

Assessing Stability of Gene Selection in Microarray Data Analysis

SIMULDIST

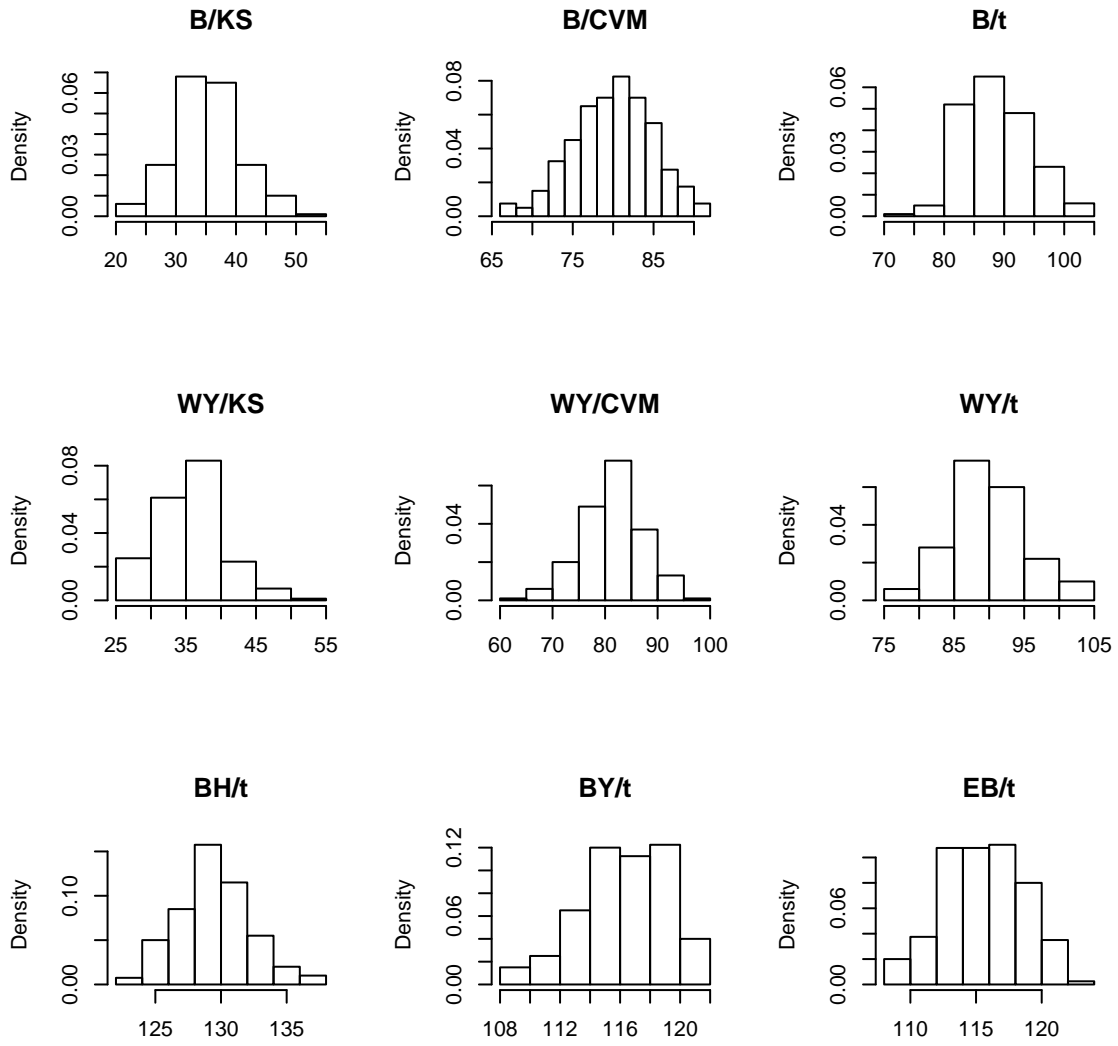


Figure 1: Histograms of the number of selected genes across 200 simulation runs constructed for S15.

