

logo

		MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	
NTDB id 1306 DR RS10060 WP 027479822.1		MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	80
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 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 1039021 ACCQ23 RS17525 WP 372177099.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus			

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		SIRPQTALNLQAFPFALEGDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGASDA	
NTDB id 1306 DR RS10060 WP 027479822.1		SIRPQTALNLQAFPFALEGDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGASDA	160
NTDB id 1168 A1552VC RS11070 WP 000957200.1		0
NTDB id 1293 VP RS12240 WP 005479695.1		0
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NTDB id 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 1039021 ACCQ23 RS17525 WP 372177099.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus			

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		GRTGGKLGGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEEGTTFLPNPSGFHPGEEVLGSM	
NTDB id 1306 DR RS10060 WP 027479822.1		GRTGGKLGGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEEGTTFLPNPSGFHPGEEVLGSM	240
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NTDB id 1110 NGFG RS09235 WP 003689817.1		0
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NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus			

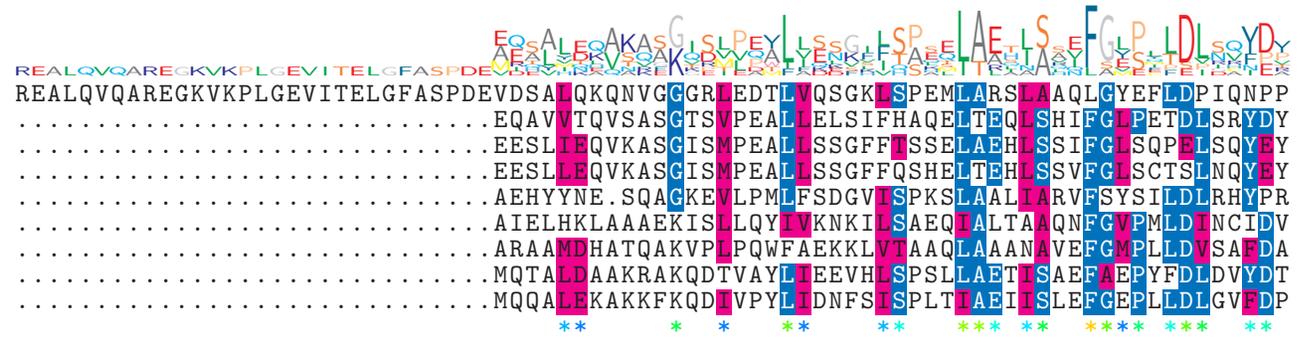
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		LRADALRLLAVPVDETEQGVTVLTS DPRKRPDI DAL I GRPVQLMLTRPRDI ER LIEQFYPQRGR LGEQLVQEGT LSRDQL	
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NTDB id 1401 DSB67 RS12665 WP 010643259.1	 MLTINLSTV LRQAGILTFSQ.	19
NTDB id 1110 NGFG RS09235 WP 003689817.1	 MSVGLLR I LVQNQV VIVEQ.	19
NTDB id 1251 GC085 RS07725 WP 011946523.1	 MALATEEYRLQ G I GQLLVLEKLLDKTK.	27
NTDB id 1039021 ACCQ23 RS17525 WP 372177099.1	 MNVAIGSNLVGI TGIARRLVQD GALDEAS.	29
NTDB id 1058 ABD1 RS01615 WP 001274986.1	 MSALHTSPKFT GFFRRLVEEKH VSAAT.	27
NTDB id 1018 ACIAD RS01685 WP 004920473.1	 MSAFTTPPKF S GFIRRLVEEGY VNAQN.	27
consensus		

