

Additional file 4. Regression analysis of GFP expression and transgene copy number scored on the Southern blots. GFP expression for each hairy root shown on Figure 7 was quantified and grayscale values correlated with the respective transgene copy number. The regression analysis showed P-Values of 0.005 and 0.039 for GmERF6- and GmERF10-containing hairy roots, respectively.

