

logo



T6335	WP	000170955.1	NC	010473:c1309809-1309702	MTLAQFAMI	F	W	H	D	L	A	A	P	I	L	A	G	I	I	T	A	A	I	V	S	W	W	R	N	R	K	35
T6329	YP	025297.1	NC	000913:c1269275-1269168	MTLAQFAMI	F	W	H	D	L	A	A	P	I	L	A	G	I	I	T	A	A	I	V	S	W	W	R	N	R	K	35
T6331	YP	025299.1	NC	000913:c1270345-1270238	MTLAQFAMI	F	W	H	D	L	A	A	P	I	L	A	G	I	I	T	A	A	I	V	S	W	W	R	N	R	K	35
T6341	WP	000170955.1	NC	007779:c1270852-1270745	MTLAQFAMI	F	W	H	D	L	A	A	P	I	L	A	G	I	I	T	A	A	I	V	S	W	W	R	N	R	K	35
T6343	WP	000170955.1	NC	007779:c1271922-1271815	MTLAQFAMI	F	W	H	D	L	A	A	P	I	L	A	G	I	I	T	A	A	I	V	S	W	W	R	N	R	K	35
T6333	WP	000170955.1	NC	010473:c1308739-1308632	MTLAQFAMI	F	W	H	D	L	A	A	P	I	L	A	G	I	I	T	A	A	I	V	S	W	W	R	N	R	K	35
T6327	CDJ71630.1	HG738867:c1157513-1157406			MTLAQFAMI	F	W	H	D	L	A	A	P	I	L	A	G	I	I	T	A	A	I	V	S	W	W	R	N	R	K	35
T6325	CDJ71628.1	HG738867:c1156443-1156336			MTLAQFAMI	F	W	H	D	L	A	A	P	I	L	A	G	I	I	T	A	A	I	V	S	W	W	R	N	R	K	35
T6347	AIZ51017.1	CP009789:c1252154-1252047			MTLAQFAMI	F	W	H	D	L	A	A	P	I	L	A	G	I	I	T	A	A	I	V	S	W	W	R	N	R	K	35
T6345	AIZ51015.1	CP009789:c1251084-1250977			MTLAQFAMI	F	W	H	D	L	A	A	P	I	L	A	G	I	I	T	A	A	I	V	S	W	W	R	N	R	K	35
T6339	WP	000170955.1	NC	020518:c1075773-1075666	MTLAQFAMI	F	W	H	D	L	A	A	P	I	L	A	G	I	I	T	A	A	I	V	S	W	W	R	N	R	K	35
T6337	WP	000170955.1	NC	020518:c1074703-1074596	MTLAQFAMI	F	W	H	D	L	A	A	P	I	L	A	G	I	I	T	A	A	I	V	S	W	W	R	N	R	K	35
T6330	YP	025298.1	NC	000913:c1269810-1269703	MTLAQFAMT	F	W	H	D	L	A	A	P	I	L	A	G	I	I	T	A	A	I	V	G	W	W	R	N	R	K	35
T6342	WP	000170963.1	NC	007779:c1271387-1271280	MTLAQFAMT	F	W	H	D	L	A	A	P	I	L	A	G	I	I	T	A	A	I	V	G	W	W	R	N	R	K	35
T6334	WP	000170963.1	NC	010473:c1309274-1309167	MTLAQFAMT	F	W	H	D	L	A	A	P	I	L	A	G	I	I	T	A	A	I	V	G	W	W	R	N	R	K	35
T6326	CDJ71629.1	HG738867:c1156978-1156871			MTLAQFAMT	F	W	H	D	L	A	A	P	I	L	A	G	I	I	T	A	A	I	V	G	W	W	R	N	R	K	35
T6346	AIZ51016.1	CP009789:c1251619-1251512			MTLAQFAMT	F	W	H	D	L	A	A	P	I	L	A	G	I	I	T	A	A	I	V	G	W	W	R	N	R	K	35
T6338	WP	000170963.1	NC	020518:c1075238-1075131	MTLAQFAMT	F	W	H	D	L	A	A	P	I	L	A	G	I	I	T	A	A	I	V	G	W	W	R	N	R	K	35
T6332	YP	026227.1	NC	000913:c3700087-3699980	MTFAELGMA	F	W	H	D	L	A	A	P	V	I	A	G	I	L	A	S	M	I	V	N	W	L	N	K	R	K	35
T6344	WP	000141634.1	NC	007779:3940328-3940435	MTFAELGMA	F	W	H	D	L	A	A	P	V	I	A	G	I	L	A	S	M	I	V	N	W	L	N	K	R	K	35
T6328	CDJ73338.1	HG738867:3081186-3081293			MTFAELGMA	F	W	H	D	L	A	A	P	V	I	A	G	I	L	A	S	M	I	V	N	W	L	N	K	R	K	35
T6348	AIZ53339.1	CP009789:c3670965-3670858			MTFAELGMA	F	W	H	D	L	A	A	P	V	I	A	G	I	L	A	S	M	I	V	N	W	L	N	K	R	K	35
T6340	WP	000141634.1	NC	020518:c3138994-3138887	MTFAELGMA	F	W	H	D	L	A	A	P	V	I	A	G	I	L	A	S	M	I	V	N	W	L	N	K	R	K	35
T6336	WP	000141634.1	NC	010473:c3795855-3795748	MTFAELGMA	F	W	H	D	L	A	A	P	V	I	A	G	I	L	A	S	M	I	V	N	W	L	N	K	R	K	35
consensus					!!*!***!	!!!!!!!**!!!!***!!	!***!!																									

- X non conserved
- X similar
- X ≥ 50% conserved