Figure S3. Distribution of conserved proteins in CoExp$^\text{Thr}$. The orthologs of many T. brucei proteins with known KEGG pathways cannot be unambiguously identified in L. infantum; this is particularly the case for ribosomal proteins whose sequence similarity to each other prevents unambiguous identification of orthologous partners. Proteins whose L. infantum orthologs can be unambiguously identified are shown by red nodes.