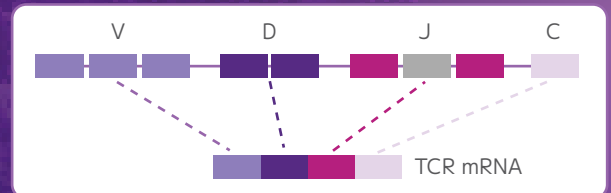


Evercode TCR

T Cell Receptor + Whole Transcriptome

More insights with more scalable single cell TCR profiling

T cell receptors (TCRs) are formed through recombination of variable (V), diversity (D), and joining (J) segments, generating the enormous range of receptors that powers adaptive immunity. Evercode™ TCR is a fixation-based single cell assay that captures paired alpha-beta TCR sequences together with whole transcriptomes—delivering unmatched scale, sensitivity, and flexibility for a complete view of T cell biology.



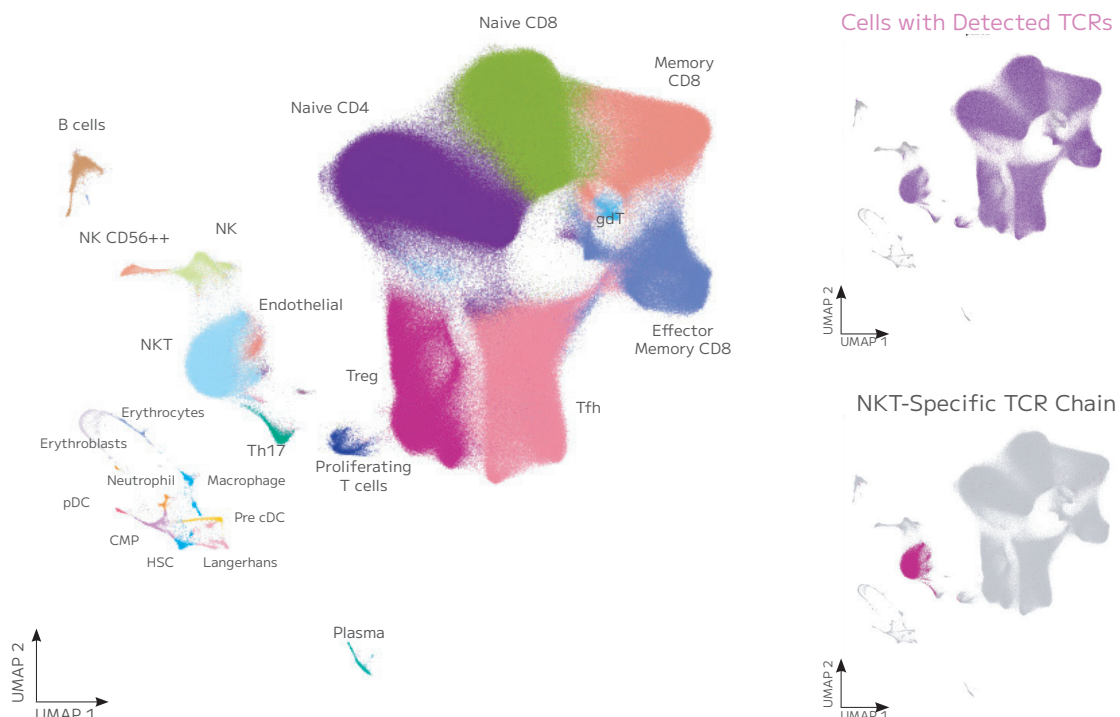
TCR Repertoire and Transcriptome in One Assay

