

Histogram of stranded vs poly-A paired analysis (k = 1)

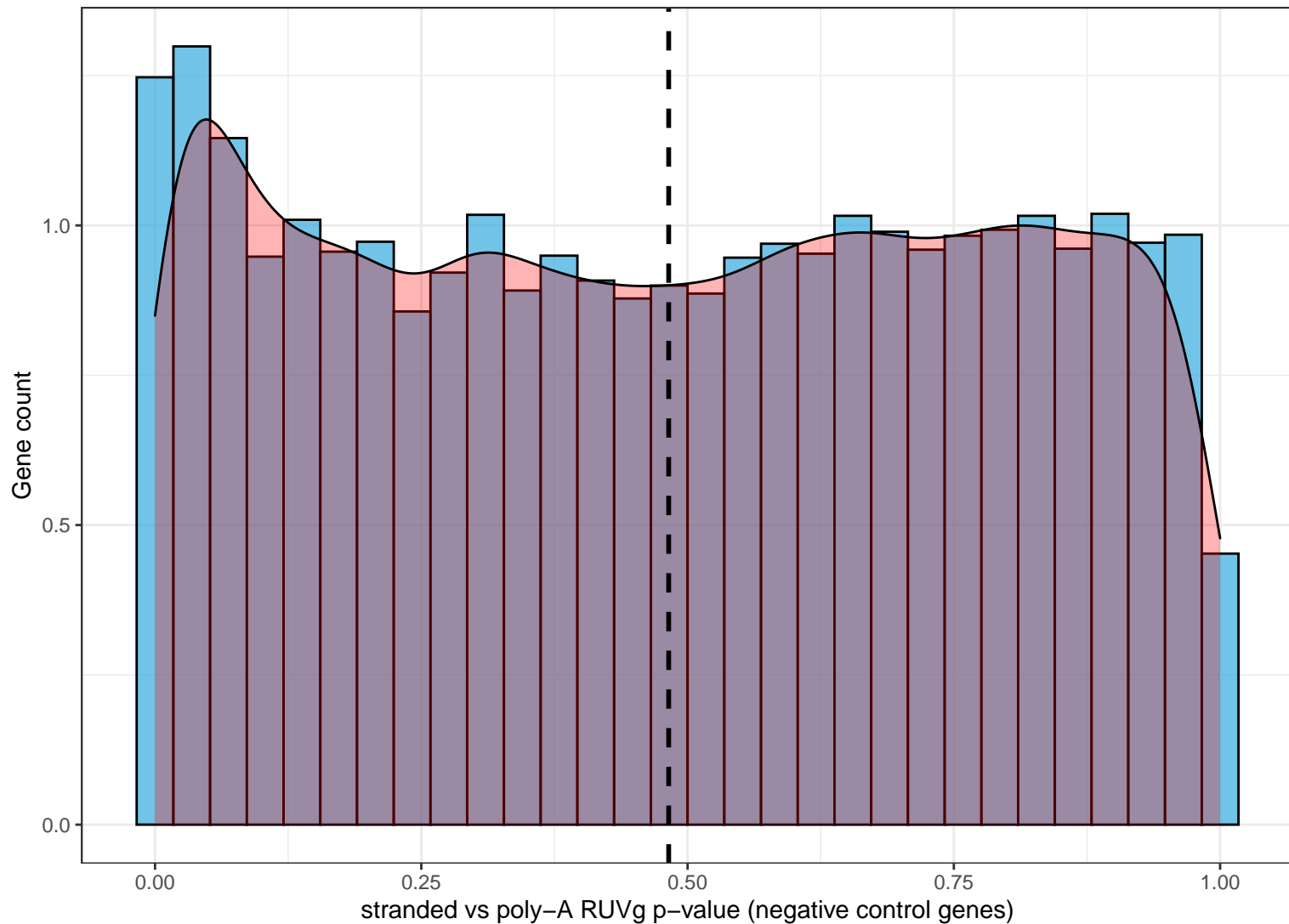
Total number of genes: 17439

1494 genes have p-value < 0.05

15945 genes have p-value >= 0.05

170 genes have BH FDR < 0.05

17269 genes have BH FDR >= 0.05



Histogram of stranded vs poly-A paired analysis ($k = 2$)

