

Resource Summary Report

Generated by PRECISE-TBI on Sep 8, 2026

Australian Medical Bioinformatics Resource

RRID:SCR_008385

Type: Tool

Proper Citation

Australian Medical Bioinformatics Resource (RRID:SCR_008385)

Resource Information

URL: <http://www.amber.org.au/>

Proper Citation: Australian Medical Bioinformatics Resource (RRID:SCR_008385)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented September 6, 2016. AMBeR's aim is to bring together Australia's unique resources for genetic epidemiology and genomics with high level expertise in bioinformatics and statistical science, conduct advanced methodological research, develop new research capacity and competitiveness in cutting-edge techniques, bring them to bear on important medical research problems, train young Australians in bioinformatics and advanced biostatistics, and transfer this expertise to the medical research community.

Synonyms: AMBeR

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

Keywords: medical, bioinformatics, statistical, science, genomics, epidemiology, research, methodological, techniques, biostatistics

Funding: Medical Bioinformatics ;
National Health ;
MRC ;
Genomics and Proteomics Program

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: Australian Medical Bioinformatics Resource

Resource ID: SCR_008385

Alternate IDs: nif-0000-30032

Record Creation Time: 20220129T080247+0000

Record Last Update: 20260905T132621+0000

Ratings and Alerts

No rating or validation information has been found for Australian Medical Bioinformatics Resource.

No alerts have been found for Australian Medical Bioinformatics Resource.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Ragan MA, et al. (2008) Genome-scale computational biology and bioinformatics in Australia. PLoS computational biology, 4(8), e1000068.