Histogram of stranded vs poly−A paired analysis (k = 1)
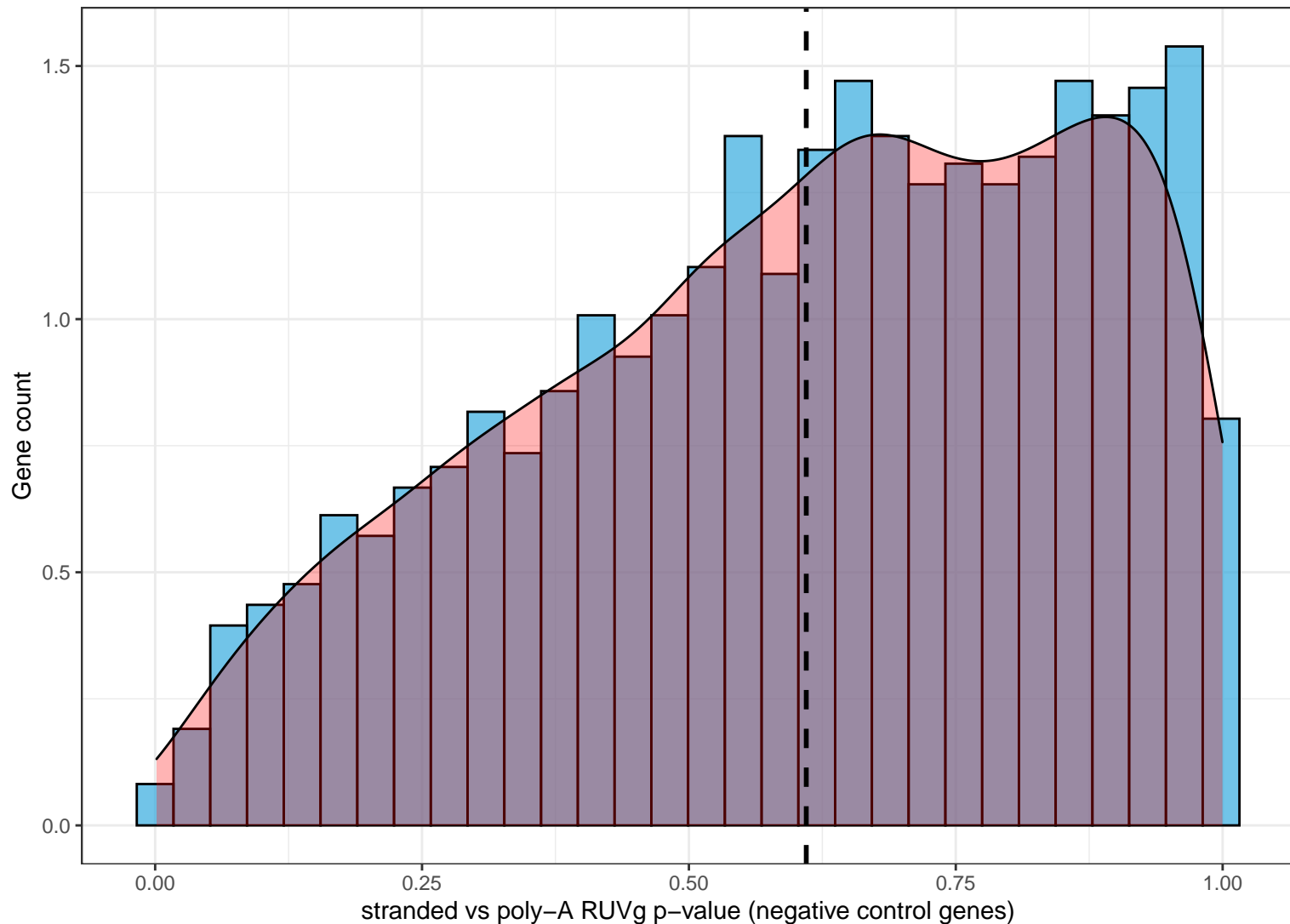Total number of genes: 2133
20 genes have p−value < 0.05
2113 genes have p−value >= 0.05
2 genes have BH FDR < 0.05
2131 genes have BH FDR >= 0.05

Histogram of stranded vs poly−A paired analysis (k = 2)
