Note: possible data quality issue - see log file RNA: Ile2-CAT-1-1 Mutation rates Read depths Reactivity distribution count nucleotide count nucleotide count Modified 0.8-0.8-0.8 Normalized Normalized Normalized Untreated 0.6 -0.6nucleotide 0.6-0.4-0.4-0.4 0.2-0.2-0.2-0.0-0.0-0.0 100 -1-0.5 0 0.5 1 1.5 2 2.5 3 3.5 4 50 50 Mutation rate (%) Normalized reactivity Effective read depth Raw background-corrected rate distribution count Modified sample: Modified sample: 0.8 95th percentile rate: 10.87% Median depth: 103 Normalized 0.62% 5th percentile depth: 10 Median rate: 0.6 nucleotide Untreated sample: Untreated sample: 0.4 95th percentile rate: Median depth: 4.5% 123 19 Median rate: 0% 5th percentile depth: 0.2