Histogram of stranded vs poly−A paired analysis (k = 1)
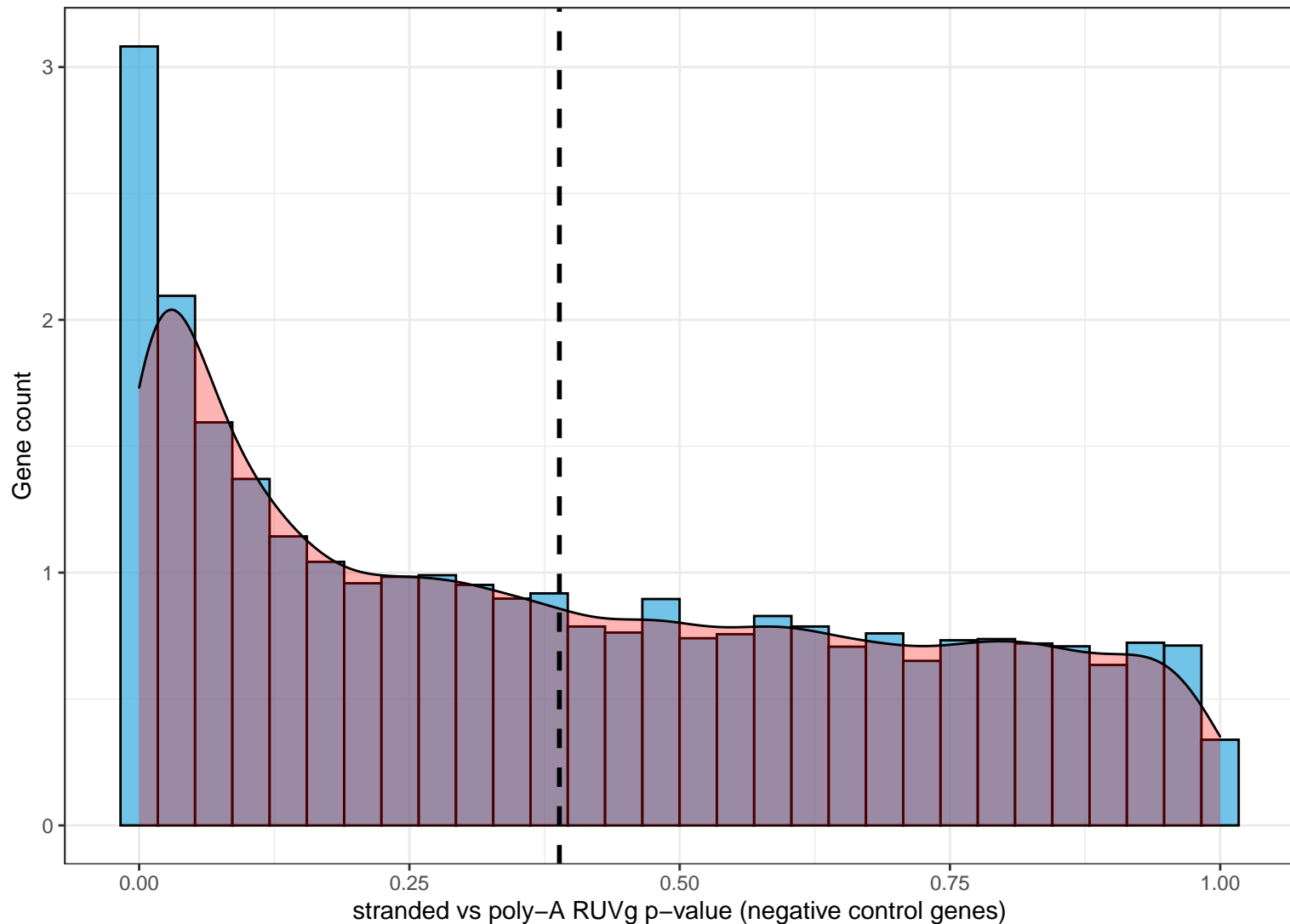Total number of genes: 18138
3187 genes have p−value < 0.05
14951 genes have p−value >= 0.05
863 genes have BH FDR < 0.05
17275 genes have BH FDR >= 0.05

Histogram of stranded vs poly−A paired analysis (k = 2)
