

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

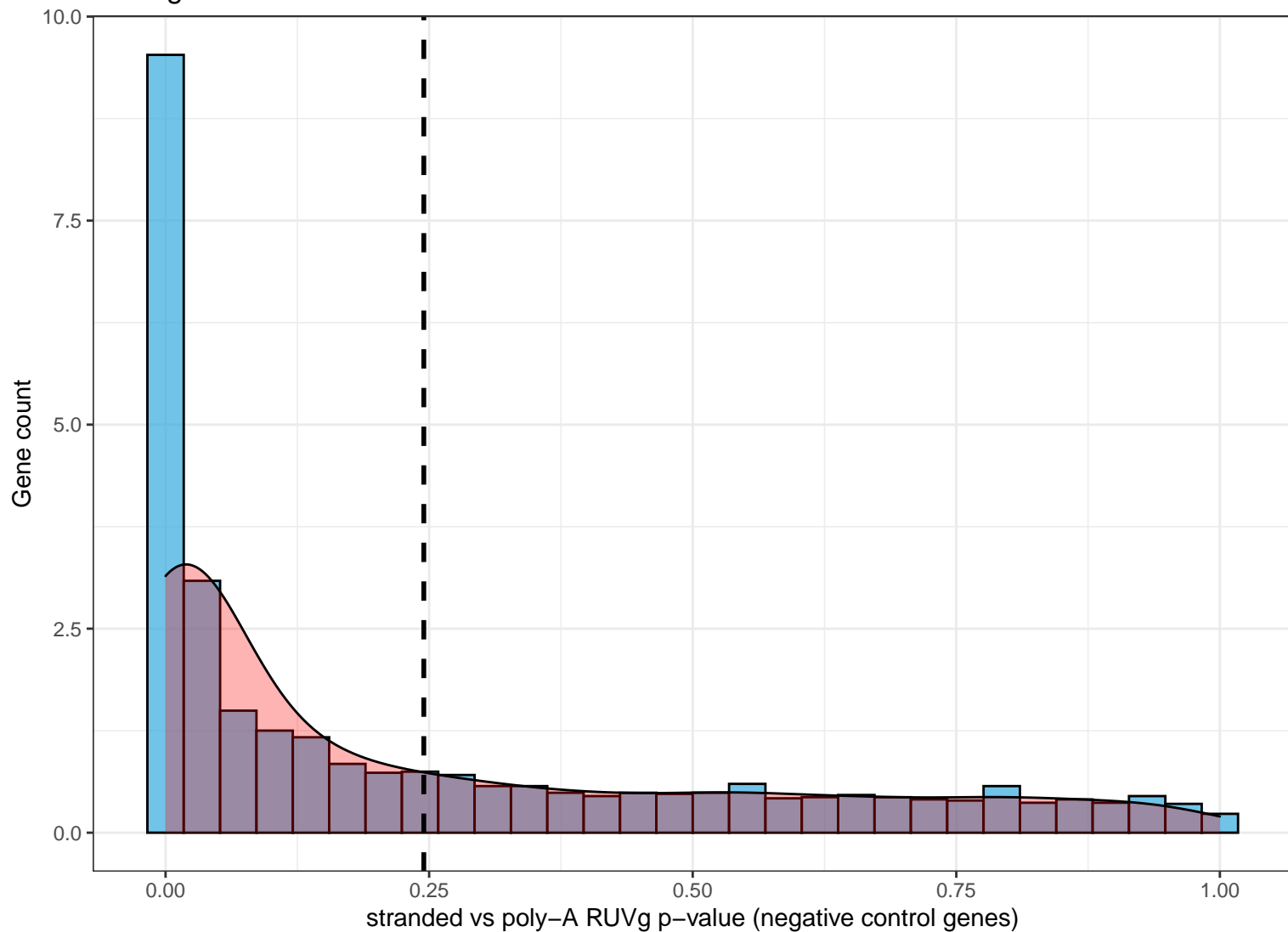
Total number of genes: 2133

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

1213 genes have $p\text{-value} \geq 0.05$

670 genes have BH FDR < 0.05

1463 genes have BH FDR ≥ 0.05



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

