

logo

		MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1028	TT RS08215 WP 011173991.1		
NTDB id 1306	DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAAL	79
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
NTDB id 1401	DSB67 RS12665 WP 010643259.1	0
NTDB id 1251	GCO85 RS07725 WP 011946523.1	0
NTDB id 317044	D6Z43 RS15025 WP 120652959.1	0
NTDB id 1058	ABD1 RS01615 WP 001274986.1	0
NTDB id 1018	ACIAD RS01685 WP 004920473.1	0
consensus			

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		ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDPLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLLEIVVPSGA	158
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NTDB id 1251	GCO85 RS07725 WP 011946523.1	0
NTDB id 317044	D6Z43 RS15025 WP 120652959.1	0
NTDB id 1058	ABD1 RS01615 WP 001274986.1	0
NTDB id 1018	ACIAD RS01685 WP 004920473.1	0
consensus			

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		SGEGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVDPDPSAA	237
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NTDB id 1306	DR RS10060 WP 027479822.1	SDAGRTGGKLGRLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	237
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NTDB id 1401	DSB67 RS12665 WP 010643259.1	0
NTDB id 1251	GCO85 RS07725 WP 011946523.1	0
NTDB id 317044	D6Z43 RS15025 WP 120652959.1	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 317044 D6Z43 RS15025 WP 120652959.1
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 NTDB id 1018 ACIAD RS01685 WP 004920473.1
 consensus



543
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208
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218
227
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NTDB id 1028 TT RS08215 WP 011173991.1
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 NTDB id 1058 ABD1 RS01615 WP 001274986.1
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 consensus



621
618
288
297
297
297
306
301
301
301

logo

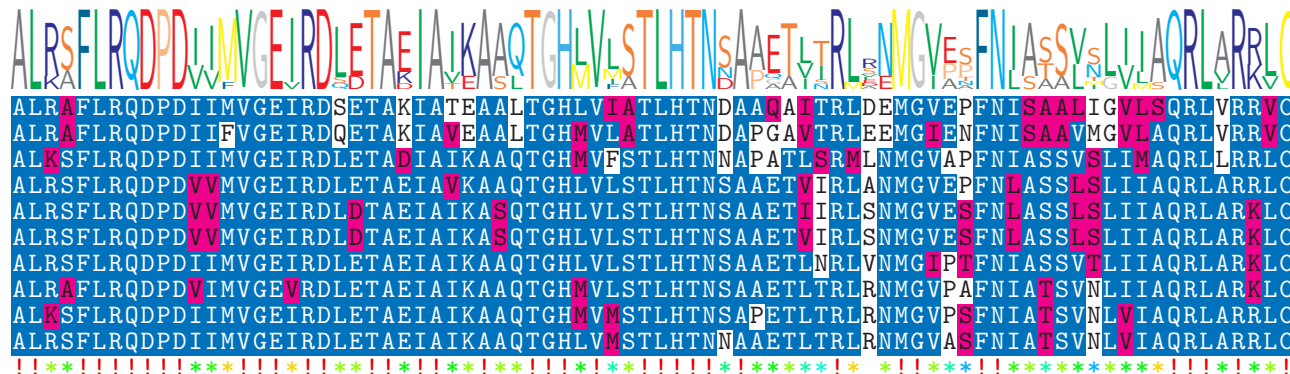
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 consensus



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698
368
377
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377
386
381
381
381

logo

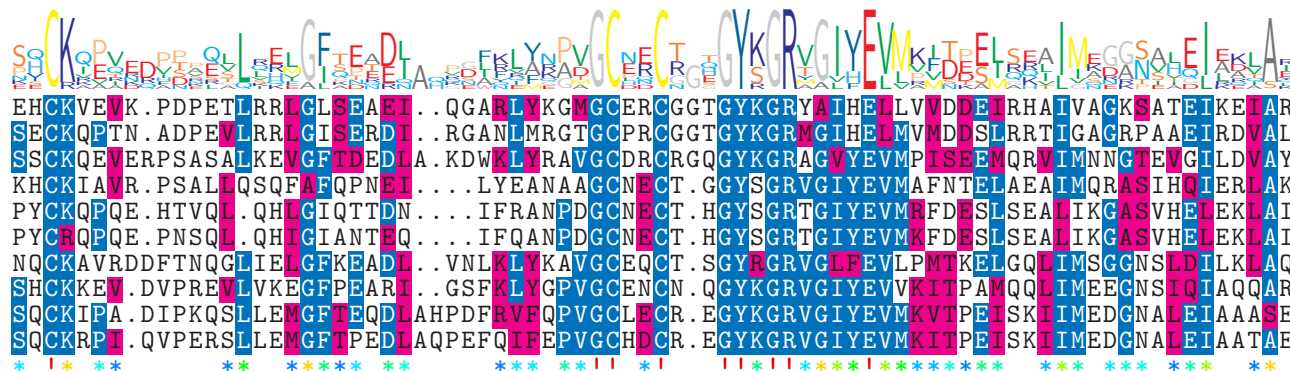
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 NTDB id 1058 ABD1 RS01615 WP 001274986.1
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781
778
448
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466
461
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461

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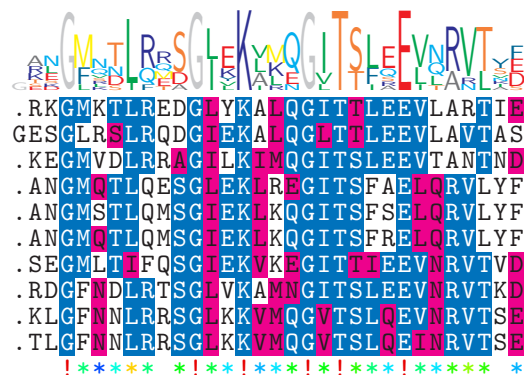
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 NTDB id 1018 ACIAD RS01685 WP 004920473.1



889
887
558
562
561
561
574
568
570
570

non conserved
 similar
 ≥ 50% conserved