

CORRECTION

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Wang, Y.-H., Garvin, D.F., and Kochian, L.V. Nitrate-Induced Genes in Tomato Roots. Array Analysis Reveals Novel Genes That May Play a Role in Nitrogen Nutrition.

On pages 357 (“Materials and Methods”) and 358 (“Acknowledgments”), the source for the tomato (*Lycopersicon esculentum*) expressed sequence tags (ESTs) used for the research in this manuscript was not adequately acknowledged. Ordered root cDNA libraries were generated as a collaboration between our lab and the laboratory of S. Tanksley at Cornell University (Ithaca, NY; supported by National Science Foundation [NSF] grant no. DBI-9872617). Resulting clones were transferred to C. Ronning at the Institute for Genomic Research (TIGR, Rockville, MD) for sequencing (supported by NSF grant no. DBI-9813392). Approximately 6,000 of the sequenced clones were included in our initial array of approximately 20,000 clones used for the primary screening that resulted in the 1,280 clones used in our final mineral nutrition array. Additional information regarding the Tomato Gene Index of tomato ESTs developed by TIGR and the tomato genome project in general can be found at the following web sites, respectively: <http://www.tigr.org/tdb/lgi/> and <http://soltdb.cit.cornell.edu/>.