

Resource Summary Report

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French Ministry of Higher Education and Research

RRID:SCR_005532

Type: Tool

Proper Citation

French Ministry of Higher Education and Research (RRID:SCR_005532)

Resource Information

URL: <http://www.enseignementsup-recherche.gouv.fr/>

Proper Citation: French Ministry of Higher Education and Research (RRID:SCR_005532)

Description: Ministry of Higher Education, Research and Innovation, France is one of the 900 Higher Education-related Organizations and Associations.

Abbreviations: MESR

Synonyms: Ministere de l'Enseignement Supereur et de la Recherche, Ministere de l'Enseignement Supérieur et de la Recherche, French Ministry of Research, The Ministry, Le ministre

Resource Type: institution

Resource Name: French Ministry of Higher Education and Research

Resource ID: SCR_005532

Alternate IDs: ISNI: 0000 0001 2369 5464, Wikidata: Q2726949, grid.425729.f, Crossref funder ID: 501100004792, nlx_143604

Alternate URLs: <https://ror.org/03sjk9a61>

Record Creation Time: 20220129T080230+0000

Record Last Update: 20260725T190605+0000

Ratings and Alerts

No rating or validation information has been found for French Ministry of Higher Education and Research.

No alerts have been found for French Ministry of Higher Education and Research.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 37 mentions in open access literature.

Listed below are recent publications. The full list is available at [nidm-terms](#).

Marcantonio W, et al. (2025) Sex-Specific Behavioral Features of Juvenile and Adult Haploinsufficient *Scn2a*^{+/-} Female Mice, Model of Autism Spectrum Disorder. *Genes, brain, and behavior*, 24(5), e70034.

Amroune K, et al. (2025) Sparse innervation and local heterogeneity in the vibrissal corticostriatal projection. *eLife*, 14.

Jablonski KP, et al. (2022) Contribution of 3D genome topological domains to genetic risk of cancers: a genome-wide computational study. *Human genomics*, 16(1), 2.

Laussel C, et al. (2022) 2-deoxyglucose transiently inhibits yeast AMPK signaling and triggers glucose transporter endocytosis, potentiating the drug toxicity. *PLoS genetics*, 18(8), e1010169.

Dehapiot B, et al. (2021) RhoA- and Cdc42-induced antagonistic forces underlie symmetry breaking and spindle rotation in mouse oocytes. *PLoS biology*, 19(9), e3001376.

Broster Reix CE, et al. (2021) Intrabody-Induced Cell Death by Targeting the T. brucei Cytoskeletal Protein TbBILBO1. *Microbiology spectrum*, 9(2), e0091521.

Mauras S, et al. (2021) Mitigating COVID-19 outbreaks in workplaces and schools by hybrid telecommuting. *PLoS computational biology*, 17(8), e1009264.

Di Mattia T, et al. (2020) FFAT motif phosphorylation controls formation and lipid transfer function of inter-organelle contacts. *The EMBO journal*, 39(23), e104369.

Foucaud L, et al. (2019) Modulation of protective reflex cough by acute immune driven inflammation of lower airways in anesthetized rabbits. *PloS one*, 14(12), e0226442.

Cau E, et al. (2019) A Notch-mediated, temporal asymmetry in BMP pathway activation promotes photoreceptor subtype diversification. *PLoS biology*, 17(1), e2006250.

Bouchard B, et al. (2019) Behavioural responses of humpback whales to food-related chemical stimuli. *PloS one*, 14(2), e0212515.

El Hajj A, et al. (2019) Age-related changes in regiospecific expression of Lipolysis Stimulated Receptor (LSR) in mice brain. *PloS one*, 14(6), e0218812.

Raieli S, et al. (2019) TLR1/2 orchestrate human plasmacytoid predendritic cell response to gram+ bacteria. *PLoS biology*, 17(4), e3000209.

Chiara V, et al. (2019) Social intolerance is a consequence, not a cause, of dispersal in spiders. *PLoS biology*, 17(7), e3000319.

Léna I, et al. (2019) Nav1.2 haploinsufficiency in Scn2a knock-out mice causes an autistic-like phenotype attenuated with age. *Scientific reports*, 9(1), 12886.

Rocha de Almeida T, et al. (2019) Domestication may affect the maternal mRNA profile in unfertilized eggs, potentially impacting the embryonic development of Eurasian perch (*Perca fluviatilis*). *PloS one*, 14(12), e0226878.

Fischer CE, et al. (2018) The multiple maternal legacy of the Late Iron Age group of Urville-Nacqueville (France, Normandy) documents a long-standing genetic contact zone in northwestern France. *PloS one*, 13(12), e0207459.

Di Mattia T, et al. (2018) Identification of MOSPD2, a novel scaffold for endoplasmic reticulum membrane contact sites. *EMBO reports*, 19(7).

Jouand N, et al. (2018) HCMV triggers frequent and persistent UL40-specific unconventional HLA-E-restricted CD8 T-cell responses with potential autologous and allogeneic peptide recognition. *PLoS pathogens*, 14(4), e1007041.

Beau A, et al. (2017) Multi-scale ancient DNA analyses confirm the western origin of Michelsberg farmers and document probable practices of human sacrifice. *PloS one*, 12(7), e0179742.