Synthetic Genomics Data Generation and Evaluation for the Use Case of Benchmarking Somatic Variant Calling Algorithms

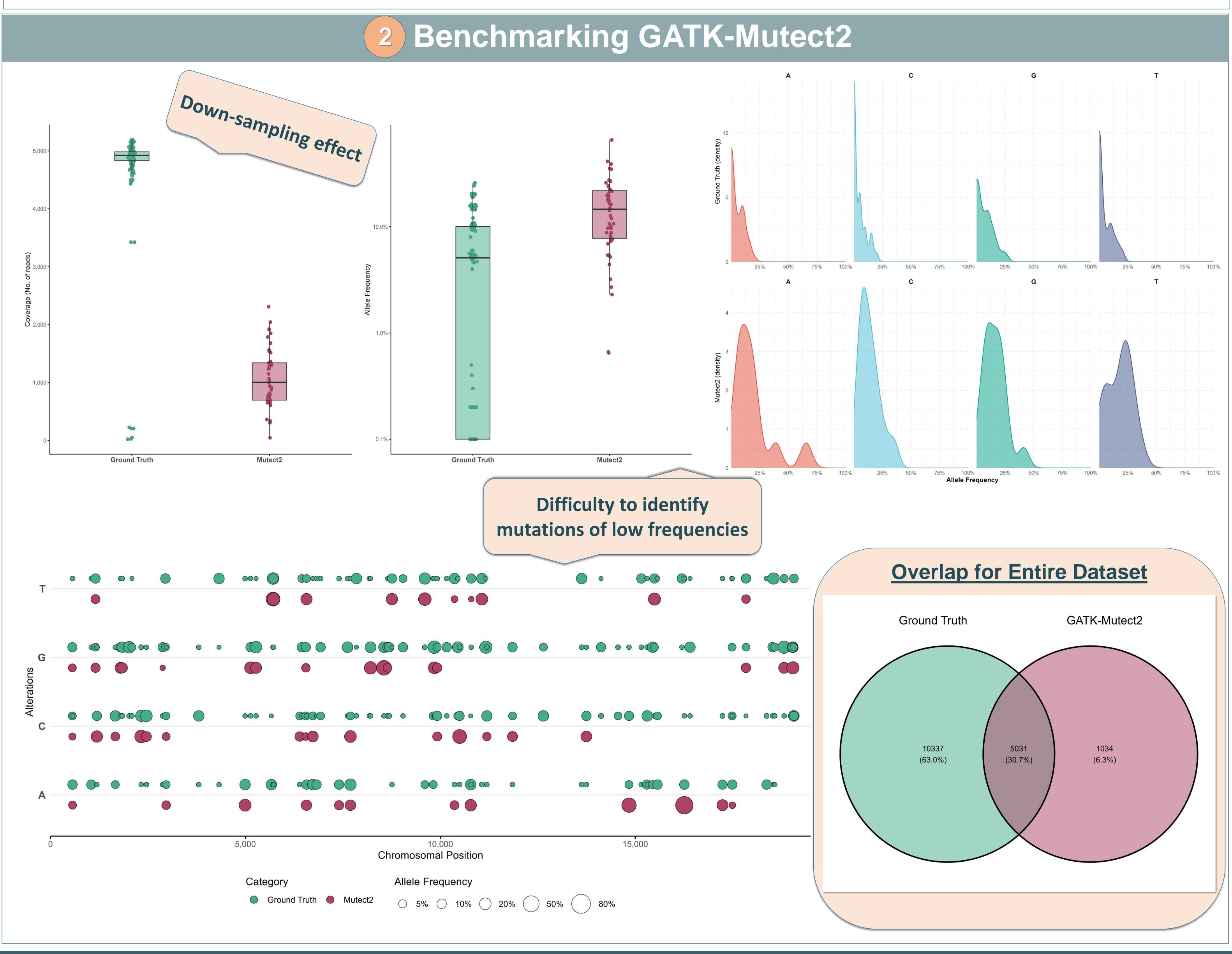
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1 Synthetic «Gold Standard» Dataset Generation NEAT To learn more AO individual barn files about our work please visit golden.bam golden.vcf.gz read1.fq.gz read2.fq.gz Ground Merge bam files 1.bam x5000 coverage x500 coverage

Gold Standard Mutations 100% Allele Frequency

Gold Standard Mutations 10% Allele Frequency











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