

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

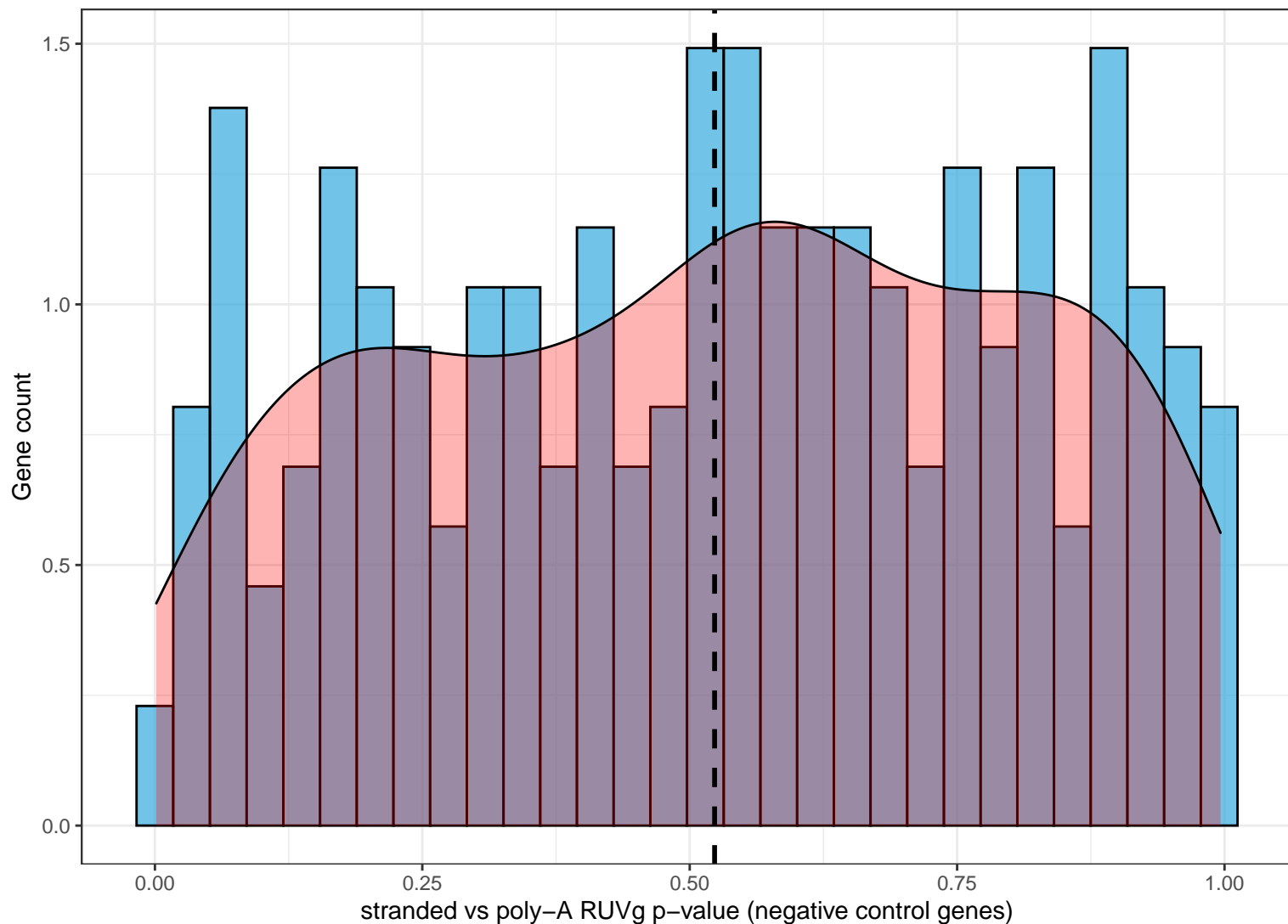
Total number of genes: 254

9 genes have p -value < 0.05

245 genes have p -value ≥ 0.05

0 genes have BH FDR < 0.05

254 genes have BH FDR ≥ 0.05



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

