

logo

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NTDB id 1366	NTHI RS01950 WP 011271972.1	0
NTDB id 268916	PH4a RS02200 WP 129037426.1	0
NTDB id 1168	A1552VC RS11070 WP 000957200.1	0
NTDB id 1293	VP RS12240 WP 005479695.1	0
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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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		KETPKKAKELRAIPFALDEEAGVVRVAFNLPLDLSLEEVEDLTGLVVEPYQTTKSFLYALAKHYPELGLPVPPP..P	
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NTDB id 1028	TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDLSLEEVEDLTGLVVEPYQTTKSFLYALAKHYPELGLPVPPP..P	158
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NTDB id 1018	ACIAD RS01685 WP 004920473.1	0
NTDB id 1058	ABD1 RS01615 WP 001274986.1	0
consensus			

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NTDB id 1410 SGL RS13950 WP 010873519.1	0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
consensus		

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NTDB id 1251 GCO85 RS07725 WP 011946523.1	0
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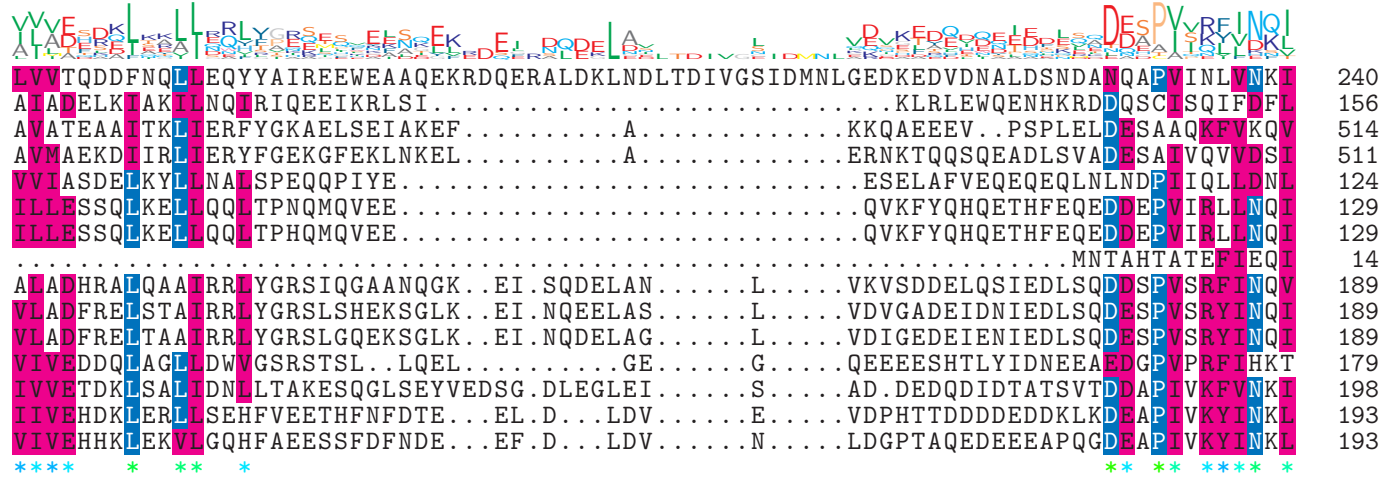
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NTDB id 1110 NGFG RS09235 WP 003689817.1MSVGLLRILVQNNQVVTVEQAEHY..NESQAGKEVLPMLFSD..GVISPKSLAAL.....LARVF	56
NTDB id 1251 GCO85 RS07725 WP 011946523.1MALATEEYRLQGIGQLVLEKLLDKTKAIELHKLAAAEEKISLLQYIVKN..KILSAEQIALT.....AAQNF	65
NTDB id 1018 ACIAD RS01685 WP 004920473.1MSAFTTPPKFSGFIRRLVEEGYVNAQNMQQALEKAKKFKQDIVPYLIDN..FSISPLTIAEI.....LSLEF	65
NTDB id 1058 ABD1 RS01615 WP 001274986.1MSALHTSPKFTGFFRRLVEEKHVSAAATMQTALDAAKRAKQDTVAYLIEE..VHISPSLLAET.....LSAEF	65
consensus*.....*.....*.....*	

