

logo

NTDB id 1410 SGL RS13950 WP 010873519.1	MSXLSIGDRRLGALLDQGYLDLQRLALERHSEVGGRLAEVLYDSQYSEERLAEALIESEFGIPLVQLHSEVETSPSSK	0
NTDB id 1306 DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNLDLQRLALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAAL	79
NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRLALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1168 A1552VC RS11070 WP 000957200.1		0
NTDB id 1401 DSB67 RS12665 WP 010643259.1		0
NTDB id 1020549 QRP18 RS12680 WP 140062768.1		0
NTDB id 1293 VP RS12240 WP 005479695.1		0
NTDB id 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1251 GC085 RS07725 WP 011946523.1		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 1410 SGL RS13950 WP 010873519.1	SETPAEKAKELRAIPFALSEEKSRVAHYNPLSLSEEDSGFYEPYQCTREAYMAAASHYPELGLLEYMPSCK	0
NTDB id 1306 DR RS10060 WP 027479822.1	RSIRPQTALNLQAFFFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLIEVVPSPA	157
NTDB id 1028 TT RS08215 WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDTLSEEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
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consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	SGSGSGSKLGEELTSGYIIBSLLEYALYQERTCGALGRTLYSBRATFEELYEXLAEQEGTEFLESTEGYMPSEEXK	0
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NTDB id 1028 TT RS08215 WP 011173991.1	SGEGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAA	237
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NTDB id 1293 VP RS12240 WP 005479695.1		0
NTDB id 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1251 GC085 RS07725 WP 011946523.1		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 1110 NGFG RS09235 WP 003689817.1		0
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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 1020549 QRP18 RS12680 WP 140062768.1	MLTNLSTVLRQKGLLTF SQEESLLEQVKASCISMPEALLS..SCFFTSS ELAEH.....LSSIF	57
NTDB id 1293 VP RS12240 WP 005479695.1	MHSNLS TILRQKGLLTF SQEESLLEQVKASCISMPEALLS..SCFFTSS ELAEH.....LSSIF	57
NTDB id 1110 NGFG RS09235 WP 003689817.1	MSVGLLRILVQNVVTV EQAHYY..NESQACKEVLPMLFS..DCVISP KSLAAL.....LARVF	56
NTDB id 1251 GC085 RS07725 WP 011946523.1	MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLAAAEKISL LQYIVK..NKILSAEQIALT.....AAQNF	65
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NTDB id 1018 ACIAD RS01685 WP 004920473.1	MSAFTTPPKFSGFIRRLV EECYVNAQIMQQA LEKAKKFKQDVPYLLID..NFSLSPLTAEI.....LSLEF	65
consensus	*****	

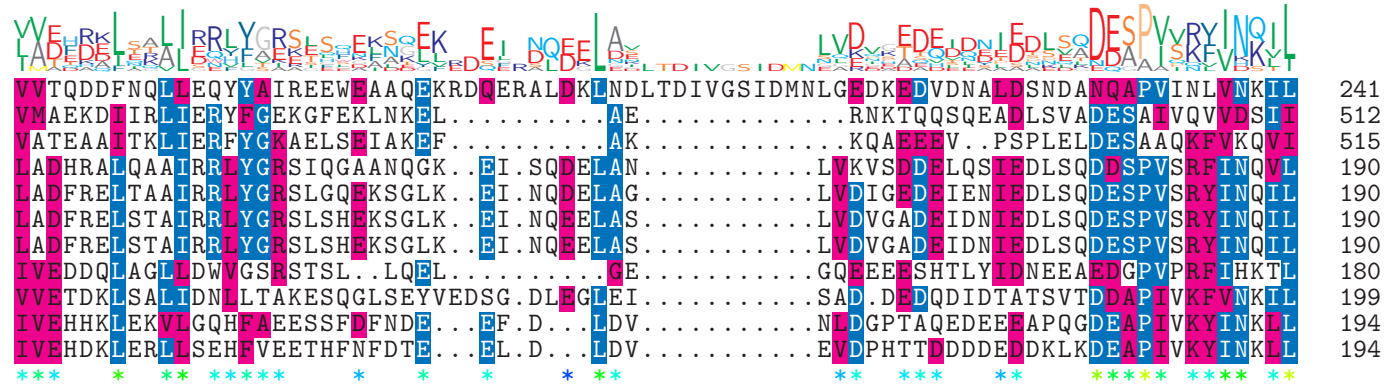
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NTDB id 1401 DSB67 RS12665 WP 010643259.1	GLSCTSLNQY EYA.....SLCQTLGLRDLITRHNALPLNRT...SSTLLAVADPTNLQAEDDFRFATG.....LQVELV	124
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NTDB id 1251 GC085 RS07725 WP 011946523.1	GVPMIDINCIDVG.....TIPVNLVNEKLIKRRHAMVPLFSR...CTNLYLATDDPSKQASLKEIQPHTG.....LNTHAI	132
NTDB id 1058 ABD1 RS01615 WP 001274986.1	AEPYFDLDVYDTS.....QIPKDLVDQKLILKHRILPLIQR...GQILYVATSNPSNTEADAI RFNSK.....LLVEPV	132
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consensus	*****	

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consensus

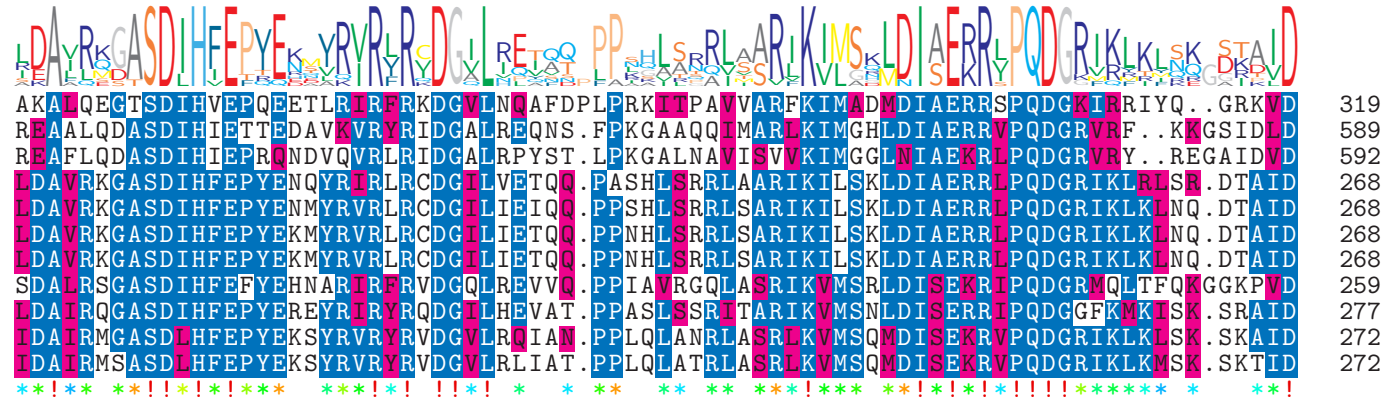
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consensus

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NTDB id 1410 SGL RS13950 WP 010873519.1
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NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

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NTDB id 1410 SGL RS13950 WP 010873519.1	EMRAKRKSALTCTCCTCAAELQQEWLDCPYCMTSRFS	672
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NTDB id 1058 ABD1 RS01615 WP 001274986.1		570
NTDB id 1018 ACIAD RS01685 WP 004920473.1		570
consensus		

- ⓧ non conserved
- ⓧ similar
- ⓧ ≥ 50% conserved