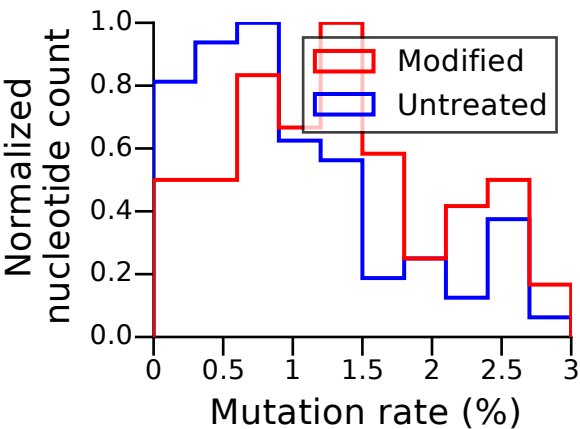


RNA: Leu-CAG-2-1

Note: possible data quality issue - see log file

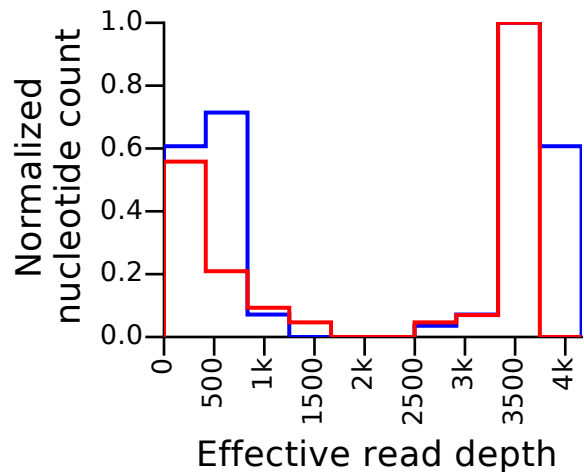
Mutation rates



Modified sample:
95th percentile rate: 11.3%
Median rate: 1.59%

Untreated sample:
95th percentile rate: 4.65%
Median rate: 0.9%

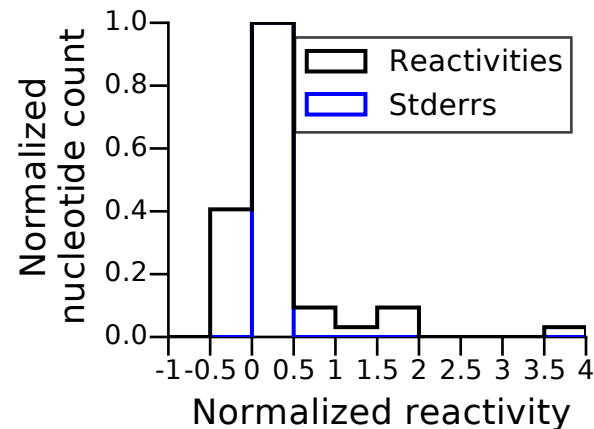
Read depths



Modified sample:
Median depth: 3,189
5th percentile depth: 238

Untreated sample:
Median depth: 3,492
5th percentile depth: 166

Reactivity distribution



Raw background-corrected rate distribution

