

MuscleWS alignment of Uniref50 Sequences - FASTA

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	80	90	100	110	120	
<i>FER_CAPAA/1-97</i>	EFD	CPDDVYI	LDQAEEAGHD	LPYSCRAGSCSS	CAGKIAGGAVD	QTDGNFLDDDDQLEEG
<i>FER_CAPAN/1-144</i>	EFD	CPDNVYI	LDQAEEAGHD	LPYSCRAGSCSS	CAGKIAGGAVD	QTDGNFLDDDDQLEEG
<i>FER1_SOLLC/1-144</i>	EFE	CPDDVYI	LDQAEEEGHD	LPYSCRAGSCSS	CAGKVTAGSVD	QSDGNFLDEDQEAAG
<i>Q93XJ9_SOLTU/1-144</i>	EFE	CPDDVYI	LDQAEEEGHD	LPYSCRAGSCSS	CAGKVTAGTVD	QSDGKFLDDDDQEAAG
<i>FER1_PEA/1-149</i>	EFE	CPSDVYI	LDHAEEVGI	LPYSCRAGSCSS	CAGKVVGGEV	QSDGSFLDDEQIEAG
<i>Q7XA98_TRIPR/1-152</i>	EFD	CPDDVYI	LDHAEEVGI	LPYSCRAGSCSS	CAGKVVNGNVN	QEDGSFLDDEQIEGG
<i>FER1_MESCR/1-148</i>	ELE	CPDDVYI	LDAAEEAGID	LPYSCRAGSCSS	CAGKVTSGSVN	QDDGSFLDDDDQIKEG
<i>FER1_SPIOL/1-147</i>	EFQ	CPDDVYI	LDAAEEEGID	LPYSCRAGSCSS	CAGKLKTGSLN	QDDQSFLDDDDQIDEG
<i>FER3_RAPSA/1-96</i>	EVE	CDDD	VYVLDAAEEAGID	LPYSCRAGSCSS	CAGKVVSGSVD	QSDQSFLDDDDQIAEG
<i>FER1_ARATH/1-148</i>	EVE	CDDD	VYVLDAAEEAGID	LPYSCRAGSCSS	CAGKVVSGSVD	QSDQSFLDDEQIGEG
<i>FER_BRANA/1-96</i>	EVE	CDDD	VYVLDAAEEAGID	LPYSCRAGSCSS	CAGKVVSGFVD	QSDQSFLDDDDQIAEG
<i>FER2_ARATH/1-148</i>	EVE	CEED	VYVLDAAEEAGLD	LPYSCRAGSCSS	CAGKVVSGSID	QSDQSFLDDEQIMSEG
<i>Q93Z60_ARATH/1-118</i>	EVE	CEED	VYVLDAAEEAGLD	LPYSCRAGSCSS	CAGKVVSGSID	QSDQSFLDD-----
<i>FER1_MAIZE/1-150</i>	ELQ	VPDDVYI	LDQAEEEDGID	LPYSCRAGSCSS	CAGKVVSGSVD	QSDQSYLDDGQIADG
<i>O80429_MAIZE/1-140</i>	ELQ	VPDDVYI	LDFAEEEGID	LPFSCRAGSCSS	CAGKVVSGSVD	QSDQSFLNDNQVADG