

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

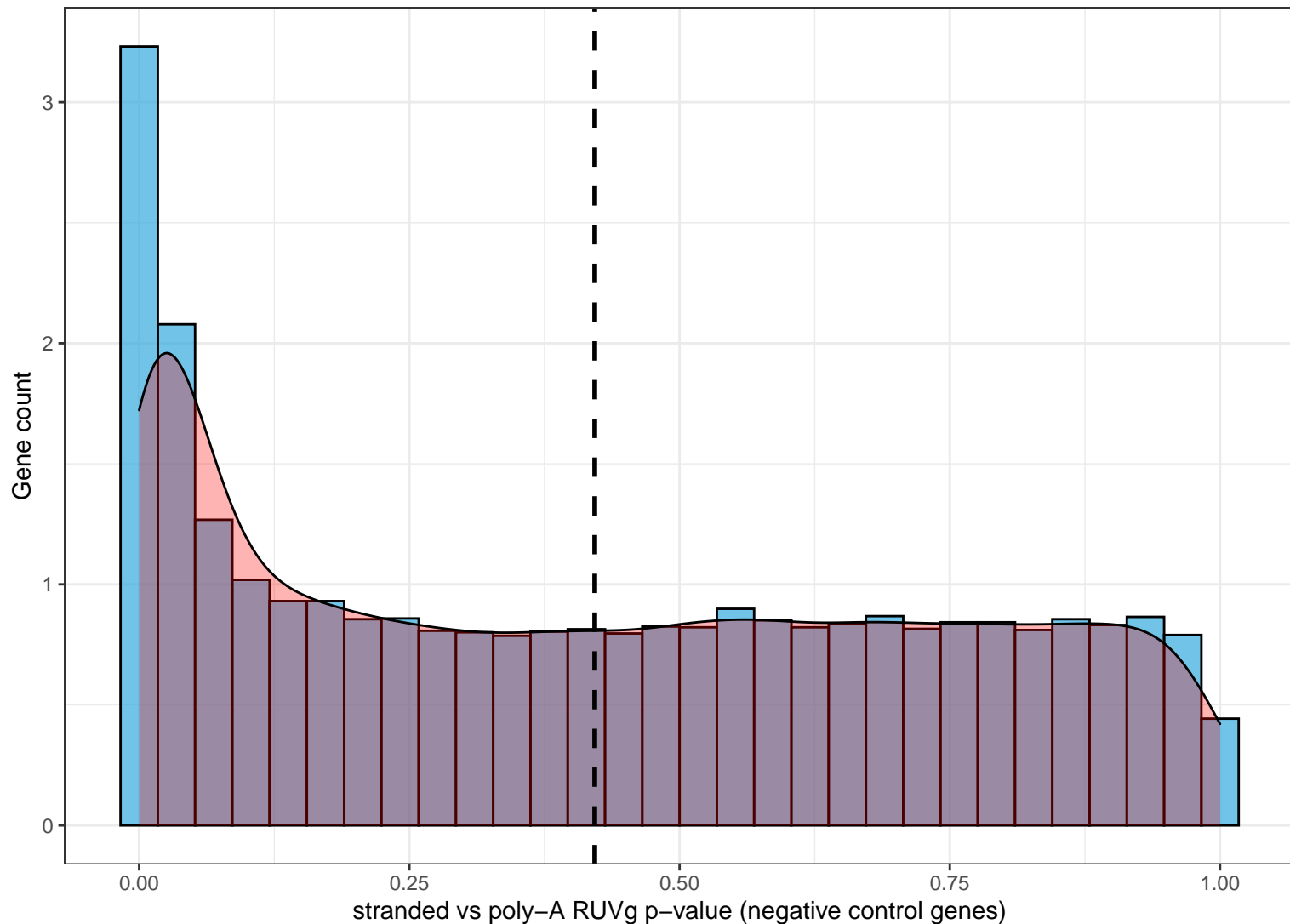
Total number of genes: 18138

3271 genes have p-value < 0.05

14867 genes have p-value ≥ 0.05

863 genes have BH FDR < 0.05

17275 genes have BH FDR ≥ 0.05



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

