

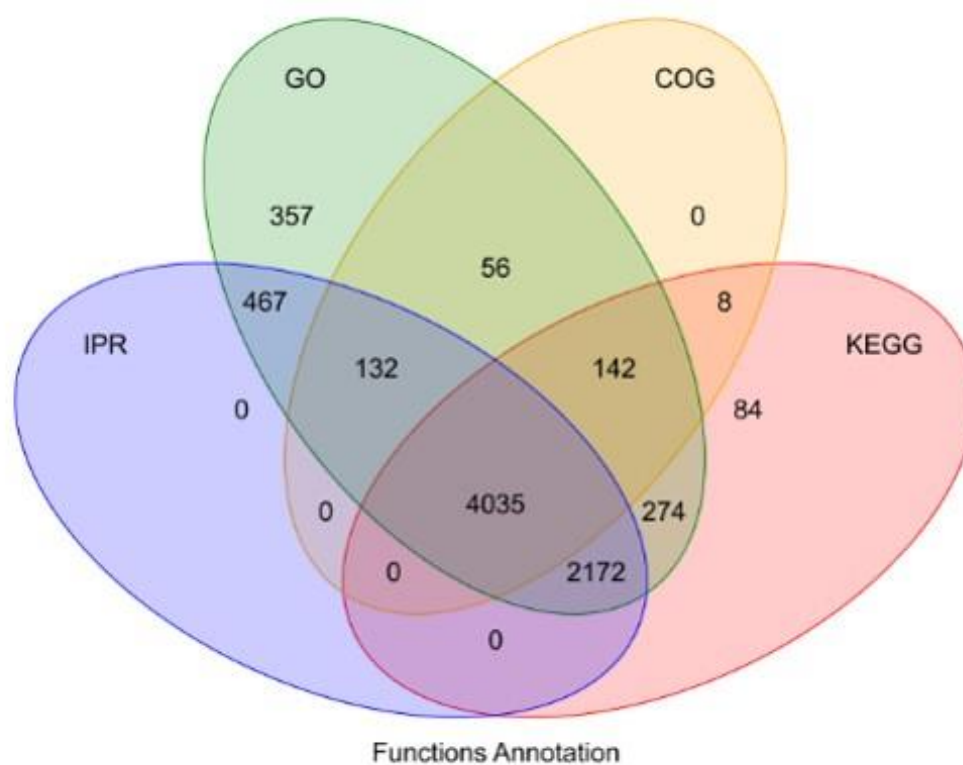


Fig. S3. The number of proteins annoated in four databases of proteome. GO, Gene Ontology; COG, Cluster of Orthologous Groups of Proteins; IPR, iProclass Protein Database; KEGG, Kyoto Encyclopedia of Genes and Genomes.