

SUPPLEMENTAL FIGURES AND TABLE FOR

Measurements of human CD8+ T-cell functions using high-throughput integrated single-cell analysis

Navin Varadarajan^{1,2*}, Boris Julg^{3,*}, Yvonne J. Yamanaka⁴, Huabiao Chen^{3,5},
Adebola O. Ogunniyi¹, Elizabeth McAndrew³, Lindsay C. Porter³, Alicja Piechocka-Trocha^{3,5},
Brenna J. Hill⁶, Daniel C. Douek⁶, Florencia Pereyra³,
Bruce D. Walker^{3,5}, and J. Christopher Love^{1,3,7†}

¹*Department of Chemical Engineering, Massachusetts Institute of Technology,
77 Massachusetts Ave., Cambridge, MA 02139, USA*

²*Current address: Department of Chemical & Biomolecular Engineering,
University of Houston, Houston, TX 77204, USA*

³*The Ragon Institute of MGH, MIT, and Harvard
Charlestown Navy Yard, Boston, MA 02129, USA*

⁴*Department of Biological Engineering, Massachusetts Institute of Technology,
77 Massachusetts Ave., Cambridge, MA 02139, USA*

⁵*Howard Hughes Medical Institute, Chevy Chase, MD 20815, USA*

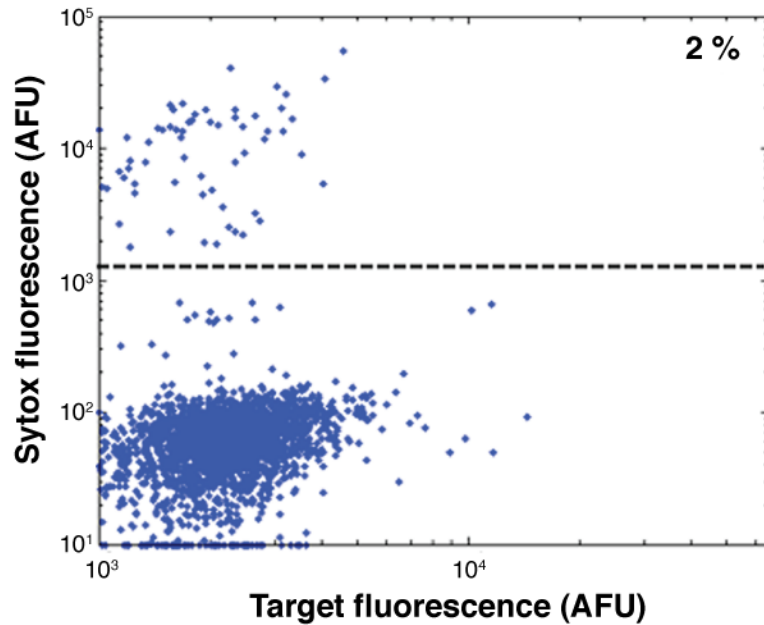
⁶*Human Immunology Section, Vaccine Research Center, National Institutes of Allergy and
Infectious Diseases, National Institutes of Health, Bethesda, MD 20892, USA*

⁷*The Eli and Edythe L. Broad Institute, Seven Cambridge Center, Cambridge, MA 02142, USA*

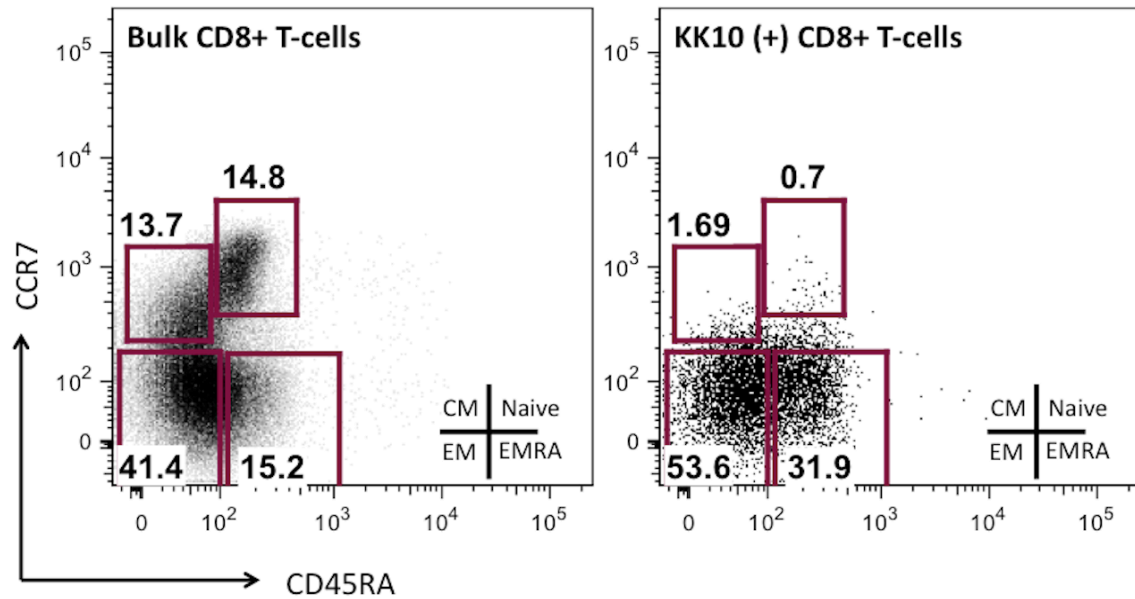
* These authors contributed equally to this work.

