

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

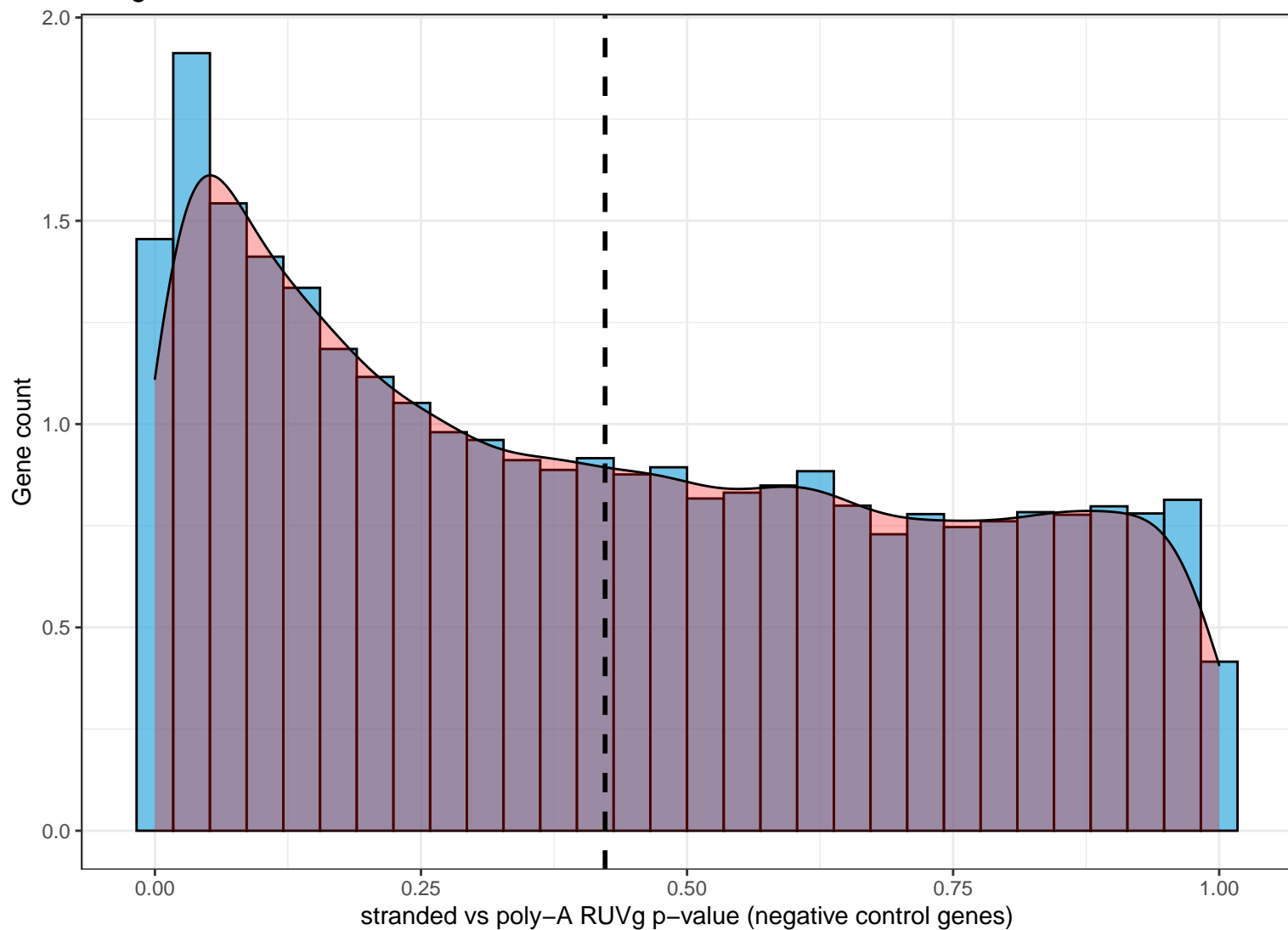
Total number of genes: 18138

2057 genes have p-value < 0.05

16081 genes have p-value ≥ 0.05

NA genes have BH FDR < 0.05

NA genes have BH FDR ≥ 0.05



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

