

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

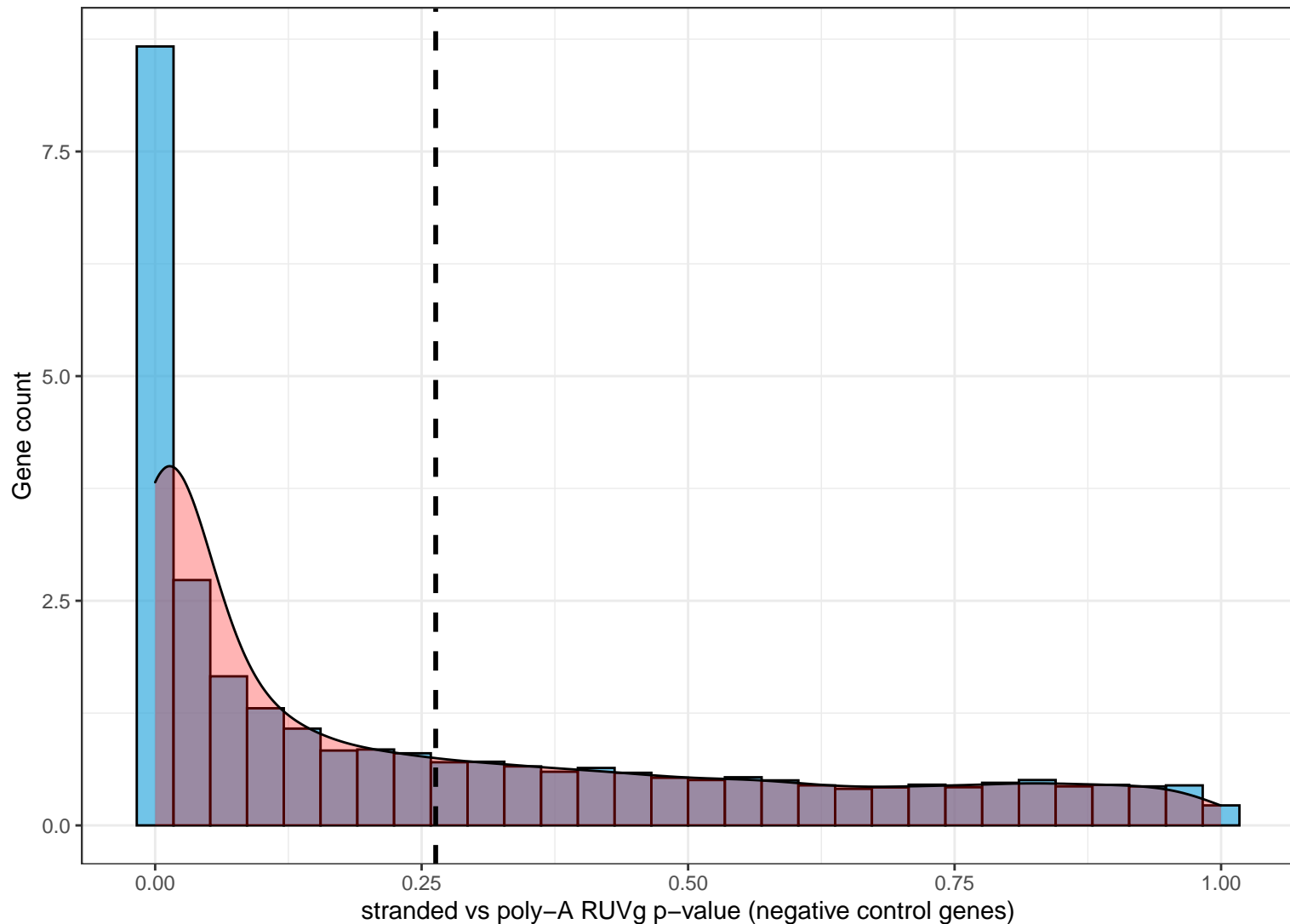
Total number of genes: 15104

5881 genes have p-value < 0.05

9223 genes have p-value ≥ 0.05

4324 genes have BH FDR < 0.05

10780 genes have BH FDR ≥ 0.05



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