		Targets per well					
		1	2	3	4	5+	
Effectors per well	1	5437	2748	1067	352	117	E:T = 1:n 9,721
	2	1253	517	123	36	4	
	3	232	78	23	6	0	
	4	45	8	0	0	0	
	5+	9	0	4	1	0	
		E:T = n:1 6,796					

Supplemental Figure 1. Distribution of effectors and targets in individual 50 μm x 50 μm x 50 μm microwells. A density matrix indicating the number of wells containing at least a single B-cell target and a single CD8+ T-cell. The arrays comprise 84,672 microwells in total. The conditions for loading the arrays were chosen to maximize the number of wells occupied with E:T=1:1. These data are representative of typical distributions measured; data shown here are from a clinical sample (CTR40).



Supplemental Figure 2. Spontaneous loss of viability for target B-cells. Scatterplot of the percentage of dead (SYTOX) KK10-loaded, labeled B-cell targets observed in arrays of microwells in the absence of effector cells (B27-restricted T cell clones). The dashed line indicates the threshold for dead cells. These data are representative of at least three independent experiments.

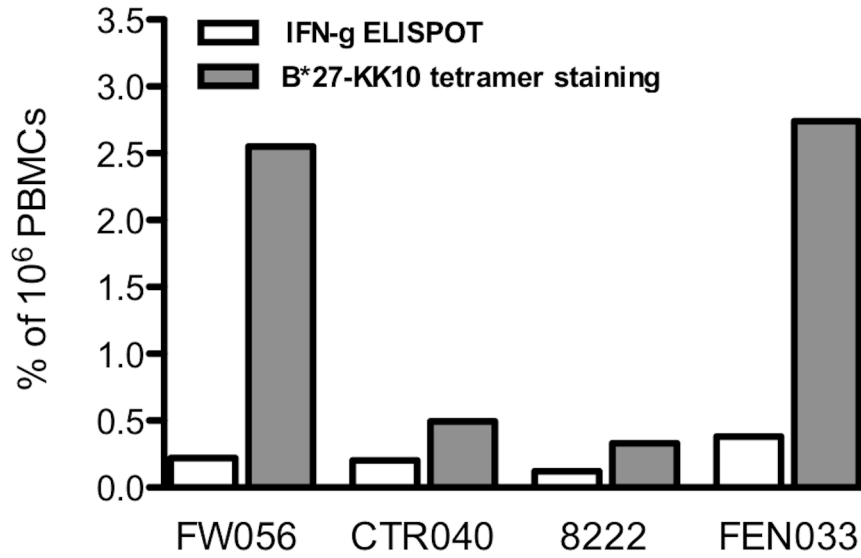


Supplemental Figure 3. Differentiated phenotypes of bulk and KK10 tetramer (+) CD8+ T-cells from Fw056 determined by flow cytometry. The state of differentiation was assessed by the surface expression of CCR7 and CD45RA, measured using fluorescent antibodies directed against these proteins and profiled on the flow-cytometer: EM = effector-memory (CD45RA⁻/CCR7⁻); CM = central-memory (CD45RA⁻/CCR7⁺); N = Naïve (CD45RA⁺/CCR7⁺); EMRA = effector-memory RA (CD45RA⁺/CCR7⁻). The relative percentages of each are indicated.

