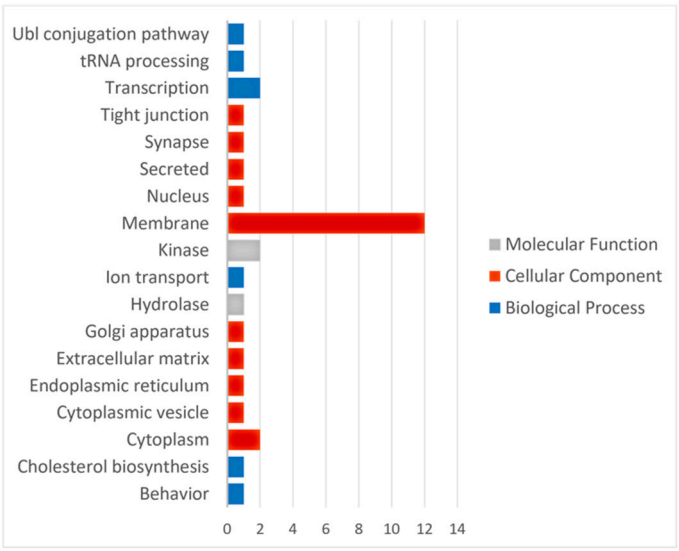

Supplementary Figure S2. Functional annotation visualization of 42 prioritized candidate genes (PCGs) for gene ontology (GO) terms relative to three main GO aspects, i.e., biological process, cellular component, and molecular function. *x*-axis, number of PCGs per GO term.