Note: possible data quality issue - see log file RNA: Ile2-CAT-2-1 Mutation rates Read depths Reactivity distribution nucleotide count count count Modified 0.8-0.8-0.8 Normalized Normalized Normalized Untreated 0.6 -0.6 -0.6nucleotide nucleotide 0.4 -0.4-0.4 0.2-0.2-0.2 0.0 0.0 -0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 -1-0.5 0 0.5 1 1.5 2 2.5 3 3.5 4 Effective read depth Mutation rate (%) Normalized reactivity Raw background-corrected rate distribution count Modified sample: Modified sample: 0.8 95th percentile rate: 0% Median depth: Normalized Median rate: 0% 5th percentile depth: 0.6 nucleotide Untreated sample: Untreated sample: 0.4 95th percentile rate: 0% Median depth: 1 Median rate: 0% 5th percentile depth: 0.2 0.0 0 In(modified rate / untreated rate)