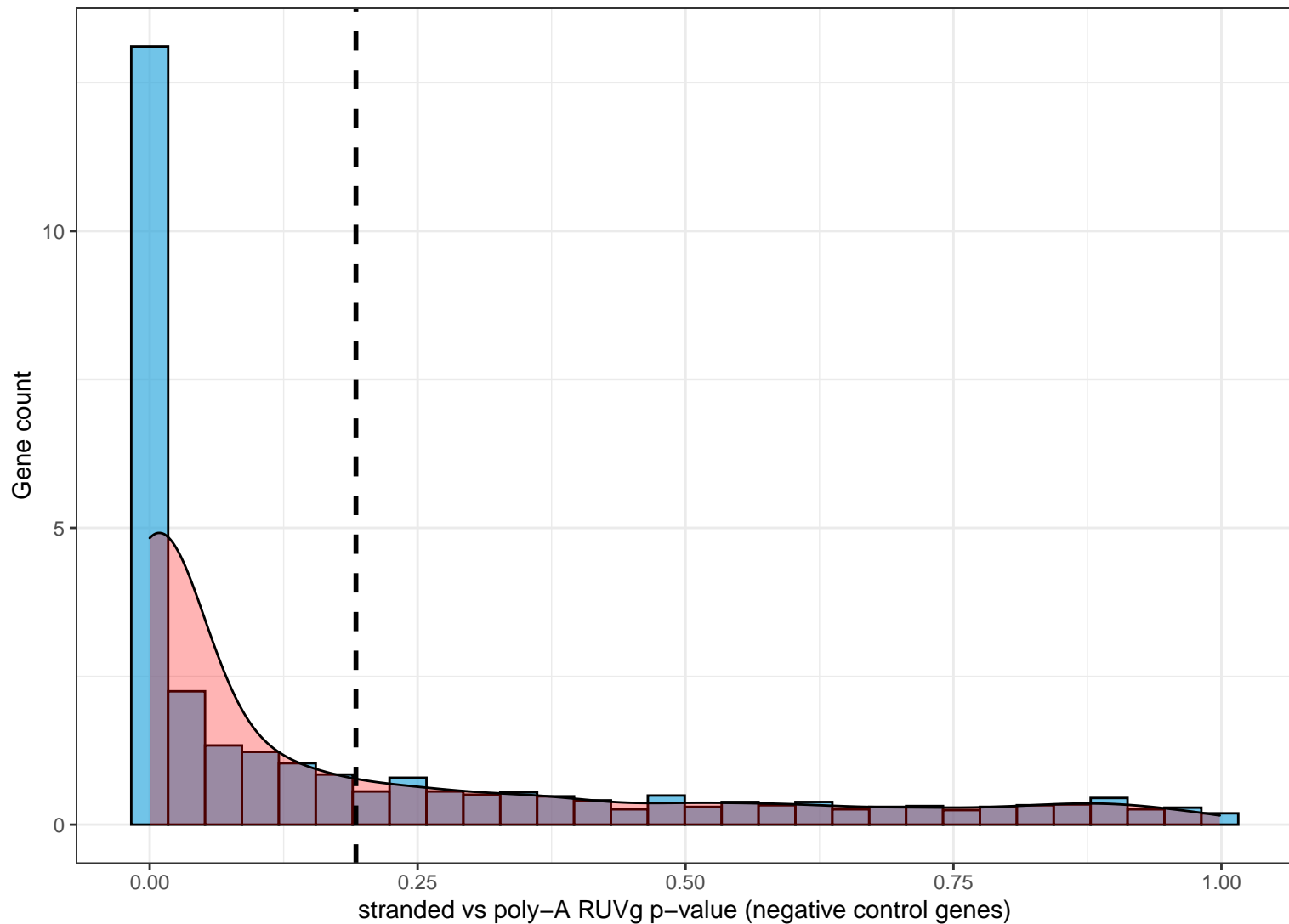
Total number of genes: 2133

1119 genes have $p\text{-value} < 0.05$

1014 genes have $p\text{-value} \geq 0.05$

988 genes have BH FDR < 0.05

1145 genes have BH FDR ≥ 0.05



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

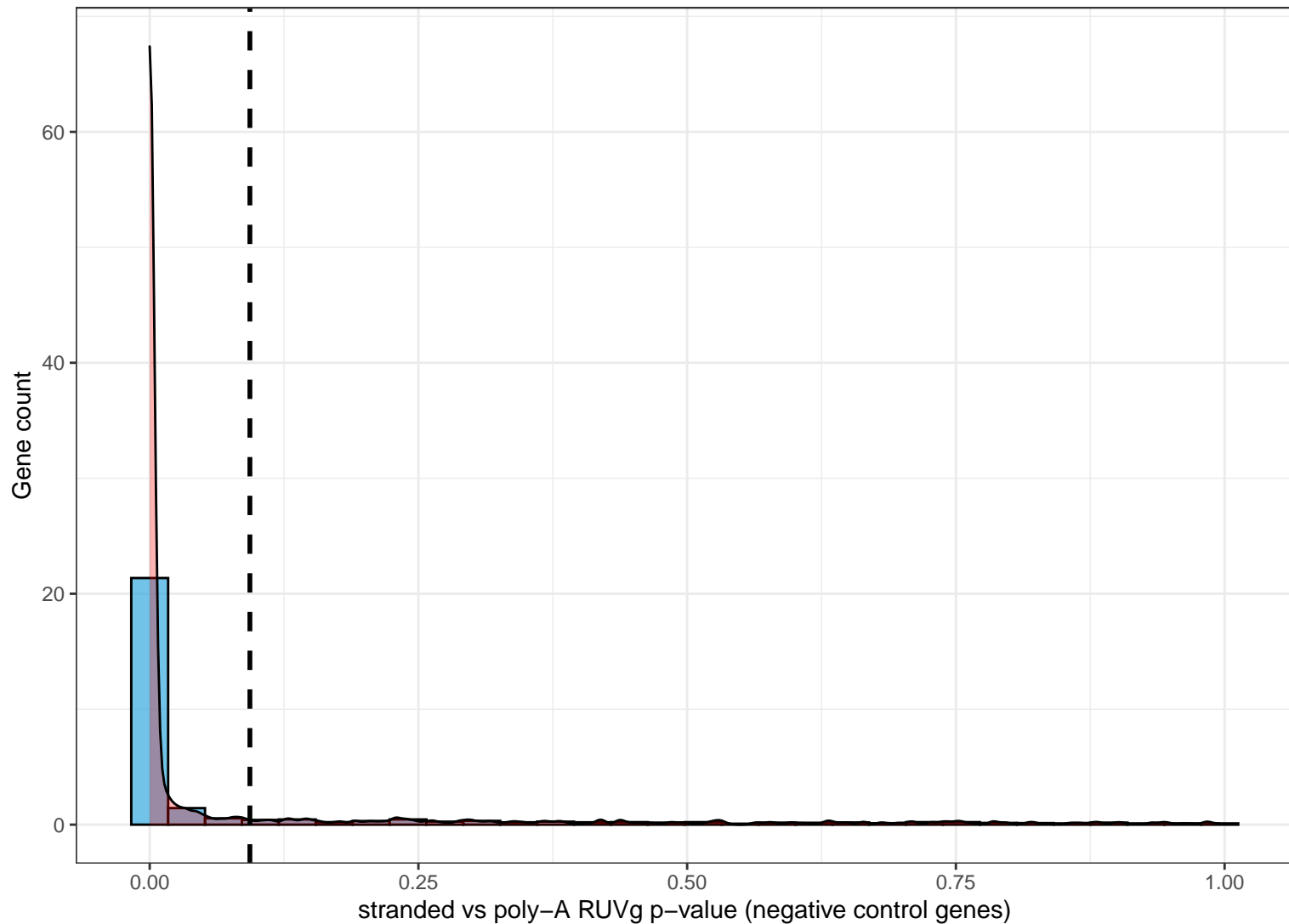
