

# Real-time, web-based genomic big data analysis

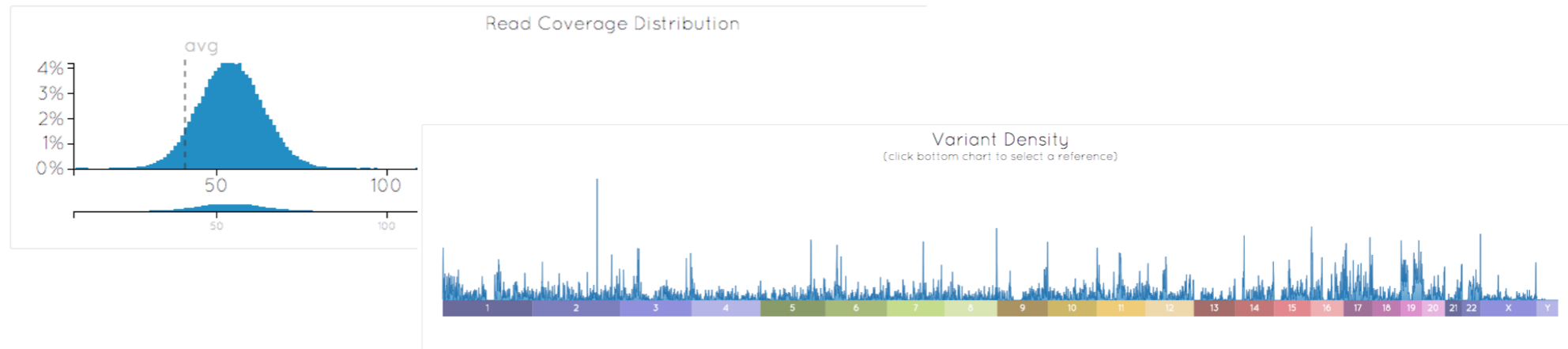
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# How can this be useful?

## 1. Help ensure quality of data



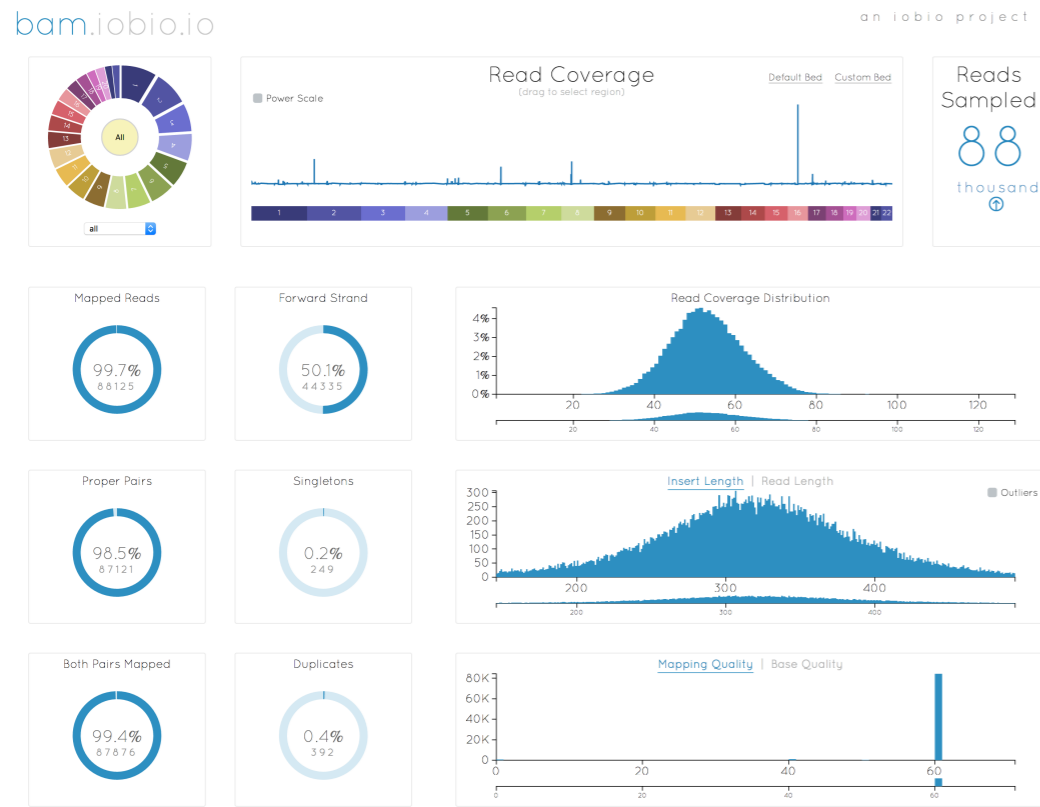
The screenshot shows a complex genomic analysis interface. On the left, there are several filter panels: "De novo VUS", "Recessive VUS", "High or moderate impact", "Inheritance mode", "Coverage (minimum)", "ClinVar", "VEP", "Allele Frequency ExAC", "Allele Frequency 1000G", "PolyPhen", "SIFT", "Zygosity", and "Type". The main panel displays "SELECTED GENE" as BCS1L on chromosome 2 (219,523,487-219,528,166). Below this, there are "RANKED VARIANTS" for three individuals: PROBAND 15-0015069, MOTHER 15-0015068, and FATHER 15-0015067. Each individual's section shows a genomic track with variants and coverage plots.

## 2. Allow non-bioinformaticians access to analysis

## 3. Aid, and speed the diagnosis process

# Project development

## Proof of concept



## Move to clinical relevance



Is the innovation useful?



# Project development



Where is money coming from?



U S T A R



# Project development

## Understand available resources

### TVC / USTAR

- Grant writing and resource advice
- Sources of funding
- Connections to collaborators/investors



U S T A R

Legal advice

