Histogram of stranded vs poly−A paired analysis (k = 1)
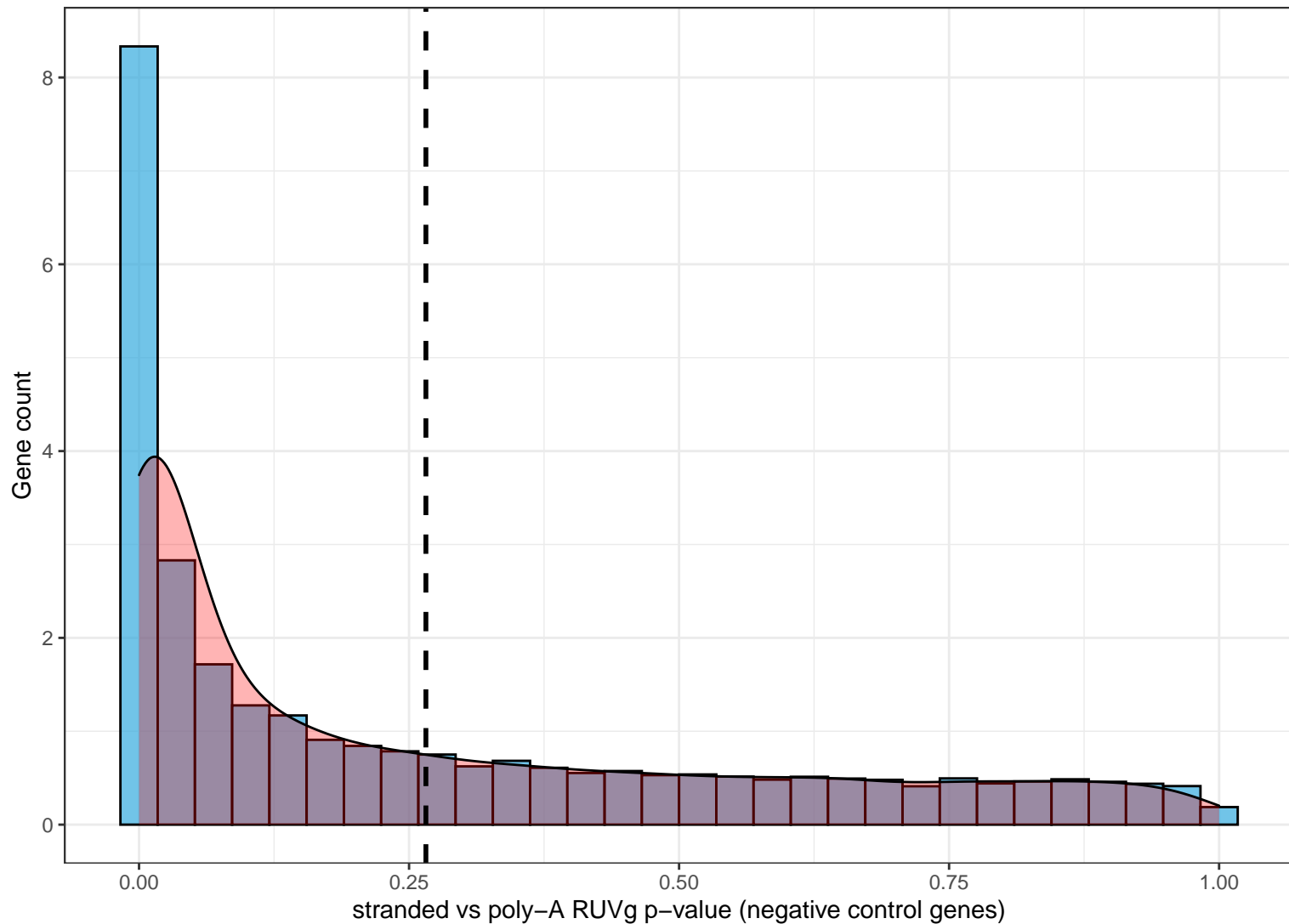Total number of genes: 15104
5755 genes have p−value < 0.05
9349 genes have p−value >= 0.05
4124 genes have BH FDR < 0.05
10980 genes have BH FDR >= 0.05

Histogram of stranded vs poly−A paired analysis (k = 2)
