# PrivaSeq2 Figures

### Change the ICI-\Pi figure for reporting one dot for each deletion?

--Attacker Strikes Back--



Sample ID	Genome-wide Signal Profiles with Predicted Deletions	HIV Status	
Pheno-1	Teu Genomic Coordinate		
Pheno-2			
• •	• • •	•	
Pheno-n		+	

Geno-3

Pheno-n



SV-2

•

SV-3

•

 $\bullet \bullet \bullet$ 

•

### b) NA12878 large deletion

248,730 kb 248,750 kb 24 CTCF ALC: CONTRACTOR OF A STATE Control Ezh2 dents with motion to a state of the big of the state of t H3K27ac H3K27me3 H3K36me3 <u>, in a hilling as a</u> and the 1.6 H3K4me1 and the state of the second state of the secon H3K4me2 <u>ind Malan the in</u> H3K4me3 and the second state of th Lagr H3K79me2 and the second state of the state of the second state of the secon H3K9ac at the property of the propert H3K9me3 and there is a second of the manufacture of the second state of the history of the second state of the sec H4K20me1 Pooled ويتأكرون والرباد والأطر المراشطة

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	<b>.</b>		

Large Deletion

b) Detection and Genotyping of small and large deletions from signal profiles



### *Fig 2* a) GEUVADIS ICI-\Pi\_GW estimates:

a) Genotyping known deletions



c, d) GEUVADIS extremity attack: c) Genotyping only d) Joint Discovery + Genotyping



800

800

1000

### e) Indel length vs vulnerability statistics



f,g) Min # of identifying variants



### a, b) Gtex ICI-\Pi\_GW estimates: a) known indels, b) De-novo indels (1kG indels, GTex mapped reads)

a) Deletion Genotyping



Gtex extremity attack: known indels (1kG MAF>0.01) c) Genotyping d) Existence of Deletion Genotype







c,d) Extremity Attack Accuracy: c) Genotyping Only d) Joint Discovery and Genotyping









### a) Hi-C analysis b) Per SV ICI-\Pi\_GW estimates: NA12878 Hi-C data (different resolutions)





### Supplementary Figures

### Fig S1

### Sample-wide predictability of 1kG SVs



b)  $\pi_{SW}$  vs ICI for 1kG Structural Variants

ICI in 1000 Genomes SVs and Predictability ( $\pi$ ) of SV genotypes from GEUVADIS expression levels of



## **Fig S3** Allele Frequency Distribution of Large and Small Deletions



