

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

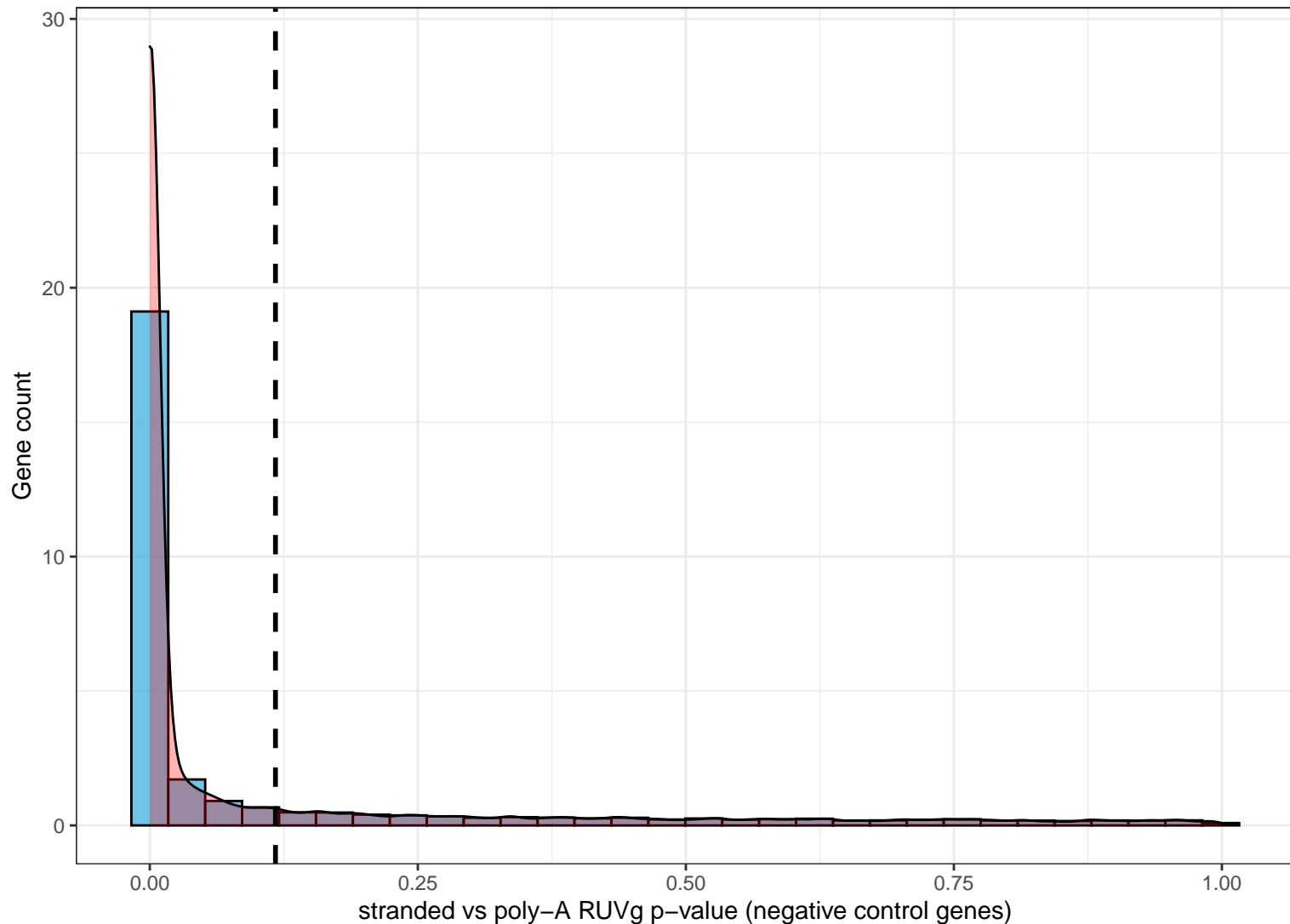
Total number of genes: 15104

10808 genes have p-value < 0.05

4296 genes have p-value ≥ 0.05

10481 genes have BH FDR < 0.05

4623 genes have BH FDR ≥ 0.05



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

