

logo

NTDB id 1306 DR RS10060 WP 027479822.1	MALSIGDRRLGAILLDQGYLGDNLDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	80
NTDB id 1168 A1552VC RS11070 WP 000957200.1		0
NTDB id 1293 VP RS12240 WP 005479695.1		0
NTDB id 1401 DSB67 RS12665 WP 010643259.1		0
NTDB id 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1251 GCO85 RS07725 WP 011946523.1		0
NTDB id 404654 DX0091 RS15310 WP 011408072.1		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 1306 DR RS10060 WP 027479822.1	SIRPQTALNLQAFPPFALEGDRRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIIVVPSGASDA	160
NTDB id 1168 A1552VC RS11070 WP 000957200.1		0
NTDB id 1293 VP RS12240 WP 005479695.1		0
NTDB id 1401 DSB67 RS12665 WP 010643259.1		0
NTDB id 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1251 GCO85 RS07725 WP 011946523.1		0
NTDB id 404654 DX0091 RS15310 WP 011408072.1		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 1306 DR RS10060 WP 027479822.1	GRTGGKLGGERLITHGYITDAQLQVALDAQQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSCGFHPGEEVLGSM	240
NTDB id 1168 A1552VC RS11070 WP 000957200.1		0
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NTDB id 1401 DSB67 RS12665 WP 010643259.1		0
NTDB id 1110 NGFG RS09235 WP 003689817.1		0
NTDB id 1251 GCO85 RS07725 WP 011946523.1		0
NTDB id 404654 DX0091 RS15310 WP 011408072.1		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 1306 DR RS10060 WP 027479822.1	LRADALRLLAVPVDETEQGVTVLTSDPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPQGRIGEQLVQECTLSRDQL	320
NTDB id 1168 A1552VC RS11070 WP 000957200.1		19
NTDB id 1293 VP RS12240 WP 005479695.1		19
NTDB id 1401 DSB67 RS12665 WP 010643259.1		19
NTDB id 1110 NGFG RS09235 WP 003689817.1		19
NTDB id 1251 GCO85 RS07725 WP 011946523.1		27
NTDB id 404654 DX0091 RS15310 WP 011408072.1		29
NTDB id 1058 ABD1 RS01615 WP 001274986.1		27
NTDB id 1018 ACIAD RS01685 WP 004920473.1		27
consensus		

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NTDB id 1306 DR RS10060 WP 027479822.1	REALQVQARECKVKPLGCEVITELGCFASPDEVD	400
NTDB id 1168 A1552VC RS11070 WP 000957200.1	EQAVVTQVSASGTSVPEALLSLSIFHAQELTEQLSHIFGPETDLSRYDY	69
NTDB id 1293 VP RS12240 WP 005479695.1	EESLIEQVKKASGISMPEALLSSGFFTSSELAEHLSSIFGSLSQPEYSQVEY	69
