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NTDB id 617 LCA RS06480 WP 011375001.1	.MST	EEKLTTLLKHCAKQVRQDLYFTPTIGGWRLTERRQSDLVTNQELDKPTGTRYLNRLKYMAGMDISETRRAQTGRSE	79	
NTDB id 586 KW2 RS10855 WP 011836048.1	..	MVQKKAKELIQKAIEMEASDIYLTASGNLYKIYIRQQLGRTLIEEVNQEIGLALLTHFKFLAGMNTGERRRVQLGACW	78	
NTDB id 584 DLJ51 RS00890 WP 019777327.1	MVEVQELGKELIRQA	VEEQAQDIYLLPKPAVYKIYLRFGKRLHIQTCPPEQAAGLISHFKFLAGMNVGEKRR	AQLGACD	80
NTDB id 385 SMU RS09035 WP 002263435.1	..	MVQKLGRKIIAEAVQNKAQDIYIIPKKESYELMRIGNERRFIDVYDHHFMANLISHFKFVAGMNVGEKRRSQLGSCD	78	
NTDB id 394 FSA28 RS09405 WP 002291652.1	..	MVQKLGRKIIAEAVQNKAQDIYIIPKKESYELMRIGNERRFIDVYDHHFMANLISHFKFVAGMNVGEKRRSQLGSCD	78	
NTDB id 411838 GRI05 RS00755 WP 011921718.1	..	MIQEKARKMIEEAADRVSADIYLVPRGECYQVYHRIMDDREFVQDLAAEEVTAIISHFKFLAGLVGEKRRSQQGSCD	78	
NTDB id 429 SGO RS09420 WP 012130942.1	..	MVQKIAQAIIVRQAKEECAQDIYFVPKDDCYELMRIGDERRFIQTYDFDQLAAVISHFKFLAGMNVGEKRRSQLGSCD	78	
NTDB id 500 SM12261 RS00945 WP 000249545.1	..	MVQEIQAQEIIRSARKKGAQDIYFVPKLDAYELHMRVGDERCKIGCYDFEKF	AAVISHFKFVAGMNVGEKRRSQLGSCD	78
NTDB id 270 KZH43 RS09265 WP 000249564.1	..	MVQEIQAQEIIRSARKKGTQDIYFVPKLDAYELHMRVGDERCKIGSYDFEKF	AAVISHFKFVAGMNVGEKRRSQLGSCD	78
NTDB id 229 SPD RS09895 WP 000249564.1	..	MVQEIQAQEIIRSARKKGTQDIYFVPKLDAYELHMRVGDERCKIGSYDFEKF	AAVISHFKFVAGMNVGEKRRSQLGSCD	78
NTDB id 195 SPR RS09430 WP 000249564.1	..	MVQEIQAQEIIRSARKKGTQDIYFVPKLDAYELHMRVGDERCKIGSYDFEKF	AAVISHFKFVAGMNVGEKRRSQLGSCD	78
NTDB id 160 SP RS10380 WP 000249564.1	..	MVQEIQAQEIIRSARKKGTQDIYFVPKLDAYELHMRVGDERCKIGSYDFEKF	AAVISHFKFVAGMNVGEKRRSQLGSCD	78
consensus		***** * !!***** *****		

