Histogram of stranded vs poly−A paired analysis (k = 1)
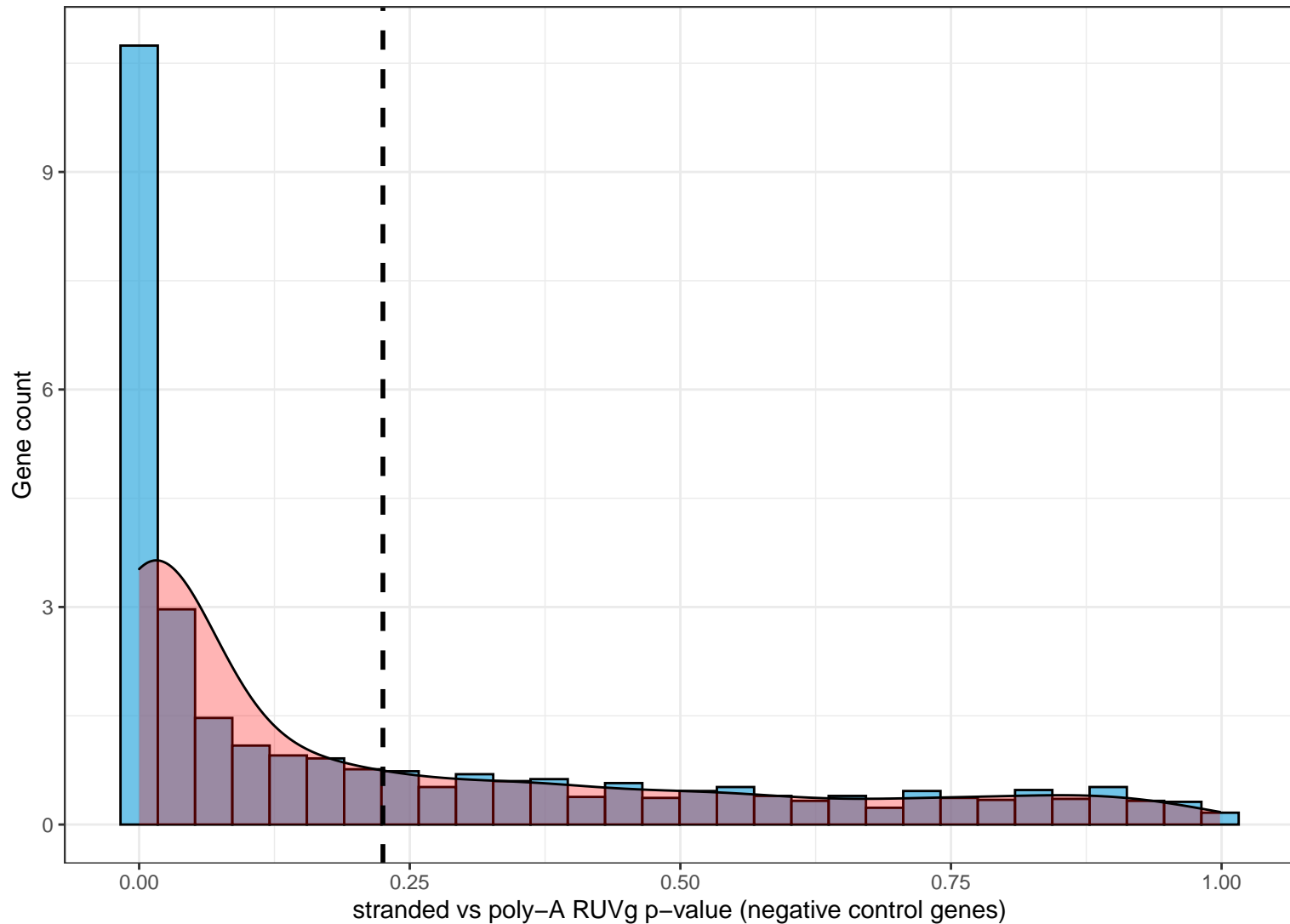Total number of genes: 2133
998 genes have p−value < 0.05
1135 genes have p−value >= 0.05
756 genes have BH FDR < 0.05
1377 genes have BH FDR >= 0.05

Histogram of stranded vs poly−A paired analysis (k = 2)
