

Histogram of stranded vs poly-A paired analysis ($k = 1$)

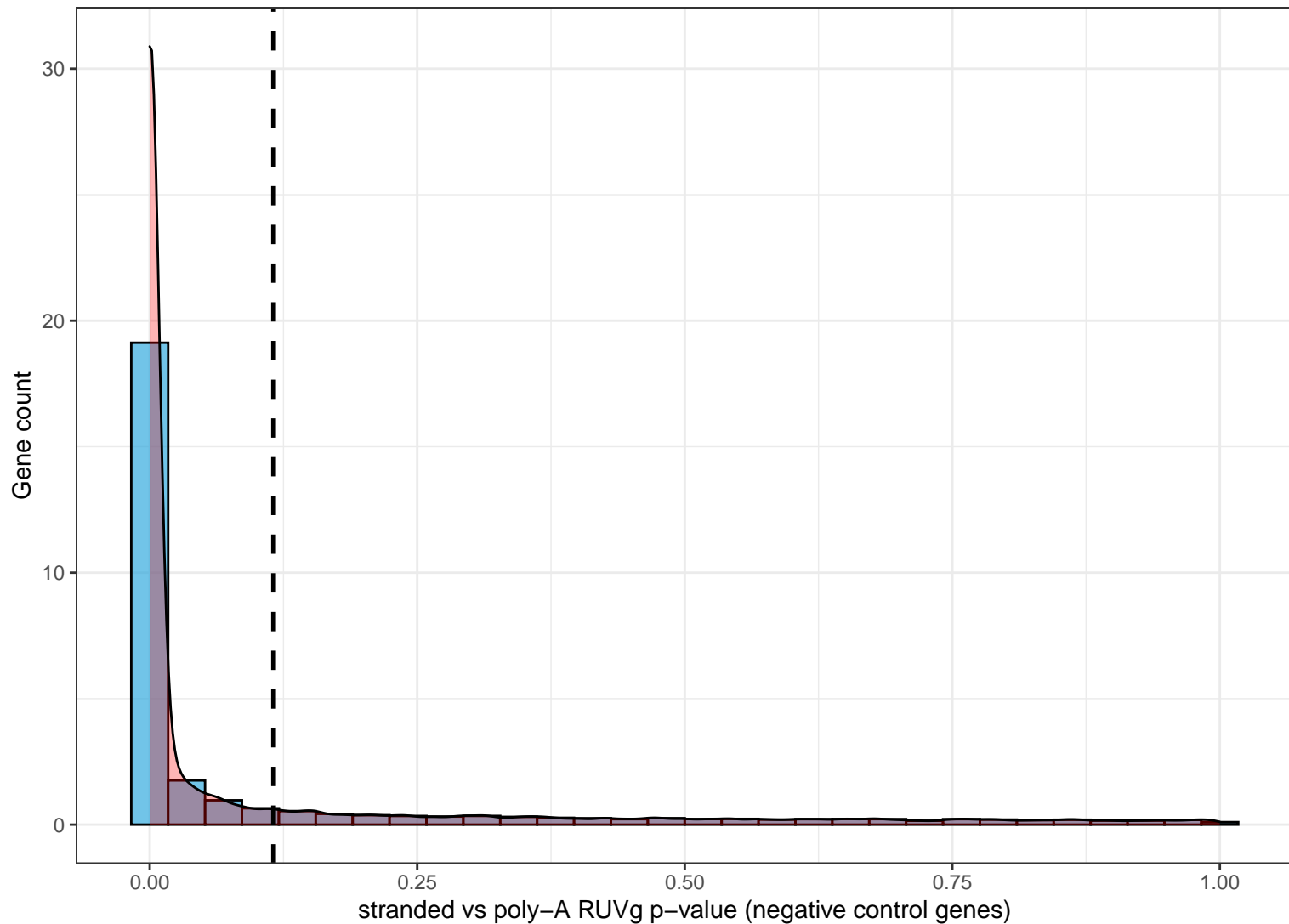
Total number of genes: 15104

10847 genes have p-value < 0.05

4257 genes have p-value ≥ 0.05

10515 genes have BH FDR < 0.05

4589 genes have BH FDR ≥ 0.05



Histogram of stranded vs poly-A paired analysis ($k = 2$)

