

NTDB id 587 KW2 RS10850 WP 042211510.1	MQTDLRLRLRPKGGKKLALIKQAKLIQLMGNLLNNGFNLGVEINFLALS	KLVEKEFVFKMHQGLSSGKSLSEILESLSFSKN	81
NTDB id 395 FSA28 RS09400 WP 257638827.1	.MTFLQQDISLKEKHKQKKLKIKKQKRV	QILFNFLFTSGFNLTETIIDFLERSHLLAEIYTKKMRQGLLNLGLHLAEIMSSSLGFSDD	84
NTDB id 386 SMU RS09030 WP 255262721.1	.MTFLQQDISLKEKHKQKKLKIKKQKRV	QILFNFLFTSGFNLTETIIDFLERSHLLAEIYTKKMRQGLLNLGLHLAEIMSSSLGFSDD	84
NTDB id 430 SGO RS09415 WP 012130941.1	MISFLQQDISILSKKQKQKKLGTSSKQKQV	IELFNFLFSSGFHLAETVDFLGRSALLEKNYVQQMRQGLANGQAFASEIMASLGFSDA	85
NTDB id 98930 R8570 RS09850 WP 077141332.1	MDISQVFRLRRKKLATAKQKNIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQMRV	GLSQGKSFSEMMESLGCSSA	79
NTDB id 271 KZH43 RS09260 WP 074196785.1	MDISQVFRLRRKKLATAKQKNIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQMRV	GLSQGKSFSEMMESLGCSSA	79
NTDB id 230 SPD RS09890 WP 074196785.1	MDISQVFRLRRKKLATAKQKNIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQMRV	GLSQGKSFSEMMESLGCSSA	79
NTDB id 196 SPR RS09425 WP 074196785.1	MDISQVFRLRRKKLATAKQKNIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQMRV	GLSQGKSFSEMMESLGCSSA	79
NTDB id 161 SP RS10375 WP 074196785.1	MDISQVFRLRRKKLATAKQKNIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQMRV	GLSQGKSFSEMMESLGCSSA	79
NTDB id 529 SMSK321 RS09235 WP 080550783.1	MDISQVFRLRRKKLATAKQKNIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQMRV	GLSQGKSFSEMMESLGFSSA	79
NTDB id 501 SM12261 RS00950 WP 004238896.1	MDISQVFRLRRKKLATAKQKNIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQMRV	GLSQGKSFSEMMESLGFSSA	79
consensus	*****	*****	79

[illegible]

NTDB id 587 KW2 RS10850 WP 042211510.1	GALIVLLTSLIIIFKHISALINFTFLVKVPLVSSFIRLYLTAYFSREWGNI	TAQGVELRQIIELMKKQKSRIFA	EVGIKLDLGL	251
NTDB id 395 FSA28 RS09400 WP 257638827.1	CCFFILLVGIFTFFSRKISRLHLIGFC	SHLPFVGKYLQLYLTAYYAREWGNI	IGQGVEMAQIVRLMQDQKSQLFREIGYDMEKAL	253
NTDB id 386 SMU RS09030 WP 255262721.1	CCFFILLVGIFTFFSRKISRLHLIGFC	SHLPFAGKYLQLYLTAYYAREWGNI	IGQGVEMAQIVRLMQDQKSQLFREIGYDMEKAL	253
NTDB id 430 SGO RS09415 WP 012130941.1	VLMLLGLTGAIYLVFKGQKRIPVYSFLARLPFVGSFVRI	YLTAYYAREWGNMIGQGLELSQIFQIMQEQRSVL	LFQEIGQDLGQAL	254
NTDB id 98930 R8570 RS09850 WP 077141332.1	VGLVSVLALLALTFYKRSSKMSVFSILARLPF	IGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMQEQQSQLFKEVGQDLAQT	TL	248
NTDB id 271 KZH43 RS09260 WP 074196785.1	VGLVSVLALLALTFYKRSSKMSVFSILARLPF	IGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMQEQQSQLFKEVGQDLAQT	TL	248
NTDB id 230 SPD RS09890 WP 074196785.1	VGLVSVLALLALTFYKRSSKMSVFSILARLPF	IGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMQEQQSQLFKEVGQDLAQT	TL	248
NTDB id 196 SPR RS09425 WP 074196785.1	VGLVSVLALLALTFYKRSSKMSVFSILARLPF	IGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMQEQQSQLFKEVGQDLAQT	TL	248
NTDB id 161 SP RS10375 WP 074196785.1	VGLVSVLALLALTFYKRSSKMSVFSILARLPF	IGIFVQTYLTAYYAREWGNMISQGMELTQIFQMMQEQQSQLFKEVGQDLAQT	TL	248
NTDB id 529 SMSK321 RS09235 WP 080550783.1	VGLVSLTLLALTFYKRSSKMRVFSILARIPFL	GIFVQTYLTAYYAREWGNMISQGMELTQIFQIMQEQQSQLFKEIGQDLAQSL		248
NTDB id 501 SM12261 RS00950 WP 004238896.1	VGFFVSVGVLLALTFYKRGSKMRVFSILARIPFL	GIFVQTYLTAYYAREWGNMISQGMELTQIFQIMQEQQSQLFKEIGQDMAHAL		248
consensus	*****	*****	*****	*****

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NTDB id 98930 R8570 RS09850 WP 077141332.1
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



logo

[illegible]

- X non conserved
- X similar
- X > 50% conserved