

## MuscleWS alignment of Uniref50 Sequences - FASTA

File Edit Select View Format Colour Calculate Web Service

	80	90	100	110	120	
<i>FER_CAPAA/1-97</i>	EFD	CPDDVYI	LDQAE EAGHD	LPYSCRAGSCSS	CAGKIAGGAVDQTDGNFL	DDDQLEEG
<i>FER_CAPAN/1-144</i>	EFD	CPDNVYI	LDQAE EAGHD	LPYSCRAGSCSS	CAGKIAGGAVDQTDGNFL	DDDQLEEG
<i>FER1_SOLLC/1-144</i>	EFE	CPDDVYI	LDQAE EAGHD	LPYSCRAGSCSS	CAGKVTAGSVDQSDGNFL	DEDQEAAG
<i>Q93XJ9_SOLTU/1-144</i>	EFE	CPDDVYI	LDQAE EAGHD	LPYSCRAGSCSS	CAGKVTAGTVDQSDGKFL	DDDQEAAG
<i>FER1_PEA/1-149</i>	EFE	CPSDVYI	LDHAE E VGID	LPYSCRAGSCSS	CAGKVVGGEV DQSDGS	FLDDEQIEAG
<i>Q7XA98_TRIPR/1-152</i>	EFD	CPDDVYI	LDHAE E VGIE	LPYSCRAGSCSS	CAGKVVNGNVNQEDGS	FLDDEQIEGG
<i>FER1_MESCR/1-148</i>	ELE	CPDDVYI	LDAAE EAGID	LPYSCRAGSCSS	CAGKVTSGSVNQDDGS	FLDDDQIKEG
<i>FER1_SPIOL/1-147</i>	EFQ	CPDDVYI	LDAAE E EGID	LPYSCRAGSCSS	CAGKLKTGSLNQDDQS	FLDDDQIDEG
<i>FER3_RAPSA/1-96</i>	EVE	CDDDVYV	LDAAE EAGID	LPYSCRAGSCSS	CAGKVVSGSVDQSDQS	FLDDDQIAEG
<i>FER1_ARATH/1-148</i>	EVE	CDDDVYV	LDAAE EAGID	LPYSCRAGSCSS	CAGKVVSGSVDQSDQS	FLDDEQIGEG
<i>FER_BRANA/1-96</i>	EVE	CDDDVYV	LDAAE EAGID	LPYSCRAGSCSS	CAGKVVSGFVDQSDES	FLDDDQIAEG
<i>FER2_ARATH/1-148</i>	EVE	CEEDVYV	LDAAE EAGLD	LPYSCRAGSCSS	CAGKVVSGSIDQSDQS	FLDDEQIMSEG
<i>Q93Z60_ARATH/1-118</i>	EVE	CEEDVYV	LDAAE EAGLD	LPYSCRAGSCSS	CAGKVVSGSIDQSDQS	FLDD-----
<i>FER1_MAIZE/1-150</i>	ELQ	VPDDVYI	LDQAE EDGID	LPYSCRAGSCSS	CAGKVVSGSVDQSDQSY	LDDGQIADG
<i>O80429_MAIZE/1-140</i>	ELQ	VPDDVYI	LDFAE E EGID	LPFSCRAGSCSS	CAGKVVSGSVDQSDQS	FLNDNQVADG