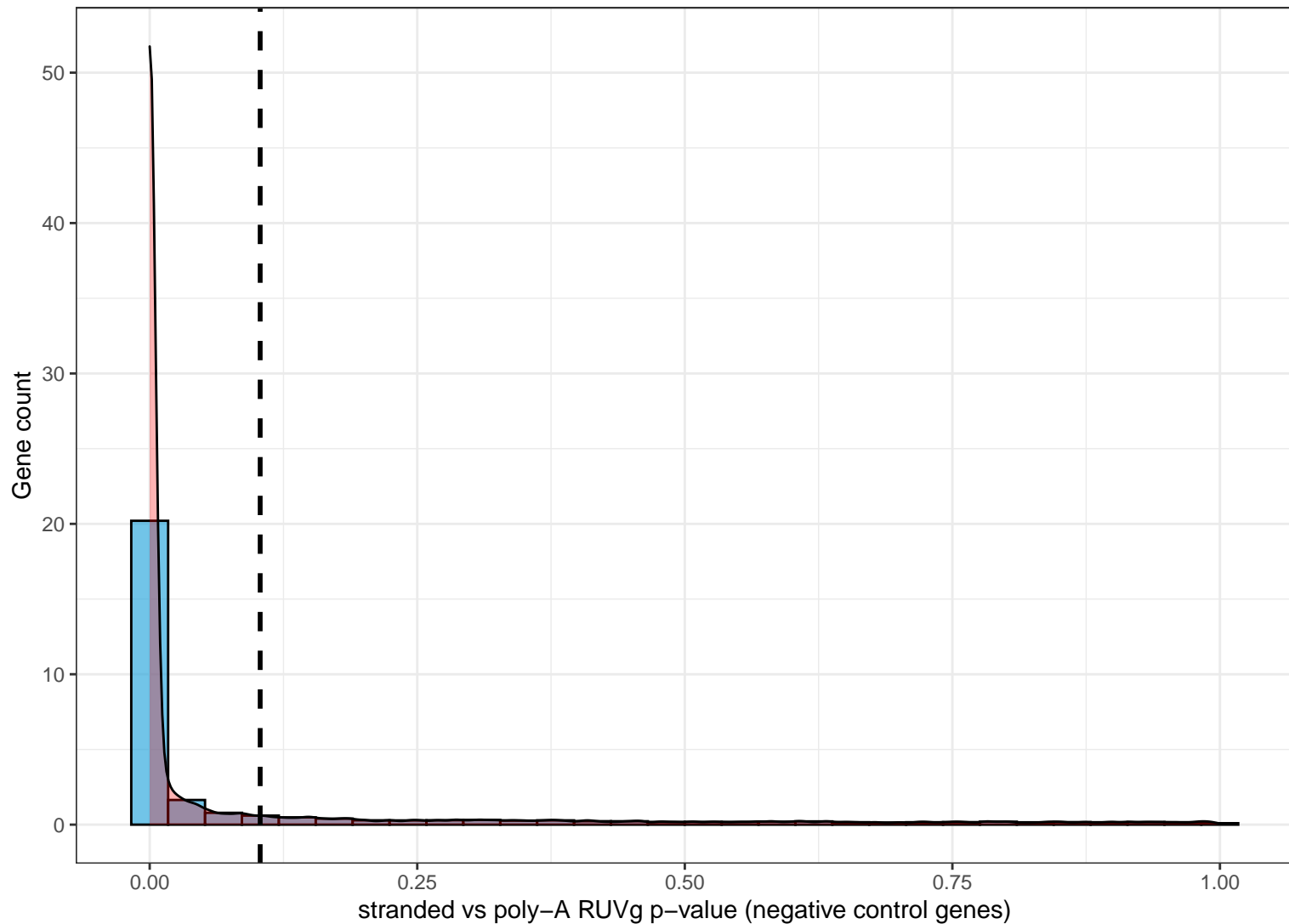
Total number of genes: 15104

11344 genes have $p\text{-value} < 0.05$

3760 genes have $p\text{-value} \geq 0.05$

11070 genes have BH FDR < 0.05

4034 genes have BH FDR ≥ 0.05



Histogram of stranded vs poly-A paired analysis ($k = 3$)

