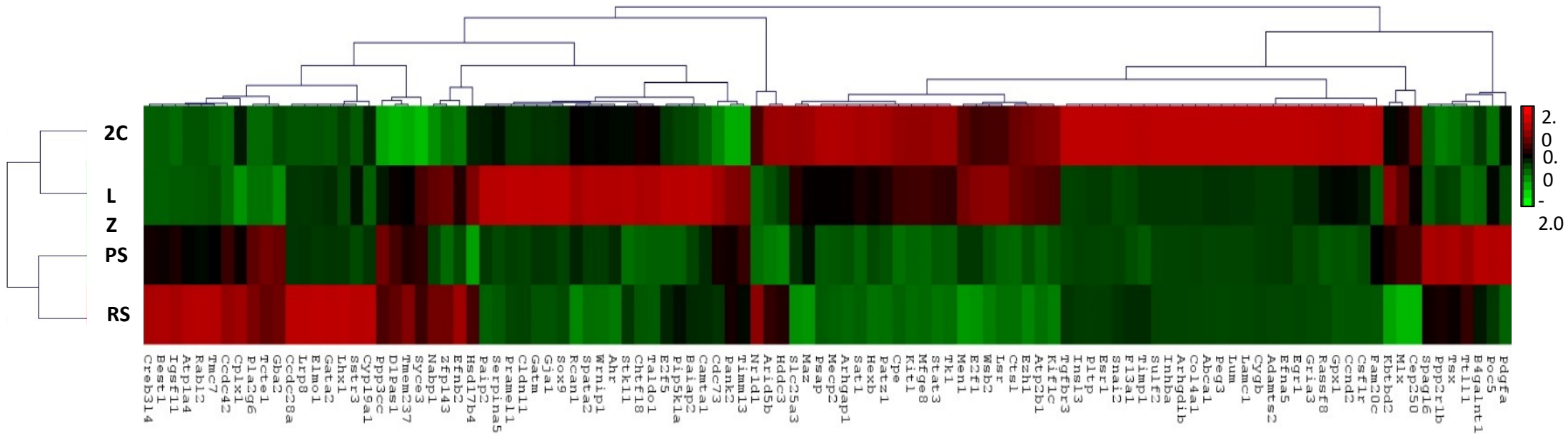




Supplementary Figure 3: Unsupervised hierarchical clustering of Cruz *et al.*'s dataset using 8 weeks DEGs (up-regulated) with male reproductive phenotypes. 106 genes found in Cruz *et al.* dataset. Samples are denoted in rows and genes are denoted in columns. Red and green denote highly and weakly expressed genes, respectively.