



Gene expression data visualization showing a heatmap with a dendrogram on the left side, indicating hierarchical clustering of samples. The color scale ranges from dark blue (low expression) to dark red (high expression). The samples are listed on the right side of the heatmap, grouped into several categories: Somatic.Primary, Somatic.Tissue, iPS, ES, and ES.parthenote. The dendrogram shows that samples within the same category are highly similar, while samples from different categories show distinct expression patterns. The heatmap is divided into several distinct blocks of color, indicating clusters of genes or samples with similar expression profiles.