

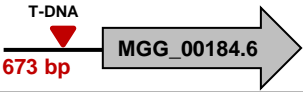

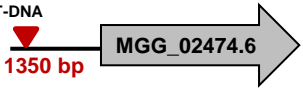

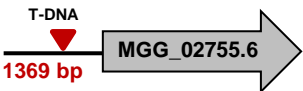
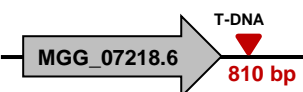
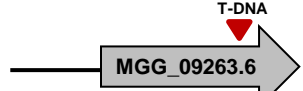
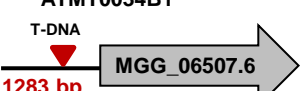
T-DNA insertion mutants	No. TFs	Conidiation	Conidial morphology	Conidial germination	Appressorium Formation	Pathogenicity
ATMT0651A4 	HOX2	No conidia	-	-	-	-
ATMT0094A6 	TF220	Reduced	Normal	<50%	<50%	No disease
ATMT0104A6 	TF116	Reduced	Normal	>50%	No App.	Reduced
ATMT0068B3 	TF041	Reduced	Normal	>50%	No App.	-
ATMT0349D2 	TF031	Reduced	no record	>50%	>50%	Normal
ATMT0052B2 	TF233	Normal	Abnormal	<50%	<50%	No disease
ATMT0591D1 	TF260	Normal	Abnormal	>50%	<50%	Reduced
ATMT0034B1 	TF150	Normal	Abnormal	>50%	>50%	No disease

Figure S5. T-DNA insertion sites in eight mutants defective in conidiation-specific TF genes and resulting phenotypes. Phenotypes of these mutants are derived from *Magnaporthe oryzae* T-DNA insertion mutant library (<http://atmt.snu.ac.kr>).