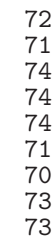


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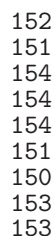
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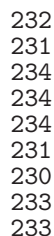
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NTDB id 1059 ABD1 RS01610 WP 000279215.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
consensus

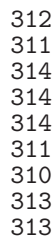
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logo

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Sequence 2	K	V	R	T	S	A	L	P	L	N	F	S	E	S	N	E	L	A	V	S	S	M	A	G	E	E	S	G	L	D	M	L	N	K	V	A	T	E	F	E	V	D	N	A	V	D	L	S	A	M	M	E	P	L	I	V	L	G	W	G	L																																							
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Sequence 9	K	I	R	E	D	V	A	T	G	Q	Q	L	Q	F	A	M	R	I	S	N	R	F	P	S	M	A	I	Q	M	V	A	G	E	E</																																																																		

V V A M Y L P I F N L M S V L G . 408
V T A M Y L P I F N L M S V L G . 407
L V A M Y L P I F N L G N V V A . 410
L V A M Y L P I F N L G N V V A . 410
L V A M Y L P I F N L G N V V A * 410
V V A M Y L P I F K L G S A V . . 406
I T A M Y L P I F Q M G S V V . . 405
V T A M Y L P I F Q M G S V V . . 408
I T A M Y L P I F Q M G S V V . . 408

* * * | ! | ! * | * ! * * * * * *

- ☐ non conserved
- ☒ similar
- ☒ $\geq 50\%$ conserved