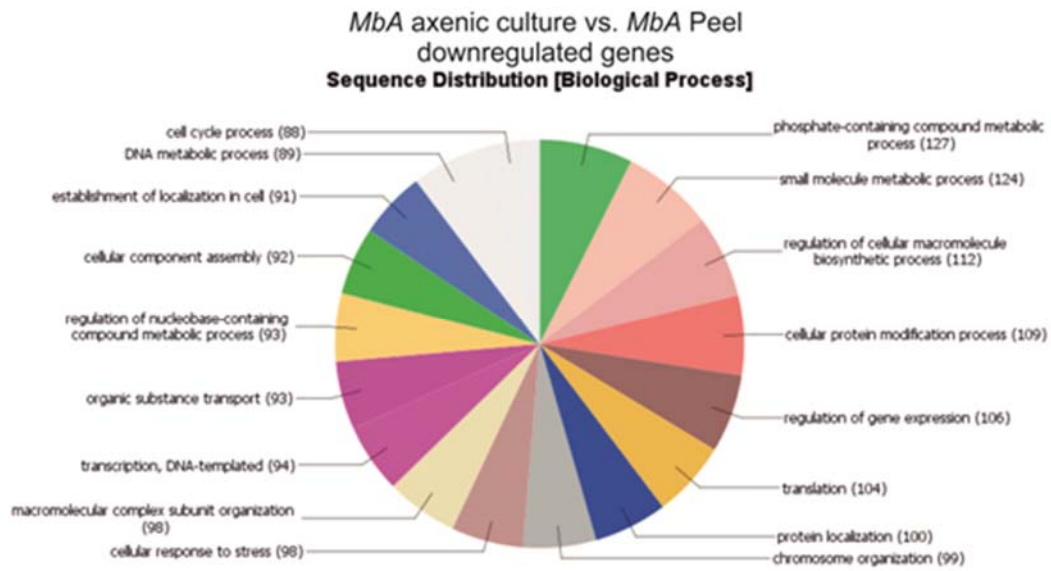


A



B

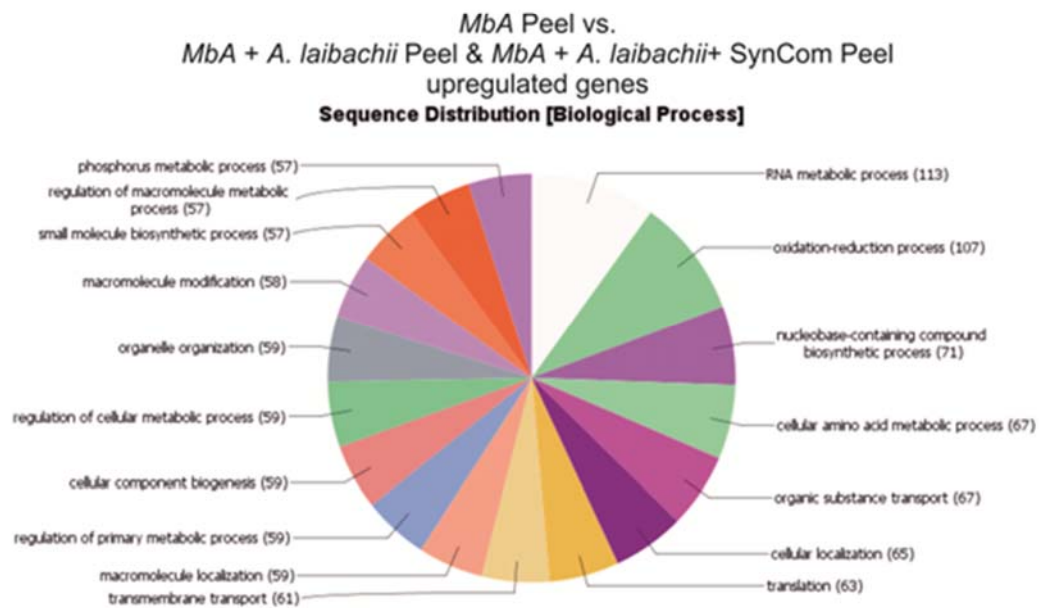


Figure 6 – figure supplement 2: (A) Sequence distribution of Gene Ontology terms. 60% of all genes, downregulated in *MbA* on plant compared to axenic culture growth, can be assigned to GO-terms related to metabolism and cell cycle. (B) In contrast, presence of *A. laibachii* leads to transcriptional activation of metabolic processes. 52% of all GO-terms associated with genes upregulated in presence of *A. laibachii* are related to metabolic processes.