Total number of genes: 2133
17 genes have p−value < 0.05
2116 genes have p−value >= 0.05
2 genes have BH FDR < 0.05
2131 genes have BH FDR >= 0.05
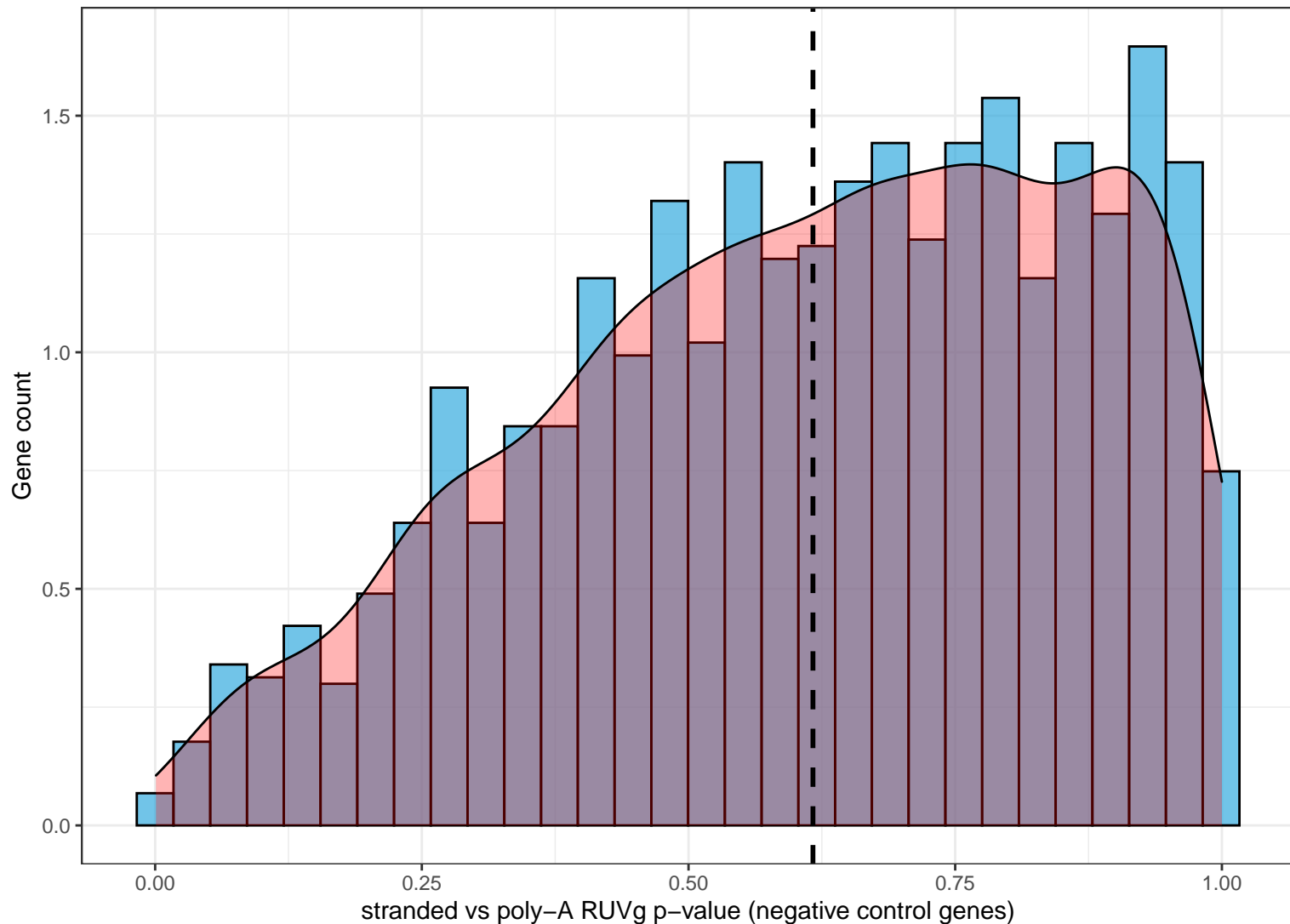
Histogram of stranded vs poly−A paired analysis (k = 3)
Total number of genes: 2133
270 genes have p−value < 0.05
