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NTDB id 1306 DR RS10060 WP 027479822.1	MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	80
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	.....	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	.....	0
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NTDB id 1306 DR RS10060 WP 027479822.1	SIRPQTALNLQAFPFALEGDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLLEIVVPSGASDA	160
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NTDB id 1306 DR RS10060 WP 027479822.1	GRTGGKLGGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVLGSM	240
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NTDB id 1306 DR RS10060 WP 027479822.1	REALQVQAREGKVKPLGEVITELGFASPDEVDSA LQKQNVGGGRLEDTLVQSGKLSPEMLARSLAAQLGYEFLDPIQNPP	400
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NTDB id 311997 D1108 RS04405 WP 237593620.1	..... MQY..... SVCDVNNR..... QVFRISE	18
NTDB id 1384 A4U84 RS08525 WP 010787006.1	..... MSQY..... TICEQQTE..... RIFDISP	19
NTDB id 1168 A1552VC RS11070 WP 000957200.1	..... MLTNLVAI LRQAEL ISATQEQAV VTQVSASGTSVPEALLELSIFHAQELTEQLSHIFGLPETDL SRYDY	69
NTDB id 1401 DSB67 RS12665 WP 010643259.1	..... MLTNLSTV LRQAGI LTFSQEESL IEQVKASGISMPEALLSSGFFQSHL EHLSSV FGLSCTSLNQYDY	69
NTDB id 1293 VP RS12240 WP 005479695.1	..... MHSNLSTI LRQKGL LTFSQEESL IEQVKASGISMPEALLSSGFFTSSELAEHLSSIFGLSQPELSQYDY	69
