

NTDB id 587 KW2 RS10850 WP 042211510.1	MQTDLRLRLRPKGGKKLALIKQAKLIQLMGNLLNNGFNLGVEINFLALS	SKLVEKEFVFKMHQGLSSGKSLSEILESLFSKN	81
NTDB id 31351 SEZ RS00730 WP 228275455.1	MISFLQRDISIPSRQGLKKLSSKQQEKLIIQLLYNLLSNGFNLT	TEVIAFLRKSQLLLPVYVTSMEASLLKCNGLADMLAALGYSDA	85
NTDB id 395 FSA28 RS09400 WP 257638827.1	.MTFLQQDISLKEKHKQKKLKIKKQKRV	QIQLFNNLFTSGFNLTETIIDFLERSHLLAEIYTKKMRQG	84
NTDB id 386 SMU RS09030 WP 255262721.1	.MTFLQQDISLKEKHKQKKLKIKKQKRV	QIQLFNNLFTSGFNLTETIIDFLERSHLLAEIYTKKMRQG	84
NTDB id 430 SGO RS09415 WP 012130941.1	MISFLQQDISILSKKQKQKKLGTSSKQKQV	IELFNLFSSGFHLAETVDFLGRSALLKKNYVQMRQGLANGQAFSEIMASLGFSDA	85
NTDB id 271 KZH43 RS09260 WP 074196785.1	MDISQVFRLRRKKKLATAKQKNIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQMRV	GLSQGKSFSEMME	79
NTDB id 230 SPD RS09890 WP 074196785.1	MDISQVFRLRRKKKLATAKQKNIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQMRV	GLSQGKSFSEMME	79
NTDB id 196 SPR RS09425 WP 074196785.1	MDISQVFRLRRKKKLATAKQKNIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQMRV	GLSQGKSFSEMME	79
NTDB id 161 SP RS10375 WP 074196785.1	MDISQVFRLRRKKKLATAKQKNIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQMRV	GLSQGKSFSEMME	79
NTDB id 529 SMSK321 RS09235 WP 080550783.1	MDISQVFRLRRKKKLATAKQKNIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQMRV	GLSQGKSFSEMME	79
NTDB id 501 SM12261 RS00950 WP 004238896.1	MDISQVFRLRRKKKLATAKQKNIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQMRV	GLSQGKSFSEMME	79
consensus	*****	*****	

[illegible]

NTDB id 587 KW2 RS10850 WP 042211510.1	GALIVLLTSLSIIFFKHISALTNFTFLVKVPLVSSFIRLYLTAYFSREWGNIAGQVELRQTIELMKKQKSRIFAEVGIKLDLGL	251
NTDB id 31351 SEZ RS00730 WP 228275455.1	CLLLLSLIGSFYLYARSCSRLRLFSQLSYYPITIGSLLRQYLTAYYAREWGNLIGQGLELITILEMMSNEKSQLMRELAKDIKQSL	254
NTDB id 395 FSA28 RS09400 WP 257638827.1	CCFFILLVGIPTFFSRKISRRLHLIGFCSHLPFVGKYQLYLTAAYYAREWGNLIGQGVEMAQIVRLMQDQKSQLFREIGYDMEKAL	253
NTDB id 386 SMU RS09030 WP 255262721.1	CCFFILLVGIPTFFSRKISRRLHLIGFCSHLPFAGKYQLYLTAAYYAREWGNLIGQGVEMAQIVRLMQDQKSQLFREIGYDMEKAL	253
NTDB id 430 SGO RS09415 WP 012130941.1	VLMILGLTGAIYLVFKGQKRIPVYSFLARLPFVGSFVRITYLTAYYAREWGNMIGQGLELQIFQIMQEQRSVLFQEIQQDLQCAL	254
NTDB id 271 KZH43 RS09260 WP 074196785.1	VGLVSVLALLALTFYKRSKMSVFSILARLPFIGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMEQEQSQFLFKEVGQDLAQTL	248
NTDB id 230 SPD RS09890 WP 074196785.1	VGLVSVLALLALTFYKRSKMSVFSILARLPFIGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMEQEQSQFLFKEVGQDLAQTL	248
NTDB id 196 SPR RS09425 WP 074196785.1	VGLVSVLALLALTFYKRSKMSVFSILARLPFIGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMEQEQSQFLFKEVGQDLAQTL	248
NTDB id 161 SP RS10375 WP 074196785.1	VGLVSVLALLALTFYKRSKMSVFSILARLPFIGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMEQEQSQFLFKEVGQDLAQTL	248
NTDB id 529 SMSK321 RS09235 WP 080550783.1	VGLVSLTLLALTFYKRSKMRRVFSILARIPFLGIFVQTYLTAYYAREWGNMISQGMELTQIFQIMQEQSQFLFKEIQGDLAQS	248
NTDB id 501 SM12261 RS00950 WP 004238896.1	VGFVSVGVLLALTFYKRGSKMRVFSILARIPFLGIFVQTYLTAYYAREWGNMISQGMELTQIFQIMQEQSQFLFKEIQGDMAHAL	248
consensus	***** *****	