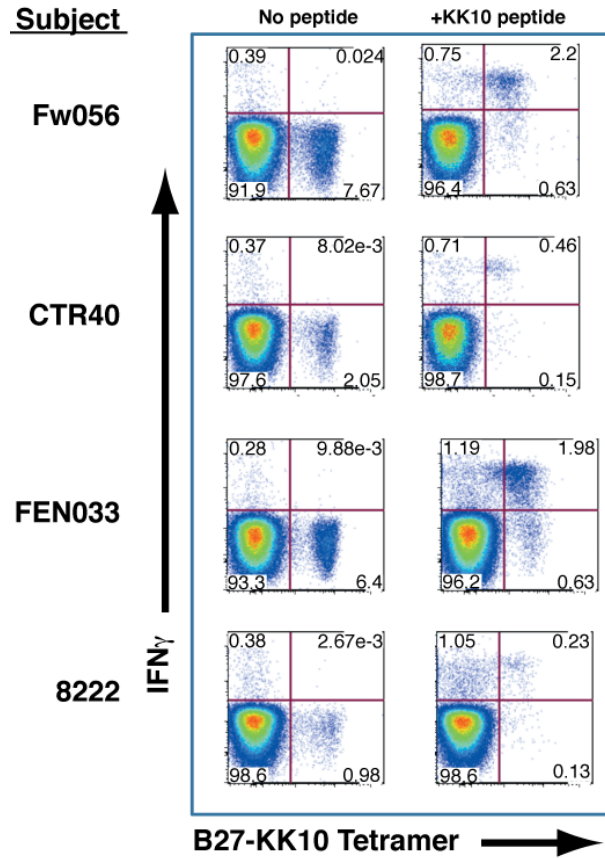
Supplemental Table 1. Clonotypic analysis of three B27-KK10-specific T-cell clones generated by screening cytolytic activity.

Clone	V region	CDR3 Sequence	J region
FW056 #2	TRBV6-5*01	CASRPGQGATEAFFGQGTRLTVV	TRBJ1-1
	TRAV30	AGRTAVLPR_SLDQGP	N/A
FW056 #3	TRBV20-1*01	CSARDRGTREVADNYGYTFGSGTRLTVV	TRBJ1-2
	TRBV15*01	CATSETGTTLEQYFGPGTRLTVT	TRBJ2-7
	TRAV14/DV4	CAMREVSSGNTGKLIFGQGTTLQVKP	TRAJ37
FW056 #4	TRBV4-3*01	CASRPGLASNEQFFGPGTRLTVL	TRBJ2-1
	TRAV14/DV4	CAMSPATARQLTFGSGTQLTVLP	TRAJ22
	TRAV8-2	CPSYRDDKIIFGKGTRLHILP	TRAJ30