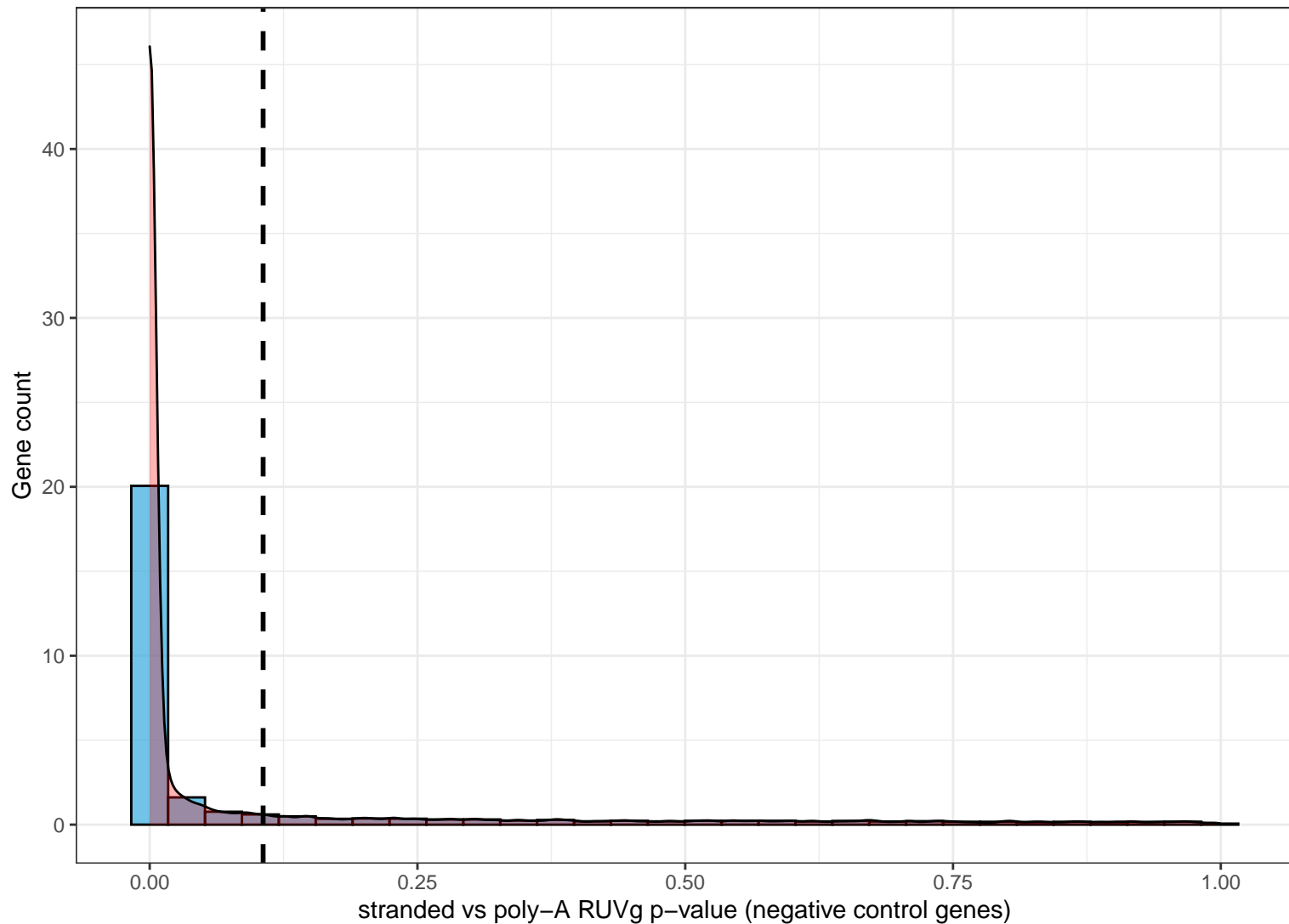
Total number of genes: 15104

11249 genes have $p\text{-value} < 0.05$

3855 genes have $p\text{-value} \geq 0.05$

10984 genes have BH FDR < 0.05

4120 genes have BH FDR ≥ 0.05



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

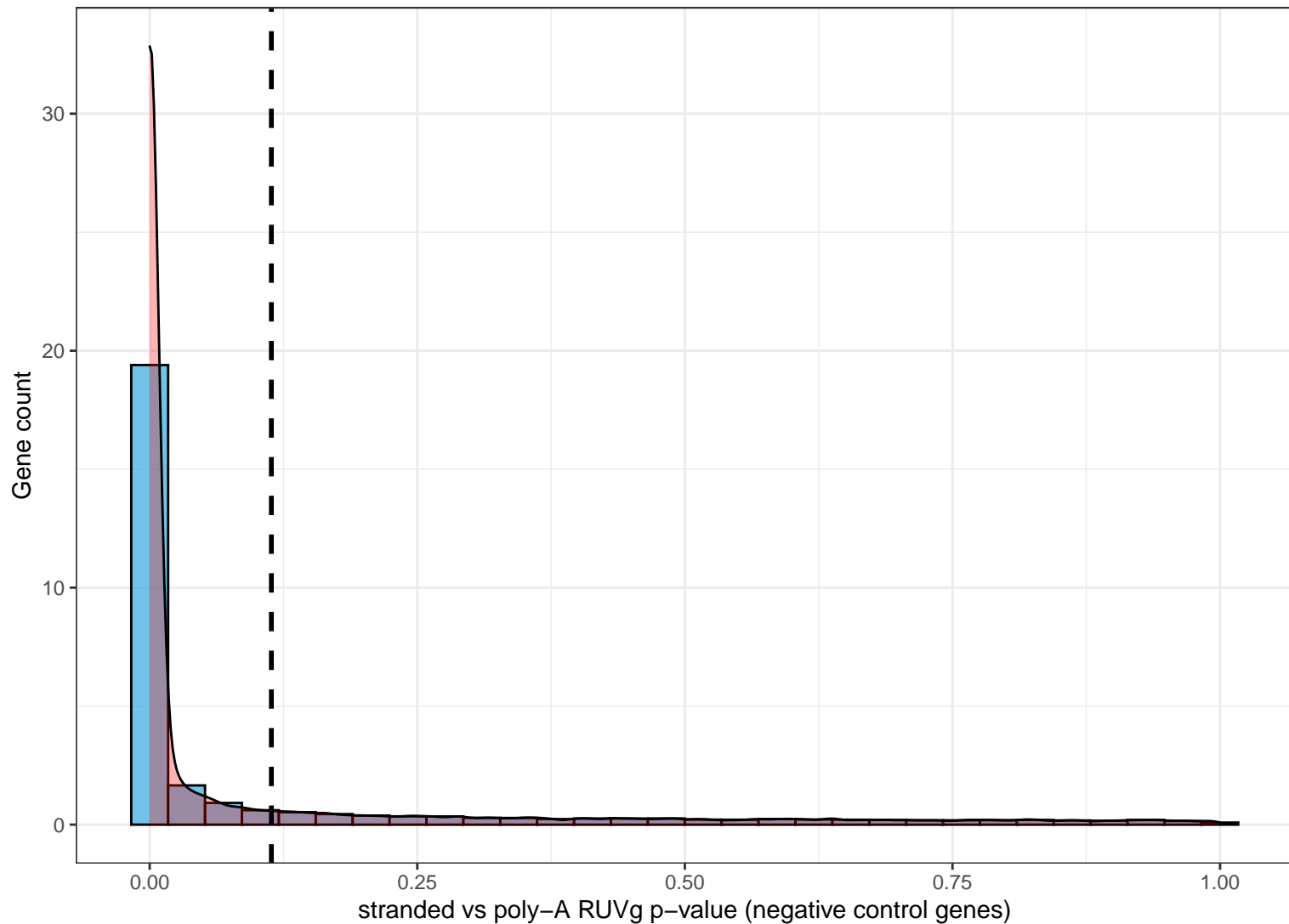
