

Resource Summary Report

Generated by [RRID](#) on Sep 18, 2026

T Cell ExTRECT

RRID:SCR_027742

Type: Tool

Proper Citation

T Cell ExTRECT (RRID:SCR_027742)

Resource Information

URL: <https://github.com/McGranahanLab/TcellExTRECT>

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Description: Software R package to calculate T cell fractions from WES data from hg19 or hg38 aligned genomes.

Synonyms: , T cell exome TREC, T cell exome T cell Receptor Excision Circle

Resource Type: software resource, software toolkit, source code

Defining Citation: [PMID:34497419](#)

Keywords: T-cell, T cell receptor excision circle, WES data, hg19 or hg38 aligned genomes,

Funding: NCATS UL1TR000100;
NCI P30CA023100;
NCI R21CA177519;
NCI U01CA196406;
NHLBI U54HL108460;
NIH Office of the Director DP5OD017937;
NLM T15LM011271;
NSF

Availability: Free, Available for download, Freely available

Resource Name: T Cell ExTRECT

Resource ID: SCR_027742

License URLs: <https://github.com/McGranahanLab/TcellExTRECT/blob/main/LICENSE>

Record Creation Time: 20251209T055453+0000

Record Last Update: 20260912T200535+0000

Ratings and Alerts

No rating or validation information has been found for T Cell ExTRECT.

No alerts have been found for T Cell ExTRECT.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 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.