NTDB id 1251 GC085 RS07725 WP 011946523.1	... MALATEEYRLQGIGQL LVLEKLLDKTKAIELHKLAAAEKISLLQYIVKNKILSAEQIALTAAQNFQVPMLDINCIDV	77
NTDB id 1058 ABD1 RS01615 WP 001274986.1	... MSALHTSPKFTGFFRRLVVEEKH VSAATMQTALDAAKRAKQDTVAYLIEEVHLSPLSLAETISAFAEPYFDLDVYDT	77
NTDB id 1018 ACIAD RS01685 WP 004920473.1	... MSAFTTPPKFSGFIRRLVVEEGYVNAQNMQAL EKAKKFKQDTPVYLLDNFSLISPLTIAEILSLEFGEPLLDLGVFDP	77

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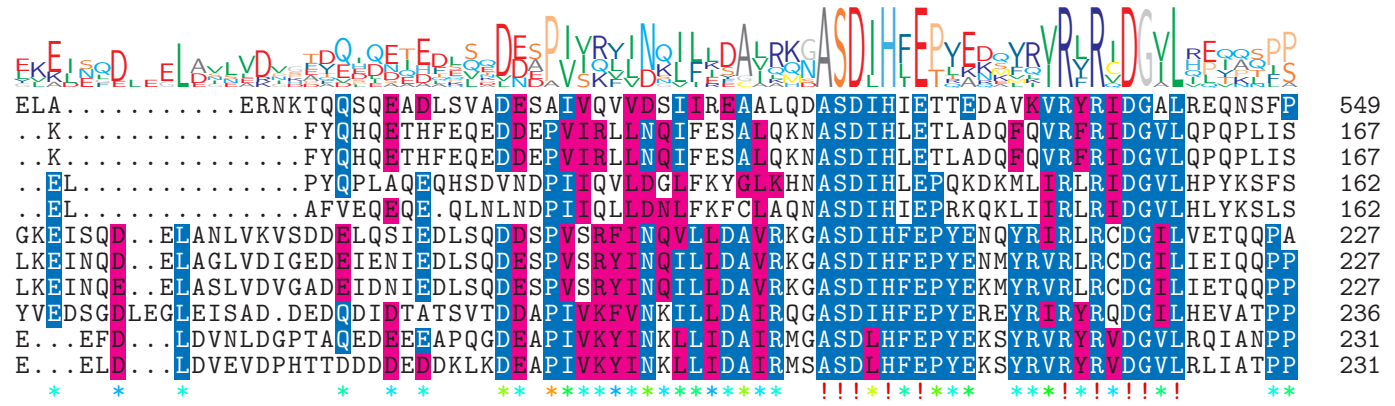
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NTDB id 1366 NTHI RS01950 WP 011271972.1	DLWERNQQQQS L L LRYFALPLKEEN N RLW LGVDSL SNLSACET IAFITGKPV EPI LLESSQLKELLQQLTPHQMQVEEQV	104
NTDB id 311997 D1108 RS04405 WP 237593620.1	EQWQKNYDERH L L LRYLTVPLREDQ T L W L A I D D M N N L S A C E I F A F L T H K R I E P V L V S S E E L K Y L L G L M P E Q Q S V Y D E T	98
NTDB id 1384 A4U84 RS08525 WP 010787006.1	SRWQKNCEEKQ L L LRYLAVPVQETEHK L W L A V D D E N N L T A C E I F A F M T H K Q I E P V V I A S D E L K Y L L N A L S P E Q Q I Y E E S	99
NTDB id 1168 A1552VC RS11070 WP 000957200.1	ANLCQQLGLREL I T R Y D A L P I A K Q G N L L L L A V S D P T L L Q A E E F R F A T G L Q V E L A L A D H R A L Q A A I R R L Y G R S I Q G A A N Q	149
