

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

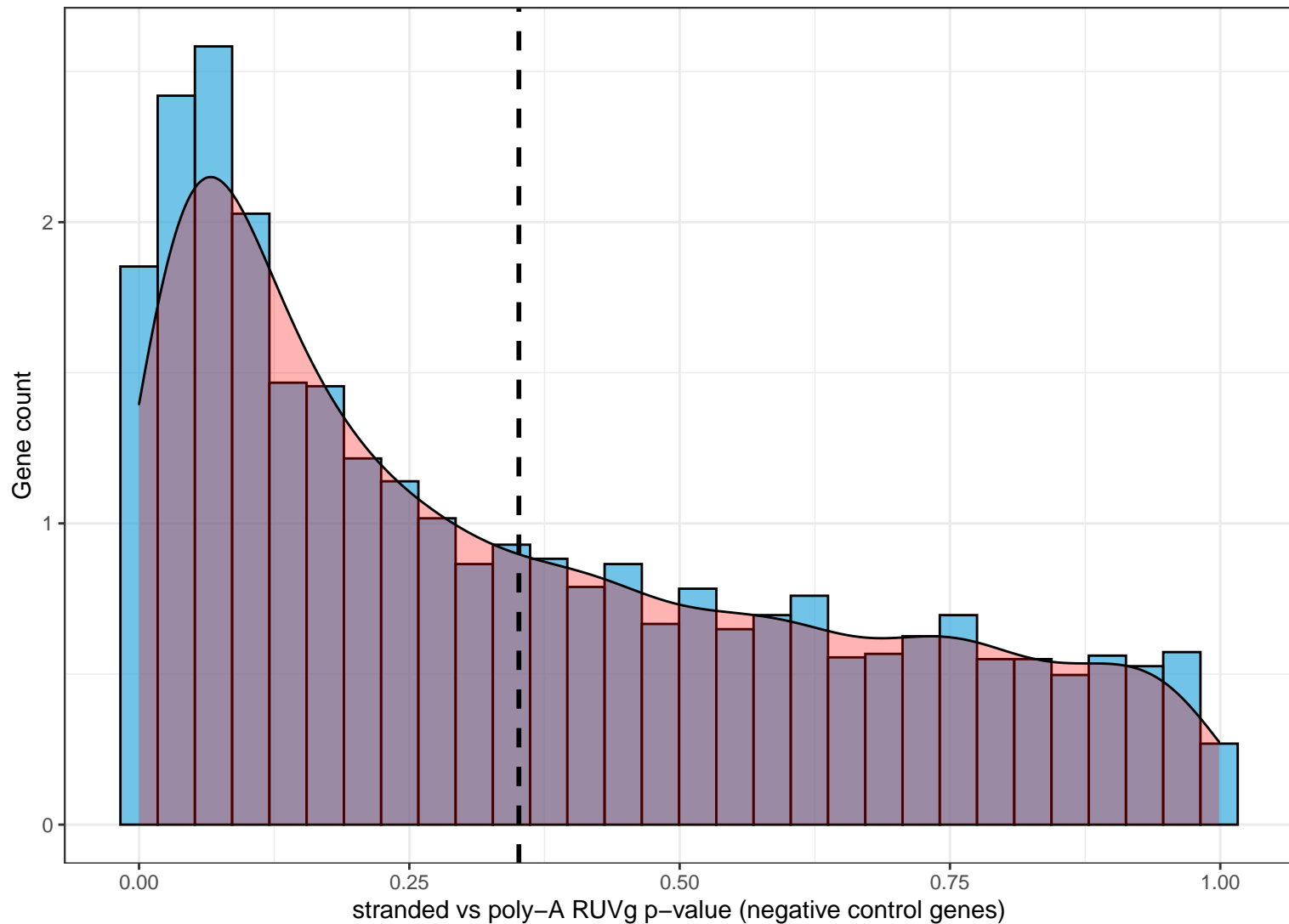
Total number of genes: 4967

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

4257 genes have $p\text{-value} \geq 0.05$

503 genes have BH FDR < 0.05

4464 genes have BH FDR ≥ 0.05



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

