

logo

NTDB id 1410 SGL RS13950 WP 010873519.1
NTDB id 1366 NTHI RS01950 WP 011271972.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 51696 MPUCK001 RS04810 WP 104869078.1
NTDB id 1436 BW25113 RS00530 WP 001025145.1
NTDB id 1168 A1552VC RS11070 WP 000957200.1
NTDB id 1401 DSB67 RS12665 WP 010643259.1
NTDB id 1293 VP RS12240 WP 005479695.1
NTDB id 1110 NGFG RS09235 WP 003689817.1
NTDB id 1251 GC085 RS07725 WP 011946523.1
NTDB id 1018 ACIAD RS01685 WP 004920473.1
NTDB id 1058 ABD1 RS01615 WP 001274986.1

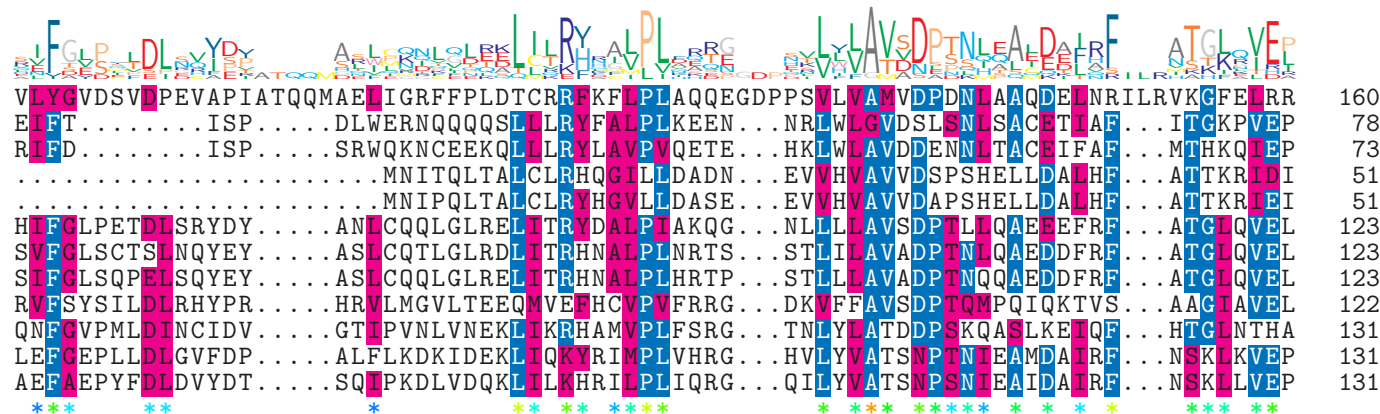
consensus



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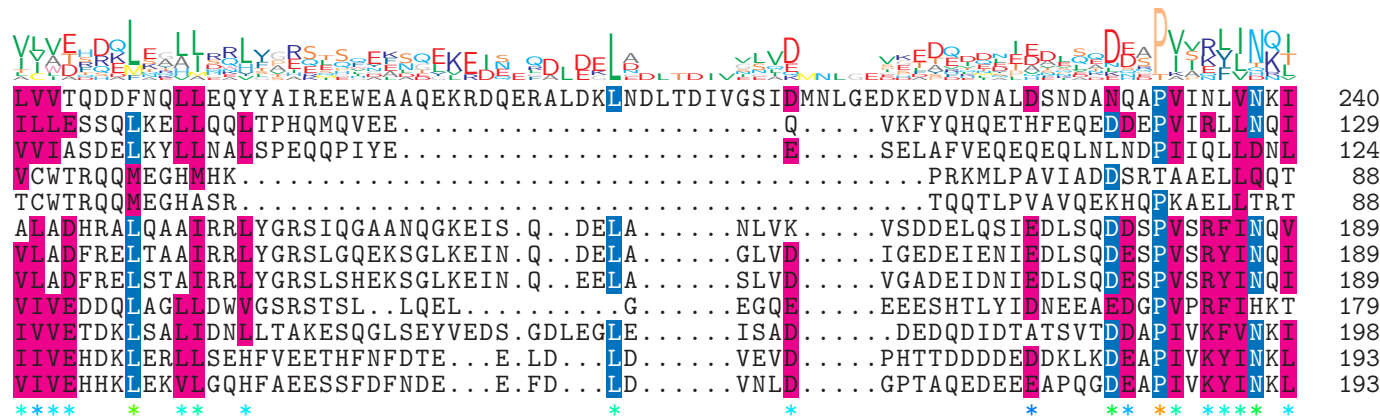
consensus



