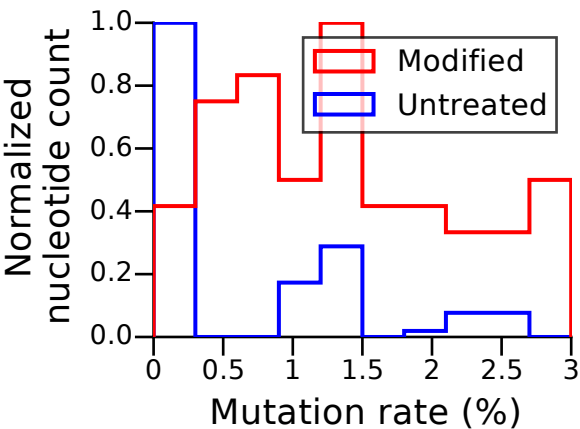


# RNA: SeC-TCA-1-1

Note: possible data quality issue - see log file

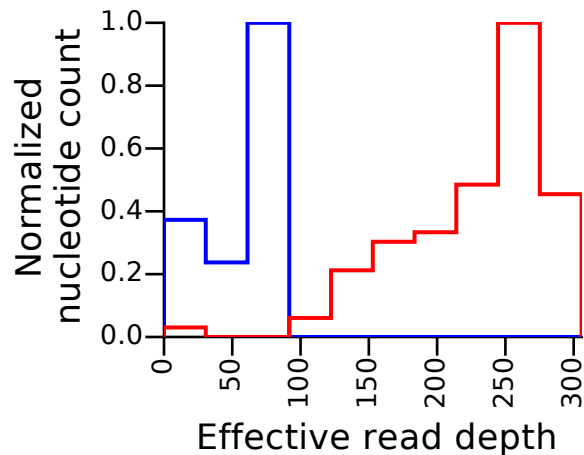
## Mutation rates



Modified sample:  
95th percentile rate: 6.68%  
Median rate: 1.81%

Untreated sample:  
95th percentile rate: 3.33%  
Median rate: 0%

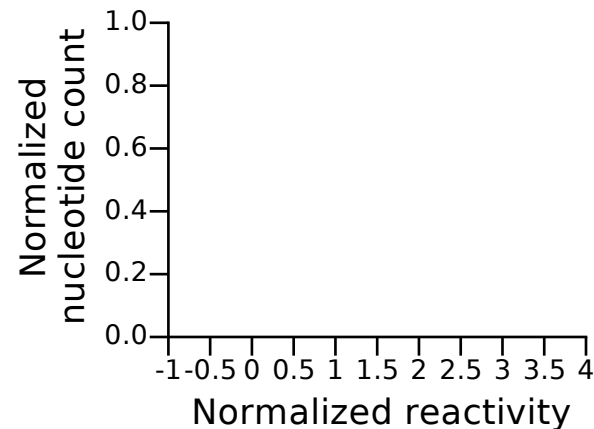
## Read depths



Modified sample:  
Median depth: 245  
5th percentile depth: 135

Untreated sample:  
Median depth: 75  
5th percentile depth: 24

## Reactivity distribution



## Raw background-corrected rate distribution