Total number of genes: 18138
4150 genes have p−value < 0.05
13988 genes have p−value >= 0.05
2122 genes have BH FDR < 0.05
16016 genes have BH FDR >= 0.05
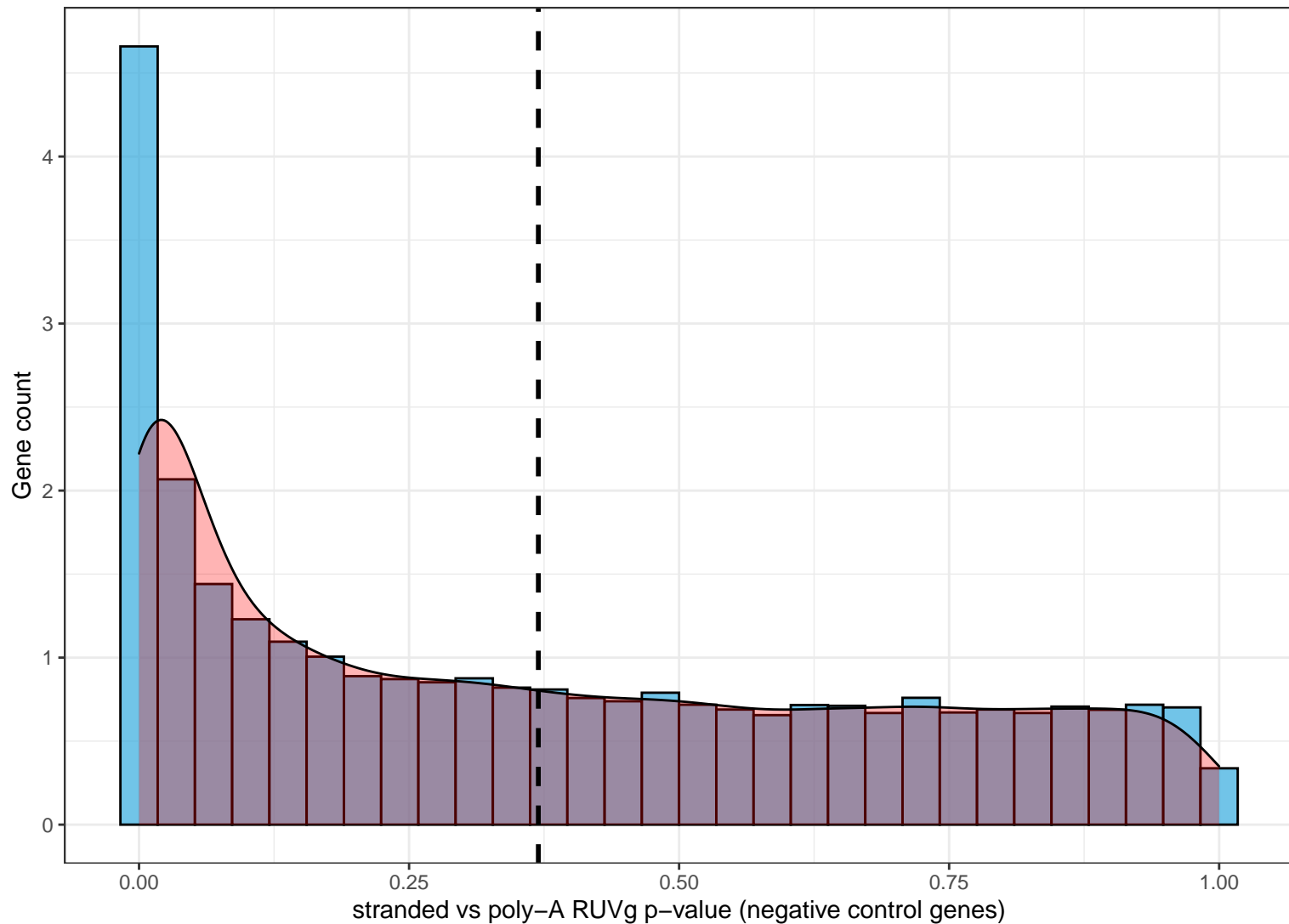
Histogram of stranded vs poly−A paired analysis (k = 3)
Total number of genes: 18138
5232 genes have p−value < 0.05