Total number of genes: 15104
7158 genes have p−value < 0.05
7946 genes have p−value >= 0.05
6062 genes have BH FDR < 0.05
9042 genes have BH FDR >= 0.05
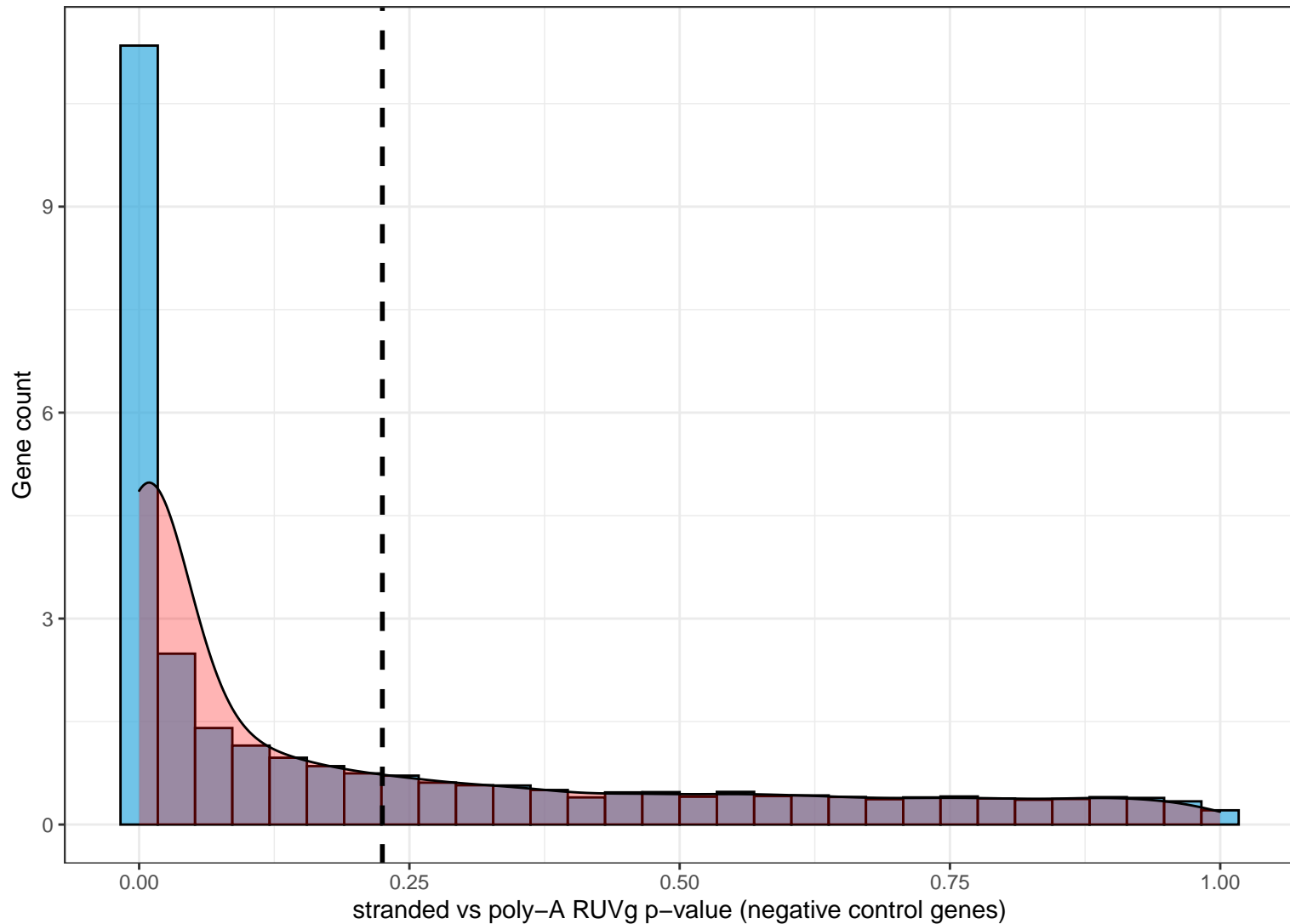
Histogram of stranded vs poly−A paired analysis (k = 3)
Total number of genes: 15104
10191 genes have p−value < 0.05
