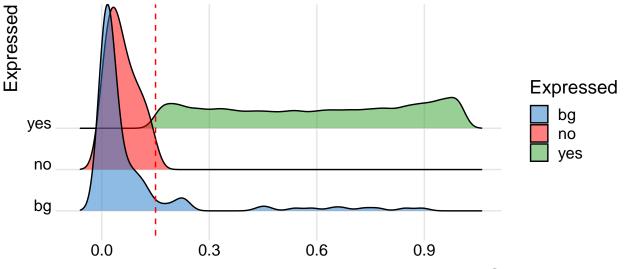
## **Gene detection distribution**

Threshold=0.15 Expr=12451 Filtered=6225 Recall=0.67



Frac detectable above BG

