Table S1. Quantification of the average number of ear primary branches in *rel2;ra2* double mutants

	n	Number of primary branches
rel2; ra2	11	23.1 (5.4)*
rel2;+/ra2	18	0
+/rel2;ra2	13	0.5 (0.4)*
+/rel2;+/ra2	15	0

Data are mean (±s.e.m.).
*t-test P=0.0002.