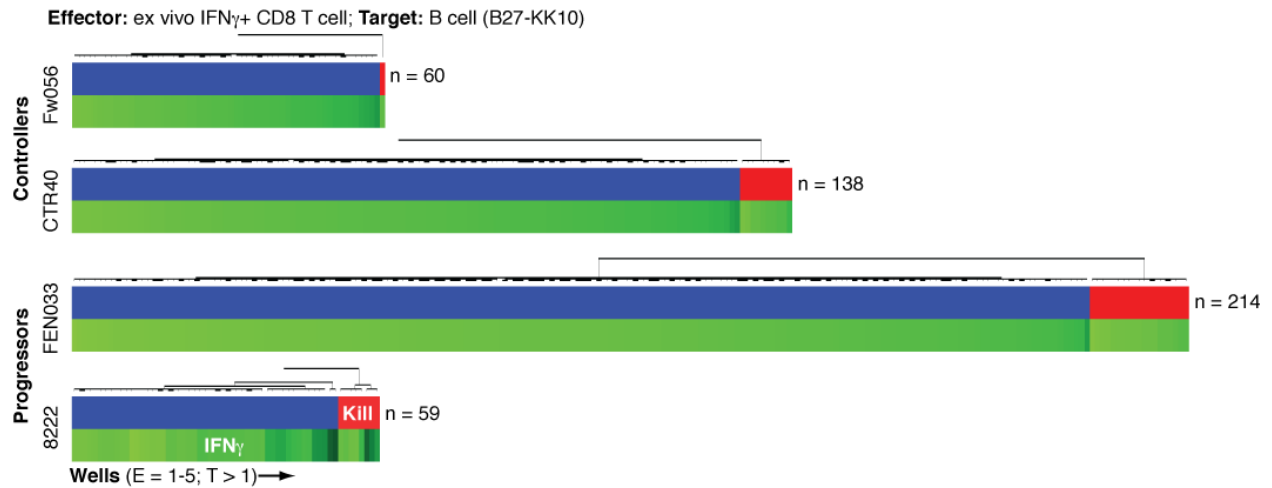
_ = not determined



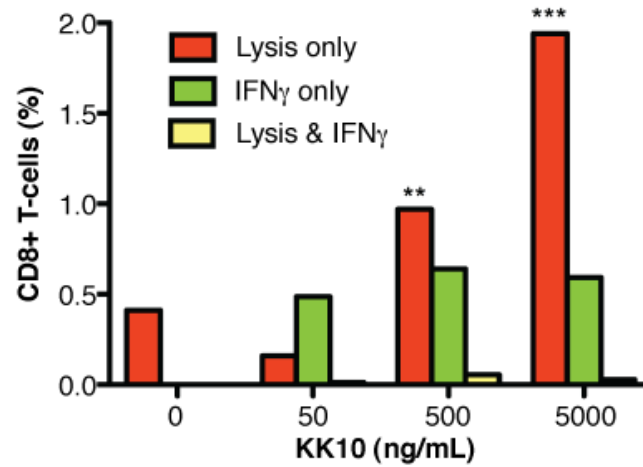
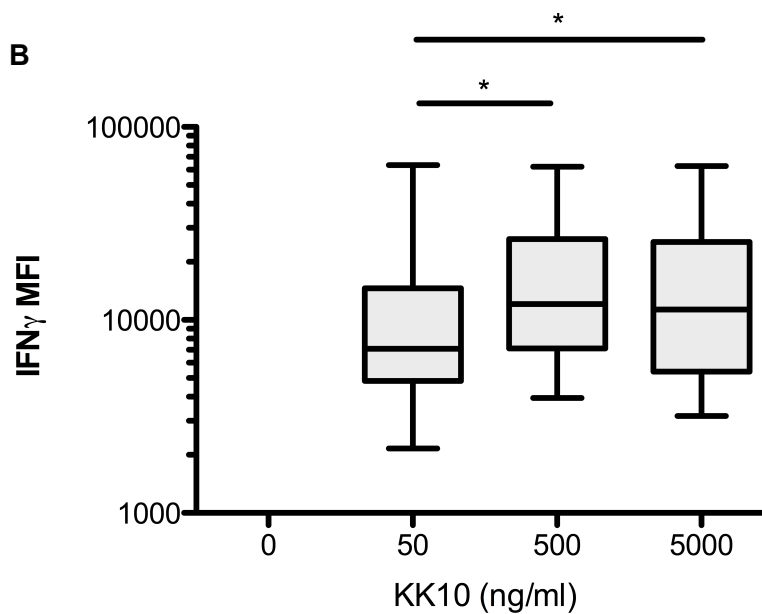
Supplemental Figure 4. Comparison of the frequencies of PBMCs specific for the HLA-B*27-restricted epitope KK10 measured by IFN γ ELISpot and CD8⁺ T-cells binding B27-KK10 tetramers measured by flow cytometry for four HIV-infected subjects.



Supplemental Figure 5. Analysis of intracellular IFN γ in KK10 tetramer (+) CD8 $^+$ T-cells of 2 elite controllers and 2 viremic progressors upon stimulation with the cognate peptide. CD8 $^+$ T-cells were stained with fluorescently labeled tetramers and incubated for 5 h in the presence or absence of KK10 peptide, antibodies against the co-stimulatory molecules CD49d and CD28, and the degranulation-inhibitors Brefeldin A and Monensin.

