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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	EESLIEQVKKASGISMPEALLSSGFFTSSELAEHLSSIFGTSQPELSQVEY	69
NTDB id 1401 DSB67 RS12665 WP 010643259.1	EESLIEQVKKASGISMPEALLSSGFFQSHELTEHSSVFCISCTSLNQVEY	69
NTDB id 1110 NGFG RS09235 WP 003689817.1	AEHYYNESQA..CKEVLPLMLFSDGVI..SPKSLAALIARVFSYSILDLRHYP	68
NTDB id 1251 GC085 RS07725 WP 011946523.1	AIELHKLAAAEKISLLQYIVKNKILSAEQIALTAQAQNFVPMIDINCIDV	77
NTDB id 1102079 Q9273 RS07770 WP 011408072.1	ARTAMEQAAA..AKVPLPQWFAENRLVSASQLAAANA..VEFCMPLMDVSVFDA	79
NTDB id 1058 ABD1 RS01615 WP 001274986.1	MQTALDAAKRAKQDVTAYLIEEVHLSPSLAEETISAEFAEPYFDLDVYDT	77
NTDB id 1018 ACIAD RS01685 WP 004920473.1	MQQALEKAKKFQKDVPVYLLIDNFS..SPLTTIAETISLEFGPEPLDLGVFDP	77
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NTDB id 1306 DR RS10060 WP 027479822.1	DPKVALMIPETARRVYVVPVRLQGNLSLVVAMKDPRNVFALDDLKLTITGKEILPAVMAEKDITIRITERYFGEKGFEKLNK	480
NTDB id 1168 A1552VC RS11070 WP 000957200.1	ANLCQQQLGLRELITRYDALPIAKQCENLALLAVSDPTLLQAEEEFRRFATGLQVELALADHRAALQAATIRRLYGRSISQGAANQ	149
NTDB id 1293 VP RS12240 WP 005479695.1	ASLCQQQLGLRELITRHNALPLHRTPTSTLLLAVADPTNQQAEDDFRFATGLQVELVLADFRELESTAIRRLYGRSISHEKSG	149
NTDB id 1401 DSB67 RS12665 WP 010643259.1	ASLCQTLGLRDLITRHNALPLNRTSSTLILAVADPTNIQAEDDFRFATGLQVELVLADFRELESTAIRRLYGRSLGQEKSG	149
NTDB id 1110 NGFG RS09235 WP 003689817.1	HRVLMGVLTIEHQMVFEHCVVFFRRGDKVFFAVSDPTQMPQITQKTVSAACTAVELVIVEDDQLAGLLDWVGRSSTSLQL	148
NTDB id 1251 GC085 RS07725 WP 011946523.1	CTIPVNLVNEKLIKRRHAMVPLFSRGTNLYLATDDPSKQASLKEITQFHTGLNTHAIVVETDKLSALIDNLLTAKESQGLSE	157
NTDB id 1102079 Q9273 RS07770 WP 011408072.1	NQNAVKLVSEELIQKYHVLPPLFKRCNRLFVGVSNPTQITRAIDDIKFHTNLVVEPTILVDEDQIRRTLEQWQASNAFSGSSL	159
NTDB id 1058 ABD1 RS01615 WP 001274986.1	SQIPKDLVDQKLIKLRHILPLIQRGQILYVATSNPSNTEADDAIRFNSKLLVEPVIVEHHKLEKVLGQHFAEESSFDFND	157
NTDB id 1018 ACIAD RS01685 WP 004920473.1	ALFLKDKIDKELIQKYRIMPLVHRGHVLYVATSNPTNTEADDAIRFNSKLLVEPTILVHDKLERLLSEHVEETHFNFD	157
