Histogram of stranded vs poly−A paired analysis (k = 1)
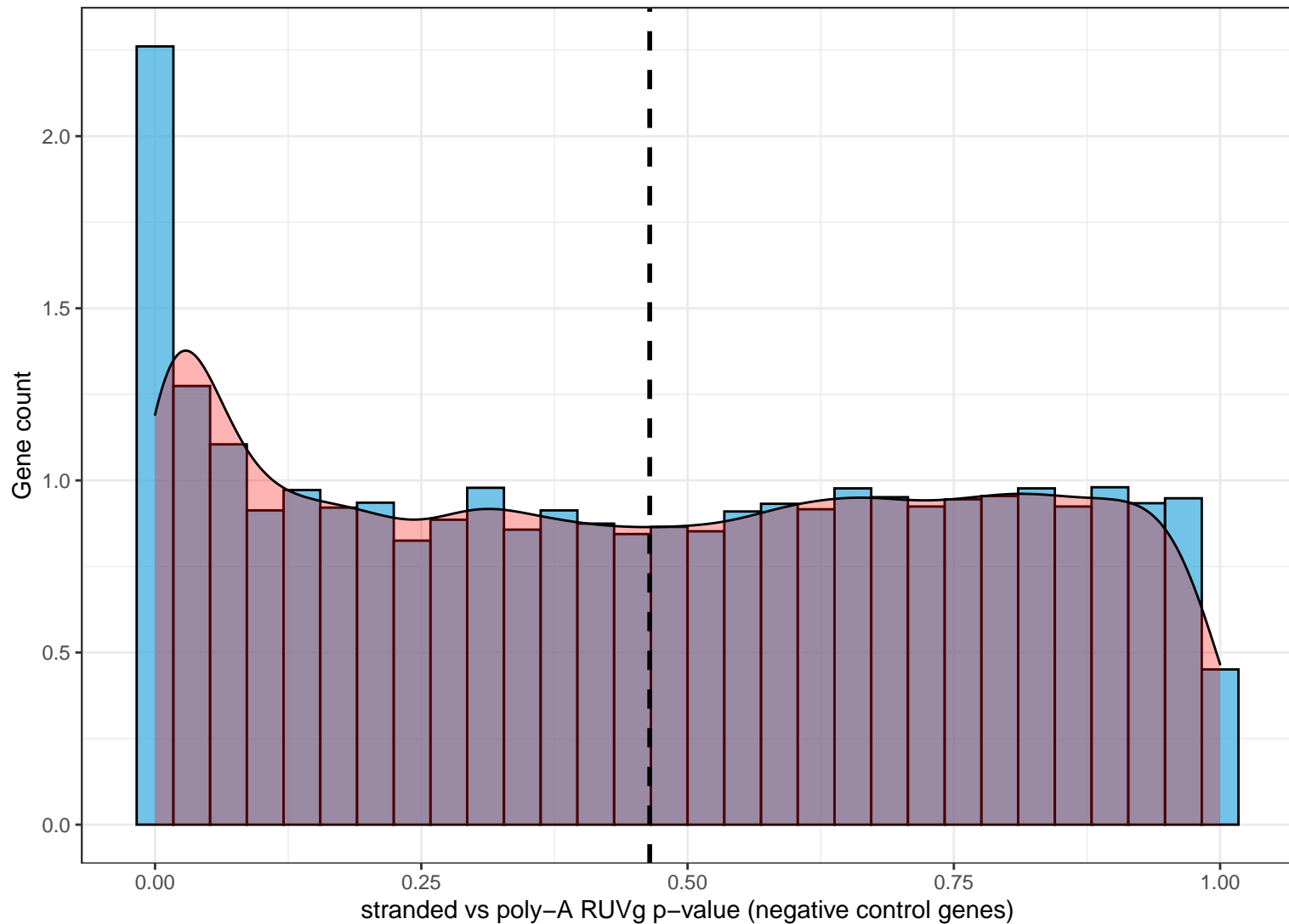Total number of genes: 18138
2174 genes have p−value < 0.05
15964 genes have p−value >= 0.05
718 genes have BH FDR < 0.05
17420 genes have BH FDR >= 0.05

Histogram of stranded vs poly−A paired analysis (k = 2)
