

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

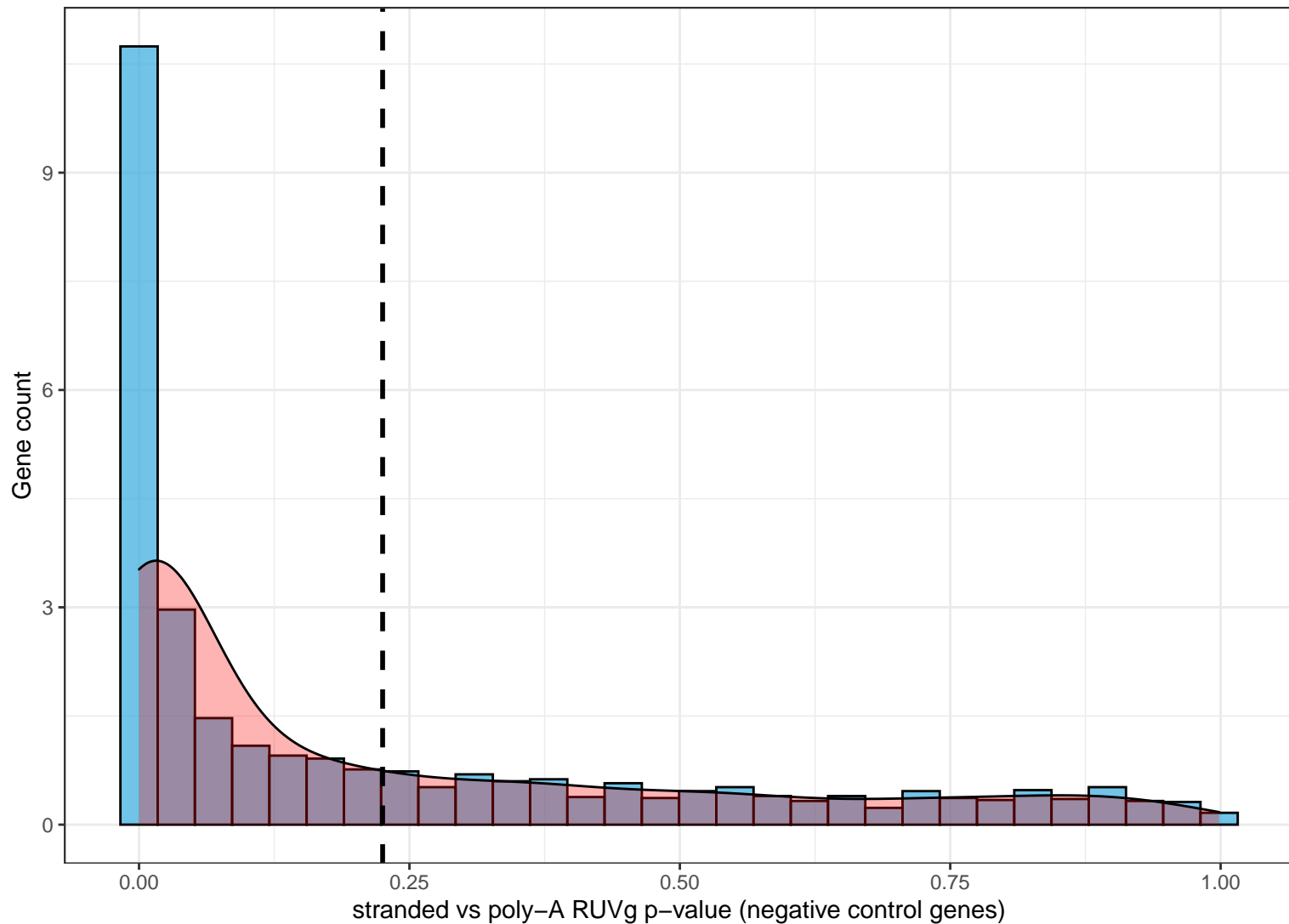
Total number of genes: 2133

998 genes have $p\text{-value} < 0.05$

1135 genes have $p\text{-value} \geq 0.05$

756 genes have BH FDR < 0.05

