

logo

		MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1028	TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1306	DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAAL	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 1072189	ACI3KK RS19210 WP 405118997.1	0
NTDB id 1058	ABD1 RS01615 WP 001274986.1	0
NTDB id 1018	ACIAD RS01685 WP 004920473.1	0
consensus			

logo

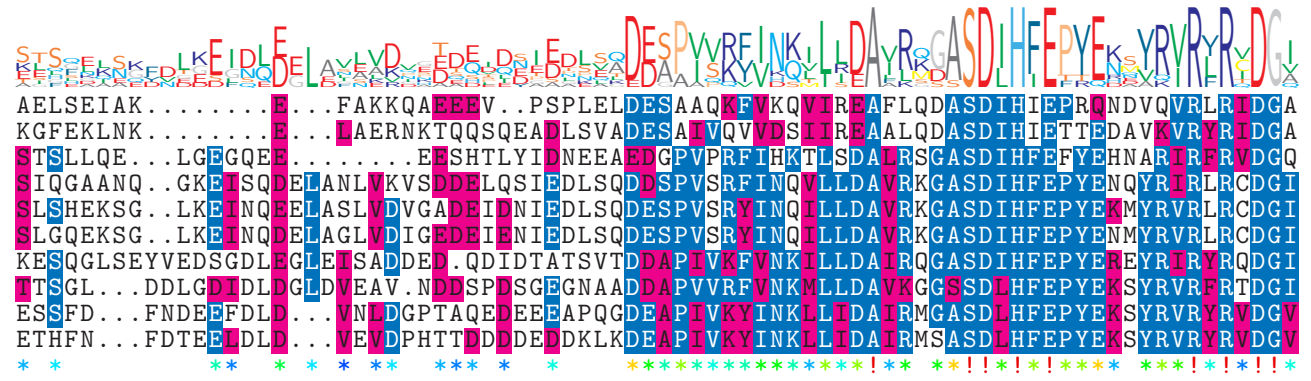
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NTDB id 1251	GCO85 RS07725 WP 011946523.1	0
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NTDB id 1058	ABD1 RS01615 WP 001274986.1	0
NTDB id 1018	ACIAD RS01685 WP 004920473.1	0
consensus			

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		SGEGQKDLKLGELLQKQWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGFLESTEGIVPDPSAA	237
NTDB id 1028	TT RS08215 WP 011173991.1	SGEGQKDLKLGELLQKQWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGFLESTEGIVPDPSAA	237
NTDB id 1306	DR RS10060 WP 027479822.1	SDAGRTGGKLGRLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	237
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consensus			

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 NTDB id 1018 ACIAD RS01685 WP 004920473.1
 consensus



544
541
209
219
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219
228
222
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logo

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



622
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logo

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



702
699
369
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387
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382

logo

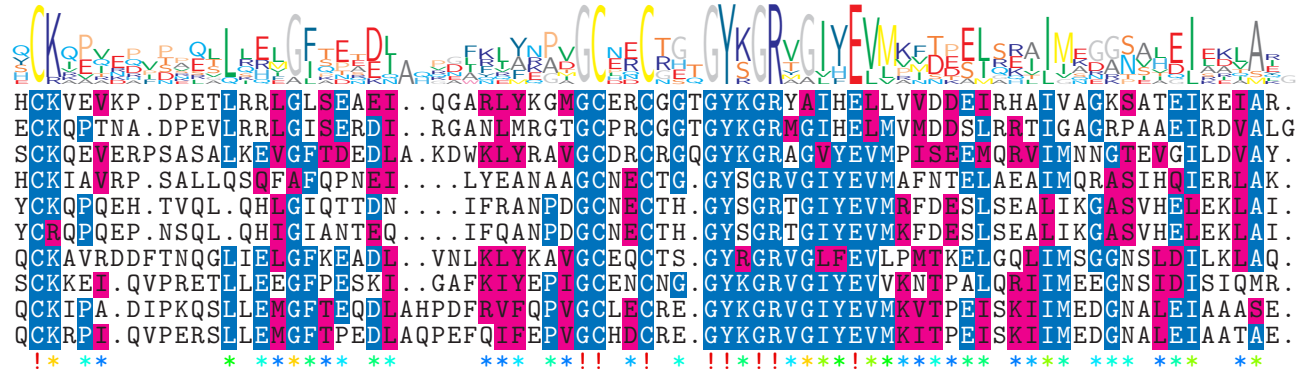
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 NTDB id 1018 ACIAD RS01685 WP 004920473.1
 consensus



782
779
449
458
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458
467
461
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462

logo

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 NTDB id 1018 ACIAD RS01685 WP 004920473.1
 consensus



858
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527
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543
536
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539

logo

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 NTDB id 1018 ACIAD RS01685 WP 004920473.1
 consensus



889
887
558
562
561
561
574
567
570
570

non conserved
 similar
 ≥ 50% conserved