

Resource Summary Report

Generated by PRECISE-TBI on Sep 8, 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: 20260905T132711+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 [PRECISE-TBI](#).

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.