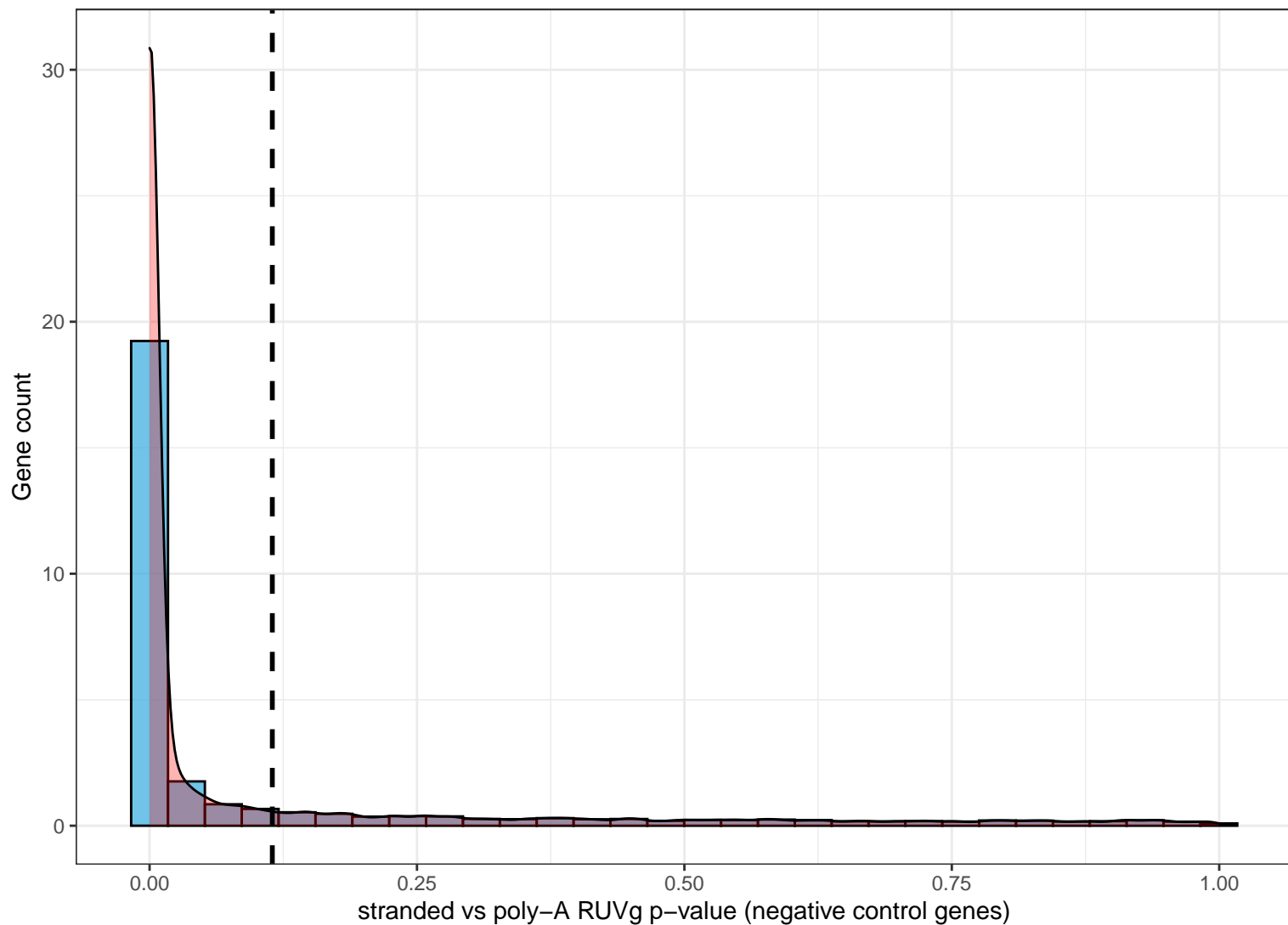
Total number of genes: 15104

10891 genes have p-value < 0.05

4213 genes have p-value ≥ 0.05

10583 genes have BH FDR < 0.05

4521 genes have BH FDR ≥ 0.05



Histogram of stranded vs poly-A paired analysis ($k = 4$)

