

Resource Summary Report

Generated by PRECISE-TBI on Aug 1, 2026

BD FACS Software sorter software

RRID:SCR_016722

Type: Tool

Proper Citation

BD FACS Software sorter software (RRID:SCR_016722)

Resource Information

URL: <http://www.bdbiosciences.com/us/instruments/research/cell-sorters/bd-influx/m/744777/features/software>

Proper Citation: BD FACS Software sorter software (RRID:SCR_016722)

Description: Software to provide control of the cell sorter from configuration and compensation setup to acquisition, sorting, and analysis, making it easy to customize the instrument for specific applications. BD implements software solutions that integrate data, devices and systems.

Abbreviations: BD FACS Software software

Synonyms: , BD FACS Software, Becton Dickinson FACS Software, BD FACS Software software

Resource Type: data visualization software, data analysis software, software application, software resource, data processing software

Keywords: control, cell, sorter, acquisition, analysis, integrate, data, instrument, system, Becton Dickinson and Company

Availability: Commercially available

Resource Name: BD FACS Software sorter software

Resource ID: SCR_016722

Record Creation Time: 20220129T080332+0000

Record Last Update: 20260801T042801+0000

Ratings and Alerts

No rating or validation information has been found for BD FACS Software sorter software.

No alerts have been found for BD FACS Software sorter software.

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 [PRECISE-TBI](#) .

Chocarro-Calvo A, et al. (2025) Fatty acid uptake activates an AXL-CAV1- β -catenin axis to drive melanoma progression. *Genes & development*, 39(7-8), 463.

Chocarro-Calvo A, et al. (2024) Phenotype-specific melanoma uptake of fatty acid from human adipocytes activates AXL and CAV1-dependent β -catenin nuclear accumulation. *bioRxiv : the preprint server for biology*.

Li S, et al. (2023) Seasonal and spatial variations of *Synechococcus* in abundance, pigment types, and genetic diversity in a temperate semi-enclosed bay. *Frontiers in microbiology*, 14, 1322548.

Errington TM, et al. (2021) Experiments from unfinished Registered Reports in the Reproducibility Project: Cancer Biology. *eLife*, 10.

Essex A, et al. (2019) Replication Study: Wnt activity defines colon cancer stem cells and is regulated by the microenvironment. *eLife*, 8.