

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

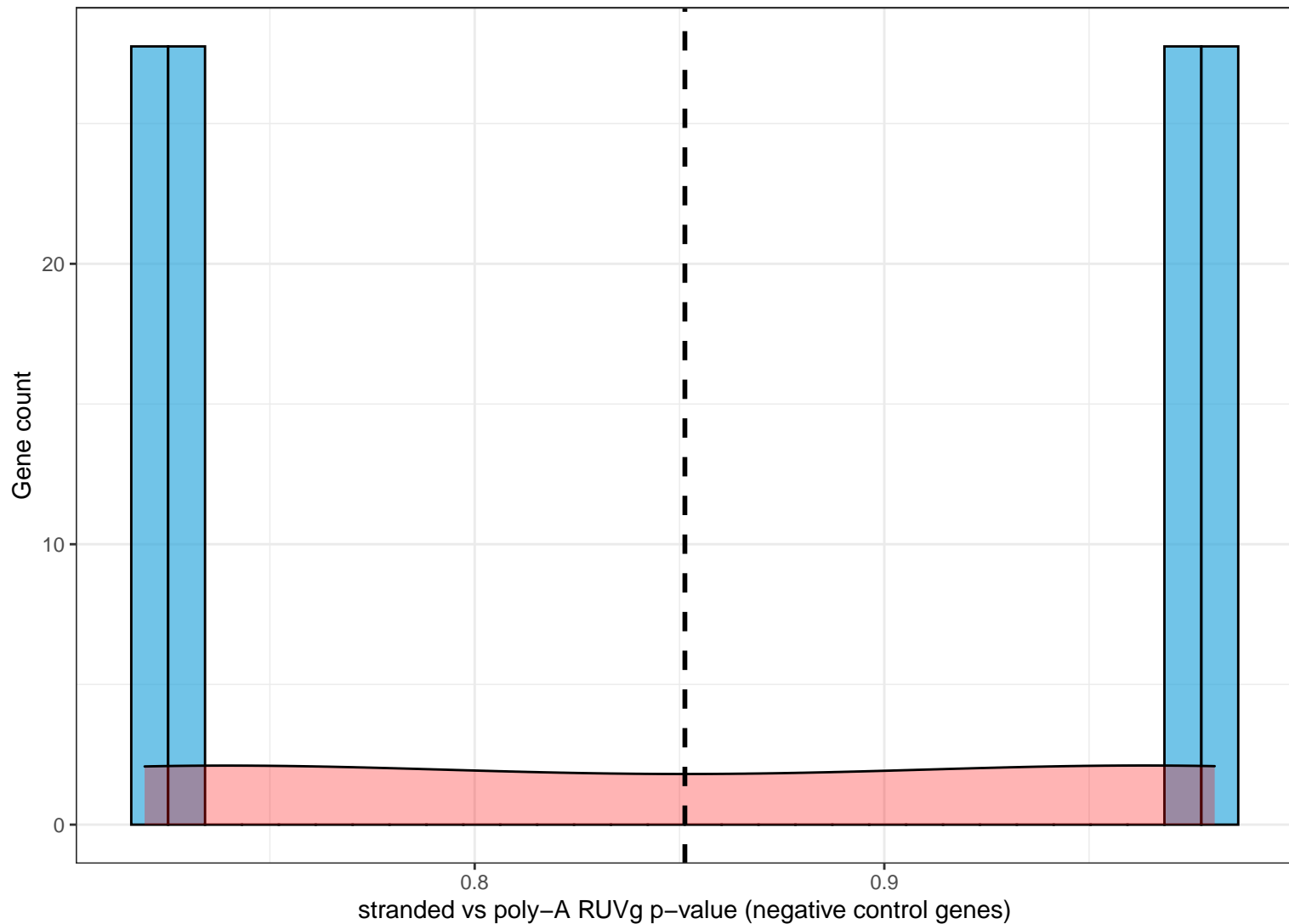
Total number of genes: 4

0 genes have p-value < 0.05

4 genes have p-value ≥ 0.05

0 genes have BH FDR < 0.05

4 genes have BH FDR ≥ 0.05



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

