



# MuscleWS alignment of Uniref50 Sequences

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	80	90	100	110	120													
<i>FER_CAPAA/1-97</i>	EFD	CPDD	VYIL	DQAE	EEAG	HDL	YPY	SCR	AGSC	SSC	AGK	IAG	GA	VD	QT	DGN	FL	DDD
<i>FER_CAPAN/1-144</i>	EFD	CPDN	VYIL	DQAE	EEAG	HDL	YPY	SCR	AGSC	SSC	AGK	IAG	GA	VD	QT	DGN	FL	DDD
<i>FER1_SOLLC/1-144</i>	EFE	CPDD	VYIL	DQAE	EEEG	HDL	YPY	SCR	AGSC	SSC	AGK	VTAG	SVD	QSD	DGN	FL	DED	
<i>Q93XJ9_SOLTU/1-144</i>	EFE	CPDD	VYIL	DQAE	EEEG	HDL	YPY	SCR	AGSC	SSC	AGK	VTAG	TVD	QSD	DGK	FL	DDD	
<i>FER1_PEA/1-149</i>	EFE	CPSD	VYIL	DHAE	EEVG	IDLP	YPY	SCR	AGSC	SSC	AGK	VVGG	EVD	QSD	DGS	FL	DDE	
<i>Q7XA98_TRIPR/1-152</i>	EFD	CPDD	VYIL	DHAE	EEVG	IEL	YPY	SCR	AGSC	SSC	AGK	VVNG	NVN	QED	DGS	FL	DDE	
<i>FER1_MESCR/1-148</i>	ELE	CPDD	VYIL	DAAE	EEAG	IDLP	YPY	SCR	AGSC	SSC	AGK	VTSG	SVN	QDD	DGS	FL	DDD	
<i>FER1_SPIOL/1-147</i>	EFE	QCPD	DDVY	ILDA	AAEE	EGID	LPY	SCR	AGSC	SSC	AGK	LKT	GS	LN	QDD	QSF	DDD	
<i>FER3_RAPSA/1-96</i>	EVE	CDDD	VYV	LDAA	EEAG	IDLP	YPY	SCR	AGSC	SSC	AGK	VVSG	SVD	QSD	QSF	FL	DDD	
<i>FER2_ARATH/1-148</i>	EVE	CDDD	VYV	LDAA	EEAG	IDLP	YPY	SCR	AGSC	SSC	AGK	VVSG	SVD	QSD	QSF	FL	DDE	
<i>FER_BRANA/1-96</i>	EVE	CDDD	VYV	LDAA	EEAG	IDLP	YPY	SCR	AGSC	SSC	AGK	VVSG	FVD	QSD	DES	FL	DDD	
<i>FER1_ARATH/1-148</i>	EVE	CEE	EDVY	VLDA	AAEE	AGLD	LPY	SCR	AGSC	SSC	AGK	VVSG	SID	QSD	QSF	FL	DDE	
<i>Q93Z60_ARATH/1-118</i>	EVE	CEE	EDVY	VLDA	AAEE	AGLD	LPY	SCR	AGSC	SSC	AGK	VVSG	SID	QSD	QSF	FL	DD-	
<i>FER1_MAIZE/1-150</i>	ELQ	VPD	DDVY	ILDQ	AEE	EDGI	DLPY	SCR	AGSC	SSC	AGK	VVSG	SVD	QSD	QSY	L	DDG	
<i>O80429_MAIZE/1-140</i>	ELQ	VPD	DDVY	ILDF	AEE	EGID	LPF	SCR	AGSC	SSC	AGK	VVSG	SVD	QSD	QSF	L	NDN	