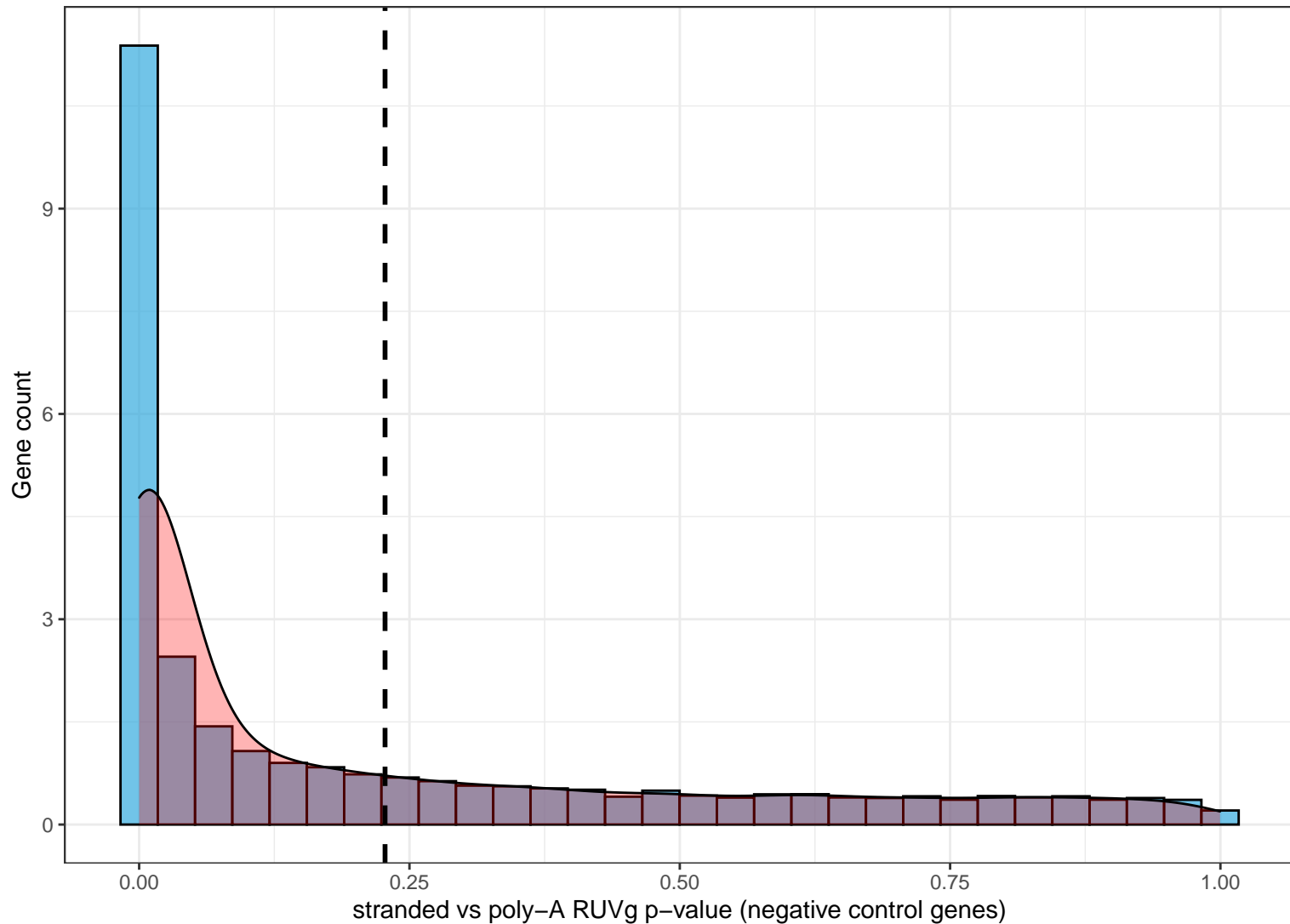
Total number of genes: 15104

7160 genes have p-value < 0.05

7944 genes have p-value ≥ 0.05

6074 genes have BH FDR < 0.05

9030 genes have BH FDR ≥ 0.05



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

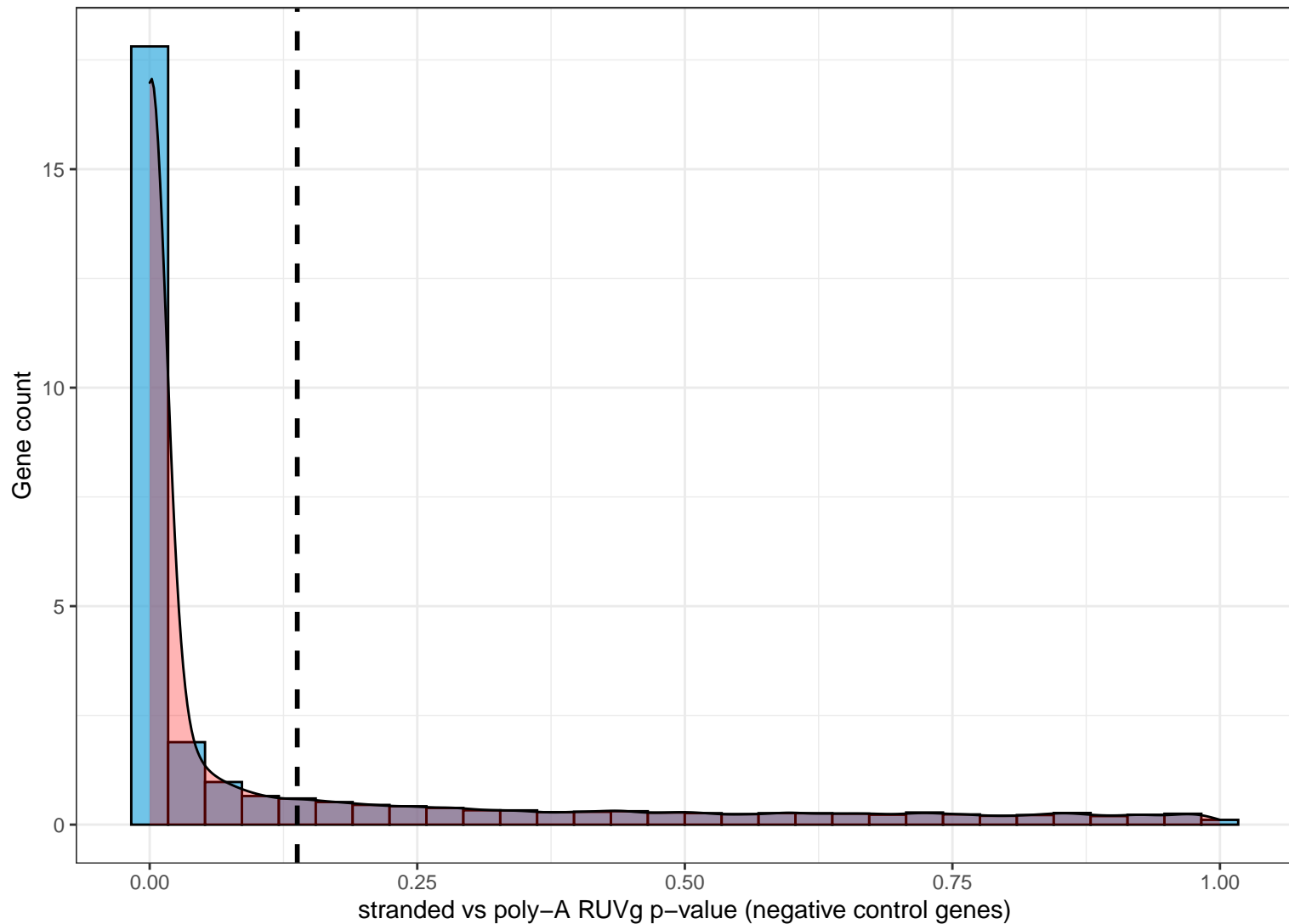
