**DELLY**

**RRID:** SCR_004603

**Type:** Tool

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**Proper Citation**

DELLY (RRID:SCR_004603)

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**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
- **Proper Citation:** DELLY (RRID:SCR_004603)
- **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
- **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.


