GENOME ANALYSIS

StainedGlass: Making colorful dot-plots of genomic sequence

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Abstract

Summary: Visualization of genomic repeats is often accomplished through the use of dot plots; however, the emergence of telomere-to-telomere assemblies with multi-megabase repeats requires new visualization strategies. Here, we introduce StainedGlass which can generate publication quality figures that communicate the identity and orientation of multi-megabase repeats while scaling to entire genomes. **Availability and implementation:** StainedGlass is implemented using snakemake and is available open source under the MIT license at mrvollger.github.io/StainedGlass/.

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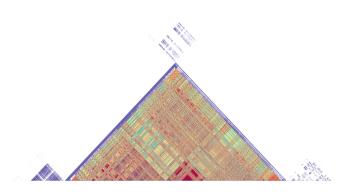


Fig. 1. Caption, caption

3 Conclusion Acknowledgements Funding

References

Bofelli, F., Name 2, Name 3 (2003) Article title, Journal Name, 199, 133-154.

1 Introduction

2 Usage and examples

Bofelli *et al.*, 2000 might want to know about text text text Figure 1

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