

Resource Summary Report

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Cancer Research Institute IFOM FC Flow Cytometry Core Facility

RRID:SCR_026865

Type: Tool

Proper Citation

Cancer Research Institute IFOM FC Flow Cytometry Core Facility (RRID:SCR_026865)

Resource Information

URL: <https://www.ifom.eu/en/cancer-research/technological-units/flow-cytometry.php>

Proper Citation: Cancer Research Institute IFOM FC Flow Cytometry Core Facility (RRID:SCR_026865)

Description: Core Facility provides technology, instrumentation, and expert support for applications, including Immunophenotyping, Cell cycle and proliferation analysis, Metabolic studies, Apoptosis detection, Fluorescent protein expression analysis, Cytokine, chemokine, and intra/extracellular protein detection.

Synonyms: IFOM FC - Flow Cytometry Core Facility

Resource Type: access service resource, core facility, service resource

Keywords: ABRF, Immunophenotyping, Cell cycle and proliferation analysis, Metabolic studies, apoptosis detection, fluorescent protein expression analysis, cytokine, chemokine, protein detection,

Resource Name: Cancer Research Institute IFOM FC Flow Cytometry Core Facility

Resource ID: SCR_026865

Alternate IDs: ABRF_3227

Alternate URLs: <https://coremarketplace.org/?FacilityID=3227&citation=1>

Record Creation Time: 20250430T055337+0000

Record Last Update: 20260912T200513+0000

Ratings and Alerts

No rating or validation information has been found for Cancer Research Institute IFOM FC Flow Cytometry Core Facility.

No alerts have been found for Cancer Research Institute IFOM FC Flow Cytometry Core Facility.

Data and Source Information

Source: [SciCrunch Registry](#)

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](#) .

Oppezzo A, et al. (2026) Therapeutic inhibition of telomeric DNA damage response rescues hematopoietic dysfunction driven by telomere shortening and aging. Nature aging, 6(7), 1377.

di Lillo A, et al. (2026) Site-specific DNA double-strand break induces local transcription in cis and protein expression. Communications biology, 9(1).