

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

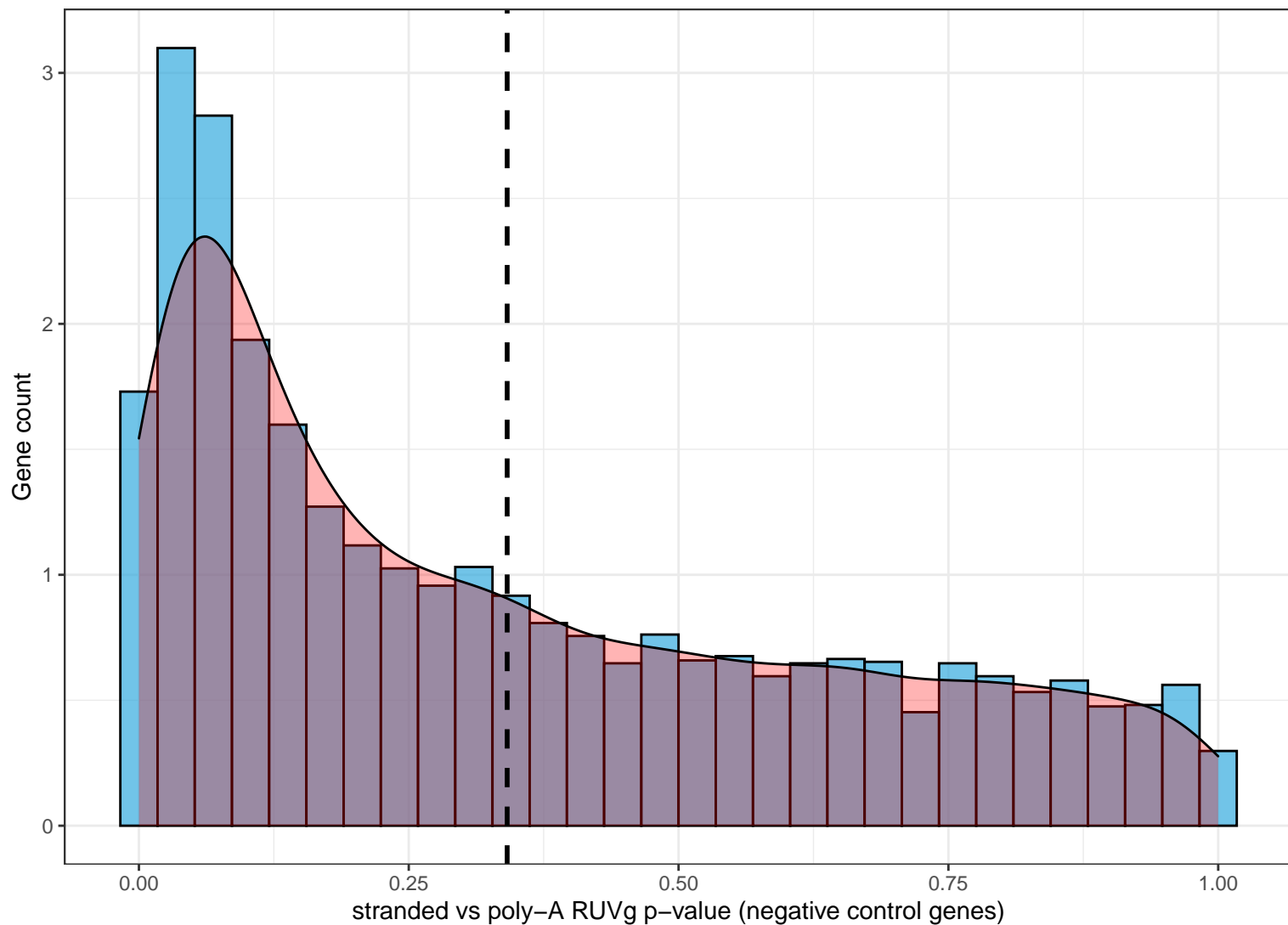
Total number of genes: 5063

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

4244 genes have $p\text{-value} \geq 0.05$

568 genes have BH FDR < 0.05

4495 genes have BH FDR ≥ 0.05



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