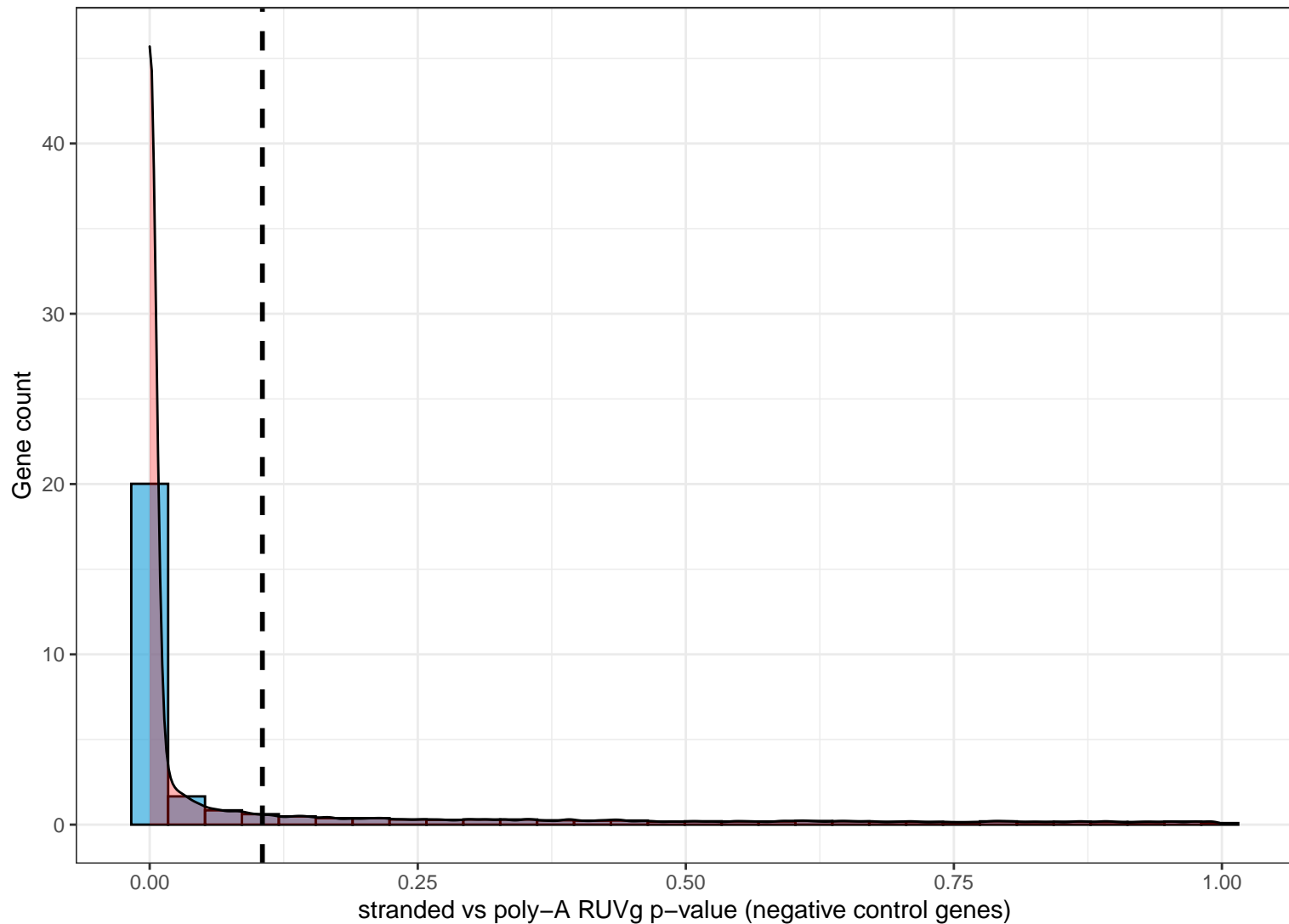
Total number of genes: 15104

11242 genes have p-value < 0.05

3862 genes have p-value ≥ 0.05

10963 genes have BH FDR < 0.05

4141 genes have BH FDR ≥ 0.05



Histogram of stranded vs poly-A paired analysis ($k = 5$)

Total number of genes: 15104

11250 genes have p-value < 0.05

3854 genes have p-value ≥ 0.05

10965 genes have BH FDR < 0.05

4139 genes have BH FDR ≥ 0.05

