Histogram of stranded vs poly−A paired analysis (k = 1)
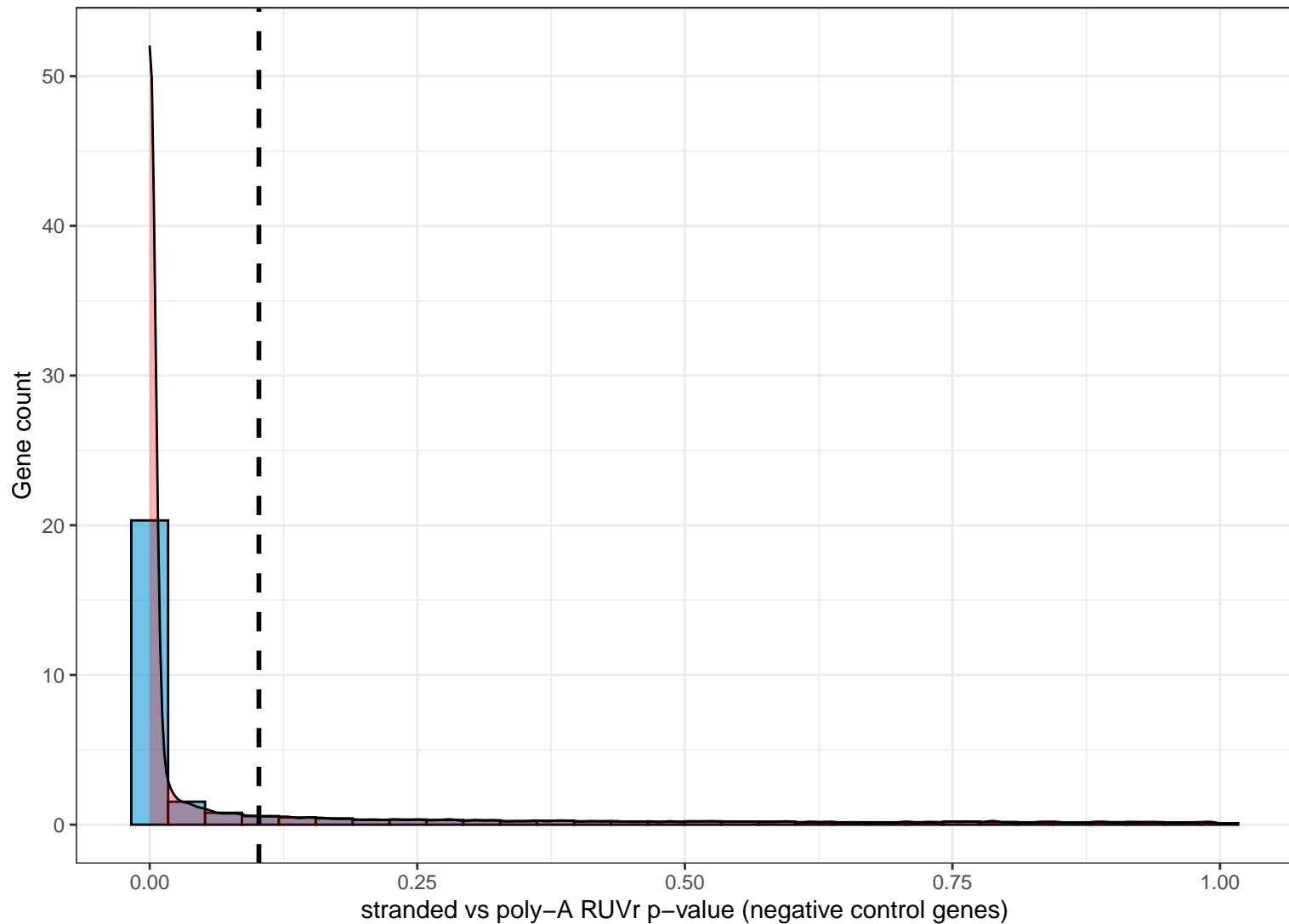Total number of genes: 15104
11344 genes have p−value < 0.05
3760 genes have p−value >= 0.05
11097 genes have BH FDR < 0.05
4007 genes have BH FDR >= 0.05

Histogram of stranded vs poly−A paired analysis (k = 2)