1863 genes have p−value >= 0.05
64 genes have BH FDR < 0.05
2069 genes have BH FDR >= 0.05
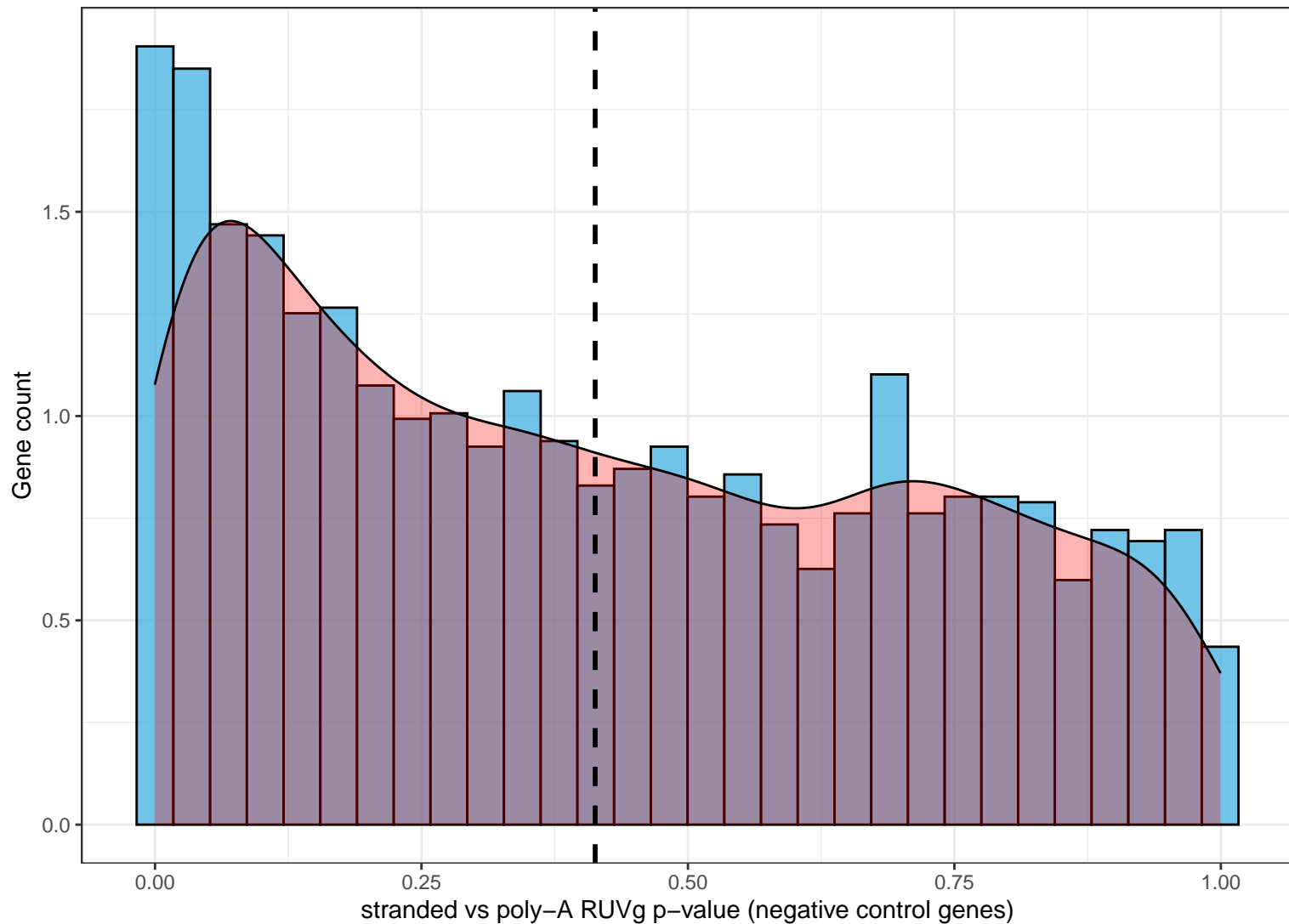
Histogram of stranded vs poly−A paired analysis (k = 4)
Total number of genes: 2133
327 genes have p−value < 0.05
1806 genes have p−value >= 0.05