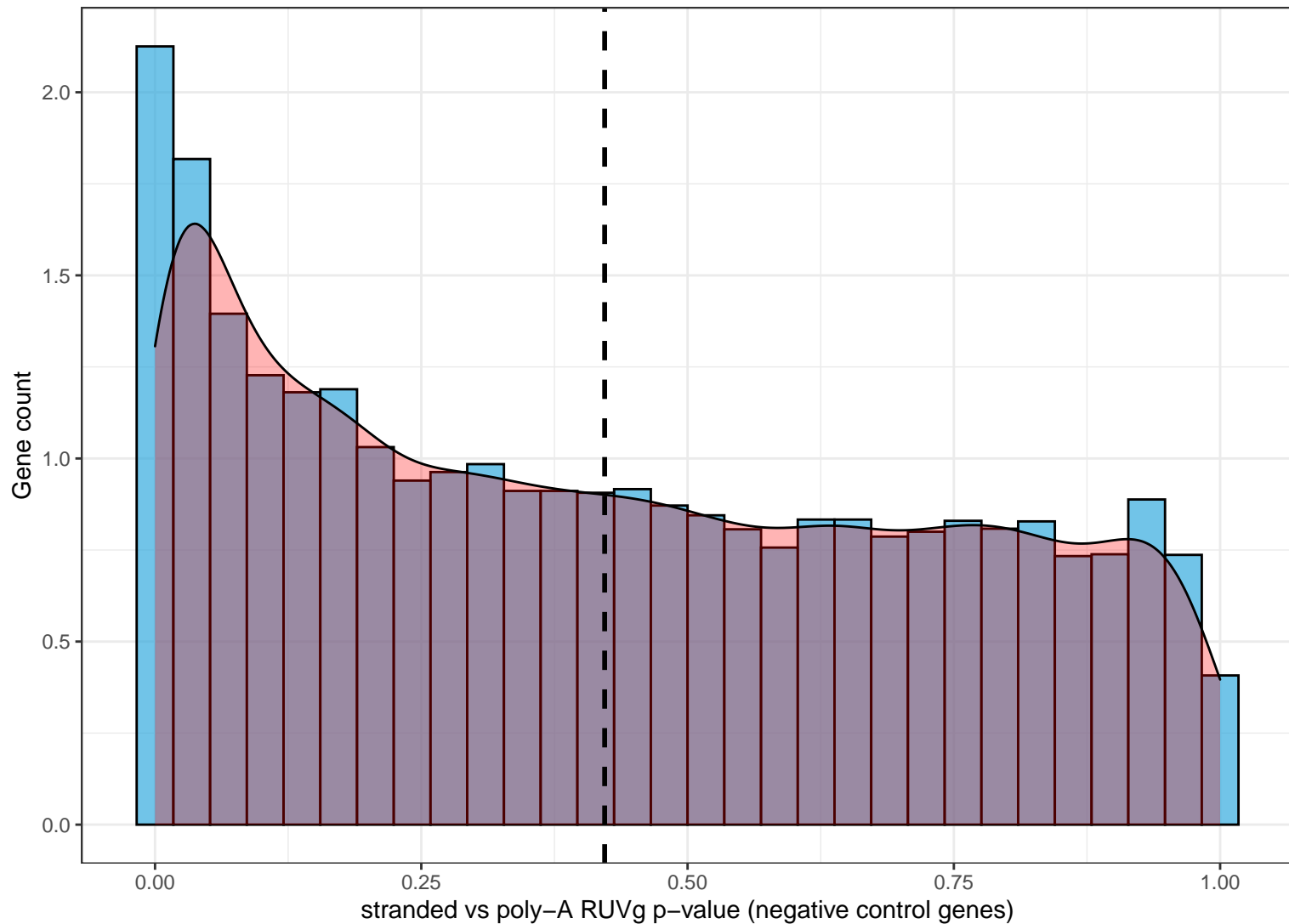
Total number of genes: 17439

2321 genes have p-value < 0.05

15118 genes have p-value ≥ 0.05

557 genes have BH FDR < 0.05

16882 genes have BH FDR ≥ 0.05



Histogram of stranded vs poly-A paired analysis (k = 3)

