

File Edit Select View Annotations Format Colour Calculate Web Service

	10	20	30	40	50	60	70	80
FER_CAPAA/1-97	-----ASYKVKLITPDGPIEFDCPDDVYILDQ							
FER_CAPAN/1-144	MA-----SVSATMISTSFMPRKPAVTS L-KPIPNVGE--ALFGLKS-A--NGGKVT CMASYKVKLITPDGPIEFDCPDDVYILDQ							
FER1_SOLLC/1-144	MA-----SISGTMISTSFLPRKPAVTS L-KAISNVGE--ALFGLKS-G--RNGRITCMASYKVKLITPDGPIEFDCPDDVYILDQ							
Q93XJ9_SOLTU/1-144	MA-----SISGTMISTSFLPRKPVVTS L-KAISNVGE--ALFGLKS-G--RNGRITCMASYKVKLITPDGPIEFDCPDDVYILDQ							
FER1_PEA/1-149	MATT---PALYGTAVSTSLRRTQMPMSV-TTTKAFSN--GFLGLKT-SLKRGLAVAMASYKVKLVTPDGTQEFECPSDVYILDH							
Q7XA98_TRIPR/1-152	MATT---PALYGTAVSTSLRRTQMPMSV-ATTTTAKAFPSGFLGLKSVSTKRGLAVAMATYKVKLITPEGPQEFDCPDDVYILDH							
FER1_MESCR/1-148	MAAT--TAALS GATMSTAFAPK--TPPMTAAL							
FER1_SPIOL/1-147	MAAT--TTTMMG--MATT FVPKPQAPPMAAL							
FER3_RAPSA/1-96	-----							
FER2_ARATH/1-148	MAST----ALS SAIVGTS FIRRS PAPIS LRS LPSANTQ--SLFGL							
FER_BRANA/1-96	-----							
FER1_ARATH/1-148	MAST----ALS SAIVSTSLRRTQTPIS LRS LPSANTQ--SLFGL							
Q93Z60_ARATH/1-118	MAST----ALS SAIVSTSLRRTQTPIS LRS LPSANTQ--SLFGL							
FER1_MAIZE/1-150	MATV LGS PRAPAFFSSSLRAAPAPTAV--ALPAAKV--GIMGR							
O80429_MAIZE/1-140	MAAT-----ALSMSILR--APPPCFSSPLRLRV--AVAKF							

Selection ▶

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Create Sequence Feature...

Create Group

Edit New Group ▶

Conservation



Quality



Consensus



MA+T LGT PALSGT+VSTSLRRTQAPTS LRS LPS+VG+FP+L FGLKSVSTARGGRVTAMATYKVKLITPEGEQEFECPDDVYILDA