

logo

NTDB id 416110 E3S98 RS00465 WP 000153535.1
NTDB id 4 SA RS09755 WP 000153535.1
NTDB id 83 BSU 35490 NP 391429.1
consensus

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Sequence	142	143	144	145	146	147	148	149	150	151	152	153	154	155	156	157	158	159	160																																																									
142	L	M	T	S	F	I	E	D	K	V	Y	R	A	D	A	G	V	D	S	Y	I	L	K	T	T	S	A	K	D	I	A	D	A	V	R	K	T	S	R	G	E	S	V	F	E	P	E	V	L	V	K	M	R	N	R	M	K	K	R	A																
142	L	M	T	S	F	I	E	D	K	V	Y	R	A	D	A	G	V	D	S	Y	I	L	K	T	T	S	A	K	D	I	A	D	A	V	R	K	T	S	R	G	E	S	V	F	E	P	E	V	L	V	K	M	R	N	R	M	K	K	R	A																
160	I	I	L	S	I	H	D	E	N	Y	T	H	A	L	K	T	G	A	R	G	I	L	L	K	E	M	D	A	D	T	L	I	E	A	V	K	V	V	A	E	G	S	Y	L	H	P	K	V	T	H	N	L	V	E	F	R	R	L	A	T	S	G	V	S	A	H	P	Q	H	E	V	Y	P	E	I	R

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[illegible]

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