



Supplemental Figure 1. Results of weighted gene coexpression network analysis (WGCNA) in A. lateral (LH), B. dorsomedial (DMH) and C. arcuate nucleus (ARC) subregions of the hypothalamus. Presented are each module and the number of genes in that module. Values in the table represent the Pearson correlation interaction term between Control and Stress groups, with asterisks marking statistically significant differences: * = $p < 0.05$, ** = $p < 0.01$. Positive values are higher expression of the module in the Stress group, negative values are lower expression in the Stress group.