

[illegible]

NTDB id	47590 CVCAS RS10980 WP 014570824.1	SELLESLSFSKNVVTQLALVEVHGNLSTGTMQVLVELHLKKQLKVKNKLVLEVATYPIILLIFLVGIMWGLKNYLLPQLNSGSNFATL	170
NTDB id	587 KW2 RS10850 WP 042211510.1	SEILESLSFSKNVVTQLALVEVHGNLVGTMTQVLVELHLKKQLKVKKKLIEVATYPVVLLVFLLIGIMWGLKNYLLPQLDKGNNFATL	154
NTDB id	395 FSA28 RS09400 WP 257638827.1	AEIMSSSLGFSDDVVTQLALSEVHGNTQKSILLKIESYLNNLILVKKKLVEVITYPEIILLIFLVVIMLGKKNYLLPQLERKGNFATR	156
NTDB id	386 SMU RS09030 WP 255262721.1	AEIMSSSLGFSDDVVTQLALSEVHGNTQKSILLKIESYLNNLILVKKKLVEVITYPEIILLIFLVVIMLGKKNYLLPQLERGKFNATR	156
NTDB id	430 SGO RS09415 WP 012130941.1	SEIMASLGFSDAVVTQLSLAELHGNLSLALLKIEEYLDNLAKVKKKLIEVATYPMMLLGLFLVLMITGLRNYLLPQLSSQNFAATQ	157
NTDB id	271 KZH43 RS09260 WP 074196785.1	SEMMEESLGCSSAIVTQLSLAEVHGNLHLSLKGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLLIMLGLRNYLLPQLDSSNIATQ	151
NTDB id	230 SPD RS09890 WP 074196785.1	SEMMEESLGCSSAIVTQLSLAEVHGNLHLSLKGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLLIMLGLRNYLLPQLDSSNIATQ	151
NTDB id	196 SPR RS09425 WP 074196785.1	SEMMEESLGCSSAIVTQLSLAEVHGNLHLSLKGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLLIMLGLRNYLLPQLDSSNIATQ	151
NTDB id	161 SP RS10375 WP 074196785.1	SEMMEESLGCSSAIVTQLSLAEVHGNLHLSLKGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLLIMLGLRNYLLPQLDSSNIATQ	151
NTDB id	529 SMSK321 RS09235 WP 080550783.1	SEMMEESLGFSSAIVTQLSLAEVHGNLHLSLKGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLSIMLGLRNYLLPQLDSSNIATQ	151
NTDB id	501 SM12261 RS00950 WP 004238896.1	SEMMEESLGFSSAIVTQLSLAEVHGNLHLSLKGKIEEYLDNLAKVKKKLIEVATYPLILLGFLLLIMLGLRNYLLPQLDSSNIATQ	151
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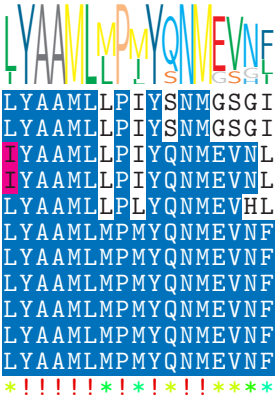
logo

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NTDB id 395 FSA28 RS09400 WP 257638827.1
NTDB id 386 SMU RS09030 WP 255262721.1
NTDB id 430 SGO RS09415 WP 012130941.1
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
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consensus



logo

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NTDB id 529 SMSK321 RS09235 WP 080550783.1
NTDB id 501 SM12261 RS00950 WP 004238896.1
consensus



☐ non conserved
☒ similar
☒ ≥ 50% conserved