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NTDB id 1307 DR RS10055 WP 010888596.1	TIDRIIDVFPAAEELMVRSMSESQAIVISQITLLKKNKGGRRVAAHEIMIGTPAIRNLI	REKVAQMYSAIQTGANHG	319
NTDB id 1283 ACE17W RS02630 WP 010947729.1	TIDRIIDVFPAAEELMVRSMSESQAIVISQITLLKKNKGGRRVAAHEIMIGTPAIRNLI	REKVAQMYSAIQTGANHG	314
NTDB id 1259 GC085 RS10305 WP 010947729.1	TIDRIIDVFPAAEELMVRSMSESQAIVISQITLLKKNKGGRRVAAHEIMIGTPAIRNLI	REKVAQMYSAIQTGANHG	314
NTDB id 1194 PAKAF RS01995 WP 003084552.1	TIDRIIDVFPAAEELMVRSMSESQAIVISQITLLKKNKGGRRVAAHEIMIGTPAIRNLI	REKVAQMYSAIQTGANHG	314
NTDB id 1202 PSJM300 01600 AFN76400.1	TIDRIIDVFPAAEELMVRSMSESQAIVISQITLLKKNKGGRRVAAHEIMIGTPAIRNLI	REKVAQMYSAIQTGANHG	314
NTDB id 1020 ACIAD RS04210 WP 011182164.1	TIDRIIDVFPAAEELMVRSMSESQAIVISQITLLKKNKGGRRVAAHEIMIGTPAIRNLI	REKVAQMYSAIQTGANHG	314
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	TIDRIIDVFPAAEELMVRSMSESQAIVISQITLLKKNKGGRRVAAHEIMIGTPAIRNLI	REKVAQMYSAIQTGANHG	314
NTDB id 1060 ABD1 RS04380 WP 000355489.1	TIDRIIDVFPAAEELMVRSMSESQAIVISQITLLKKNKGGRRVAAHEIMIGTPAIRNLI	REKVAQMYSAIQTGANHG	314
NTDB id 1040 HON27 RS13360 WP 000355489.1	TIDRIIDVFPAAEELMVRSMSESQAIVISQITLLKKNKGGRRVAAHEIMIGTPAIRNLI	REKVAQMYSAIQTGANHG	314
NTDB id 1000662 ABFU58 RS17955 WP 014506546.1	TIDRIIDVFPAAEELMVRSMSESQAIVISQITLLKKNKGGRRVAAHEIMIGTPAIRNLI	REKVAQMYSAIQTGANHG	314
NTDB id 1177 A1552VC RS01165 WP 000422572.1	TIDRIIDVFPAAEELMVRSMSESQAIVISQITLLKKNKGGRRVAAHEIMIGTPAIRNLI	REKVAQMYSAIQTGANHG	313
NTDB id 1203 PSJM300 01605 AFN76401.1	TIDRIIDVFPAAEELMVRSMSESQAIVISQITLLKKNKGGRRVAAHEIMIGTPAIRNLI	REKVAQMYSAIQTGANHG	315
NTDB id 1019 ACIAD RS04205 WP 004922051.1	TIDRIIDVFPAAEELMVRSMSESQAIVISQITLLKKNKGGRRVAAHEIMIGTPAIRNLI	REKVAQMYSAIQTGANHG	316
consensus	***!***** ** *!*!!*****!*** * ***** !*** *****!* ***** ***** !**		

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NTDB id 1307 DR RS10055 WP 010888596.1	TLDQHLTELVNKNIISKHTAHEAALNKSLF	...GASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAA	395
NTDB id 1283 ACE17W RS02630 WP 010947729.1	TLDQHLTELVNKNIISKHTAHEAALNKSLF	...GASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAA	344
NTDB id 1259 GC085 RS10305 WP 010947729.1	TLDQHLTELVNKNIISKHTAHEAALNKSLF	...GASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAA	344
NTDB id 1194 PAKAF RS01995 WP 003084552.1	TLDQHLTELVNKNIISKHTAHEAALNKSLF	...GASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAA	344
NTDB id 1202 PSJM300 01600 AFN76400.1	TLDQHLTELVNKNIISKHTAHEAALNKSLF	...GASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAA	344
NTDB id 1020 ACIAD RS04210 WP 011182164.1	TLDQHLTELVNKNIISKHTAHEAALNKSLF	...GASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAA	344
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	TLDQHLTELVNKNIISKHTAHEAALNKSLF	...GASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAA	345
NTDB id 1060 ABD1 RS04380 WP 000355489.1	TLDQHLTELVNKNIISKHTAHEAALNKSLF	...GASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAA	345
NTDB id 1040 HON27 RS13360 WP 000355489.1	TLDQHLTELVNKNIISKHTAHEAALNKSLF	...GASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAA	345
NTDB id 1000662 ABFU58 RS17955 WP 014506546.1	TLDQHLTELVNKNIISKHTAHEAALNKSLF	...GASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAA	377
NTDB id 1177 A1552VC RS01165 WP 000422572.1	TLDQHLTELVNKNIISKHTAHEAALNKSLF	...GASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAA	368
NTDB id 1203 PSJM300 01605 AFN76401.1	TLDQHLTELVNKNIISKHTAHEAALNKSLF	...GASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAA	381
NTDB id 1019 ACIAD RS04205 WP 004922051.1	TLDQHLTELVNKNIISKHTAHEAALNKSLF	...GASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAA	383
consensus	!*!* ! ** ***!* * ! * *		

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NTDB id 1307 DR RS10055 WP 010888596.1	PATPAAPAAGRSEPGRAGGFGR	418
NTDB id 1283 ACE17W RS02630 WP 010947729.1	.....	344
NTDB id 1259 GC085 RS10305 WP 010947729.1	.....	344
NTDB id 1194 PAKAF RS01995 WP 003084552.1	.....	344
NTDB id 1202 PSJM300 01600 AFN76400.1	.....	344
NTDB id 1020 ACIAD RS04210 WP 011182164.1	.....	344
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	.....	345
NTDB id 1060 ABD1 RS04380 WP 000355489.1	.....	345
NTDB id 1040 HON27 RS13360 WP 000355489.1	.....	345
NTDB id 1000662 ABFU58 RS17955 WP 014506546.1	.....	377
NTDB id 1177 A1552VC RS01165 WP 000422572.1	.....	368
NTDB id 1203 PSJM300 01605 AFN76401.1	.....	381
NTDB id 1019 ACIAD RS04205 WP 004922051.1	.....	383
consensus		

 non conserved  
 similar  
  $\geq 50\%$  conserved