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NTDB id 1410 SGL RS13950 WP 010873519.1	GVDSVDPEVAPIATQQMAELIGRFFPLDTCRRIFKFIPTAQQEGDPPSILVAVVDPDNLAAQDEIN.RILRVKG..FETRR	160
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NTDB id 1028 TT RS08215 WP 011173991.1	GYPYVDPEEDPPD...PGAPLLLPEDLCCRRYGVFPHRLEG..NRIVLLMKDPRNLLALDDVR.LALKRKGLNYEVAP	459
NTDB id 1306 DR RS10060 WP 027479822.1	GYEFLDPIQNPPD....PKVALMIPEATARRVVVVPVRLQG...NSLVVAMKDPNRFALDDLK.LITG.....KETLP	454
NTDB id 1384 A4U84 RS08525 WP 010787006.1	D.....ISPS....RWQKNCEEKQLLLRYLAVPVQETE...HKLWLAVDDENNLTACEIFA.FMTH.....KQIEP	73
NTDB id 1355 HI 0298 AAC21962.1	T.....ISPD....LWERNQQQSSLLRYFALPIKEEN...NRLWLGVDLSLSNISACETIA.FITG.....KPVEP	78
NTDB id 1366 NTHI RS01950 WP 011271972.1	T.....ISPD....LWERNQQQSSLLRYFALPIKEEN...NRLWLGVDLSLSNISACETIA.FITG.....KPVEP	78
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NTDB id 1401 DSB67 RS12665 WP 010643259.1	GLSCTSLNQYEYA....SLCQTLGLRDLITRHNALPINRTS...STLLAVADPTNQAEDDFR.FATG.....LQVEL	123
NTDB id 1110 NGFG RS09235 WP 003689817.1	SYSILDLRHYPRH....RVLMGVLTTEEQVVEHCVVFRRG...DKVFFAVSDPTQMPPIQKTV.SAAG.....IAVEL	122
NTDB id 1251 GCO85 RS07725 WP 011946523.1	GVPMLDINCIDVG....TIPVNLVNEKLIKRRHAMVPLFSRG...TNLYLATDDPSKQASLKEITQ.FHTG.....LNTHA	131
NTDB id 1018 ACIAD RS01685 WP 004920473.1	CEPLLDLGVFDPA....LFLKDKIDEKLIQKVRIMPIVHRG...HVLYVATSNPTNTEAMDATR.FNSK.....LKVEP	131
NTDB id 1058 ABD1 RS01615 WP 001274986.1	AEPYFDLDVYDTS....QIPKDLVDQKLLKHRILPIIQRG...QILYVATSNPSNTEAIDATR.FNSK.....LLVEP	131
consensus	*.....*.....** ** ** ** **.....*.....*.....*	

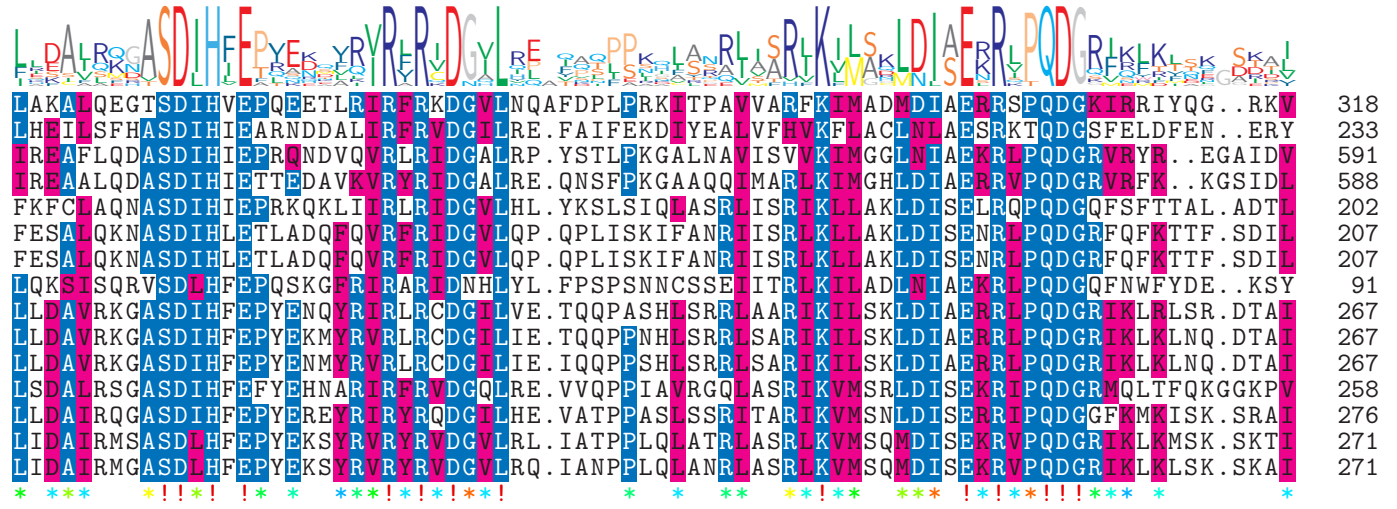
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 consensus



