Histogram of stranded vs poly−A paired analysis (k = 1)
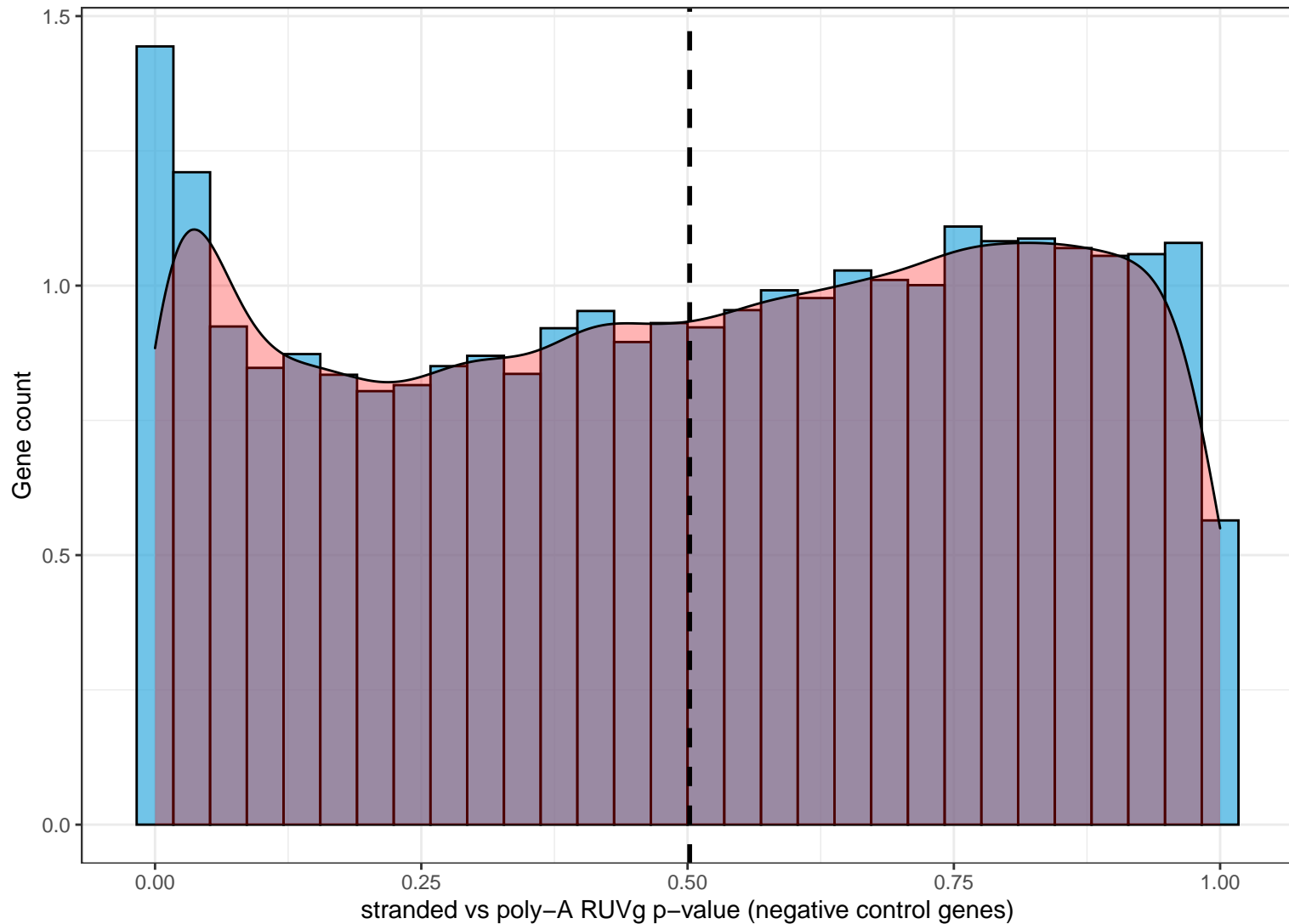Total number of genes: 18138
1620 genes have p−value < 0.05
16518 genes have p−value >= 0.05
245 genes have BH FDR < 0.05
17893 genes have BH FDR >= 0.05

Histogram of stranded vs poly−A paired analysis (k = 2)
