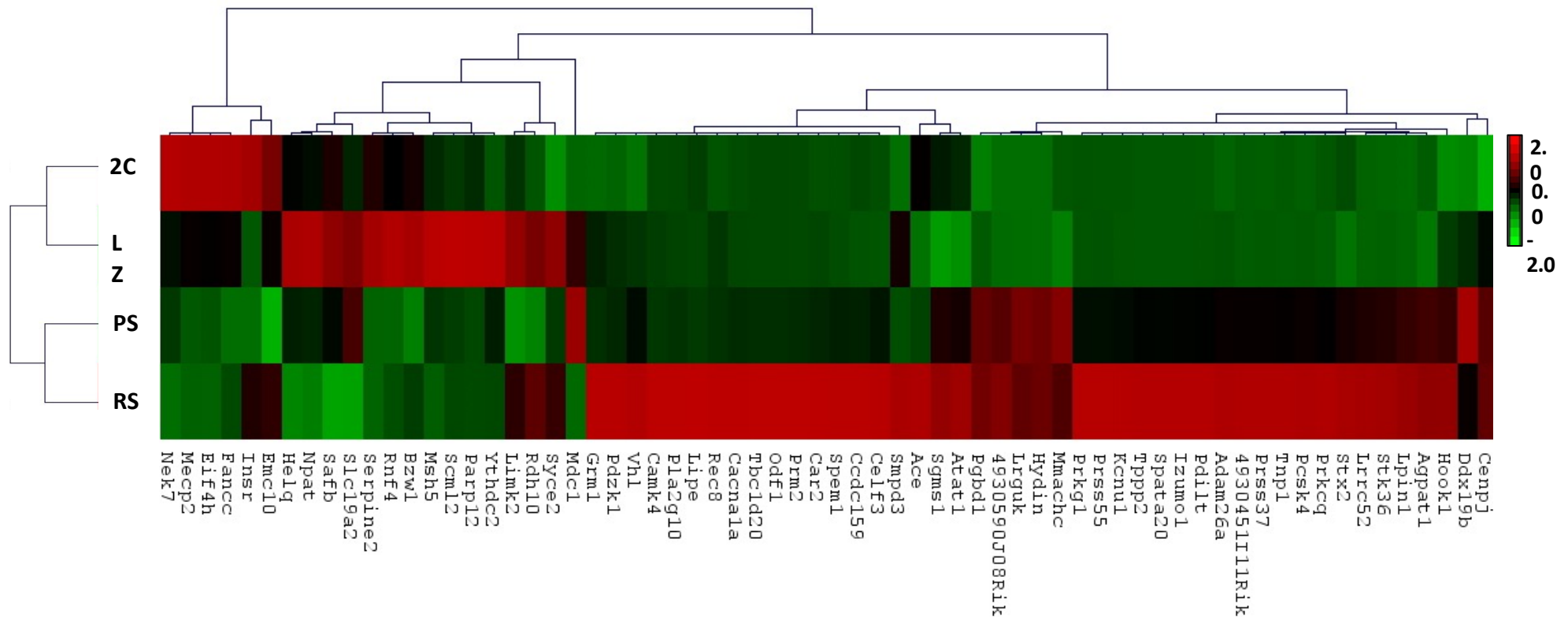




Supplementary Figure 4: Unsupervised hierarchical clustering of Cruz et al.'s dataset using 8 weeks DEGs (down-regulated) with male reproductive phenotypes. 66 genes found in Cruz *et al* dataset. Samples are denoted in rows and genes are denoted in columns. Red and green denote highly and weakly expressed genes, respectively.