Supplemental data

Shoot regeneration is not a single cell event

Patharajan Subban, Yaarit Kutsher, Dalia Evenor, Eduard Belausov, Hanita Zemach, Adi Faigenboim, Samuel Bocobza, Mike Timko, Moshe Reuveni.



Figure S1: Number of genes that were up or down-regulated during the induction period on Reg medium.



Fig. S2: Transcript changes during shoot induction period on Reg medium show five distinct sub cluster. **A**: sub cluster of genes that upregulate mostly on day 4; **B**: sub cluster of genes that upregulate moderately on day 1; **C**: sub cluster of genes that upregulate sharply on day 1; **D**: sub cluster of genes that downregulate on day 1; **E**: sub cluster of genes that downregulate on day 4.



Figure S3. transcript analyses of biological processes that upregulate at day 1 compared to day 0 of the induction of shoot regeneration.



Figure S4. transcript analyses of biological processes that upregulate at day 4 compared to day 0 of the induction of shoot regeneration.



Figure S5. Transcript analyses of biological processes that upregulate at day 7 compared to day 0 of the induction of shoot regeneration.



Figure S6. Transcript analyses of MADS box gene family during the 7 c days ompared to day 0 of the induction of shoot regeneration.



Figure S7. Transcript analyses of NAC gene family during the 7 c days ompared to day 0 of the induction of shoot regeneration.



Figure S8. Transcript analyses of histone related genes during the 7 c days ompared to day 0 of the induction of shoot regeneration.



Figure S9. Transcript analyses of phytohormones signal transduction phatways at the 1 day ompared to day 0 of the induction of shoot regeneration. Genes that are boxed in red show change.