1377 genes have BH FDR ≥ 0.05



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

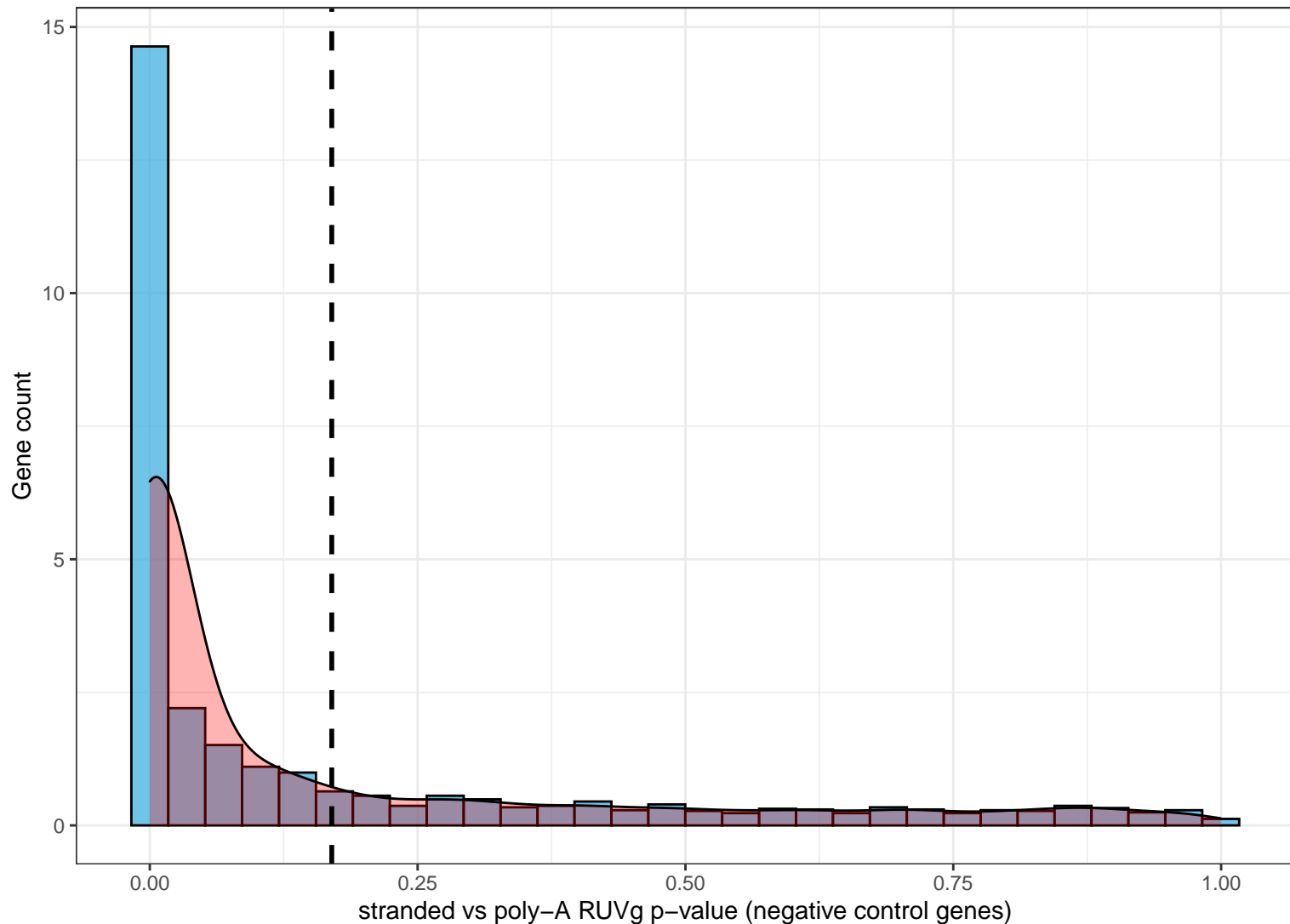
Total number of genes: 2133

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)

