

# Resource Summary Report

Generated by SPARC SAWG on Aug 8, 2026

## IGG

RRID:SCR\_009114

Type: Tool

### Proper Citation

IGG (RRID:SCR\_009114)

### Resource Information

**URL:** <https://github.com/gaow/genetic-analysis-software/blob/master/pages/IGG.md>

**Proper Citation:** IGG (RRID:SCR\_009114)

**Description:** THIS RESOURCE IS NO LONGER IN SERVICE, documented September 22, 2016.

**Abbreviations:** IGG

**Synonyms:** Integrate Genotypes for genome-wide Genetic studies

**Resource Type:** software resource, software application

**Keywords:** gene, genetic, genomic, java, web-based

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

**Resource Name:** IGG

**Resource ID:** SCR\_009114

**Alternate IDs:** nlx\_154214

**Old URLs:** <http://bioinfo.hku.hk:13080/iggweb/>

**Record Creation Time:** 20220129T080251+0000

**Record Last Update:** 20260808T190505+0000

### Ratings and Alerts

No rating or validation information has been found for IGG.

No alerts have been found for IGG.

## 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 [SPARC SAWG](#) .

Yap Y, et al. (2004) Classification between normal and tumor tissues based on the pair-wise gene expression ratio. BMC cancer, 4, 72.

Danchin A, et al. (2004) The bag or the spindle: the cell factory at the time of systems' biology. Microbial cell factories, 3(1), 13.