

logo

NTDB id 1028	TT	RS08215	WP	011173991.1	MSXLSIGDRLGAALLDAGLLTDEELQRALERHREVGGS	80
NTDB id 1306	DR	RS10060	WP	027479822.1	LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	79
NTDB id 1384	A4U84	RS08525	WP	010787006.1	.MALSIGDRLGAILLDQGYLGDNLDQRALERHSEVGGR	0
NTDB id 1355	HI	0298	AAC21962.1		LADVLIDSGMVGEKRIARAIEEALGIPLVNL	0
NTDB id 1366	NTHI	RS01950	WP	011271972.1	LAVQPDPAAL	0
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 1058	ABD1	RS01615	WP	001274986.1		0
NTDB id 1251	GC085	RS07725	WP	011946523.1		0
consensus						0

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NTDB id 1028	TT	RS08215	WP	011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLD	158
NTDB id 1306	DR	RS10060	WP	027479822.1	TLSEEEVDTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	157
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NTDB id 1251	GC085	RS07725	WP	011946523.1		0
consensus						0

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NTDB id 1028	TT	RS08215	WP	011173991.1	SGEGQKDLKLGELLQKGWISREALEEALVEQ	237
NTDB id 1306	DR	RS10060	WP	027479822.1	KTGDLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVDPDSAA	237
NTDB id 1384	A4U84	RS08525	WP	010787006.1	SDAGRTGGKLGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	0
NTDB id 1355	HI	0298	AAC21962.1			0
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NTDB id 1251	GC085	RS07725	WP	011946523.1		0
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[illegible]

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EALKEALEVQKGLPRAKPIGELVELGLARPEDVEEAIQKQRRCGRREDTLVQSCKLRPEALAQAVATQLGYPYVDPEE 397
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MSQYTI.....CEQQTERIFDIS..... 18
MTSYAIIHTQRVTAKNGEVFTIS..... 23
MTSYALLHTQRTVAQNGEIFTIS..... 23
MLTNIVAILRQAELISATQEAVVTQVSASCTSVPEALIELSIFHAQELTEQLSHIFGPETDISR 66
MHSNISTILRQKGLLIFSQUEESLIEQVKASCISMPEALLSSGFFTSSELAHELSSIFGLSQPELSQ 66
MLTNISTVL RQAGILTF SQUEESLIEQVKASCISMPEALLSSCFFQSHLTHELSSVFGLSCTSINQ 66
MSALHTSPKFTGGFRRLVEEKHVS AATMQTADAAKRAKQDTVAYLIEEVHLSPSLLAETISAFAEPYFDLDV 74
MALATEEYRLQGILGLELVLEKLLDKTKAIELHKLA AEKISILQYIVKNKILSAEQALTAAQNFGVPMLDINC 74
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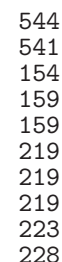
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[illegible]

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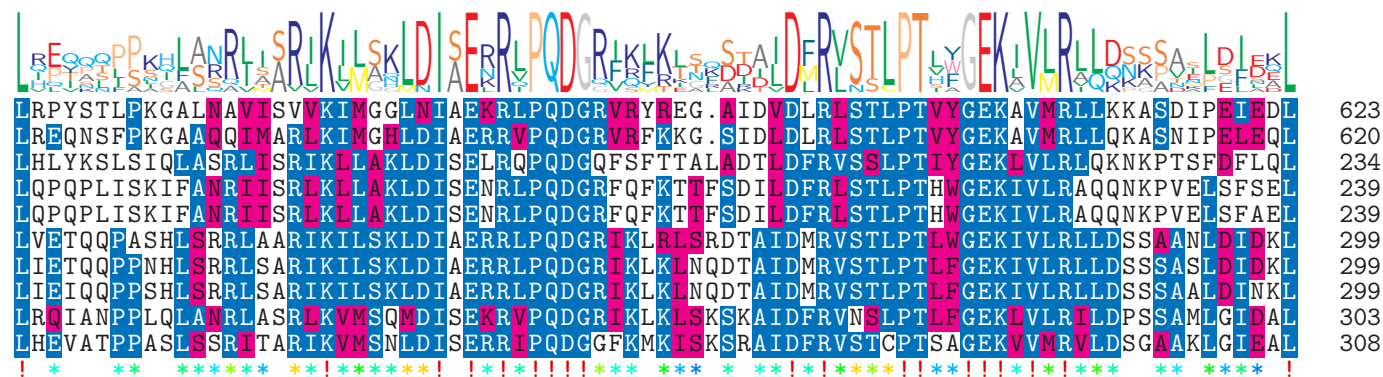
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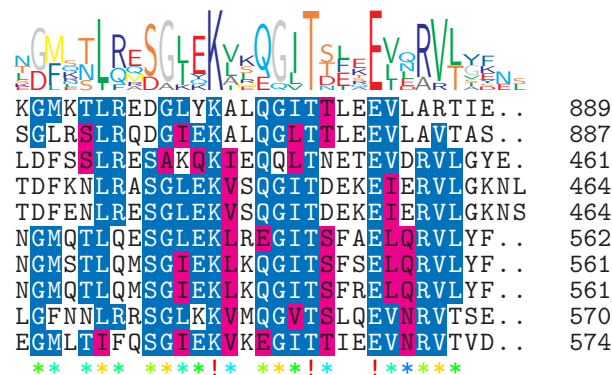
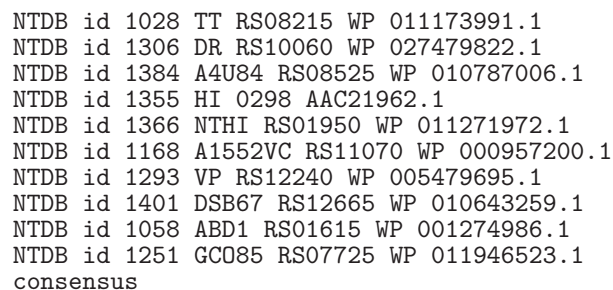
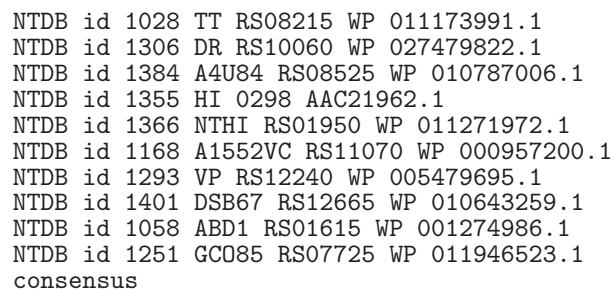
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- X non conserved
- X similar
- X > 50% conserved