

[illegible]

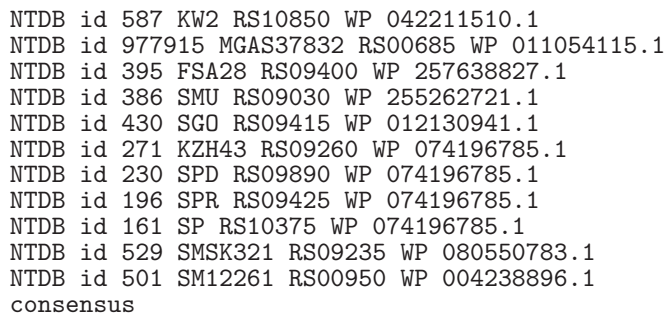
NTDB id	587 KW2 RS10850 WP 042211510.1	SFSKNVVTQLALVEVHGNLVGTMQVLVELHLKKQLKVKKKLIEVATYPVVLLVFLIGIMWGLKNYLLPQLDKGNFATLLI	156
NTDB id	977915 MGAS37832 RS00685 WP 011054115.1	GFSDAILTQISLADRHGNIETTLVAIQHYLNQMARRRKTVEVITYPLILLFLFVMMLGLRRYLVPQLETQNQITYFL	159
NTDB id	395 FSA28 RS09400 WP 257638827.1	GFSDDVVTQLALSEVHGNTQKSLLKIESYLNLIILVKKKLVEVITYPTIILLIFLVVIMLGLKNYLLPQLKKG.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 587 KW2 RS10850 WP 042211510.1
NTDB id 977915 MGAS37832 RS00685 WP 011054115.1
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 230 SPD RS09890 WP 074196785.1
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