Total number of genes: 15104
10753 genes have p−value < 0.05
4351 genes have p−value >= 0.05
10438 genes have BH FDR < 0.05
4666 genes have BH FDR >= 0.05
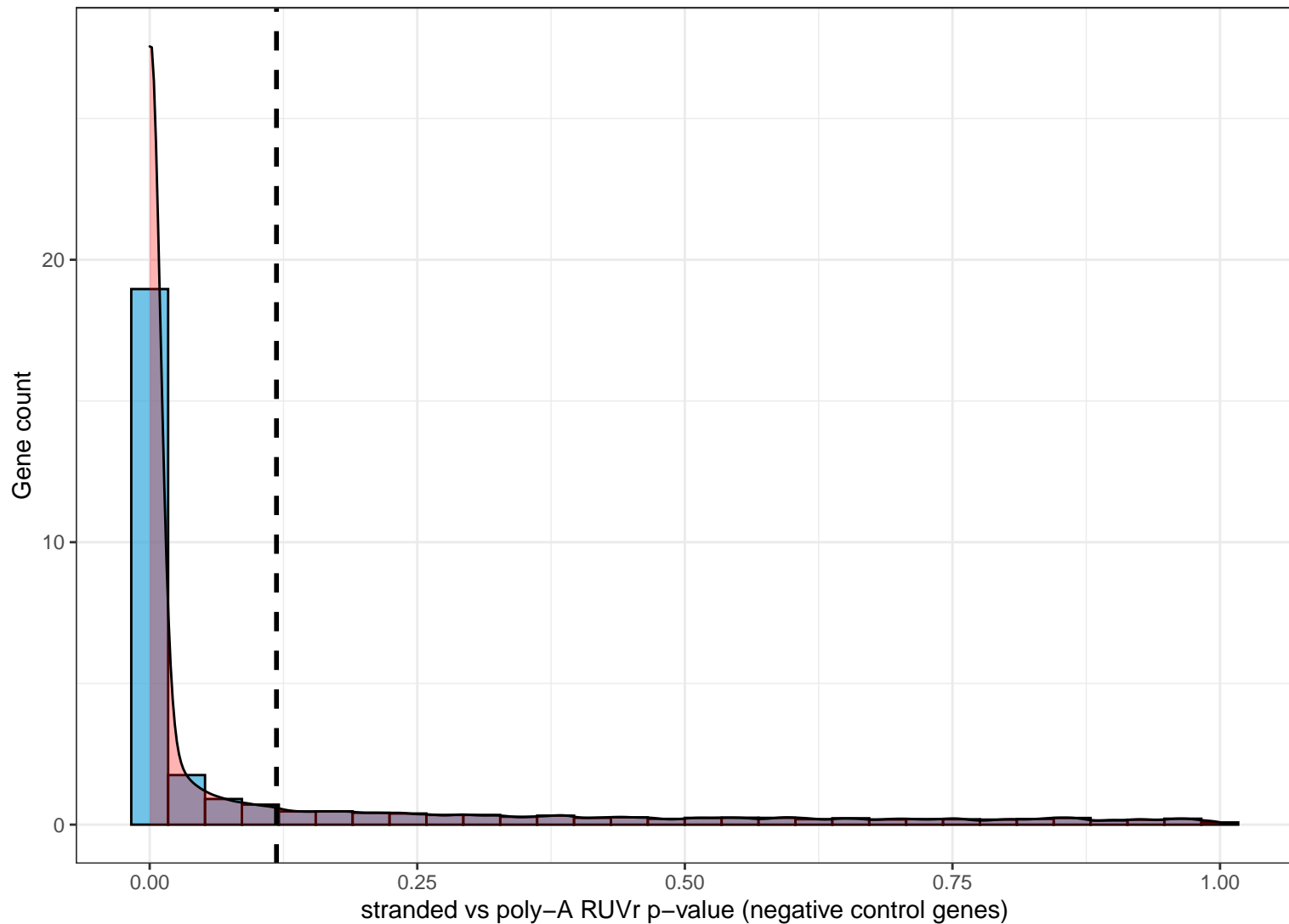
Histogram of stranded vs poly−A paired analysis (k = 3)
Total number of genes: 15104
10408 genes have p−value < 0.05
