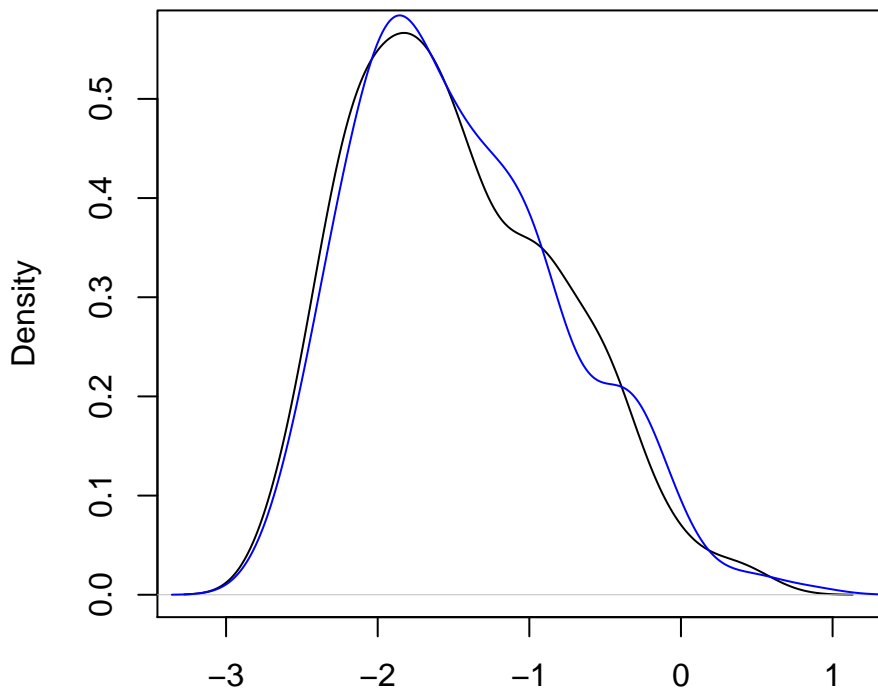


```
.default(x = log10(mySample(subset(dat_means, gene_
protein_coding"), n = 339, EmpDist = learned.Inc.den)$m
```



N = 339 Bandwidth = 0.1922