

Resource Summary Report

Generated by [SPARC SAWG](#) on Aug 9, 2026

PRIDE Converter 2

RRID:SCR_012051

Type: Tool

Proper Citation

PRIDE Converter 2 (RRID:SCR_012051)

Resource Information

URL: <https://code.google.com/p/pride-converter-2/>

Proper Citation: PRIDE Converter 2 (RRID:SCR_012051)

Description: Suite of software tools that allows users to convert search result files into PRIDE XML, generate mzTab skeleton files that can be used as a basis to submit quantitative and gel-based MS data, and post-process PRIDE XML files by filtering out contaminants and empty spectra.

Synonyms: PRIDE, PRoteomics IDentification (PRIDE) Converter 2

Resource Type: software resource

Defining Citation: [PMID:22949509](#)

Keywords: standalone software, mac os x, unix/linux, windows, java, xml, mass spectrometry, pride, ols, proteomics, psi, bio.tools

Availability: Apache License, v2

Resource Name: PRIDE Converter 2

Resource ID: SCR_012051

Alternate IDs: biotools:pride_converter, OMICS_03344

Alternate URLs: https://bio.tools/pride_converter

Record Creation Time: 20220129T080308+0000

Record Last Update: 20260808T190005+0000

Ratings and Alerts

No rating or validation information has been found for PRIDE Converter 2.

No alerts have been found for PRIDE Converter 2.

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 [SPARC SAWG](#) .

Monti M, et al. (2026) Integrating interferon gamma receptor pathways, antigenicity, and immune contexture as predictors of immunotherapeutic strategies for mucosal melanomas. Journal for immunotherapy of cancer, 14(3).