Histogram of stranded vs poly−A paired analysis (k = 1)
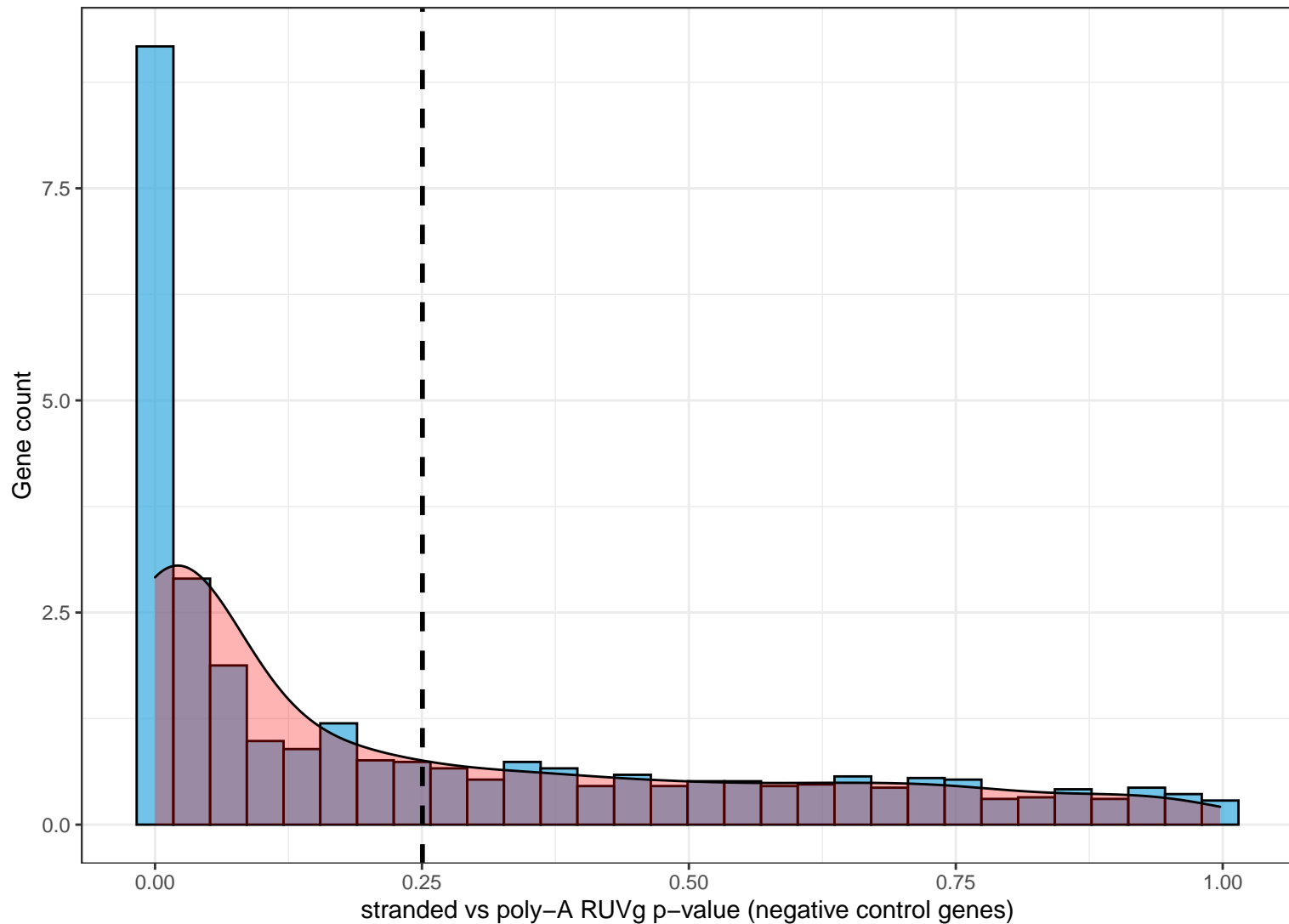Total number of genes: 1534
636 genes have p−value < 0.05
898 genes have p−value >= 0.05
454 genes have BH FDR < 0.05
1080 genes have BH FDR >= 0.05

Histogram of stranded vs poly−A paired analysis (k = 2)
