

logo

NTDB id	587 KW2 RS10850 WP 042211510.1	 MQT	DILRLLRPKGKKLALIKQAKLIQLMGNNLNNGFNLGVEVINFLALSKLVEKEFVFKMHQGLSSGKSLSEILESLSFSKN	81
NTDB id	54279 SDSE RS11295 WP 015057201.1	.MTFLQR	DISVRSKQGLKKLSSEKEYKLIQLLANILLTGFNLAEIIAFILTSSHLLIPEPYVLRMEEALLKGHGLADMLAGITGFSDA	84
NTDB id	395 FSA28 RS09400 WP 257638827.1	.MTFLQQ	DISLKEKHKKQKKLKIKQKRKIQLFNNFLTSGFNLTETIIDFLERSHLLAEIYTKMKMRQGLLNGLHLAEIMSSLGFSDDD	84
NTDB id	386 SMU RS09030 WP 255262721.1	.MTFLQQ	DISLKEKHKKQKKLKIKQKRKIQLFNNFLTSGFNLTETIIDFLERSHLLAEIYTKMKMRQGLLNGLHLAEIMSSLGFSDDD	84
NTDB id	430 SGO RS09415 WP 012130941.1	MISFLQQ	DISILSKQKQKKLGTSTKQKVIELFNNLFSSGFHLAETVDFLGRSALLEKNYYVQQMRRQLANGQAFAFEIMASLGFSDA	85
NTDB id	271 KZH43 RS09260 WP 074196785.1		MDISQVFRLRKKKLATAKQNIIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQMRRVGLSQGGSFSEMMESELGCSSA	79
NTDB id	230 SPD RS09890 WP 074196785.1		MDISQVFRLRKKKLATAKQNIIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQMRRVGLSQGGSFSEMMESELGCSSA	79
NTDB id	196 SPR RS09425 WP 074196785.1		MDISQVFRLRKKKLATAKQNIIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQMRRVGLSQGGSFSEMMESELGCSSA	79
NTDB id	161 SP RS10375 WP 074196785.1		MDISQVFRLRKKKLATAKQNIIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQMRRVGLSQGGSFSEMMESELGCSSA	79
NTDB id	529 SMSK321 RS09235 WP 080550783.1		MDISQVFRLRKKKLATAKQNIIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQMRTDLSQGGSFSEMMESELGFSSA	79
NTDB id	501 SM12261 RS00950 WP 004238896.1		MDISQVFRLRKKKLATAKQNIIITLFNNLFSSGFHLVETISFLDRSALLDKQCVTQMRAGLSQGGSFSEMMESELGFSSA	79
consensus				* ! * ! * * * * * * * ! ! * * * * ! * * * ! * * * ! * * * ! * ! * * * * ! * ! * ! * * * * ! * * ! * * ! * * * * * * * * * ! * *

logo

[illegible]

logo

[illegible]

