

logo

		MSXLIIGDRRLGALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVK	
NTDB id 1028	TT RS08215 WP 011173991.1	MSVLTIGDKRLGAAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVK	80
NTDB id 1306	DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNLDLQRALERHSEVGGRLADVLDIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAAL	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 1128225	WDLP6 RS05850 WP 162591591.1	0
NTDB id 1058	ABD1 RS01615 WP 001274986.1	0
NTDB id 1018	ACIAD RS01685 WP 004920473.1	0
consensus			

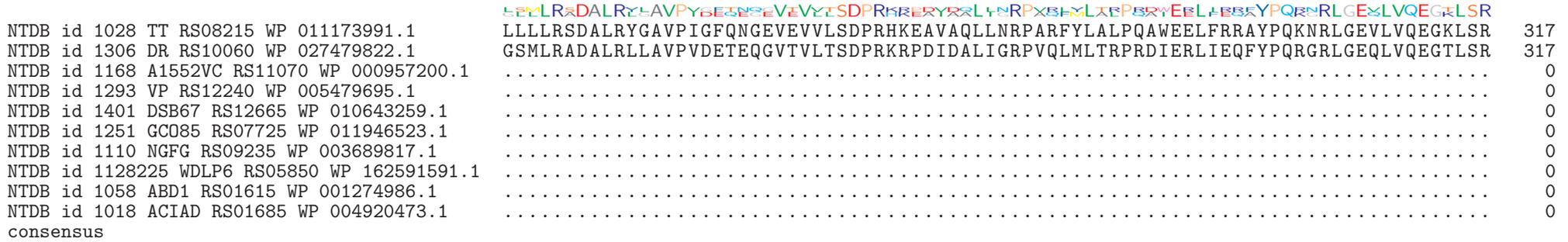
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NTDB id 1306	DR RS10060 WP 027479822.1	RSIRPQTALNLQAFPPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGA	157
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consensus			

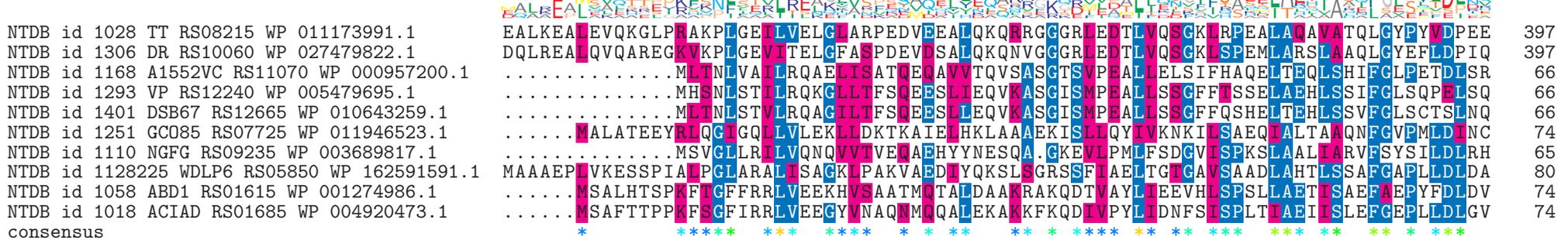
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		SAGSRSKLCERLTHGYIIRKLYALYQRTGELCRLTYSRKTFEERLYEXLAEQEGTFLFLESTEGIVPDPSSAA	
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NTDB id 1306	DR RS10060 WP 027479822.1	SDAGRTGGKLGGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTFLFNPSPGFHPGEEVL	237
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consensus			

