Females

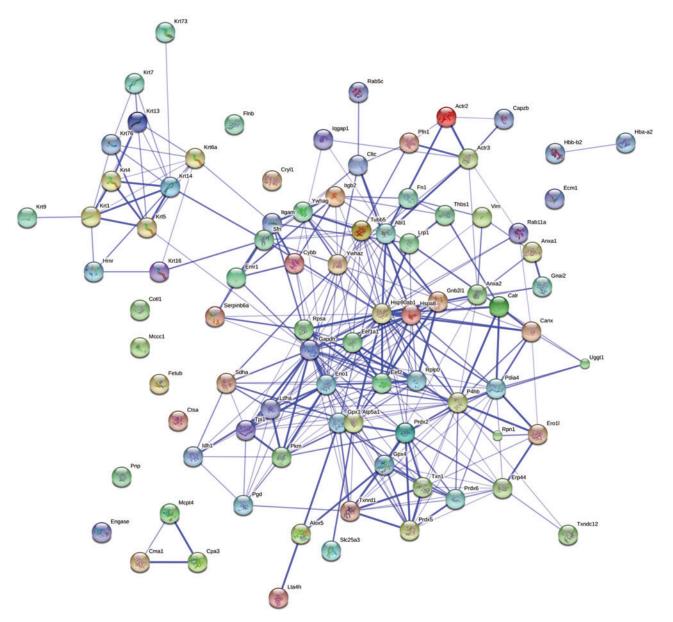**SUPPLEMENTARY FIG. S3.** **Functional cluster and gene ontology of all macrophage *S*-glutathionylated proteins identified in dyslipidemic female mice.** STRING (http://string-db/org) (58) analysis was utilized to identify the most significant processes showing the highest enrichment scores for *S*-glutathionylation in macrophages of female mice. The *top* two clusters were *Cell Differentiation* ($p = 8.9 \times 10^{-5}$) and *Inflammatory Response* ($p = 8.8 \times 10^{-4}$). STRING, Search Tool for the Retrieval of Interacting Genes/Proteins.