



MuscleWS alignment of Uniref50 Sequences - FASTA

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	80	90	100	110	120
<i>FER_CAPAA/1-97</i>	E FDCPDDVY I LDQAE E AGHD LPY SCRAG SCS SCAGK I AGGAVDQTDGN F LDDDDQLE EG				
<i>FER_CAPAN/1-144</i>	E FDCPDNVY I LDQAE E AGHD LPY SCRAG SCS SCAGK I AGGAVDQTDGN F LDDDDQLE EG				
<i>FER1_SOLLC/1-144</i>	E FEC PDDVY I LDQAE E EGHD LPY SCRAG SCS SCAGKVTAG SVDQSDGN F LDEDQE AAG				
<i>Q93XJ9_SOLTU/1-144</i>	E FEC PDDVY I LDQAE E EGHD LPY SCRAG SCS SCAGKVTAG TVDQSDGK F LDDDDQE AAG				
<i>FER1_PEA/1-149</i>	E FEC P SDVY I LDHAE E VG I DLPY SCRAG SCS SCAGKVVGG EVDQSDG S F LDDEQ I EAG				
<i>Q7XA98_TRIPR/1-152</i>	E FDCPDDVY I LDHAE E VG I E LPY SCRAG SCS SCAGKVVNGNVNQEDG S F LDDEQ I EGG				
<i>FER1_MESCR/1-148</i>	E L ECPDDVY I LDAAE E AG I DLPY SCRAG SCS SCAGKVT SG SVNQDDG S F LDDDDQ I KEG				
<i>FER1_SPIOL/1-147</i>	E FQCPDDVY I LDAAE E EG I DLPY SCRAG SCS SCAGK LKTG S LNQDDQ S F LDDDDQ I DEG				
<i>FER3_RAPSA/1-96</i>	E VE CDDDDVYV LDAAE E AG I DLPY SCRAG SCS SCAGKVV SG SVDQSDQ S F LDDDDQ I AEG				
<i>FER1_ARATH/1-148</i>	E VE CDDDDVYV LDAAE E AG I DLPY SCRAG SCS SCAGKVV SG SVDQSDQ S F LDDEQ I GEG				
<i>FER_BRANA/1-96</i>	E VE CDDDDVYV LDAAE E AG I DLPY SCRAG SCS SCAGKVV SG FVDQSD E S F LDDDDQ I AEG				
<i>FER2_ARATH/1-148</i>	E VE CE EDVYV LDAAE E AG LDLPY SCRAG SCS SCAGKVV SG S IDQSDQ S F LDDEQMSEG				
<i>Q93Z60_ARATH/1-118</i>	E VE CE EDVYV LDAAE E AG LDLPY SCRAG SCS SCAGKVV SG S IDQSDQ S F LDD - - - - -				