## **Resource Summary Report**

Generated by RRID on Apr 8, 2025

# **DownSample**

RRID:SCR\_021056

Type: Tool

### **Proper Citation**

DownSample (RRID:SCR\_021056)

#### **Resource Information**

**URL:** https://www.flowjo.com/exchange/#/plugin/profile?id=25

Proper Citation: DownSample (RRID:SCR\_021056)

**Description:** Software tool to subset samples in FlowJo software.Reduces number of events in data matrix by generating subpopulation containing cells distributed regularly or randomly throughout selected parent population.

Resource Type: software resource

**Keywords:** subset samples, generating subpopulation containing cells, reduce event number, data matrix, flow cytometry

**Funding:** 

Resource Name: DownSample

Resource ID: SCR\_021056

Alternate URLs: https://docs.flowjo.com/seqgeq/dimensionality-reduction/downsample/

**Record Creation Time:** 20220129T080353+0000

**Record Last Update:** 20250214T183421+0000

### Ratings and Alerts

No rating or validation information has been found for DownSample.

No alerts have been found for DownSample.

#### Data and Source Information

Source: SciCrunch Registry

## **Usage and Citation Metrics**

We found 5 mentions in open access literature.

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

Sanlorenzo M, et al. (2025) Systemic IFN-I combined with topical TLR7/8 agonists promotes distant tumor suppression by c-Jun-dependent IL-12 expression in dendritic cells. Nature cancer, 6(1), 175.

Szabó E, et al. (2024) Identification of immune subsets with distinct lectin binding signatures using multi-parameter flow cytometry: correlations with disease activity in systemic lupus erythematosus. Frontiers in immunology, 15, 1380481.

Finlay CM, et al. (2023) T helper 2 cells control monocyte to tissue-resident macrophage differentiation during nematode infection of the pleural cavity. Immunity, 56(5), 1064.

Baracho GV, et al. (2022) Functional phenotyping of circulating human cytotoxic T cells and NK cells using a 16-color flow cytometry panel. STAR protocols, 3(1), 101069.

Tomaszewski WH, et al. (2021) Broad immunophenotyping of the murine brain tumor microenvironment. Journal of immunological methods, 499, 113158.