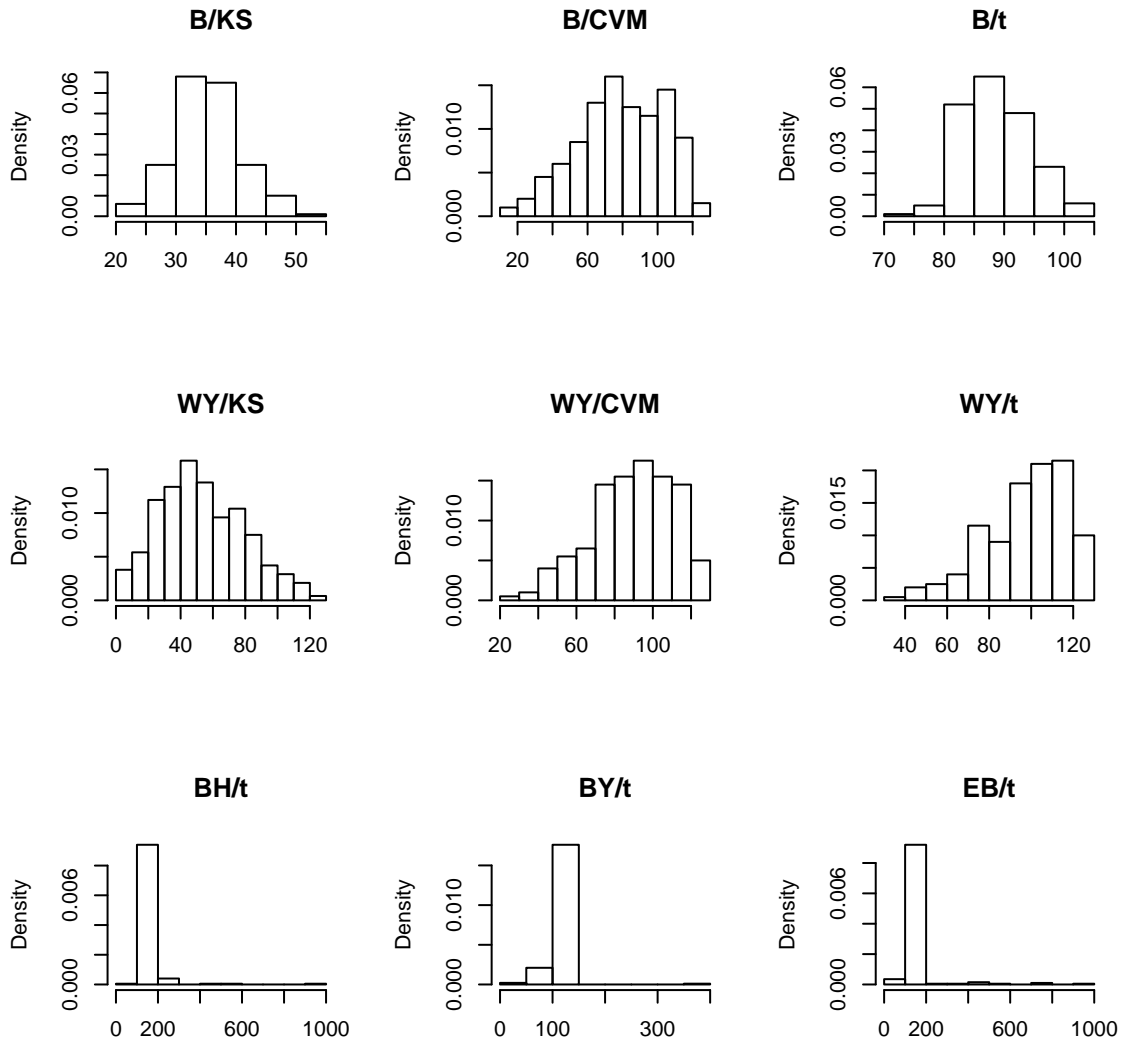


Figure 2: Histograms of the number of selected genes across 200 simulation runs constructed for S15COR.

