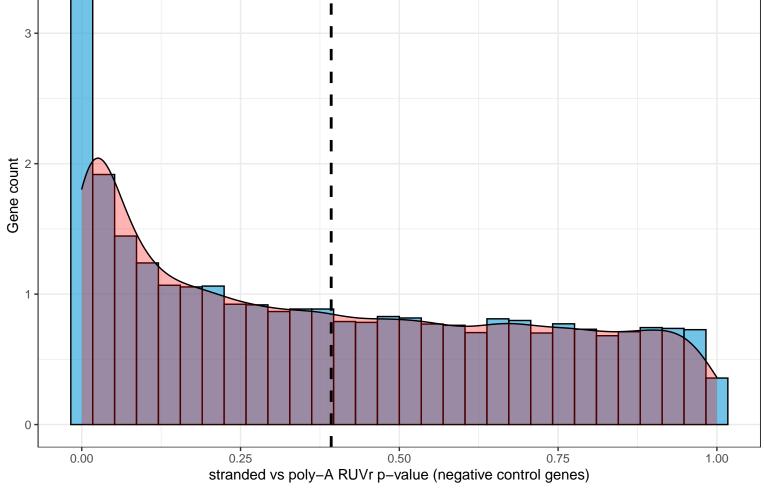
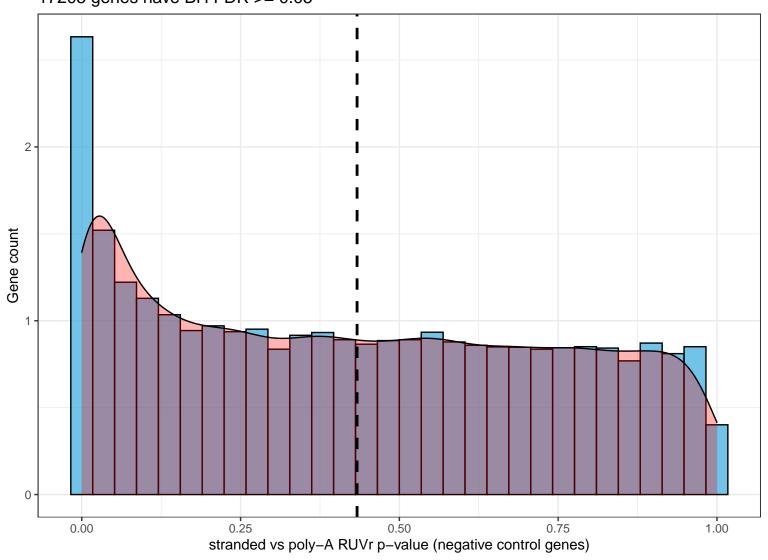
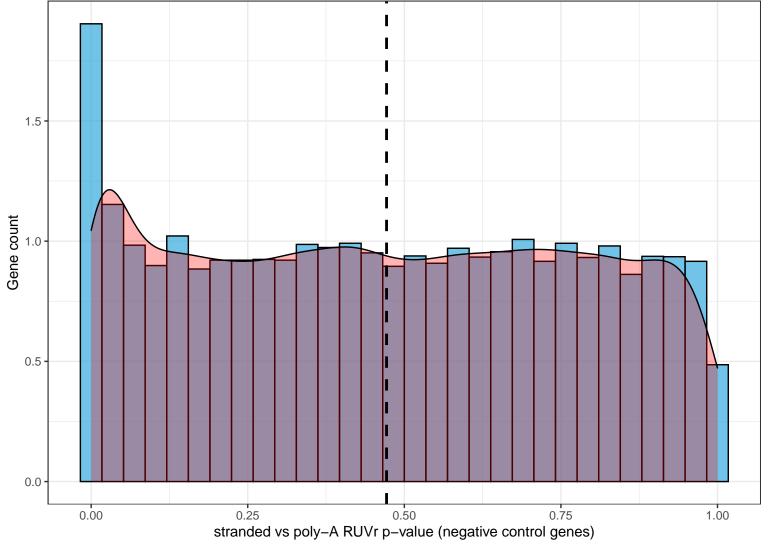
Histogram of stranded vs poly–A paired analysis (k = 1)Total number of genes: 18138 3342 genes have p-value < 0.05 14796 genes have p-value >= 0.05 1398 genes have BH FDR < 0.05 16740 genes have BH FDR >= 0.05 3 -



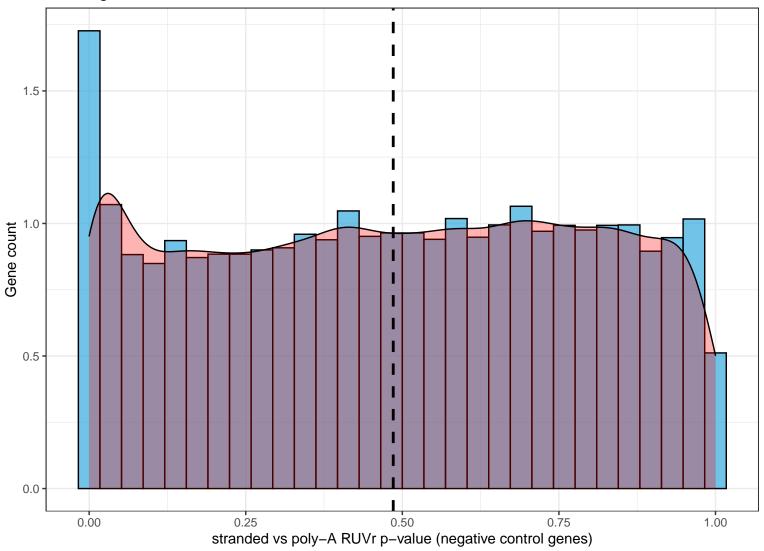
Histogram of stranded vs poly–A paired analysis (k = 2) Total number of genes: 18138 2550 genes have p-value < 0.05 15588 genes have p-value >= 0.05 933 genes have BH FDR < 0.05 17205 genes have BH FDR >= 0.05



Histogram of stranded vs poly–A paired analysis (k = 3) Total number of genes: 18138
1878 genes have p-value < 0.05
16260 genes have p-value >= 0.05
685 genes have BH FDR < 0.05
17453 genes have BH FDR >= 0.05



Histogram of stranded vs poly–A paired analysis (k = 4) Total number of genes: 18138 1720 genes have p-value < 0.05 16418 genes have p-value >= 0.05 616 genes have BH FDR < 0.05 17522 genes have BH FDR >= 0.05



Histogram of stranded vs poly–A paired analysis (k = 5) Total number of genes: 18138 1032 genes have p-value < 0.05 17106 genes have p-value >= 0.05 134 genes have BH FDR < 0.05 18004 genes have BH FDR >= 0.05

