

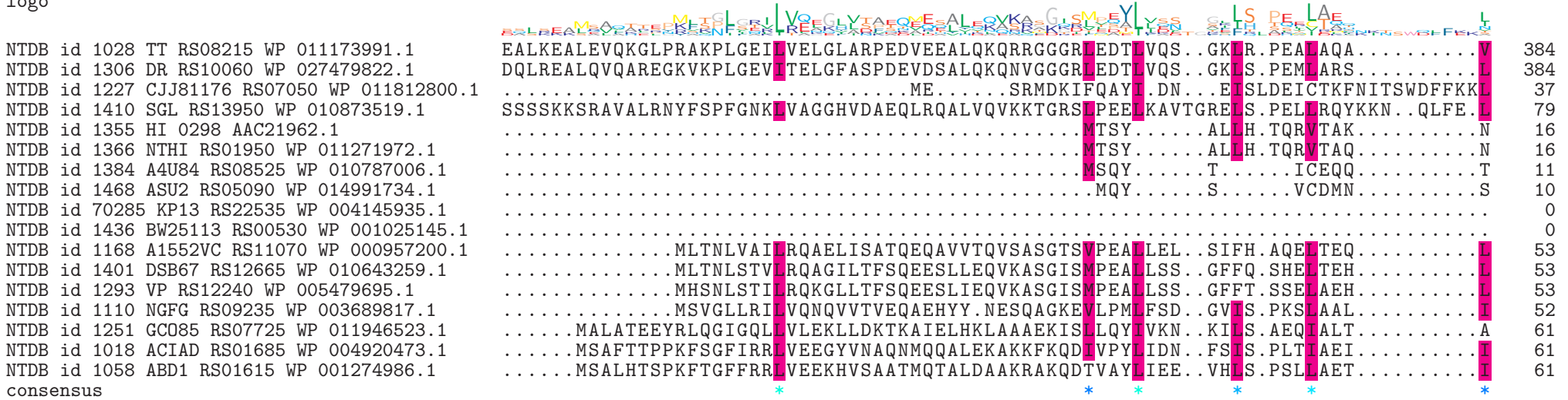
logo

		MSXLRITCDRLGAAALLDQGLDRLQRALERHREVGGLAEVLYDGLQYERRIASXIEERFGIPLVQLHRVENSPPKXK	
NTDB id 1028	TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGLAEVLYDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVK	80
NTDB id 1306	DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNLDLQRALERHSEVGGRLADVLDIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAAL	79
NTDB id 1227	CJJ81176 RS07050 WP 011812800.1	0
NTDB id 1410	SGL RS13950 WP 010873519.1	0
NTDB id 1355	HI 0298 AAC21962.1	0
NTDB id 1366	NTHI RS01950 WP 011271972.1	0
NTDB id 1384	A4U84 RS08525 WP 010787006.1	0
NTDB id 1468	ASU2 RS05090 WP 014991734.1	0
NTDB id 70285	KP13 RS22535 WP 004145935.1	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
consensus			

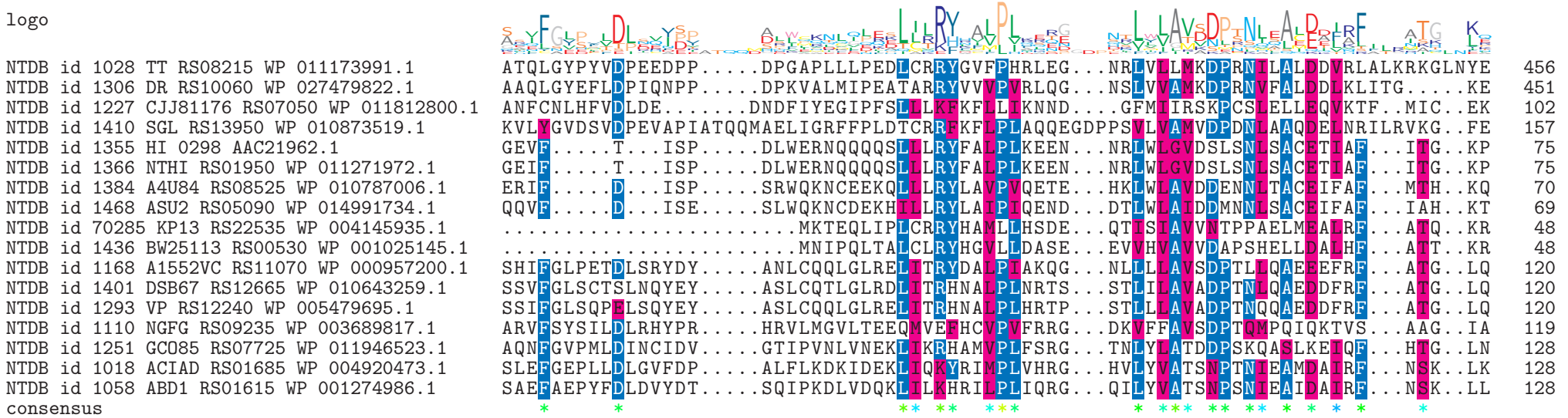
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		REYKAKELKAIPFALDEEAGVVRVAFNLPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLEIVVPSGA	
NTDB id 1028	TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLEIVVPSGA	158
NTDB id 1306	DR RS10060 WP 027479822.1	RSIRPQTALNLQAFPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGA	157
NTDB id 1227	CJJ81176 RS07050 WP 011812800.1	0
NTDB id 1410	SGL RS13950 WP 010873519.1	0
NTDB id 1355	HI 0298 AAC21962.1	0
NTDB id 1366	NTHI RS01950 WP 011271972.1	0
NTDB id 1384	A4U84 RS08525 WP 010787006.1	0
NTDB id 1468	ASU2 RS05090 WP 014991734.1	0
NTDB id 70285	KP13 RS22535 WP 004145935.1	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
consensus			

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 consensus



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NTDB id 1028 TT RS08215 WP 011173991.1
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 NTDB id 1018 ACIAD RS01685 WP 004920473.1
 NTDB id 1058 ABD1 RS01615 WP 001274986.1
 consensus



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NTDB id 1028 TT RS08215 WP 011173991.1	889
NTDB id 1306 DR RS10060 WP 027479822.1	887
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NTDB id 1468 ASU2 RS05090 WP 014991734.1	G.....	461
NTDB id 70285 KP13 RS22535 WP 004145935.1	PHGD.....	461
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NTDB id 1401 DSB67 RS12665 WP 010643259.1	561
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NTDB id 1110 NGFG RS09235 WP 003689817.1	558
NTDB id 1251 GC085 RS07725 WP 011946523.1	574
NTDB id 1018 ACIAD RS01685 WP 004920473.1	570
NTDB id 1058 ABD1 RS01615 WP 001274986.1	570
consensus		

DTGLESEMRAKRKKSALTCRTCAAELQQEWLDCPYCMTSRFS

- non conserved
- similar
- ≥ 50% conserved