

Transcriptome and genome sequencing elucidates the molecular basis for the high yield and good quality of the hybrid rice variety Chuanyou6203

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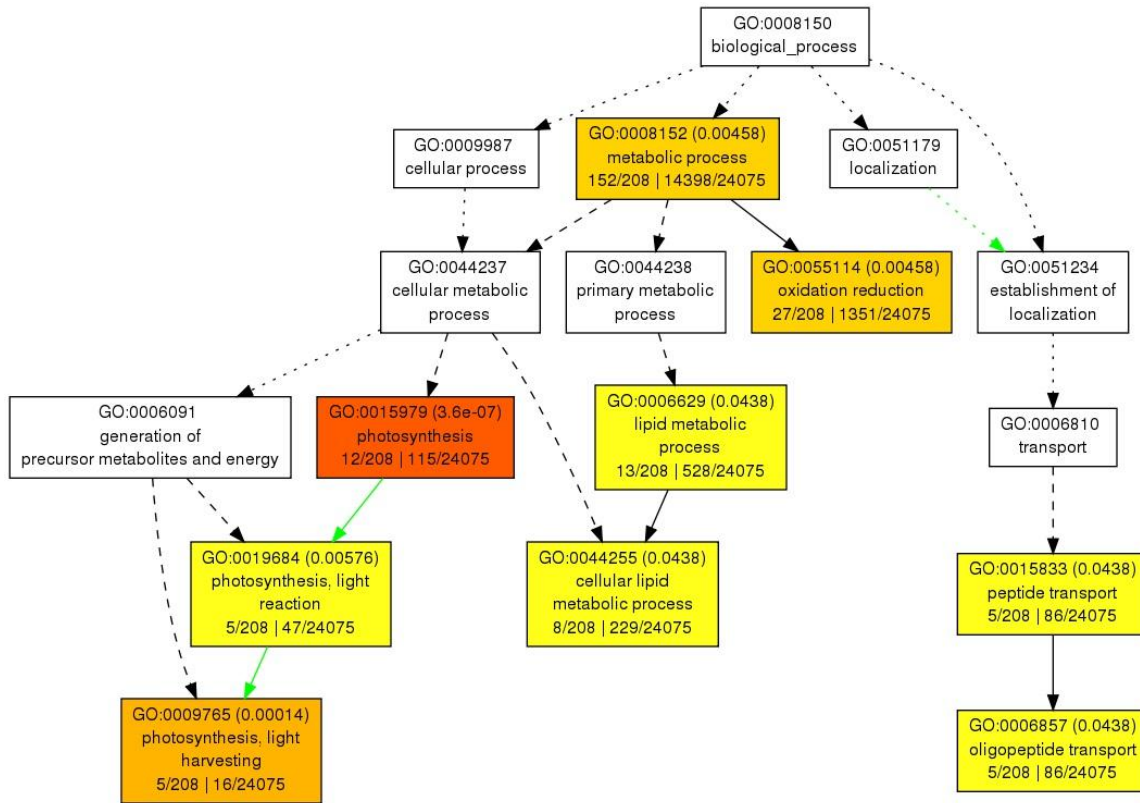


Figure S5. Gene ontology analysis of DEGs belonging to green-yellow module by agriGO v2 (<http://systemsbiology.cau.edu.cn/agriGOv2/>). The green arrow represents negative regulation.