4913 genes have p−value >= 0.05
9774 genes have BH FDR < 0.05
5330 genes have BH FDR >= 0.05
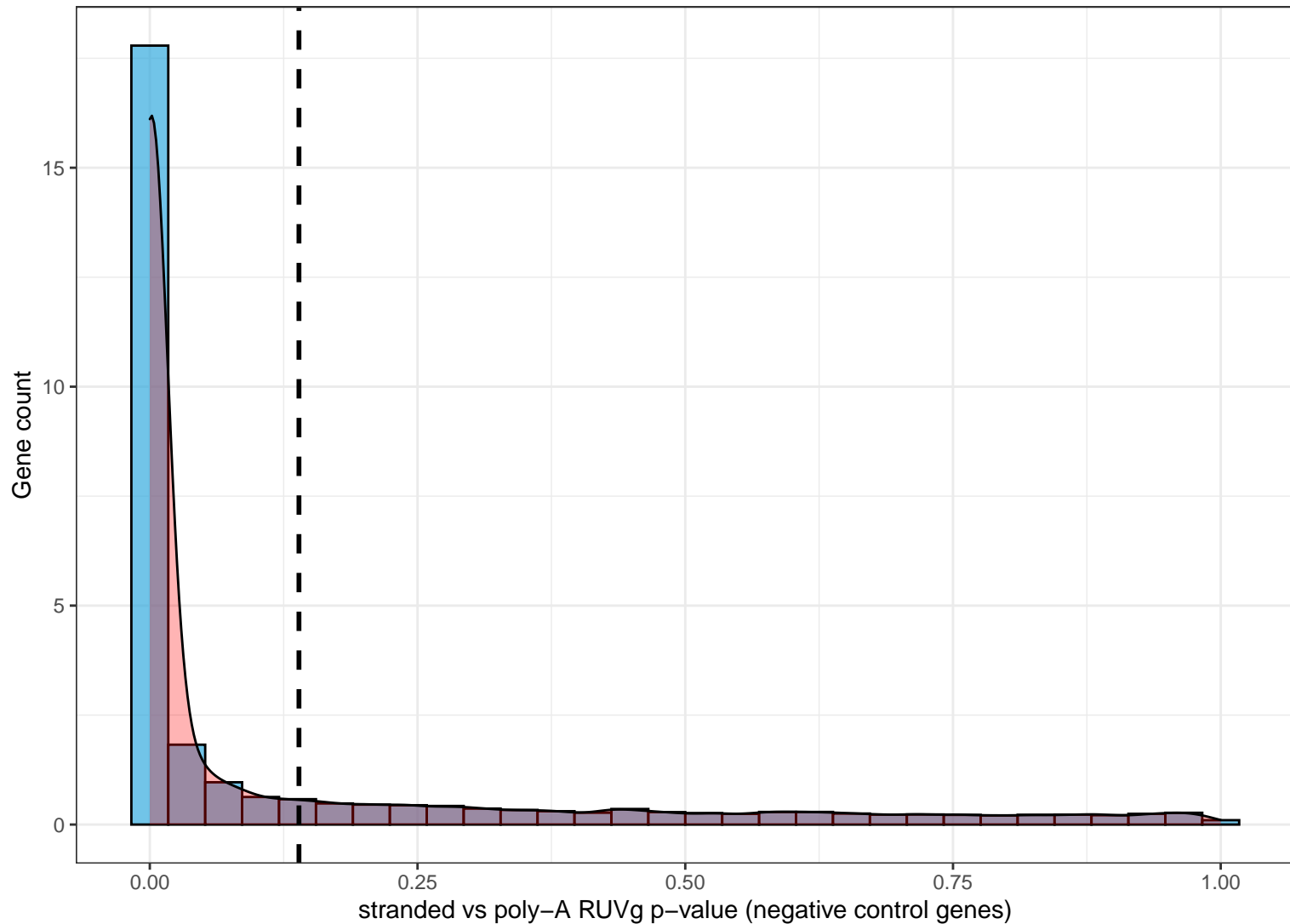
Histogram of stranded vs poly−A paired analysis (k = 4)
Total number of genes: 15104
9357 genes have p−value < 0.05
5747 genes have p−value >= 0.05