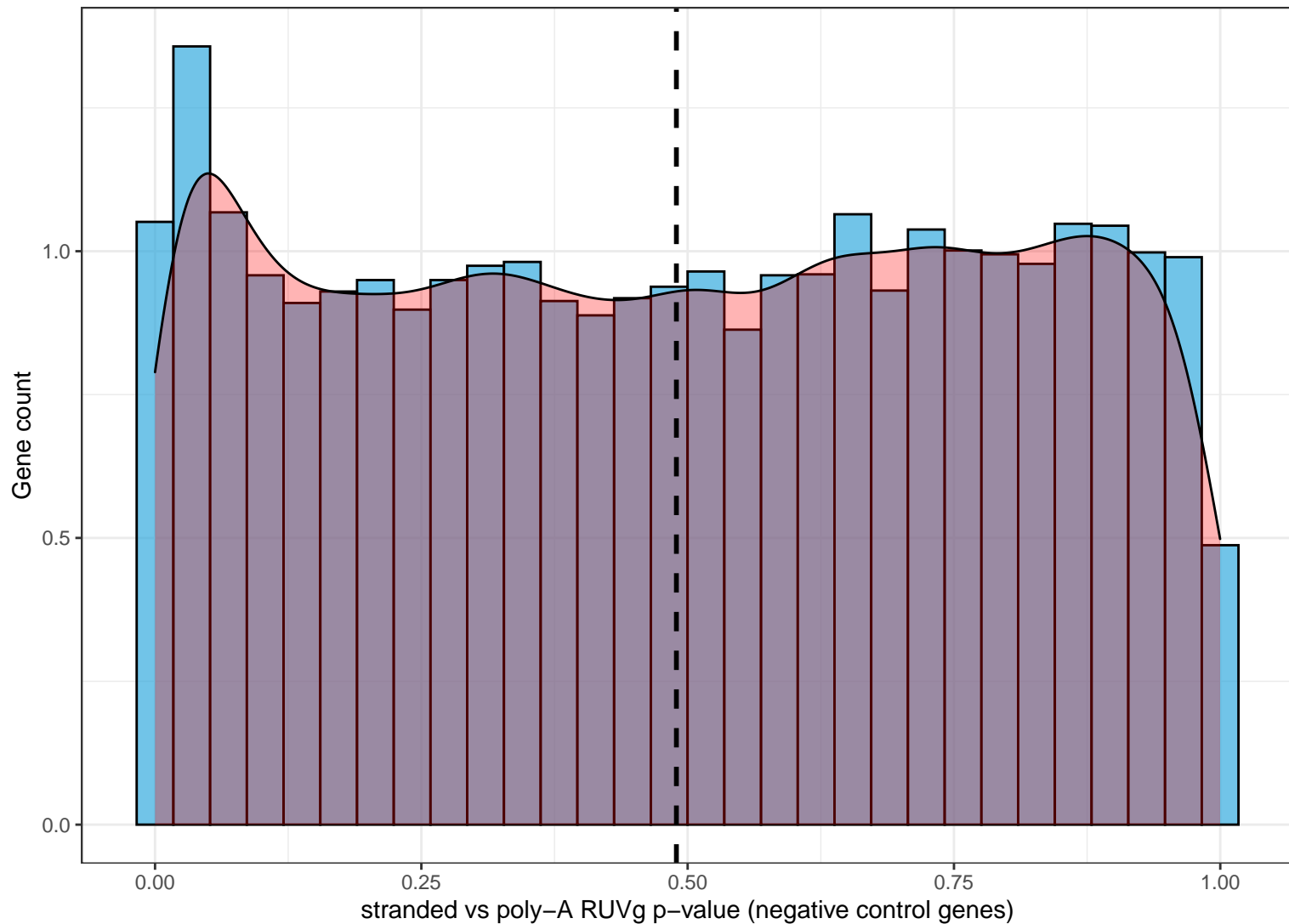
Total number of genes: 17439

1410 genes have p-value < 0.05

16029 genes have p-value ≥ 0.05

117 genes have BH FDR < 0.05

17322 genes have BH FDR ≥ 0.05



Histogram of stranded vs poly-A paired analysis (k = 4)

