Figure S2. Pathway analysis of the effects of HFD on liver gene transcription in C57BL/6J and 129S6 mice. Only KEGG biological pathways identified by GSEA as significantly affected by HFD in at least one of the strains are reported. False discovery rate (FDR) q-values (<0.05) were used to test for statistically significant pathway upregulation (red) or downregulation (green) in response to fat feeding in 129S6 (*) and C57BL/6J ($) mice.