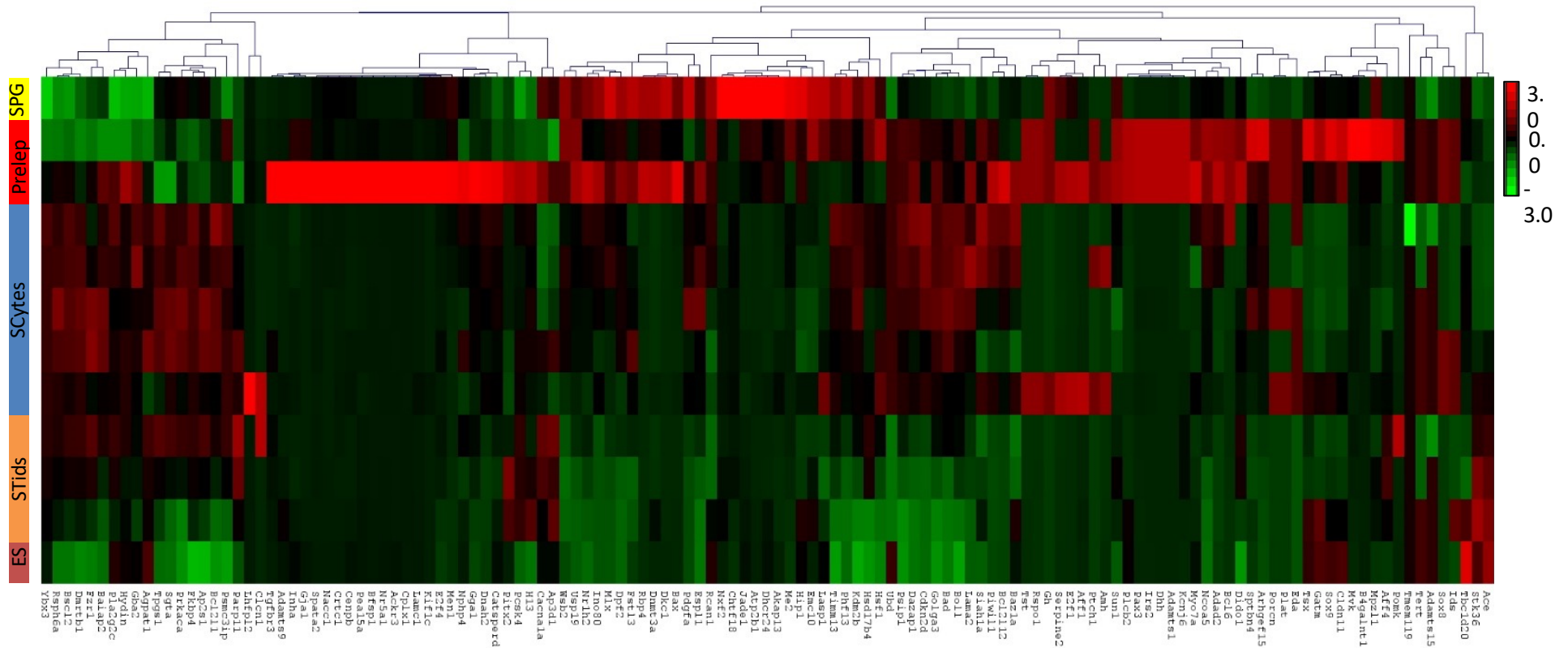




Supplementary Figure 5: Unsupervised hierarchical clustering of Green et al.'s dataset using 3 weeks DEGs (up-regulated) with male reproductive phenotypes. 129 genes found in Green *et al* dataset. Samples are denoted in rows and genes are denoted in columns. Red and green denote highly and weakly expressed genes, respectively.