

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>	E FDCPDDVY I LDQAE E AGHD LPY SCRAG SCSSCAGKI AGGAVDQTDGN F LDDDQLEEG				
<i>FER_CAPAN/1-144</i>	E FDCPDNVY I LDQAE E AGHD LPY SCRAG SCSSCAGKI AGGAVDQTDGN F LDDDQLEEG				
<i>FER1_SOLLC/1-144</i>	E FEC PDDVY I LDQAE E EGHDL PY SCRAG SCSSCAGKVTAG SVDQSDGN F LDEDQEAAG				
<i>Q93XJ9_SOLTU/1-144</i>	E FEC PDDVY I LDQAE E EGHDL PY SCRAG SCSSCAGKVTAG TVDQSDGKF LDDDQEAAG				
<i>FER1_PEA/1-149</i>	E FEC P SDVY I LDHAE E VG I DLPY SCRAG SCSSCAGKVVGGEVDQSDGS F LDDEQIEAG				
<i>Q7XA98_TRIPR/1-152</i>	E FDCPDDVY I LDHAE E VG I E LPY SCRAG SCSSCAGKV VNGNVNQEDGS F LDDEQIEGG				
<i>FER1_MESCR/1-148</i>	E LEC PDDVY I LDAAE E AG I DLPY SCRAG SCSSCAGKVTSG SVNQDDGS F LDDDQIKEG				
<i>FER1_SPIOL/1-147</i>	E FQCPDDVY I LDAAE E EG I DLPY SCRAG SCSSCAGK LKTGS LNQDDQS F LDDDQIDEG				
<i>FER3_RAPSA/1-96</i>	E VEC DDDVY V LDAAE E AG I DLPY SCRAG SCSSCAGKVVSG SVDQSDQS F LDDDQIAEG				
<i>FER1_ARATH/1-148</i>	E VEC DDDVY V LDAAE E AG I DLPY SCRAG SCSSCAGKVVSG SVDQSDQS F LDDEQIGEG				
<i>FER_BRANA/1-96</i>	E VEC DDDVY V LDAAE E AG I DLPY SCRAG SCSSCAGKVVSG FVDQSDE SF LDDDQIAEG				
<i>FER2_ARATH/1-148</i>	E VEC EEDVY V LDAAE E AG LD LPY SCRAG SCSSCAGKVVSG S IDQSDQS F LDDEQMSEG				
<i>Q93Z60_ARATH/1-118</i>	E VEC EEDVY V LDAAE E AG LD LPY SCRAG SCSSCAGKVVSG S IDQSDQS F LDD-----				