Tree scale: 0.1
MpMAX2 Mapoly0113s0006
MAX2 Q9SIM9|AT2G42620
SpTIR Cluster 8427.1 TRINITY DN11823 c2 g4 i4|g.8509|m.8509
KfTIR Cluster 31363.1 TRINITY DN20215 c0 g2 i1|g.49437|m.49437
KfTIR Cluster 25758.4 TRINITY DN20971 c0 g1 i2|g.37033|m.37033
KfTIR Cluster 26784.0 TRINITY DN20971 c0 g2 i1|g.39180|m.39180
SpTIR Cluster 17496.0 TRINITY DN10300 c3 g1 i1|g.21336|m.21336
SpTIR Cluster 17496.0 TRINITY DN10300 c3 g1 i1|g.21335|m.21335
SpTIR1COI1 comp5783 c0 seq2
MpVFB Mapoly0026s0081
VFB2|A0A178VA62 ARATH|AT3G50080
VFB4|A0A178UK44 ARATH|AT5G67250
VFB1|A0A178W976 ARATH|AT1G47056
VFB3|A0A178U6I0 ARATH|AT4G07400
SpTIR Cluster 16895.0 TRINITY DN10360 c11 g4 i1|g.20501|m.20501
KfTIR Cluster 33039.0 TRINITY DN49742 c0 g1 i1|g.53075|m.53075
kfl00290 0040 v1.1
KfTIR Cluster 10648.0 TRINITY DN26013 c0 g1 i1|g.11331|m.11331
EBF2 Q708Y0|AT5G25350
EBF1 Q9SKK0|AT2G25490
MpEBF Mapoly0013s0206
SpTIR Cluster 20457.2 TRINITY DN9981 c8 g1 i3|g.26078|m.26078
SpTIR Cluster 20457.0 TRINITY DN9981 c8 g1 i5|g.26076|m.26076
SpTIR Cluster 16742.3 TRINITY DN13893 c4 g2 i1|g.20273|m.20273
kfl00036 0340 v1.1
KfTIR Cluster 3073.0 TRINITY DN15887 c0 g1 i1|g.2514|m.2514
KfTIR Cluster 33234.0 TRINITY DN22639 c4 g1 i3|g.53506|m.53506
SpTIR Cluster 12610.0 TRINITY DN9516 c0 g1 i5|g.13832|m.13832
kfl00028 0560 v1.1
KfTIR Cluster 15171.0 TRINITY DN1688 c0 g1 i1|g.18457|m.18457
KfTIR Cluster 14568.0 TRINITY DN11605 c0 g1 i1|g.17572|m.17572
KfTIR Cluster 14568.0 TRINITY DN11605 c0 g1 i1|g.17571|m.17571
KfTIR1COI1like kfl00434 0100 v1.1
GA XRTZ 2005787 Roya obtusa|m.1026
GA JOJQ 2003471 Cylindrocystis cushleckae|m.744
GA YOXI 2051391 Cylindrocystis brebissonii M2213|m.1054
GA VAZE 2053808 Cylindrocystis sp.|m.956
KfTIR1COI1 kfl00434 0030 v1.1
KfTIR Cluster 30851.0 TRINITY DN20170 c6 g2 i4|g.48275|m.48275
GA VQBJ 2001144 Coleochaete scutata|m.971
GA VQBJ 2011792 Coleochaete scutata|m.967
GA QPDY 2030019 Coleochaete irregularis|m.872
Pp3c18 16630V3.1 COI1
Pp3c22 6020V3.1 COI1
PpCOI1b Cluster 18008.0 TRINITY DN12432 c2 g8 i3|g.28231|m.28231
CrCOI1 Cluster 15372.0 TRINITY DN24371 c0 g1 i2|g.15278|m.15278
AmTr v1.0 scaffold00029.327 COI1