Total number of genes: 18138
3019 genes have p−value < 0.05
15119 genes have p−value >= 0.05
1252 genes have BH FDR < 0.05
16886 genes have BH FDR >= 0.05
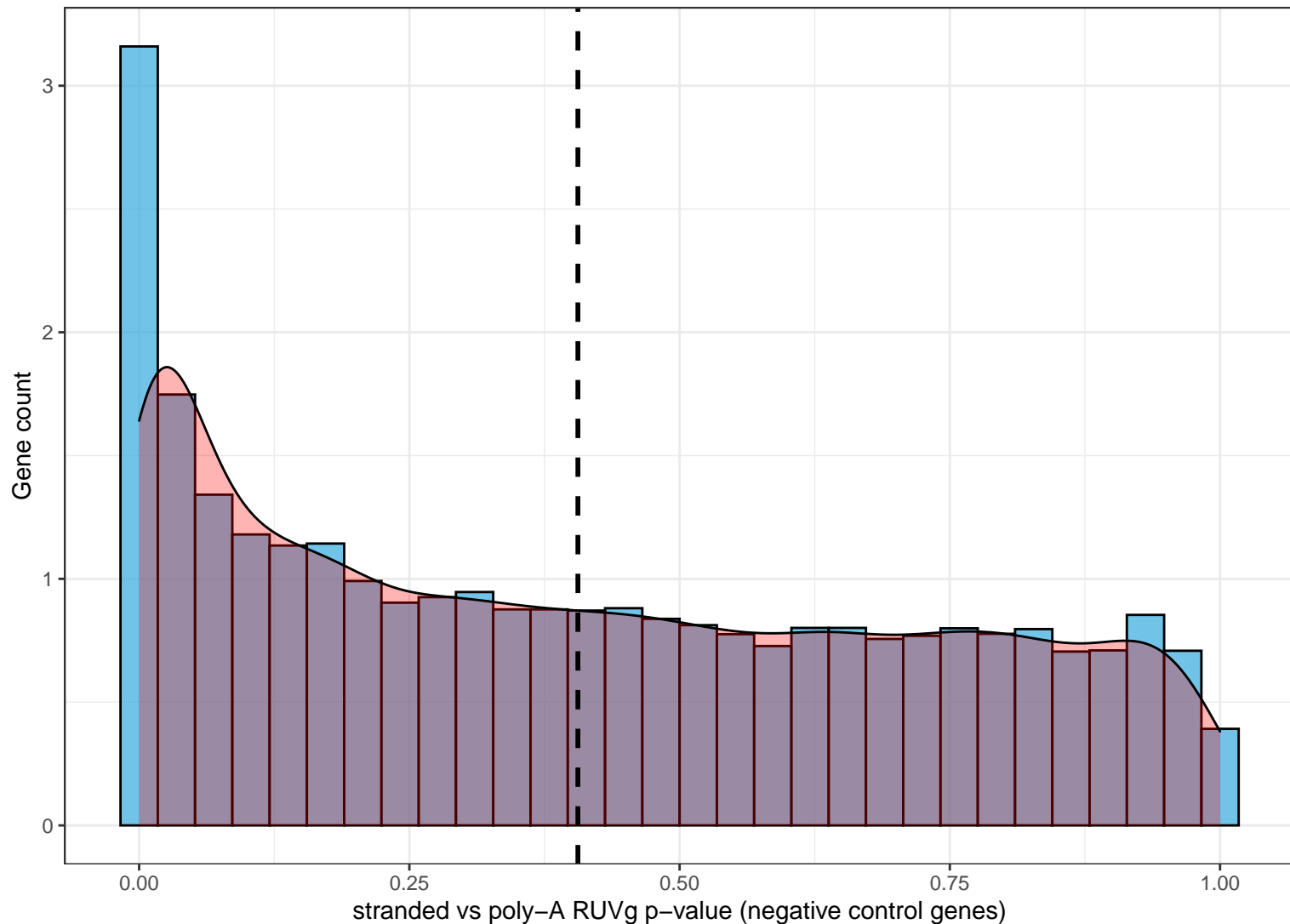
Histogram of stranded vs poly−A paired analysis (k = 3)
Total number of genes: 18138
2108 genes have p−value < 0.05
