



# MuscleWS alignment of Uniref50 Sequences

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	80	90	111	121	131																																																					
<i>FER_CAPAA/1-97</i>	E	F	D	C	P	D	D	V	I	L	D	Q	A	E	E	A	G	H	D	L	P	Y	S	K	I	A	G	G	A	V	D	Q	T	D	G	N	F	L	D	D	D	Q	L	E	E	G	W	V	L	T	C	V	A	Y	P	Q	S	
<i>FER_CAPAN/1-144</i>	E	F	D	C	P	D	N	V	I	L	D	Q	A	E	E	A	G	H	D	L	P	Y	S	K	I	A	G	G	A	V	D	Q	T	D	G	N	F	L	D	D	D	Q	L	E	E	G	W	V	L	T	C	V	A	Y	P	Q	S	
<i>FER1_SOLLC/1-144</i>	E	F	E	C	P	D	D	V	I	L	D	Q	A	E	E	E	G	H	D	L	P	Y	S	K	V	T	A	G	S	V	D	Q	S	D	G	N	F	L	D	E	D	Q	E	A	A	G	F	V	L	T	C	V	A	Y	P	K	G	
<i>Q93XJ9_SOLTU/1-144</i>	E	F	E	C	P	D	D	V	I	L	D	Q	A	E	E	E	G	H	D	L	P	Y	S	K	V	T	A	G	T	V	D	Q	S	D	G	K	F	L	D	D	D	Q	E	A	A	G	F	V	L	T	C	V	A	Y	P	K	C	
<i>FER1_PEA/1-149</i>	E	F	E	C	P	S	D	V	I	L	D	H	A	E	E	V	G	I	D	L	P	Y	S	K	V	V	G	G	E	V	D	Q	S	D	G	S	F	L	D	D	E	Q	I	E	A	G	F	V	L	T	C	V	A	Y	P	T	S	
<i>Q7XA98_TRIPR/1-152</i>	E	F	D	C	P	D	D	V	I	L	D	H	A	E	E	V	G	I	E	L	P	Y	S	K	V	V	N	G	N	V	N	Q	E	D	G	S	F	L	D	D	E	Q	I	E	G	W	V	L	T	C	V	A	F	P	T	S		
<i>FER1_MESCR/1-148</i>	E	L	E	C	P	D	D	V	I	L	D	A	A	E	E	A	G	I	D	L	P	Y	S	K	V	T	S	G	S	V	N	Q	D	D	G	S	F	L	D	D	D	Q	I	K	E	G	W	V	L	T	C	V	A	Y	P	T	G	
<i>FER1_SPIOL/1-147</i>	E	F	Q	C	P	D	D	V	I	L	D	A	A	E	E	E	G	I	D	L	P	Y	S	K	L	K	T	G	S	L	N	Q	D	D	Q	S	F	L	D	D	D	Q	I	D	E	G	W	V	L	T	C	A	A	Y	P	V	S	
<i>FER3_RAPSA/1-96</i>	E	V	E	C	D	D	D	V	Y	V	L	D	A	A	E	E	A	G	I	D	L	P	Y	S	K	V	V	S	G	S	V	D	Q	S	D	Q	S	F	L	D	D	D	Q	I	A	E	G	F	V	L	T	C	A	A	Y	P	T	S
<i>FER2_ARATH/1-148</i>	E	V	E	C	D	D	D	V	Y	V	L	D	A	A	E	E	A	G	I	D	L	P	Y	S	K	V	V	S	G	S	V	D	Q	S	D	Q	S	F	L	D	D	E	Q	I	G	E	G	F	V	L	T	C	A	A	Y	P	T	S