

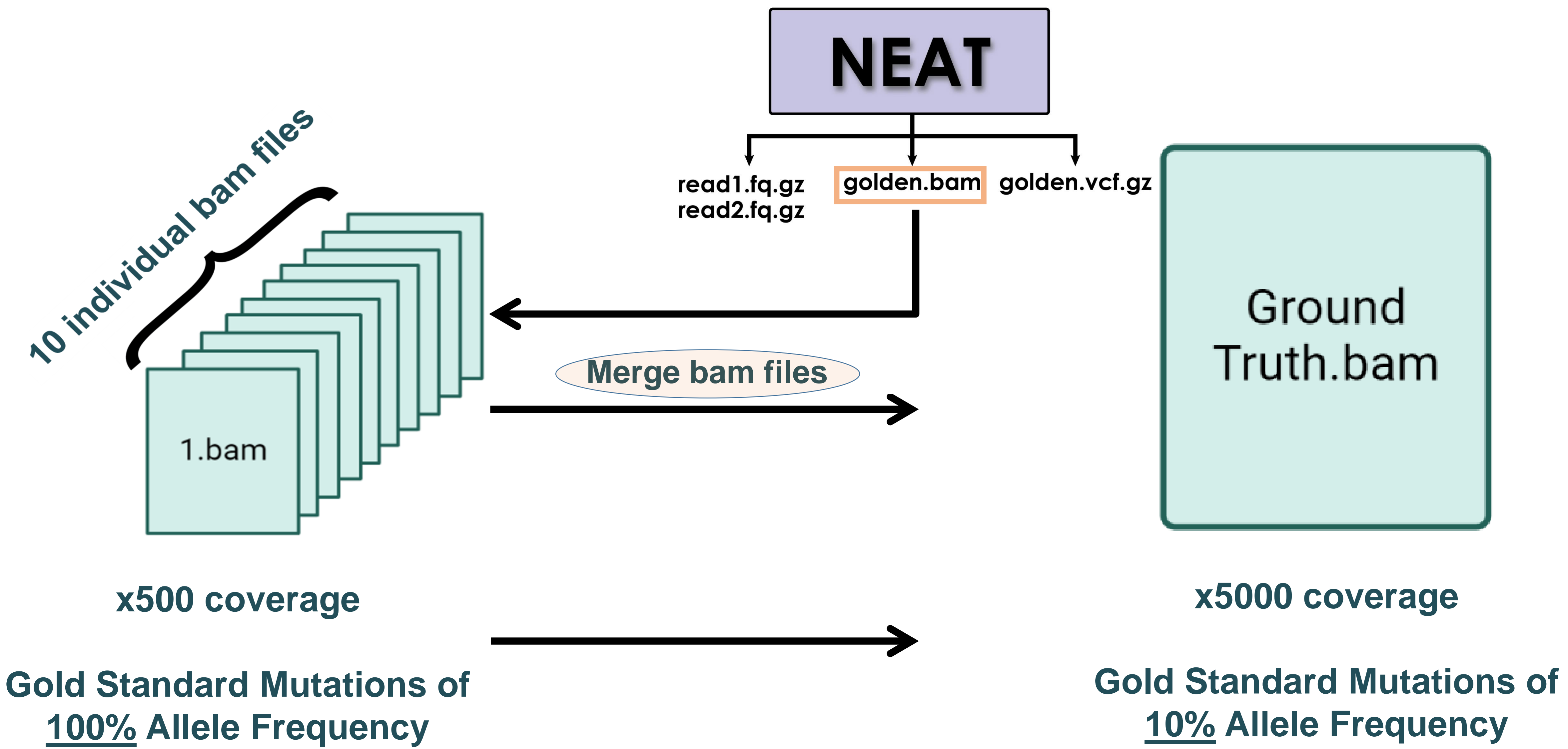
Synthetic Genomics Data Generation and Evaluation for the Use Case of Benchmarking Somatic Variant Calling Algorithms

Fragkouli Styliani-Christina^{1, 2}, Pechlivanis Nikos¹, Agathangelidis Andreas², Psomopoulos Fotis¹