4696 genes have p−value >= 0.05
10026 genes have BH FDR < 0.05
5078 genes have BH FDR >= 0.05
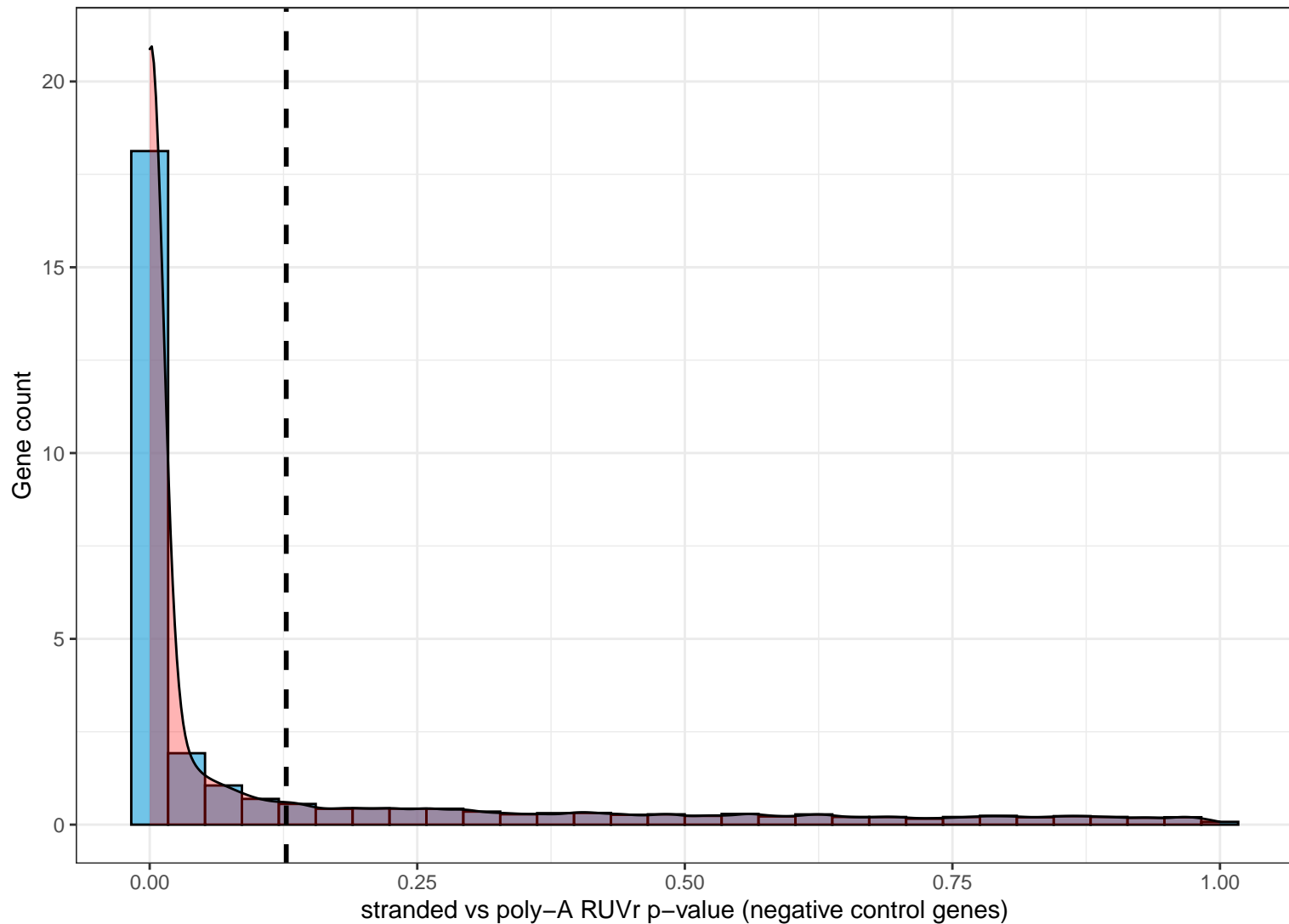
Histogram of stranded vs poly−A paired analysis (k = 4)
Total number of genes: 15104
10370 genes have p−value < 0.05
4734 genes have p−value >= 0.05
