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NTDB id 1169 A1552VC RS11075 WP 000648511.1
NTDB id 1402 DSB67 RS12670 WP 010643257.1
NTDB id 1112 NGFG RS09215 WP 003689811.1
NTDB id 1113 AAA85695.1 219..1451( )
NTDB id 1252 GCO85 RS07730 WP 011213805.1
NTDB id 1198 PSJM300 03950 AFN76868.1
NTDB id 158662 AOT82 RS08235 WP 071002607.1
NTDB id 1016 ACIAD RS01680 WP 004920476.1
NTDB id 1059 ABD1 RS01610 WP 000279215.1
consensus
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logo

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Sequence	Position 1	Position 2	Position 3	Position 4	Position 5	Position 6	Position 7	Position 8	Position 9	Position 10	Score
VAMYLPIFNLMSV	V	A	M	L	P	I	F	N	L	M	408
TAMYLPIFNLMSV	T	A	M	L	P	I	F	N	L	M	407
VAMYLPLFNLGNV	V	A	M	L	P	L	F	N	L	G	410
VAMYLPLFNLGNV*	V	A	M	L	P	L	F	N	L	G	410
VAMYLPIFKLGS	V	A	M	L	P	I	F	K	L	G	406
IAMYLPIFQMGSV	I	A	M	L	P	I	F	Q	M	G	405
IAMYLPIFQMGSV	I	A	M	L	P	I	F	Q	M	G	407
IAMYLPIFQMGSV	I	A	M	L	P	I	F	Q	M	G	408
IAMYLPIFQMGSV	I	A	M	L	P	I	F	Q	M	G	408

* ! ! ! ! * ! * * * * * *

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