81 genes have BH FDR < 0.05
2052 genes have BH FDR >= 0.05
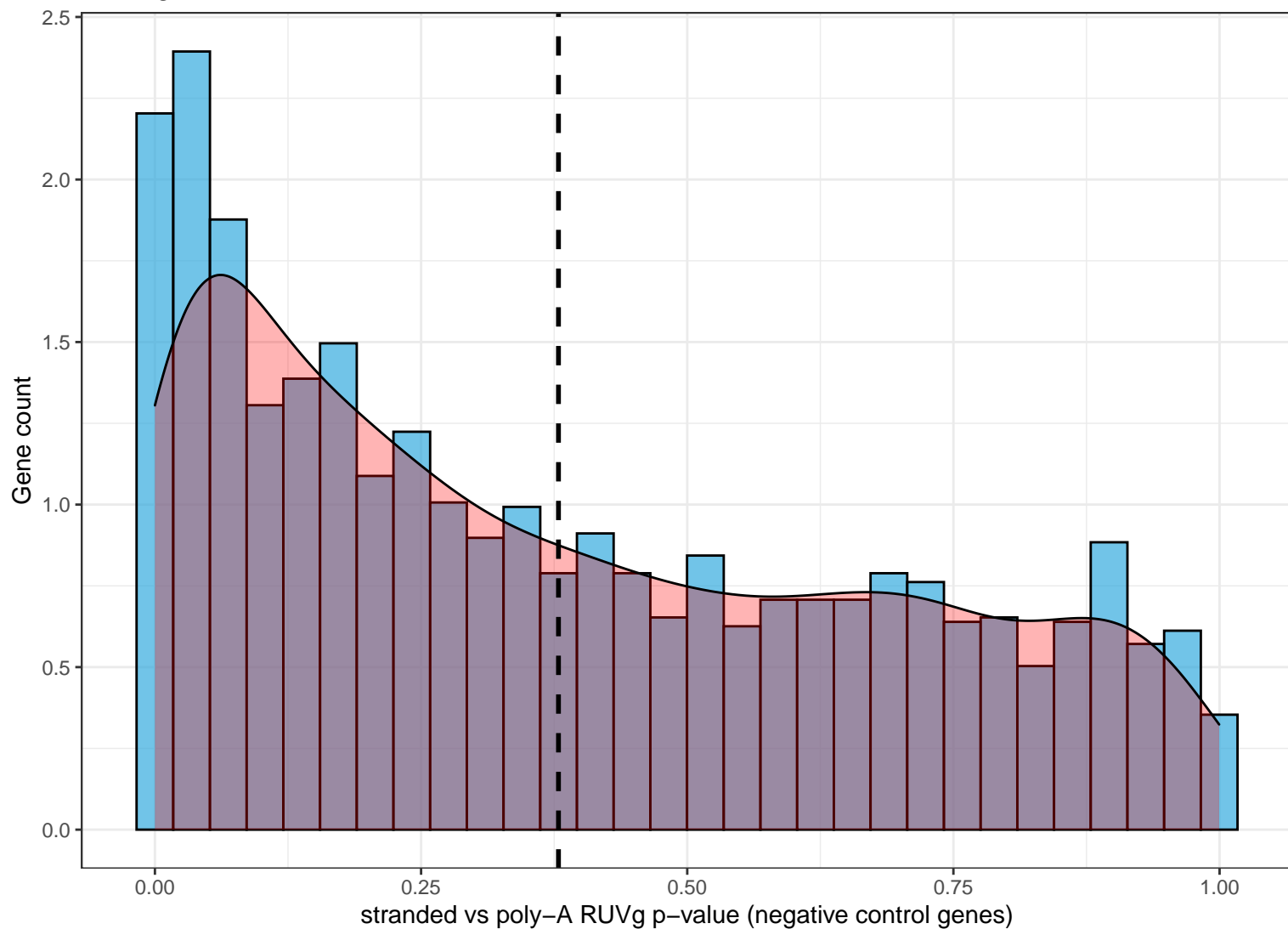
Histogram of stranded vs poly−A paired analysis (k = 5)
Total number of genes: 2133
175 genes have p−value < 0.05
1958 genes have p−value >= 0.05
18 genes have BH FDR < 0.05
2115 genes have BH FDR >= 0.05