9945 genes have BH FDR < 0.05
5159 genes have BH FDR >= 0.05
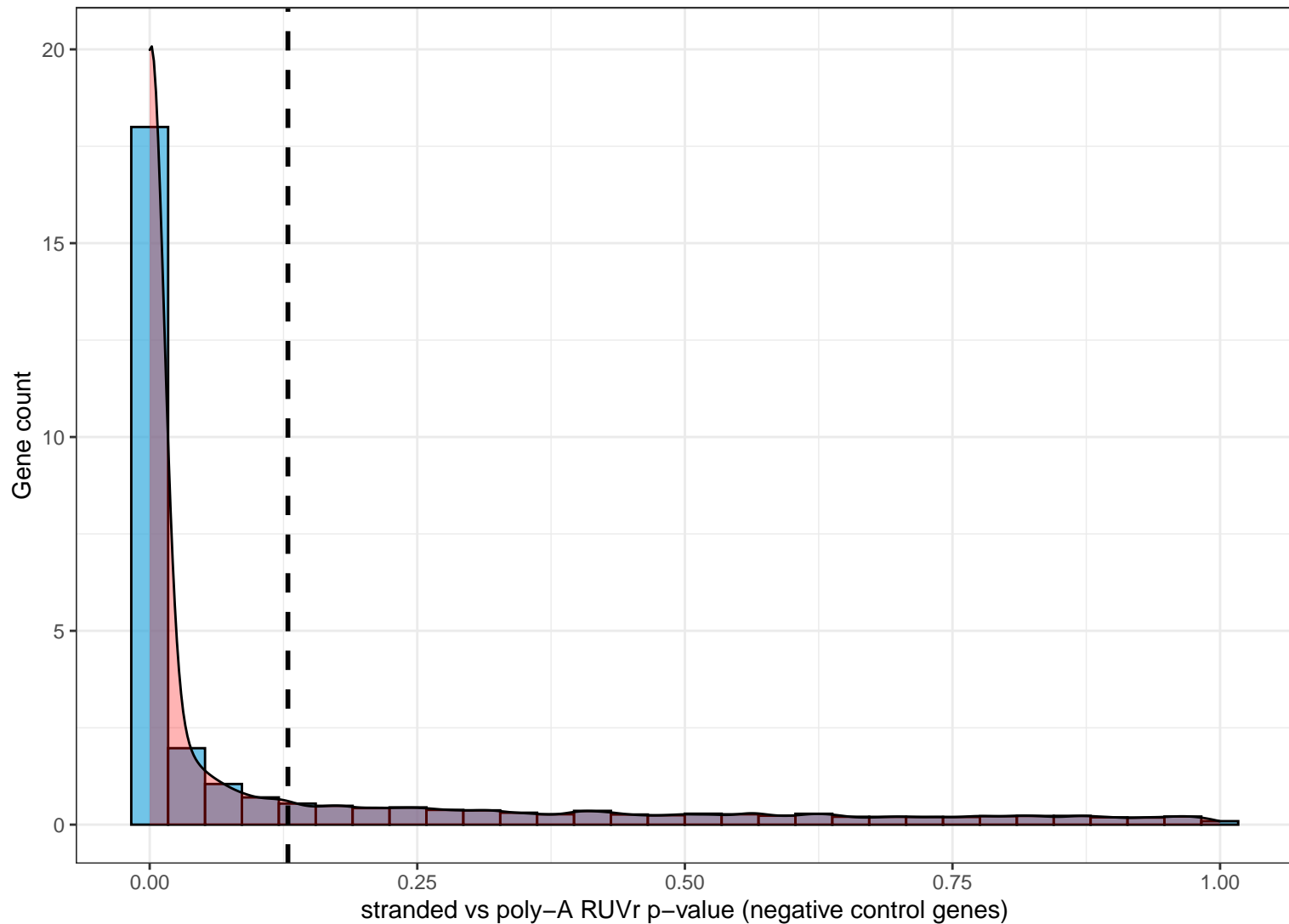
Histogram of stranded vs poly−A paired analysis (k = 5)
Total number of genes: 15104
10301 genes have p−value < 0.05
4803 genes have p−value >= 0.05
9910 genes have BH FDR < 0.05
5194 genes have BH FDR >= 0.05