## **Resource Summary Report**

Generated by NIF on Apr 29, 2025

# **OCplus**

RRID:SCR\_001342

Type: Tool

### **Proper Citation**

OCplus (RRID:SCR\_001342)

#### **Resource Information**

URL: http://www.bioconductor.org/packages/release/bioc/html/OCplus.html

**Proper Citation:** OCplus (RRID:SCR\_001342)

**Description:** Software package that allows to characterize the operating characteristics of a microarray experiment, i.e. the trade-off between false discovery rate and the power to detect truly regulated genes. The package includes tools both for planned experiments (for sample size assessment) and for already collected data (identification of differentially expressed genes).

Abbreviations: OCplus

Synonyms: Ocplus - Operating characteristics plus sample size and local fdr for microarray

experiments

**Resource Type:** software resource

**Defining Citation: PMID:16368770** 

**Keywords:** differential expression, microarray, multiple comparison

Funding:

Availability: GNU Lesser General Public License

Resource Name: OCplus

Resource ID: SCR\_001342

Alternate IDs: OMICS\_01999

**Record Creation Time:** 20220129T080207+0000

Record Last Update: 20250420T014027+0000

## **Ratings and Alerts**

No rating or validation information has been found for OCplus.

No alerts have been found for OCplus.

#### Data and Source Information

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at NIF.

Karathanasis N, et al. (2016) omicsNPC: Applying the Non-Parametric Combination Methodology to the Integrative Analysis of Heterogeneous Omics Data. PloS one, 11(11), e0165545.