

Resource Summary Report

Generated by T1D on Aug 1, 2026

NIAID

RRID:SCR_016598

Type: Tool

Proper Citation

NIAID (RRID:SCR_016598)

Resource Information

URL: <https://www.niaid.nih.gov/>

Proper Citation: NIAID (RRID:SCR_016598)

Description: National Institute of Allergy and Infectious Diseases is a leading research institution to understand, treat, and prevent infectious, immunologic, and allergic diseases.

Abbreviations: NIAID

Synonyms: National Institute of Allergy and Infectious Diseases

Resource Type: data or information resource, portal, topical portal, disease-related portal, organization portal

Keywords: immunologic, allergic, infectious, disease, institute, treat, prevent

Funding: NIH

Resource Name: NIAID

Resource ID: SCR_016598

Record Creation Time: 20220129T080331+0000

Record Last Update: 20260801T042800+0000

Ratings and Alerts

No rating or validation information has been found for NIAID.

No alerts have been found for NIAID.

Data and Source Information

Usage and Citation Metrics

We found 510 mentions in open access literature.

Listed below are recent publications. The full list is available at [T1D](#) .

Rist M, et al. (2025) In vitro HIV DNA integration in STAT3 drives T cell persistence-A model of HIV-associated T cell lymphoma. PLoS pathogens, 21(7), e1013087.

Jones MA, et al. (2025) Multiple marginalized identities and internalized HIV stigma among people living with HIV in South Florida: An intersectional approach. PloS one, 20(8), e0329966.

Loveday EK, et al. (2025) Starve a cold or feed a fever? Identifying cellular metabolic changes following infection and exposure to SARS-CoV-2. PloS one, 20(2), e0305065.

de Paula FM, et al. (2025) Dynamics of anti-Strongyloides IgG antibody responses and implications for strongyloidiasis surveillance in rural Amazonians: A population-based panel data analysis. PLoS neglected tropical diseases, 19(4), e0012967.

Becker JT, et al. (2025) Mammalian antiviral proteins ZAP and KHNYN can independently restrict CpG-enriched avian viruses. PLoS biology, 23(10), e3003471.

Nemzow L, et al. (2025) Intracellular lymphocyte protein biomarkers for early radiological triage in the human population. PloS one, 20(9), e0331230.

Derby N, et al. (2025) Liver biopsies obtained throughout SIV infection reveal evolving interferon stimulated protein expression within distinct monocyte/macrophage subsets. PLoS pathogens, 21(9), e1013175.

Coelho MA, et al. (2025) The complex evolution and genomic dynamics of mating-type loci in Cryptococcus and Kwoniella. PLoS biology, 23(10), e3003417.

Speth ZJ, et al. (2025) Performance of two low-threshold population replacement gene drives in cage populations of the yellow fever mosquito, Aedes aegypti. PLoS genetics, 21(6), e1011757.

Picker L, et al. (2025) Identification of initial sites of SIV rebound after treatment cessation. Research square.

Williamson C, et al. (2025) Influence of the broadly neutralizing antibody VRC01 on HIV breakthrough virus populations in antibody-mediated prevention trials. bioRxiv : the preprint server for biology.

Inagaki T, et al. (2025) KSHV TR deletion episomes uncover enhancer-promoter dynamics in gene regulation. PLoS pathogens, 21(12), e1013061.

Horejsi RV, et al. (2025) Quantitative ethology of schistosome miracidia characterizes a conserved snail peptide that inhibits host recognition. PLoS pathogens, 21(12), e1013766.

Li Y, et al. (2025) Identification of a protective antigen reveals the trade-off between iron acquisition and antigen exposure in a global fungal pathogen. *Proceedings of the National Academy of Sciences of the United States of America*, 122(7), e2420898122.

Miller D, et al. (2025) Pooled PPIseq: Screening the SARS-CoV-2 and human interface with a scalable multiplexed protein-protein interaction assay platform. *PloS one*, 20(1), e0299440.

Nzeli CO, et al. (2025) Detection of *Leishmania* metacyclogenesis within the sand fly vector employing a real-time PCR for *sherp* gene expression: A tool for *Leishmania* surveillance and transmission potential. *PLoS neglected tropical diseases*, 19(3), e0012915.

Martin MA, et al. (2025) Quantifying prevalence and risk factors of HIV multiple infection in Uganda from population-based deep-sequence data. *PLoS pathogens*, 21(4), e1013065.

Wiens KE, et al. (2025) Correction: Estimating the proportion of clinically suspected cholera cases that are true *Vibrio cholerae* infections: A systematic review and meta-analysis. *PLoS medicine*, 22(9), e1004743.

Melo G, et al. (2025) Activation of B1 B cells by *F. tularensis* atypical LPS depends on classical complement and C3a. *PLoS pathogens*, 21(12), e1013799.

Blackburn EE, et al. (2025) Host-induced climate change: Carbon dioxide tolerance as a *Cryptococcus neoformans* virulence trait. *PLoS pathogens*, 21(8), e1013351.