Total number of genes: 18138
3188 genes have p−value < 0.05
14950 genes have p−value >= 0.05
1551 genes have BH FDR < 0.05
16587 genes have BH FDR >= 0.05
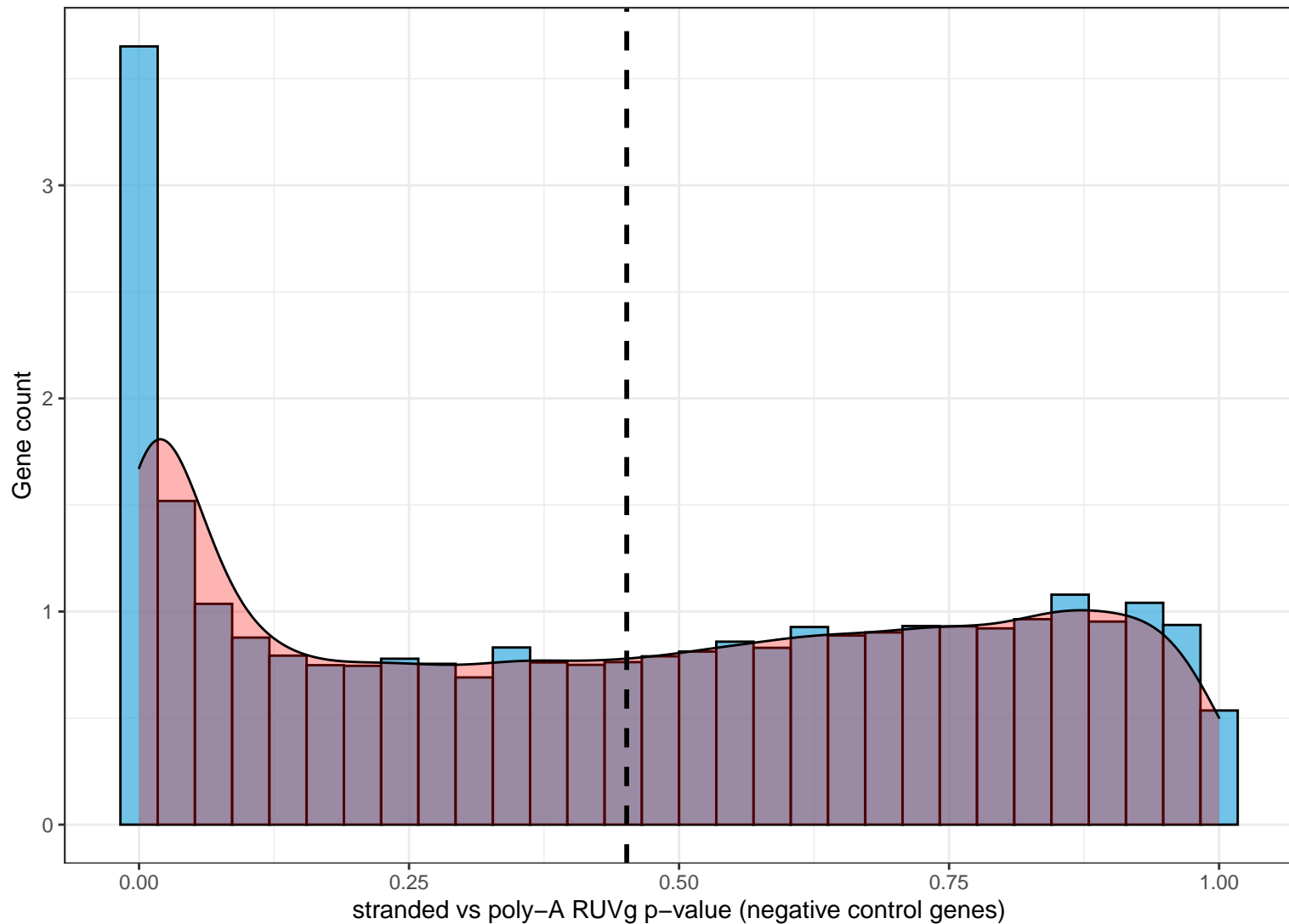
Histogram of stranded vs poly−A paired analysis (k = 3)
Total number of genes: 18138
2687 genes have p−value < 0.05