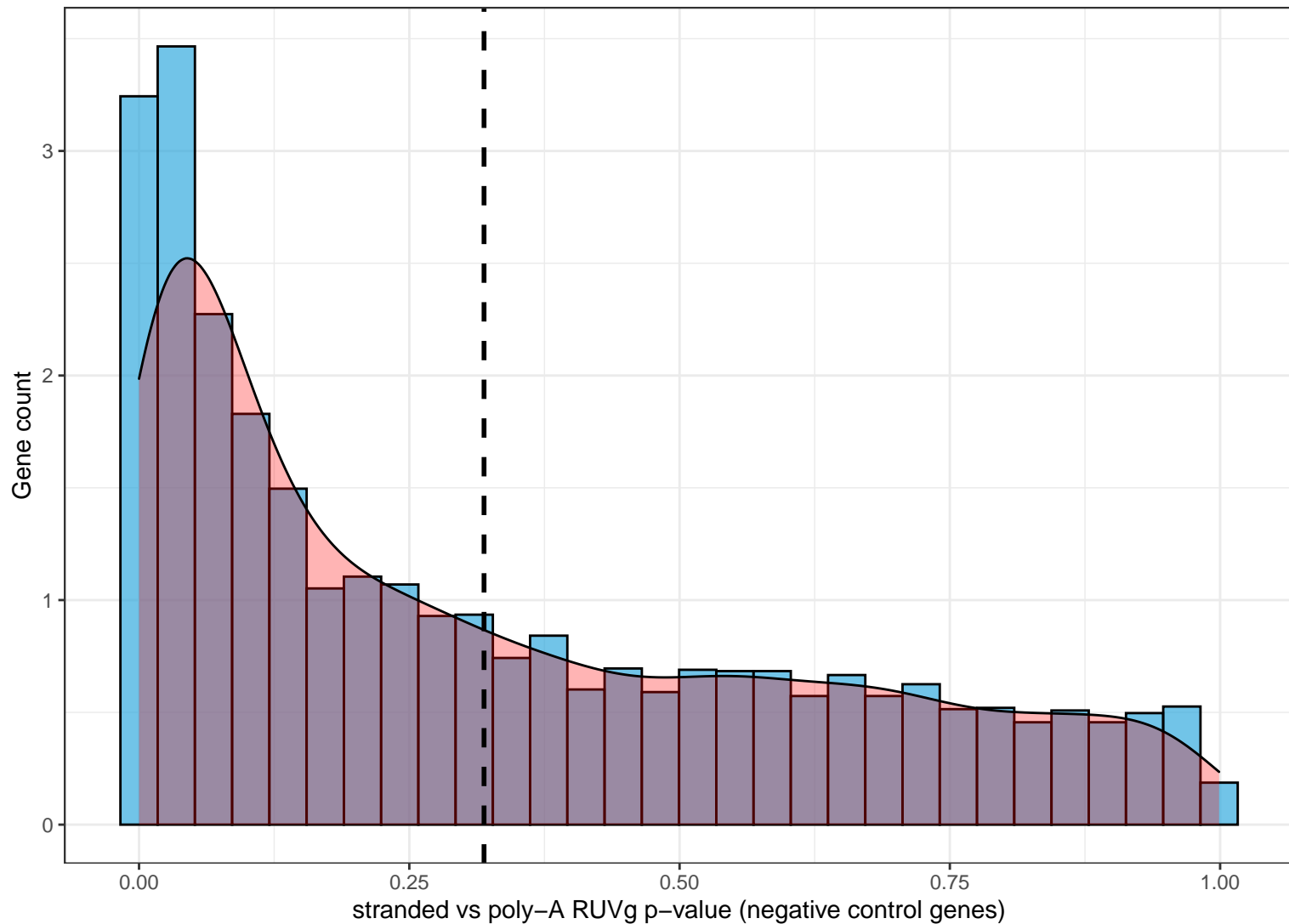
Total number of genes: 4967

1122 genes have $p\text{-value} < 0.05$

3845 genes have $p\text{-value} \geq 0.05$

888 genes have BH FDR < 0.05

4079 genes have BH FDR ≥ 0.05



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

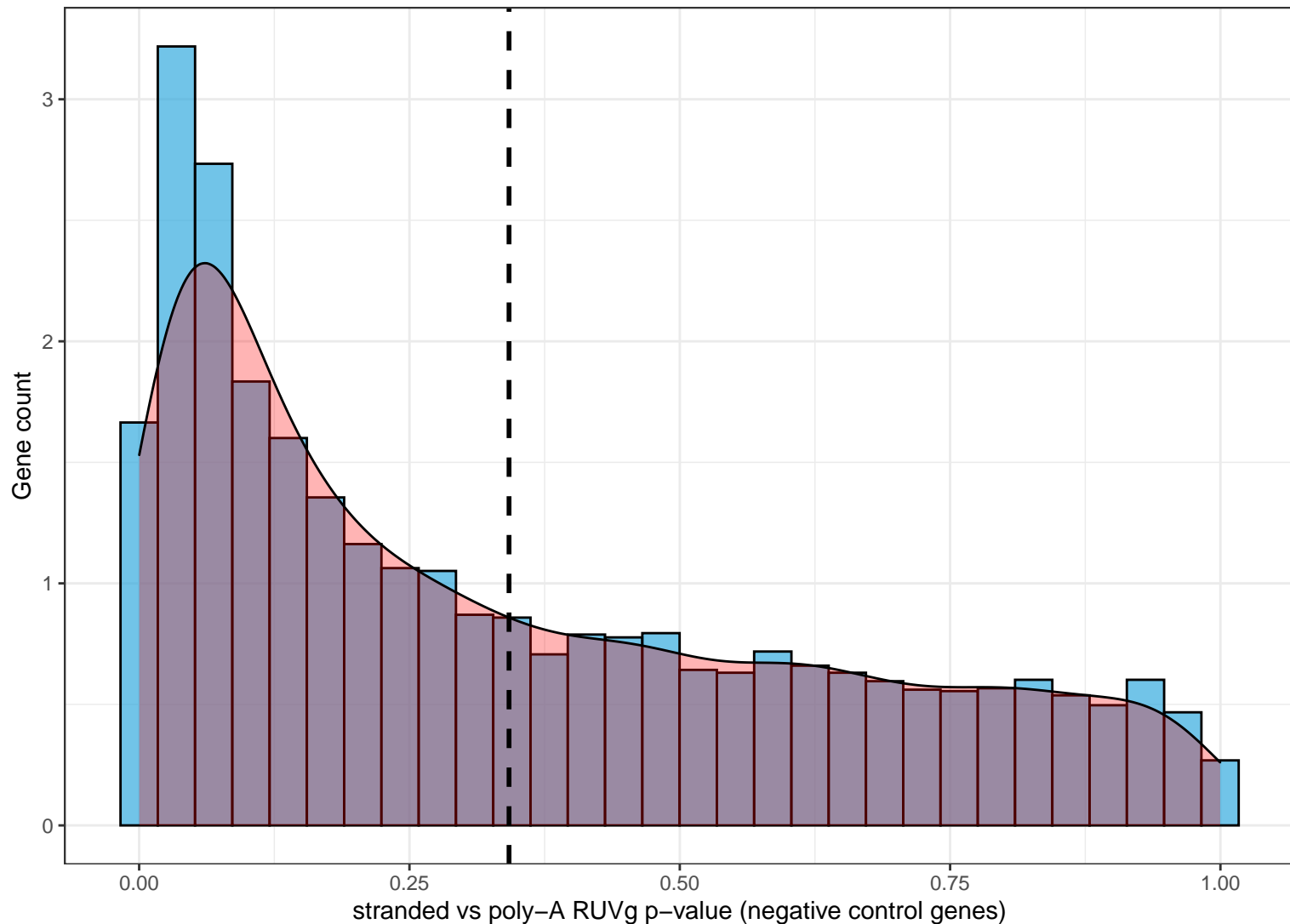