Total number of genes: 15104

10226 genes have p-value < 0.05

4878 genes have p-value ≥ 0.05

9801 genes have BH FDR < 0.05

5303 genes have BH FDR ≥ 0.05



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

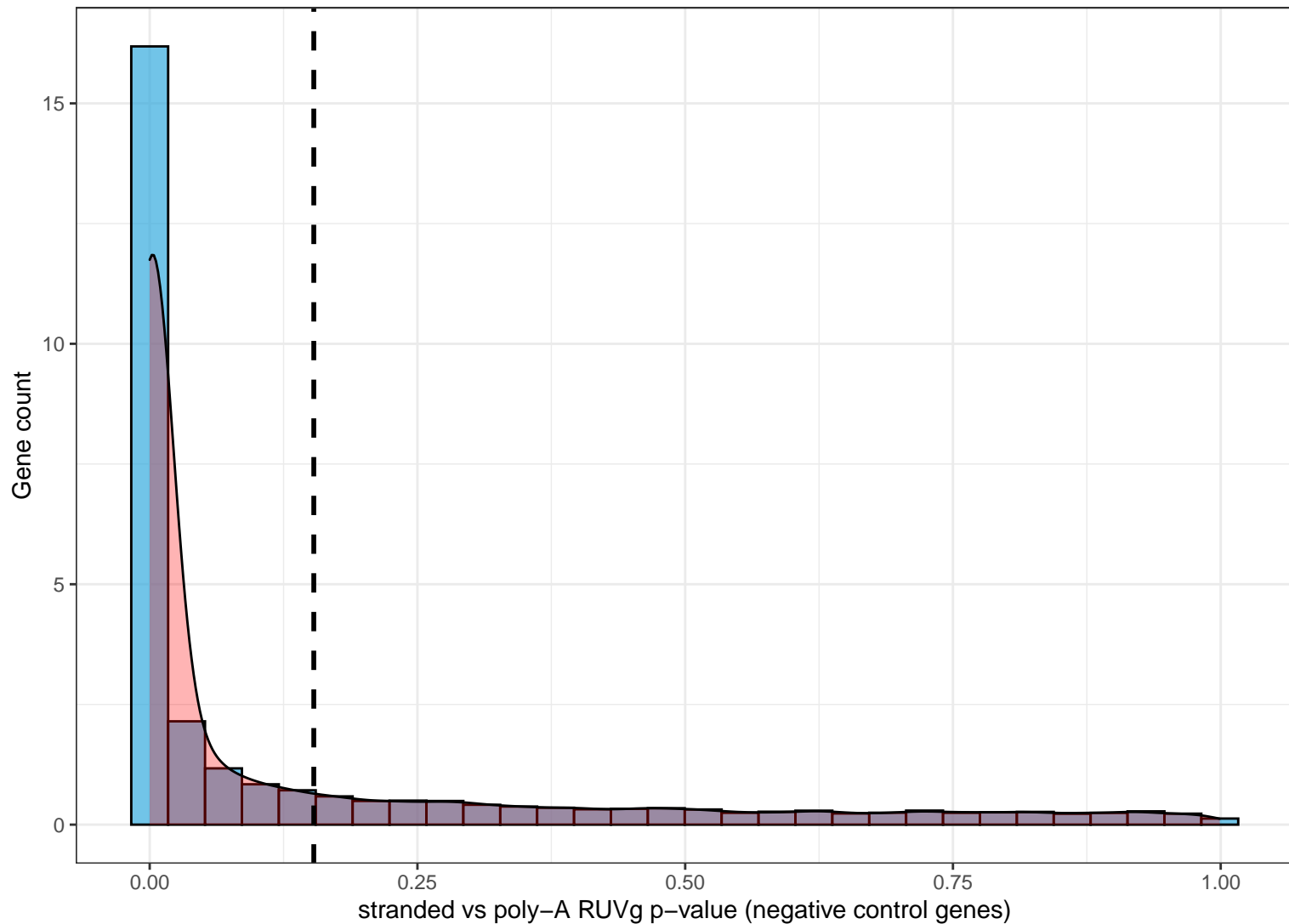
Total number of genes: 15104

9498 genes have p-value < 0.05

5606 genes have p-value ≥ 0.05

8960 genes have BH FDR < 0.05

6144 genes have BH FDR ≥ 0.05



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

Total number of genes: 15104

10864 genes have p-value < 0.05

4240 genes have p-value ≥ 0.05

10580 genes have BH FDR < 0.05

4524 genes have BH FDR ≥ 0.05

