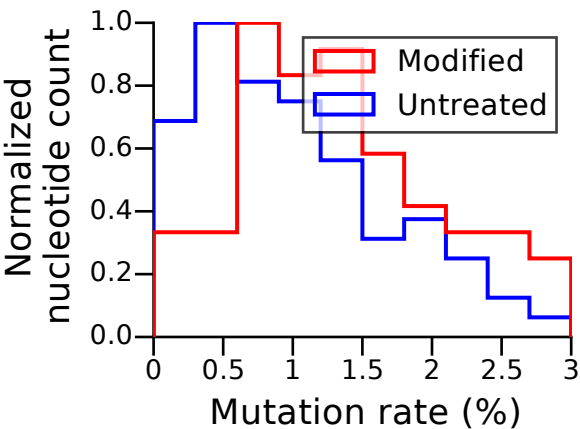


RNA: Leu-CAG-2-1

Note: possible data quality issue - see log file

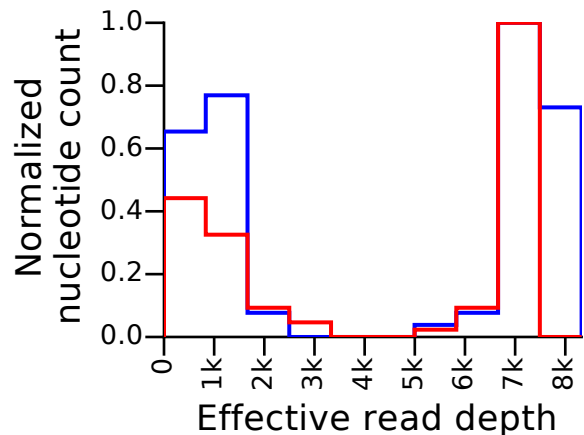
Mutation rates



Modified sample:
95th percentile rate: 11.44%
Median rate: 1.62%

Untreated sample:
95th percentile rate: 4.29%
Median rate: 0.96%

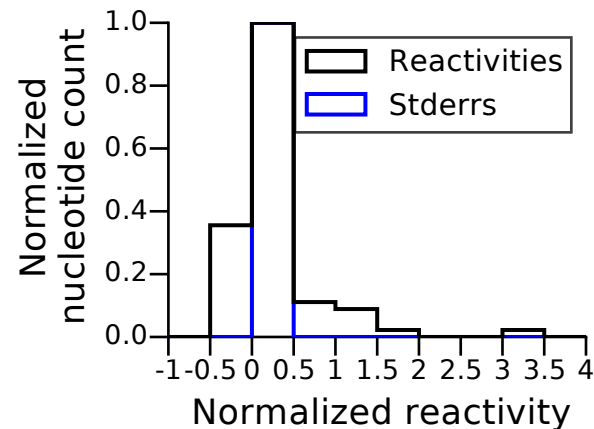
Read depths



Modified sample:
Median depth: 6,459
5th percentile depth: 517

Untreated sample:
Median depth: 7,019
5th percentile depth: 345

Reactivity distribution



Raw background-corrected rate distribution