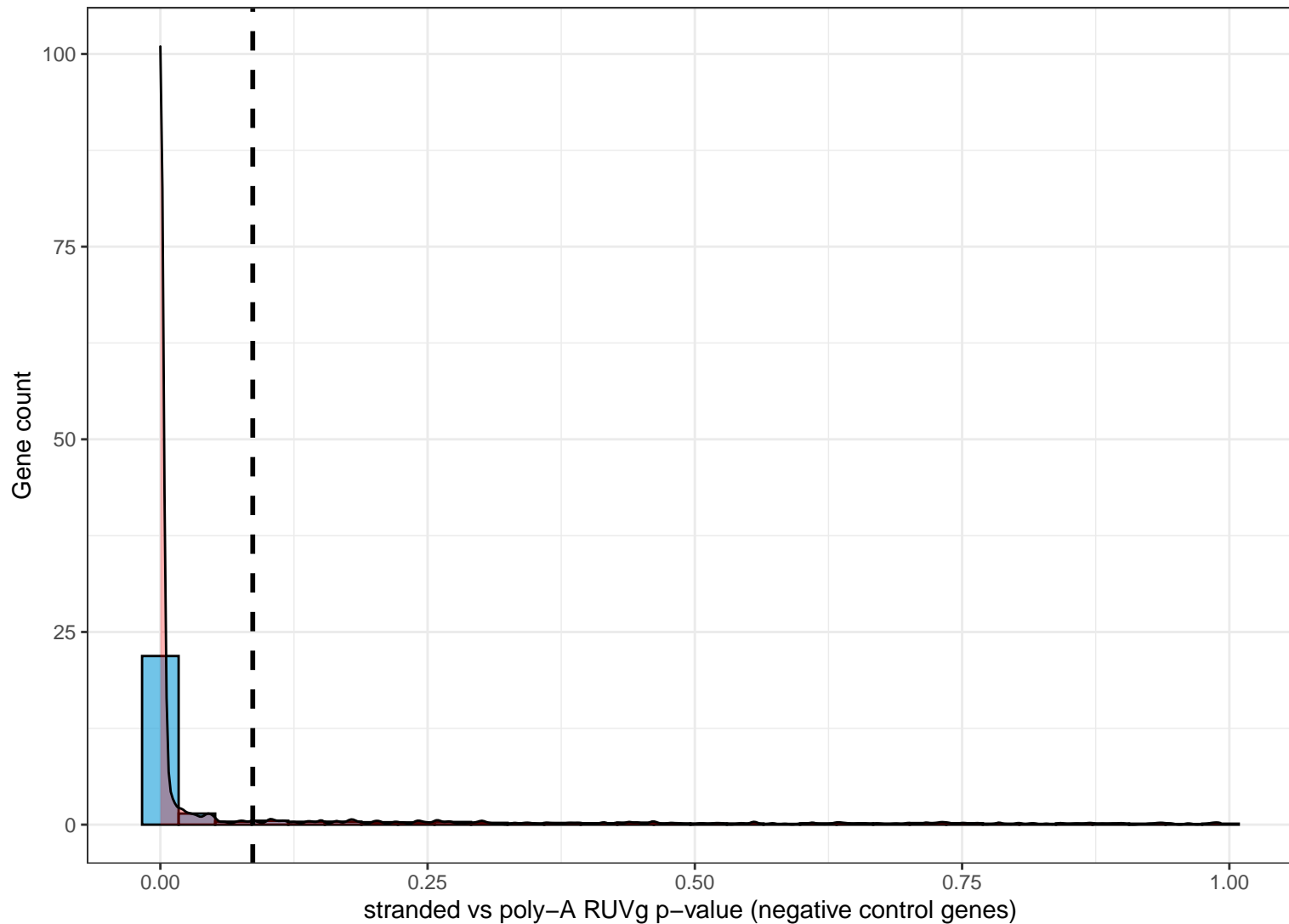
Total number of genes: 2133

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 ($k = 4$)

