

MuscleWS alignment of Uniref50 Sequences

File	Edit	Select	View	Annotations	Format	Colour	Calculate	Web Service
				80	90	100	110	120
<i>FER_CAPAA/1-97</i>								
<i>FER_CAPAN/1-144</i>								
<i>FER1_SOLLC/1-144</i>								
<i>Q93XJ9_SOLTU/1-144</i>								
<i>FER1_PEA/1-149</i>								
<i>Q7XA98_TRIPR/1-152</i>								
<i>FER1_MESCR/1-148</i>								
<i>FER1_SPIOL/1-147</i>								
<i>FER3_RAPSA/1-96</i>								
<i>FER2_ARATH/1-148</i>								
<i>FER_BRANA/1-96</i>								
<i>FER1_ARATH/1-148</i>								
<i>Q93Z60_ARATH/1-118</i>								

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EFDCPDDVYILDQAE EAGHDLPYSCRAGSCSSCAGKIAGGAVDQTDGNFLDDDQLEEG
EFDCPDNVYILDQAE EAGHDLPYSCRAGSCSSCAGKIAGGAVDQTDGNFLDDDQLEEG
EF ECPDDVYILDQAE E EGHDLPYSCRAGSCSSCAGKVTAGSVDQSDGNFLDEDQEAAG
EF ECPDDVYILDQAE E EGHDLPYSCRAGSCSSCAGKVTAGTVDQSDGKFLDDDQEAAG
EF ECP SDVYILDHAE EVGIDLPLYSCRAGSCSSCAGKVVGGEVDQSDGSFLDDEQIEAG
EFDCPDDVYILDHAE EVGIELPLYSCRAGSCSSCAGKV VNGNVNQEDGSFLDDEQIEGG
EL ECPDDVYILDAAE EAGIDLPLYSCRAGSCSSCAGKVTSGSVNQDDGSFLDDDQIKEG
EFQCPDDVYILDAAE E E GIDLPLYSCRAGSCSSCAGK LKTGSLNQDDQSFLDDDQIDEG
EVE CDDD VYVLDAAE EAGIDLPLYSCRAGSCSSCAGKV VSGSVDQSDQSFLDDDQIAEG
EVE CDDD VYVLDAAE EAGIDLPLYSCRAGSCSSCAGKV VSGSVDQSDQSFLDDEQIGEG
EVE CDDD VYVLDAAE EAGIDLPLYSCRAGSCSSCAGKV VSGFVDQSDESFLDDDQIAEG
EVE CE EDVYVLDAAE EAGLDPLYSCRAGSCSSCAGKV VSGSIDQSDQSFLDDEQMSEG
EVE CE EDVYVLDAAE EAGLDPLYSCRAGSCSSCAGKV VSGSIDQSDQSFLDD - - - - -

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