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NTDB id 1410 SGL RS13950 WP 010873519.1	MALSIIGDRRLGAILLDQGYLGDNLDLRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	0
NTDB id 1306 DR RS10060 WP 027479822.1	MALSIGDRRLGAILLDQGYLGDNLDLRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	80
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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
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NTDB id 409593 GNX18 RS00650 WP 166214794.1		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 1410 SGL RS13950 WP 010873519.1	SIRPQTALNLQAFPPFALEGDRRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGASDA	0
NTDB id 1306 DR RS10060 WP 027479822.1	SIRPQTALNLQAFPPFALEGDRRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGASDA	160
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consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	GRTGGKLGGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVLGSM	0
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consensus		

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NTDB id 1410 SGL RS13950 WP 010873519.1	LRADALRLLAVPVDETEQGVTVLTS DPRKRPDI DAL I GRPVQLMLTRPRDI ER L I EQFYPPQRGR LGEQLVQEGT LSRSSS	6
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NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		0

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NTDB id 1410 SGL RS13950 WP 010873519.1	SKKSRAVALRNYFSPFGNKL VAGGHVDAEQ L RQAL VQV KKTGRSLPEELKAVTGRELSPELLRQYKKNQLFELKVLYGV	86
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NTDB id 1168 A1552VC RS11070 WP 000957200.1	MLTNLVAILRQAELISATQEQAVVTQVSASGTSVPEALLE..LSIFHAQELTEQ.....LSHIFGLP	60
NTDB id 1293 VP RS12240 WP 005479695.1	MHSNLSLTLRQKGLLTFSCQ EESLIEQVKASGISMPEALLS..SGFFTSS ELAEH.....LSSIFGLS	60
NTDB id 1401 DSB67 RS12665 WP 010643259.1	MLTNLSTVLRQAGILTFSCQ EESLIEQVKASGISMPEALLS..SGFFQSHELTEH.....LSSVFGLS	60
NTDB id 1251 GC085 RS07725 WP 011946523.1	...MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLAAAEKISLLQYIVK..NKILSAEQIALT.....AAQNFGVP	68
NTDB id 409593 GNX18 RS00650 WP 166214794.1	MSSTPLSGLAKRLVIDNAIDEATAQSAVQA AKRERQSFAQHAVE..SKLIK SRELATI.....ASTAFGSP	64
NTDB id 1058 ABD1 RS01615 WP 001274986.1	...MSALHTSPKFTGFFRRLVEEKHVSAA TMQTALDA AKRAKQDTVAYLIE..EVHLSPSLLAET.....ISAEFAEP	68
NTDB id 1018 ACIAD RS01685 WP 004920473.1	...MSAFTTPPKFSGFIRRLVEEGYVNAQNMQQALEKAKKFKQDTPVPLID..NFSISPLTTAEI.....ISLEFGEP	68
consensus	* * *** ** * * * * * ** ** ** ** ** **	

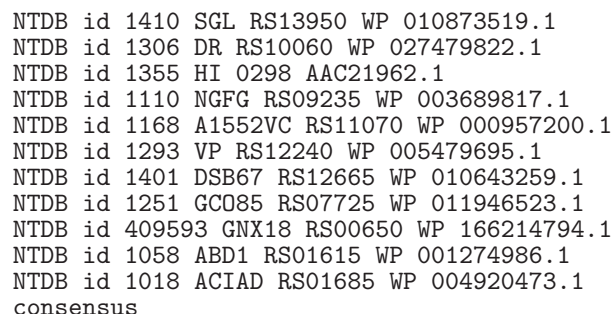
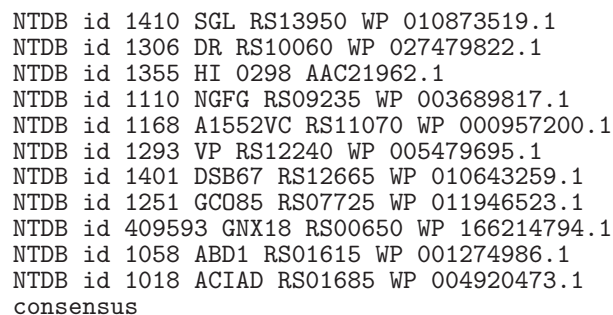
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	YFDLDVYDTSQ.....IPKDLVDQKLILKHRILPLIQRG...QILYVATSNPSNIEAIDAIRF...NSKLLVEPVIVEHH	137
NTDB id 1018 ACIAD RS01685 WP 004920473.1	LLDLGVFDPAI.....FLDKDIDEKLIQKYRIMPLVHRG...HVLVYATSNPTNIEAMDAIRF...NSKLLKVEPIIVEHD	137
consensus	** ** * * ** ** *! * * ***** ***** ***** ***** ***** *****	

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consensus

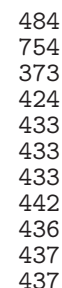
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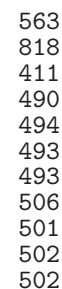
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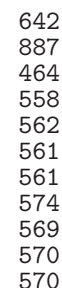
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consensus



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consensus



logo

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NTDB id 1018 ACIAD RS01685 WP 004920473.1		570
consensus		

-  non conserved
-  similar
-  ≥ 50% conserved