

The figure displays a hierarchical clustering dendrogram at the top, showing the relationship between various samples. The samples are labeled on the left side of the dendrogram, including HSB168.V1C, HSB168.ATC, HSB168.SBC, and HSB168.TIC. Below the dendrogram is a heatmap representing the expression levels of various genes across different samples. The color scale ranges from red (high expression) to blue (low expression). The bottom part of the figure shows a bar chart representing the expression levels of specific genes across different samples, with a color scale from red to blue.

GrossRegion