

logo

		MSXLRIGDRRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1028	TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1306	DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAAL	79
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 1110	NGFG RS09235 WP 003689817.1	0
NTDB id 1129795	AACL56 RS23135 WP 339092131.1	0
NTDB id 1058	ABD1 RS01615 WP 001274986.1	0
NTDB id 1018	ACIAD RS01685 WP 004920473.1	0
consensus			

logo

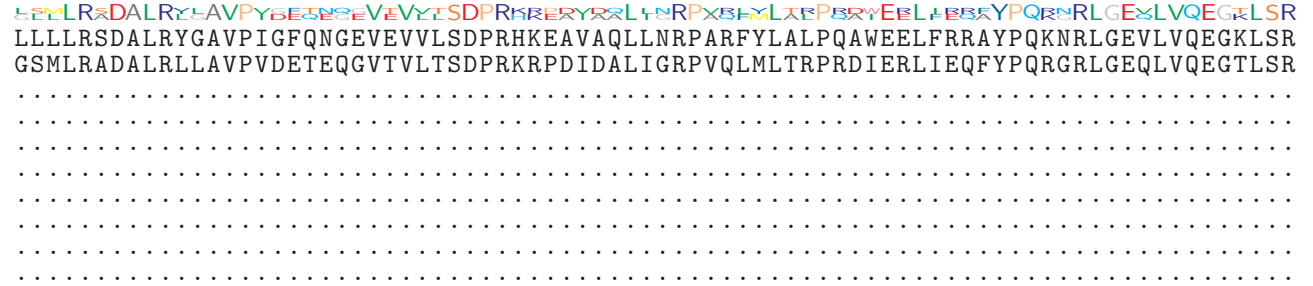
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consensus			

logo

		SGEGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGFLESTEGIVPDPSAA	237
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NTDB id 1018	ACIAD RS01685 WP 004920473.1	0
consensus			

logo

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 NTDB id 1058 ABD1 RS01615 WP 001274986.1
 NTDB id 1018 ACIAD RS01685 WP 004920473.1
 consensus



317
317
0
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logo

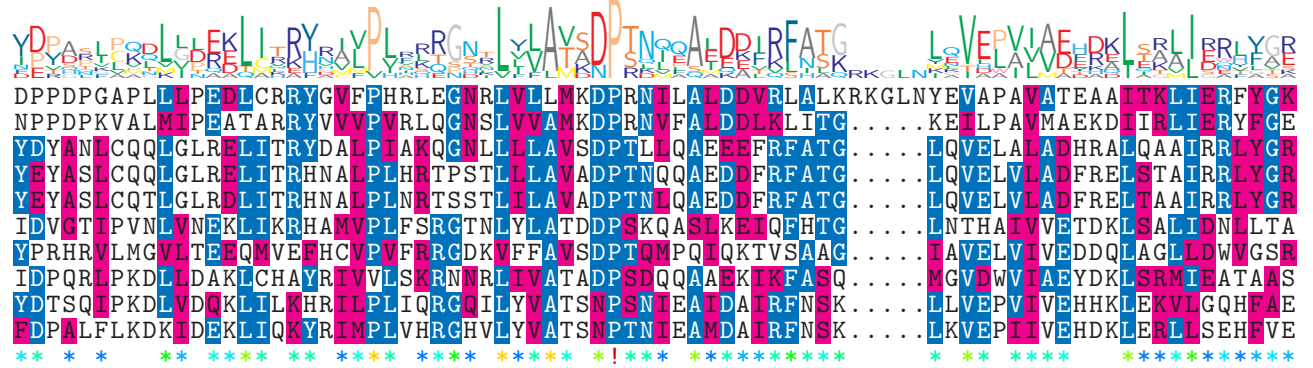
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 NTDB id 1129795 AACL56 RS23135 WP 339092131.1
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 NTDB id 1018 ACIAD RS01685 WP 004920473.1
 consensus



397
397
66
66
74
65
80
74
74

logo

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 consensus



477
472
141
141
141
149
140
155
149
149

logo

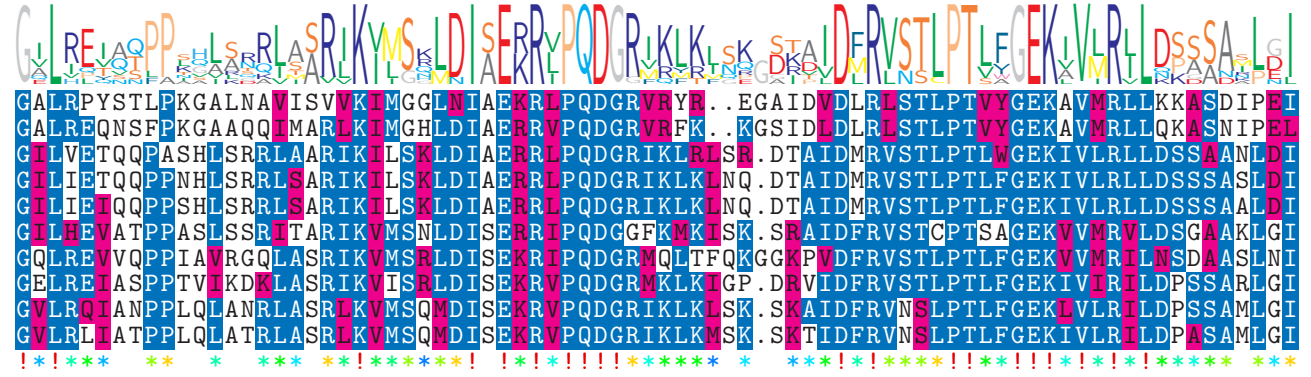
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 NTDB id 1129795 AACL56 RS23135 WP 339092131.1
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 consensus



