX	79
NTDB id 1028 TT RS08215 WP 011173991.1	.MALSIGDRRRLGAILLDQGYLGDNLDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAAL	80
NTDB id 1366 NTHI RS01950 WP 011271972.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	0
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NTDB id 1293 VP RS12240 WP 005479695.1		0
NTDB id 1110 NGFG RS09235 WP 003689817.1		0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
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NTDB id 1306 DR RS10060 WP 027479822.1	RSIRPQTALNLQAFPPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGA	157
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NTDB id 1306 DR RS10060 WP 027479822.1	SDAGRTGGKLGGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEEQEGTTFLPNPSGFGHPGEEVL	237
NTDB id 1028 TT RS08215 WP 011173991.1	SGEGQKDLKLGELLQKQWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVDPDSAA	237
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NTDB id 1306 DR RS10060 WP 027479822.1	GSMLRADALRLLAVPVDETEQGVTVLTSDPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRGLGEQLVQEGTLSR	317
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consensus					

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NTDB id 1293 VP RS12240 WP 005479695.1	YEYASLCQQLGLRELITRHNALPLHRTPTST	LLLAVADPTNQAEDDFRFATG.....LQVEL	VLADFRELTAAIRRLYGR	141
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	YDTSQIPKDLVDQKLLILKHRIPLIQRGQIL	YVATSNPSNIEAIDAIRFNSK.....LLVEP	VIVEHHKLEKVLGQHFAE	149
consensus				

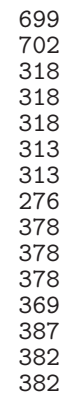
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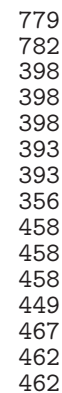
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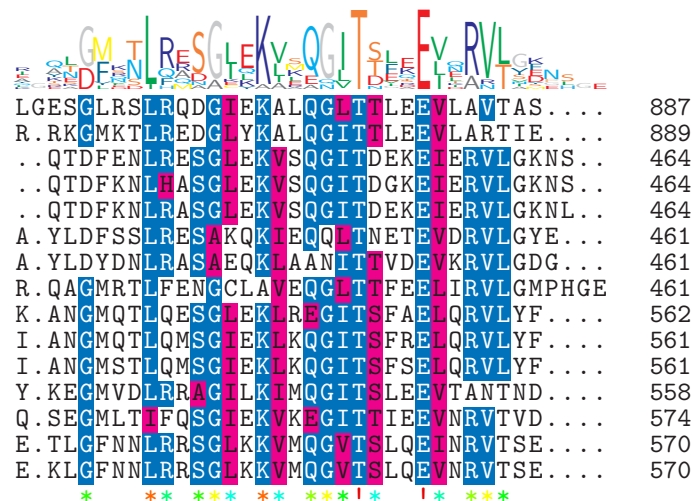
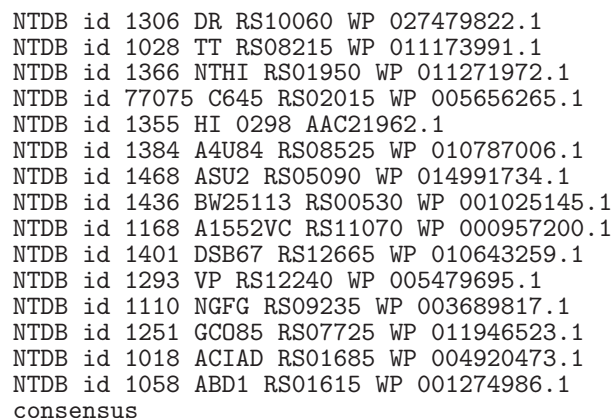
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consensus

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- ☐ non conserved
- ☒ similar
- ☒ > 50% conserved