biology

Supplementary Materials:

Advances in Omics Approaches for Abiotic Stress Tolerance in Tomato

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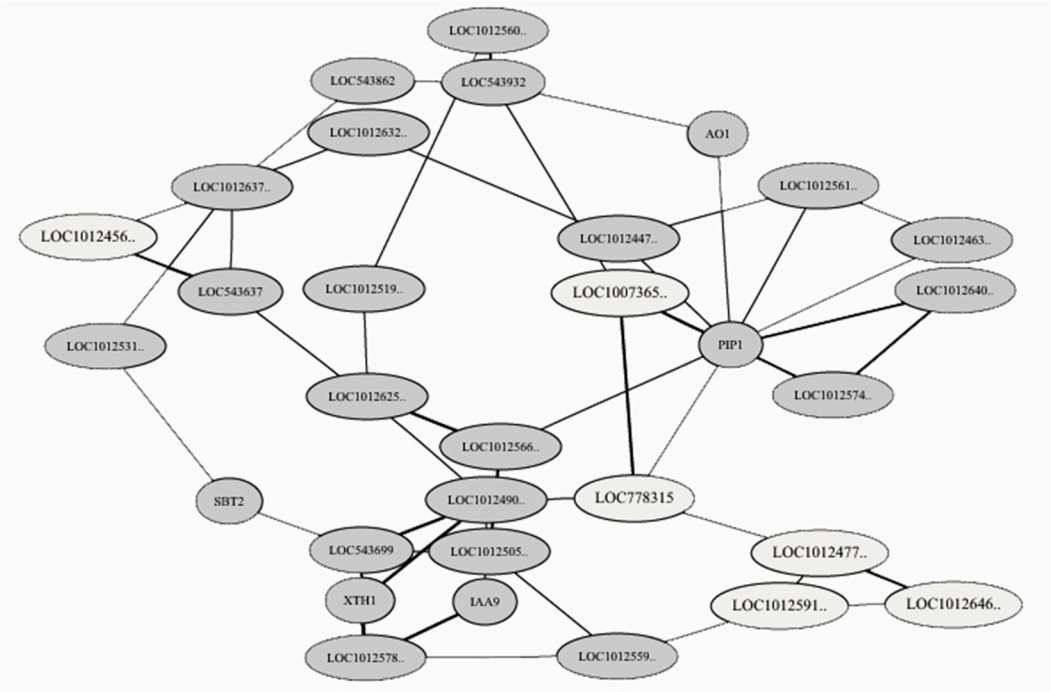

Supplementary Figure S1. Co-expression correlation network of tomato genes developed using an online tool ATTED-II (http://atted.jp/). The tool involves expression data of 401 microarray-based sample and 282 RNA-seq based samples. The figure depicts the correlation of Plasma Membrane Intrinsic Protein with other genes governing basic physiological function in tomato.