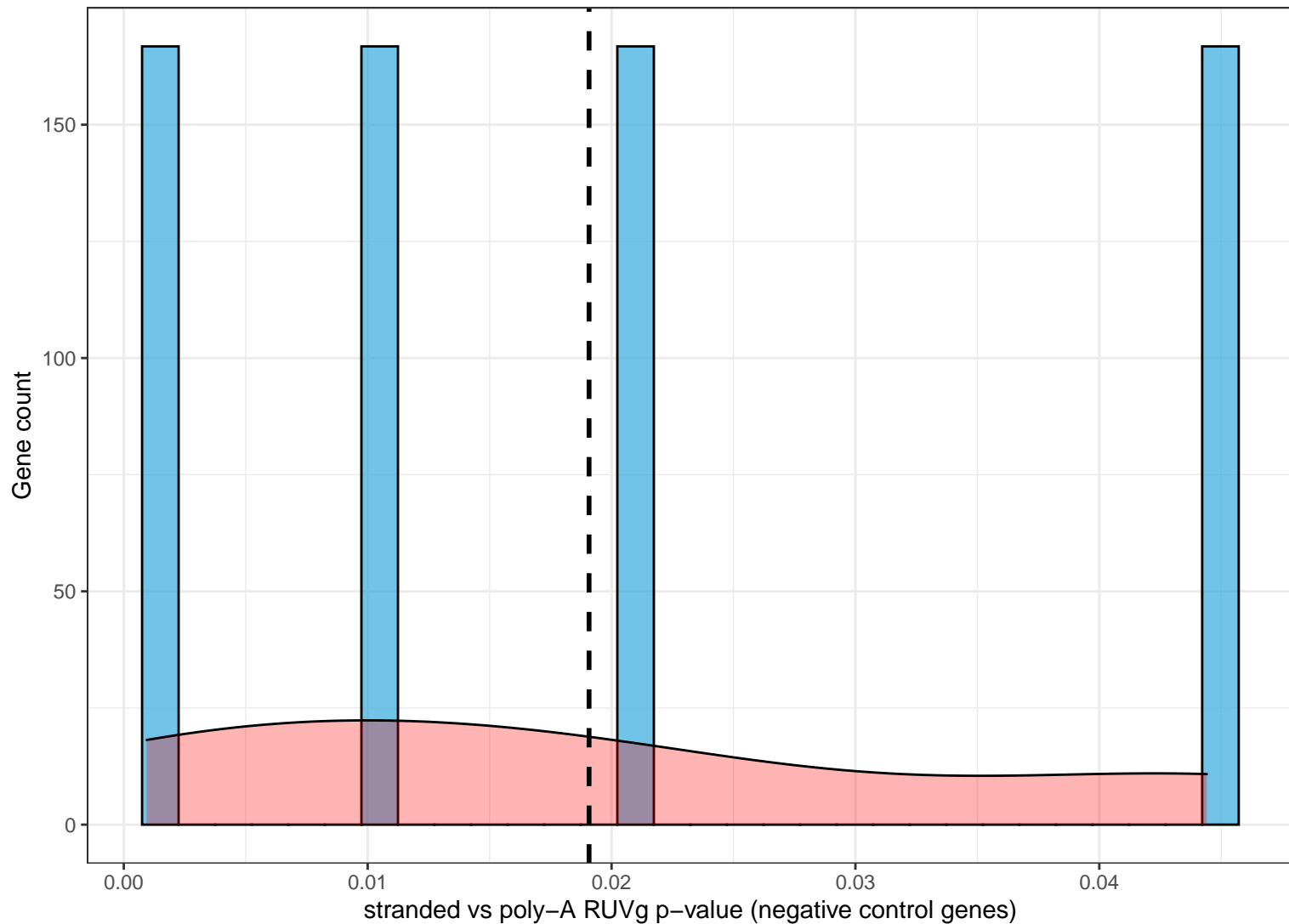
Total number of genes: 4

4 genes have p-value < 0.05

0 genes have p-value >= 0.05

0 genes have BH FDR < 0.05

4 genes have BH FDR >= 0.05



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

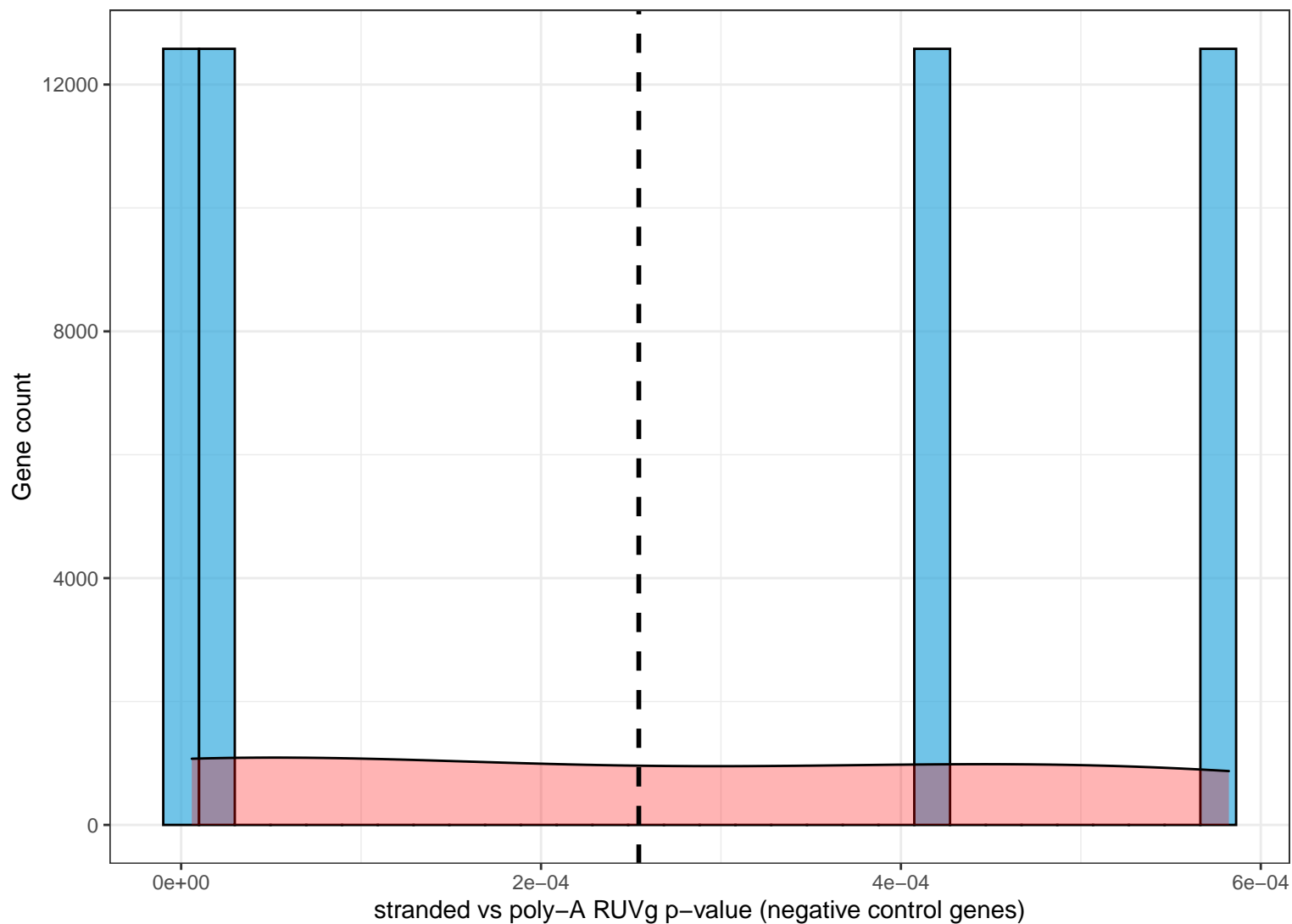
Total number of genes: 4

4 genes have p-value < 0.05

0 genes have p-value >= 0.05

4 genes have BH FDR < 0.05

0 genes have BH FDR >= 0.05



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

Total number of genes: 4

4 genes have p-value < 0.05

0 genes have p-value ≥ 0.05

4 genes have BH FDR < 0.05

0 genes have BH FDR ≥ 0.05