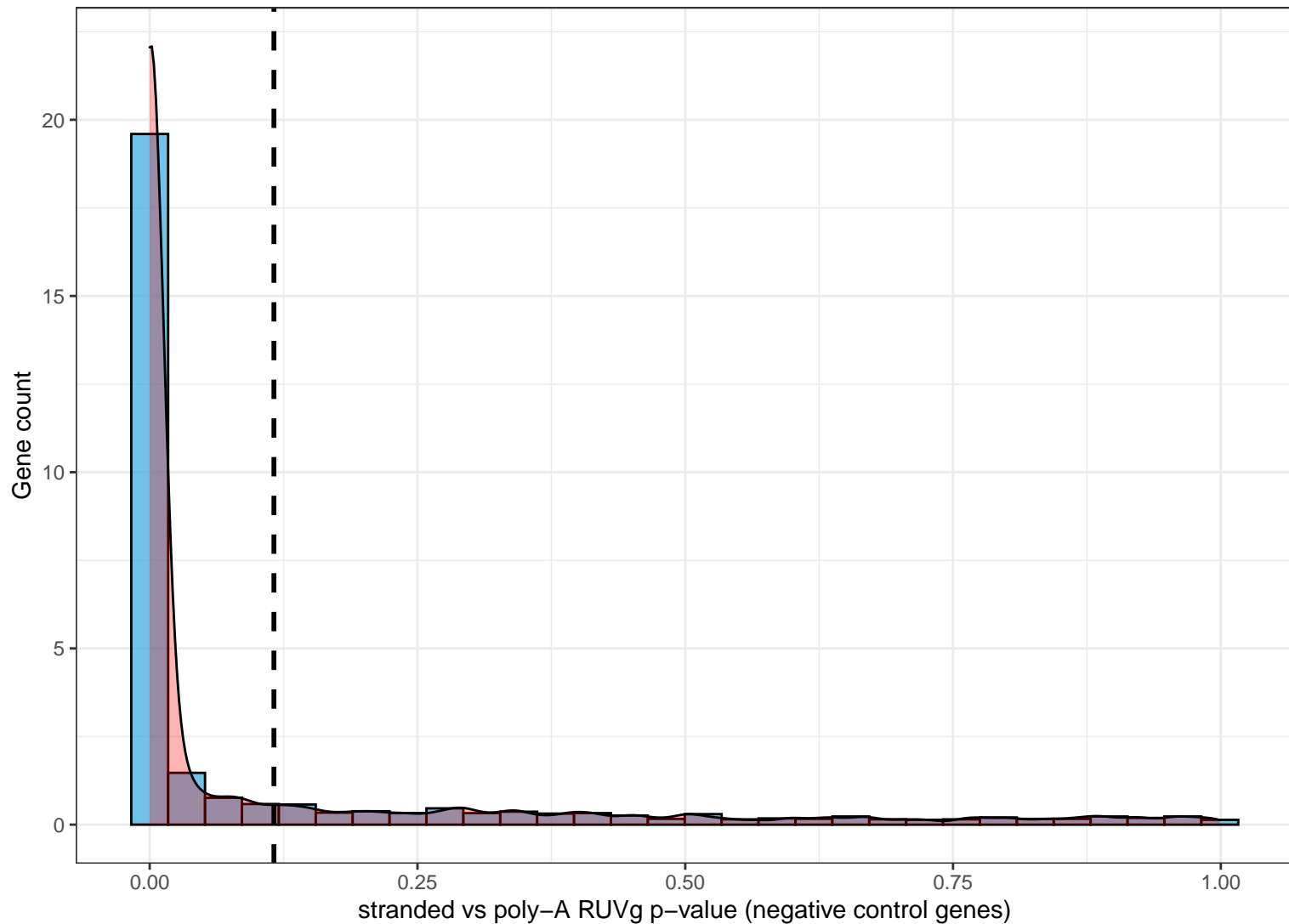
Total number of genes: 2133

1544 genes have $p\text{-value} < 0.05$

589 genes have $p\text{-value} \geq 0.05$

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

