

Resource Summary Report

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Source BioScience LifeSciences

RRID:SCR_003344

Type: Tool

Proper Citation

Source BioScience LifeSciences (RRID:SCR_003344)

Resource Information

URL: <http://www.lifesciences.sourcebioscience.com/welcome-to-source-bioscience-lifesciences.aspx>

Proper Citation: Source BioScience LifeSciences (RRID:SCR_003344)

Description: Source BioScience LifeSciences are European leaders in DNA sequencing, genomic services, bioinformatic analyses and offers a comprehensive portfolio of genomic reagents and antibodies. Source BioScience LifeScience is a CPA, GLP/GCP accredited and Illumina NGS CPro certified genomic service provider and distributes a comprehensive range of biological products to companies and institutions worldwide. Source BioScience is proud to announce that Source BioScience imaGenes has joined it's LifeSciences group. This will enable us to bring our customers one unified website with our complete range of Products, Clones and Services. We have merged our clone libraries together so that can now offer you over 20 million clones! Add to this our excellent sequencing service and our range of over 100,000 antibodies and Source BioScience LifeSciences is your ideal outsourcing partner.

Abbreviations: RZPD

Synonyms: imaGenes, Source BioScience imaGenes

Resource Type: biomaterial supply resource, material resource

Keywords: clones, libraries, microarray, dna sequencing, rna interference, next generation sequencing, proteomics

Availability: Free, Freely available

Resource Name: Source BioScience LifeSciences

Resource ID: SCR_003344

Alternate IDs: nif-0000-31984

Alternate URLs: <https://sourcebioscience.com/>

Record Creation Time: 20220129T080218+0000

Record Last Update: 20260829T183052+0000

Ratings and Alerts

No rating or validation information has been found for Source BioScience LifeSciences.

No alerts have been found for Source BioScience LifeSciences.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 72 mentions in open access literature.

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

Wildsmith C, et al. (2025) Genomic and phenotypic characterization of staphylococci isolated from the skin of non-human primates. *Microbiology (Reading, England)*, 171(3).

Azeem A, et al. (2024) Investigating the genetic basis of hereditary spastic paraplegia and cerebellar Ataxia in Pakistani families. *BMC neurology*, 24(1), 354.

Sadeh TT, et al. (2023) Cav1.4 congenital stationary night blindness is associated with an increased rate of proteasomal degradation. *Frontiers in cell and developmental biology*, 11, 1161548.

Marwan M, et al. (2023) Unravelling the genetic basis of retinal dystrophies in Pakistani consanguineous families. *BMC ophthalmology*, 23(1), 205.

Liu QR, et al. (2023) Novel Hominid-Specific IAPP Isoforms: Potential Biomarkers of Early Alzheimer's Disease and Inhibitors of Amyloid Formation. *Biomolecules*, 13(1).

Vozdek R, et al. (2022) Fluorescent reporter of *Caenorhabditis elegans* Parkin: Regulators of its abundance and role in autophagy-lysosomal dynamics. *Open research Europe*, 2, 23.

Thornton EL, et al. (2022) Self-Assembling Protein Surfaces for In Situ Capture of Cell-Free-Synthesized Proteins. *Frontiers in bioengineering and biotechnology*, 10, 915035.

Sui Y, et al. (2017) Paradoxical myeloid-derived suppressor cell reduction in the bone marrow of SIV chronically infected macaques. *PLoS pathogens*, 13(5), e1006395.

Herschhorn A, et al. (2017) The 20-21 of gp120 is a regulatory switch for HIV-1 Env conformational transitions. *Nature communications*, 8(1), 1049.

Hollenbaugh JA, et al. (2017) Substrates and Inhibitors of SAMHD1. *PloS one*, 12(1), e0169052.

Goetze RW, et al. (2017) A CRISPR/Cas9 approach reveals that the polymerase activity of DNA polymerase γ is dispensable for HIV-1 infection in dividing and nondividing cells. *The Journal of biological chemistry*, 292(34), 14016.

Ernst D, et al. (2017) Anti-MYC-associated zinc finger protein antibodies are associated with inflammatory atherosclerotic lesions on 18F-fluorodeoxyglucose positron emission tomography. *Atherosclerosis*, 259, 12.

Robertson HE, et al. (2017) The mitochondrial genomes of the acoelomorph worms *Paratomella rubra*, *Isodiametra pulchra* and *Archaphanostoma ylvae*. *Scientific reports*, 7(1), 1847.

Vargas-Inchaustegui DA, et al. (2016) Evaluation of Functional NK Cell Responses in Vaccinated and SIV-Infected Rhesus Macaques. *Frontiers in immunology*, 7, 340.

Yuan T, et al. (2016) An HIV-1 capsid binding protein TRIM11 accelerates viral uncoating. *Retrovirology*, 13(1), 72.

Chevalier B, et al. (2015) miR-34/449 control apical actin network formation during multiciliogenesis through small GTPase pathways. *Nature communications*, 6, 8386.

Hollevoet K, et al. (2015) Methylation-associated partial down-regulation of mesothelin causes resistance to anti-mesothelin immunotoxins in a pancreatic cancer cell line. *PloS one*, 10(3), e0122462.

Bagley K, et al. (2015) The catalytic A1 domains of cholera toxin and heat-labile enterotoxin are potent DNA adjuvants that evoke mixed Th1/Th17 cellular immune responses. *Human vaccines & immunotherapeutics*, 11(9), 2228.

Marchesin V, et al. (2015) ARF6-JIP3/4 regulate endosomal tubules for MT1-MMP exocytosis in cancer invasion. *The Journal of cell biology*, 211(2), 339.

Mothe B, et al. (2015) A human immune data-informed vaccine concept elicits strong and broad T-cell specificities associated with HIV-1 control in mice and macaques. *Journal of translational medicine*, 13, 60.