

logo

NTDB id 1028 TT RS08215 WP 011173991.1	MSXLIIGDRLGAALLDRLDNLQRALERHREVGGLAELVLYDSQYSEERIAELIEELGILPLVLLHVVLPSPKXK	80
NTDB id 1306 DR RS10060 WP 027479822.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV	79
NTDB id 1110 NGFG RS09235 WP 003689817.1	.MALSIGDRRLGAILLDQGYLGNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAL	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 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 27169 SPUTW3181 RS02685 WP 011787903.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
consensus		

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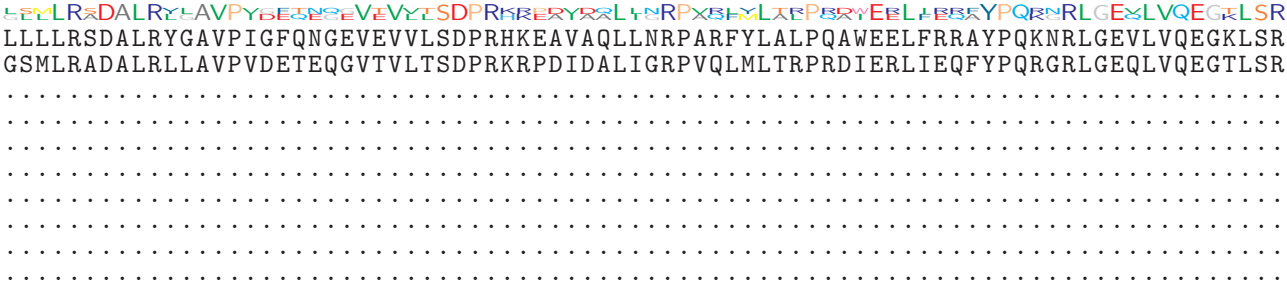
NTDB id 1028 TT RS08215 WP 011173991.1	RLTPEKELAKELRAIPFALDEEAGVVRVAFLNPLDTLSLEEVEDLTGLVVPEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
NTDB id 1306 DR RS10060 WP 027479822.1	RSIRPQTALNLQAFPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGA	157
NTDB id 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1168 A1552VC RS11070 WP 000957200.1		0
NTDB id 1293 VP RS12240 WP 005479695.1		0
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NTDB id 27169 SPUTW3181 RS02685 WP 011787903.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
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NTDB id 1028 TT RS08215 WP 011173991.1	SGEGQKDLKLGELLLQKGWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAA	237
NTDB id 1306 DR RS10060 WP 027479822.1	SDAGRTGGKLGRLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	237
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NTDB id 27169 SPUTW3181 RS02685 WP 011787903.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
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NTDB id	1028	TT	RS08215	WP	011173991.1
NTDB id	1306	DR	RS10060	WP	027479822.1
NTDB id	1110	NGFG	RS09235	WP	003689817.1
NTDB id	1168	A1552VC	RS11070	WP	000957200.1
NTDB id	1293	VP	RS12240	WP	005479695.1
NTDB id	1401	DSB67	RS12665	WP	010643259.1
NTDB id	1251	GC085	RS07725	WP	011946523.1
NTDB id	27169	SPUTW3181	RS02685	WP	011787903.1
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