COI1 AT2G39940.1
Solyc05g052620.2 COI1
AaCOI1 Cluster 14646.0 TRINITY DN5778 c0 g5 i11|g.39358|m.39358
MpCOI1 Mapoly0025s0025.1
Pp3c23 18960V3.1 COI1
PpCOI1d Cluster 25569.0 TRINITY DN12124 c1 g1 i4|g.46718|m.46718
Pp3c3 8580V3.1 COI1
AaCOI1 Cluster 14646.0 TRINITY DN5778 c0 g5 i11|g.39360|m.39360
Pp3c9 10120V3.1 COI1
Pp3c17 7680V3.1 COI1
PpCOI1c Cluster 2314.0 TRINITY DN11836 c1 g3 i1|g.2086|m.2086
PpCOI1a Cluster 10449.0 TRINITY DN11718 c5 g3 i10|g.13339|m.13339
Pp3c15 1200V3.1 COI1
CrTIR1e Cluster 43370.0 TRINITY DN21417 c0 g1 i3|g.51190|m.51190
Solyc02g079190.2
AmTr v1.0 scaffold00003.426 TIR1AFB
AaTIR1 Cluster 8963.4916 TRINITY DN4627 c4 g1 i2|g.20270|m.20270
CrTIR1e Cluster 43370.0 TRINITY DN21417 c0 g1 i3|g.51189|m.51189
MpTIR1 Mapoly0035s0062.1
Pp3c23 5870V3.1 TIR1AFB
PpTIR1b Cluster 22752.1 TRINITY DN12554 c3 g2 i1|g.39552|m.39552
Pp3c20 17150V3.1 TIR1AFB
PpTIR1a Cluster 22752.0 TRINITY DN12554 c3 g3 i2|g.39549|m.39549
Pp3c23 11300V3.1 TIR1AFB
PpTIR1c Cluster 26757.1 TRINITY DN12554 c3 g5 i3|g.50004|m.50004
PpTIR1d Cluster 26757.2 TRINITY DN12604 c0 g3 i2|g.50008|m.50008
PpTIR1d Cluster 26757.2 TRINITY DN12604 c0 g3 i2|g.50007|m.50007
Pp3c24 9800V3.1 TIR1AFB
Pp3c24 9814V3.1 TIR1AFB
CrTIR1b Cluster 28618.0 TRINITY DN18266 c0 g1 i1|g.30469|m.30469
CrTIR1d Cluster 3237.365 TRINITY DN27305 c0 g1 i3|g.1853|m.1853
CrTIR1c Cluster 31007.0 TRINITY DN28153 c7 g3 i1|g.33551|m.33551
CrTIR1f Cluster 6319.0 TRINITY DN28153 c7 g1 i1|g.6735|m.6735
AaTIR1 Cluster 8963.4916 TRINITY DN4627 c4 g1 i2|g.20269|m.20269
Sm170974 TIR1 like
Sm168175 TIR1 like
AmTr v1.0 scaffold00135.41 TIR1AFB
Solyc04g074980.2
AFB5 AT5G49980.1
AFB4 AT4G24390.1
LOC Os03g08850.1
GRMZM2G095786 T01
GRMZM2G398848 T01
LOC Os02g52230.1
GRMZM2G427529 T01
GRMZM2G024180 T01
CrTIR1a Cluster 24229.0 TRINITY DN3861 c0 g1 i1|g.25020|m.25020
AFB2 AT3G26810.1
AFB3 AT1G12820.1
GRMZM2G155849 T01
LOC Os11g31620.1
LOC Os04g32460.1
GRMZM2G137451 T02
GRMZM5G848945 T01
AmTr v1.0 scaffold00016.85 TIR1AFB
GRMZM2G135978 T01
LOC Os05g05800.1
AFB1 AT4G03190.1
TIR1 AT3G62980.1
Solyc09g074520.2
Solyc06g008460.1
Solyc06g008780.1
Solyc06g008810.2