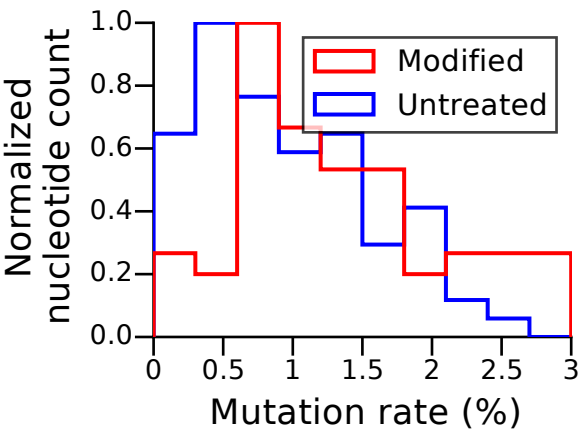


# RNA: Leu-CAG-2-1

Note: possible data quality issue - see log file

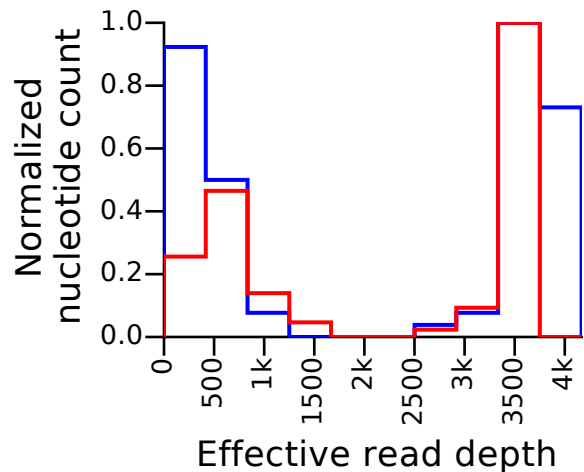
## Mutation rates



Modified sample:  
 95th percentile rate: 10.45%  
 Median rate: 1.6%

Untreated sample:  
 95th percentile rate: 3.93%  
 Median rate: 0.95%

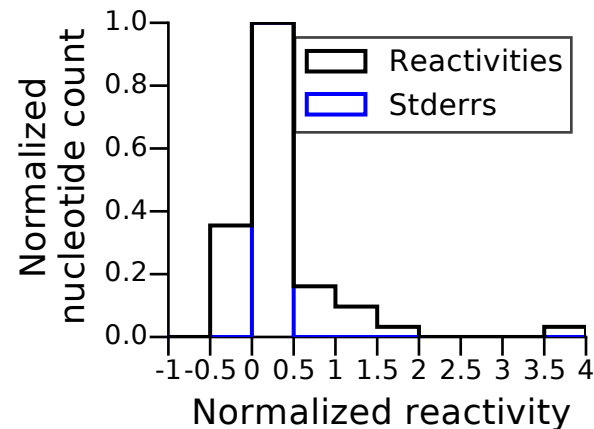
## Read depths



Modified sample:  
 Median depth: 3,271  
 5th percentile depth: 256

Untreated sample:  
 Median depth: 3,527  
 5th percentile depth: 178

## Reactivity distribution



## Raw background-corrected rate distribution