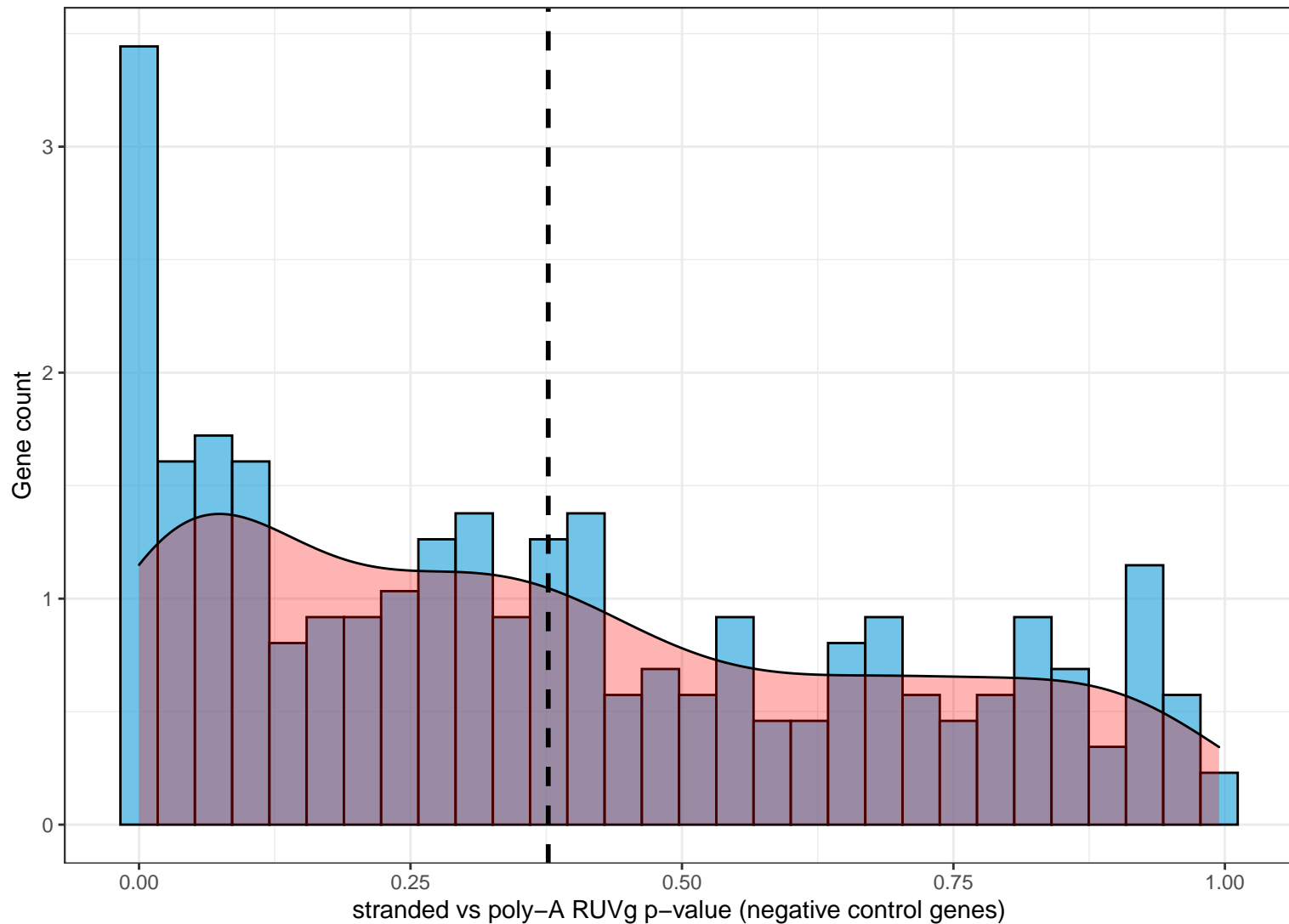
Total number of genes: 254

42 genes have p -value < 0.05

212 genes have p -value ≥ 0.05

10 genes have BH FDR < 0.05

244 genes have BH FDR ≥ 0.05



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

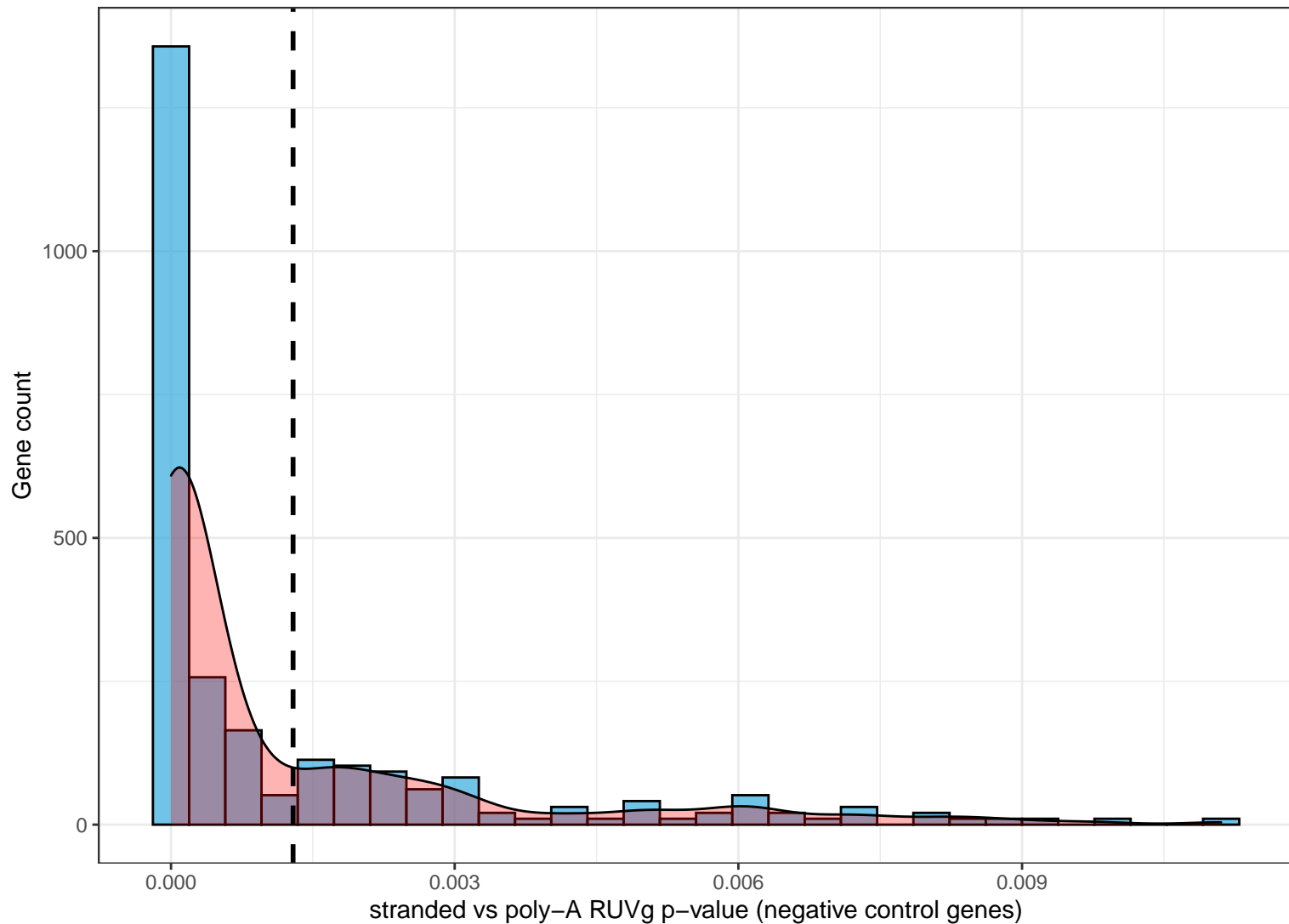
