

Resource Summary Report

Generated by SPARC SAWG on Aug 12, 2026

CivicDb

RRID:SCR_028055

Type: Tool

Proper Citation

CivicDb (RRID:SCR_028055)

Resource Information

URL: <https://CivicDb.org>

Proper Citation: CivicDb (RRID:SCR_028055)

Description: Open-access, community-driven knowledgebase designed to crowdsource and curate evidence on the clinical significance of cancer-related genomic variants. It helps researchers and clinicians interpret tumor DNA mutations to guide precision medicine.

Synonyms: , Clinical Interpretation of Variants in Cancer, CIViCdb

Resource Type: data or information resource, knowledge base, database

Defining Citation: [PMID:36373660](#)

Keywords: clinical significance of cancer-related genomic variants, cancer-related genomic variants, genomic variants,

Funding: NCI U01CA209936;
NCI U24CA237719;
NCATS UL1TR002345;
NHGRI U41 HG008735;
NHGRI R00HG010157;
NHGRI R00HG007940;
NCI K22CA188163

Availability: Free, Freely available,

Resource Name: CivicDb

Resource ID: SCR_028055

Record Creation Time: 20260306T054431+0000

Record Last Update: 20260812T044537+0000

Ratings and Alerts

No rating or validation information has been found for CivicDb.

No alerts have been found for CivicDb.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

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

Ritter DI, et al. (2025) Multi-Platform Curation in the Development of ACMG/AMP Specifications for Von Hippel Lindau (VHL) Disease. medRxiv : the preprint server for health sciences.

Choochuen P, et al. (2024) Discovery of Novel Potential Prognostic Markers and Targeted Therapy to Overcome Chemotherapy Resistance in an Advanced-Stage Wilms Tumor. Cancers, 16(8).

Pires SF, et al. (2023) Analysis of the Mutational Landscape of Osteosarcomas Identifies Genes Related to Metastasis and Prognosis and Disrupted Biological Pathways of Immune Response and Bone Development. International journal of molecular sciences, 24(13).

Gu W, et al. (2022) MarkerGenie: an NLP-enabled text-mining system for biomedical entity relation extraction. Bioinformatics advances, 2(1), vbac035.

Akolkar D, et al. (2021) Development and validation of a multigene variant profiling assay to guide targeted and immuno therapy selection in solid tumors. PloS one, 16(2), e0246048.