

[illegible][illegible]

NTDB id 587 KW2 RS10850 WP 042211510.1	GALIVLLTSLSIIFKHHISALTNFTFLVKVPLVSSFIRLYLTAYFSREWGNLIAQGVELRQIIELMKKKQSRIFAEVGIKLDLGL	251
NTDB id 72861 C1102 RS00665 WP 011054115.1	CSSLILLFGMVWLRWRSQRRLKLYSRLSRYPFLGKLILKQYLTSYAREWGTLIGQGLDMLTILDMAIEKSSLMKELAEDIRMSL	254
NTDB id 395 FSA28 RS09400 WP 257638827.1	CCFFILLVGIPTTFFSRKISRHLHIGFCSHLPFVGKYVLQLYLTAYYAREWGNLIQGQVEMAIQIVRLMQDQKSQLFREICGYDEKAL	253
NTDB id 386 SMU RS09030 WP 255262721.1	CCFFILLVGIPTTFFSRKISRHLHIGFCSHLPFAGKYVLQLYLTAYYAREWGNLIQGQVEMAIQIVRLMQDQKSQLFREICGYDEKAL	253
NTDB id 430 SGO RS09415 WP 012130941.1	VLMMLGLTGAIYLVFKGQKRIPVYSFLARLPFVGSFVRITLYTAYYAREWGNMISQGMELTQIFQMMEQQSGQLFKEVGGDLAQTL	254
NTDB id 271 KZH43 RS09260 WP 074196785.1	VGLVSVALALLALTFYKRSKMSVFSILARLPFIGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMEQQSGQLFKEVGGDLAQTL	248
NTDB id 230 SPD RS09890 WP 074196785.1	VGLVSVALALLALTFYKRSKMSVFSILARLPFIGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMEQQSGQLFKEVGGDLAQTL	248
NTDB id 196 SPR RS09425 WP 074196785.1	VGLVSVALALLALTFYKRSKMSVFSILARLPFIGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMEQQSGQLFKEVGGDLAQTL	248
NTDB id 161 SP RS10375 WP 074196785.1	VGLVSVALALLALTFYKRSKMSVFSILARLPFIGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMEQQSGQLFKEVGGDLAQTL	248
NTDB id 529 SMSK321 RS09235 WP 080550783.1	VGLVLSLTLALLATFYKRSKMRRVFSILARIPFLGIFVQTYLTAYYAREWGNMISQGMELTQIFQIMQEQQSGQLFKETGGDLAQS	248
NTDB id 501 SM1261 RS00950 WP 004238896.1	VGFVSVGVLLALTFYKRGSKMRVFSILARIPLFGIFVQTYLTAYYAREWGNMISQGMELTQIFQIMQEQQSGQLFKETGGDMAHAL	248
consensus	***** *****	

