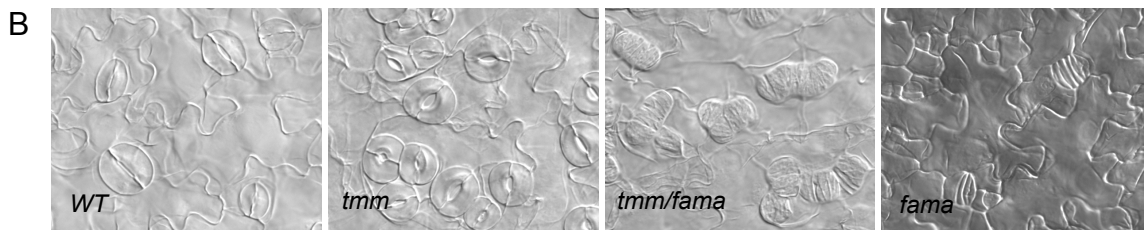


A

parent		progeny				results
		WT	mutant1	fama (-/-)	muant1 fama	
fama/+	tmm/tmm	-	109	-	17	additive
fama/+	sdd1/+	41	21	8	3	additive
fama/+	er/erl1/erl2	-	11*	-	1	additive
fama/+	flp/flp	-	153	-	29	additive
fama/+	yoda	46	10	8	2	additive



**Supplemental Figure 3: Genetic interactions between *fama* and mutations in other stomatal regulators.** (A) Table of crosses tested, all phenotypes were additive with respect to number of cells in the tumors, but the *fama-1* differentiation phenotype was always epistatic. (B) Example of *tmm; fama-1* double mutant phenotypes.