Top Positive and Negative Enriched GO Pathways (log2FC) GO:0019684 GO_BP:photosynthesis,_light_reaction GO_MF:phosphatidylinositol_phosphate_phosphatase_activity GO:0052866 GO:0006367 GO_BP:transcription_initiation_at_RNA_polymerase_II_promoter GO_BP:negative_regulation_of_endopeptidase_activity GO:0010951 GO_BP:negative_regulation_of_proteolysis GO:0045861 GO:0005795 GO_CC:Golgi_stack GO_MF:transition_metal_ion_transmembrane_transporter_activity GO:0046915 GO:0071489 GO_BP:cellular_response_to_red_or_far_red_light GO:0010017 GO_BP:red_or_far-red_light_signaling_pathway direction ☐ GO:0016102 O GO:0007017 GO_BP:diterpenoid_biosynthetic_process Downregulated GO:0007017 Upregulated GO_BP:microtubule-based_process GO:0015630 GO_CC:microtubule_cytoskeleton GO_CC:spindle GO:0005819 GO:0033037 GO_BP:polysaccharide_localization GO:0032182 GO_MF:ubiquitin-like_protein_binding GO:0003774 GO_MF:cytoskeletal_motor_activity GO:0007018 GO_BP:microtubule-based_movement GO:0005871 GO_CC:kinesin_complex GO:0003777 GO_MF:microtubule_motor_activity GO:0005875 GO_CC:microtubule_associated_complex

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