Multiplexed antigen screening for rapid B- and T-cell clonotype discovery

Single Cell Immune Profiling with Barcode Enabled Antigen Mapping (BEAM)

Despite the huge therapeutic potential of the functional adaptive immune repertoire, it remains difficult to match antigen targets with specific B- and T-cell receptors (BCRs/TCRs) using conventional methods. BEAM empowers rapid discovery of antigen-specific B-cell (BEAM-Ab) and T-cell (BEAM-T) clonotypes with unparalleled cellular resolution. Built on the proven Chromium Single Cell Immune Profiling workflow, BEAM enables screening of BCRs/TCRs against putative antigens in conjunction with gene expression, V(D)J sequencing, and cell surface protein expression. Leverage multiplexed antigen screening to generate tens to hundreds of high-quality hits from the same sample, augmenting the diversity and magnitude of clonotypes/receptors for downstream validation in therapeutic applications.

Highlights

- Generate tens to hundreds of high-quality antigen-specific hits from the same sample
- Discover antigen-specific clonotypes with full-length, paired V(D)J sequences, gene expression, and surface protein expression from the same single cell
- Go from sample to antigen-specific immune repertoire sequencing data in a week
- Explore and interpret antigen-clonotype relationships with easy-to-use analysis and visualization software

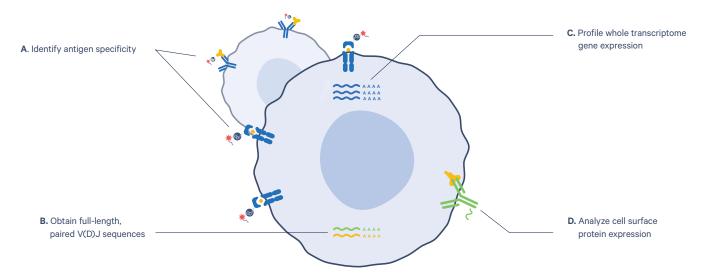


Figure 1. Comprehensive profiling of the immune repertoire with antigen-specific clonotype characterization. A. Feature Barcode technology with unique BEAM-Ab and BEAM-T reagent assemblies allows antigen-specificity determination of target B and T cells, respectively. B. Obtain full-length, paired V(D)J sequences of antigen-specific B or T cells alongside whole transcriptome gene expression (C) and cell surface proteins (D) at the single cell level.



Product features

BEAM-Ab

- Utilize a seamlessly integrated, end-to-end workflow to go from sample to high-quality antibody hits in under a week
- Screen as many as 15 antigens in parallel to generate tens to hundreds of hits per sample and increase your funnel size for antibody candidates
- Leverage intuitive software solutions to identify expanded antigen-specific clonotypes for efficient selection of the ideal antibody candidates for downstream validation

BEAM-T

- Leverage kitted, custom loadable MHC monomers to easily prepare peptide–MHC complexes, offering flexibility to design and source peptides from any vendor
- Detect rare and/or therapeutically relevant TCR clonotypes with gene expression at single cell resolution for a comprehensive understanding of the immune response
- Explore antigen-clonotype relationships with easyto-use software solutions to access the full diversity of antigen-reactive T cells from a single sample

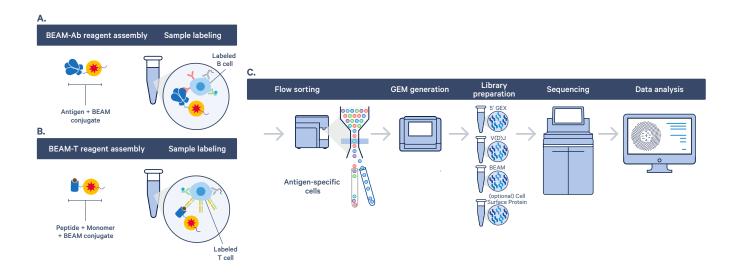


Figure 2. Fast and robust workflow for antigen-specific BCR and TCR discovery. The BEAM-Ab and BEAM-T workflows begin with user-supplied antigens, which are first assembled and uniquely barcoded using 10x Genomics BEAM reagents. A. BEAM-Ab reagent assemblies couple biotinylated antigen to a barcoded BEAM Conjugate complex that contains PE fluorescent marker. B. BEAM-T reagent assemblies couple antigenic peptide to empty loadable MHC, which are themselves coupled to BEAM conjugate. BEAM-T offers flexibility to design peptides of interest and choose from five different MHC Class I alleles provided through 10x Genomics Chromium Human/Mouse MHC Class I Monomer Kits. C. The barcoded assemblies are then used to stain B or T cells prior to flow sorting for enrichment (flow sorting is recommended). Normal library preparation steps are then carried out on the sample, generating multiple library types, including antigen, V(D)J, gene expression, and, optionally, cell surface protein. After sequencing, raw FASTQ files are processed for downstream analysis with Cell Ranger and Loupe V(D)J Browser, our fully integrated and easy-to-use analysis and visualization software tools. (Note: BEAM-Ab and BEAM-T are performed in separate workflows with unique reagents).

