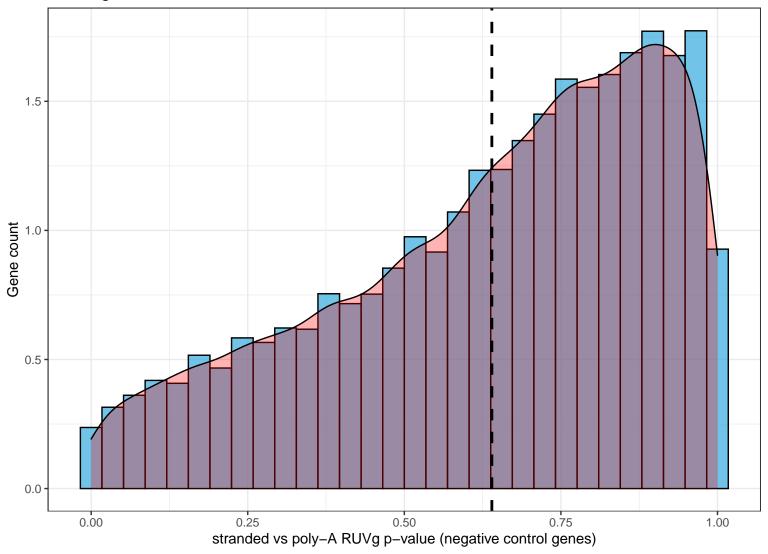
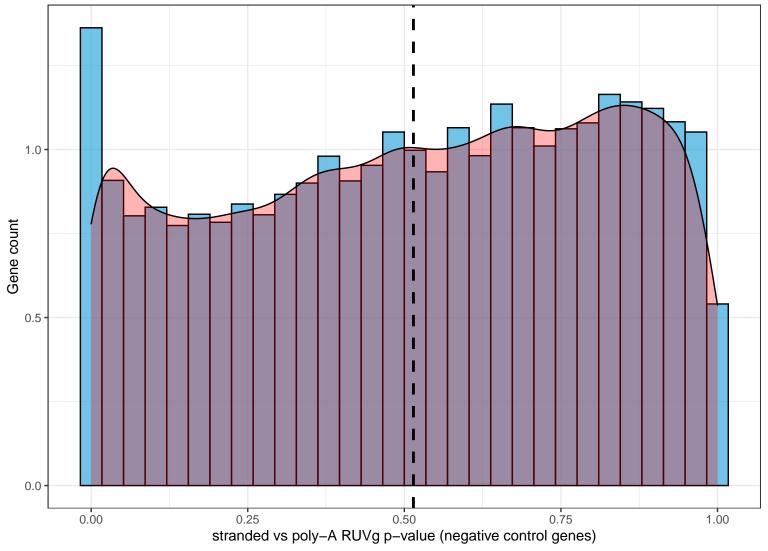
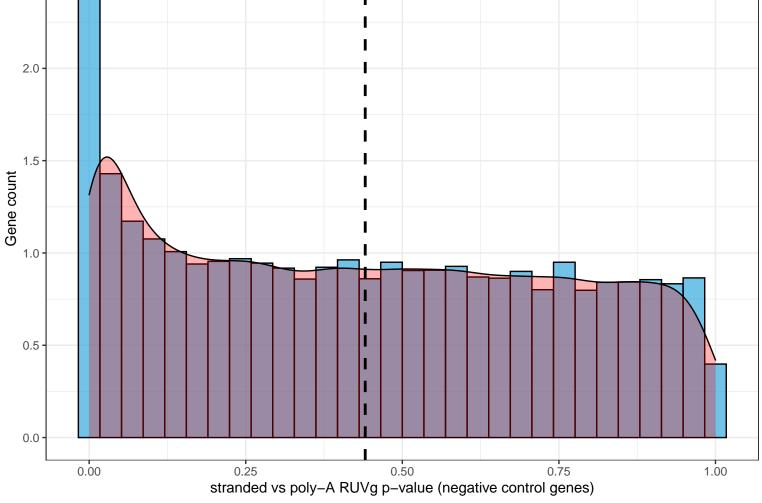
Histogram of stranded vs poly–A paired analysis (k = 1) Total number of genes: 18138 342 genes have p-value < 0.05 17796 genes have p-value >= 0.05 0 genes have BH FDR < 0.05 18138 genes have BH FDR >= 0.05



Histogram of stranded vs poly–A paired analysis (k = 2) Total number of genes: 18138 1392 genes have p-value < 0.05 16746 genes have p-value >= 0.05 332 genes have BH FDR < 0.05 17806 genes have BH FDR >= 0.05



Histogram of stranded vs poly–A paired analysis (k = 3)Total number of genes: 18138 2403 genes have p-value < 0.05 15735 genes have p-value >= 0.05 896 genes have BH FDR < 0.05 17242 genes have BH FDR >= 0.05 2.5 2.0 1.5



Histogram of stranded vs poly–A paired analysis (k = 4)
Total number of genes: 18138
2536 genes have p–value < 0.05
15602 genes have p–value >= 0.05
905 genes have BH FDR < 0.05
17233 genes have BH FDR >= 0.05

