

Resource Summary Report

Generated by PRECISE-TBI on Sep 8, 2026

QFAB Bioinformatics

RRID:SCR_012513

Type: Tool

Proper Citation

QFAB Bioinformatics (RRID:SCR_012513)

Resource Information

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

Proper Citation: QFAB Bioinformatics (RRID:SCR_012513)

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

Ratings and Alerts

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

No alerts have been found for QFAB Bioinformatics.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at [PRECISE-TBI](#) .

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.