

MuscleWS alignment of Uniref50 Sequences - FASTA

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	80	90	100	110	120				
<i>FER_CAPAA/1-97</i>	EFD	CPDDVY I LDQA	EEAGHDL PYS	CRAGS	SSCAGKI	AGGAVDQT	DGNFL	LDDDQLE	EEG
<i>FER_CAPAN/1-144</i>	EFD	CPDNVY I LDQA	EEAGHDL PYS	CRAGS	SSCAGKI	AGGAVDQT	DGNFL	LDDDQLE	EEG
<i>FER1_SOLLC/1-144</i>	EFE	CPDDVY I LDQA	EEEGHDL PYS	CRAGS	SSCAGKV	TAGSVDQSD	DGNFL	LDEDQE	AAG
<i>Q93XJ9_SOLTU/1-144</i>	EFE	CPDDVY I LDQA	EEEGHDL PYS	CRAGS	SSCAGKV	TAGTVDQSD	DGNFL	LDDDQE	AAG
<i>FER1_PEA/1-149</i>	EFE	CPSDVY I LDHA	EEVGI DLPYS	CRAGS	SSCAGKV	VGGVDQSD	DGNFL	LDDEQI	EAG
<i>Q7XA98_TRIPR/1-152</i>	EFD	CPDDVY I LDHA	EEVGI ELPYS	CRAGS	SSCAGKV	VNGNVNQED	DGNFL	LDDEQI	EAG
<i>FER1_MESCR/1-148</i>	ELE	CPDDVY I LDA	AAEEAGI DLPYS	CRAGS	SSCAGKV	TSGSVNQDD	DGNFL	LDDDQI	KEG
<i>FER1_SPIOL/1-147</i>	EFQ	CPDDVY I LDA	AAEEEGI DLPYS	CRAGS	SSCAGKL	KTGS LNQDD	DGNFL	LDDDQI	DEG
<i>FER3_RAPSA/1-96</i>	EVE	CDDDVYV LDA	AAEEAGI DLPYS	CRAGS	SSCAGKV	VSGSVDQSD	DGNFL	LDDDQI	AEG
<i>FER1_ARATH/1-148</i>	EVE	CDDDVYV LDA	AAEEAGI DLPYS	CRAGS	SSCAGKV	VSGSVDQSD	DGNFL	LDDEQI	GEG
<i>FER_BRANA/1-96</i>	EVE	CDDDVYV LDA	AAEEAGI DLPYS	CRAGS	SSCAGKV	VSGFVDQSD	DGNFL	LDDDQI	AEG
<i>FER2_ARATH/1-148</i>	EVE	CEEDVYV LDA	AAEEAGLDLPYS	CRAGS	SSCAGKV	VSGSIDQSD	DGNFL	LDDEQM	SEG
<i>Q93Z60_ARATH/1-118</i>	EVE	CEEDVYV LDA	AAEEAGLDLPYS	CRAGS	SSCAGKV	VSGSIDQSD	DGNFL	LDD	---
<i>FER1_MAIZE/1-150</i>	ELQVP	PDDVY I LDQA	EEEDGIDLPYS	CRAGS	SSCAGKV	VSGSVDQSD	DGNFL	LDD	GQIADG
<i>O80429_MAIZE/1-140</i>	ELQVP	PDDVY I LDFA	EEEGIDLPFS	CRAGS	SSCAGKV	VSGSVDQSD	DGNFL	LNDNQV	ADG