

# Resource Summary Report

Generated by [ASWG](#) on Aug 5, 2026

## University of Liege Interdisciplinary Cluster for Applied Genoproteomics Bioinformatics Core Facility

RRID:SCR\_017163

Type: Tool

### Proper Citation

University of Liege Interdisciplinary Cluster for Applied Genoproteomics Bioinformatics Core Facility (RRID:SCR\_017163)

### Resource Information

**URL:** [http://www.giga.uliege.be/cms/c\\_151663/fr/giga-genomcis-lab-manager](http://www.giga.uliege.be/cms/c_151663/fr/giga-genomcis-lab-manager)

**Proper Citation:** University of Liege Interdisciplinary Cluster for Applied Genoproteomics Bioinformatics Core Facility (RRID:SCR\_017163)

**Description:** Core provided by interdisciplinary center of biomedical research of Liege University. Develops and applies computational methods to extract information from large sets of biological and biomedical data.

**Abbreviations:** GIGA

**Synonyms:** , Bioinformatics Facility, Bioinformatics Core Facility, University of Liege, GIGA, Interdisciplinary Cluster for Applied Genoproteomics

**Resource Type:** access service resource, core facility, service resource

**Keywords:** bioinformatics, center, biomedical, data, computational, service

**Resource Name:** University of Liege Interdisciplinary Cluster for Applied Genoproteomics Bioinformatics Core Facility

**Resource ID:** SCR\_017163

**Record Creation Time:** 20220129T080333+0000

**Record Last Update:** 20260805T044406+0000

### Ratings and Alerts

No rating or validation information has been found for University of Liege Interdisciplinary Cluster for Applied Genoproteomics Bioinformatics Core Facility.

No alerts have been found for University of Liege Interdisciplinary Cluster for Applied Genoproteomics Bioinformatics Core Facility.

---

## Data and Source Information

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

---

## Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Alhindi T, et al. (2021) Genome-Wide Identification and Analysis of the MADS-Box Gene Family in American Beautyberry (*Callicarpa americana*). *Plants* (Basel, Switzerland), 10(9).

Lin X, et al. (2021) Chromosomal-Level Genome Assembly of Silver Sillago (*Sillago sihama*). *Genome biology and evolution*, 13(2).

Jiang JJ, et al. (2016) Insights into long noncoding RNAs of naked mole rat (*Heterocephalus glaber*) and their potential association with cancer resistance. *Epigenetics & chromatin*, 9, 51.