Total number of genes: 254

254 genes have p-value < 0.05

0 genes have p-value ≥ 0.05

251 genes have BH FDR < 0.05

3 genes have BH FDR ≥ 0.05



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

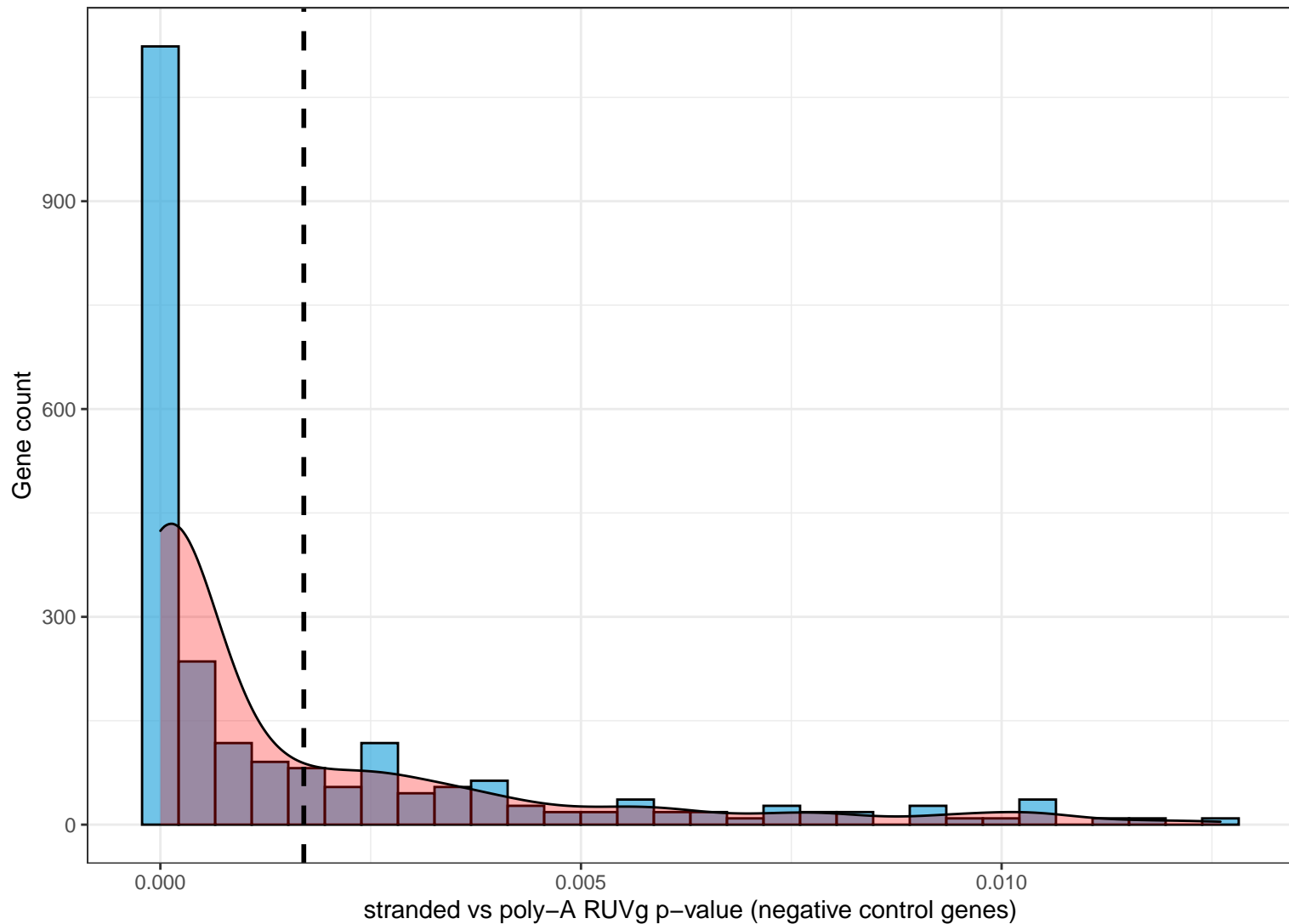
Total number of genes: 254

254 genes have p-value < 0.05

0 genes have p-value >= 0.05

238 genes have BH FDR < 0.05

16 genes have BH FDR >= 0.05



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

Total number of genes: 254

254 genes have p-value < 0.05

0 genes have p-value >= 0.05

249 genes have BH FDR < 0.05

5 genes have BH FDR >= 0.05

