DELLY
RRID:SCR_004603
Type: Tool

Proper Citation

DELLY (RRID:SCR_004603)

Resource Information

**URL:** [https://tobiasrausch.com/delly/](https://tobiasrausch.com/delly/)

**Description:** Integrated structural variant prediction software that can detect deletions, tandem duplications, inversions and translocations at single-nucleotide resolution in short-read massively parallel sequencing data. It uses paired-ends and split-reads to sensitively and accurately delineate genomic rearrangements throughout genome.

**Resource Name:** DELLY

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**Resource Type:** Resource, software resource

**Keywords:** structural variant, genomic rearrangement, deletion, tandem duplication, inversion, translocation

**Resource ID:** SCR_004603

**Parent Organization:** European Molecular Biology Laboratory

**References:** [PMID:22962449](https://www.ncbi.nlm.nih.gov/pubmed/22962449)

**Website Status:** Last checked down

**Alternate IDs:** OMICS_00313

**Alternate URLs:** [https://github.com/dellytools/delly/](https://github.com/dellytools/delly/)

**Abbreviations:** DELLY
Ratings and Alerts

No rating or validation information has been found for DELLY.

No alerts have been found for DELLY.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 124 mentions in open access literature.

**Listed below are recent publications.** The full list is available at FDI Lab - SciCrunch Infrastructure.