NTDB id 1401 DSB67 RS12665 WP 010643259.1	EESLIEQVKKASGISMPEALLSSGFFQSHLTHSSVFCISCTSLNQVEY	69
NTDB id 1110 NGFG RS09235 WP 003689817.1	AEHYYNESQA..GKEVLPLMFLSDGVI..SPKSLAALIARVFSYSILDLRHYP	68
NTDB id 1251 GC085 RS07725 WP 011946523.1	AIELHKLAAAEKISLLQYIVKNKILSAEQIALTAQAQNFVPMIDINCIDV	77
NTDB id 404654 DX0091 RS15310 WP 011408072.1	ARTAMEQAAA..AKVPLPQWFAENRLVSASQLAAANA..VEFCMPLMDVSVFDA	79
NTDB id 1058 ABD1 RS01615 WP 001274986.1	MQTALDAAKRAKQDVTAYL..EEVHLSPSLAETISAEFAEPYFDLDVYDT	77
NTDB id 1018 ACIAD RS01685 WP 004920473.1	MQQALEKAKKFQKDTPVPLYIDNFS..SPLTTABEISLEFGPEPLDLGVFDP	77
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NTDB id 1306 DR RS10060 WP 027479822.1	DPKVALMIPETARRVYVVPVRLQGNLSLVVAMKDPRNVFALDDLKLTITGKEILPAVMAEKDITIRITERYFGGKGFKEKLNK	480
NTDB id 1168 A1552VC RS11070 WP 000957200.1	ANLCQQQLGLRELITRYDALPIAKQCENLALLAVSDPTLLQAEEEFRRFATGLQVELALADHRAALQAATIRRLYGRSISQGAANQ	149
NTDB id 1293 VP RS12240 WP 005479695.1	ASLCQQQLGLRELITRHNALPLHRTPTSTLLLAVADPTNQQAEDDFRFATGLQVELVLADFRELSTAIRRLYGRSLSHEKSG	149
NTDB id 1401 DSB67 RS12665 WP 010643259.1	ASLCQTLGLRDLITRHNALPLNRTSSTLILAVADPTNIQAEDDFRFATGLQVELVLADFRELTAATIRRLYGRSLGQKEKSG	149
NTDB id 1110 NGFG RS09235 WP 003689817.1	HRVLMGVLTETEQMVETHCVPVFRGDKVFFAVSDPTQMPQITQKTVSAACTAVELVIVEDDQLAGLLDWVGRSSTSLQLQEL	148
NTDB id 1251 GC085 RS07725 WP 011946523.1	GTIPVNLVNEKLIKRRHAMVPLFSRGTNLYLATDDPSKQASLKEITQFHTGLNTHAIVVETDKLSALIDNLLTAKESQGLSE	157
NTDB id 404654 DX0091 RS15310 WP 011408072.1	NQNAVKLIVSEELIQKYHVLPLFKRCNRLFVGVSNPTQTRALDDIKFHTNLVVEPTILVDEDQIRRTLEQWQASNAFSGSSL	159
NTDB id 1058 ABD1 RS01615 WP 001274986.1	SQIPKDLVDQKLIKLRHRLPLIQRGQILYVATSNPSNTEADDAIRFNSKLLVEPVIVEHHKLEKVLGQHFAEESSFDFND	157
NTDB id 1018 ACIAD RS01685 WP 004920473.1	ALFLKDKIDKELIQKYRIMPLVHRGHVLYVATSNPTNTEADDAIRFNSKLLVEPTIIVHDKLERLLSEHVEETHFNFD	157
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NTDB id 1306 DR RS10060 WP 027479822.1	ELA.....ERNKTQSSQEA..LSVADESALVQVVDSTIREAALQDASDIHETTEDAVKVRVRLDGALREQNS	547
NTDB id 1168 A1552VC RS11070 WP 000957200.1	GKELISQDELANLVK...VSDDELQSIEDLSQDDSPVSRFINQVLLDAVRKGASDIHFEPYENQYRIRRCDCGTLVETQQ	225
