



Figure S7. MAPMAN-based functional classification of differentially expressed transcripts in *PXVE:NF-YA* and *PXVE:miR169nm* lines.

MAPMAN classification of the transcripts according to MAPMAN categories using the web-based tool Classification Supersviewer (<http://bar.utoronto.ca>) with normalized class score option. One, two and three asterisks indicate a p-value < 0.05, 0.001 and 0.0001, respectively.