

logo

	MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	
NTDB id 1306 DR RS10060 WP 027479822.1	MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	80
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NTDB id 1147359 CKV78 RS01740 WP 005764663.1	0
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NTDB id 1366 NTHI RS01950 WP 011271972.1	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 1018 ACIAD RS01685 WP 004920473.1	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
consensus		

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	SIRPQTALNLQAFPPFALEGDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLLEIIVVPSGASDA	
NTDB id 1306 DR RS10060 WP 027479822.1	SIRPQTALNLQAFPPFALEGDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLLEIIVVPSGASDA	160
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consensus		

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	GRTGGKLGGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITDQLYEVLAEQEGTTFLPNPSGFHPGEEVLGSM	
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NTDB id 1401 DSB67 RS12665 WP 010643259.1	0
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NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
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NTDB id 1306 DR RS10060 WP 027479822.1	LRADALRLLAVPVDETEQGVTVLTSDFRKRDPIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGLGEQLVQEGTLSRDQL	320
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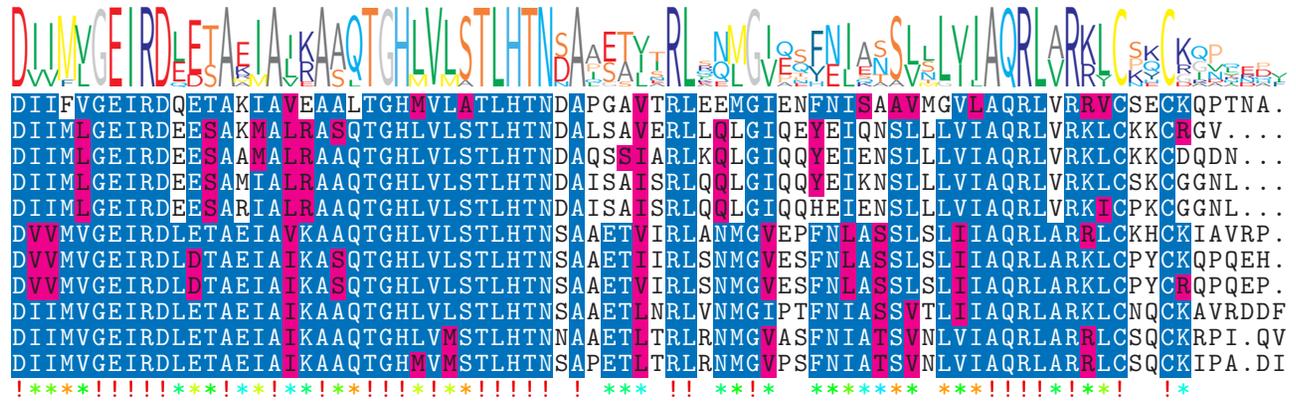
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consensus		

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NTDB id 1306 DR RS10060 WP 027479822.1	DPKVALMPEATARRYVVVVRQLQGNSLVVAMKDPNRVFDLDDKLIITGKEILPAVMAEKDIIIRLIERYFGEKGFELKLNK	480
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NTDB id 1355 HI 0298 AAC21962.1	DLWERNQQQQSLLRYFALPLKEENRNLWLGVDLSNLSACETIAFITGKPVPEILLLESSQLKELLQQLTPNQMQVVEEQV	104
NTDB id 1366 NTHI RS01950 WP 011271972.1	DLWERNQQQQSLLRYFALPLKEENRNLWLGVDLSNLSACETIAFITGKPVPEILLLESSQLKELLQQLTPHQMQVVEEQV	104
NTDB id 1168 A1552VC RS11070 WP 000957200.1	ANLCQQIGLRELITRYDALPIAKQGNLLAVSDPTLLQAEFEFRFATGLQVELALADHRAQAATRRLLYGRSIIQGAANQ	149
NTDB id 1293 VP RS12240 WP 005479695.1	ASLCQQIGLRELITRHNALPLHRTPTSTLLAVADPTNQQAEEDDFRFATGLQVELVLAADFRELSTAIRRLYGRSISHEKSG	149
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NTDB id 1018 ACIAD RS01685 WP 004920473.1	ALFLKDKIDEKLIQKYRIMPLVHRGHVLYVATSNPTEAMDAIRFNSKLLKVEPIIVEHDKLERLLSEHVEETHFNFD	157
NTDB id 1058 ABD1 RS01615 WP 001274986.1	SQIPKDIIVDQKLLILKHRILPLIQRGQILYVATSNPTEAMDAIRFNSKLLKVEPIIVEHDKLERLLSEHVEETHFNFD	157
consensus		

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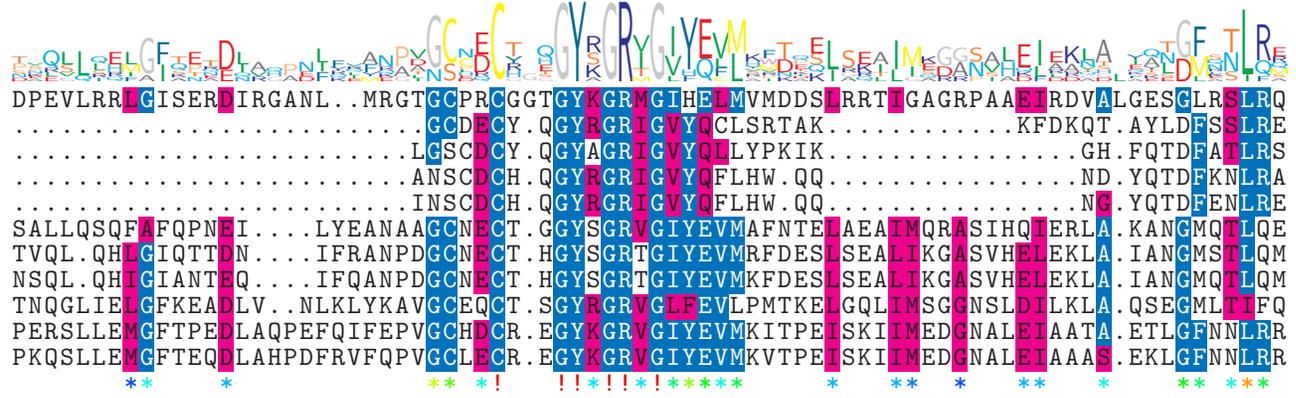
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 consensus



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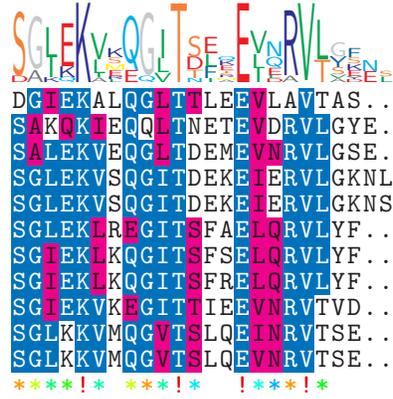
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 consensus



887
461
465
464
464
562
561
561
574
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570

⊗ non conserved
 ⊗ similar
 ⊗ ≥ 50% conserved