

[illegible]

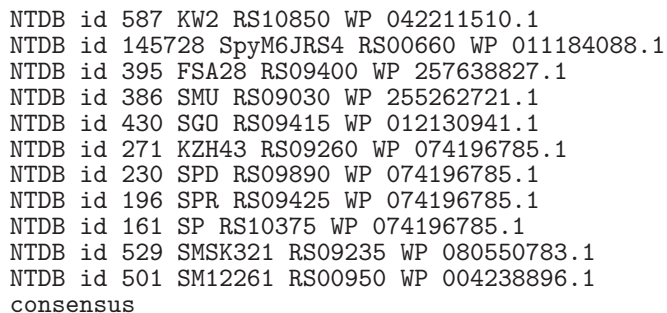
NTDB id 587 KW2 RS10850 WP 042211510.1	SFSKNVVTQLALVEVHGNLVGTMQVLVELHLKKQLKVKKKLIEVATYPVVLLVFLIGIMWGLKNYLLPQLDKGNFATLLI	156
NTDB id 145728 SpyM6JRS4 RS00660 WP 011184088.1	GFSDAILTQISLADRHGNIETTLVAIQHYLNQMARRRKTVEVITYPLILLFLFVMMLGLRRYLVPQLETQNQITYFL	159
NTDB id 395 FSA28 RS09400 WP 257638827.1	GFSDDVVTQLALSEVHGNTQKSLLKIESYLNLIILVKKKLVEVITYPTIILLIFLVVIMLGLKNYLLPQLKKG.NFATRLI	158
NTDB id 386 SMU RS09030 WP 255262721.1	GFSDDVVTQLALSEVHGNTQKSLLKIESYLNLIILVKKKLVEVITYPTIILLIFLVVIMLGLKNYLLPQLERG.NFATRLI	158
NTDB id 430 SGO RS09415 WP 012130941.1	GFSDAVVTQLSLAELHGNLSLALLKIEEYLDNLAKVKKKLIEVATYPMMLLGFLVLMIGLRNYLLPQLSSQ.NFATQLI	159
NTDB id 271 KZH43 RS09260 WP 074196785.1	GCSSAIVTQLSLAEVHGNLHLSLGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLIMLGLRNYYLLPQLDSS.NIATQII	153
NTDB id 230 SPD RS09890 WP 074196785.1	GCSSAIVTQLSLAEVHGNLHLSLGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLIMLGLRNYYLLPQLDSS.NIATQII	153
NTDB id 196 SPR RS09425 WP 074196785.1	GCSSAIVTQLSLAEVHGNLHLSLGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLIMLGLRNYYLLPQLDSS.NIATQII	153
NTDB id 161 SP RS10375 WP 074196785.1	GCSSAIVTQLSLAEVHGNLHLSLGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLIMLGLRNYYLLPQLDSS.NIATQII	153
NTDB id 529 SMSK321 RS09235 WP 080550783.1	GFSSAIVTQLSLAEVHGNLHLSLGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLSIMLGLRNYYLLPQLDSS.NIATQII	153
NTDB id 501 SM12261 RS00950 WP 004238896.1	GFSSAIVTQLSLAEVHGNLHLSLGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLIMLGLRNYYLLPQLDSS.NIATQII	153
consensus	* * * * *	

[illegible]

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NTDB id 145728 SpyM6JRS4 RS00660 WP 011184088.1
NTDB id 395 FSA28 RS09400 WP 257638827.1
NTDB id 386 SMU RS09030 WP 255262721.1
NTDB id 430 SGO RS09415 WP 012130941.1
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NTDB id 196 SPR RS09425 WP 074196785.1
NTDB id 161 SP RS10375 WP 074196785.1
NTDB id 529 SMSK321 RS09235 WP 080550783.1
NTDB id 501 SM12261 RS00950 WP 004238896.1
consensus

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- X non conserved
- X similar
- X $\geq 50\%$ conserved