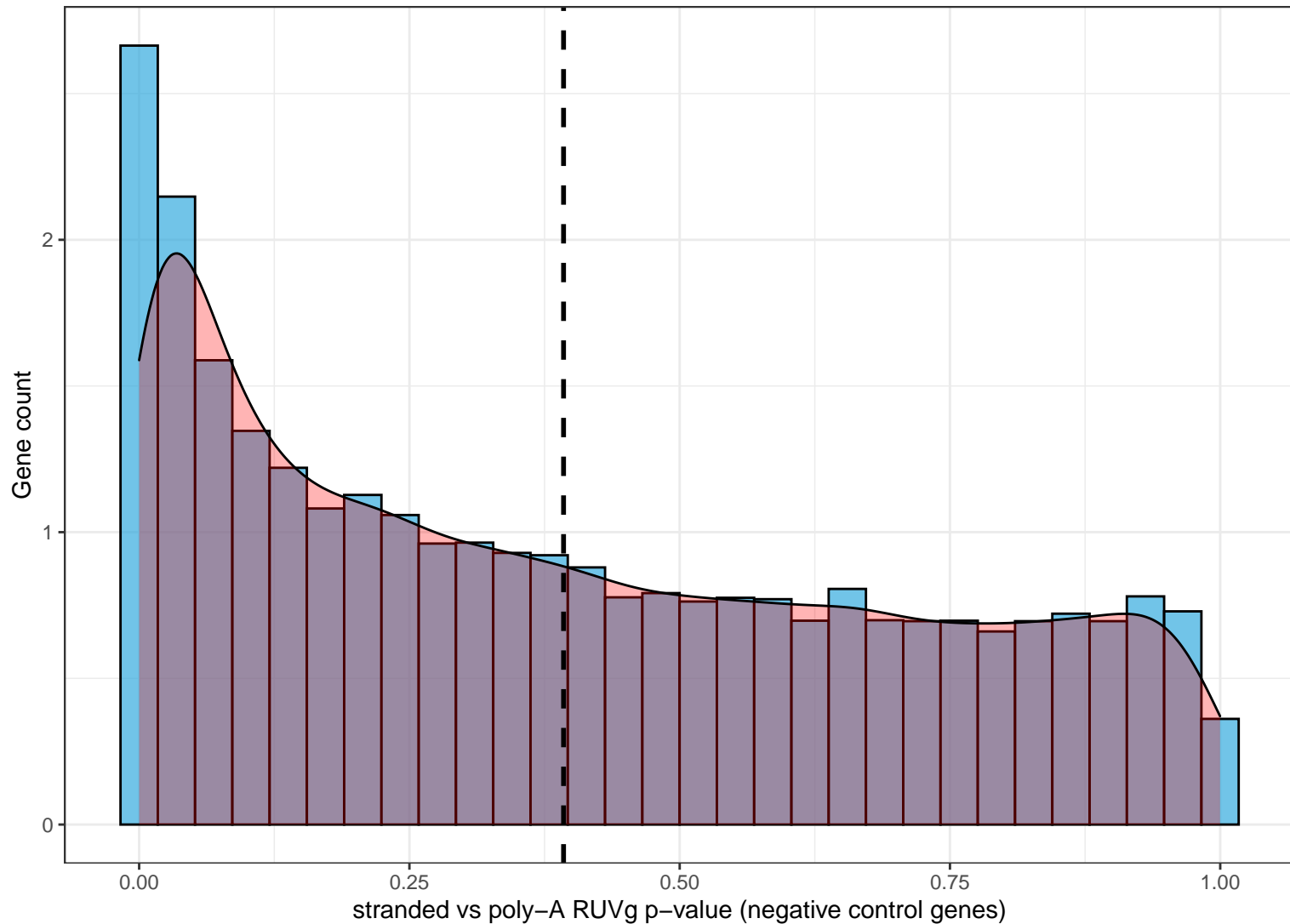
Total number of genes: 18138

2930 genes have p-value < 0.05

15208 genes have p-value ≥ 0.05

NA genes have BH FDR < 0.05

NA genes have BH FDR ≥ 0.05



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

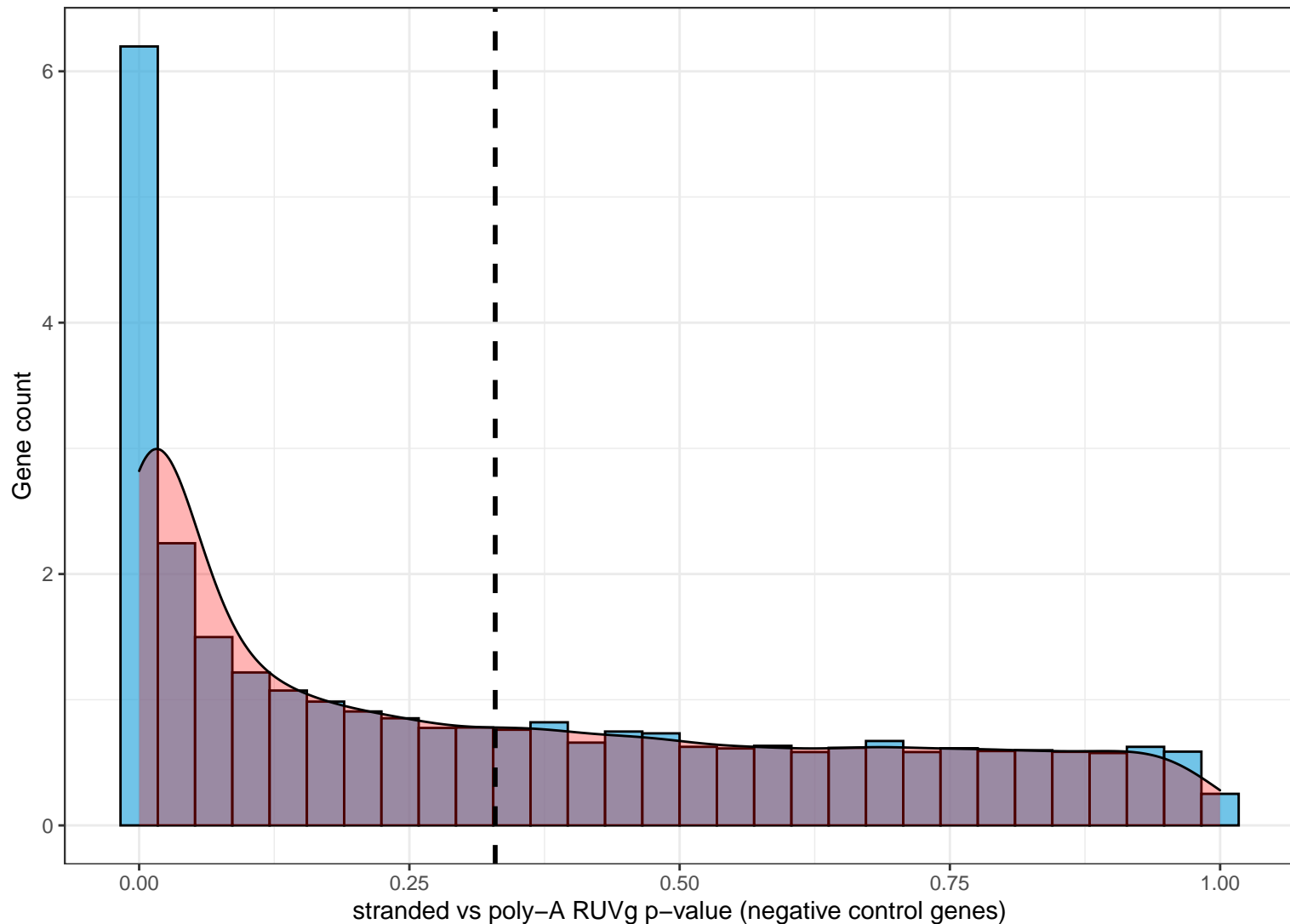
