



MuscleWS alignment of Uniref50 Sequences - FASTA

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	80	90	100	110
<i>FER_CAPAA/1-97</i>	E FDCPDDVY I LDQAE E AGHD LPY SCRAG SC	S SCAGK I AGGAVDQ		
<i>FER_CAPAN/1-144</i>	E FDCPDNVY I LDQAE E AGHD LPY SCRAG SC	S SCAGK I AGGAVDQ		
<i>FER1_SOLLC/1-144</i>	E F ECPDDVY I LDQAE E EGHDL PY SCRAG SC	S SCAGKVTAG SVDQ		
<i>Q93XJ9_SOLTU/1-144</i>	E F ECPDDVY I LDQAE E EGHDL PY SCRAG SC	S SCAGKVTAG TVDQ		
<i>FER1_PEA/1-149</i>	E F ECP SDVY I LDHAE E VG I DLPY SCRAG SC	S SCAGK VVGGE V DQ		
<i>Q7XA98_TRIPR/1-152</i>	E FDCPDDVY I LDHAE E VG I E LPY SCRAG SC	S SCAGK VVNGNVNQ		
<i>FER1_MESCR/1-148</i>	E L ECPDDVY I LDAAE E AG I DLPY SCRAG SC	S SCAGK V TSG S V NQ		
<i>FER1_SPIOL/1-147</i>	E FQCPDDVY I LDAAE E E G I DLPY SCRAG SC	S SCAGK LKTG S LNQ		
<i>FER3_RAPSA/1-96</i>	E V E CDDDVY V LDAAE E AG I DLPY SCRAG SC	S SCAGK V VSG S V DQ		
<i>FER1_ARATH/1-148</i>	E V E CDDDVY V LDAAE E AG I DLPY SCRAG SC	S SCAGK V VSG S V DQ		
<i>FER_BRANA/1-96</i>	E V E CDDDVY V LDAAE E AG I DLPY SCRAG SC	S SCAGK V VSG F V DQ		
<i>FER2_ARATH/1-148</i>	E V E C E D V Y V LDAAE E AG L DLPY SCRAG SC	S SCAGK V VSG S I DQ		
<i>Q93Z60_ARATH/1-118</i>	E V E C E D V Y V LDAAE E AG L DLPY SCRAG SC	S SCAGK V VSG S I DQ		