

Resource Summary Report

Generated by NIF on Aug 9, 2026

Walter and Eliza Hall Institute of Medical Research; Victoria; Australia

RRID:SCR_011773

Type: Tool

Proper Citation

Walter and Eliza Hall Institute of Medical Research; Victoria; Australia
(RRID:SCR_011773)

Resource Information

URL: <http://www.wehi.edu.au/>

Proper Citation: Walter and Eliza Hall Institute of Medical Research; Victoria; Australia
(RRID:SCR_011773)

Description: Medical research institute in Australia that researches and develops treatments for cancer, infectious diseases and chronic inflammatory diseases.

Abbreviations: WEHI

Synonyms: Walter and Eliza Hall Institute of Medical Research, Walter and Eliza Hall Institute

Resource Type: institution

Resource Name: Walter and Eliza Hall Institute of Medical Research; Victoria; Australia

Resource ID: SCR_011773

Alternate IDs: Wikidata: Q7966487, nlx_157660, grid.1042.7

Alternate URLs: <https://ror.org/01b6kha49>

Record Creation Time: 20220129T080306+0000

Record Last Update: 20260808T185935+0000

Ratings and Alerts

No rating or validation information has been found for Walter and Eliza Hall Institute of Medical Research; Victoria; Australia.

No alerts have been found for Walter and Eliza Hall Institute of Medical Research; Victoria; Australia.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 10 mentions in open access literature.

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

Thung TY, et al. (2024) Genetic variation in individuals from a population of the minimalist bacteriophage Merri-merri-uth nyilam marra-natj driving evolution of the virus. mBio, 15(12), e0256424.

Han J, et al. (2023) AmpSeqR: an R package for amplicon deep sequencing data analysis. F1000Research, 12, 327.

Han J, et al. (2022) Population-level genome-wide STR discovery and validation for population structure and genetic diversity assessment of Plasmodium species. PLoS genetics, 18(1), e1009604.

Almuslehi MSM, et al. (2022) Histological and Top-Down Proteomic Analyses of the Visual Pathway in the Cuprizone Demyelination Model. Journal of molecular neuroscience : MN, 72(6), 1374.

Singh RP, et al. (2020) Disrupting Mitochondrial Copper Distribution Inhibits Leukemic Stem Cell Self-Renewal. Cell stem cell, 26(6), 926.

Jones DC, et al. (2016) Allele-specific recognition by LILRB3 and LILRA6 of a cytokeratin 8-associated ligand on necrotic glandular epithelial cells. Oncotarget, 7(13), 15618.

Lesterlin C, et al. (2014) RecA bundles mediate homology pairing between distant sisters during DNA break repair. Nature, 506(7487), 249.

Ruffini PA, et al. (2014) Targeted DNA vaccines eliciting crossreactive anti-idiotypic antibody responses against human B cell malignancies in mice. Journal of translational medicine, 12, 207.

Markham JF, et al. (2010) A minimum of two distinct heritable factors are required to explain correlation structures in proliferating lymphocytes. Journal of the Royal Society, Interface, 7(48), 1049.

Wykes MN, et al. (2007) Plasmodium strain determines dendritic cell function essential for survival from malaria. PLoS pathogens, 3(7), e96.