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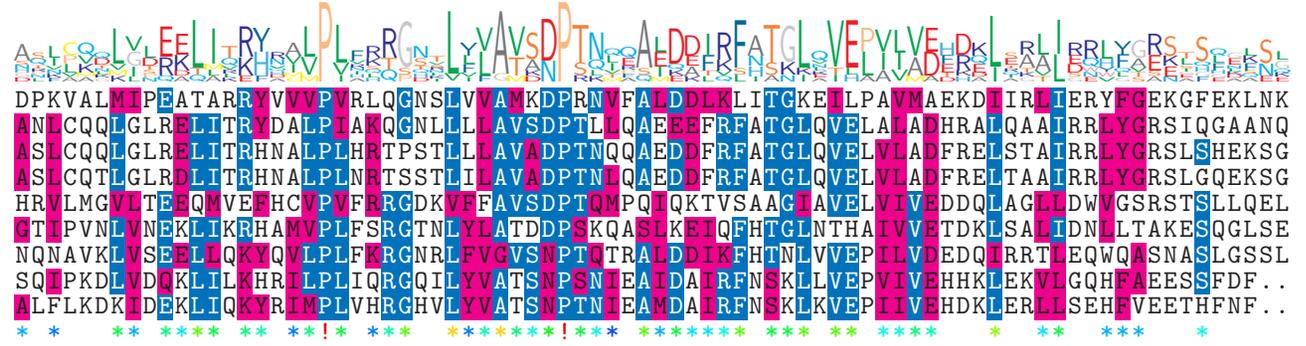
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 consensus



400
69
69
69
68
77
79
77
77

logo

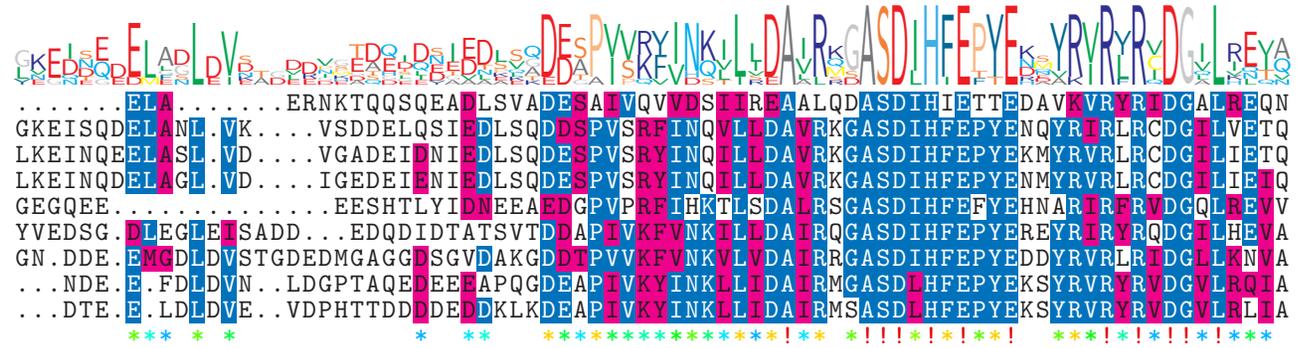
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 consensus



480
149
149
149
148
157
159
155
155

logo

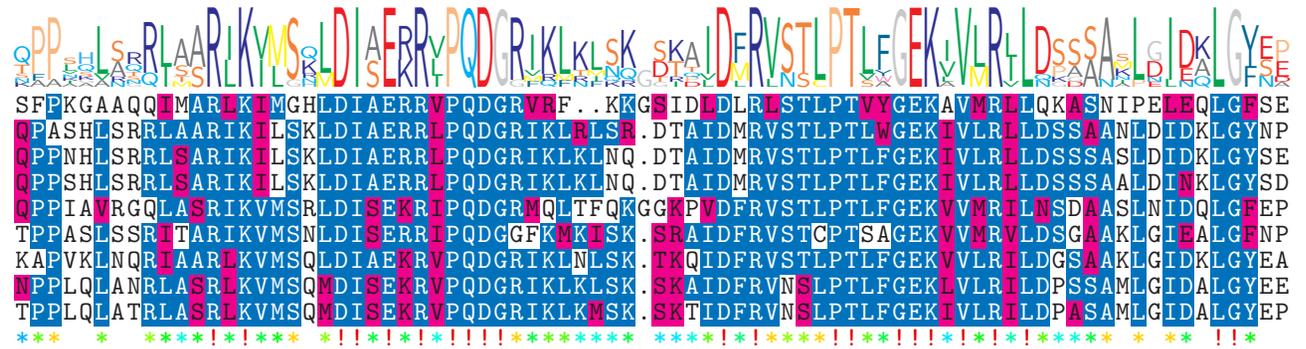
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 NTDB id 1058 ABD1 RS01615 WP 001274986.1
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 consensus



546
224
224
224
214
233
237
228
228

logo

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 NTDB id 1018 ACIAD RS01685 WP 004920473.1
 consensus



624
303
303
303
294
312
316
307
307

 non conserved
 similar
 $\geq 50\%$ conserved