

# Resource Summary Report

Generated by [dkNET](#) on Sep 9, 2026

## QFAB Bioinformatics

RRID:SCR\_012513

Type: Tool

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### Proper Citation

QFAB Bioinformatics (RRID:SCR\_012513)

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### Resource Information

**URL:** <http://www.sciexnceexchange.com/facilities/qfab-bioinformatics>

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**Description:** Core facility providing support in bioinformatics, biostatistics and biodata to life sciences and health researchers across Queensland and beyond. Services include microarrays, next generation sequencing and proteomics profiling, biostatistical support from design of experiments, to analysis and interpretation, help High Degree Research students and Early Career Researchers in management, mining and visualization of small and big data, leveraging the cloud and hpc research infrastructure powered by the Queensland Cyber Infrastructure Foundation., THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

**Abbreviations:** QFAB

**Synonyms:** University of Queensland QFAB Bioinformatics, University of Queensland Queensland Facility for Advanced Bioinformatics, UQ QFAB, University of Queensland Facility for Advanced Bioinformatics, UQ QFAB Bioinformatics

**Resource Type:** access service resource, analysis service resource, core facility, data analysis service, production service resource, service resource, training service resource

**Keywords:** microarray, next generation sequencing, proteomics profiling, biostatistic,

**Availability:** THIS RESOURCE IS NO LONGER IN SERVICE

**Resource Name:** QFAB Bioinformatics

**Resource ID:** SCR\_012513

**Alternate IDs:** SciEx\_409

**Alternate URLs:** <https://www.qfab.org/>

**Record Creation Time:** 20220129T080310+0000

**Record Last Update:** 20260905T132724+0000

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## Ratings and Alerts

No rating or validation information has been found for QFAB Bioinformatics.

No alerts have been found for QFAB Bioinformatics.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [dkNET](#) .

Jia L, et al. (2026) EFO2 promotes triple-negative breast cancer progression and reduces CD8+ T cell-mediated killing by acetylating USF1 to enhance SLC2A1-mediated glycolysis. Biochemical pharmacology, 253(Pt 1), 118260.

de Freitas GPA, et al. (2022) Centromere protein J is overexpressed in human glioblastoma and promotes cell proliferation and migration. Journal of neurochemistry, 162(6), 501.