# Resource Summary Report

Generated by SPARC SAWG on Sep 17, 2026

## Apache

RRID:SCR_025967
Type: Tool

### Proper Citation

Apache (RRID:SCR_025967)

### Resource Information

**URL:** https://httpd.apache.org/

**Proper Citation:** Apache (RRID:SCR_025967)

**Description:** Apache HTTP Server Project is collaborative software development effort aimed at creating freely-available source code implementation of HTTP (Web) server. Project is jointly managed by group of volunteers located around the world, using the Internet and the Web to communicate, plan, and develop the server and its related documentation.

**Synonyms:** Apache HTTP Server Project

**Resource Type:** data or information resource, portal, project portal

**Keywords:** collaborative software development, volunteers located around the world,

**Availability:** Free, Freely available

**Resource Name:** Apache

**Resource ID:** SCR_025967

**Record Creation Time:** 20241024T053301+0000

**Record Last Update:** 20260912T200454+0000

### Ratings and Alerts

No rating or validation information has been found for Apache.

No alerts have been found for Apache.

### Data and Source Information

**Source:** SciCrunch Registry

## Usage and Citation Metrics

We found 16 mentions in open access literature.

**Listed below are recent publications.** The full list is available at SPARC SAWG .


Baskaran SP, et al. (2026) sCentInDB: a database of essential oil chemical profiles of Indian medicinal plants. Molecular diversity, 30(1), 789.

Moretti S, et al. (2026) MetaNetX: a bridge between metabolic resources for enhanced curation and multi-omics data harmonization. Nucleic acids research, 54(D1), D617.

Koh H, et al. (2025) MiCML: a causal machine learning cloud platform for the analysis of treatment effects using microbiome profiles. BioData mining, 18(1), 10.

Wang ?? C, et al. (2025) PSIA: A Comprehensive Knowledgebase of Plant Self-incompatibility. Genomics, proteomics & bioinformatics, 23(3).

Karthikeyan SK, et al. (2025) MammOnc-DB, an integrative breast cancer data analysis platform for target discovery. NPJ breast cancer, 11(1), 35.

Ospina OE, et al. (2025) spatialGE Is a User-Friendly Web Application That Facilitates Spatial Transcriptomics Data Analysis. Cancer research, 85(5), 848.

Varambally S, et al. (2024) MammOnc-DB, an integrative breast cancer data analysis platform for target discovery. Research square.

Paik H, et al. (2024) ELiAH: the atlas of E3 ligases in human tissues for targeted protein degradation with reduced off-target effect. Database : the journal of biological databases and curation, 2024.

Sun L, et al. (2024) GEPREP: A comprehensive data atlas of RNA-seq-based gene expression profiles of exercise responses. Journal of sport and health science, 14, 100992.

Kumar B, et al. (2023) Genome-wide identification and characterization of tissue-specific non-coding RNAs in black pepper (Piper nigrum L.). Frontiers in plant science, 14, 1079221.

Kim J, et al. (2023) MiTree: A Unified Web Cloud Analytic Platform for User-Friendly and Interpretable Microbiome Data Mining Using Tree-Based Methods. Microorganisms, 11(11).

Huang CW, et al. (2022) Elegancy: Digitizing the wisdom from laboratories to the cloud with free no-code platform. iScience, 25(8), 104710.

Kaur S, et al. (2022) MGTdb: a web service and database for studying the global and local genomic epidemiology of bacterial pathogens. Database : the journal of biological databases and curation, 2022.

Pike M, et al. (2022) TAFFIES: Tailored Automated Feedback Framework for Developing Integrated and Extensible Feedback Systems. SN computer science, 3(2), 159.

Ras-Carmona A, et al. (2022) Prediction of B cell epitopes in proteins using a novel sequence similarity-based method. Scientific reports, 12(1), 13739.

Vivek-Ananth RP, et al. (2021) MeFSAT: a curated natural product database specific to secondary metabolites of medicinal fungi. RSC advances, 11(5), 2596.