Total number of genes: 1534
867 genes have p−value < 0.05
667 genes have p−value >= 0.05
763 genes have BH FDR < 0.05
771 genes have BH FDR >= 0.05
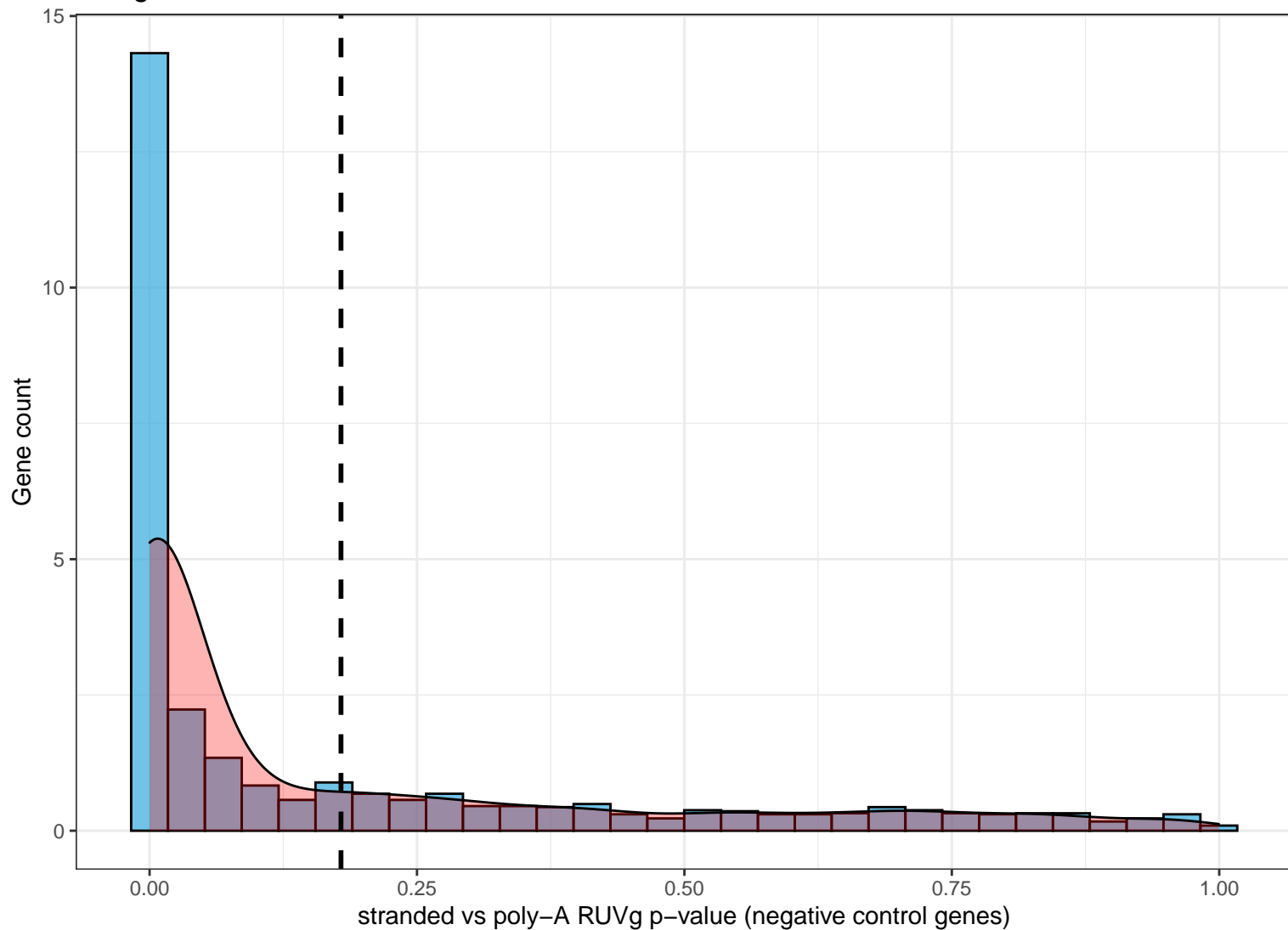
Histogram of stranded vs poly−A paired analysis (k = 3)
Total number of genes: 1534
1435 genes have p−value < 0.05
