

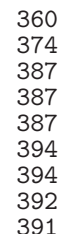
NTDB id	85	BSU 00860 NP 387967.1	..MMFGRFTEAAQKVLALAQEEALRLGHNNIGTEHILLGLVREGEGTAAKALQALGLGSEKIQKEVESLIGRG.....	71
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NTDB id	287	SP RS11210 WP 001109712.1	MNYSKALNECIESAYMVAGHFAGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLEE.....VALELTETDYS	70
NTDB id	290	KZH43 RS10025 WP 001109677.1	MNYSKALNECIESAYMVAGHFAGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLEE.....VALELTETDYS	70
NTDB id	289	SPD RS10700 WP 001109677.1	MNYSKALNECIESAYMVAGHFAGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLEE.....VALELTETDYS	70
NTDB id	329	STU RS10020 WP 011225298.1	MTIYSRKMQAIFHRAQLAEARFESPFLFTWHVLLAMVEVPGSVAYLTFTDFEDRIHSEEIETAAVLAAMEKRPKDLS	76
NTDB id	297	STER RS00545 WP 011680614.1	...MTIYSRKMQAIFHRAQLAEARFESPFLFTWHVLLAMVEVTGSVAYLTFTDFEDRIHSEEIETAAVLAAMEKRPKDLS	76
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NTDB id	377	SMU RS09275 WP 002262344.1	MTDYSLMKQEVFRLLAQFEAARFESSYLESWHVLLAMVEIDSVAAGLSFAEFEAQDVRFEDIYQAAAIIAIGKKPKSSS	76
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NTDB id 287 SP RS11210 WP 001109712.1	QDETFTTELPSRRLLQVLFD EAEYVASVVHAQVLTGETHVLYAILHDGNALATRILERAGFSSYEDKKDQ.VKIIAALRRNLEE	149
NTDB id 290 KZH43 RS10025 WP 001109677.1	QDETFTTELPSRRLLQILFDEAEY VASVVHAQVLTGETHVLYAILHDGNALATRILERAGFSSYEDKKDQ.VKIIAALRRNLEE	149
NTDB id 289 SPD RS10700 WP 001109677.1	QDETFTTELPSRRLLQILFDEAEY VASVVHAQVLTGETHVLYAILHDGNALATRILERAGFSSYEDKKDQ.VKIIAALRRNLEE	149
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NTDB id 297	STER RS00545 WP 011680614.1	NAGLSKQDLK.	AIHD LRKP KKS KASANF ANMM QPPQSSTGELADYTKDLTALAESGNL D PVIGRDEEISRM IQVLSRKT	234
NTDB id 232127	CCX85 RS08540 WP 002991294.1	HACYSKEMIK.	AIHEL RKPK PTK TQGTFSDMMKPP.STAGELSDFTRDLTEMARQGLLESVIGRDQEVSRMIQVLSRKT	232
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consensus		!	*!*****!* * * * *	

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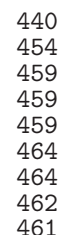
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consensus
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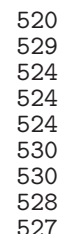
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