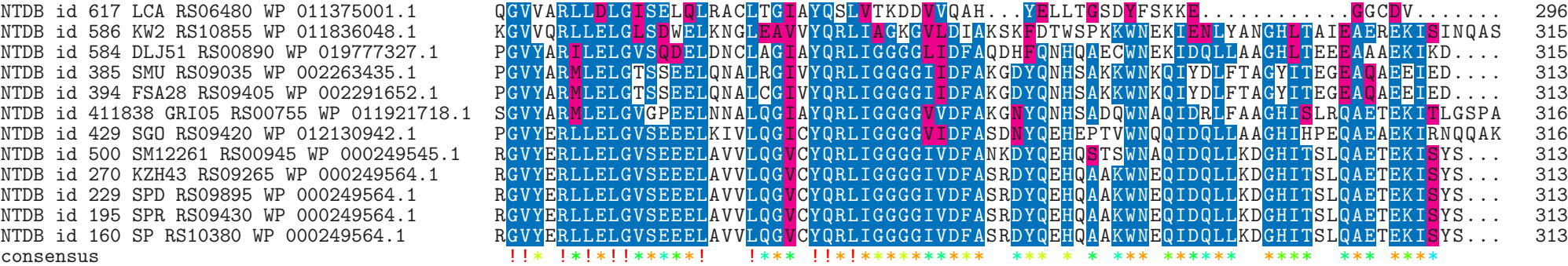
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NTDB id 617 LCA RS06480 WP 011375001.1	LQLADQ.TIYLRRLATVGDFLNRESLVIRFTYPIGAIIYHCDDQQLAQLTQMSRQSGLIILFAGPTGSGKTTSLYHLAQQSM	158
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NTDB id 385 SMU RS09035 WP 002263435.1	YSYDEGKVISLRLSVVGDRGYESLVIRLLFSGRHDLEKFWFN.SMDLILNEIKGRGLYLFSGPVGSGKTSLMYQLVKLRF	157
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NTDB id 429 SGO RS09420 WP 012130942.1	YRYDDK.ETSIRLSTVGDRGYESLVIRLLHDEETELKFWFT.HFPELREKFKDRGLYLFSGPVGSGKTTLMHQLAQLKF	156
NTDB id 500 SM12261 RS00945 WP 000249545.1	YEYDQK.MASRLSTVGDRGHESLVIRLLHDEEQDLHFWFQ.DIEELGKQYRQRGLYLFAGPVGSGKTTLMHLEAKSLF	156
NTDB id 270 KZH43 RS09265 WP 000249564.1	YAYDHK.IASRLSTVGDRGHESLVIRLLHDEEQDLHFWFQ.DIEELGKQYRQRGLYLFAGPVGSGKTTLMHLELSKSLF	156
NTDB id 229 SPD RS09895 WP 000249564.1	YAYDHK.IASRLSTVGDRGHESLVIRLLHDEEQDLHFWFQ.DIEELGKQYRQRGLYLFAGPVGSGKTTLMHLELSKSLF	156
NTDB id 195 SPR RS09430 WP 000249564.1	YAYDHK.IASRLSTVGDRGHESLVIRLLHDEEQDLHFWFQ.DIEELGKQYRQRGLYLFAGPVGSGKTTLMHLELSKSLF	156
NTDB id 160 SP RS10380 WP 000249564.1	YAYDHK.IASRLSTVGDRGHESLVIRLLHDEEQDLHFWFQ.DIEELGKQYRQRGLYLFAGPVGSGKTTLMHLELSKSLF	156
consensus	* ** * ** ! * * ! ! * * * ! ! ! ! ! * * * * * * * * * * * * * * ! * * * * * ! * * *	

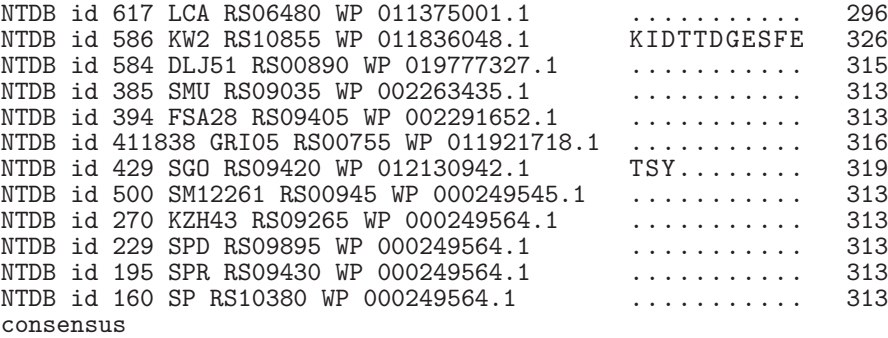
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NTDB id 617 LCA RS06480 WP 011375001.1	SAKMVVAIEDPIEIVAPEFLQLQVNDNAGLSYAELLKISLRLRPDTLIIGEIRDLETAQYVVSAALS	GHVLVSTIHARST	238
NTDB id 586 KW2 RS10855 WP 011836048.1	SNAQVITIEEPVELIDSDFIQLQVNDVIGNSYDELIKLSLRHRPDLLIIGEIRDQQTARAVLRASLTGY	TVFSTIHAASV	235
NTDB id 584 DLJ51 RS00890 WP 019777327.1	PDSQIITIEDPVEIKQDNMLQLQLNPSIGMTYDHLIKLSLRHRPDVLIIGEIRDRETAQAVIRASLT	GIRVFSTVHAKSI	239
NTDB id 385 SMU RS09035 WP 002263435.1	ETKQVITIEDPVEIKQDNMLQLQLNESIDMTYDNLIKLSLRHRPDILIIGEIRDKETARAVIRASLT	GAMVFSTIHAKSI	237
NTDB id 394 FSA28 RS09405 WP 002291652.1	ETKQVITIEDPVEIKQDNMLQLQLNESIDMTYDNLIKLSLRHRPDILIIGEIRDKETARAVIRASLT	GAMVFSTIHAKSI	237
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consensus	* **** ! *		

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non conserved
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 ≥ 50% conserved