99 genes have p−value >= 0.05
1422 genes have BH FDR < 0.05
112 genes have BH FDR >= 0.05
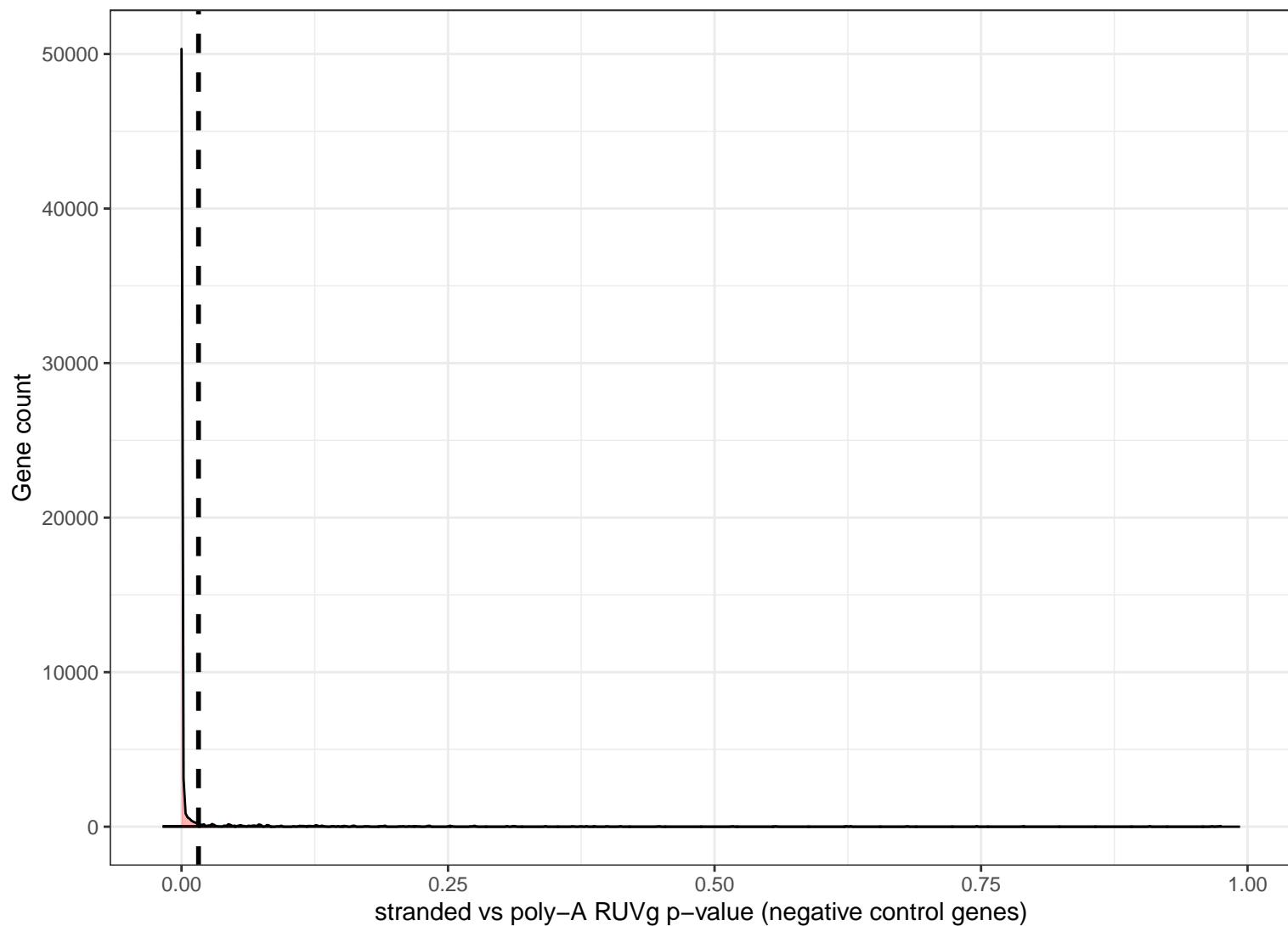
Histogram of stranded vs poly−A paired analysis (k = 4)
Total number of genes: 1534
1360 genes have p−value < 0.05
174 genes have p−value >= 0.05
