

Resource Summary Report

Generated by [FDI Lab - SciCrunch.org](#) on May 13, 2025

Stitchr

RRID:SCR_022139

Type: Tool

Proper Citation

Stitchr (RRID:SCR_022139)

Resource Information

URL: <https://github.com/JamieHeather/stitchr>

Proper Citation: Stitchr (RRID:SCR_022139)

Description: Software Python tool for stitching coding T cell receptors nucleotide sequences from V,J,CDR3 information. Produces complete coding sequences representing fully spliced TCR cDNA given minimal V,J,CDR3 information.

Resource Type: data processing software, software application, software resource, sequence analysis software, data analysis software

Defining Citation: [PMID:35325179](#)

Keywords: Stitch together coding TCR nucleotide sequences, Python, T cell receptors nucleotide, V and J gene symbols, hypervariable CDR3 amino acid sequence, fully spliced TCR cDNA

Funding: NCI R01 CA164273;
NIAID R43 AI120313;
NCI R43 CA232942;
Emily Venanzi Fund

Availability: Free, Available for download, Freely available

Resource Name: Stitchr

Resource ID: SCR_022139

License: BSD 3-Clause "New" or "Revised" License

Record Creation Time: 20220421T050138+0000

Record Last Update: 20250513T062233+0000

Ratings and Alerts

No rating or validation information has been found for Stitchr.

No alerts have been found for Stitchr.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Kirk AM, et al. (2024) DNAJB1-PRKACA fusion neoantigens elicit rare endogenous T cell responses that potentiate cell therapy for fibrolamellar carcinoma. Cell reports. Medicine, 5(3), 101469.

Heather JM, et al. (2022) Stitchr: stitching coding TCR nucleotide sequences from V/J/CDR3 information. Nucleic acids research, 50(12), e68.