



**Figure S5. COG analysis of the IscR regulon, functional regulon, and indirect regulon.** Clusters of Orthologous Groups of proteins (COG) analysis of genes with a IscR ChIP-Seq peak (A), genes with a IscR ChIP-Seq peak and are differentially expressed between WT and the *iscR* mutant under at least one tested condition (B), and genes differentially expressed between WT and the *iscR* mutant but have no IscR ChIP-Seq peak (C).