



MuscleWS alignment of Uniref50 Sequences - FASTA

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	80	90	100	110	120
<i>FER_CAPAA/1-97</i>	E FDCPDDVY I LDQ	AEE AGHD LPY	SCRAG SCS SCAGK I	AGG AVDQTDGN	N FLDDDQLE EG
<i>FER_CAPAN/1-144</i>	E FDCPDNVY I LDQ	AEE AGHD LPY	SCRAG SCS SCAGK I	AGG AVDQTDGN	N FLDDDQLE EG
<i>FER1_SOLLC/1-144</i>	E FEC PDDVY I LDQ	AEE EGHDL	LPY SCRAG SCS	SCAGKVTAG SVDQ	SDGN FLDEDQE AAG
<i>Q93XJ9_SOLTU/1-144</i>	E FEC PDDVY I LDQ	AEE EGHDL	LPY SCRAG SCS	SCAGKVTAG TVDQ	SDGK FLDDDQE AAG
<i>FER1_PEA/1-149</i>	E FEC P SDVY I LDH	AEE VGI DLPY	SCRAG SCS SCAGK	VVGG EVDQSDG	S FLDDDEQI EAG
<i>Q7XA98_TRIPR/1-152</i>	E FDCPDDVY I LDH	AEE VGI ELPY	SCRAG SCS SCAGK	VVNGNVNQ EDG	S FLDDDEQI EGG
<i>FER1_MESCR/1-148</i>	E L ECPDDVY I LDA	AEE AGIDLPY	SCRAG SCS SCAGK	VTSG SVNQDDG	S FLDDDQI KEG
<i>FER1_SPIOL/1-147</i>	E FQCPDDVY I LDA	AEE EGIDLPY	SCRAG SCS SCAGK	LKTGSLNQDDQ	S FLDDDQI DEG
<i>FER3_RAPSA/1-96</i>	E V E C D D D V Y V	L D A A E E A G I D	L P Y S C R A G S C S	S C A G K V V S G S	V D Q S D Q S F L D D D Q I A E G
<i>FER1_ARATH/1-148</i>	E V E C D D D V Y V	L D A A E E A G I D	L P Y S C R A G S C S	S C A G K V V S G S	V D Q S D Q S F L D D E Q I G E G
<i>FER_BRANA/1-96</i>	E V E C D D D V Y V	L D A A E E A G I D	L P Y S C R A G S C S	S C A G K V V S G F	V D Q S D E S F L D D D Q I A E G
<i>FER2_ARATH/1-148</i>	E V E C E E D V Y V	L D A A E E A G I D	L P Y S C R A G S C S	S C A G K V V S G S	I D Q S D Q S F L D D E Q M S E G
<i>Q93Z60_ARATH/1-118</i>	E V E C E E D V Y V	L D A A E E A G I D	L P Y S C R A G S C S	S C A G K V V S G S	I D Q S D Q S F L D D - - - - -
<i>FER1_MAIZE/1-150</i>	E L Q V P D D V Y I	L D Q A E E D G I D	L P Y S C R A G S C S	S C A G K V V S G S	V D Q S D Q S Y L D D G Q I A D G
<i>O80429_MAIZE/1-140</i>	E L Q V P D D V Y I	L D F A E E E G I D	L P F S C R A G S C S	S C A G K V V S G S	V D Q S D Q S F L N D N Q V A D G