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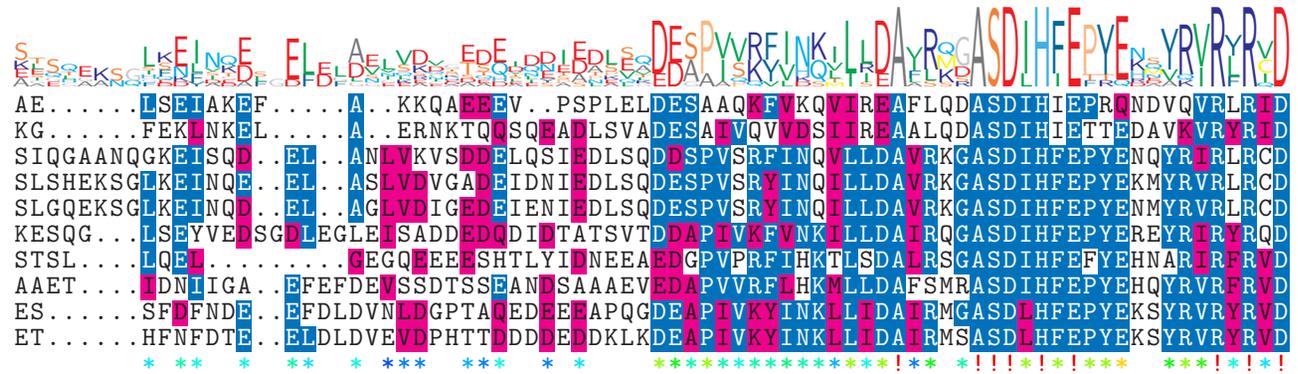


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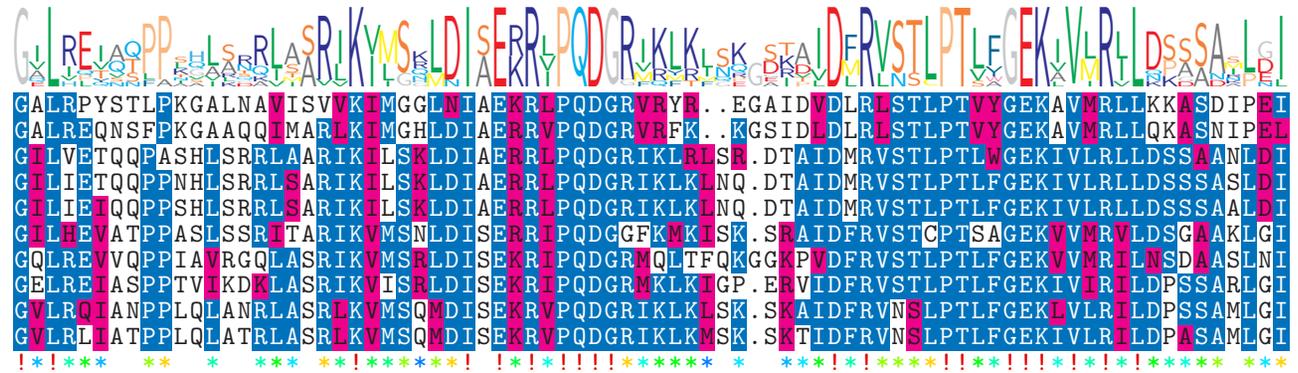
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 consensus



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logo

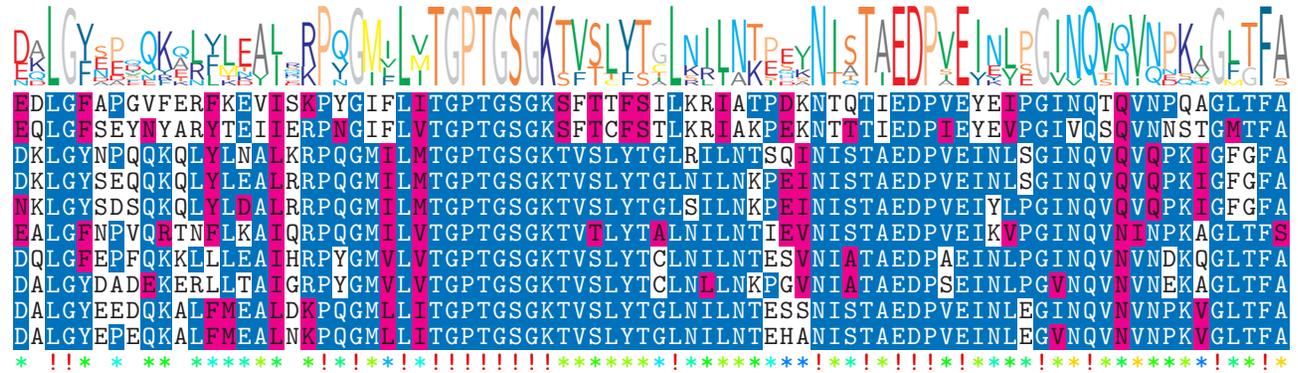
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 consensus