16030 genes have p−value >= 0.05
723 genes have BH FDR < 0.05
17415 genes have BH FDR >= 0.05
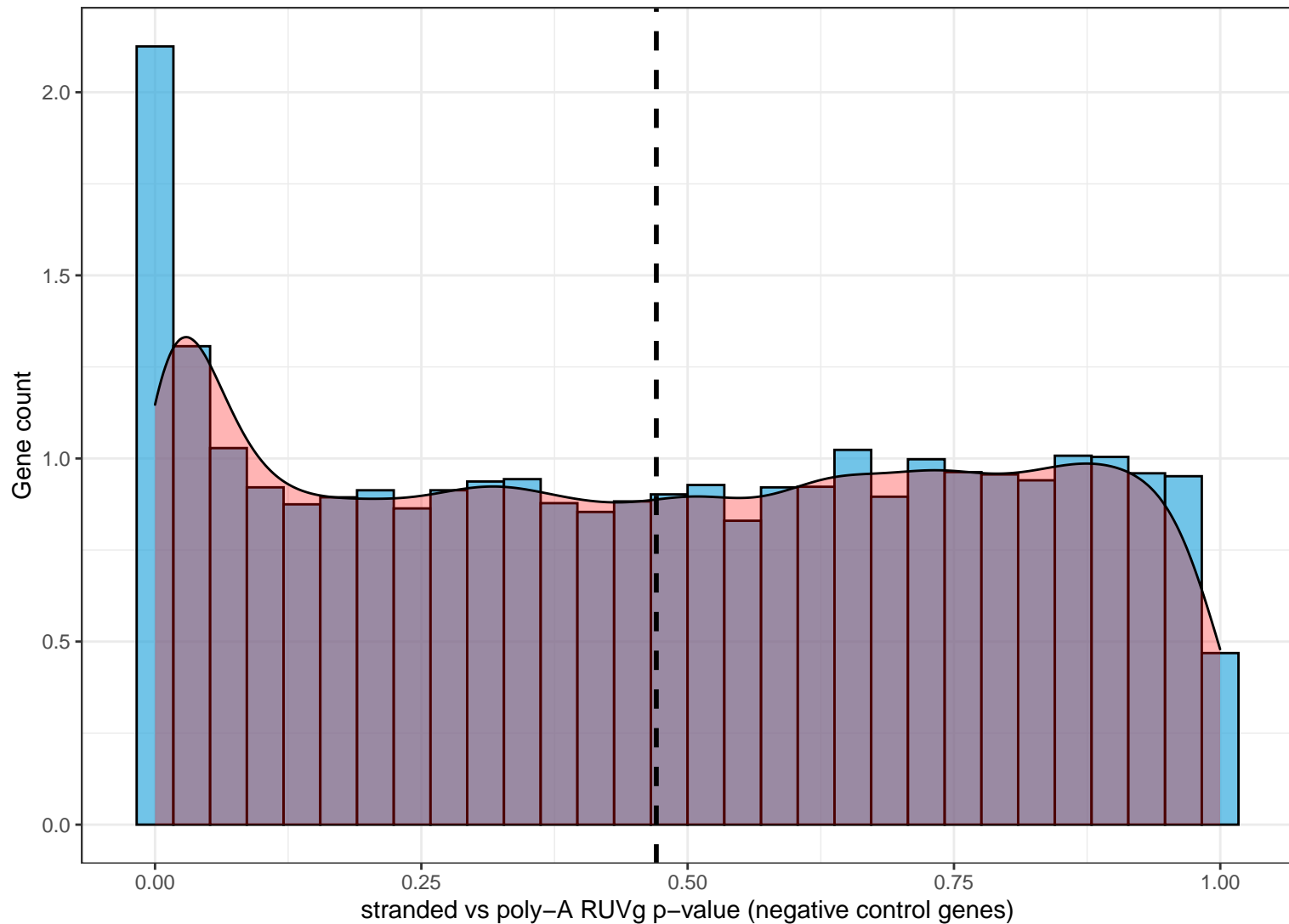
Histogram of stranded vs poly−A paired analysis (k = 4)
Total number of genes: 18138
2430 genes have p−value < 0.05
15708 genes have p−value >= 0.05