Total number of genes: 2133
1231 genes have p−value < 0.05
902 genes have p−value >= 0.05
1095 genes have BH FDR < 0.05
1038 genes have BH FDR >= 0.05
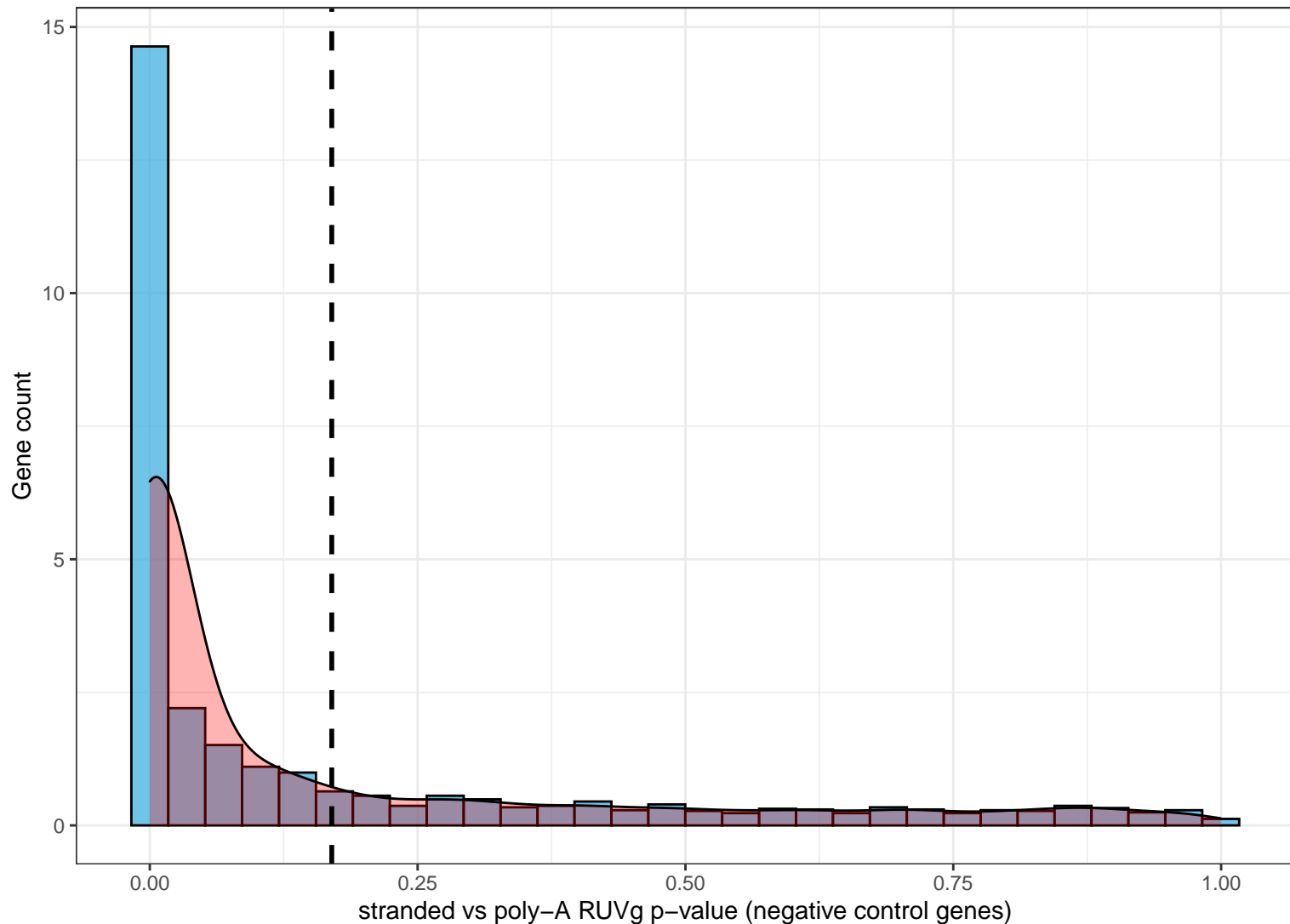
Histogram of stranded vs poly−A paired analysis (k = 3)
Total number of genes: 2133
1698 genes have p−value < 0.05
