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				M S X L I C D R L G A L L D R G L L D E L Q R A L E R H R E V G G R L A E V L D S G M V G E K R I A R A I E E A L G I P L V N L L A V Q P D P A L	
NTDB id 1306	DR	RS10060	WP 027479822.1	.MALSIGDRRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLDISGMVGEKRIARAIEEALGIPLVNLAVQPDPAAL	79
NTDB id 1028	TT	RS08215	WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	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 1123908	EL297	RS00195	WP 002221967.1	0
NTDB id 1110	NGFG	RS09235	WP 003689817.1	0
NTDB id 1251	GCO85	RS07725	WP 011946523.1	0
NTDB id 1018	ACIAD	RS01685	WP 004920473.1	0
NTDB id 1058	ABD1	RS01615	WP 001274986.1	0
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				R S I R P Q T A L N L Q A F P F A L E E A G V V R V A F L N P L D T L S L E E V E D L T G L V V E P Y Q T T K S A F L Y A L A K H Y P E L G L P V P P P P P P P	
NTDB id 1306	DR	RS10060	WP 027479822.1	RSIRPQTALNLQAFPPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGA	157
NTDB id 1028	TT	RS08215	WP 011173991.1	ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
NTDB id 1168	A1552VC	RS11070	WP 000957200.1	0
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NTDB id 1110	NGFG	RS09235	WP 003689817.1	0
NTDB id 1251	GCO85	RS07725	WP 011946523.1	0
NTDB id 1018	ACIAD	RS01685	WP 004920473.1	0
NTDB id 1058	ABD1	RS01615	WP 001274986.1	0
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				S D A G R T G G K L G E R L I T H G Y I T D A Q L Q V A L D A Q Q T G E A L G A T L I S Q R A I T E D Q L Y E V L A E Q E G T T F L P N P S G F H P G E E V L	
NTDB id 1306	DR	RS10060	WP 027479822.1	SDAGRTGGKLGRLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEEQEGTTFLPNPSGFHPGEEVL	237
NTDB id 1028	TT	RS08215	WP 011173991.1	SGEGQKDLKLGELLQKQGWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAA	237
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NTDB id 1110	NGFG	RS09235	WP 003689817.1	0
NTDB id 1251	GCO85	RS07725	WP 011946523.1	0
NTDB id 1018	ACIAD	RS01685	WP 004920473.1	0
NTDB id 1058	ABD1	RS01615	WP 001274986.1	0
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NTDB id 1306 DR RS10060 WP 027479822.1	0
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NTDB id 1110 NGFG RS09235 WP 003689817.1	0
NTDB id 1251 GCO85 RS07725 WP 011946523.1	0
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NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
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NTDB id 1306 DR RS10060 WP 027479822.1	DQLREALQVQAREGKVKPLGEVITELCFASPDVDSALQKQNVCCGRIEDTLVQSCKLRPEALARSAAQLGYEFLDPIQ	397
NTDB id 1028 TT RS08215 WP 011173991.1	EALKEALVQKGLPRAKPLGELVLELGLARPEDVDEALQKQRRCGRIEDTLVQSCKLRPEALAGAVATQIGYPVVDPEE	397
NTDB id 1168 A1552VC RS11070 WP 000957200.1MLTNLVAIIIRQAELISATGEAVVTQVSASGTSVPEALLELSIFHAQELTEQLSHIFGLPETDLR	66
NTDB id 1293 VP RS12240 WP 005479695.1MHSNLSITLRQKGLLTFSSQESLIEGVKASCISMPEALLSSCFFTSSELAPHLSSTFGLSQPELSQ	66
NTDB id 1401 DSB67 RS12665 WP 010643259.1MLTNLSTVLRQAGILTFSQEBLLIEGVKASCISMPEALLSSCFFQSHETHELSSVFGLSCTSLNQ	66
NTDB id 1123908 EL297 RS00195 WP 002221967.1MSVGLLRILVQNVVIVEQAHHY.NESQACKEVLPMLFSDCVISPKSLAALIRVFSYSILDIRH	65
NTDB id 1110 NGFG RS09235 WP 003689817.1MSVGLLRILVQNVVIVEQAHHY.NESQACKEVLPMLFSDCVISPKSLAALIRVFSYSILDIRH	65
NTDB id 1251 GCO85 RS07725 WP 011946523.1MALATEEYRLQCTGGQLVLEKLLDKTKATELHKLAAAEKISLLQYIVKNIKLSAEQIALTAAQNFVGMPLDINC	74
NTDB id 1018 ACIAD RS01685 WP 004920473.1MSAFTTPPKFSGFIRRLVEEGYVNAQNMQALAKKFKQDIVPYLDNFSTISPLTIAETISLEFGEPILLDIGV	74
NTDB id 1058 ABD1 RS01615 WP 001274986.1MSALHTSPKFTGFFRRLVEEKHVSAAATMTALDAAKRAKQDTWAYLIEEVHLSPSLLAETISAFAEPYFDLIV	74
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NTDB id 1306 DR RS10060 WP 027479822.1	NPPDPKVALMPEATARRVVVVRRLQGNLSVAMKDPKRVFALDDKLLITG.....KEILPAVMAEKDIRLIERYFGE	472
NTDB id 1028 TT RS08215 WP 011173991.1	DPPDPGAPLLLPEDLCRRYGVVPHRLEGNRLVLLMKDPRNIALDDVRLALKRKGLNVEVAPAVATEAAITKLIBFYGK	477
NTDB id 1168 A1552VC RS11070 WP 000957200.1	YDYANLCQQLGLRELITRYDALPTAKQCMLLLAVSDPTLLQAEDEFRFATG.....LQVELALADHRALQAIRRLYGR	141
NTDB id 1293 VP RS12240 WP 005479695.1	YEYASICQQLGLRELITRHNAIPLHRTPTSTLLAVADPTNQAEDDFRFATG.....LQVELVLADEFRELTAIRRLYGR	141
NTDB id 1401 DSB67 RS12665 WP 010643259.1	YEYASICQQLGLRDLITRHNAIPLNRTSSTLLAVADPTNQAEDDFRFATG.....LQVELVLADEFRELTAIRRLYGR	141
NTDB id 1123908 EL297 RS00195 WP 002221967.1	YPRHRVLMGVLTTEEQMVEFHCVVFRGDKVFFAVSDPTQMPQIQKTVSAAG.....TEVELVIVEDDQLAGLLDWGSR	140
NTDB id 1110 NGFG RS09235 WP 003689817.1	YPRHRVLMGVLTTEEQMVEFHCVVFRGDKVFFAVSDPTQMPQIQKTVSAAG.....TAVELVIVEDDQLAGLLDWGSR	140
NTDB id 1251 GCO85 RS07725 WP 011946523.1	IDVGTIPVNLVNEKLIKRHAMVPLFSRGTNLYLATDDPSKQASIKETQFHTG.....LNTHAVVETDKLSALINDLTA	149
NTDB id 1018 ACIAD RS01685 WP 004920473.1	FDPALFLKDKIDEKLIQKYRIMPLVHRGHVLYVATSNPTNTEAMDARFNISK.....LKVEPIIVEHDKLERLSEHFVE	149
NTDB id 1058 ABD1 RS01615 WP 001274986.1	YDTSQIPKDLVDQKLLKHLRILPLIQRGQILYVATSNPSNTEADARFNISK.....LLVEPVIVEHDKLERLSEHFVE	149
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