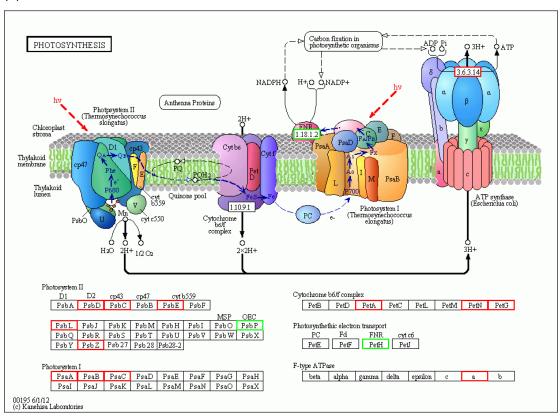
(A) Significantly enriched pathways of differentially expressed genes by 1-h cold stress in Anthurium

Pathway	DEGs with pathway annotation (721)	DEGs with pathway annotation (721)	Pvalue	Qvalue	Pathway ID
Photosynthesis	15 (2.08%)	65 (0.45%)	5.05E-07	5.65E-05	ko00195
Oxidative	22 (3.05%)	214 (1.48%)	0.001032	5.78E-02	ko00190
phosphorylation					

(B)



Detailed information of photosynthesis pathway in KEGG database. In the figure, up-regulated genes are marked with red borders while down-regulated genes are marked with green borders. Non-change genes are marked with black borders.