

logo

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NTDB id 1028	TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVEGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
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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 323759	D8668 RS05455 WP 003116249.1	0
NTDB id 1058	ABD1 RS01615 WP 001274986.1	0
NTDB id 1018	ACIAD RS01685 WP 004920473.1	0
consensus			

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		RSIRPQTALNLQAFPPFALE..GDRRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLLEIVVPSGA	
NTDB id 1028	TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDPLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLLPVPPP..P	158
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NTDB id 1251	GCO85 RS07725 WP 011946523.1	0
NTDB id 323759	D8668 RS05455 WP 003116249.1	0
NTDB id 1058	ABD1 RS01615 WP 001274986.1	0
NTDB id 1018	ACIAD RS01685 WP 004920473.1	0
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		SGEGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVDPDPSAA	
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NTDB id 1401	DSB67 RS12665 WP 010643259.1	0
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 NTDB id 1058 ABD1 RS01615 WP 001274986.1
 NTDB id 1018 ACIAD RS01685 WP 004920473.1

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 GMSLRADALRLLAVPVDETEQGVTVLTS DPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPPQRGRLLGEQLVQEGTLSR

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NTDB id 1028 TT RS08215 WP 011173991.1
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 NTDB id 1018 ACIAD RS01685 WP 004920473.1

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MSVGLLRILVQNRVVTVEQAEEHYNESQ.AGKEVLPMLFSDGVI SPKSLAAL IARVFSYSILDLRH
MLTNLVAILRQAE LISATQEQAVVTQVSASGTSVPEALLELSIFHAQELTEQLSHIFGLPETDLSR
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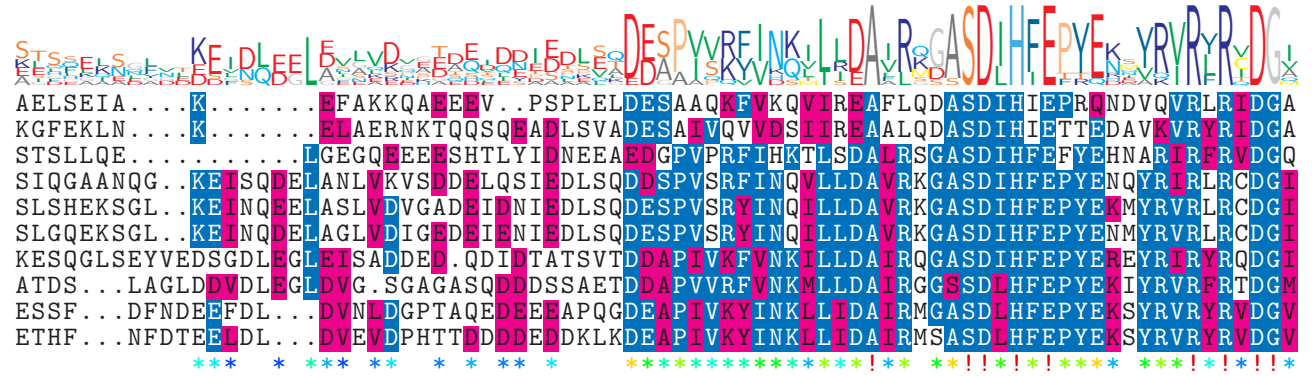
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 YPRHRVLMGVLTEEQMVEFHCVVFRRGDKVFFAVSDPTQMPQIQKTVSAAG.....TAVELVIVEDDQLAGLLDWVGSR
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 YEYASLCQQLGLRELITRHNALPLHRTPTSTLLLAVADPTNQAEDDFRFATG.....LQVELVLADFRELSTAIRRLYGR
 YEYASLCQTLGLRDLITRHNALPLNRTSSTLILAVADPTNLQAEDDFRFATG.....LQVELVLADFRELTAAIRRLYGR
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 LERDSL PKDLISEKLVQRHRVPLWRGRGNKLFVGISDPANHQAINDVQFSTG.....LTTEAILVEDDKLGIITIEKLFEN
 YDTSQIPKDLVDQKLLKHRILPLIQRGQILYVATSNPSNTEAIDAIRFNSK.....LLVEPVIVEHHKLEKVLGQHFAE
 FDPALFLKDKIDEKLIQKYRIMPLVHRGHVLYVATSNPTNIEAMDAIRFNSK.....LKVEPIIVEHDKLERLLSEHFVE

