

Resource Summary Report

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Canadian Bioinformatics Workshops

RRID:SCR_006774

Type: Tool

Proper Citation

Canadian Bioinformatics Workshops (RRID:SCR_006774)

Resource Information

URL: <http://www.bioinformatics.ca/>

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Description: Offers one and two week short courses in bioinformatics, genomics and proteomics in response to an identified need for a skilled bioinformatics workforce in Canada. For eight years, the series offered short courses in bioinformatics, genomics and proteomics in various cities across Canada. Taught by top faculty from Canada and the US, the courses offered small classes and hands-on instruction. The CBW initiated development of a new format and series of exciting workshops focusing on training the researchers of these advanced technologies on the latest approaches being used in computational biology to deal with the new data. Past workshop content is available under a Creative Commons License.

Abbreviations: CBW

Synonyms: bioinformatics.ca

Resource Type: data or information resource, narrative resource, short course, training material, training resource, workshop

Keywords: education, bioinformatics, course, genomics, proteomics

Funding: Canadian Institutes of Health Research ;
Ontario Institute for Cancer Research ;
Institute of Genetics ;
Genome British Columbia ;
PrioNet Canada ;
MathWorks ;
ActiveState

Availability: Creative Commons License

Resource Name: Canadian Bioinformatics Workshops

Resource ID: SCR_006774

Alternate IDs: nif-0000-10185

Record Creation Time: 20220129T080238+0000

Record Last Update: 20260905T132600+0000

Ratings and Alerts

No rating or validation information has been found for Canadian Bioinformatics Workshops.

No alerts have been found for Canadian Bioinformatics Workshops.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Taj B, et al. (2023) MetaPro: a scalable and reproducible data processing and analysis pipeline for metatranscriptomic investigation of microbial communities. Microbiome, 11(1), 143.

, et al. (2017) hackseq: Catalyzing collaboration between biological and computational scientists via hackathon. F1000Research, 6, 197.

Cuff WR, et al. (2010) A novel interpretation of structural dot plots of genomes derived from the analysis of two strains of Neisseria meningitidis. Genomics, proteomics & bioinformatics, 8(3), 159.