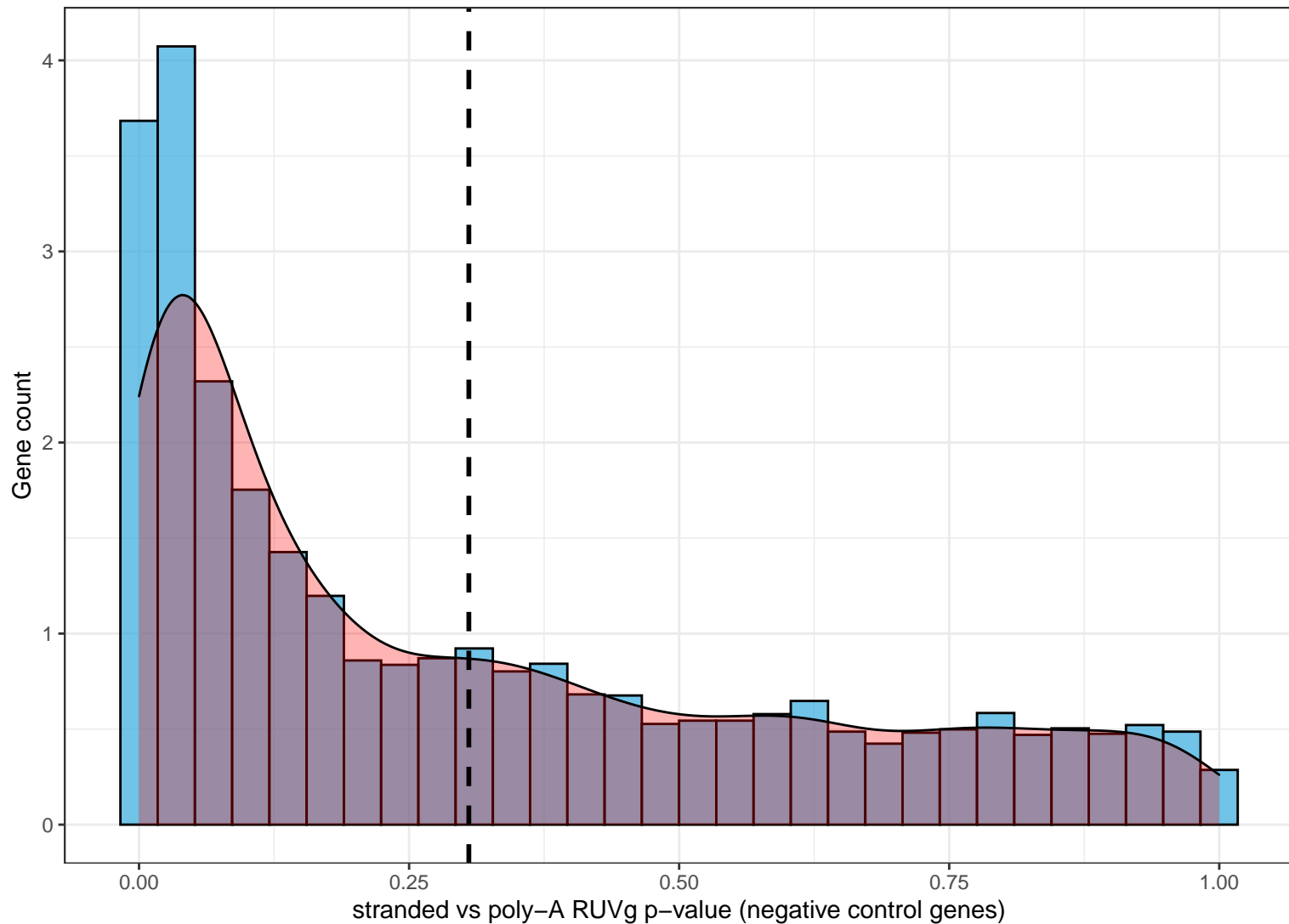
Total number of genes: 5063

1324 genes have p-value < 0.05

3739 genes have p-value ≥ 0.05

1068 genes have BH FDR < 0.05

3995 genes have BH FDR ≥ 0.05



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

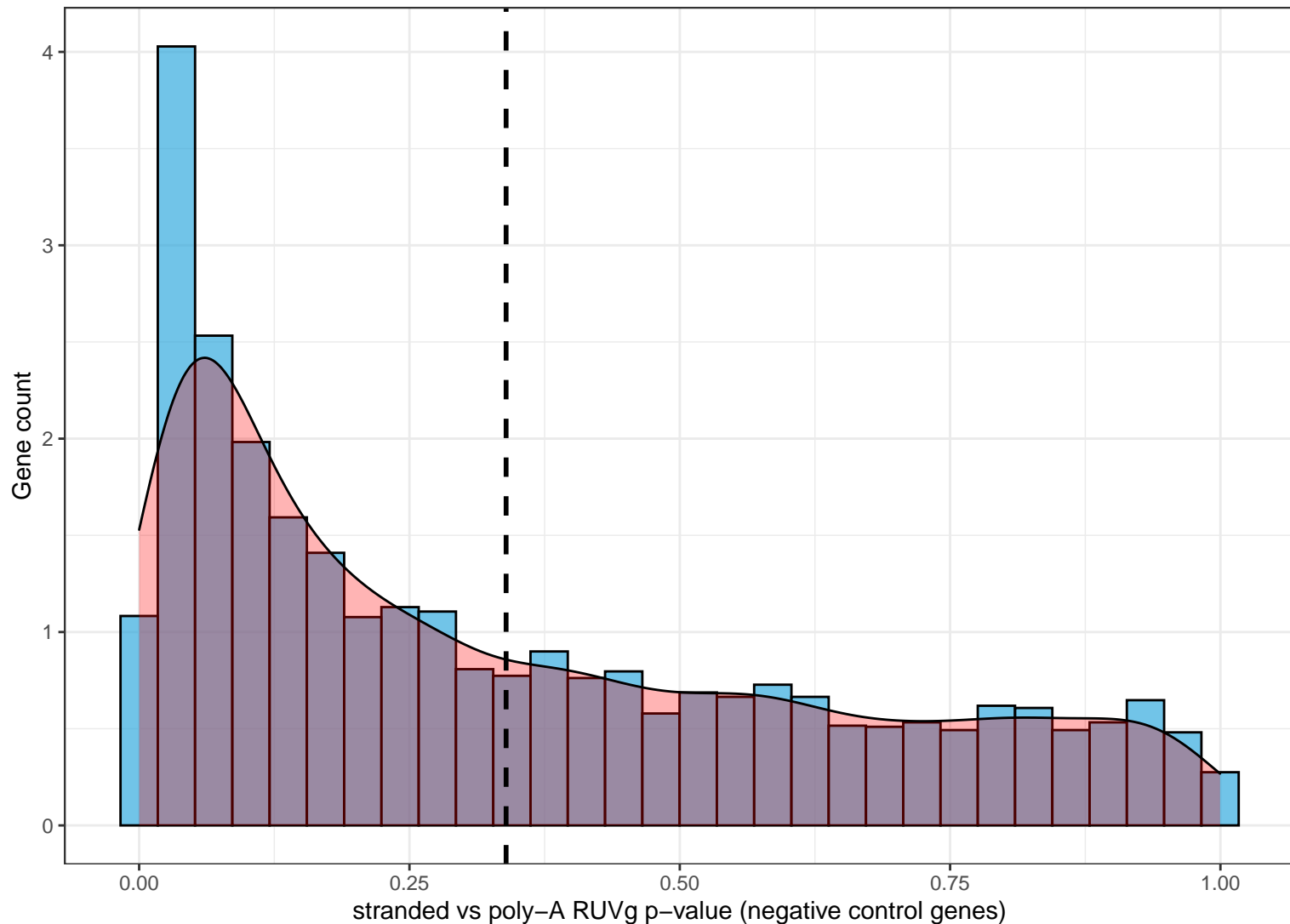
