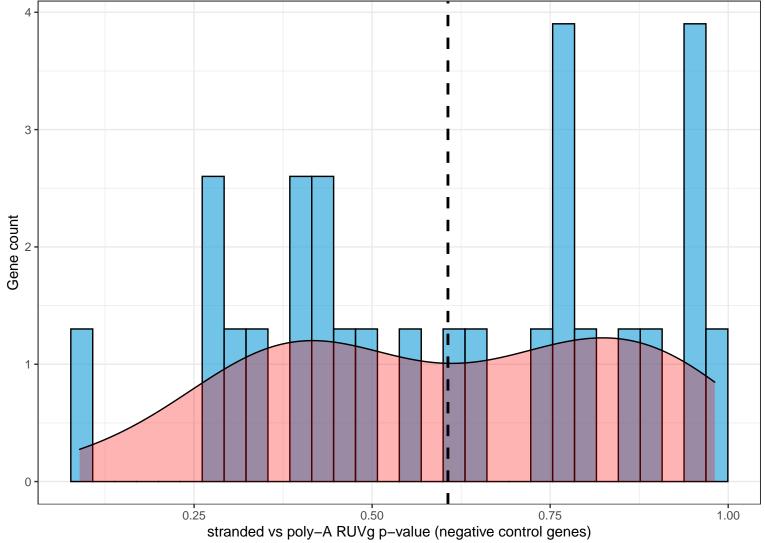
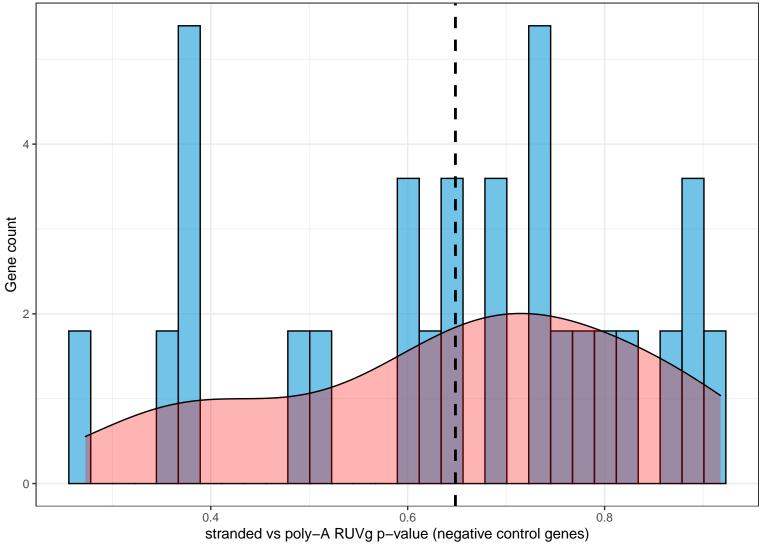
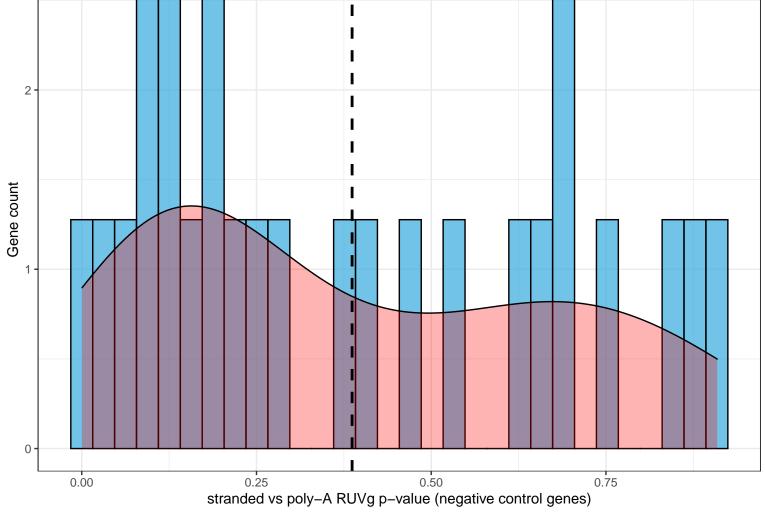
Histogram of stranded vs poly–A paired analysis (k = 1)
Total number of genes: 25
0 genes have p–value < 0.05
25 genes have p–value >= 0.05
0 genes have BH FDR < 0.05
25 genes have BH FDR >= 0.05



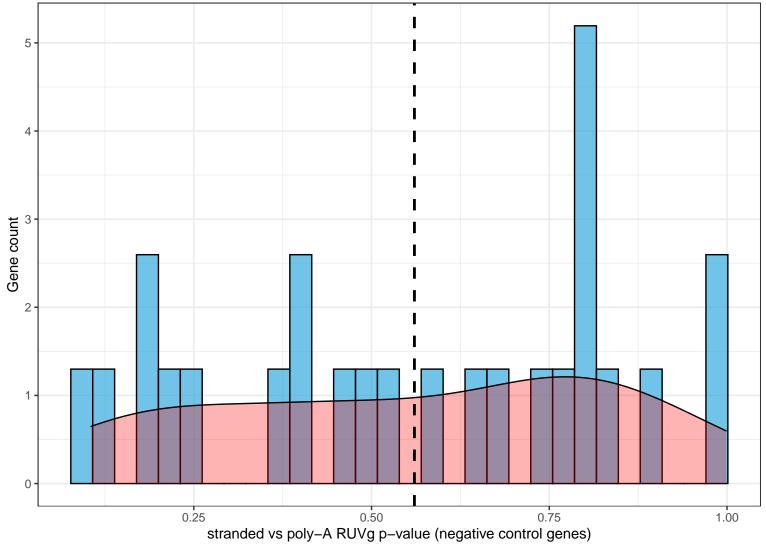
Histogram of stranded vs poly–A paired analysis (k = 2)
Total number of genes: 25
0 genes have p-value < 0.05
25 genes have p-value >= 0.05
0 genes have BH FDR < 0.05
25 genes have BH FDR >= 0.05



Histogram of stranded vs poly–A paired analysis (k = 3)Total number of genes: 25 2 genes have p-value < 0.05 23 genes have p-value >= 0.05 1 genes have BH FDR < 0.05 24 genes have BH FDR >= 0.05 2



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



Histogram of stranded vs poly–A paired analysis (k = 5)Total number of genes: 25 2 genes have p-value < 0.05 23 genes have p-value >= 0.05 1 genes have BH FDR < 0.05 24 genes have BH FDR >= 0.05 3