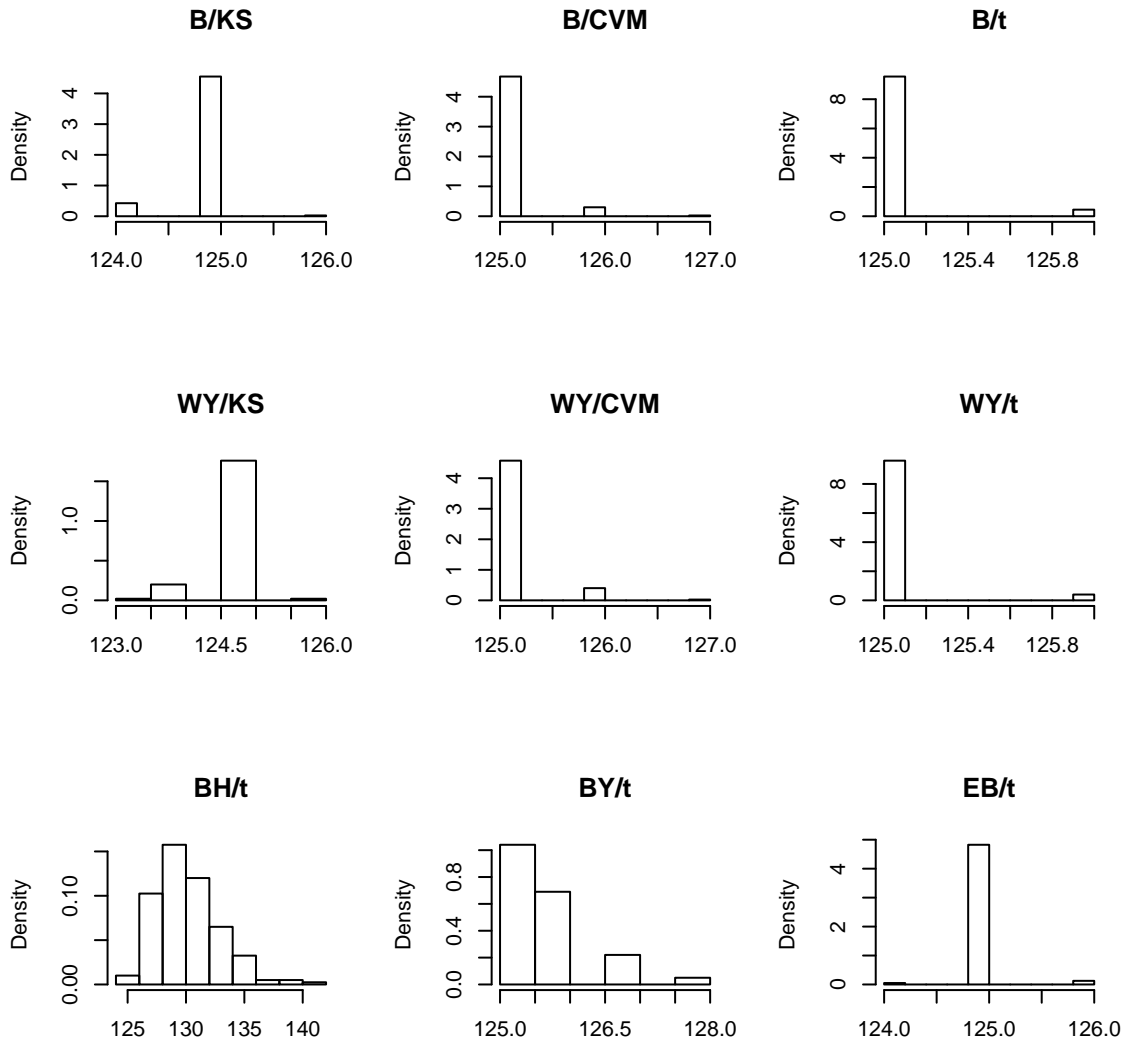


Figure 3: Histograms of the number of selected genes across 200 simulation runs constructed for S43.

