

Plant line	% of roots with tumors	% of roots showing GUS activity 6 days after infection
Wild-type (Ws)	82.7± 3.0	48.3± 3.9
RNAi BTI1-22	20.3± 3.2	12.2± 1.7
RNAi BTI1-23	17.7± 4.4	11.3± 1.5
RNAi BTI1-24	12.3± 3.2	20.5± 2.5
RNAi BTI1-27	26.3± 2.7	17.3± 1.8
RNAi BTI2-42	19.1± 2.3	25.5± 6.7
RNAi BTI2-45	10.3± 0.8	10.9± 1.1
RNAi BTI2-47	15.3± 2.0	20.6± 2.7
RNAi BTI3-7	14.3± 2.5	13.8± 2.5
RNAi BTI3-16	8.3± 1.5	17.0± 2.3
RNAi BTI3-18	10.6± 1.9	17.8± 2.4
RNAi BTI3-83	17.2± 4.2	38.4± 5.8
RNAi AtRAB8-9	10.2± 2.7	14.3± 2.5
RNAi AtRAB8-11	12.1± 2.0	20.1± 1.0
RNAi AtRAB8-22	12.4± 3.3	11.3± 2.4

Supplemental Table 2. Stable and transient transformation assays of T2 generation transgenic *BTI1*, *BTI2*, *BTI3*, and *AtRAB8* RNA interference (RNAi) plants. Data are indicated as average values of results from 20 plants ± standard error. At least 80 root segments were examined for each plant.