

Genomics for Australian Plants FAQ

1. How can I get involved?

All Consortium members are required to agree and sign the GAP Collaborative Agreement.

GAP Agreements and Policies can be found here:

<https://www.genomicsforaustralianplants.com/agreements-policies/>

GAP webpage:

<https://www.genomicsforaustralianplants.com/>

GAP Data Portal:

<https://data.bioplatforms.com/organization/about/bpa-plants>

Note: Access to GAP data is restricted to Consortium members for period of 12 months.

Please see the Data Policy for more details:

<https://www.genomicsforaustralianplants.com/agreements-policies/>

2. How is the GAP project funded? How long is the GAP project funded?

GAP is funded by Bioplatforms through the Australian Government National Collaborative Research Infrastructure Strategy (NCRIS), with additional contributions from the Botanic Gardens and Herbaria.

Bioplatforms funding for the project is for 3 years, from December 2018.

The Ian Potter Foundation has also provided substantial funds for 5 years (bioinformatics position, bioinformatics training and some funds for wet lab).

3. What are Bioplatforms' investments in GAP?

Bioplatforms investment is in genomics data generation, raw data storage and project management.

4. What in-kind contributions are expected of the researchers?

In-kind contributions from the wider plant community include sourcing plant material, preparation of material for genomic sequencing (e.g. DNA extractions), development of bioinformatics pipelines for data analysis and data sharing (e.g. contributing to and benefiting from collaborations with other international initiatives).

5. Can I apply for funding from GAP for my own research?

GAP is a nationally-inclusive and collaborative initiative. We do not fund individual research projects. The aim of the initiative is to build resources which will benefit the plant community.

6. What is the GAP Data Policy? Who can access data from my samples if I am part of GAP?

Data access falls broadly into two phases: a "mediated-access" phase, where access to the data will be limited to members of the Consortium and other authorised parties; and an "open-

access” phase where the data will be made openly available from resources including International Data Repositories.

Consortium members who have agreed and signed the GAP Collaborative Agreement will be given access upon registration at the Data Portal.

The “open-access” phase is set at 12 months from deposition of data into Bioplatforms Data Portal to allow the members of the Consortium and other authorised parties to progress analysis and publications, but not hold up the release of the data to be made openly available from resources including International Data Repositories.

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7. Will I be acknowledged if other Consortium members use data from my samples in their publication?

All communications (scientific or general publications and presentations) that arise from the Consortium’s work will appropriately acknowledge the input of all relevant contributions.

The publications should specify the collaborative nature of the project, and authorship is expected to include all those contributing significantly to the work.

Authorship for publications will be determined on a case-by-case basis.

For more details, please see the GAP Agreements and Policies:

<https://www.genomicsforaustralianplants.com/agreements-policies/>

8. How should I acknowledge the data used from the GAP consortium?

When using Genomics for Australian Plants Framework Initiative data, please acknowledge the Consortium and Bioplatforms Australia support using the wording in the following link:

<https://www.genomicsforaustralianplants.com/acknowledging-the-consortium/>

Please forward your publications and/or other communications to the [GAP Project Manager](#).