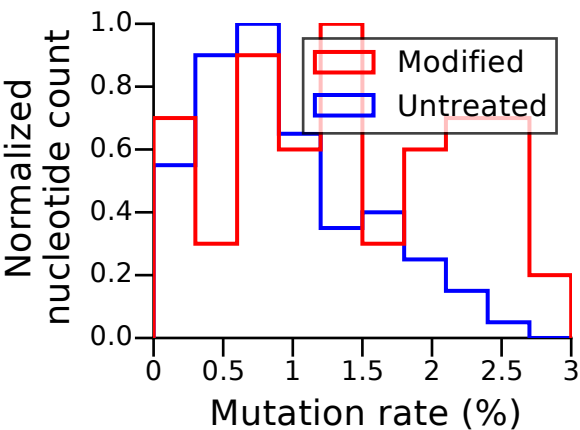


# RNA: Ser-CGA-1-1

Note: possible data quality issue - see log file

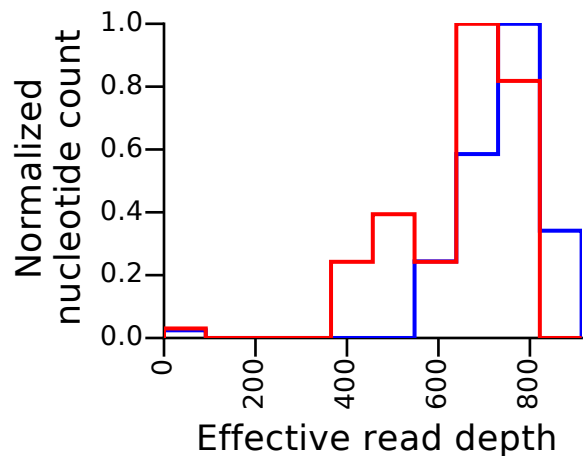
## Mutation rates



Modified sample:  
95th percentile rate: 8.58%  
Median rate: 2.11%

Untreated sample:  
95th percentile rate: 2.33%  
Median rate: 0.82%

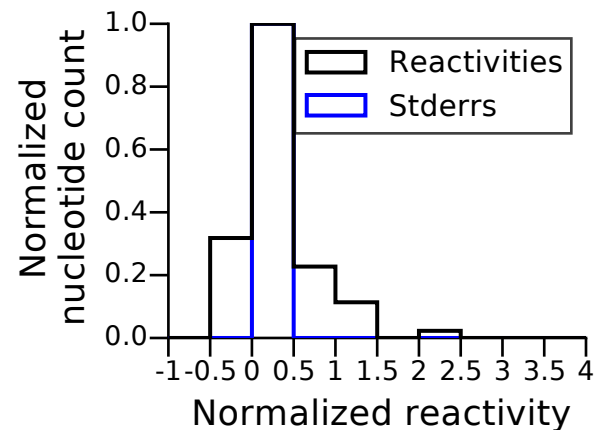
## Read depths



Modified sample:  
Median depth: 706  
5th percentile depth: 439

Untreated sample:  
Median depth: 754  
5th percentile depth: 619

## Reactivity distribution



## Raw background-corrected rate distribution