NTDB id 1401 DSB67 RS12665 WP 010643259.1	ASLCQTLGLRD L I T R H N A L P L N R T S S T L I L A V A D P T N L Q A E D D F R F A T G L Q V E L V L A D F R E L T A A I R R L Y G R S L G Q E K S G	149
NTDB id 1293 VP RS12240 WP 005479695.1	ASLCQQLGLREL I T R H N A L P L H R T P S T L L L A V A D P T N Q Q A E D D F R F A T G L Q V E L V L A D F R E L S T A I R R L Y G R S L S H E K S G	149
NTDB id 1251 GC085 RS07725 WP 011946523.1	GTIPVNLVNEK L I K R H A M V P L F S R G T N L Y L A T D D P S K Q A S L K E I Q F H T G L N T H A I V V E T D K L S A L I D N L L T A K E S Q G L S E	157
NTDB id 1058 ABD1 RS01615 WP 001274986.1	SQIPKDLVDQK L I L K H R I L P L I Q R G Q I L Y V A T S N P S N I E A I D A I R F N S K L L V E P V I V E H H K L E K V L G Q H F A E E S S F D F N D	157
NTDB id 1018 ACIAD RS01685 WP 004920473.1	ALFLKDKIDEK L I Q K Y R I M P L V H R G H V L Y V A T S N P T N I E A M D A I R F N S K L K V E P I I V E H D K L E R L L S E H F V E E T H F N F D T	157

consensus

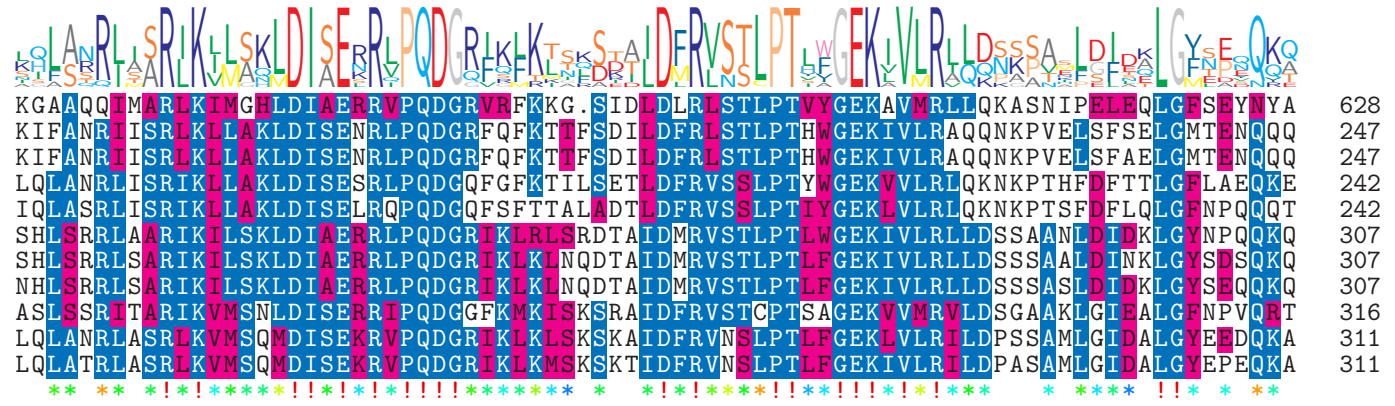
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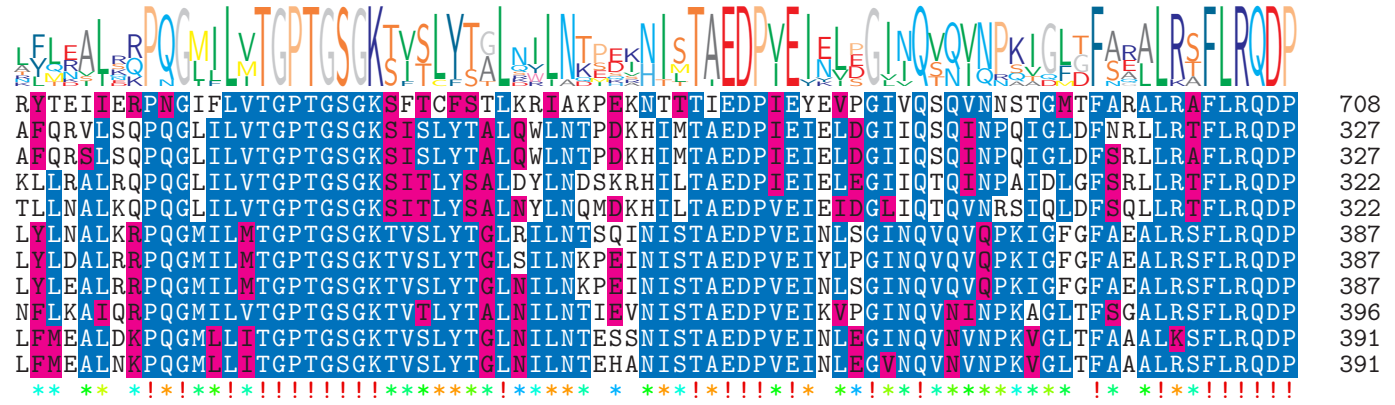
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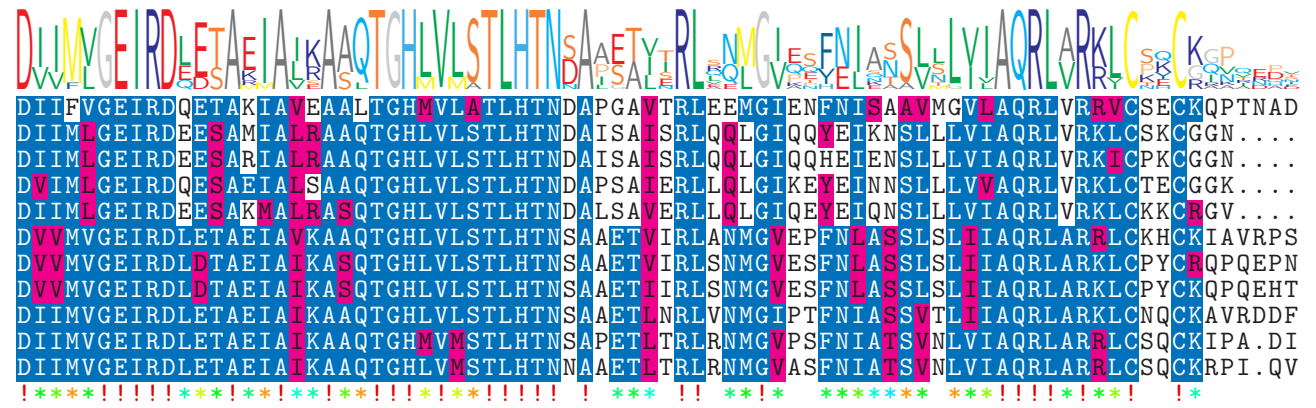
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 consensus



887  
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X non conserved  
 X similar  
 X ≥ 50% conserved