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NTDB id 1168 A1552VC RS11070 WP 000957200.1	MSALTTSPKLTGFRRVVEKLYSATIMQALAKAKQSvYLIENYLSPSTLAETISSEFGEPYLDLVYDAIT	72
NTDB id 1251 GC085 RS07725 WP 011946523.1	MLINLVAILRQAELISATQEQAIVTQVSASGTSVPEALLELSIFHAQELTEQLSHIFGLPETDLSRYDYANL	80
NTDB id 126217 AMD07 RS05710 WP 005915819.1	MALATEEYRLQGIGQLLVLEKLLDKTKAIELHKLAAAEEKISLLQYIVKNKILSAEQIALTAQAQNFVPMPLDINCIDVGTI	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	MSAFTTPPKFSGFIRRLVEEGYVNAQNMQQALEKAKKFKQDTPYPLIDNFSISPLTTAEIISLEFGPELLDLGVFDPALF	80
NTDB id 1058 ABD1 RS01615 WP 001274986.1	MSALHTSPKFTGFFRRLVEEKHVSAAATMQTALDAAKRAKQDTPVAYLIEEVHLSPSLLAETISAEFAEPYFDLDVYDTSQI	80
consensus	* *	

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NTDB id 1251 GC085 RS07725 WP 011946523.1	CQQLGLRELITRYDALPIAKQGNLLLLAVSDPTLLQAEEEFRFATGLQVELALADHRALQAAIRRLYGRSIQGAANQGKE	160
NTDB id 126217 AMD07 RS05710 WP 005915819.1	PVNLYVNEKLIKPHAMVPLFSRGTNLYLATDDPSKQASLIKEIQFHTGLNTHAIVVETDKLSALIDNLLTAKESQGLSEYVE	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	LKDKIDEKLIQKYRIMPLVHRGHVLYVATSNPTNIEAMDAIRFNSKCLKVEPTIVEHDKLERLLSEHFVEETHFNFDTE..	158
NTDB id 1058 ABD1 RS01615 WP 001274986.1	PKDLVDQKLILKHRILPLIQRGQILYVATSNPSNIEAIDAIRFNSKLLVEPVIVEHHKLEKVLGQHF AEESSFDFNDE..	158
consensus	* *	

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NTDB id 126217 AMD07 RS05710 WP 005915819.1	ELD...LDVEVDPHTTDDDEDKCLKDEAPIVKYINKLLIDAIRMSASDLHFEPYEKSYRVRVYRVDGVLRLIATPPLQL	0
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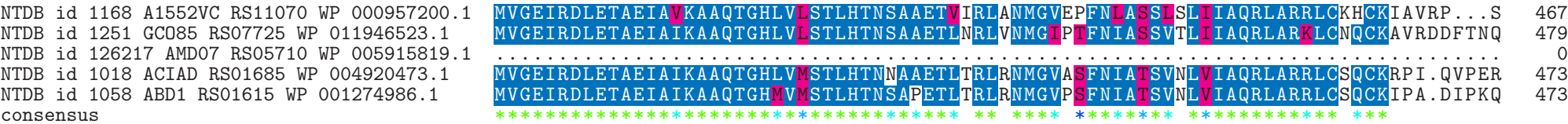
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NTDB id 126217 AMD07 RS05710 WP 005915819.1	ATRLASRLKVMSQMDISEKRVPPQDGRIKLRKMSKSKTIDFRVNSLPTLFGEKIVLRILDPASAMLGIDALGYEPEQKALFM	0
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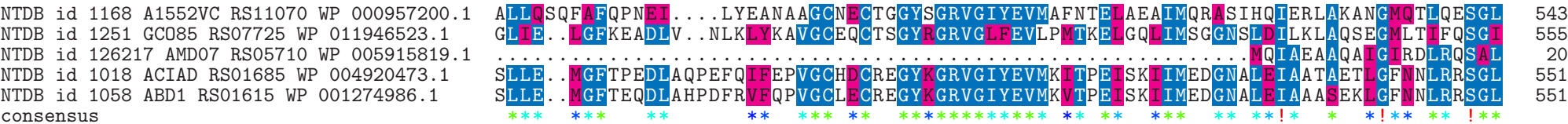
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NTDB id 126217 AMD07 RS05710 WP 005915819.1	EALNKPQGMILLITGPTGSGKTVSLYTGLNINLNTTEHANISTAEDPVEINLEGINQVNVNPKVGLTFAAALRSFLRQDPDI	0
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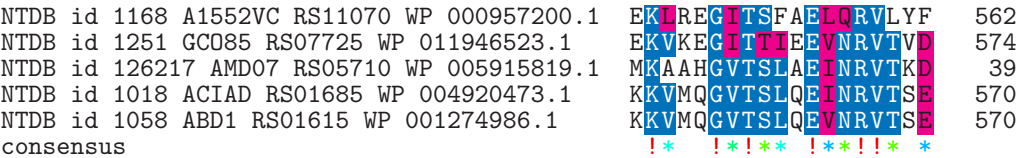
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