435 genes have p−value >= 0.05
1653 genes have BH FDR < 0.05
480 genes have BH FDR >= 0.05
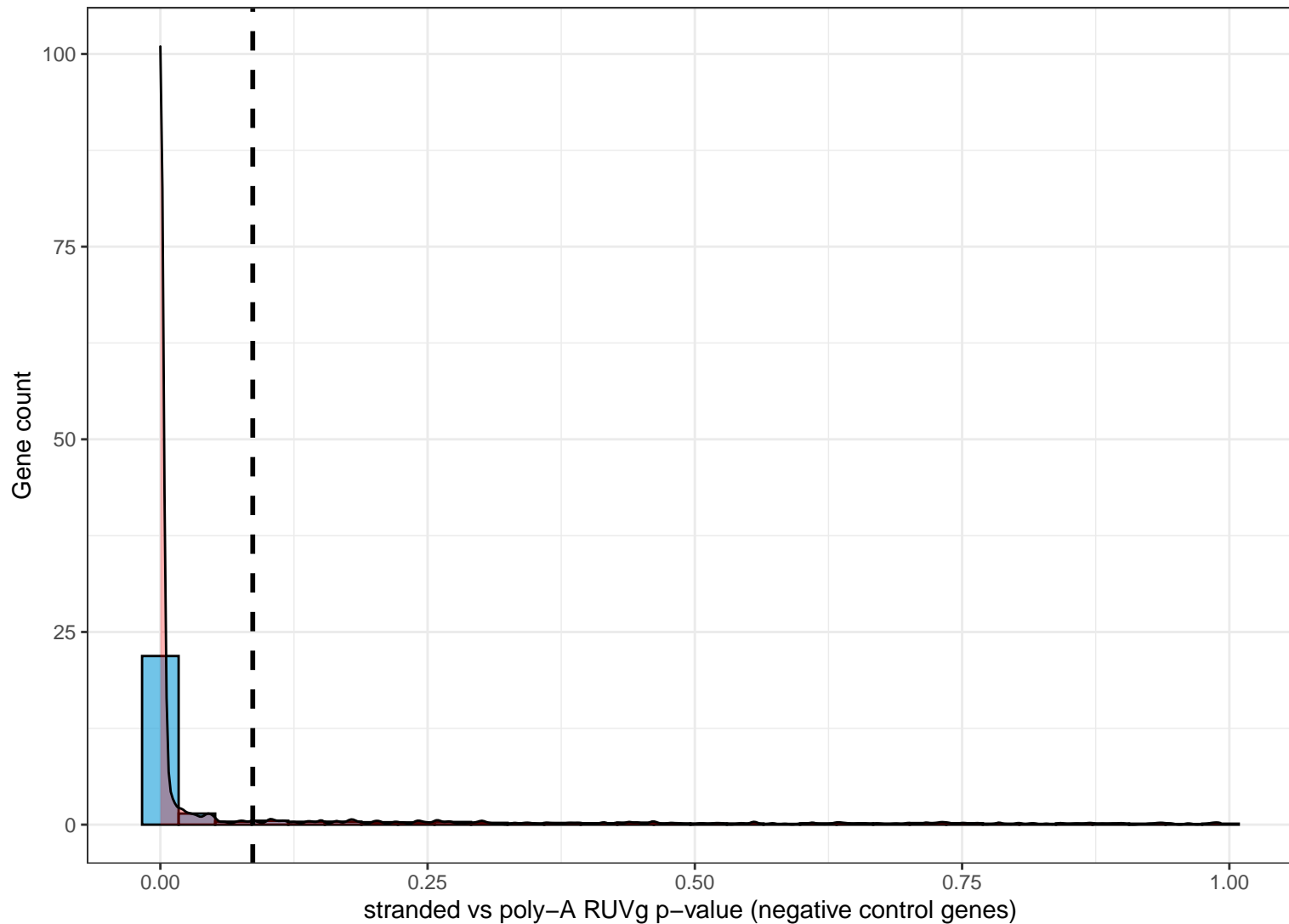
Histogram of stranded vs poly−A paired analysis (k = 4)
Total number of genes: 2133
1544 genes have p−value < 0.05
589 genes have p−value >= 0.05
