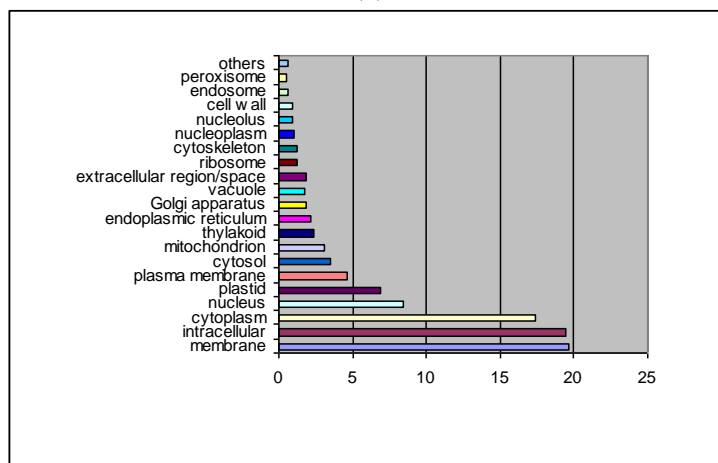


Supplementary data

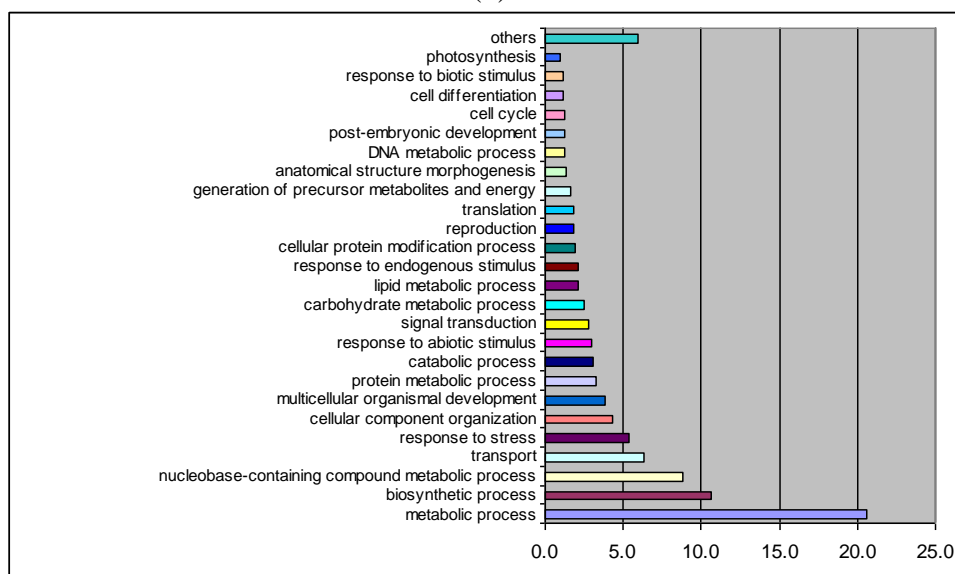
Analysis of expressed sequence tags and alternative splicing genes in sacred lotus (*Nelumbo nucifera* Gaertn.)

Robert VanBuren, Braden Walters, Ray Ming, Xiang Jia Min

(a)



(b)



Supplementary Fig 1. Distribution of predicted proteins in ESTs of sacred lotus based on Gene Ontology classifications. (a) Cellular component; and (b) Biological process. The x-axis indicates the percentage of each category.