Total number of genes: 5063

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

4210 genes have $p\text{-value} \geq 0.05$

559 genes have BH FDR < 0.05

4504 genes have BH FDR ≥ 0.05



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

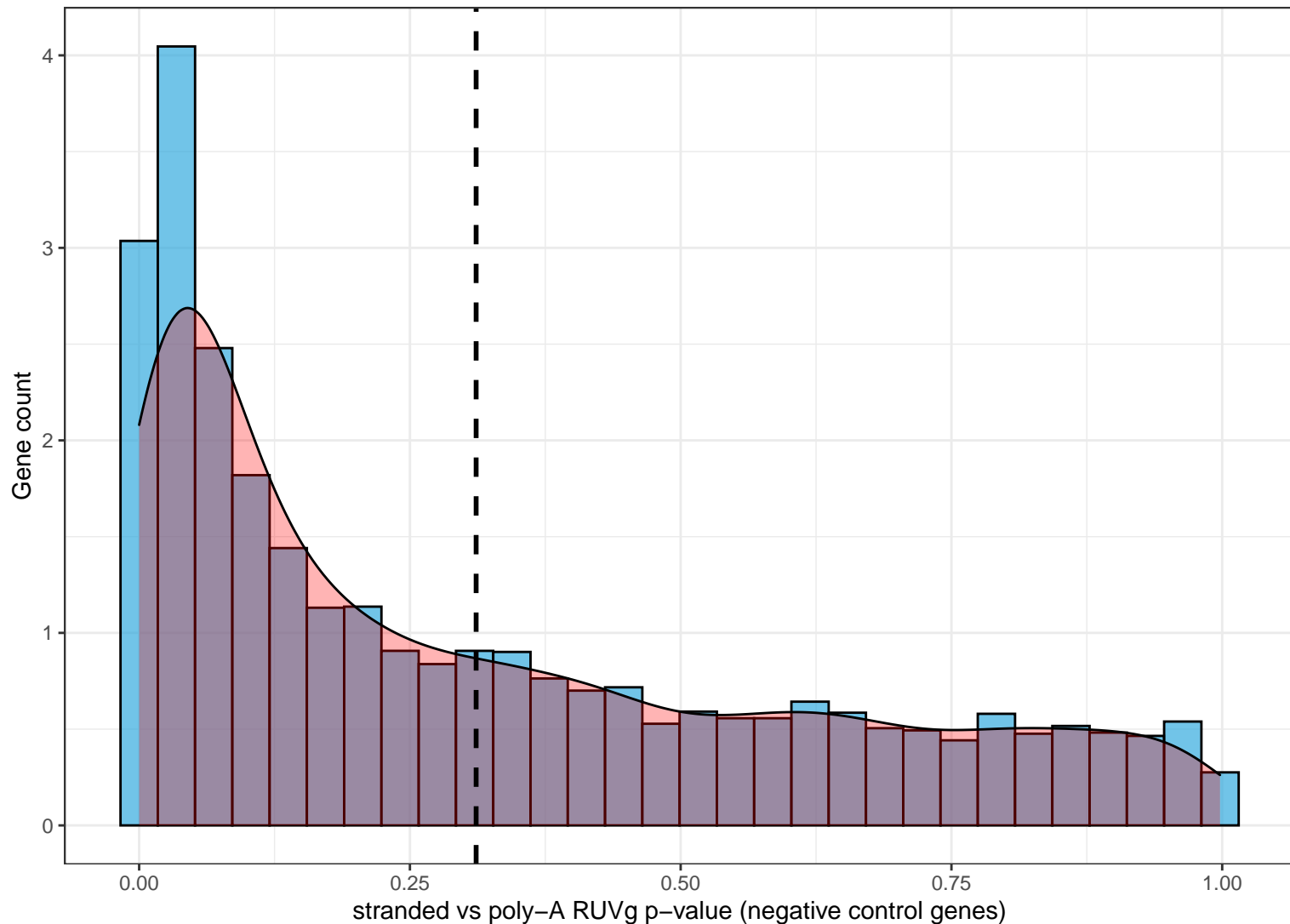
Total number of genes: 5063

1210 genes have p-value < 0.05

3853 genes have p-value ≥ 0.05

944 genes have BH FDR < 0.05

4119 genes have BH FDR ≥ 0.05



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

Total number of genes: 5063

1226 genes have p-value < 0.05

3837 genes have p-value ≥ 0.05

957 genes have BH FDR < 0.05

4106 genes have BH FDR ≥ 0.05

