

Resource Summary Report

Generated by [RRID](#) on Aug 15, 2026

PRROC

RRID:SCR_026113

Type: Tool

Proper Citation

PRROC (RRID:SCR_026113)

Resource Information

URL: <https://github.com/cran/PRROC>

Proper Citation: PRROC (RRID:SCR_026113)

Description: Software package for computing and visualizing precision-recall and receiver operating characteristic curves in R for weighted and unweighted data.

Synonyms: Precision-Recall and Receiver Operating Characteristic

Resource Type: software toolkit, source code, software resource

Defining Citation: [PMID:25810428](#)

Keywords: Computing and visualizing precision-recall, receiver operating characteristic curves, weighted and unweighted data,

Availability: Free, Available for download, Freely available

Resource Name: PRROC

Resource ID: SCR_026113

Record Creation Time: 20241203T053255+0000

Record Last Update: 20260815T042911+0000

Ratings and Alerts

No rating or validation information has been found for PRROC.

No alerts have been found for PRROC.

Data and Source Information

Usage and Citation Metrics

We found 10 mentions in open access literature.

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

Hassan AZ, et al. (2026) Orobas: A computational approach for scoring and analysis of quantitative chemical-genetic interactions from CRISPR-Cas9 screens. STAR protocols, 7(2), 104594.

Cheng Y, et al. (2025) Investigating the Amyloid-Tau-Neurodegeneration Framework in Alzheimer's Disease Using Semi-Supervised Multimodal Imaging Data Fusion. medRxiv : the preprint server for health sciences.

Vorontsov IE, et al. (2025) Cross-platform motif discovery and benchmarking to explore binding specificities of poorly studied human transcription factors. Communications biology, 8(1), 1545.

Golov AK, et al. (2024) A genome-wide nucleosome-resolution map of promoter-centered interactions in human cells corroborates the enhancer-promoter looping model. eLife, 12.

Devillers J, et al. (2023) Nonlinear SAR Modelling of Mosquito Repellents for Skin Application. Toxics, 11(10).

Lee K, et al. (2022) ASpediaFI: Functional Interaction Analysis of Alternative Splicing Events. Genomics, proteomics & bioinformatics, 20(3), 466.

Marotz C, et al. (2021) SARS-CoV-2 detection status associates with bacterial community composition in patients and the hospital environment. Microbiome, 9(1), 132.

Badet T, et al. (2021) Machine-learning predicts genomic determinants of meiosis-driven structural variation in a eukaryotic pathogen. Nature communications, 12(1), 3551.

Noguchi Y, et al. (2020) Improved Detection Criteria for Detecting Drug-Drug Interaction Signals Using the Proportional Reporting Ratio. Pharmaceuticals (Basel, Switzerland), 14(1).

Li X, et al. (2018) Systematic Analysis and Biomarker Study for Alzheimer's Disease. Scientific reports, 8(1), 17394.