



Figure S1. Scatterplot of the number of metaproteins of each taxon (92) with the number of non-redundant human proteins found by the PSI-blast algorithm using the metaproteins. The color-coded information represents the number of Gene ontologies (GOs) found by the enrichment analysis.

Flores-Saiffe, et al. An ontological systems approach on human brain expression and metaproteomics. *Frontiers in Microbiology* (2017).