

logo

NTDB id 1306 DR RS10060 WP 027479822.1	REALQVQAREGKVKPLGEVITELGFASPD	400
NTDB id 1168 A1552VC RS11070 WP 000957200.1	EQAVVTQVSASGTSVPEALLSIFHAQELTEQLSHIFG	69
NTDB id 1293 VP RS12240 WP 005479695.1	EESLIEQVKASGISMPEALLSSGFFISSELAEHLSSIF	69
NTDB id 1401 DSB67 RS12665 WP 010643259.1	EESSLIEQVKASGISMPEALLSSGFFQSHLTHSSVFC	69
NTDB id 1110 NGFG RS09235 WP 003689817.1	AEHYYNESQA.CKEVLPMLFSDGVI.SPKSLAALIA	68
NTDB id 1251 GC085 RS07725 WP 011946523.1	AIELHKLAAAEKISLLQYIVKNKILSAEQIALTAAQ	77
NTDB id 1101889 Q9286 RS07775 WP 011408072.1	ARTAMEQAAA.AKVPLPQWFAENRLVSASQLAAANA	79
NTDB id 1058 ABD1 RS01615 WP 001274986.1	MQTALDAAKRAKQDVTAYLIEEVHLSPSLAEETIS	77
NTDB id 1018 ACIAD RS01685 WP 004920473.1	MQQALEKAKKFQKD.VPPLYLDNFS.SPLTTABETIS	77
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NTDB id 1306 DR RS10060 WP 027479822.1	DPKVALMIPETARRVYVVPVRLQGNLSLVAMKDPRN	480
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NTDB id 1293 VP RS12240 WP 005479695.1	ASLCQQQLGLRELITRHNALPLHRTPTSTLLLAVADPT	149
NTDB id 1401 DSB67 RS12665 WP 010643259.1	ASLCQTLGLRDLITRHNALPLNRTSSTLLIIVADPTNI	149
NTDB id 1110 NGFG RS09235 WP 003689817.1	HRVLMGVLTETEQMVEFHCVPVFRRGDKVFFAVSDPT	148
NTDB id 1251 GC085 RS07725 WP 011946523.1	GTIPVNLVNEKLIKRRHAMVPLFSRGTNLYLATDDPS	157
NTDB id 1101889 Q9286 RS07775 WP 011408072.1	NQNAVKLVSEELIQKYHVLPFLKRCNRLFVGVSNPTQ	159
NTDB id 1058 ABD1 RS01615 WP 001274986.1	SQIPKDLVDQKLIKLRHRLPLIQRGQILYVATSNPSN	157
NTDB id 1018 ACIAD RS01685 WP 004920473.1	ALFLKDKIDKELIQKYRIMPLVHRGHVLYVATSNPTN	157
consensus	*****	

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NTDB id 1306 DR RS10060 WP 027479822.1	ELA.....ERNKTQQSQEADLSVADESALVQVVD	547
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NTDB id 1293 VP RS12240 WP 005479695.1	LKEINQEELASLVD...VGADEINIEDLSQDES	225
NTDB id 1401 DSB67 RS12665 WP 010643259.1	LKEINQDELAGLVD...IGDEIENIEDLSQDES	225
NTDB id 1110 NGFG RS09235 WP 003689817.1	GEGQ.E.....EESHTLYIDNEEAEDGPVPRFI	215
NTDB id 1251 GC085 RS07725 WP 011946523.1	YVEDSGDLEGLETSADD...EDQDIDTATSVTD	234
NTDB id 1101889 Q9286 RS07775 WP 011408072.1	GNDD.EEMGDLDVSTGDEDMGAGGDSGVDAKGDD	238
NTDB id 1058 ABD1 RS01615 WP 001274986.1	EE.FDLVDN..LDGPTAQEDDEEAPQGD	229
NTDB id 1018 ACIAD RS01685 WP 004920473.1	EE.LDLDVD..VDPHTTDDDEDKCLKD	229
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NTDB id 1306 DR RS10060 WP 027479822.1	FPKGAAQQIMARLIKIMGHLDIAERRVPQDGRVRF	625
NTDB id 1168 A1552VC RS11070 WP 000957200.1	PASHLSRRRLAARIKILSKLDIAERRLPQDGRIKL	304
NTDB id 1293 VP RS12240 WP 005479695.1	PPNHLSRRLSARIKILSKLDIAERRLPQDGRIKL	304
NTDB id 1401 DSB67 RS12665 WP 010643259.1	PPSHLSRRLSARIKILSKLDIAERRLPQDGRIKL	304
NTDB id 1110 NGFG RS09235 WP 003689817.1	PPIAVRGQLASRIKIVMSRLDISEKRIPQDGRMQL	295
NTDB id 1251 GC085 RS07725 WP 011946523.1	PPASLSSRITARIKIVMSNLDISERRIPQDGGFKM	313
NTDB id 1101889 Q9286 RS07775 WP 011408072.1	APVKLSQRIIAARIKIVMSQLDIAEKRPQDGRIKL	317
NTDB id 1058 ABD1 RS01615 WP 001274986.1	PPLQLANRLASRIKIVMSQMDISEKRVPPQDGRIK	308
NTDB id 1018 ACIAD RS01685 WP 004920473.1	PPLQLATRLASRIKIVMSQMDISEKRVPPQDGRIK	308
consensus	*****	

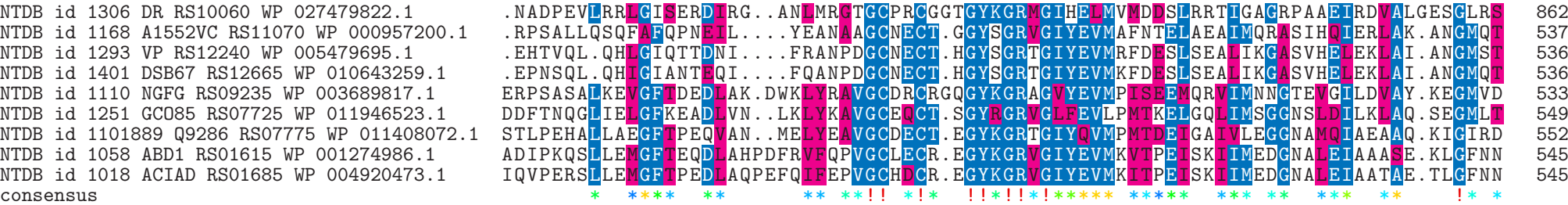
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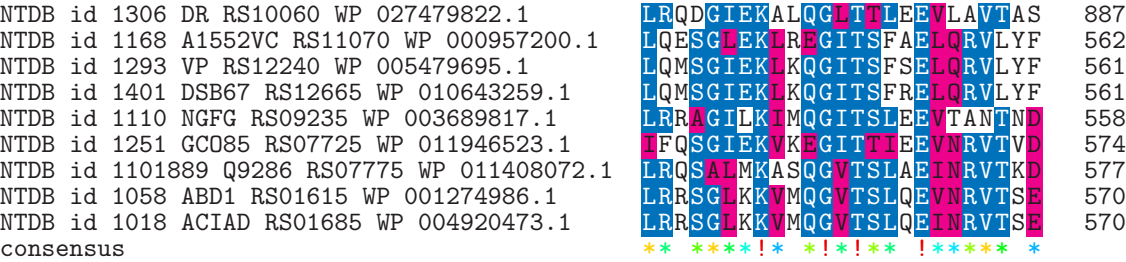
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 non conserved
 similar
 $\geq 50\%$ conserved