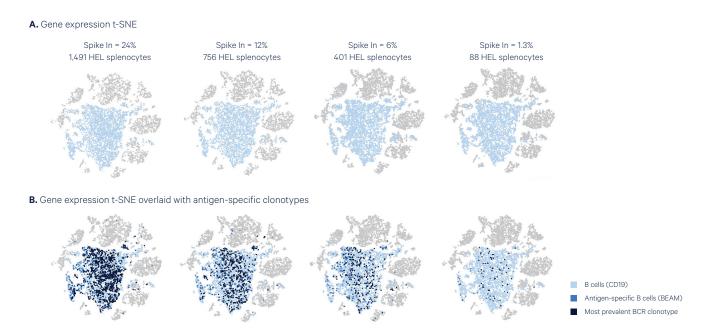


Figure 3. BEAM-Ab provides a precise view of antigen-specific, expanded clonotypes. Transgenic mouse splenocytes that recognize the foreign antigen hen egg lysozyme (HEL) were labeled with a 2-plex BEAM-Ab reagent assembly of HEL and a negative control. These cells were flow sorted to purity and then spiked into a background of mouse splenocytes at the following percentages: 24%, 12%, 6%, and 1.3%. A. Aggregated t-SNE projections show splenocyte cells clustered by gene expression. Light blue dots indicate a cell expressing CD19 (a marker for B cells), while gray dots indicate non-B-cell populations clustered via 5′ gene expression profiles. B. Antigen-specific B cells, identified by the presence of a BEAM Conjugate, were overlaid with gene expression and cells from the most prevalent BCR clonotypes in the B-cell cluster. As shown, the most prevalent BCR clonotypes were found to be specific for the HEL antigen.

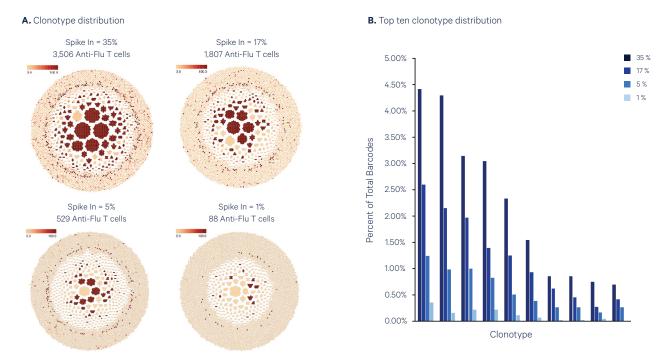


Figure 4. Highly sensitive, consistent clonotype-antigen assignments with BEAM-T. HLA-A*02:01 anti-flu expanded human T cells were labeled with a 2-plex BEAM-T reagent assembly of FLU and the HLA-A*02:01 negative control. These cells were flow sorted to purity and then spiked into a background of HLA-A*02:01 peripheral blood mononuclear cells (PBMCs) at the following percentages: 35%, 17%, 5%, and 1%. A. Visualization of anti-gen-specific T cells identified via BEAM-T. Cells are represented as individual points and grouped by clonotype assignment. Cells are colored according to their antigen specificity, where dark red denotes higher antigen specificity. B. Top ten TCR clonotype frequency distribution demonstrates consistent clonotype detection and abundance consistent with the spike-in percentage.

Product specifications

- Multiplex up to 15 antigens per sample plus a negative control
- Diverse sample type compatibility, including PBMCs, splenocytes, lymph node aspirates, and enriched B or T cells
- Antigen specificity score computed for screened antigens against controls
- Low doublet rates of under 0.8% per 1,000 cells
- High cell capture rates of up to 65%

Gene expression repertoire prof	on and / or immune iling	Product code
Chromium Next GEM Single Cell 5' Kit v2, 16 rxns		1000263
Chromium Next GEM Single Cell 5' Kit v2, 4 rxns		1000265
Chromium Next GEM Chip K Single Cell Kit, 48 rxns		1000286
Chromium Next GEM Chip K Single Cell Kit, 16 rxns		1000287
Library Construction Kit, 16 rxns		1000190
Dual Index Kit TT Set A, 96-rxns		1000215
Dual Index Kit TS Set A, 96-rxns		1000251
V(D)J amplification kits		Product code
Human T cell	Chromium Single Cell Human TCR Amplification Kit, 16 rxns	1000252
Human T cell	<u> </u>	1000252
	TCR Amplification Kit, 16 rxns Chromium Single Cell Human	
Human B cell	TCR Amplification Kit, 16 rxns Chromium Single Cell Human BCR Amplification Kit, 16 rxns Chromium Single Cell Mouse	1000253
Human B cell Mouse T cell Mouse B cell	TCR Amplification Kit, 16 rxns Chromium Single Cell Human BCR Amplification Kit, 16 rxns Chromium Single Cell Mouse TCR Amplification Kit, 16 rxns Chromium Single Cell Mouse	1000253
Human B cell Mouse T cell Mouse B cell	TCR Amplification Kit, 16 rxns Chromium Single Cell Human BCR Amplification Kit, 16 rxns Chromium Single Cell Mouse TCR Amplification Kit, 16 rxns Chromium Single Cell Mouse BCR Amplification Kit, 16 rxns	1000253 1000254 1000255

BEAM-Ab reagents	Product code
Chromium Single Cell 5' BEAM Core Kit, PE, Set A 128 rxns	1000539
BEAM-T reagents	Product code
Chromium Single Cell 5' BEAM Core Kit, PE, Set A 128 rxns	1000539
Chromium Human MHC Class I A0201 Monomer Kit, 32 rxns	1000542
Chromium Human MHC Class I A1101 Monomer Kit, 32 rxns	1000543
Chromium Human MHC Class I B0702 Monomer Kit, 32 rxns	1000544
Chromium Human MHC Class I A2402 Monomer Kit, 32 rxns	1000545
Chromium Mouse MHC Class I H2Kb Monomer Kit, 32 rxns	1000546
Instrument compatibility	Product code
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Chromium iX & Accessory Kit, 12 Mo. Warranty	1000328
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