

MuscleWS alignment of Uniref50 Sequences - FASTA

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	80	90	100	110	120						
<i>FER_CAPAA/1-97</i>	EFD	CPDD	VYILD	QAEEAGHD	LPYSCRAG	SSCAGKI	AGGAVD	QTDGN	FLDDD	QLEEG	
<i>FER_CAPAN/1-144</i>	EFD	CPDN	VYILD	QAEEAGHD	LPYSCRAG	SSCAGKI	AGGAVD	QTDGN	FLDDD	QLEEG	
<i>FER1_SOLLC/1-144</i>	EFE	CPDD	VYILD	QAEEEGHD	LPYSCRAG	SSCAGK	VTAGS	VDQSDGN	FLDED	QEAAG	
<i>Q93XJ9_SOLTU/1-144</i>	EFE	CPDD	VYILD	QAEEEGHD	LPYSCRAG	SSCAGK	VTAGT	VDQSDGK	FLDDD	QEAAG	
<i>FER1_PEA/1-149</i>	EFE	CP	SDVYILD	HAEEVGID	LPYSCRAG	SSCAGK	VVGG	VDQSDGS	FLDDE	QIEAG	
<i>Q7XA98_TRIPR/1-152</i>	EFD	CPDD	VYILD	HAEEVGI	ELPYSCRAG	SSCAGK	VVNGN	VNQEDGS	FLDDE	QIEGG	
<i>FER1_MESCR/1-148</i>	ELE	CPDD	VYILD	AAEEAGID	LPYSCRAG	SSCAGK	VTSG	SVNQDDGS	FLDDD	QIKEG	
<i>FER1_SPIOL/1-147</i>	EFE	QCPDD	VYILD	AAEEEGID	LPYSCRAG	SSCAGK	LKTG	SLNQDDQS	FLDDD	QIDEG	
<i>FER3_RAPSA/1-96</i>	EVE	CDDD	VYVLD	AAEEAGID	LPYSCRAG	SSCAGK	VVSG	SVDQSDQS	FLDDD	QIAEG	
<i>FER1_ARATH/1-148</i>	EVE	CDDD	VYVLD	AAEEAGID	LPYSCRAG	SSCAGK	VVSG	SVDQSDQS	FLDDE	QIGEG	
<i>FER_BRANA/1-96</i>	EVE	CDDD	VYVLD	AAEEAGID	LPYSCRAG	SSCAGK	VVSG	FVDQSDQS	FLDDD	QIAEG	
<i>FER2_ARATH/1-148</i>	EVE	CEED	VYVLD	AAEEAGLD	LPYSCRAG	SSCAGK	VVSG	SIDQSDQS	FLDDE	QIMSEG	
<i>Q93Z60_ARATH/1-118</i>	EVE	CEED	VYVLD	AAEEAGLD	LPYSCRAG	SSCAGK	VVSG	SIDQSDQS	FLDD	-----	
<i>FER1_MAIZE/1-150</i>	ELQ	VPDD	VYILD	QAEE	DGIDLPY	SCRAG	SSCAGK	VVSG	SVDQSDQS	YLDDG	QIADG
<i>O80429_MAIZE/1-140</i>	ELQ	VPDD	VYILD	FAEE	EGIDLP	FSRAG	SSCAGK	VVSG	SVDQSDQS	FLNDN	NQVADG