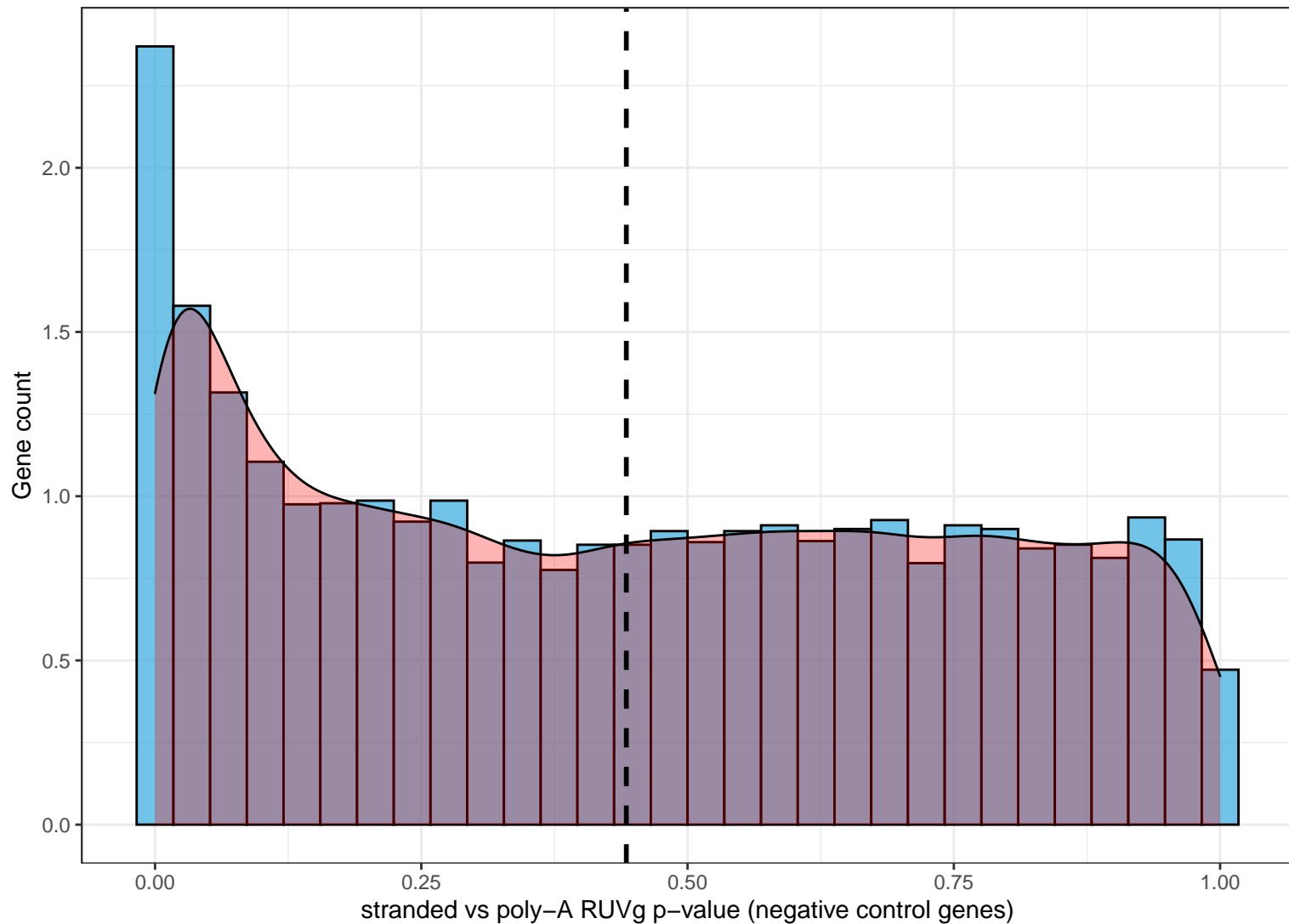
Total number of genes: 18138

2427 genes have p-value < 0.05

15711 genes have p-value ≥ 0.05

635 genes have BH FDR < 0.05

17503 genes have BH FDR ≥ 0.05



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

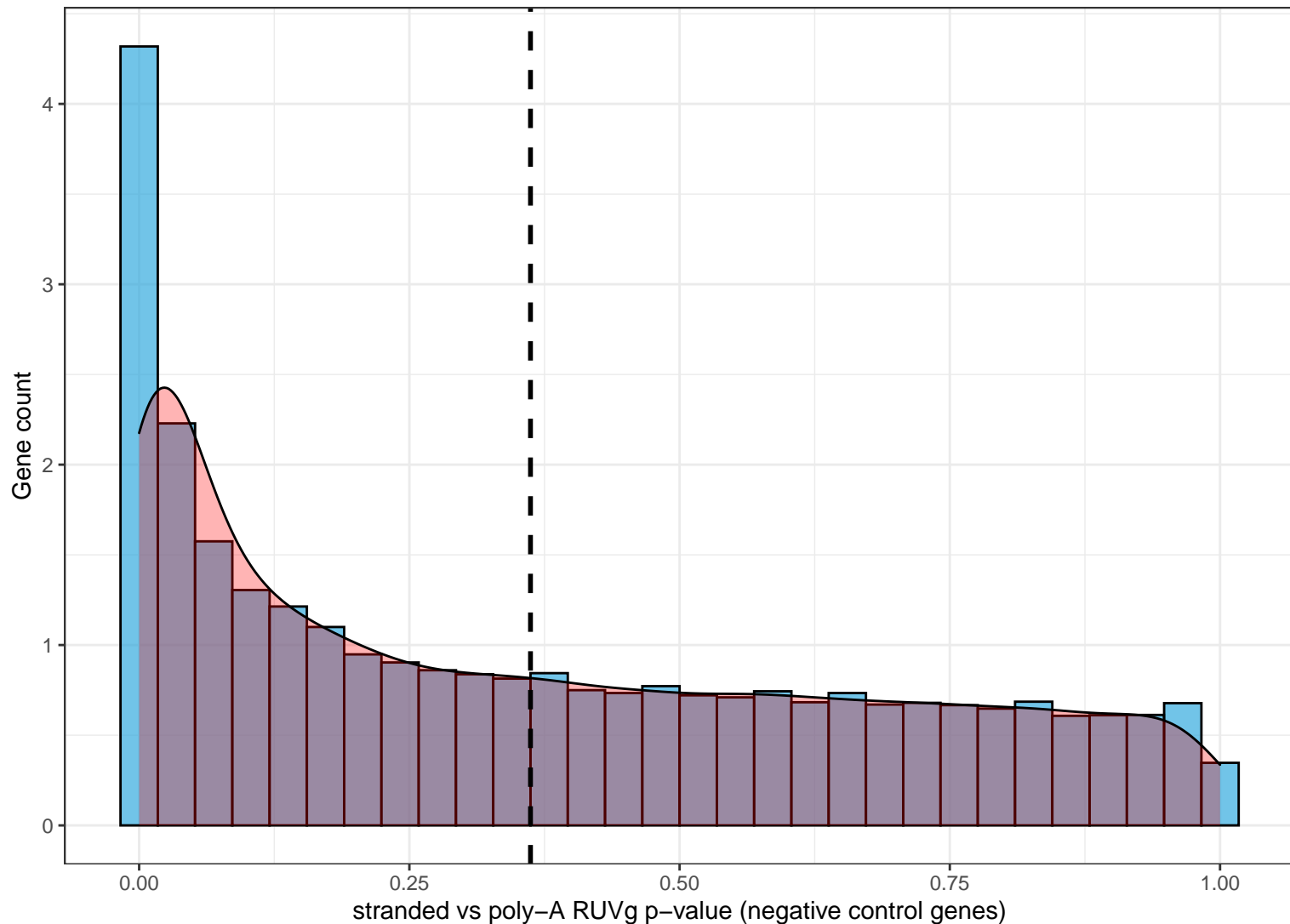
