

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

Annmap

RRID:SCR_011783

Type: Tool

Proper Citation

Annmap (RRID:SCR_011783)

Resource Information

URL: <http://annmap.picr.man.ac.uk/>

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Description: A genome browser that includes mappings between genomic features and Affymetrix microarrays. Associated with annmap is: * a Bioconductor package, annmap that provides programmatic access to the underlying MySQL database tables (which are freely available for download on this site) * xmapbridge, a Bioconductor package that outputs numeric data in a form suitable for presentation in the browser. This is supported by XMapBridge, a Java client that sits on the local desktop and performs the graph rendering for the browser.

Abbreviations: Annmap

Resource Type: database, data or information resource, software resource

Funding: Cancer Research UK ;
Cancer Research UK Manchester Institute

Resource Name: Annmap

Resource ID: SCR_011783

Alternate IDs: OMICS_00900

Record Creation Time: 20220129T080306+0000

Record Last Update: 20260728T043357+0000

Ratings and Alerts

No rating or validation information has been found for Annmap.

No alerts have been found for Annmap.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 6 mentions in open access literature.

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

Memon D, et al. (2019) In silico prediction of housekeeping long intergenic non-coding RNAs reveals HKlincR1 as an essential player in lung cancer cell survival. Scientific reports, 9(1), 7372.

Parras A, et al. (2018) Autism-like phenotype and risk gene mRNA deadenylation by CPEB4 mis-splicing. Nature, 560(7719), 441.

Bazot Q, et al. (2018) Epstein-Barr Virus (EBV) Latent Protein EBNA3A Directly Targets and Silences the STK39 Gene in B Cells Infected by EBV. Journal of virology, 92(7).

Cuvertino S, et al. (2016) SOX7-enforced expression promotes the expansion of adult blood progenitors and blocks B-cell development. Open biology, 6(7).

Hoff AM, et al. (2015) Novel RNA variants in colorectal cancers. Oncotarget, 6(34), 36587.

Bitton DA, et al. (2011) Programmed fluctuations in sense/antisense transcript ratios drive sexual differentiation in *S. pombe*. Molecular systems biology, 7, 559.