

[illegible][illegible]

NTDB id 587	KW2 RS10850 WP 042211510.1	GALIVLLTSLSIIFKHISALINFTFLVKVPLVSSFIRLYLTAIFYSREWGNLIAQGVELRQIIELMKKQKSRIFAEGVGIKLDLGL	251
NTDB id 395	FSA28 RS09400 WP 257638827.1	CCFFILLVGIIFTFFSRKISRHLHIGFCSHLPFVGKYLQLYLTAIFYAREWGNLIQGGVEMAIQIVRLMQDQKSQLFREICYDMEKAL	253
NTDB id 386	SMU RS09030 WP 255262721.1	CCFFILLVGIIFTFFSRKISRHLHIGFCSHLPFAGKYLQLYLTAIFYAREWGNLIQGGVEMAIQIVRLMQDQKSQLFREICYDMEKAL	253
NTDB id 430	SGO RS09415 WP 012130941.1	VLMLLGLTGAIYLVFKGQKRIPVYSFLARLPFVGSFVRIYLTAIFYAREWGNMIGQGLLELSQIFQIMQEQRSVLFQEIQGDILGQAL	254
NTDB id 98622	R8623 RS09750 WP 074017570.1	VGLVSVLALLALTIFYKRSSKMSVFSILARLPFIGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMQEQQSQLFKEIGQDLAQTL	248
NTDB id 271	KZH43 RS09260 WP 074196785.1	VGLVSVLALLALTIFYKRSSKMSVFSILARLPFIGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMQEQQSQLFKEVQGDLAQTL	248
NTDB id 230	SPD RS09890 WP 074196785.1	VGLVSVLALLALTIFYKRSSKMSVFSILARLPFIGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMQEQQSQLFKEVQGDLAQTL	248
NTDB id 196	SPR RS09425 WP 074196785.1	VGLVSVLALLALTIFYKRSSKMSVFSILARLPFIGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMQEQQSQLFKEVQGDLAQTL	248
NTDB id 161	SP RS10375 WP 074196785.1	VGLVSVLALLALTIFYKRSSKMSVFSILARLPFIGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMQEQQSQLFKEVQGDLAQTL	248
NTDB id 529	SMSK321 RS09235 WP 080550783.1	VGLVSLTLLALTIFYKRSSKMRVFSILARIPFLGIFVQTYLTAYYAREWGNMISQGMELTQIFQIMQEQQSQLFKEIGQDLAQSL	248
NTDB id 501	SM12261 RS00950 WP 004238896.1	VGFVSVGVLLALTIFYKRGSKMRVFSILARIPFLGIFVQTYLTAYYAREWGNMISQGMELTQIFQIMQEQQSQLFKEIGQDMAHAL	248
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NTDB id 386 SMU RS09030 WP 255262721.1
NTDB id 430 SGO RS09415 WP 012130941.1
NTDB id 98622 R8623 RS09750 WP 074017570.1
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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[illegible]

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