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NTDB id 1009 ACIAD RS15175 WP 004923730.1	MNLYEYRRQFSMGAKIYVAIAIMQ.....VASAQVAITNVVPMNIPGQGTEIRVMFNGLPP.QPQAYQLEQPARLILDFDKA	71
NTDB id 1039 HON27 RS01460 WP 031993779.1	MNHVFRQFSMGAVAIAIMQ.....AASAQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRLILDFDKA	71
NTDB id 1052 ABD1 RS15850 WP 001017033.1	MNHVFRQFSMGAVAIAIMQ.....AASAQVSMTNIVPMQIAGQGTEIRVMFNGLPP.QPQAYQLENPSRLILDFDKA	71
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NTDB id 1175 A1552VC RS12105 WP 000788426.1		0
NTDB id 1408 DSB67 RS13980 WP 010643500.1		0
NTDB id 1284 ACE17W RS12370 WP 010946666.1	MR.....KIITAILILISASLSMAYSKDNSLISVKVLPLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDFIST	70
NTDB id 1257 GC085 RS04855 WP 163621000.1	MR.....KIITAILILISASLSLAYSKDNSLISVKVLPLPENRVRIDFQFSKPLKQLPASFITQKPPRLVLDFIST	70
NTDB id 93423 AK972 RS28940 WP 063033466.1		0
NTDB id 1193 PA5040 NP 253727.1	MNSGLSR.....LGIALLAAMF....APALLAADLEKLDVAALPGDRVELKLQFDEPVAA.PRGYTIEQPARTALDLPGV	70
consensus	***** * * * * *	

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NTDB id 1009 ACIAD RS15175 WP 004923730.1	QQQLPSSTIYQINEASSVDVSDDDRRLIYNLKRSLYTRVEGNELVYNLAKTANKP	133
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NTDB id 1052 ABD1 RS15850 WP 001017033.1	QQGLKQ..SKISVATNEASSVDVTSDDQRSRLTVNLKDTGAFTTRVEGNTFILKINSAQTSNKP.....	133
NTDB id 1186 GTF74 RS01320 WP 000788426.1	QQGLKQ..SKISVATNEASSVDVTSDDQRSRLTVNLKDA GAFTTRVEGNTFILKINSAQTSNKP.....	25
NTDB id 1175 A1552VC RS12105 WP 000788426.1	MKNGLKT..YVAQTWLTLWVGLALCA.....S.	25
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NTDB id 1257 GC085 RS04855 WP 163621000.1	DMQIPSDQKTKTITQLGSLHSYNIVAVGDRVRAILELDRSVSYSGSTAGSIYSLVLNGKSNDL.F.....ENSK	137
NTDB id 93423 AK972 RS28940 WP 063033466.1	DMQIPSDQKTKTITQLGSLHSYNIVAVGDRVRAILELDRSVSYSGSTAGSVYSLVLNGKSNDL.F.....ENSK	0
NTDB id 1193 PA5040 NP 253727.1	QNKLI..GTKNRELISVGNTRSVTVVEAKDRTRLIINLTALSSYTTTRVEGNLFFVVGNSPAGASVASAAPVKASPAPASYA	148
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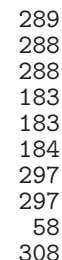
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NTDB id 1257 GC085 RS04855 WP 163621000.1	EVFITNQVVNAKHEIKRIDFRGIEKQGGRLVIDVSDTGIPIDVTQVGKEIVVNFLNTKIPLNLMKRYDVSDFHSPVQIIT	217
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NTDB id 1193 PA5040 NP 253727.1	QPIKPKPYVPAGRAIRNIDFQRGEKGEGNVVIDLSDPTLSPDIQEQQGKIRLIDFAKTQLPDA LRVRLDVKDFATPVQFVN	228
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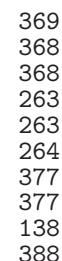
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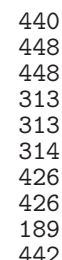
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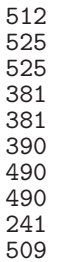
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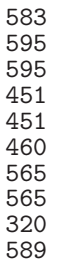
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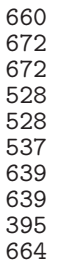
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