

# Resource Summary Report

Generated by [FDI Lab - SciCrunch.org](https://fdi-lab.sci-crunch.org) on Apr 27, 2025

## Inference of CRISPR Edits

RRID:SCR\_024508

Type: Tool

### Proper Citation

Inference of CRISPR Edits (RRID:SCR\_024508)

### Resource Information

**URL:** <https://ice.synthego.com/#/>

**Proper Citation:** Inference of CRISPR Edits (RRID:SCR\_024508)

**Description:** Software tool that offers analysis of CRISPR editing data. Used for inference of CRISPR edits from Sanger trace data.

**Abbreviations:** ICE

**Synonyms:** Synthego Inference of CRISPR Edits

**Resource Type:** service resource, production service resource, software resource, analysis service resource

**Defining Citation:** [PMID:35119294](https://pubmed.ncbi.nlm.nih.gov/35119294/)

**Keywords:** analysis of CRISPR editing data, CRISPR editing analysis,

**Funding:**

**Availability:** Free, Freely available

**Resource Name:** Inference of CRISPR Edits

**Resource ID:** SCR\_024508

**Alternate URLs:** <https://github.com/synthego-open/ice#ref2>

**Record Creation Time:** 20231002T161336+0000

**Record Last Update:** 20250426T061016+0000

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## Ratings and Alerts

No rating or validation information has been found for Inference of CRISPR Edits.

No alerts have been found for Inference of CRISPR Edits.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 40 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [FDI Lab - SciCrunch.org](#).

Blanco E, et al. (2025) Dominant negative variants in ITPR3 impair T cell Ca<sup>2+</sup> dynamics causing combined immunodeficiency. *The Journal of experimental medicine*, 222(1).

Nurmi K, et al. (2024) Truncating NFKB1 variants cause combined NLRP3 inflammasome activation and type I interferon signaling and predispose to necrotizing fasciitis. *Cell reports. Medicine*, 5(4), 101503.

Jové V, et al. (2024) Type I interferon regulation by USP18 is a key vulnerability in cancer. *iScience*, 27(4), 109593.

LaCoursiere CM, et al. (2024) Zebrafish models of candidate human epilepsy-associated genes provide evidence of hyperexcitability. *iScience*, 27(7), 110172.

Yang Q, et al. (2024) T4 DNA polymerase prevents deleterious on-target DNA damage and enhances precise CRISPR editing. *The EMBO journal*, 43(17), 3733.

Jansen RA, et al. (2024) Small-molecule inhibition of MAP2K4 is synergistic with RAS inhibitors in KRAS-mutant cancers. *Proceedings of the National Academy of Sciences of the United States of America*, 121(9), e2319492121.

Janecki DM, et al. (2024) LINE-1 mRNA 3' end dynamics shape its biology and retrotransposition potential. *Nucleic acids research*.

Lu J, et al. (2024) RAB18 regulates extrahepatic siRNA-mediated gene silencing efficacy. *Molecular therapy. Nucleic acids*, 35(4), 102335.

Tudorica DA, et al. (2024) A RAB7A phosphoswitch coordinates Rubicon Homology protein regulation of Parkin-dependent mitophagy. *The Journal of cell biology*, 223(7).

Choy L, et al. (2024) Inhibition of Aurora Kinase Induces Endogenous Retroelements to Induce a Type I/III IFN Response via RIG-I. *Cancer research communications*, 4(2), 540.

Yashar WM, et al. (2024) Predicting transcription factor activity using prior biological information. *iScience*, 27(3), 109124.

Schlapp G, et al. (2024) Zygote cryobanking applied to CRISPR/Cas9 microinjection in mice. *PloS one*, 19(7), e0306617.

Shin JH, et al. (2024) Enhanced efficacy of glycoengineered rice cell-produced trastuzumab. *Plant biotechnology journal*, 22(11), 3068.

Wardlaw CP, et al. (2024) Protocol for the quantitative identification of endogenously ISGylated proteins from mammalian cell lines. *STAR protocols*, 5(1), 102843.

Carrillo-Gálvez AB, et al. (2024) NLRP3 and AIM2 inflammasomes expression is modified by LPS and titanium ions increasing the release of active IL-1 $\beta$  in alveolar bone-derived MSCs. *Stem cells translational medicine*, 13(8), 826.

Campbell CA, et al. (2024) p65 signaling dynamics drive the developmental progression of hematopoietic stem and progenitor cells through cell cycle regulation. *Nature communications*, 15(1), 7787.

Naigles B, et al. (2024) Genetic Knock-Ins of Endogenous Fluorescent Tags in RAW 264.7 Murine Macrophages Using CRISPR/Cas9 Genome Editing. *Bio-protocol*, 14(6), e4960.

Dhaliwal NK, et al. (2024) Protocol for the efficient and inducible generation of CRISPR-Cas9-edited human cortical neurons from the iCas9-iNgn2 hPSCs. *STAR protocols*, 5(4), 103352.

Ugur B, et al. (2024) VPS13B is localized at the interface between Golgi cisternae and is a functional partner of FAM177A1. *The Journal of cell biology*, 223(12).

Chen CY, et al. (2024) Rescue of the endogenous FVIII expression in hemophilia A mice using CRISPR-Cas9 mRNA LNPs. *Molecular therapy. Nucleic acids*, 35(4), 102383.