

Histogram of stranded vs poly–A paired analysis (k = 2)Total number of genes: 25 0 genes have p-value < 0.05 25 genes have p-value >= 0.05 0 genes have BH FDR < 0.05 25 genes have BH FDR >= 0.05 4 3 Gene count 0

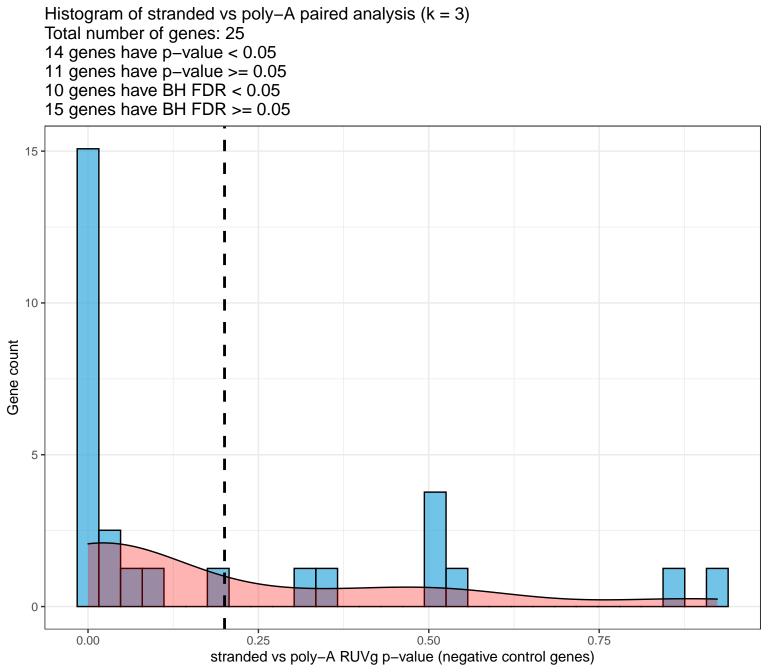
0.50

0.25

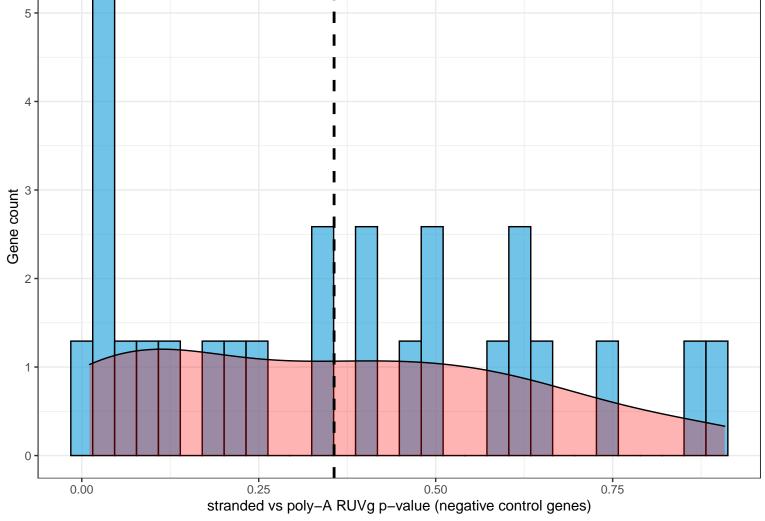
0.00

stranded vs poly–A RUVg p–value (negative control genes)

0.75



Histogram of stranded vs poly–A paired analysis (k = 4)Total number of genes: 25 5 genes have p-value < 0.05 20 genes have p-value >= 0.05 0 genes have BH FDR < 0.05 25 genes have BH FDR >= 0.05 5 4



Histogram of stranded vs poly–A paired analysis (k = 5)Total number of genes: 25 9 genes have p-value < 0.05 16 genes have p-value >= 0.05 8 genes have BH FDR < 0.05 17 genes have BH FDR >= 0.05 7.5