Total number of genes: 4967

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

4162 genes have $p\text{-value} \geq 0.05$

549 genes have BH FDR < 0.05

4418 genes have BH FDR ≥ 0.05



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

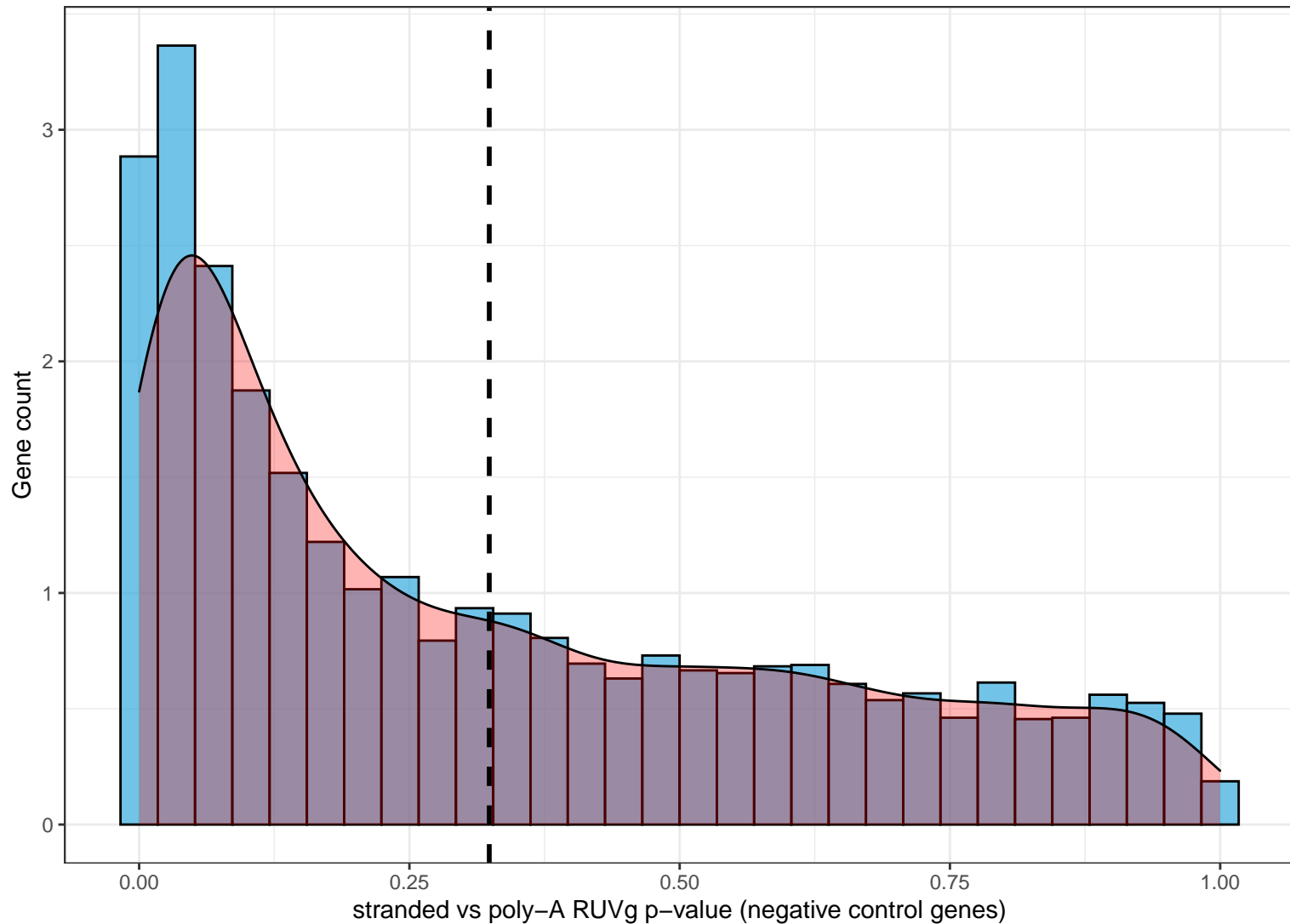
Total number of genes: 4967

1041 genes have p-value < 0.05

3926 genes have p-value ≥ 0.05

804 genes have BH FDR < 0.05

4163 genes have BH FDR ≥ 0.05



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

Total number of genes: 4967

1038 genes have $p\text{-value} < 0.05$

3929 genes have $p\text{-value} \geq 0.05$

803 genes have BH FDR < 0.05

4164 genes have BH FDR ≥ 0.05

