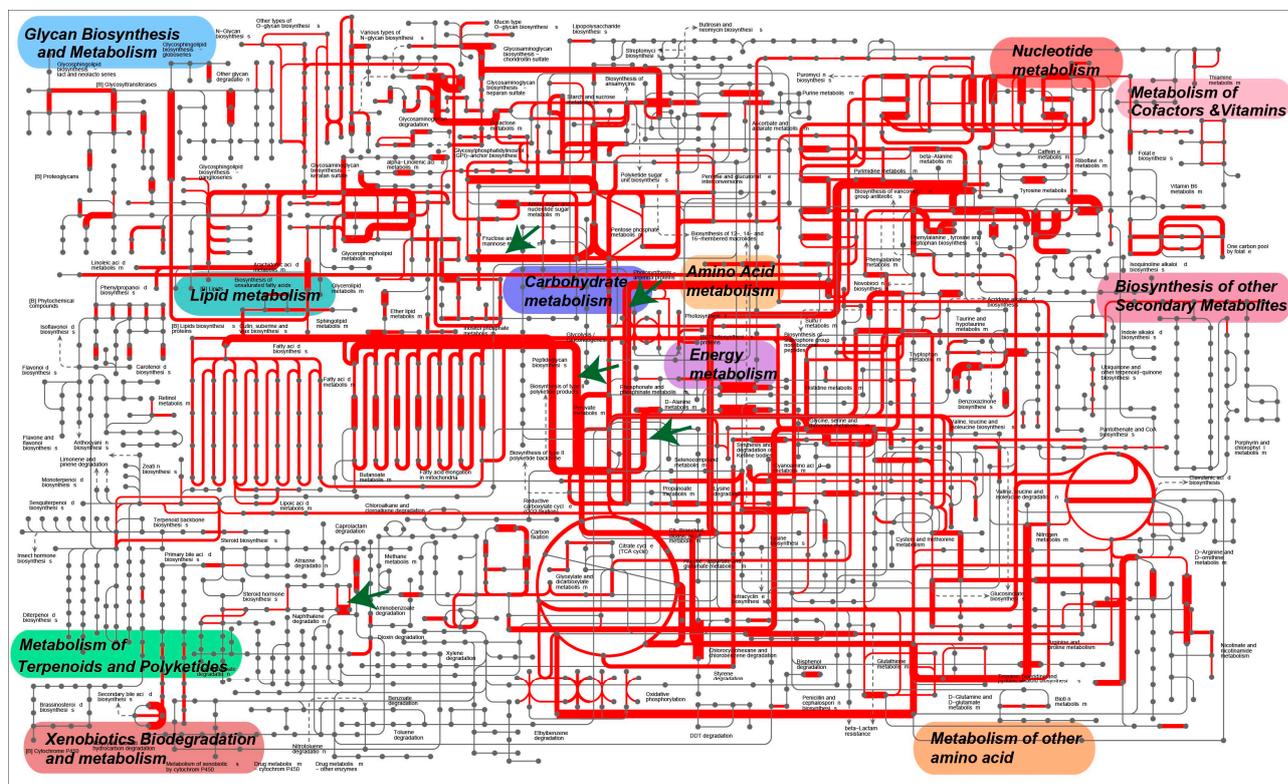


# Supplementary Materials: Transcriptome Sequencing and De Novo Assembly of Golden Cuttlefish *Sepia esculenta* Hoyle

Changlin Liu, Fazhen Zhao, Jing-ping Yan, Chun-sheng Liu, Siwei Liu and Siqing Chen



**Figure S1.** Metabolic pathways in KEGG encoded by unigenes and gene dose of individual KO ID. The metabolic enzymes encoded by expanded copies of unigenes were indicated with green arrows.