

# Resource Summary Report

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## Arpege Core Facility

RRID:SCR\_023092

Type: Tool

### Proper Citation

Arpege Core Facility (RRID:SCR\_023092)

### Resource Information

**URL:** <http://www.arpege.cnrs.frCore FacilityArpege>

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**Description:** Offers functional and structural analysis of protein complexes. Offers possibility to characterize all types of molecular interactions: ligand-target binding, protein-protein interactions, signal transduction, and signalling complexes. Staff members are skilled specialists in luminescence-based techniques and in receptor biology. Application portfolio includes Analysis of cell signalling, by measuring signal transduction; Ligand-target binding (pharmacological analysis, affinity, selectivity); Protein-protein interactions (oligomerization, effectors...); Activation of signalling pathways (G proteins, GRK, arrestins & internalization); Intracellular production of second messengers (cAMP, IP1, Ca<sup>2+</sup>); Phosphorylation of signalling proteins (ERK, AKT...); Measurement of global cell functions and of different cell parameters (proliferation, migration, invasion...).

**Synonyms:** Arpege

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

**Keywords:** USEDit, ABRF, functional and structural analysis of protein complexes,

**Resource Name:** Arpege Core Facility

**Resource ID:** SCR\_023092

**Alternate IDs:** ABRF\_1654

**Alternate URLs:** <https://coremarketplace.org/?FacilityID=1654&citation=1>

**Record Creation Time:** 20221230T050203+0000

**Record Last Update:** 20260919T195953+0000

### Ratings and Alerts

No rating or validation information has been found for Arpege Core Facility.

No alerts have been found for Arpege Core Facility.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.