12906 genes have p−value >= 0.05
3442 genes have BH FDR < 0.05
14696 genes have BH FDR >= 0.05
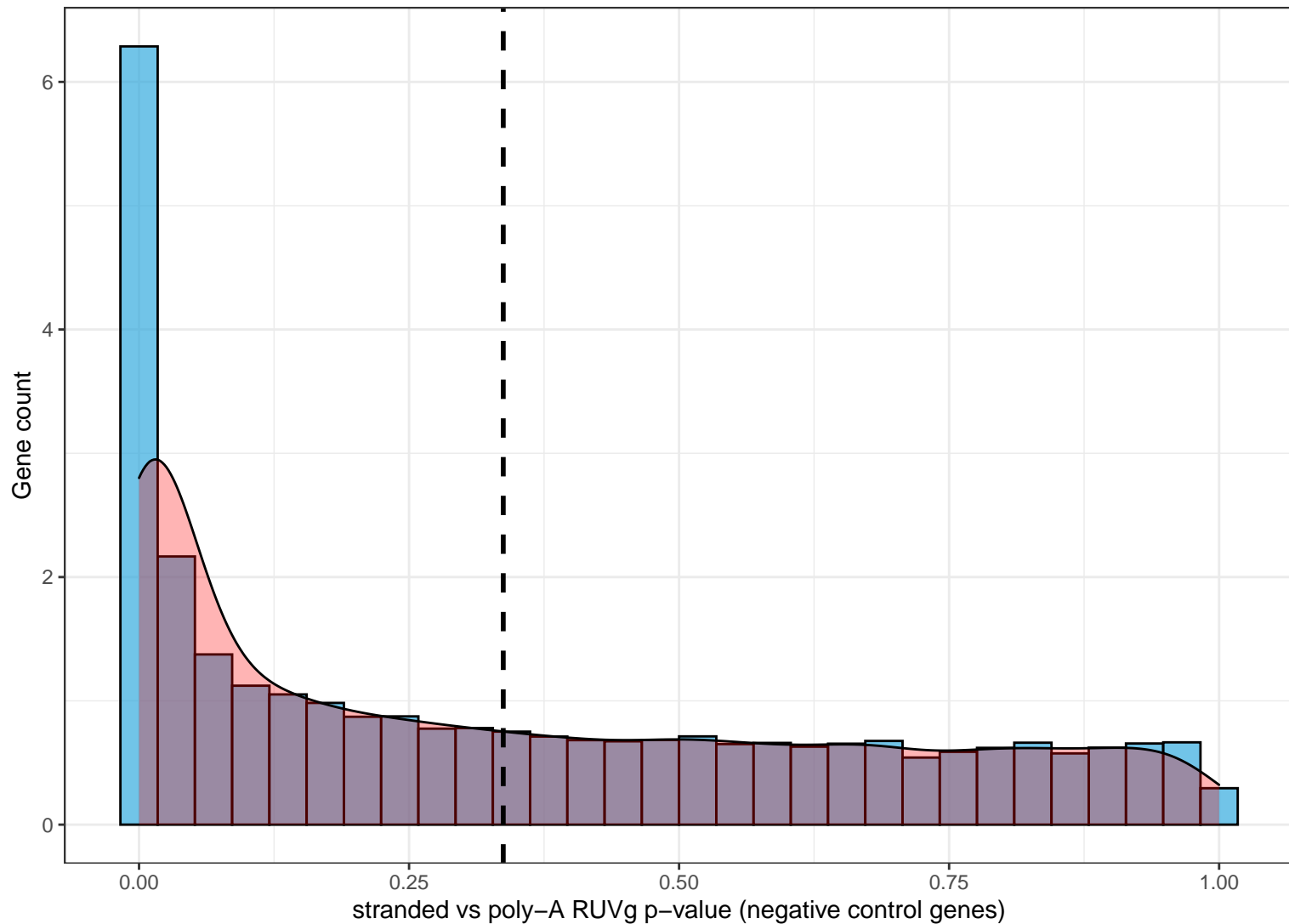
Histogram of stranded vs poly−A paired analysis (k = 4)
Total number of genes: 18138
5067 genes have p−value < 0.05
13071 genes have p−value >= 0.05