8765 genes have BH FDR < 0.05
6339 genes have BH FDR >= 0.05
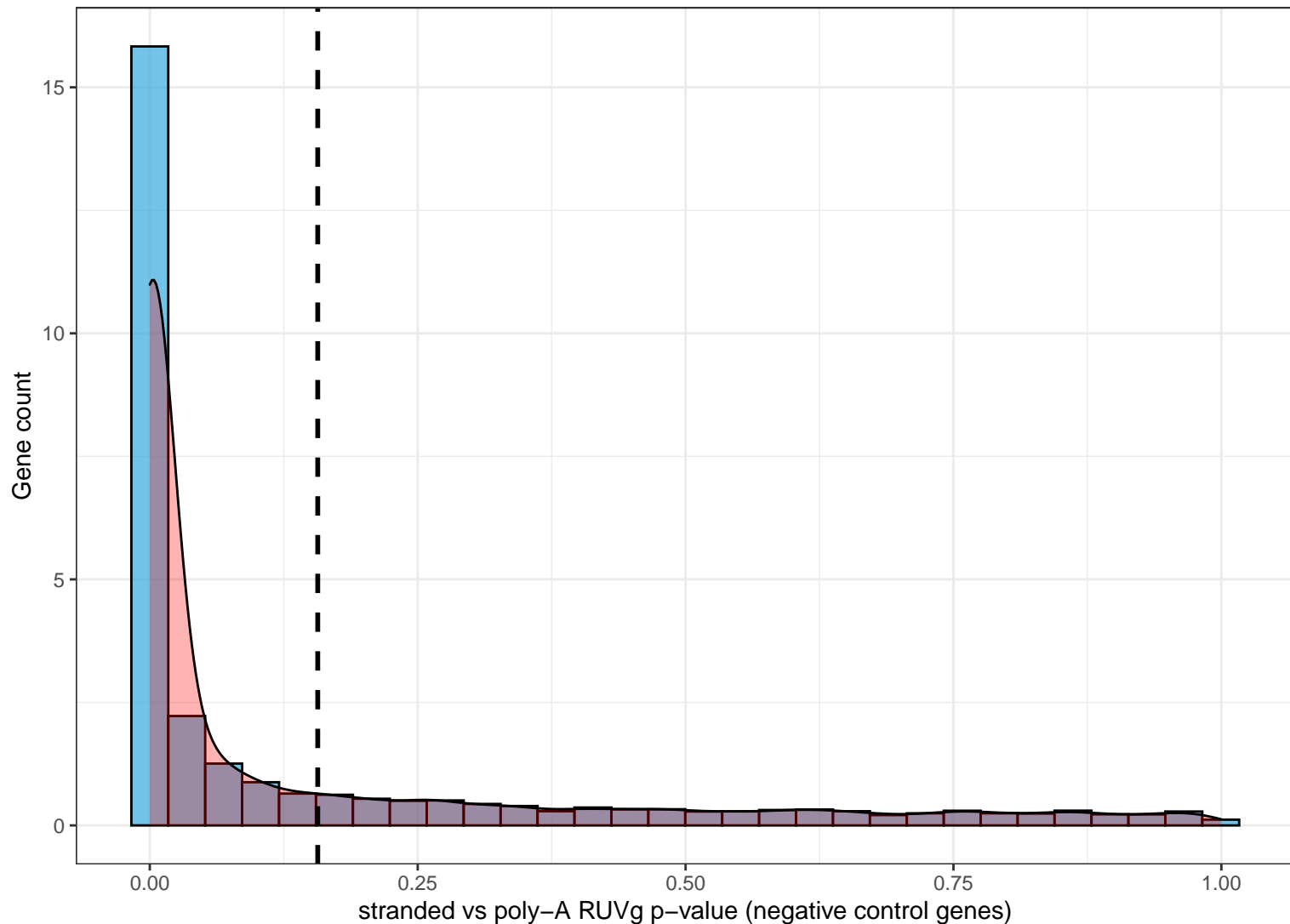
Histogram of stranded vs poly−A paired analysis (k = 5)
Total number of genes: 15104
10884 genes have p−value < 0.05
4220 genes have p−value >= 0.05
10575 genes have BH FDR < 0.05
4529 genes have BH FDR >= 0.05