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Correction to: Carbon nanotube biocompatibility in plants is determined by their surface chemistry

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Following publication of the original article [1], the authors identified an error in Fig. 2. The correct Fig. 2 is given in this erratum.

The original article can be found online at https://doi.org/10.1186/s12951-021-01178-8.

Full list of author information is available at the end of the article



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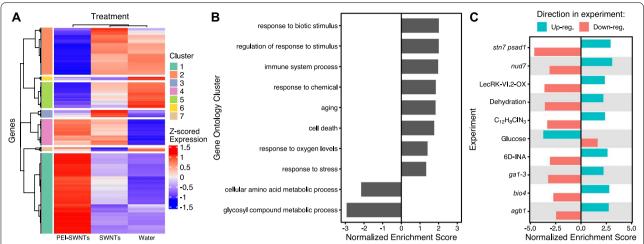


Fig. 2 PEI-SWNT responding genes are involved in stress responses, immune system, and programmed cell death. **A** Gene expression heatmap of genes with statistically significant twofold expression change in at least one of the three treatments, compared to non-infiltrated samples. Clusters 1 and 2 show PEI-SWNT specific up- and down-regulated genes. **B**, **C** Gene Set Enrichment Analysis of Cluster 1 and 2 genes using Biological Process Gene Ontology categories (**B**), and Arapath and PlantGSEA databases (**C**). Up- or down-regulation genes in the original experiment were used as independent gene sets to calculate their normalized enriched score. Details corresponding to each experiment can be found in Additional file 4: Table S4C

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