2744 genes have BH FDR < 0.05
15394 genes have BH FDR >= 0.05
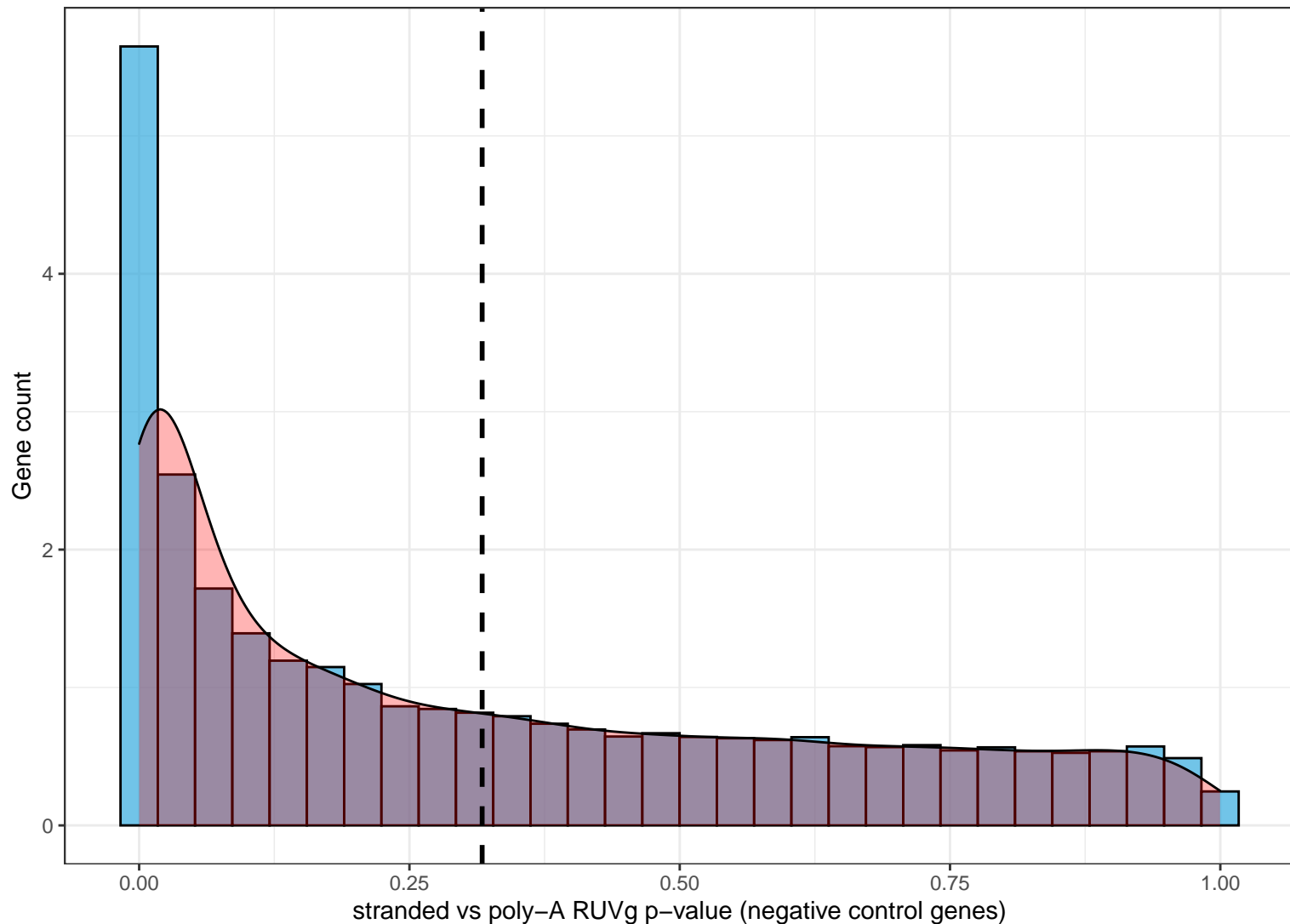
Histogram of stranded vs poly−A paired analysis (k = 5)
Total number of genes: 18138
4579 genes have p−value < 0.05
13559 genes have p−value >= 0.05
2652 genes have BH FDR < 0.05
15486 genes have BH FDR >= 0.05