¹Institute of Applied Biosciences, Centre of Research and Technology Hellas, Thessaloniki, Greece
²Department of Biology, National and Kapodistrian University of Athens, Athens 10679, GR

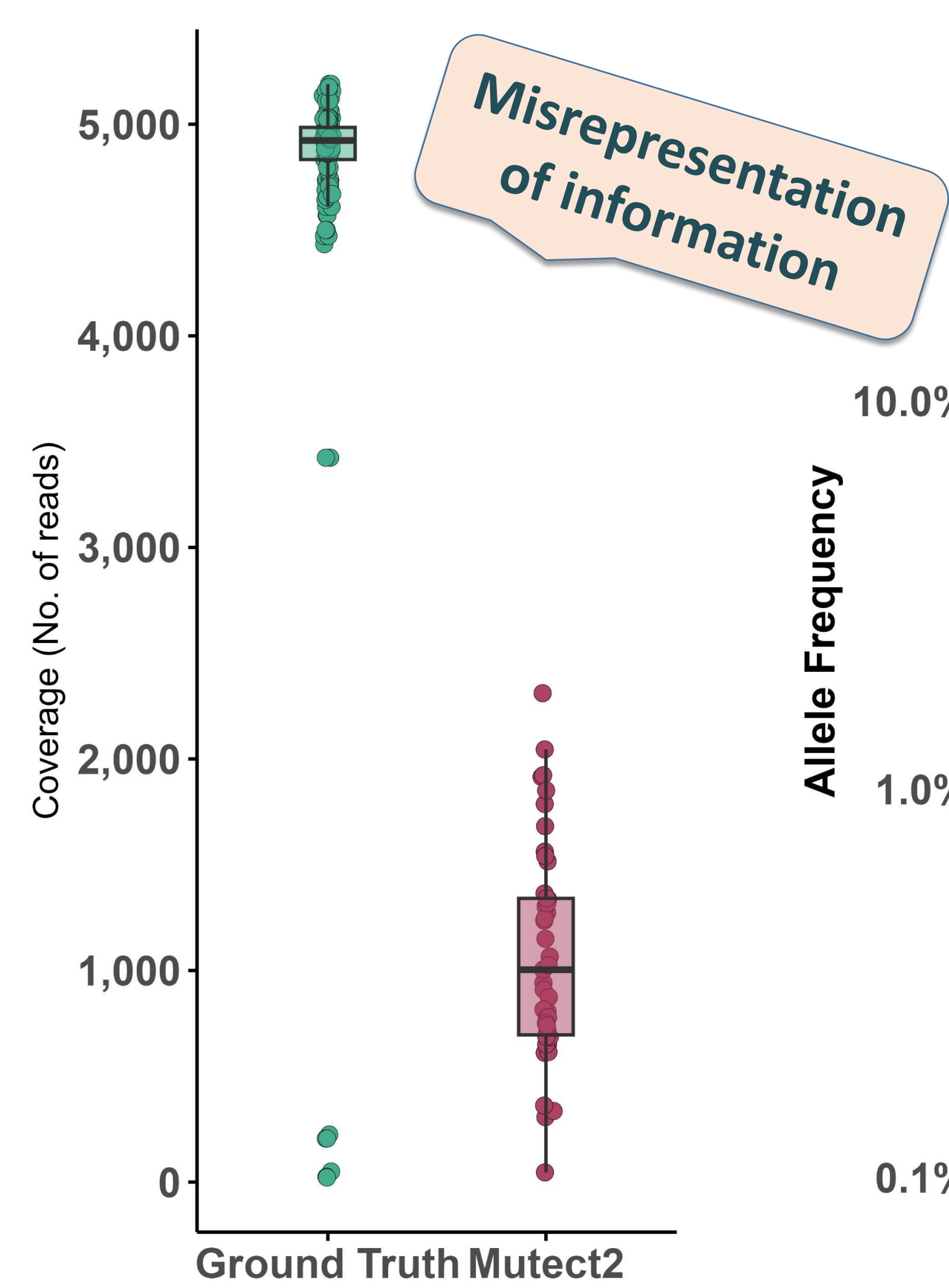
1 Synthetic «Gold Standard» Dataset Generation