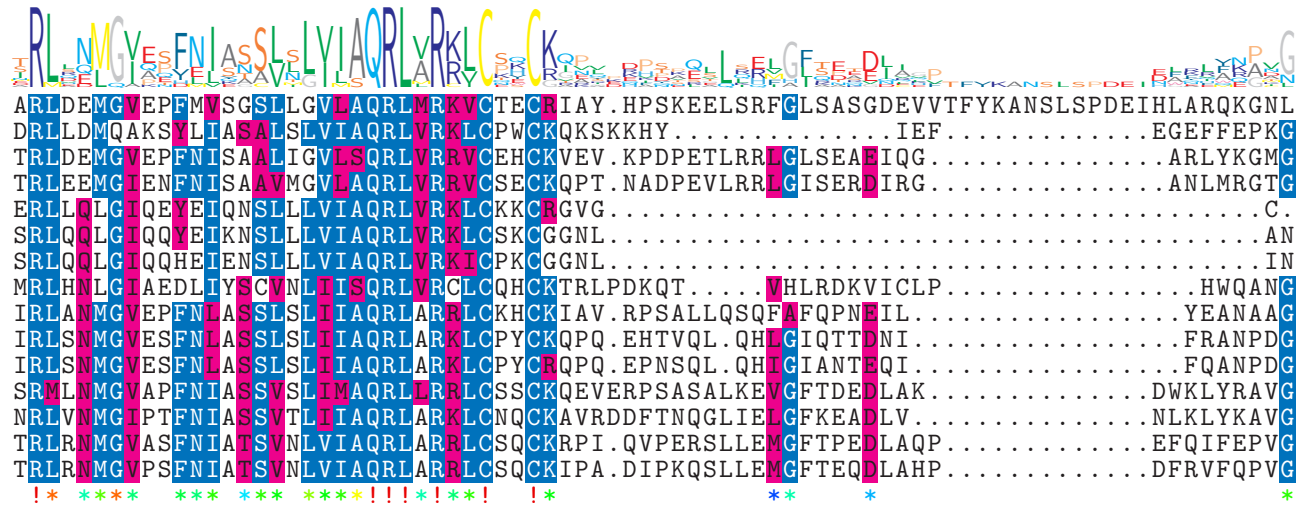
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 consensus



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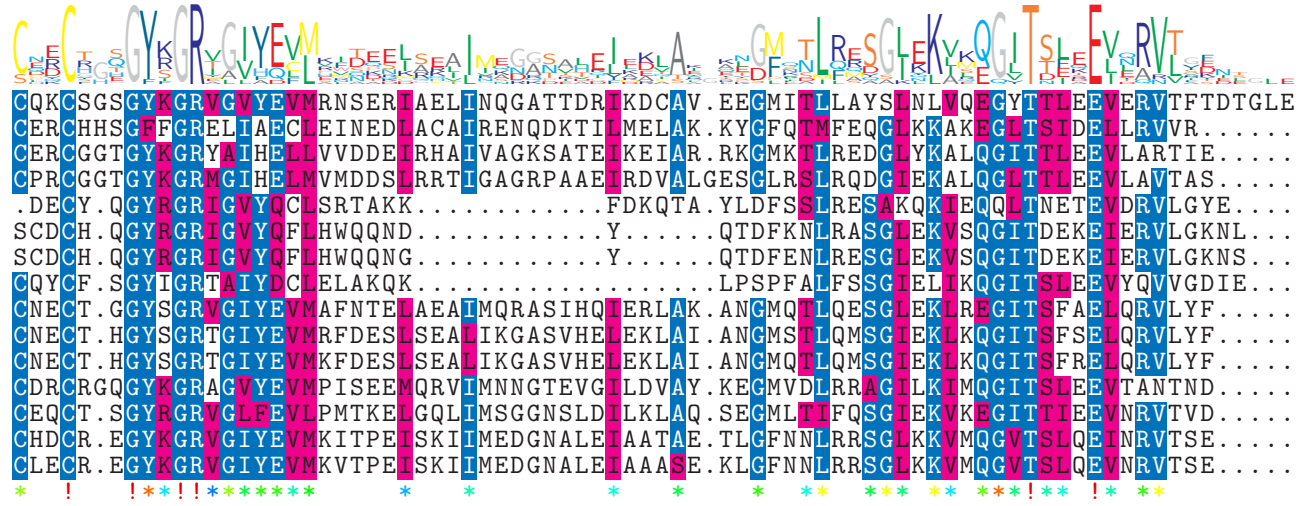
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 consensus



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NTDB id 1410 SGL RS13950 WP 010873519.1
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 consensus



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519
889
887
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logo

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NTDB id 1028 TT RS08215 WP 011173991.1	889
NTDB id 1306 DR RS10060 WP 027479822.1	887
NTDB id 1384 A4U84 RS08525 WP 010787006.1	461
NTDB id 1355 HI 0298 AAC21962.1	464
NTDB id 1366 NTHI RS01950 WP 011271972.1	464
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NTDB id 1401 DSB67 RS12665 WP 010643259.1	561
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NTDB id 1251 GCO85 RS07725 WP 011946523.1	574
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consensus		

 non conserved
 similar
 ≥ 50% conserved