Total number of genes: 2133

1665 genes have p-value < 0.05

468 genes have p-value ≥ 0.05

1620 genes have BH FDR < 0.05

513 genes have BH FDR ≥ 0.05



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

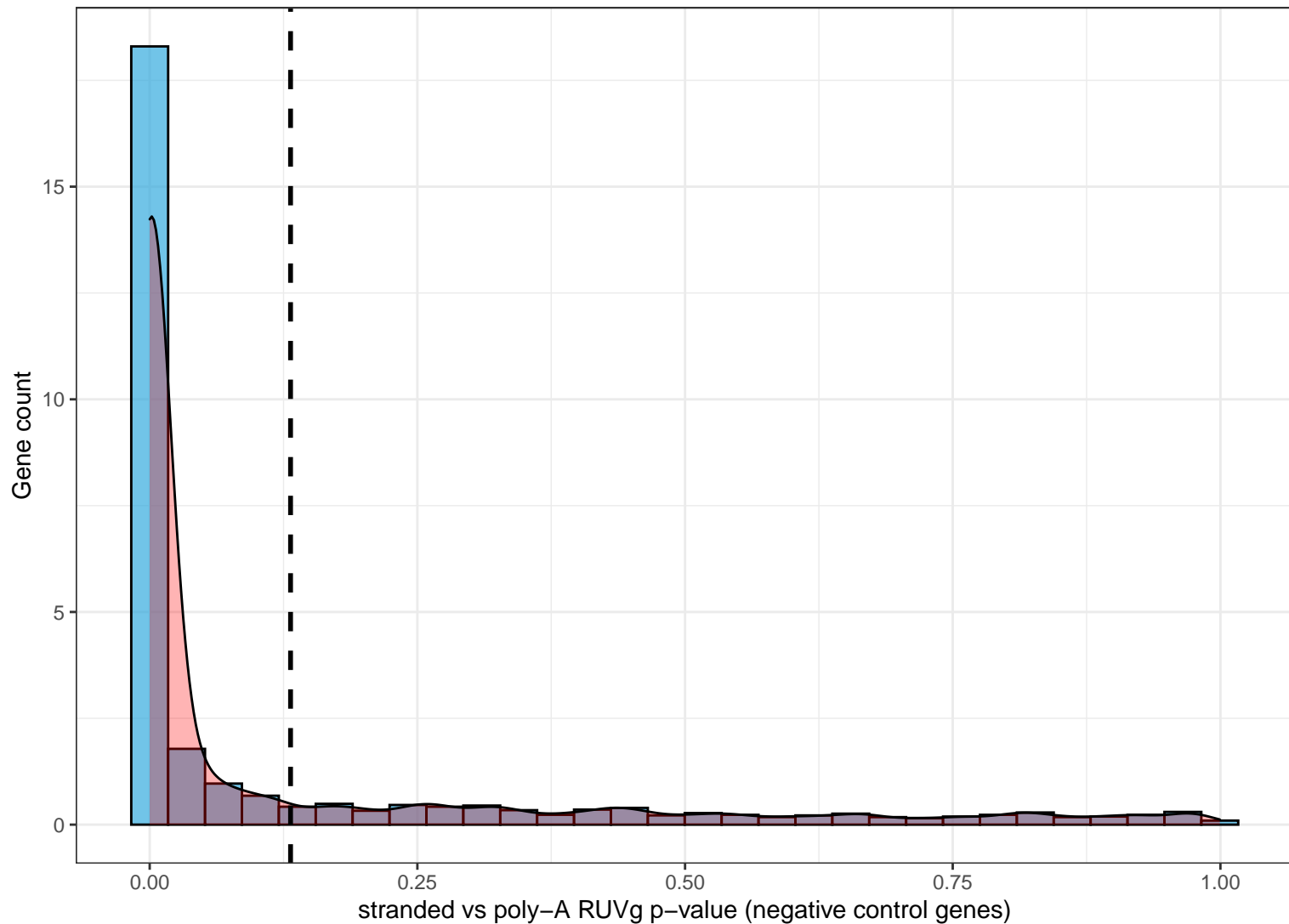
Total number of genes: 2133

1474 genes have $p\text{-value} < 0.05$

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

1403 genes have BH FDR < 0.05

730 genes have BH FDR ≥ 0.05



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

Total number of genes: 2133

1649 genes have $p\text{-value} < 0.05$

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

1616 genes have BH FDR < 0.05

517 genes have BH FDR ≥ 0.05