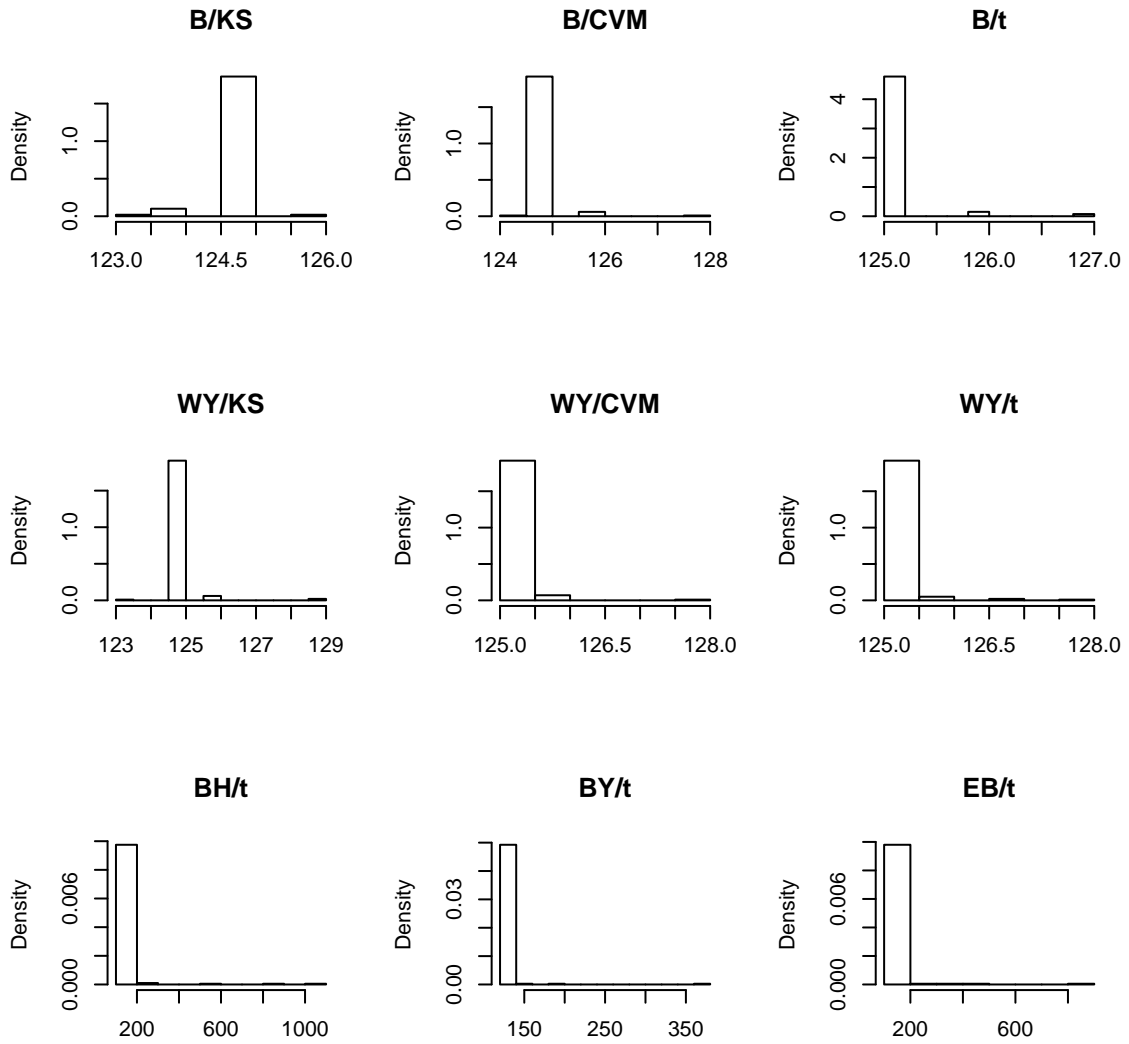


Figure 4: Histograms of the number of selected genes across 200 simulation runs constructed for S43COR.