Total number of genes: 18138

5220 genes have p-value < 0.05

12918 genes have p-value ≥ 0.05

3289 genes have BH FDR < 0.05

14849 genes have BH FDR ≥ 0.05



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

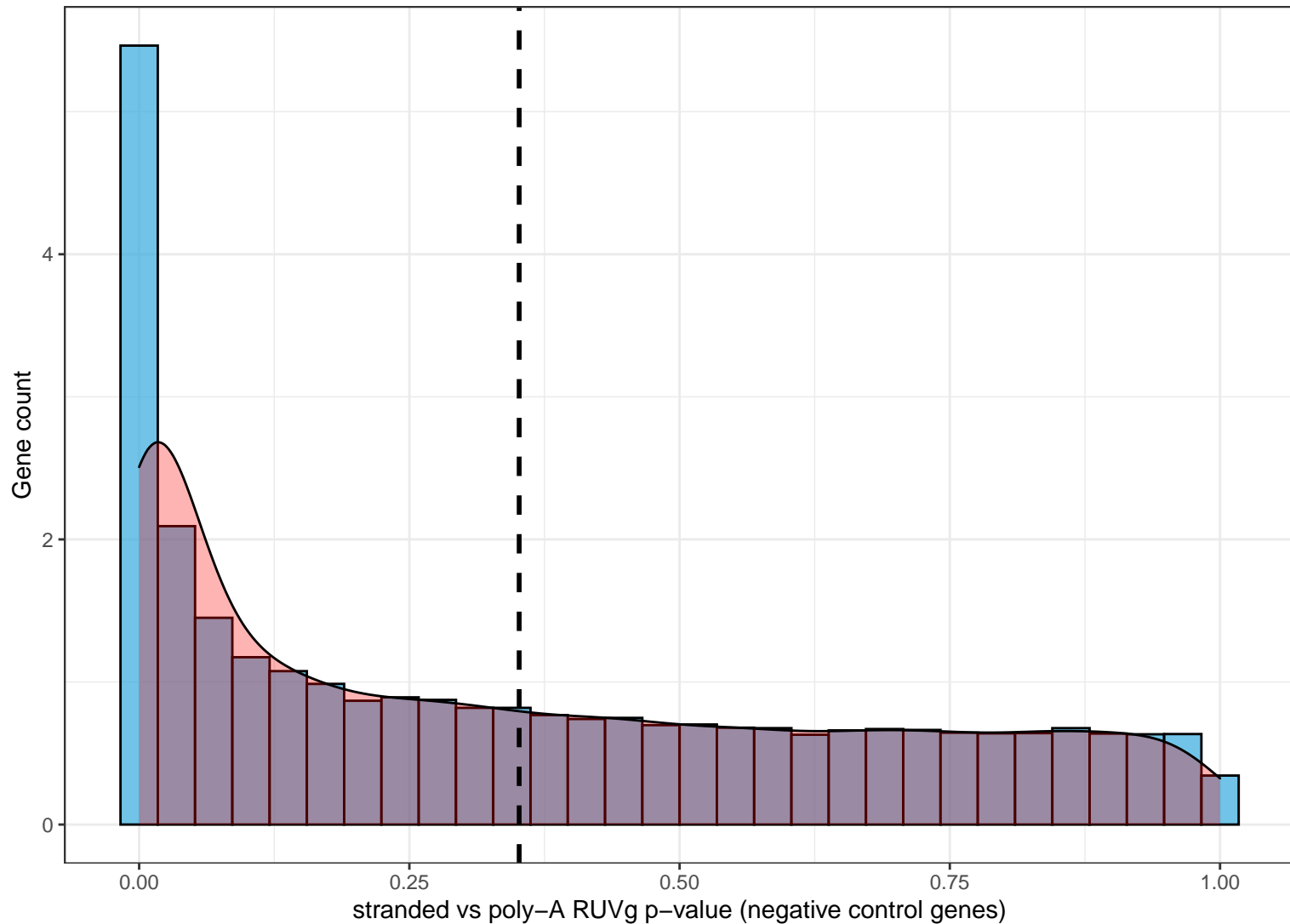
Total number of genes: 18138

4674 genes have p-value < 0.05

13464 genes have p-value ≥ 0.05

2754 genes have BH FDR < 0.05

15384 genes have BH FDR ≥ 0.05



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

Total number of genes: 18138

4925 genes have p-value < 0.05

13213 genes have p-value ≥ 0.05

2975 genes have BH FDR < 0.05

15163 genes have BH FDR ≥ 0.05