1330 genes have BH FDR < 0.05
204 genes have BH FDR >= 0.05
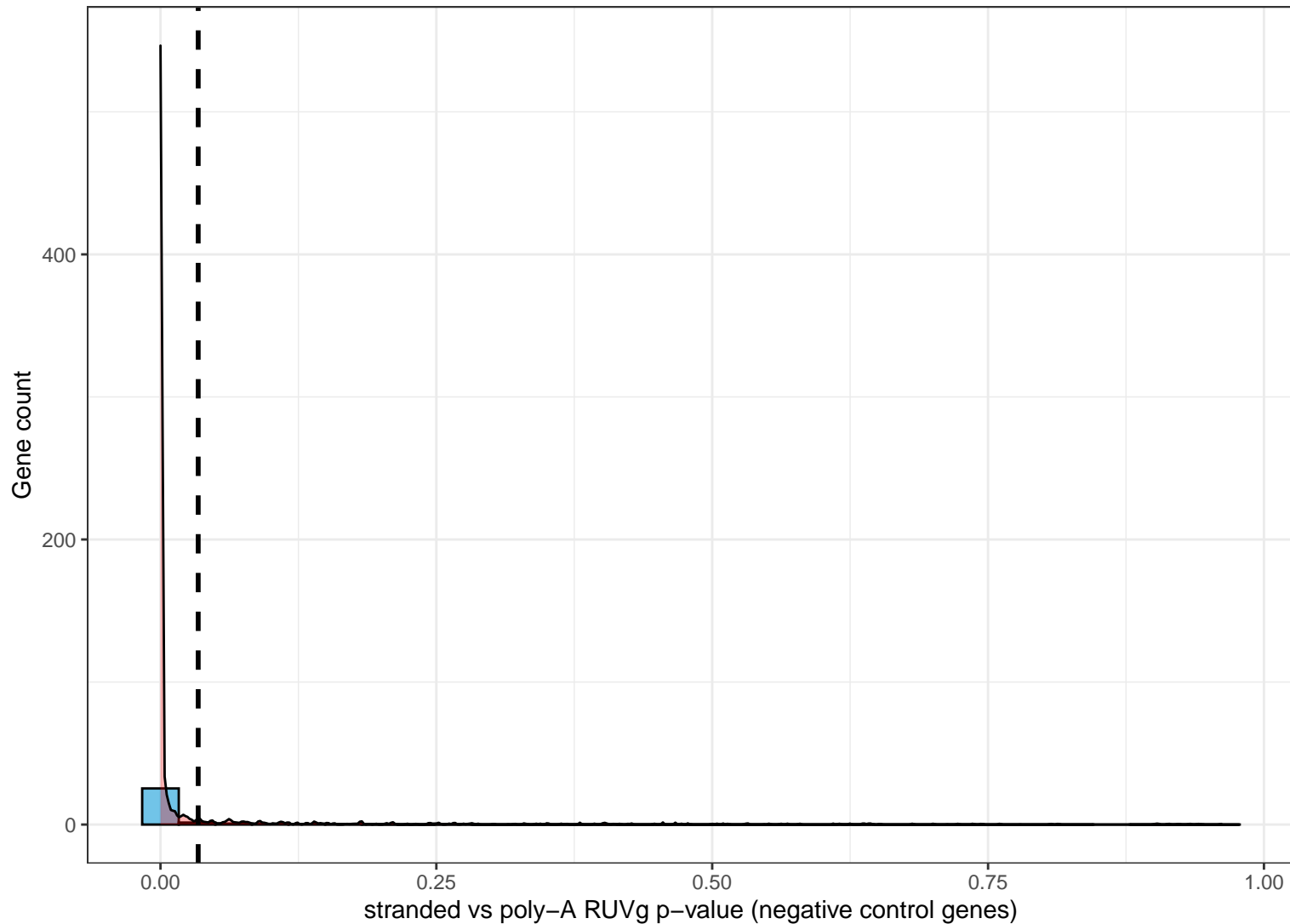
Histogram of stranded vs poly−A paired analysis (k = 5)
Total number of genes: 1534
1534 genes have p−value < 0.05
0 genes have p−value >= 0.05
1534 genes have BH FDR < 0.05
0 genes have BH FDR >= 0.05