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consensus



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NTDB id 1410 SGL RS13950 WP 010873519.1	ARLIDEMGV	557
NTDB id 1366 NTHI RS01950 WP 011271972.1	SRLQLGLGIQQHETIENSLLLVIAQRLVRKICPKCGGNL	406
NTDB id 1384 A4U84 RS08525 WP 010787006.1	ERLLQLGIQEVEIQNSLLLVIAQRLVRKLCCKCGVG	399
NTDB id 51696 MPUCK001 RS04810 WP 104869078.1	IRLQMGVARWMISSALTLVIAQRLVRKLCPCYCRHLAAESVS	384
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NTDB id 1401 DSB67 RS12665 WP 010643259.1	IRLSNMGVESFNLASSLSLIAQRLARKLCPYCRQPQ	488
NTDB id 1293 VP RS12240 WP 005479695.1	IRLSNMGVESFNLASSLSLIAQRLARKLCPYCKQPQ	488
NTDB id 1110 NGFG RS09235 WP 003689817.1	SRMLNMGVAPFNIASSVSLIAQRLRLRLCSSCKQE	484
NTDB id 1251 GC085 RS07725 WP 011946523.1	NRLVNMGIPTFNIASSVTLLIAQRLARKLCLNQCK	501
NTDB id 1018 ACIAD RS01685 WP 004920473.1	TRLRNMGVASFNIATSVNLVIAQRLARRLCSQCKRPI	497
NTDB id 1058 ABD1 RS01615 WP 001274986.1	TRLRNMGVPSFNIATSVNLVIAQRLARRLCSQCKIPA	497
consensus	!* **!* *****!***!!!! !*** !* * **	

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NTDB id 1410 SGL RS13950 WP 010873519.1	CQKCSGS	637
NTDB id 1366 NTHI RS01950 WP 011271972.1	SCDCH.QGYRGRIGVYQFLHWQQN	464
NTDB id 1384 A4U84 RS08525 WP 010787006.1	CDECY.QGYRGRIGVYQCLSRITAK	461
NTDB id 51696 MPUCK001 RS04810 WP 104869078.1	CERCY.HGFFYGRITALFEVLPVTPAIRQLLINNATVEAL	461
NTDB id 1436 BW25113 RS00530 WP 001025145.1	CVHCY.HGFFYGRITALFEVLPITPVIRQLISANTDVESLETHARQA	461
NTDB id 1168 A1552VC RS11070 WP 000957200.1	CNECT.GGYSGRVGIYEVMAFNTELAEAIMQRASIHQIERLAKANGMQTLQESG	562
NTDB id 1401 DSB67 RS12665 WP 010643259.1	CNECT.HGYSGRGTGIYEVUMKFDESLSSEALIKGASVHELEKLAIANGM	561
NTDB id 1293 VP RS12240 WP 005479695.1	CNECT.HGYSGRGTGIYEVUMRFDESLSSEALIKGASVHELEKLAIANGM	561
NTDB id 1110 NGFG RS09235 WP 003689817.1	CDRCRGQGYKGRAGVYEVMPISEEMQVRVIMNNGTEVGILDVAYKEGMVDLRR	558
NTDB id 1251 GC085 RS07725 WP 011946523.1	CEQCT.SGYRGRVGLFEVLPMTIKELGQLIMSGGNSLDILKLAQSEGMLTIFQSG	574
NTDB id 1018 ACIAD RS01685 WP 004920473.1	CHDCR.EGYKGRVGIYEVUMKITPEISKIIMEDGNALEIAATAETLGFNNLRRSGL	570
NTDB id 1058 ABD1 RS01615 WP 001274986.1	CLECR.EGYKGRVGIYEVUMKVITPEISKIIMEDGNALEIAAASEKLGFN	570
consensus	* *! !***!***** * * *** * * ** ** ** **	

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NTDB id 1410 SGL RS13950 WP 010873519.1	EMRAKRKSALT	672
NTDB id 1366 NTHI RS01950 WP 011271972.1	EMRAKRKSALT	464
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	EMRAKRKSALT	570
consensus	EMRAKRKSALT	

- X non conserved
- X similar
- X ≥ 50% conserved