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NTDB id 1306 DR RS10060 WP 027479822.1	ELA.....ERNKTQQSQEADLSVADESALVQVVDSTIREAALQDASDIHETTEDAVKVRVRLDGALREQNS	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.....EESHTLYIDNEEAEDGPVPRFIHKTLSDALRSASDIHFEPYEHMARTIRVDCGQLREVQ	215
NTDB id 1251 GC085 RS07725 WP 011946523.1	YVEDSGDLEGLETSADD...EDQDIDTATSVTDAPITVKFVNKLLDAIRQGASDIHFEPYEREYRIRVRCDCGTLHEVAT	234
NTDB id 1102079 Q9273 RS07770 WP 011408072.1	GNDD.EEMGDLDVSTGDEDMGAGGDSGVDAKGDDTPVVVKFVNKVLVDAIRRGASDIHFEPYEDDYRVRIRIDCGILKNVAK	238
NTDB id 1058 ABD1 RS01615 WP 001274986.1	EE.FDLDVN..LDGPTAQEDDEEAPQGDAPITVKYINKLLDAIRMGASDIHFEPYEKSYRVRVYRDCGVLRLQIAN	229
NTDB id 1018 ACIAD RS01685 WP 004920473.1	EE.LDLDVE..VDPHTTDDDEDKCLKDEAPITVKYINKLLDAIRMSASDIHFEPYEKSYRVRVYRDCGVLRLIAT	229
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NTDB id 1306 DR RS10060 WP 027479822.1	FPKGAAQQIMARLIKIMGHLDIAERRVPQDGRVRF..KKSIDLDLRLSTLPTVYGEKAVMRLLQKASNIPELEQLCFSEY	625
NTDB id 1168 A1552VC RS11070 WP 000957200.1	PASHLSRRRLAARIKILSKLDIAERRVPQDGRILKLSH..DTAIDMRVSTLPTLWGEKIVLRILDSSAANLIDTKLGYNPQ	304
NTDB id 1293 VP RS12240 WP 005479695.1	PPNHLSRRLSARIKILSKLDIAERRVPQDGRILKLNQ..DTAIDMRVSTLPTLFGEKIVLRILDSSASLIDIDKLGYSEQ	304
NTDB id 1401 DSB67 RS12665 WP 010643259.1	PPSHLSRRLSARIKILSKLDIAERRVPQDGRILKLNQ..DTAIDMRVSTLPTLFGEKIVLRILDSSAALDINKLGYSDS	304
NTDB id 1110 NGFG RS09235 WP 003689817.1	PPIAVRGQLASRIKIVMSRLDISEKRIPQDGRMLTFQKGGKPVDFRVSTLPTLFGEKIVMRILNSDAASLNDIQLGHEPF	295
NTDB id 1251 GC085 RS07725 WP 011946523.1	PPASLSSRITARIKIVMSNLDISERRIPQDGGFKMKISK..SRAIDFRVSTCPTSAGEKVMVRVLDSGAAKLGIEALGFNPV	313
NTDB id 1102079 Q9273 RS07770 WP 011408072.1	APVKLSQRITARIKIVMSQLDIAEKRVPPQDGRILKLSK..TKQIDFRVSTLPTLFGEKIVLRILDGSAAKLGIDKLGYEPD	317
NTDB id 1058 ABD1 RS01615 WP 001274986.1	PPLQLANRLASRIKIVMSQMDISEKRVPPQDGRILKLSK..SKAIDFRVNSLPTLFGEKIVLRILDPSSAMLGIDALGYEED	308
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consensus	*****	

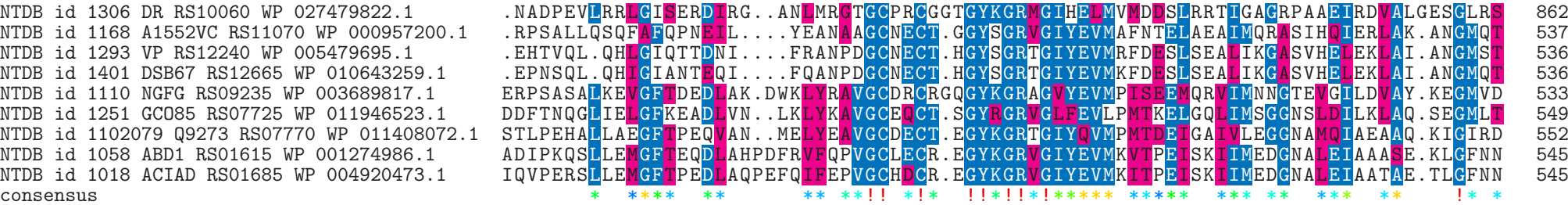
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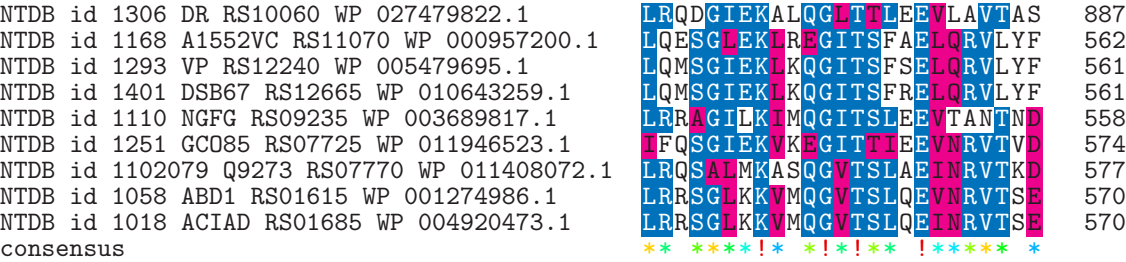
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