

MuscleWS alignment of Uniref50 Sequences - FASTA

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	80	90	100	110	120									
<i>FER_CAPAA/1-97</i>	EFD	CPDDVY	ILDQAE	EAGHD	LPYSCR	AGSCSS	CAGKI	AGGAVD	QTDGN	FLDDD	QLEEG			
<i>FER_CAPAN/1-144</i>	EFD	CPDNVY	ILDQAE	EAGHD	LPYSCR	AGSCSS	CAGKI	AGGAVD	QTDGN	FLDDD	QLEEG			
<i>FER1_SOLLC/1-144</i>	EFE	CPDDVY	ILDQAE	EAGHD	LPYSCR	AGSCSS	CAGKV	TAGSVD	QSDGN	FLDED	QEAAG			
<i>Q93XJ9_SOLTU/1-144</i>	EFE	CPDDVY	ILDQAE	EAGHD	LPYSCR	AGSCSS	CAGKV	TAGTVD	QSDGK	FLDDD	QEAAG			
<i>FER1_PEA/1-149</i>	EFE	CPSDVY	ILDHAE	EVGID	LPYSCR	AGSCSS	CAGKV	VGGEVD	QSDGS	FLDDE	QIEAG			
<i>Q7XA98_TRIPR/1-152</i>	EFD	CPDDVY	ILDHAE	EVGIE	LPYSCR	AGSCSS	CAGKV	VNGNVN	QEDGS	FLDDE	QIEGG			
<i>FER1_MESCR/1-148</i>	ELE	CPDDVY	ILDAAE	EAGID	LPYSCR	AGSCSS	CAGKV	TSGSVN	QDDGS	FLDDD	QIKEG			
<i>FER1_SPIOL/1-147</i>	EFQ	CPDDVY	ILDAAE	EAGID	LPYSCR	AGSCSS	CAGKL	TGSLN	QDDQS	FLDDD	QIDEG			
<i>FER3_RAPSA/1-96</i>	EVE	CDDD	VYVLD	AAE	EAGID	LPYSCR	AGSCSS	CAGKV	VVSGS	VDQSD	QSF	FLDDD	QIAEG	
<i>FER1_ARATH/1-148</i>	EVE	CDDD	VYVLD	AAE	EAGID	LPYSCR	AGSCSS	CAGKV	VVSGS	VDQSD	QSF	FLDDE	QIEG	
<i>FER_BRANA/1-96</i>	EVE	CDDD	VYVLD	AAE	EAGID	LPYSCR	AGSCSS	CAGKV	VVSGF	VDQSD	ES	FLDDD	QIAEG	
<i>FER2_ARATH/1-148</i>	EVE	CEED	VYVLD	AAE	EAGLD	LPYSCR	AGSCSS	CAGKV	VVSGS	IDQSD	QSF	FLDDE	EQM	SEG
<i>Q93Z60_ARATH/1-118</i>	EVE	CEED	VYVLD	AAE	EAGLD	LPYSCR	AGSCSS	CAGKV	VVSGS	IDQSD	QSF	FLDD	-----	
<i>FER1_MAIZE/1-150</i>	ELQ	VPDDVY	ILDQAE	EDGID	LPYSCR	AGSCSS	CAGKV	VVSGS	VDQSD	QSYL	DDG	QI	ADG	
<i>O80429_MAIZE/1-140</i>	ELQ	VPDDVY	ILDFAE	EEGID	LPF	SCRAG	SCSS	CAGKV	VVSGS	VDQSD	QSYL	NDN	QV	ADG