- Highlights
- Generation of **synthetic genomics** data based on TP53 gene
 - Define «**Ground Truth**» mutations in order to **benchmark** somatic variant callers
 - Investigate the impact of variant callers in mutations at **low frequencies**

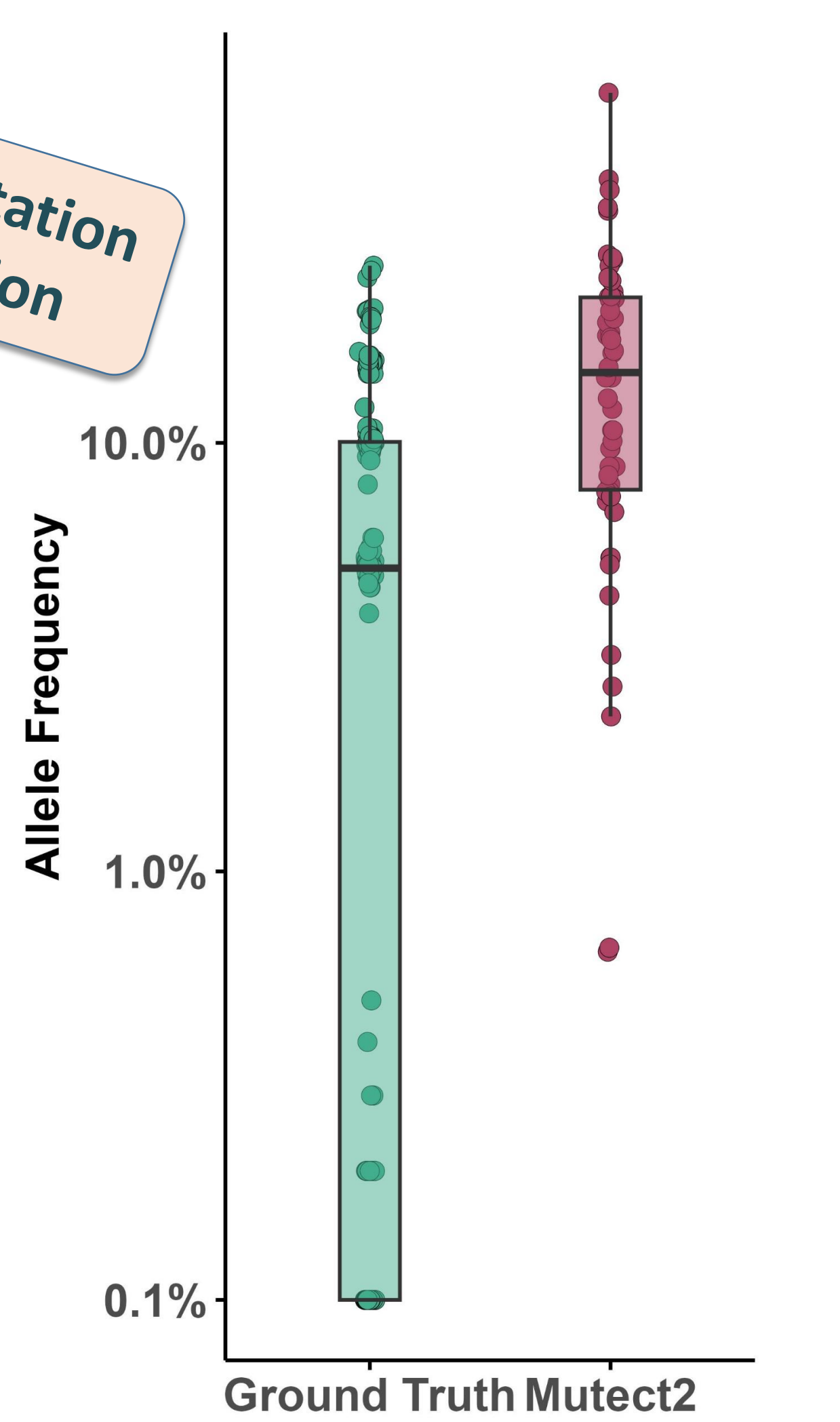


2 Benchmarking GATK-Mutect2

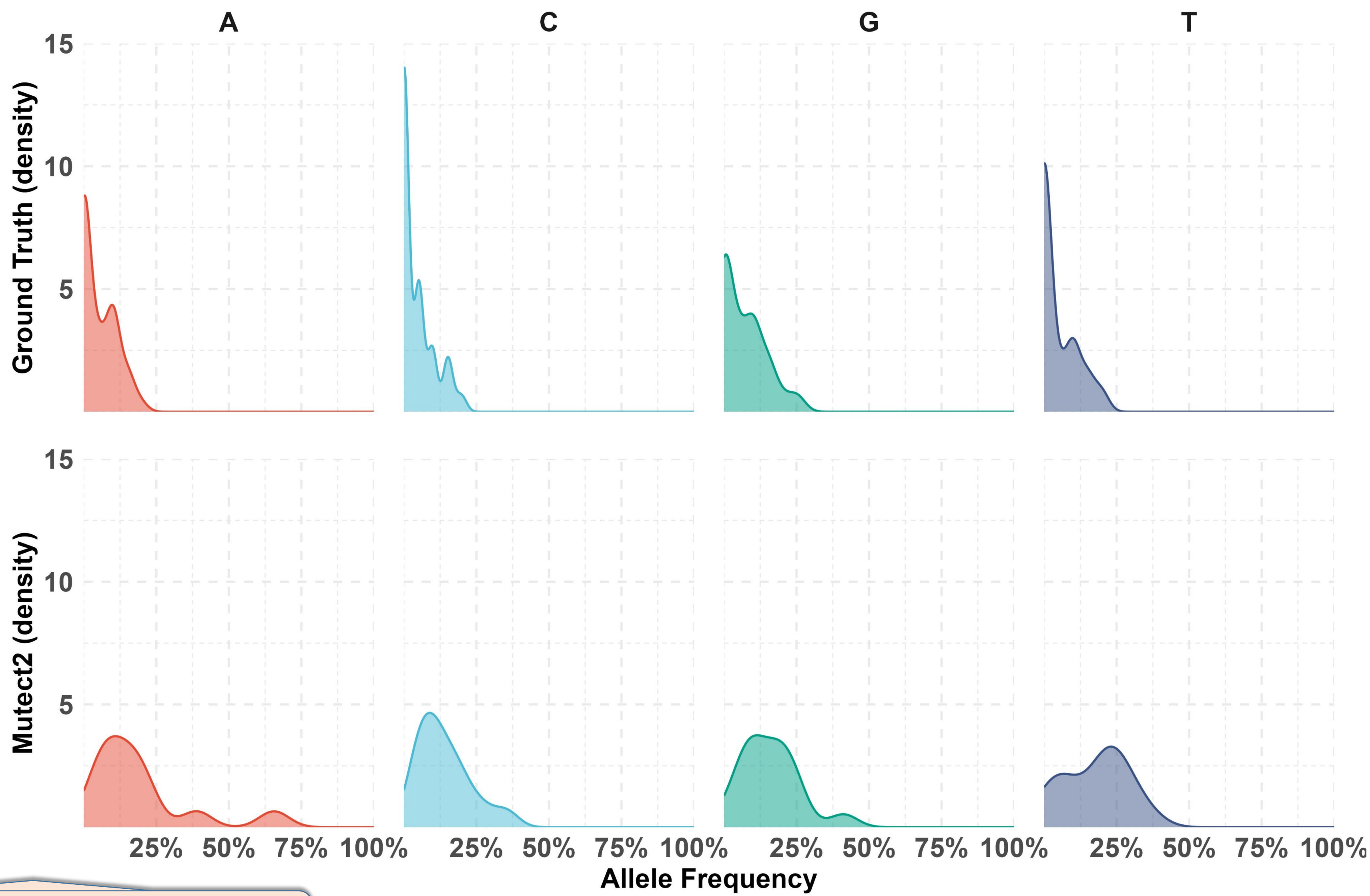
Down-sampling of coverage of «Ground Truth» Mutations



