Resource Summary Report

Generated by dkNET on May 8, 2025

BEI Resource Repository

RRID:SCR_013698

Type: Tool

Proper Citation

BEI Resource Repository (RRID:SCR_013698)

Resource Information

URL: https://www.beiresources.org/

Proper Citation: BEI Resource Repository (RRID:SCR_013698)

Description: Central data repository that supplies organisms and reagents to the broad

community of microbiology and infectious diseases researchers.

Synonyms: BEI Resource

Resource Type: biobank, service resource, storage service resource, material storage

repository

Keywords: biomaterial, material, reagent, microbiology, infectious diseases

Related Condition: infectious disease

Funding: NIAID

Resource Name: BEI Resource Repository

Resource ID: SCR_013698

License URLs: http://www.beiresources.org/TermsofUse.aspx

Record Creation Time: 20220129T080317+0000

Record Last Update: 20250508T065454+0000

Ratings and Alerts

No rating or validation information has been found for BEI Resource Repository.

No alerts have been found for BEI Resource Repository.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 193 mentions in open access literature.

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

Vosbigian KA, et al. (2025) ATF6 enables pathogen infection in ticks by inducing stomatin and altering cholesterol dynamics. bioRxiv: the preprint server for biology.

Lv H, et al. (2025) Differential antigenic imprinting effects between influenza H1N1 hemagglutinin and neuraminidase in a mouse model. Journal of virology, 99(1), e0169524.

Khan M, et al. (2025) Repurposing of lonafarnib as a treatment for SARS-CoV-2 infection. JCI insight, 10(1).

Obellianne C, et al. (2024) Interspecies co-feeding transmission of Powassan virus between a native tick, Ixodes scapularis, and the invasive East Asian tick, Haemaphysalis longicornis. Parasites & vectors, 17(1), 259.

Gualdrón-López M, et al. (2024) Proteomics of circulating extracellular vesicles reveals diverse clinical presentations of COVID-19 but fails to identify viral peptides. Frontiers in cellular and infection microbiology, 14, 1442743.

Rafferty C, et al. (2024) Loop-Mediated Isothermal Amplification Assay to Detect Invasive Malaria Vector Anopheles stephensi Mosquitoes. Emerging infectious diseases, 30(9), 1770.

Ouyang WO, et al. (2024) Rapid synthesis and screening of natively paired antibodies against influenza hemagglutinin stem via oPool+ display. bioRxiv: the preprint server for biology.

Vergara S, et al. (2024) Structural basis of deoxynucleotide addition by HIV-1 RT during reverse transcription. Nature communications, 15(1), 10553.

Faísca-Silva D, et al. (2024) High-Sensitivity RT-LAMP for Molecular Detection of O'nyong-nyong (Alphavirus onyong). Pathogens (Basel, Switzerland), 13(10).

de Cesare M, et al. (2024) Flexible and cost-effective genomic surveillance of P. falciparum malaria with targeted nanopore sequencing. Nature communications, 15(1), 1413.

Joy J, et al. (2024) Antigen specificities and proviral integration sites differ in HIV-infected cells by timing of antiretroviral treatment initiation. The Journal of clinical investigation, 134(14).

Chen Y-C, et al. (2024) Uncovering the roles of Mycobacterium tuberculosis melH in redox and bioenergetic homeostasis: implications for antitubercular therapy. mSphere, 9(4), e0006124.

Germano ER, et al. (2024) Building-level wastewater surveillance localizes interseasonal influenza variation. mSphere, 9(1), e0060023.

Keck JW, et al. (2024) Wastewater Surveillance for Identifying SARS-CoV-2 Infections in Long-Term Care Facilities, Kentucky, USA, 2021-2022. Emerging infectious diseases, 30(3), 530.

Nayak S, et al. (2024) Population genomics and transcriptomics of Plasmodium falciparum in Cambodia and Vietnam uncover key components of the artemisinin resistance genetic background. Nature communications, 15(1), 10625.

Gao C, et al. (2024) MAIVeSS: streamlined selection of antigenically matched, high-yield viruses for seasonal influenza vaccine production. Nature communications, 15(1), 1128.

Liénard MA, et al. (2024) TRPA5 encodes a thermosensitive ankyrin ion channel receptor in a triatomine insect. iScience, 27(4), 109541.

Olali AZ, et al. (2024) The anti-HIV drug abacavir stimulates ?-catenin activity in osteoblast lineage cells. JBMR plus, 8(5), ziae037.

Huang X, et al. (2024) Unveiling candidate genes for metabolic resistance to malathion in Aedes albopictus through RNA sequencing-based transcriptome profiling. PLoS neglected tropical diseases, 18(6), e0012243.

Porwollik S, et al. (2024) A genome-wide collection of barcoded single-gene deletion mutants in Salmonella enterica serovar Typhimurium. PloS one, 19(3), e0298419.