Total number of genes: 18138

4052 genes have p-value < 0.05

14086 genes have p-value ≥ 0.05

1726 genes have BH FDR < 0.05

16412 genes have BH FDR ≥ 0.05



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

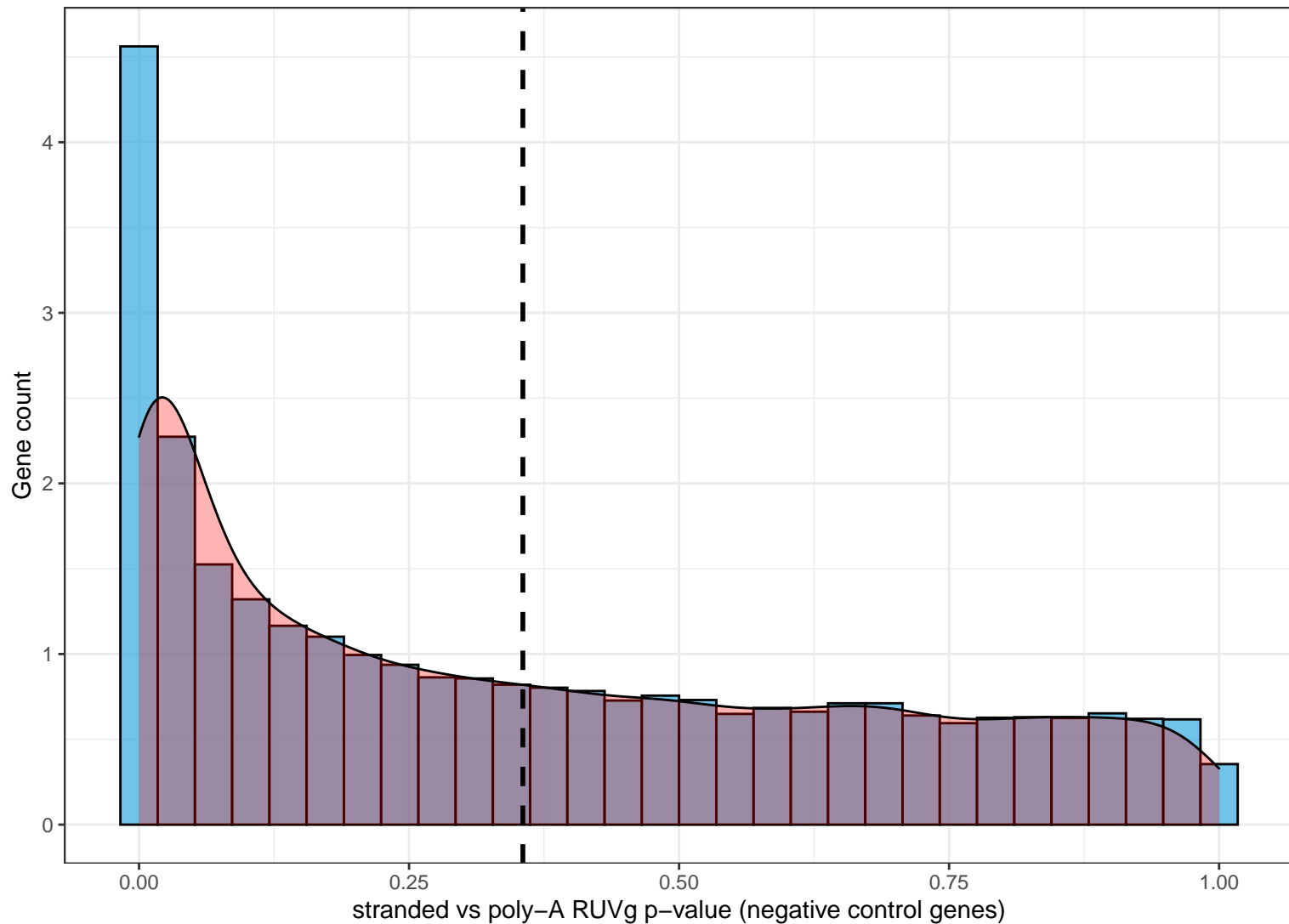
Total number of genes: 18138

4219 genes have p-value < 0.05

13919 genes have p-value ≥ 0.05

1996 genes have BH FDR < 0.05

16142 genes have BH FDR ≥ 0.05



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

Total number of genes: 18138

2736 genes have p-value < 0.05

15402 genes have p-value >= 0.05

992 genes have BH FDR < 0.05

17146 genes have BH FDR >= 0.05