Total number of genes: 15104

10930 genes have p-value < 0.05

4174 genes have p-value ≥ 0.05

10627 genes have BH FDR < 0.05

4477 genes have BH FDR ≥ 0.05



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

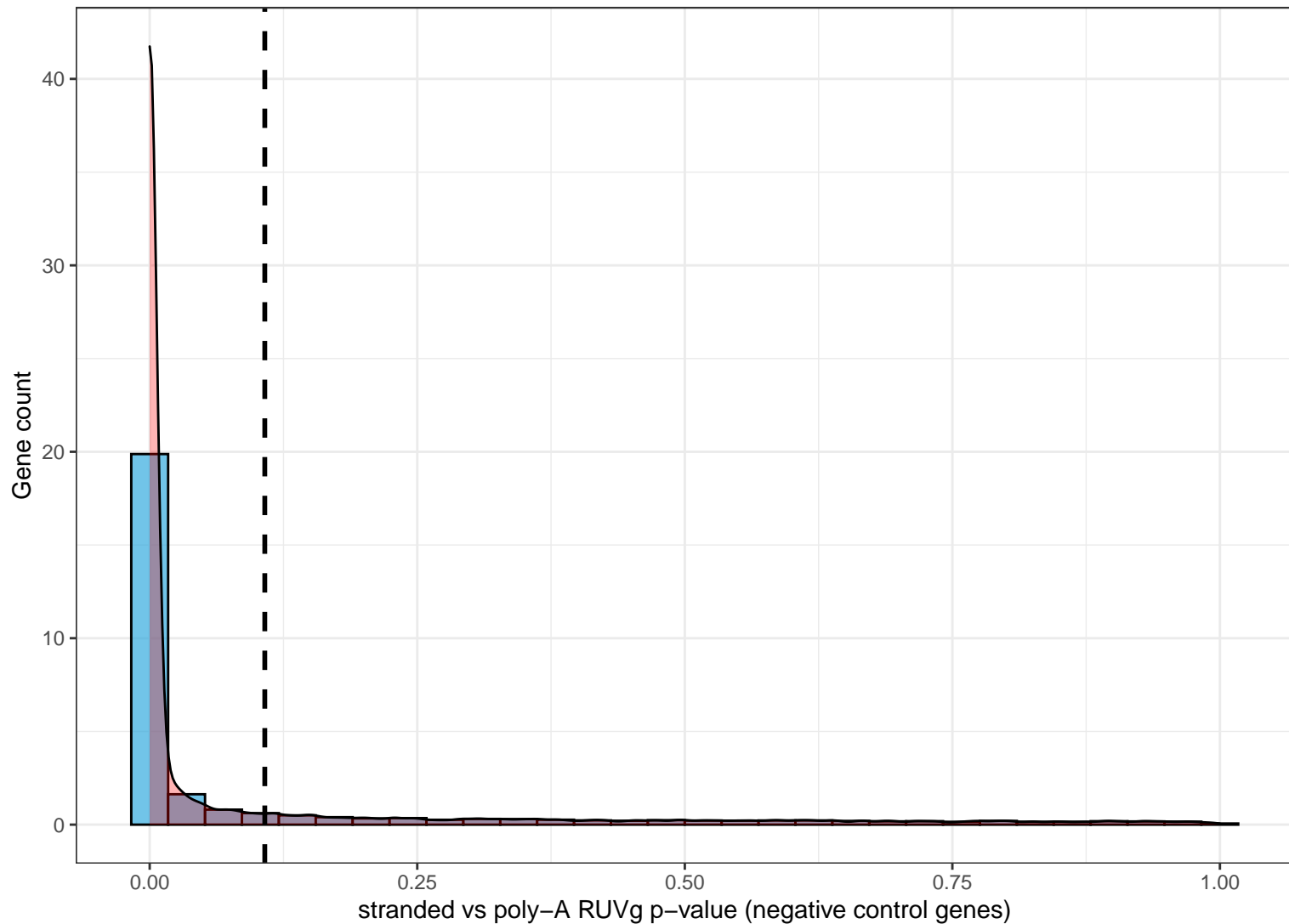
Total number of genes: 15104

11169 genes have p-value < 0.05

3935 genes have p-value ≥ 0.05

10898 genes have BH FDR < 0.05

4206 genes have BH FDR ≥ 0.05



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

Total number of genes: 15104

11165 genes have p-value < 0.05

3939 genes have p-value ≥ 0.05

10889 genes have BH FDR < 0.05

4215 genes have BH FDR ≥ 0.05

