Flowchart for multispecies analysis. An overview of the pipeline and methods used for exploring the conservation of homologous cell types and tissues among tomato, Arabidopsis, and rice. Ath=Arabidopsis thaliana; Osa=Oryza sativa; Sly=Solanum lycopersicum; IQR= interquartile range; TPM=transcript per million; GO=gene ontology; CTEG=cell type/tissue enriched genes.