542
539
217
217
226
207
229
221
221

logo

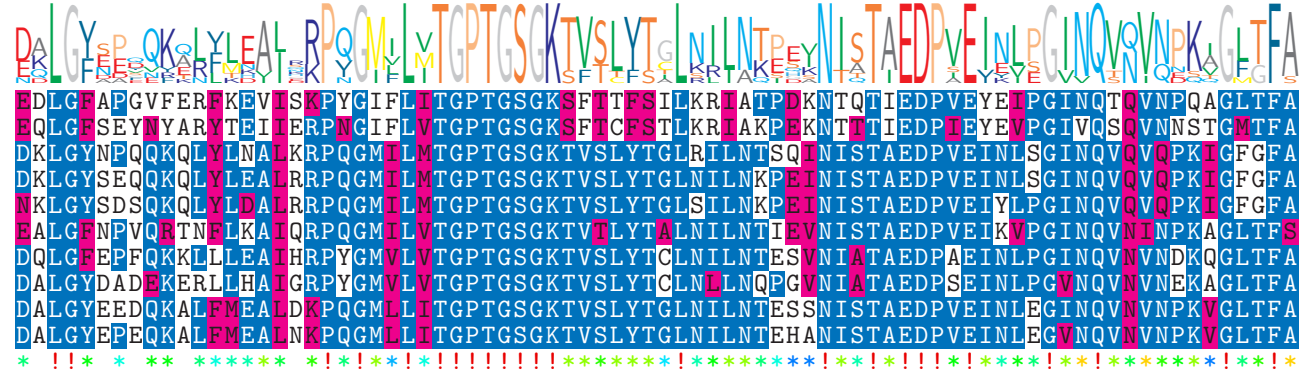
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 NTDB id 1058 ABD1 RS01615 WP 001274986.1
 NTDB id 1018 ACIAD RS01685 WP 004920473.1
 consensus



620
617
296
296
305
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308
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logo

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 NTDB id 1058 ABD1 RS01615 WP 001274986.1
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 consensus



700
697
376
376
376
385
367
388
380
380

logo

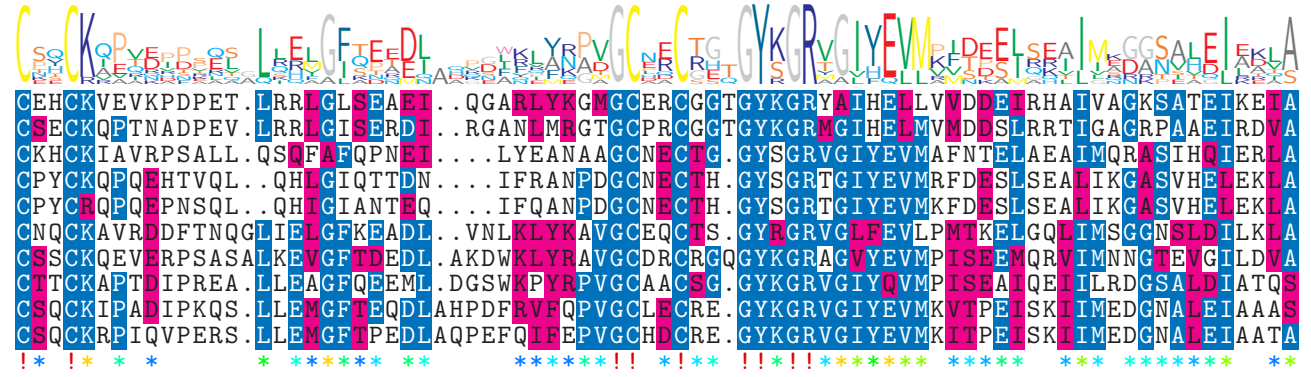
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 NTDB id 1018 ACIAD RS01685 WP 004920473.1
 consensus



780
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456
456
465
447
468
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460

logo

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 consensus



857
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529
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545
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538

logo

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 consensus



889
887
562
561
561
574
558
577
570
570

- non conserved
- similar
- ≥ 50% conserved