Resource Summary Report

Generated by RRID on Aug 12, 2026

partec flomax

RRID:SCR_014026
Type: Tool

Proper Citation

partec flomax (RRID:SCR_014026)

Resource Information

URL: http://www.partec.de

Proper Citation: partec flomax (RRID:SCR_014026)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on June 1,2023. Partec uses FloMax software on most cytometers, which produces FCS 3.0 compliant files. FloMax rescales the fluorescent data on the acquisition display without changing the stored values.

Resource Type: portal, topical portal, data or information resource

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: partec flomax

Resource ID: SCR_014026

Record Creation Time: 20230602T050123+0000

Record Last Update: 20260812T044243+0000

Ratings and Alerts

No rating or validation information has been found for partec flomax.

No alerts have been found for partec flomax.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 7 mentions in open access literature.

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

Beric A, et al. (2021) Comparative phylogenetics of repetitive elements in a diverse order of flowering plants (Brassicales). G3 (Bethesda, Md.), 11(7).

Abbate L, et al. (2020) Genetic Distinctiveness Highlights the Conservation Value of a Sicilian Manna Ash Germplasm Collection Assigned to Fraxinus angustifolia (Oleaceae). Plants (Basel, Switzerland), 9(8).

Soorni A, et al. (2019) Transcriptome Landscape Variation in the Genus Thymus. Genes, 10(8).

Harkess A, et al. (2016) Retrotransposon Proliferation Coincident with the Evolution of Dioecy in Asparagus. G3 (Bethesda, Md.), 6(9), 2679.

Teixeira H, et al. (2014) Genetic diversity and differentiation of Juniperus thurifera in Spain and Morocco as determined by SSR. PloS one, 9(2), e88996.

Hadizadeh S, et al. (2014) Cytotoxic Effects of Newly Synthesized Palladium(II) Complexes of Diethyldithiocarbamate on Gastrointestinal Cancer Cell Lines. Biochemistry research international, 2014, 813457.

Tavares S, et al. (2014) Genome size analyses of Pucciniales reveal the largest fungal genomes. Frontiers in plant science, 5, 422.