

Resource Summary Report

Generated by NIF on Jul 30, 2026

ProCon - PROteomics CONversion

RRID:SCR_016363

Type: Tool

Proper Citation

ProCon - PROteomics CONversion (RRID:SCR_016363)

Resource Information

URL: <https://www.ruhr-uni-bochum.de/mpc/software/ProCon/index.html.en>

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Description: Java based conversion tool for conversion of data from Proteomics files or a LIMS (Laboratory Information Management System) database into standard formats. Used to support wet-lab scientists in creating proteomics data files ready for upload into the public repositories.

Abbreviations: ProCon

Synonyms: PROteomics CONversion

Resource Type: software resource, software application, data processing software

Defining Citation: [PMID:26182917](#)

Keywords: data, proteomics, conversion, file, laboratory, information, management, system, database, standard, format, , bio.tools

Funding: European Union Projects ProDac ;
European Union Projects ProteomeXchange ;
the German Federal Ministry of Education and Research BMBF

Availability: Free, Available for download, Freely available

Resource Name: ProCon - PROteomics CONversion

Resource ID: SCR_016363

Alternate IDs: biotools:procon

Alternate URLs: <https://bio.tools/procon>

Record Creation Time: 20220129T080330+0000

Record Last Update: 20260731T043009+0000

Ratings and Alerts

No rating or validation information has been found for ProCon - PROteomics CONversion.

No alerts have been found for ProCon - PROteomics CONversion.

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 [NIF](#) .

Parker K, et al. (2023) Invitation methods for Indigenous New Zealand Māori in lung cancer screening: Protocol for a pragmatic cluster randomized controlled trial. PloS one, 18(8), e0281420.