15451 genes have p−value >= 0.05
1096 genes have BH FDR < 0.05
17042 genes have BH FDR >= 0.05
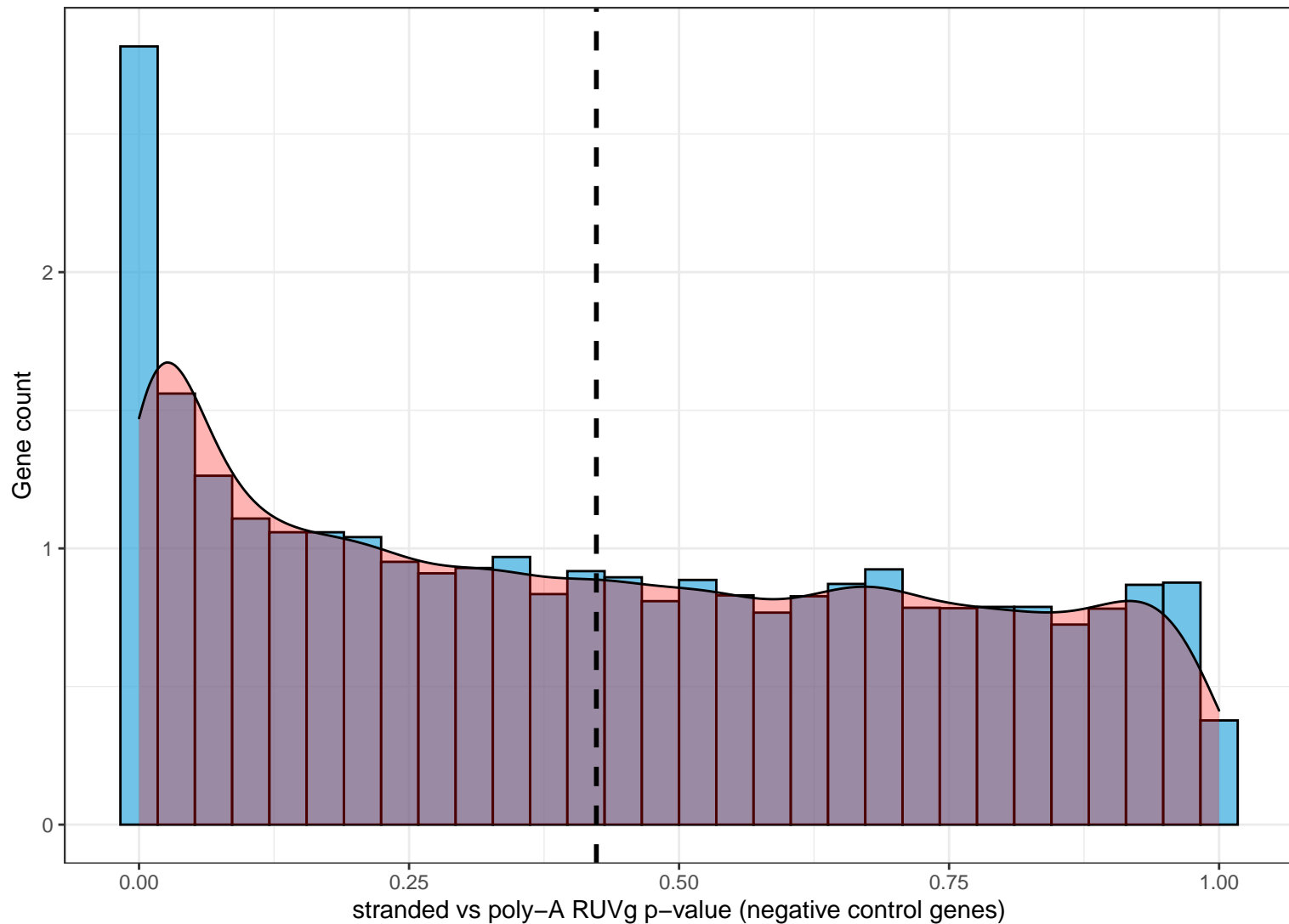
Histogram of stranded vs poly−A paired analysis (k = 4)
Total number of genes: 18138
3215 genes have p−value < 0.05
14923 genes have p−value >= 0.05
