

logo

NTDB id 462 GPW51 RS00385 WP 012774894.1
NTDB id 412 SMU RS09535 WP 002262393.1
NTDB id 170 SP RS09755 WP 000642718.1
NTDB id 280 KZH43 RS08670 WP 000642700.1
NTDB id 205 SPR RS08830 WP 000642700.1
NTDB id 239 SPD RS09270 WP 000642701.1
NTDB id 538 SMSK321 RS11060 WP 000642686.1
NTDB id 510 SM12261 RS08320 WP 000642690.1
NTDB id 32083 BCB4264 RS19080 WP 000990697.1
NTDB id 124 BSU 16930 NP 389575.2
consensus



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NTDB id 462 GPW51 RS00385 WP 012774894.1	LDILENQILDCQTFFEGISLRDFCYGYGEETSLSASIVVEELKRQCKTIAAAESLTAQLFQATVANFSGVSSIFEGGFVTYS	308
NTDB id 412 SMU RS09535 WP 002262393.1	LDKLEKKITITID.....NLADYFYGYGEENSLPQVFDLLKEKGKTTTAAESLTAAGLFQATVANFSGVSSIFKGGFVTYS	308
NTDB id 170 SP RS09755 WP 000642718.1	LDILENQILDCQTFFEGISLRDFCYGYGEETSLSASIVVEELKRQCKTIAAAESLTAAGLFQATVANFSGVSSIFKGGFVTYS	312
NTDB id 280 KZH43 RS08670 WP 000642700.1	LDILENQILDCQTFFEGISLRDFCYGYGEETSLSASIVVEELKRQCKTIAAAESLTAAGLFQATVANFSGVSSIFEGGFVTYS	312
NTDB id 205 SPR RS08830 WP 000642700.1	LDILENQILDCQTFFEGISLRDFCYGYGEETSLSASIVVEELKRQCKTIAAAESLTAAGLFQATVANFSGVSSIFEGGFVTYS	312
NTDB id 239 SPD RS09270 WP 000642701.1	LDILENQILDCQTFFEGISLRDFCYGYGEETSLSASIVVEELKRQCKTIAAAESLTAAGLFQATVANFSGVSSIFEGGFVTYS	312
NTDB id 538 SMSK321 RS11060 WP 000642686.1	LDILENQILGRQTFFEGISLRDLICYGYGEEASLASIVVEELKRQCKTITAAESLTAAGLFQATVADFSGASSIFKGGFVTYS	312
NTDB id 510 SM12261 RS08320 WP 000642690.1	LDILENQILGRQTFFEGISLRDLICYGYGEETSLSASIVVEELKRQCKTITSAAESLTAAGLFQATVADFSGASSIFKGGFVTYS	312
NTDB id 32083 BCB4264 RS19080 WP 000990697.1	IQHVEDLILE.....RVGEFFYGYDDQ.EFLHYKALELLKRKGLTLACAESLTGGLFGNQVTENAGVSSVFKGGVLCYH	306
NTDB id 124 BSU 16930 NP 389575.2	LKETEAVILE.....RVGEFFYGYDDQ.TSLVKEISLACKEKCTTISAAESFTGGLFSEWITDHSAGSKLEAGGVVCYT	311
consensus	*****!***!*** *****!*****!*****!*****!*****!*****!*****!*****!*****!*****	

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NTDB id 462 GPW51 RS00385 WP 012774894.1	LEEKSRMLDIPAKNLEEHGVSEFTAQKMAEQARSKTQSDFGISLTGVAGPDSLEGHVPAGTVFIGLAQDQGTEVIKVNIL	369
NTDB id 412 SMU RS09535 WP 002262393.1	LEEKSRMLDIPAKNLEEHGVSEFTAQKMAEQARSKTQSDFGISLTGVAGPDSLEGHVPAGTVFIGLAQDQGTEVIKVNIL	387
NTDB id 170 SP RS09755 WP 000642718.1	LEEKSRMLDIPAKNLEEHGVSEFTAQKMAEQARSKTQSDFGISLTGVAGPDSLEGHVPAGTVFIGLAQDQGTEVIKVNIL	391
NTDB id 280 KZH43 RS08670 WP 000642700.1	LEEKSRMLDIPAKNLEEHGVSEFTAQKMAEQARSKTQSDFGISLTGVAGPDSLEGHVPAGTVFIGLAQDQGTEVIKVNIL	391
NTDB id 205 SPR RS08830 WP 000642700.1	LEEKSRMLDIPAKNLEEHGVSEFTAQKMAEQARSKTQSDFGISLTGVAGPDSLEGHVPAGTVFIGLAQDQGTEVIKVNIL	391
NTDB id 239 SPD RS09270 WP 000642701.1	LEEKSRMLDIPAKNLEEHGVSEFTAQKMAEQARSKTQSDFGISLTGVAGPDSLEGHVPAGTVFIGLAQDQGTEVIKVNIL	391
NTDB id 538 SMSK321 RS11060 WP 000642686.1	LEEKSRMLDIPAKNLEEHGVSEFTAQKMAEQARSKTQSDFGISLTGVAGPDSLEGHVPAGTVFIGLAQDQGTEVIKVNIL	391
NTDB id 510 SM12261 RS08320 WP 000642690.1	LEEKSRMLDIPAKNLEEHGVSEFTAQKMAEQARSKTQSDFGISLTGVAGPDSLEGHVPAGTVFIGLAQDQGTEVIKVNIL	391
NTDB id 32083 BCB4264 RS19080 WP 000990697.1	NDVVKQHVLRVPEEVLHTDGAVSKECARYLAENVDVLKADIGISFTGVAGPDASEQKEPAGTVFVGLSIKDEPTVVFPDNL	386
NTDB id 124 BSU 16930 NP 389575.2	NDVVKQNVLCVKKETLDRFGAVSKECARELAKGVQKLTGSDIGISFTGVAGPDAQEGHEPCHVFICISANGKEEVHEFH	390
consensus	*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****	

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NTDB id 462 GPW51 RS00385 WP 012774894.1	GGRSRADVRHIAVMHAFNLVRKALLSD.....	393
NTDB id 412 SMU RS09535 WP 002262393.1	GGRSRADVRHIAVMHAFNLVRKALLSD.....	418
NTDB id 170 SP RS09755 WP 000642718.1	GGRSRADVRHIAVMHAFNLVRKALLSD.....	418
NTDB id 280 KZH43 RS08670 WP 000642700.1	GGRSRADVRHIAVMHAFNLVRKALLSD.....	418
NTDB id 205 SPR RS08830 WP 000642700.1	GGRSRADVRHIAVMHAFNLVRKALLSD.....	418
NTDB id 239 SPD RS09270 WP 000642701.1	GGRSRADVRHIAVMHAFNLVRKALLSD.....	418
NTDB id 538 SMSK321 RS11060 WP 000642686.1	GGRSRADVRHIAVMHAFNLVRKALLSD.....	418
NTDB id 510 SM12261 RS08320 WP 000642690.1	GGRSRADVRHIAVMHAFNLVRKALLSD.....	418
NTDB id 32083 BCB4264 RS19080 WP 000990697.1	SG.SRQQIRERTAKYCFYHLYKKLEEI....	412
NTDB id 124 BSU 16930 NP 389575.2	AG.SRTGIRKRGAKYGCCHLLKLLLEQK....	416
consensus	*!*!*!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****!*****	

- X non conserved
- X similar
- X ≥ 50% conserved