

Resource Summary Report

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

ClusterSV

RRID:SCR_027722

Type: Tool

Proper Citation

ClusterSV (RRID:SCR_027722)

Resource Information

URL: <https://github.com/cancerit/ClusterSV>

Proper Citation: ClusterSV (RRID:SCR_027722)

Description: Software application to group genetic structural variant rearrangements into rearrangement clusters and footprints. Used in the PCAWG-6 project.

Resource Type: source code, software application, software resource

Keywords: structural variant, rearrangement region, group genetic structural variant rearrangements, rearrangement clusters and footprints,

Availability: Free, Available for download, Freely available

Resource Name: ClusterSV

Resource ID: SCR_027722

License: AGPL-3.0 license

Record Creation Time: 20251209T055452+0000

Record Last Update: 20260815T183307+0000

Ratings and Alerts

No rating or validation information has been found for ClusterSV.

No alerts have been found for ClusterSV.

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 [RRID](#) .

Culliford R, et al. (2026) Contrasting Features of Papillary and Chromophobe Renal Cell Carcinoma Revealed by Whole-Genome Sequencing. Molecular cancer research : MCR, 24(5), 327.

Lee JJ, et al. (2026) Evolution of oncogene amplification across 86,000 cancer cell genomes. bioRxiv : the preprint server for biology.

Yang Y, et al. (2025) Panorama of Chromosomal Instability in Lung Cancer. medRxiv : the preprint server for health sciences.

Kim S, et al. (2024) A Subset of Microsatellite Unstable Cancer Genomes Prone to Short Insertions over Deletions Is Associated with Elevated Anticancer Immunity. Genes, 15(6).

Ng AWT, et al. (2022) Rearrangement processes and structural variations show evidence of selection in oesophageal adenocarcinomas. Communications biology, 5(1), 335.