Precise Paired-Chain Resolution

Instrument-Free, Fixation Powered Workflow

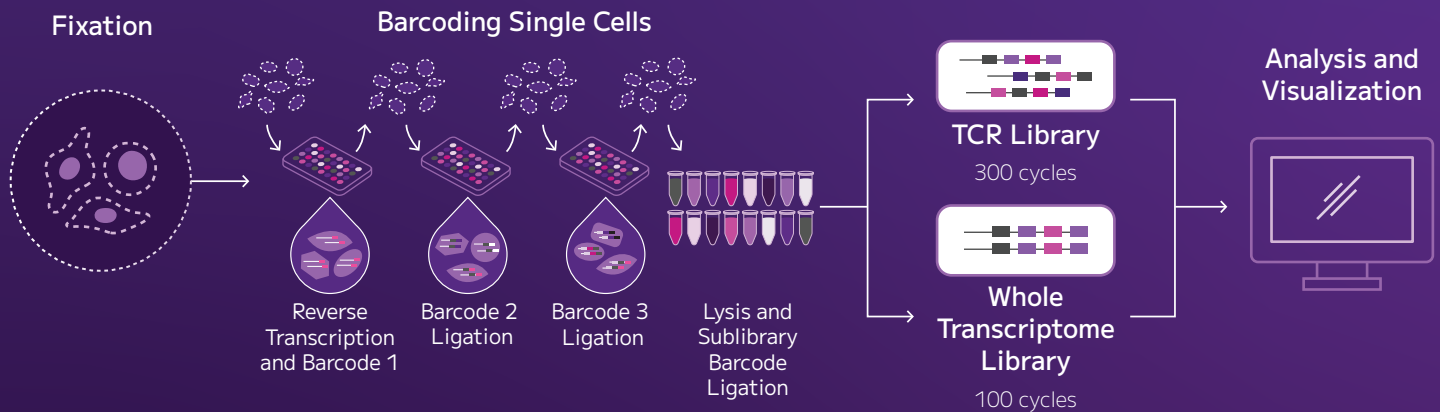
Sensitive Clonotype Detection

Go Beyond the TCR Alone and Discover the Cell Behind It



Nearly 1 million mouse T cells were profiled with Evercode TCR Mega, resolving diverse T cell populations and confirming that TCRs appear only in true T cells. An NKT-specific alpha chain was restricted to the NKT cluster, underscoring the power of Evercode TCR to place receptors in their full cellular context.

Instrument-free, fixation-powered workflow

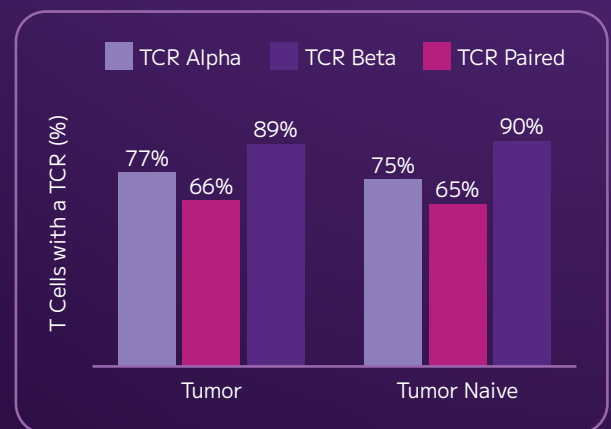


High-confidence TCR pairing in real customer data

"Evercode TCR's simple protocols and impressive throughput allowed us to make new discoveries about T cell responses to metastatic cancers."

Nathan Reticker-Flynn, PhD
Assistant Professor, Stanford University

Evercode TCR achieved up to 65% alpha-beta pairing in isolated T cells from tumor and tumor-naïve samples in real customer data. In a study by the Reticker-Flynn Lab (Stanford University) using a mouse transplant model of lymph node metastasis, Evercode TCR enabled reliable insights into TCR repertoire changes during tumor progression.



Flexible configurations to match your study scale in human or mouse samples

EVERCODE TCR

10K-100K cells,
1-48 samples

EVERCODE TCR MEGA

100K-1M cells,
1-96 samples



For research use only. Not for use in diagnostic procedures.

Evercode TCR Product Sheet - Version 1.1
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