



# MuscleWS alignment of Uniref50 Sequences - FASTA

File Edit Select View Annotations Format Colour Calculate Web Service

	80	90	100	110	120
<i>FER_CAPAA/1-97</i>	E FDCPDDVY I LDQAE E AGHDL P Y SC RAG SC S SCAGK I AGGAVDQ T DGN F LDDDQ L E EG				
<i>FER_CAPAN/1-144</i>	E FDCPDNVY I LDQAE E AGHDL P Y SC RAG SC S SCAGK I AGGAVDQ T DGN F LDDDQ L E EG				
<i>FER1_SOLLC/1-144</i>	E F E C P D D V Y I L D Q A E E E G H D L P Y S C R A G S C S S C A G K V T A G S V D Q S D G N F L D E D Q E A A G				
<i>Q93XJ9_SOLTU/1-144</i>	E F E C P D D V Y I L D Q A E E E G H D L P Y S C R A G S C S S C A G K V T A G T V D Q S D G K F L D D D Q E A A G				
<i>FER1_PEA/1-149</i>	E F E C P S D V Y I L D H A E E V G I D L P Y S C R A G S C S S C A G K V V G G E V D Q S D G S F L D D E Q I E A G				
<i>Q7XA98_TRIPR/1-152</i>	E F D C P D D V Y I L D H A E E V G I E L P Y S C R A G S C S S C A G K V V N G N V N Q E D G S F L D D E Q I E G G				
<i>FER1_MESCR/1-148</i>	E L E C P D D V Y I 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 T S G S V N Q D D G S F L D D D Q I K E G				
<i>FER1_SPIOL/1-147</i>	E F Q C P D D V Y I L D A A E E E G I D L P Y S C R A G S C S S C A G K L K T G S L N Q D D Q S F L D D D Q I D E G				
<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 L 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 L 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 - - - - -				



Sequence 3 ID: FER1\_SOLLC Residue: LYS (97)