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consensus

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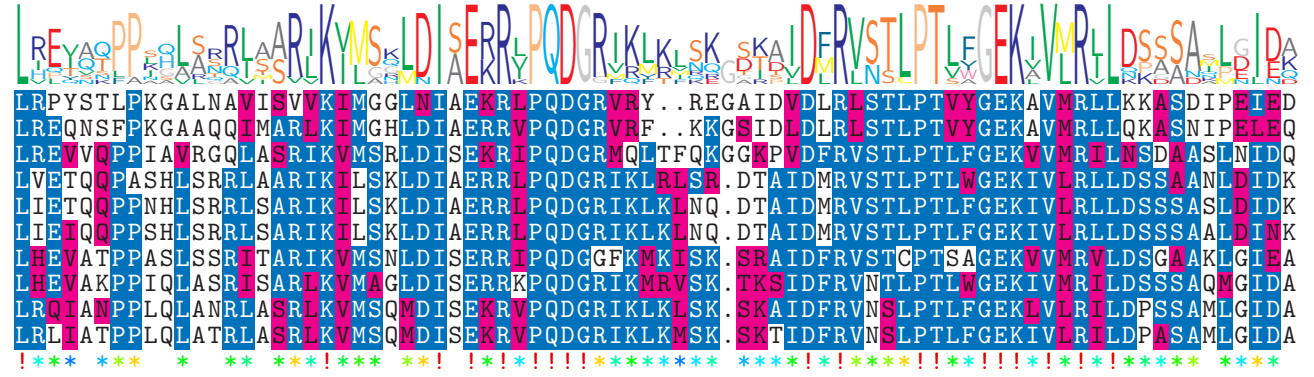
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 consensus



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 consensus



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 consensus



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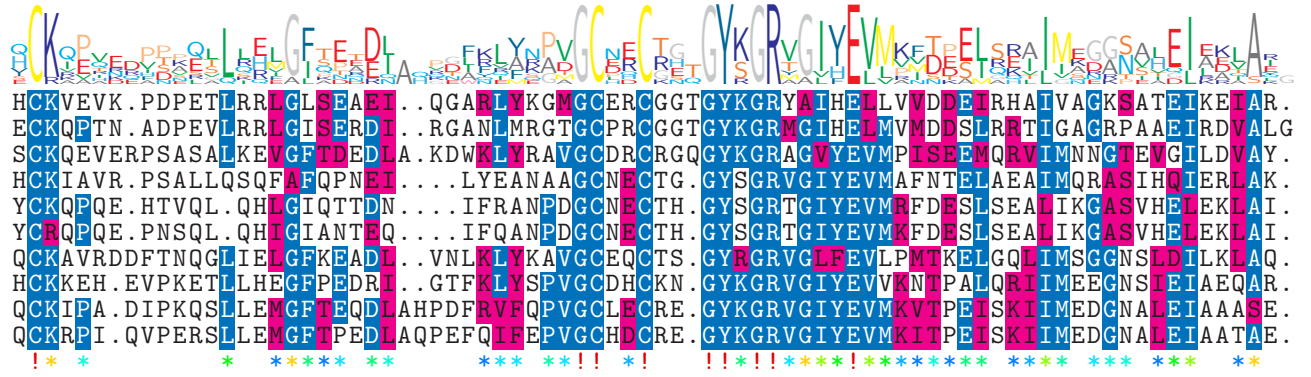
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 consensus



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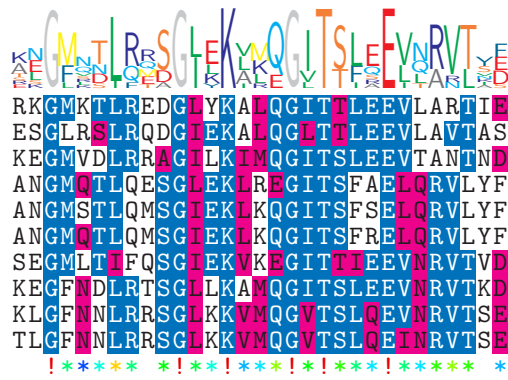
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 consensus



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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