

	10	20	30	40	50	60	70	80	90	100										
FER_CAPAA/1-97	-----ASYKVKLITPDGPIEFDCPDDVYILDQAE EAGHDL PYS CRA																			
FER_CAPAN/1-144	MA	SVSATMI	STSFMP	PKPAVT	SL-KPIP	NVGE--	ALFGL	KS-A--	NGGKV	TCMAS	YKVKL	ITPDG	PIEF	DCPD	NVYI	LDQA	EEAG	HDL	PYS	SCRA
FER1_SOLLC/1-144	MA	SVSATMI	STSFMP	PKPAVT	SL-KPIP	NVGE--	ALFGL	KS-G--	RNGRI	TCMAS	YKVKL	ITPEG	PIEF	ECPD	DDVY	ILDQA	EEEG	HDL	PYS	SCRA
Q93XJ9_SOLTU/1-144	MA	SVSATMI	STSFMP	PKPAVT	SL-KPIP	NVGE--	ALFGL	KS-G--	RNGRI	TCMAS	YKVKL	ITPDG	PIEF	ECPD	DDVY	ILDQA	EEEG	HDL	PYS	SCRA
FER1_PEA/1-149	MATT	--P	ALYG	TAT																
Q7XA98_TRIPR/1-152	MATT	--P	ALYG	TAT																
FER1_MESCR/1-148	MAAT	--T	AAL	SGAT																
FER1_SPIOL/1-147	MAAT	--T	TTMM	G--																
FER3_RAPSA/1-96	-----ATYKVKFITPEGEQEVECD										DDVY	VLDA	AAEE	EAGI	DL	PYS	SCRA			
FER2_ARATH/1-148	MAST	--A	LSSAI																	
FER_BRANA/1-96	-----																			
FER1_ARATH/1-148	MAST	--A	LSSAI	VST	SFL	RRQ	QTP	ISL	RS	LP	FA									
Q93Z60_ARATH/1-118	MAST	--A	LSSAI	VST	SFL	RRQ	QTP	ISL	RS	LP	FA									
FER1_MAIZE/1-150	MAT	VLG	SPRA	PAFF	FSS	SSL	RAAP	APT	AV--	AL	PA									
O80429_MAIZE/1-140	MAAT	--A	LSSAI																	

- Selection
- Sequence Details
- Show annotations
- Hide annotations
- Add reference annotations
- Edit
- Output to Textbox...
- Create Sequence Feature...
- Create Group
- Edit New Group
- Edit name and description of current group
- Group Colour
 - Text Colour...
 - None
 - Clustal
 - Blosum62
 - % Identity
 - Zappo
 - Taylor
 - Hydrophobic
 - Helix Propensity
 - Strand Propensity
 - Turn Propensity
 - Buried Index
 - Nucleotide
 - Purine/Pyrimidine
 - RNA Helices
 - T-Coffee Scores
- By Conservation
 - Modify Conservation Threshold...
 - Above Identity Threshold
 - Modify Identity Threshold...
- Boxes
- Text
- Colour Text
- Border colour
- Show nonconserved

Conservation
Quality
Consensus