1365 genes have BH FDR < 0.05
16773 genes have BH FDR >= 0.05
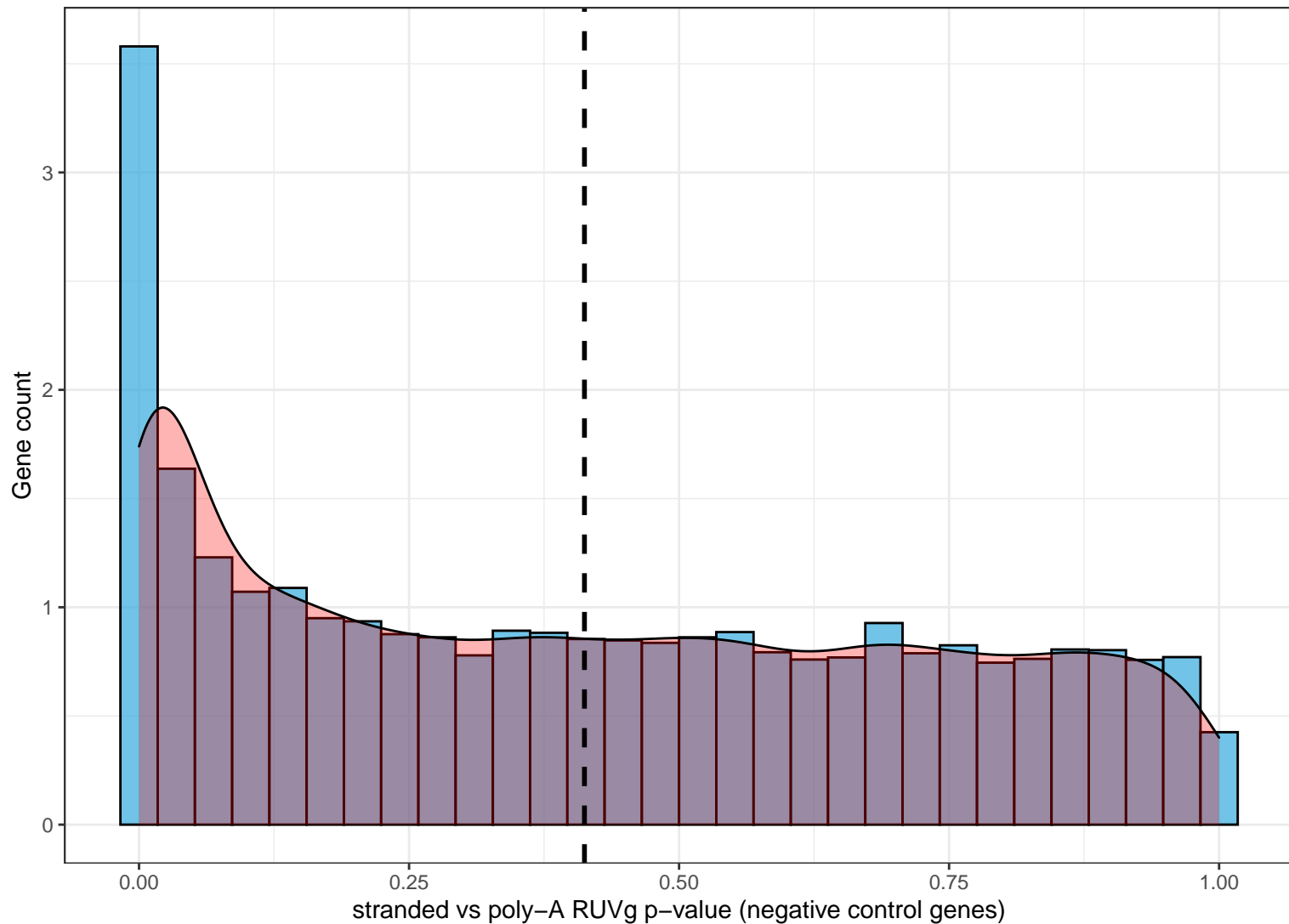
Histogram of stranded vs poly−A paired analysis (k = 5)
Total number of genes: 18138
2292 genes have p−value < 0.05
15846 genes have p−value >= 0.05
827 genes have BH FDR < 0.05
17311 genes have BH FDR >= 0.05