900 genes have BH FDR < 0.05
17238 genes have BH FDR >= 0.05
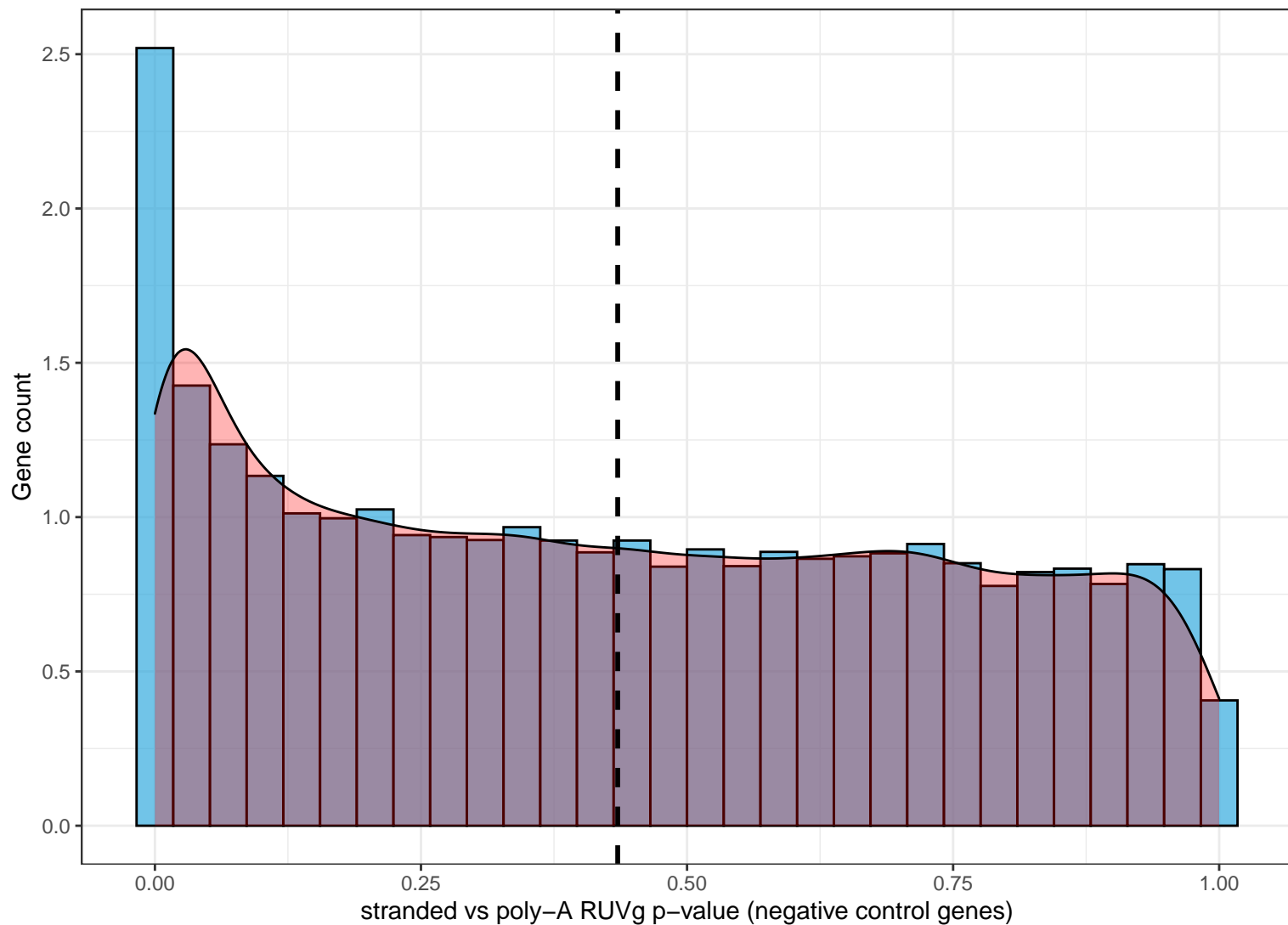
Histogram of stranded vs poly−A paired analysis (k = 5)
Total number of genes: 18138
2138 genes have p−value < 0.05
16000 genes have p−value >= 0.05
836 genes have BH FDR < 0.05
17302 genes have BH FDR >= 0.05