NTDB id 1293 VP RS12240 WP 005479695.1	LKEINQEELASLVD...VGADEINIEDLSQDESPPVSRFINQVLLDAVRKGASDIHFEPYEKMYRVRRLRCDCGTLVETQQ	225
NTDB id 1401 DSB67 RS12665 WP 010643259.1	LKEINQDELAGLVD...IGDEIENIEDLSQDESPPVSRFINQVLLDAVRKGASDIHFEPYENMYRVRRLRCDCGTLVETQQ	225
NTDB id 1110 NGFG RS09235 WP 003689817.1	GEGQ.E.....EESHTLYIDNEEAEDGPVPRFIHKTLSDALRSASDIHFEPYEHMARIRFRVDCGQLREVQ	215
NTDB id 1251 GC085 RS07725 WP 011946523.1	YVEDSGDLEGLETSADD...EDQDIDTATSVTDAPITVKFVNKLLDAIRQGASDIHFEPYEREYRIRVRQDCGTLHEVAT	234
NTDB id 404654 DX0091 RS15310 WP 011408072.1	GNDD.EEMGDLDVSTGDEDMGAGGDSGVDAKGDDTPVVVKFVNKVLVDAIRRGASDIHFEPYEDDYRVRIRIDGILLKNVAK	238
NTDB id 1058 ABD1 RS01615 WP 001274986.1	EE..FDLDVN..LDGPTAQEDEE..EAPQGDEAPITVKYINKLLDAIRMGASDIHFEPYEKSYRVRVYRVDGVLRLQIAN	229
NTDB id 1018 ACIAD RS01685 WP 004920473.1	EE..LDLDVE..VDPHTTDDDED..DKLKDAPITVKYINKLLDAIRMSASDIHFEPYEKSYRVRVYRVDGVLRLIAT	229
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NTDB id 1306 DR RS10060 WP 027479822.1	FPKGAAQQIMARLIKIMGHLDIAERRVPQDGRVRF..KKGSIDLDLRLSTLPTVYGEKAVMRLLQKASNIPELEQLCFSEY	625
NTDB id 1168 A1552VC RS11070 WP 000957200.1	PASHLSRRRLAARIKILSKLDIAERRVPQDGRILKLSH..DTAIDMRVSTLPTLWGEKIVLRILDSSAANLDTDKLGYNPQ	304
NTDB id 1293 VP RS12240 WP 005479695.1	PPNHLSRRLSARIKILSKLDIAERRVPQDGRILKLNQ..DTAIDMRVSTLPTLFGEKIVLRILDSSASLDDIDKLGYSEQ	304
NTDB id 1401 DSB67 RS12665 WP 010643259.1	PPSHLSRRLSARIKILSKLDIAERRVPQDGRILKLNQ..DTAIDMRVSTLPTLFGEKIVLRILDSSAALDINKLGYSDS	304
NTDB id 1110 NGFG RS09235 WP 003689817.1	PPIAVRGQLASRIKIVMSRLDISEKRIPQDGRMQLTFQKGGKPVDFRVSTLPTLFGEKIVMRILNSDAASLNDIQLGHEPF	295
NTDB id 1251 GC085 RS07725 WP 011946523.1	PPASLSSRITARIKIVMSNLDISERRIPQDGGFKMKISK..SRAIDFRVSTCPTSAGEKVMVRVLDSGAAKLGIEALGFNPV	313
NTDB id 404654 DX0091 RS15310 WP 011408072.1	APVKLSQRITARIKIVMSQLDIAEKRVPPQDGRILKLSK..TKQIDFRVSTLPTLFGEKIVLRILDGSAAKLGIDKLGYEPD	317
NTDB id 1058 ABD1 RS01615 WP 001274986.1	PPLQLANRLASRIKIVMSQMDISEKRVPPQDGRILKLSK..SKAIDFRVNSLPTLFGEKIVLRILDPSSAMLGIDALGYEED	308
NTDB id 1018 ACIAD RS01685 WP 004920473.1	PPLQLATRLASRIKIVMSQMDISEKRVPPQDGRILKLSK..SKTIDFRVNSLPTLFGEKIVLRILDPASAMLGIDALGYEPE	308
consensus	*****	

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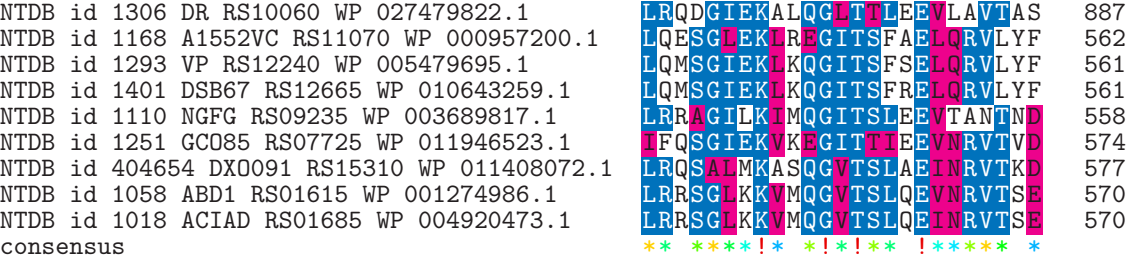
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