Differences in AF of «Ground Truth» Mutations



Variance in AF Density plots of «Ground Truth» Mutations per DNA Base

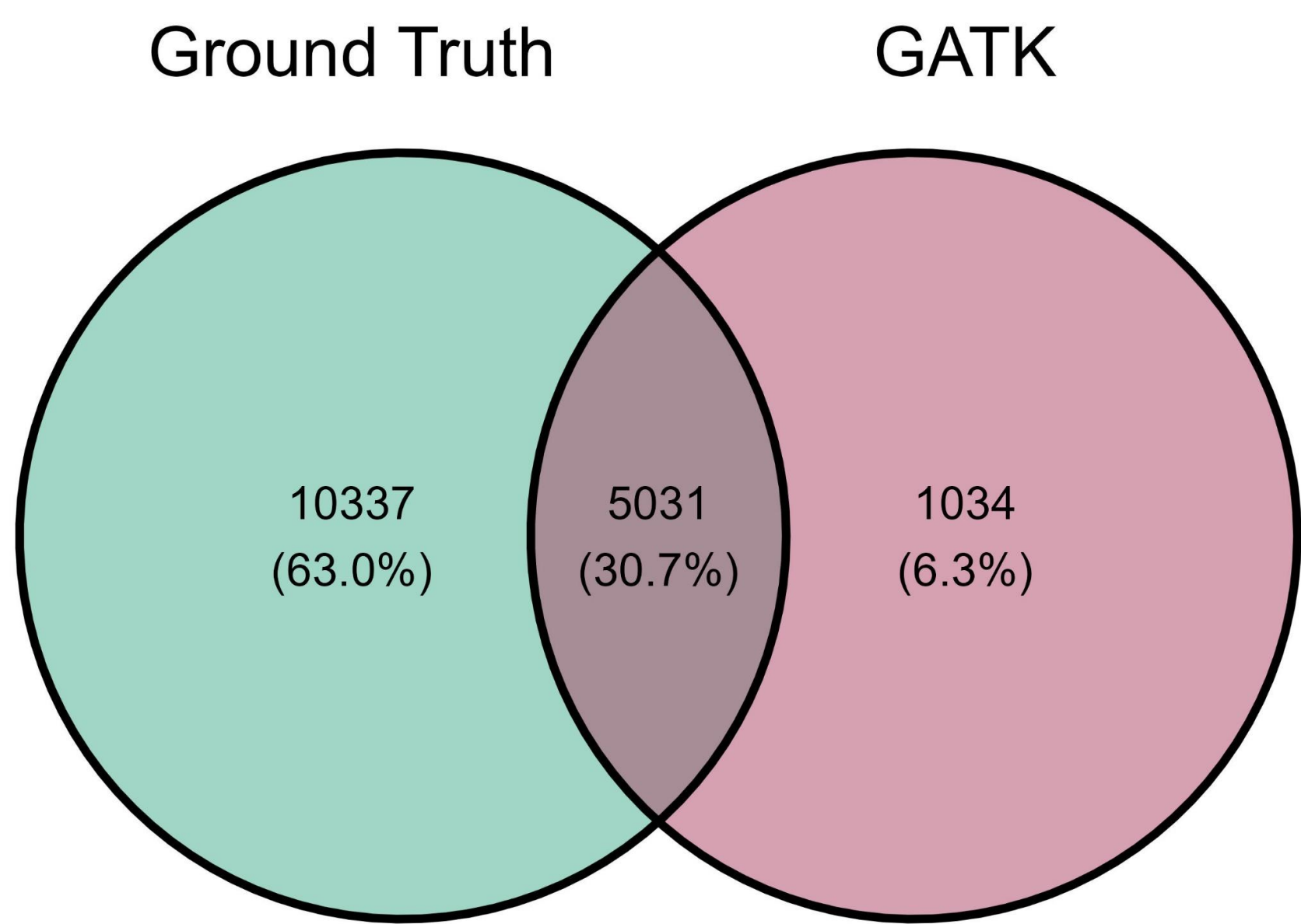


Difficulty to identify mutations at low frequencies



Divergence in the identification of SNVs and their AF of «Ground Truth» Mutations

Venn plot of the Overall Mutations



To learn more about our work please visit

