

[illegible][illegible]

NTDB id	587 KW2 RS10850 WP 042211510.1	NHLPVLVFFSVGALIVLLTSLSTIIIFKHISALTNFITFLVKVPLVSSFIRLYLTAYFSREWGNLIAGQVELRQTIELMKKIQK	236
NTDB id	977467 MGAS37844 RS00665 WP 011054115.1	NHFPAFFIGFCSGILILLFGMVWLRWRWSQSRLKLYSRLSRYPFGLKLLKQLYTSYYAREWGTLIGQGCLDLMTILDIMAIIEK	239
NTDB id	395 FSA28 RS09400 WP 257638827.1	SNFPFILFLLGCCFFILLVGIFTTFFSRKISRHLHLIGFCSHLPPGVGYLKYLQLYLTAYYAREWGNLIGGGVEMAQIIVRMQQDQK	238
NTDB id	386 SMU RS09030 WP 255262721.1	SNFPFILFLLGCCFFILLVGIFTTFFSRKISRHLHLIGFCSHLPPFACKYLVKYLQLYLTAYYAREWGNLIGGGVEMAQIIVRMQQDQK	238
NTDB id	430 SGO RS09415 WP 012130941.1	GHLPTIFLLTVLMMLLGTGAIVLFKGQRIPVVSFLARLPFVGSFVRILYTAYYAREWGNMGQGLELSQIFQMIMEEQR	239
NTDB id	271 KZH43 RS09260 WP 074196785.1	GNLPQIFLGMVGLVSVALLALTIFYKRSSKMRSVFSLARLPFIGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMQEQQ	233
NTDB id	230 SPD RS09890 WP 074196785.1	GNLPQIFLGMVGLVSVALLALTIFYKRSSKMRSVFSLARLPFIGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMQEQQ	233
NTDB id	196 SPR RS09425 WP 074196785.1	GNLPQIFLGMVGLVSVALLALTIFYKRSSKMRSVFSLARLPFIGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMQEQQ	233
NTDB id	161 SP RS10375 WP 074196785.1	GNLPQIFLGMVGLVSVALLALTIFYKRSSKMRSVFSLARLPFIGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMQEQQ	233
NTDB id	529 SMSK321 RS09235 WP 080550783.1	GNLPQIFLGMVGLVSLTLLALTIFYKRSSKMRFVSILARPFLGFIFVQTYLTAYYAREWGNMISQGMELTQIFQIMEEQG	233
NTDB id	501 SM12261 RS00950 WP 004238896.1	GNLPQIFLGMVGFVSVGVALLATIFYKRSSKMRFVSILARPFLGFIFVQTYLTAYYAREWGNMISQGMELTQIFQIMEEQG	233
consensus		* ** * ! * !! * **** ! ! * * * * * * ! * * * * * * * * *	

