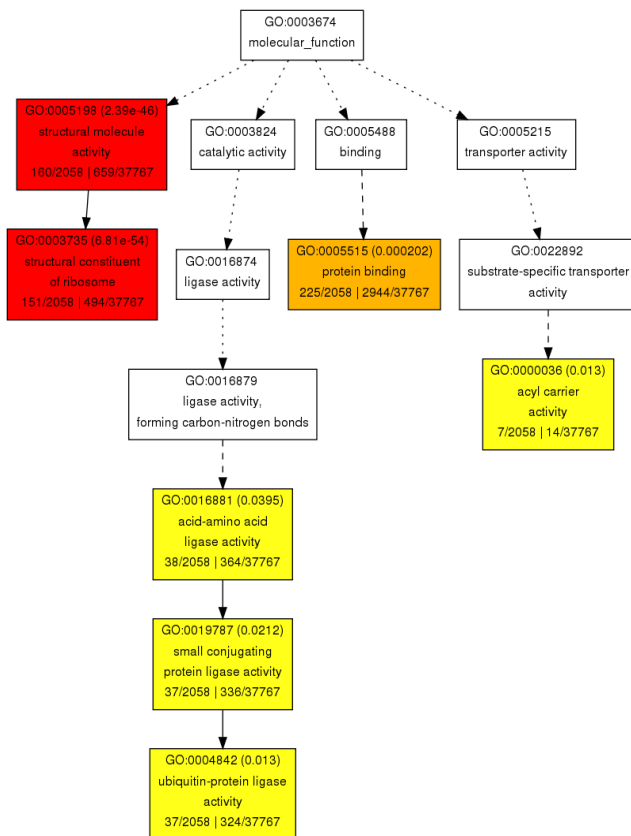


Figure S4



**Figure S4.** Gene Ontology (GO) analysis of mRNAs poorly translated during HS and affected in translation recovery in *hsp101* (2103/3309). GO groups highly enriched are presented in red. p-values are presented in parentheses for all groups.