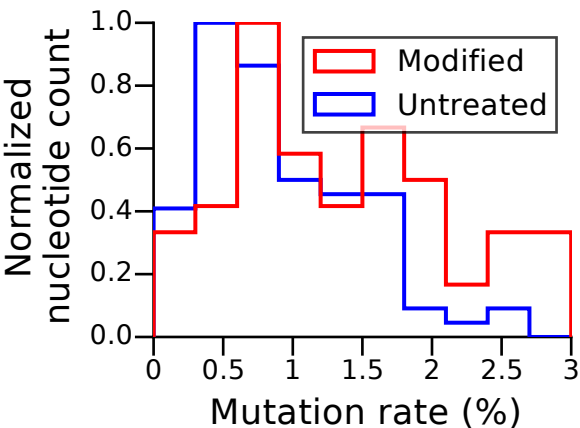


# RNA: Ser-CGA-1-1

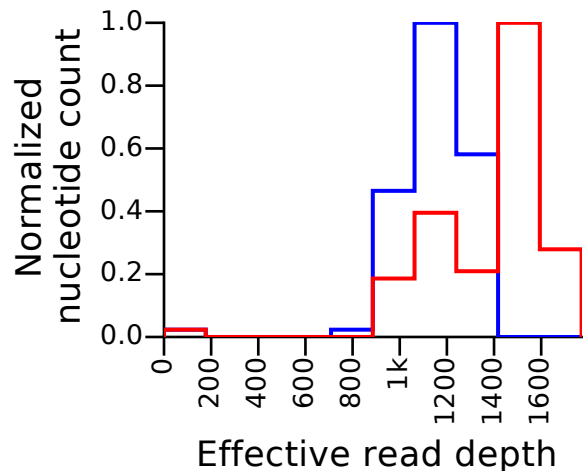
## Mutation rates



Modified sample:  
95th percentile rate: 8.21%  
Median rate: 1.99%

Untreated sample:  
95th percentile rate: 2.42%  
Median rate: 0.8%

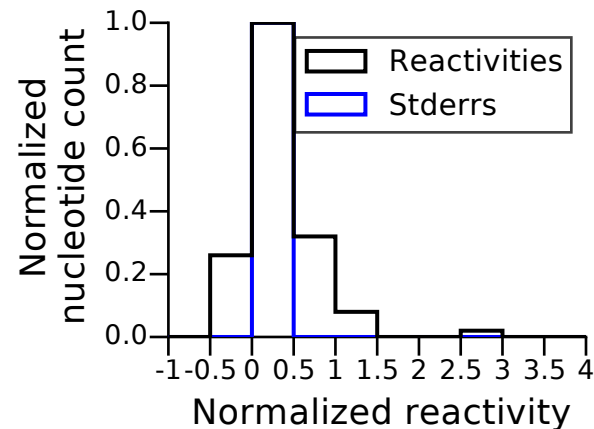
## Read depths



Modified sample:  
Median depth: 1,475  
5th percentile depth: 987

Untreated sample:  
Median depth: 1,179  
5th percentile depth: 965

## Reactivity distribution



## Raw background-corrected rate distribution

