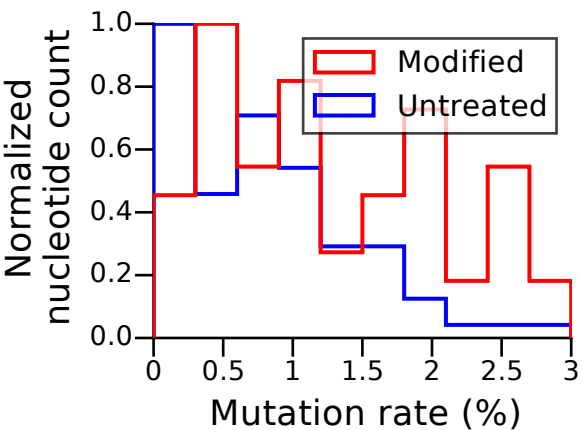


# RNA: Ser-CGA-1-1

Note: possible data quality issue - see log file

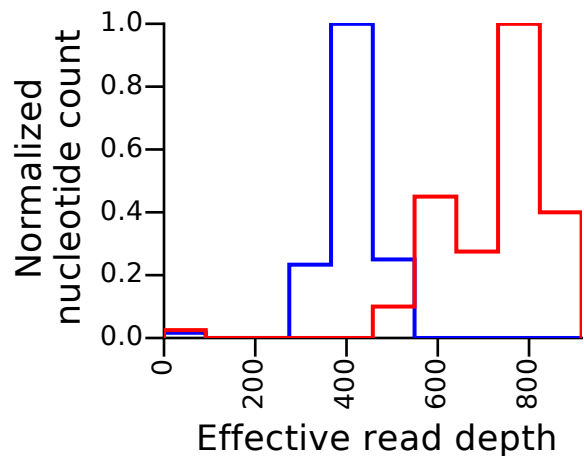
## Mutation rates



Modified sample:  
95th percentile rate: 8.78%  
Median rate: 2.06%

Untreated sample:  
95th percentile rate: 2.61%  
Median rate: 0.74%

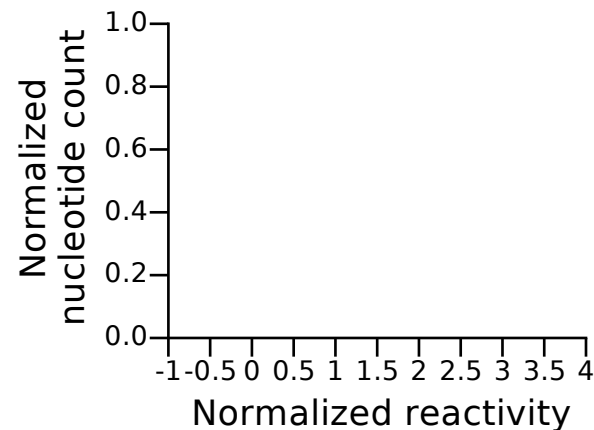
## Read depths



Modified sample:  
Median depth: 766  
5th percentile depth: 548

Untreated sample:  
Median depth: 422  
5th percentile depth: 341

## Reactivity distribution



## Raw background-corrected rate distribution