1493 genes have BH FDR < 0.05
640 genes have BH FDR >= 0.05
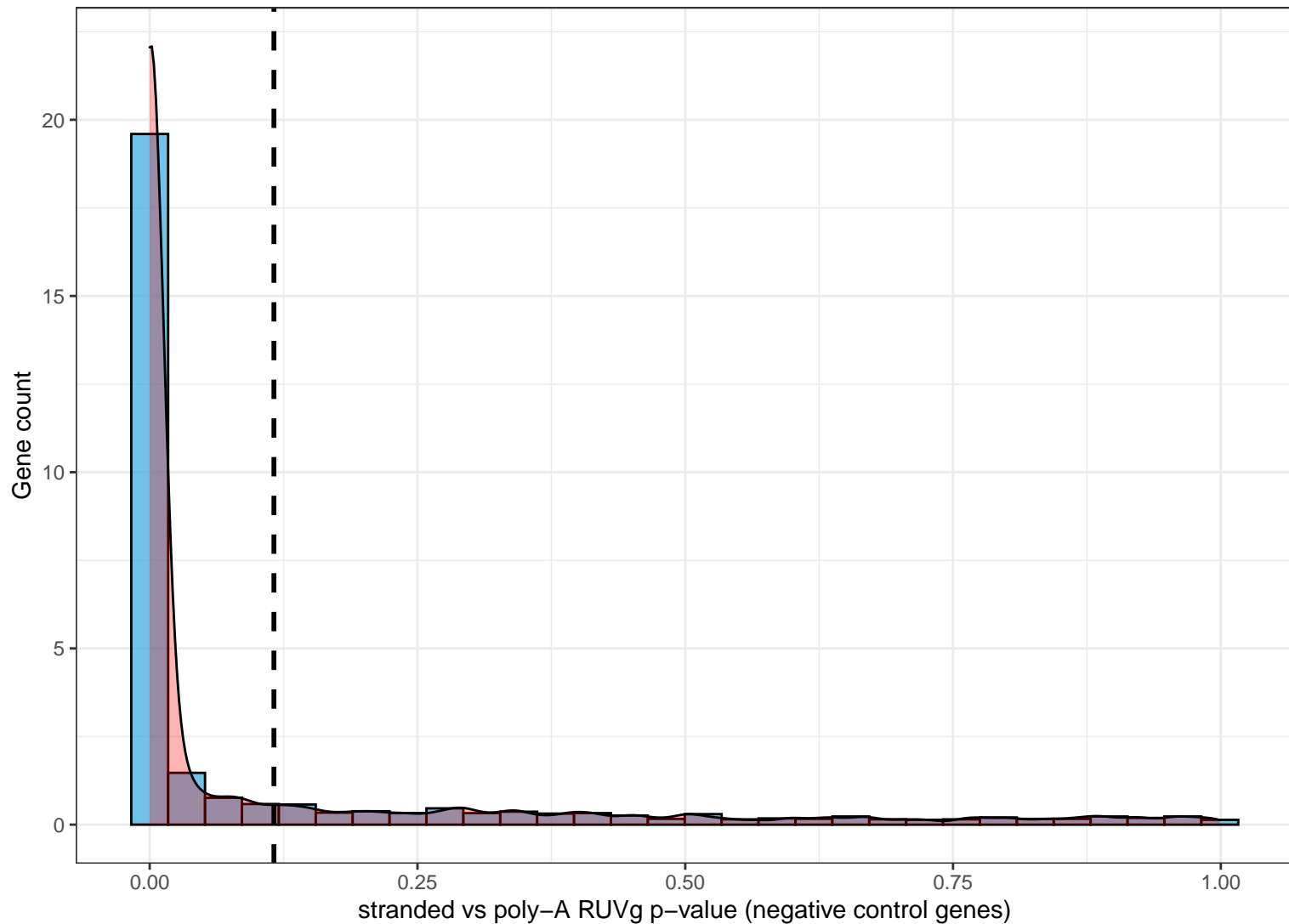
Histogram of stranded vs poly−A paired analysis (k = 5)
Total number of genes: 2133
1700 genes have p−value < 0.05
433 genes have p−value >= 0.05
1670 genes have BH FDR < 0.05
463 genes have BH FDR >= 0.05