



# MuscleWS alignment of Uniref50 Sequences

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80 90 100 110 120

<i>FER_CAPAA/1-97</i>	E	F	D	C	P	D	V	Y	I	L	D	Q	A	E	E	A	G	H	D	L	P	Y	S	C	R	A	G	S	C	S	S	C	A	G	K	I	A	G	G	A	V	D	Q	T	D	G	N	F	L	D	D	I	
<i>FER_CAPAN/1-144</i>	E	F	D	C	P	D	N	V	Y	I	L	D	Q	A	E	E	A	G	H	D	L	P	Y	S	C	R	A	G	S	C	S	S	C	A	G	K	I	A	G	G	A	V	D	Q	T	D	G	N	F	L	D	D	I
<i>FER1_SOLLC/1-144</i>	E	F	E	C	P	D	D	V	Y	I	L	D	Q	A	E	E	E	G	H	D	L	P	Y	S	C	R	A	G	S	C	S	S	C	A	G	K	V	T	A	G	S	V	D	Q	S	D	G	N	F	L	D	E	I
<i>Q93XJ9_SOLTU/1-144</i>	E	F	E	C	P	D	D	V	Y	I	L	D	Q	A	E	E	E	G	H	D	L	P	Y	S	C	R	A	G	S	C	S	S	C	A	G	K	V	T	A	G	T	V	D	Q	S	D	G	K	F	L	D	D	I
<i>FER1_PEA/1-149</i>	E	F	E	C	P	S	D	V	Y	I	L	D	H	A	E	E	V	G	I	D	L	P	Y	S	C	R	A	G	S	C	S	S	C	A	G	K	V	V	G	G	E	V	D	Q	S	D	G	S	F	L	D	D	I
<i>Q7XA98_TRIPR/1-152</i>	E	F	D	C	P	D	D	V	Y	I	L	D	H	A	E	E	V	G	I	E	L	P	Y	S	C	R	A	G	S	C	S	S	C	A	G	K	V	V	N	G	N	V	N	Q	E	D	G	S	F	L	D	D	I
<i>FER1_MESCR/1-148</i>	E	L	E	C	P	D	D	V	Y	I	L	D	A	A	E	E	A	G	I	D	L	P	Y	S	C	R	A	G	S	C	S	S	C	A	G	K	V	T	S	G	S	V	N	Q	D	D	G	S	F	L	D	D	I
<i>FER1_SPIOL/1-147</i>	E	F	Q	C	P	D	D	V	Y	I	L	D	A	A	E	E	E	G	I	D	L	P	Y	S	C	R	A	G	S	C	S	S	C	A	G	K	L	K	T	G	S	L	N	Q	D	D	Q	S	F	L	D	D	I
<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	C	R	A	G	S	C	S	S	C	A	G	K	V	V	S	G	S	V	D	Q	S	D	Q	S	F	L	D	D	I
<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	C	R	A	G	S	C	S	S	C	A	G	K	V	V	S	G	S	V	D	Q	S	D	Q	S	F	L	D	D	I
<i>FER_BRANA/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	C	R	A	G	S	C	S	S	C	A	G	K	V	V	S	G	F	V	D	Q	S	D	E	S	F	L	D	D	I
<i>FER1_ARATH/1-148</i>	E	V	E	C	E	E	D	V	Y	V	L	D	A	A	E	E	A	G	L	D	L	P	Y	S	C	R	A	G	S	C	S	S	C	A	G	K	V	V	S	G	S	I	D	Q	S	D	Q	S	F	L	D	D	I
<i>Q93Z60_ARATH/1-118</i>	E	V	E	C	E	E	D	V	Y	V	L	D	A	A	E	E	A	G	L	D	L	P	Y	S	C	R	A	G	S	C	S	S	C	A	G	K	V	V	S	G	S	I	D	Q	S	D	Q	S	F	L	D	D	I
<i>FER1_MAIZE/1-150</i>	E	L	Q	V	P	D	D	V	Y	I	L	D	Q	A	E	E	D	G	I	D	L	P	Y	S	C	R	A	G	S	C	S	S	C	A	G	K	V	V	S	G	S	V	D	Q	S	D	Q	S	Y	L	D	D	I
<i>O80429_MAIZE/1-140</i>	E	L	Q	V	P	D	D	V	Y	I	L	D	F	A	E	E	E	G	I	D	L	P	F	S	C	R	A	G	S	C	S	S	C	A	G	K	V	V	S	G	S	V	D	Q	S	D	Q	S	F	L	N	D	I