620
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logo

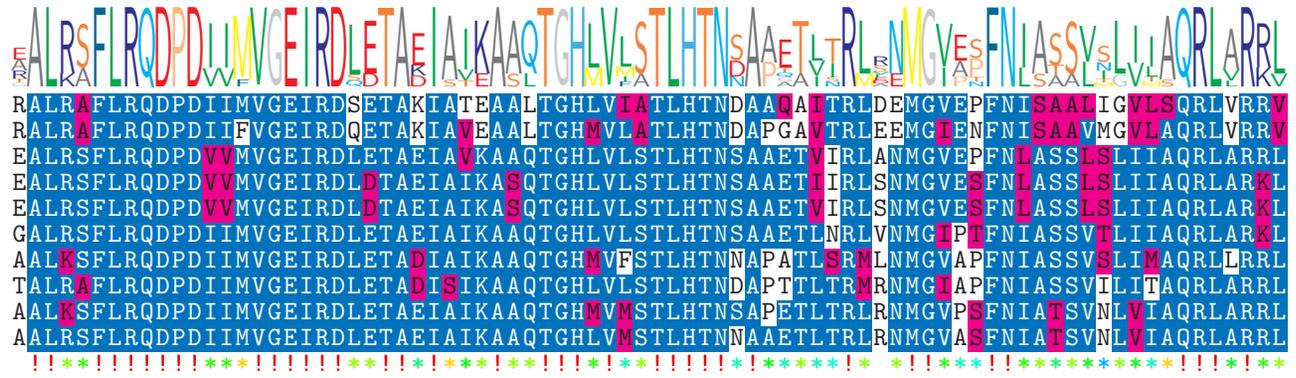
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 consensus



700
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376
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logo

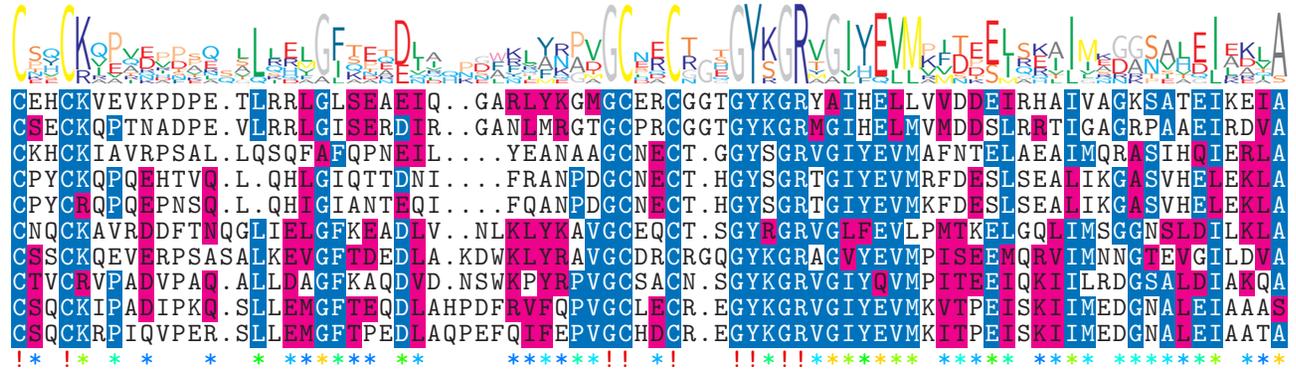
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 consensus



780
777
456
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456
465
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468
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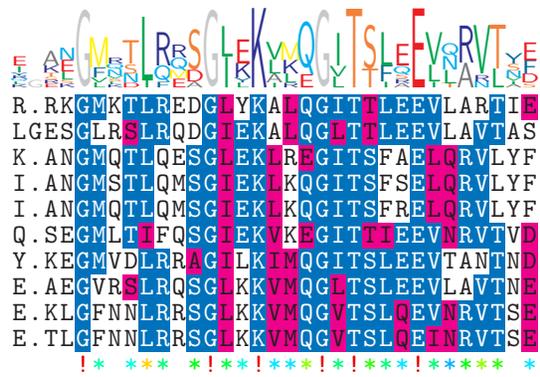
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 NTDB id 1018 ACIAD RS01685 WP 004920473.1
 consensus



857
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529
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logo

NTDB id 1028 TT RS08215 WP 011173991.1
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 consensus



889
887
562
561
561
574
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577
570
570

- non conserved
- similar
- ≥ 50% conserved