



CALIFORNIA CONSERVATION GENOMICS PROJECT

Dear CCGP Investigators,

We would like to remind you to please share any existing whole genome resequencing data with the CCGP Bioinformatics team as soon as possible. Please note that this will **not impact** your team's ability to use or access the data, but it is of utmost importance for the overall project.

WHAT ARE WE DOING WITH IT?

The WGS short read data will be uploaded to NCBI Short Read Archive (SRA) and **embargoed and remain private** until either the PI chooses to release the data, or shortly before the planned submission of the first synthetic CCGP publication (around mid-2023), whichever comes first.

Click [here](#) if you would like to review the CCGP Data Release Policies.

Once processed, the Bioinformatics team will run the data through their pipeline for alignment, filtration and variant calling. If you are curious, [click here](#) to see a visual representation of their pipeline. Your team will then receive VCF files and various quality metrics and visualizations.

HOW TO SHARE YOUR DATA

If your WGS data has already been generated or will be generated outside of the CCGP Mini-Core, it is your responsibility to share raw resequencing data and the associated metadata for all of the resequencing samples with the Bioinformatics team.

Please review our [Whole Genome Resequencing Data Ingest and Management Plan](#) for more information on how to share this material.

If you have any questions about this data sharing process, please contact our Data Wrangler, [Cade Mirchandani](#).

Thank you, as always, for your participation and collaboration. Please don't hesitate to reach out with any questions.

California Conservation Genomics Project

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