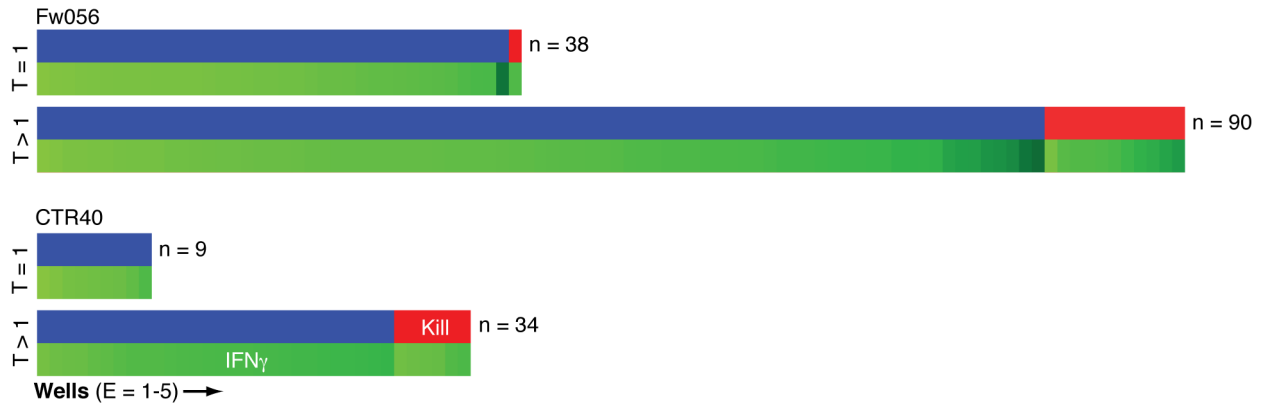


Supplemental Figure 6. Matched data for concurrent measurement of cytolysis and IFN γ secretion for ex vivo IFN γ -secreting CD8+ T-cells from four HIV+ individuals co-incubated in arrays of microwells for 6 h with multiple HLA-matched, KK10-pulsed target B-cells. The data are shown as a heatmap and organized by unsupervised hierarchical clustering (Euclidean distance, complete linkage). Cytolytic activity is shown as discrete values (blue = negative; red = death). The relative rate of secretion is shown for IFN γ (green). The dendrograms indicate the relative calculated distances and separation among clusters of wells. The number of wells scored with E = 1-5 and T > 1 with detectable IFN γ are indicated for each sample. There were no cells among any of the four samples that released either IL-2 or TNF α at a detectable rate (limit of detection: \sim 2 molecules/s).

A**B**

Supplemental Figure 7. (A) Functional response of ex vivo CD8+ T-cells (HIV+ controller) to HLA-matched B-cell targets loaded with different doses of KK10 peptide (E = 1-5 : T = 1). The lysis rate is corrected for the background death rate of target cells on the same microwell array (E = 0 : T = 1). **p < 0.01, ***p < 0.0001 for lysis rate comparison to both on-chip background death rate and 0 ng/ml KK10 killing rate (Fisher's exact test, or, for large numbers of events, chi-squared with Yate's correction). The minimum number of events scored for each condition was $n = 3,192$. (B) Boxplot of the relative rates of secretion of IFN γ as a function of KK10 concentration (secretors only; E = 1-5 : T = 1). *p < 0.05, two-tailed t-test.

Effector: ex vivo IFN γ + CD8+ T-cell; **Target:** B cell (A2-SL9)



Supplemental Figure 8. Matched data for concurrent measurement of cytolysis and IFN γ secretion for IFN γ -secreting CD8+ T-cells from two HIV+ controllers co-incubated in arrays of microwells for 6 h with HLA-matched (HLA-A2), SL9-pulsed target B-cells. The data are shown as a heatmap and organized by unsupervised hierarchical clustering (Euclidean distance, complete linkage). Cytolytic activity is shown as discrete values (blue = negative; red = death). The relative rate of secretion is shown for IFN γ (green). The heatmaps show all wells scored that had IFN γ secretion and (top) E:T = 1:1 – 5:1 or (bottom) E = 1-5; T > 1 for each sample (Fw056 and CTR040). There were no cells in either sample that released either IL-2 or TNF α at a detectable rate (limit of detection: \sim 2 molecules/s).

a) CTR084

Effector: ex vivo CD8+ T-cell; Target: B cell (Gag pool)



b) Effector: ex vivo IFN γ + CD8+ T-cell; Target: B cell (Gag pool)



Supplemental Figure 9. Matched data for concurrent measurement of cytolysis and IFN γ secretion for CD8+ T-cells from an HIV+ controller (CTR084) co-incubated in microwells for 6 h with autologous target B-cells pulsed with overlapping peptides from Gag. The data are shown as a heatmap and organized by unsupervised hierarchical clustering (Euclidean distance, complete linkage). Cytolytic activity is shown as discrete values (blue = negative; red = death). The relative rate of secretion is shown for IFN γ (green). (A) All events observed for wells scored with E:T = 1:1 – 5:1 exhibiting any functional activity; (B) All events observed for wells with E = 1-5; T > 1 and IFN γ secretion.