

[illegible]

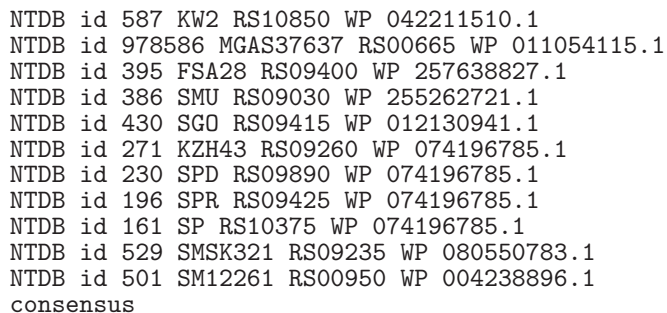
NTDB id	587	KW2	RS10850	WP	042211510.1	SFSKNVVTQLALVEVHGNLVGTMQVLVELHLKKQLKVKKKLIEVATYPVVLLVFLIGIMWGLENYLLPQLDKGNFATLLI	156
NTDB id	978586	MGA537637	RS00665	WP	011054115.1	GFSDAILTQISLADRHGNIETTLVAIQHYLNQMARIIRKTVEVITYPLILLFLFVMMLGLRRIYVPQLETQNQITYFL	159
NTDB id	395	FSA28	RS09400	WP	257638827.1	GFSDDVVTQLALSEVHGNTQKSLLKIESYLNLIILVKKKLVEVITYPTIILLIFLVVIMLGKENYLLPQLEKG.NFATRLLI	158
NTDB id	386	SMU	RS09030	WP	255262721.1	GFSDDVVTQLALSEVHGNTQKSLLKIESYLNLIILVKKKLVEVITYPTIILLIFLVVIMLGKENYLLPQLERG.NFATRLLI	158
NTDB id	430	SGO	RS09415	WP	012130941.1	GFSDAVVTQLSLAELHGNLSALLMKIEEYLDNLAKVKKKLIEVATYPMMLLGFVLIMIGLRNYLLPQLSSQ.NFATQILI	159
NTDB id	271	KZH43	RS09260	WP	074196785.1	GCSSAIVTQLSLAEVHGNLHLSLGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLLIMLGRNYYLLPQLDSS.NIATQUII	153
NTDB id	230	SPD	RS09890	WP	074196785.1	GCSSAIVTQLSLAEVHGNLHLSLGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLLIMLGRNYYLLPQLDSS.NIATQUII	153
NTDB id	196	SPR	RS09425	WP	074196785.1	GCSSAIVTQLSLAEVHGNLHLSLGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLLIMLGRNYYLLPQLDSS.NIATQUII	153
NTDB id	161	SP	RS10375	WP	074196785.1	GCSSAIVTQLSLAEVHGNLHLSLGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLLIMLGRNYYLLPQLDSS.NIATQUII	153
NTDB id	529	SMSK321	RS09235	WP	080550783.1	GFSSAIVTQLSLAEVHGNLHLSLGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLSIMLGRNYYLLPQLDSS.NIATQUII	153
NTDB id	501	SM12261	RS00950	WP	004238896.1	GFSSAIVTQLSLAEVHGNLHLSLGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLLIMLGRNYYLLPQLDSS.NIATQUII	153
consensus						* ** * * * *	

[illegible]

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NTDB id 587 KW2 RS10850 WP 042211510.1
NTDB id 978586 MGAS37637 RS00665 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
NTDB id 271 KZH43 RS09260 WP 074196785.1
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