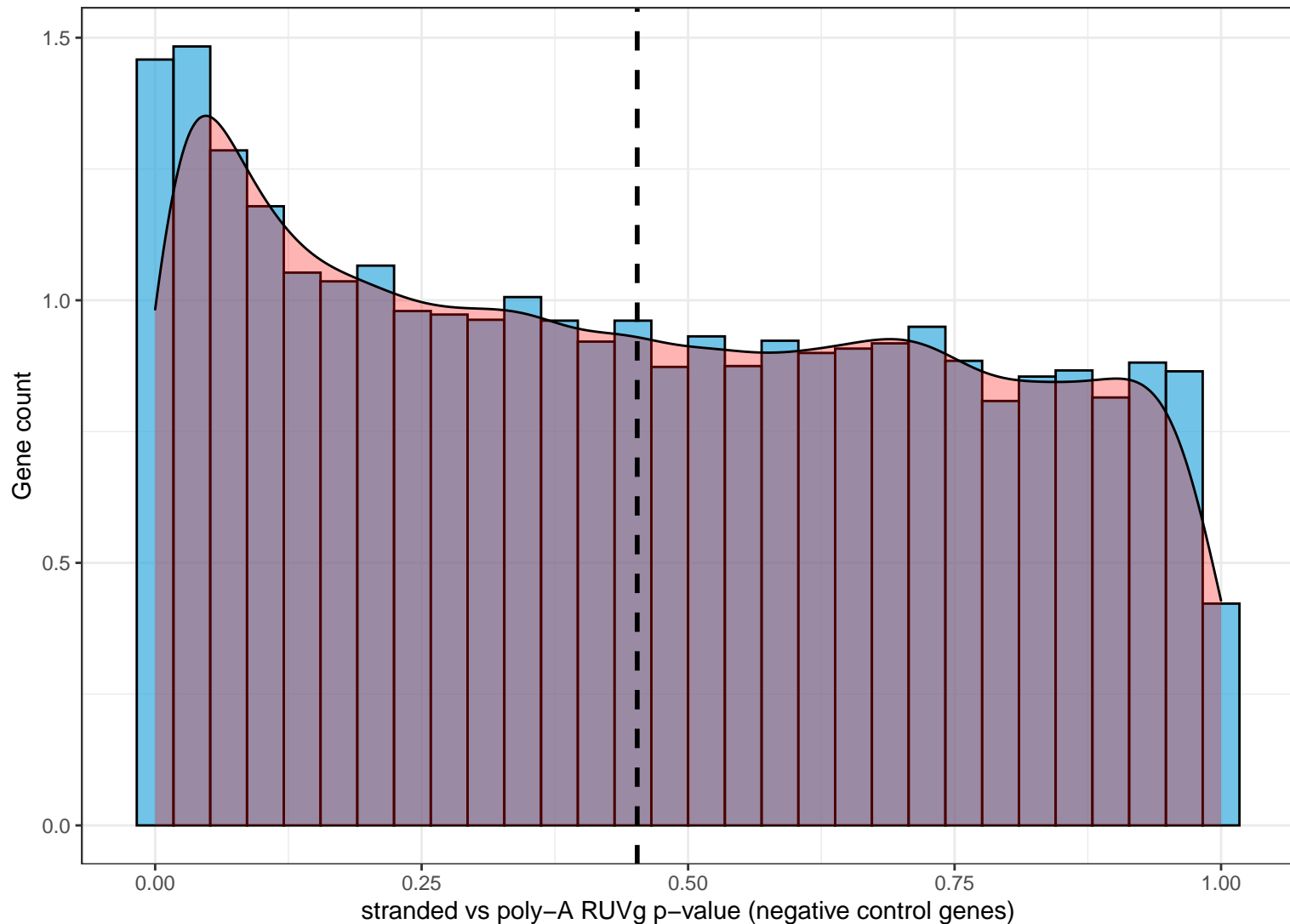
Total number of genes: 17439

1731 genes have p-value < 0.05

15708 genes have p-value ≥ 0.05

203 genes have BH FDR < 0.05

17236 genes have BH FDR ≥ 0.05



Histogram of stranded vs poly-A paired analysis (k = 5)

Total number of genes: 17439

1439 genes have p-value < 0.05

16000 genes have p-value ≥ 0.05

226 genes have BH FDR < 0.05

17213 genes have BH FDR ≥ 0.05

