Resource Summary Report

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Swiss National Science Foundation

RRID:SCR_011554
Type: Tool

Proper Citation

Swiss National Science Foundation (RRID:SCR_011554)

Resource Information

URL: http://www.snf.ch/en/Pages/default.aspx

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Description: The Swiss National Science Foundation (SNSF) is Switzerland"s leading provider of scientific research funding. The SNSF annually supports some 7200 researchers, almost 80 percent of whom are aged 35 years or younger. With its federal mandate, it supports basic research in all disciplines, from philosophy and biology to the nanosciences and medicine. It also invests in applied research in various scientific fields. The focus of its activities is the scientific endorsement of projects submitted by researchers. The best applicants are funded by the SNSF with an annual total amount equalling approximately CHF 700 million. Established in 1952 as a foundation under private law, the SNSF has the autonomy it needs to promote independent scientific research. The SNSF is committed to promoting young scientists and works to ensure that scientific research in Switzerland has the most favourable conditions for developing internationally. It also encourages dialogue between scientists and representatives in society, politics and the economy.

Abbreviations: SNSF

Resource Type: institution

Resource Name: Swiss National Science Foundation

Resource ID: SCR_011554

Alternate IDs: nlx_60722, grid.425888.b, Wikidata: Q471732, Crossref funder ID: 501100001711, ISNI: 0000 0001 1957 0992

Alternate URLs: https://ror.org/00yjd3n13

Record Creation Time: 20220129T080305+0000

Record Last Update: 20260725T190718+0000

Ratings and Alerts

No rating or validation information has been found for Swiss National Science Foundation.

No alerts have been found for Swiss National Science Foundation.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 301 mentions in open access literature.

Listed below are recent publications. The full list is available at kravitz2 .

Giacomino K, et al. (2025) Physical exercises to prevent and rehabilitate hospital-associated disability in hospitalised older people: A protocol for a living systematic review with network meta-analysis. F1000Research, 14, 414.

Croisier M, et al. (2025) Electron microscopy ascertains presence of centrioles in rectal epithelial cells of C. elegans L1 larvae. microPublication biology, 2025.

Ryckebusch F, et al. (2025) Layers of immunity: Deconstructing the Drosophila effector response. eLife, 14.

Mirkovic M, et al. (2025) Dissociation of the nuclear basket triggers chromosome loss in aging yeast. eLife, 14.

Chaguza C, et al. (2025) The capsule and genetic background, rather than specific individual loci, strongly influence in vitro pneumococcal growth kinetics. eLife, 14.

Potenza DM, et al. (2025) Cell-autonomous and non-cell-autonomous effects of Arginase 2 on cardiac aging. eLife, 13.

Morsy Y, et al. (2025) Blood-borne immune cells carry low biomass DNA remnants of microbes in patients with colorectal cancer or inflammatory bowel disease. Gut microbes, 17(1), 2530157.

López-Jiménez AT, et al. (2025) High-content high-resolution microscopy and deep learning-assisted analysis reveals host and bacterial heterogeneity during Shigella infection. eLife, 13.

Garbelli M, et al. (2025) Characterization of postsynaptic glutamate transporter functionality in the zebrafish retinal first synapse across different wavelengths. eLife, 13.

Cornelius J, et al. (2025) From literature to biodiversity data: mining arthropod organismal traits with machine learning. Biodiversity data journal, 13, e153070.

Bru S, et al. (2025) Acidocalcisome-like vacuoles constitute a feedback-controlled phosphate buffering system for the cytosol. eLife, 14.

Gini?nait? R, et al. (2025) Nucleosome wrapping energy in CpG islands and the role of epigenetic base modifications. eLife, 13.

Burdet V, et al. (2024) Ultrastructure Expansion Microscopy applied to C. elegans embryos. microPublication biology, 2024.

Merigliano C, et al. (2024) Replication forks associated with long nuclear actin filaments in mild stress conditions display increased dynamics. microPublication biology, 2024.

Schmutzer M, et al. (2024) Frustration can Limit the Adaptation of Promiscuous Enzymes Through Gene Duplication and Specialisation. Journal of molecular evolution, 92(2), 104.

Schmutzer M, et al. (2023) Not Quite Lost in Translation: Mistranslation Alters Adaptive Landscape Topography and the Dynamics of Evolution. Molecular biology and evolution, 40(6).

Mairpady Shambat S, et al. (2022) Hyperinflammatory environment drives dysfunctional myeloid cell effector response to bacterial challenge in COVID-19. PLoS pathogens, 18(1), e1010176.

Wolfe CM, et al. (2021) Systematic review of Integrated Disease Surveillance and Response (IDSR) implementation in the African region. PloS one, 16(2), e0245457.

Dietz N, et al. (2021) Structural basis for selective AMPylation of Rac-subfamily GTPases by Bartonella effector protein 1 (Bep1). Proceedings of the National Academy of Sciences of the United States of America, 118(12).

Craig AJ, et al. (2021) Transcriptomic characterization of cancer-testis antigens identifies MAGEA3 as a driver of tumor progression in hepatocellular carcinoma. PLoS genetics, 17(6), e1009589.