

**Supplementary information****Supplementary Tables****Supplementary Table 1**

<i>peptide</i>	<i>graft</i>	<i>peptide-specific T cells (total number)</i>		<i>expansion (fold)</i>
		<i>before expansion</i>	<i>after expansion</i>	
HPV	G1	1.49x10 <sup>3</sup>	4.61x10 <sup>4</sup>	31
	G2	1.06x10 <sup>4</sup>	2.86x10 <sup>5</sup>	27
	G3	2.49x10 <sup>4</sup>	1.45x10 <sup>6</sup>	58
	G4	3.60x10 <sup>3</sup>	2.36x10 <sup>4</sup>	7
	G5	8.42x10 <sup>3</sup>	1.14x10 <sup>4</sup>	1
YPL	G1	1.02x10 <sup>3</sup>	1.49x10 <sup>4</sup>	15
	G2	1.67x10 <sup>5</sup>	4.48x10 <sup>6</sup>	27
	G3	1.58x10 <sup>4</sup>	2.36x10 <sup>4</sup>	1
	G4	1.06x10 <sup>4</sup>	1.53x10 <sup>4</sup>	1
	G5	2.73x10 <sup>3</sup>	2.23x10 <sup>4</sup>	8
EPL	G1	4.09x10 <sup>4</sup>	1.67x10 <sup>6</sup>	41
	G2	1.24x10 <sup>5</sup>	1.11x10 <sup>7</sup>	89
	G3	1.13x10 <sup>5</sup>	8.67x10 <sup>6</sup>	77
	G4	3.70x10 <sup>5</sup>	1.75x10 <sup>7</sup>	47
	G5	4.10x10 <sup>4</sup>	1.18x10 <sup>7</sup>	288
GLC	G6	3.55x10 <sup>5</sup>	4.92x10 <sup>6</sup>	14
CLG	G6	1.10x10 <sup>6</sup>	9.01x10 <sup>6</sup>	8
FLY	G6	2.67x10 <sup>5</sup>	6.87x10 <sup>6</sup>	26
YVL	G6	1.16x10 <sup>6</sup>	3.10x10 <sup>7</sup>	27

**Supplementary Table 1 Peptide-specific CD8<sup>+</sup> T cell expansion in individual grafts**

Total numbers of leukocytes were determined by automatic cell counting (Sysmex Hematology Analyzer XN-350, Sysmex Corporation, Kōbe, Japan). Numbers of peptide-specific T cells were calculated by flow cytometry-determined frequencies of peptide-specific T cells among total leukocytes.

**Supplementary Table 2**

<i>reagent/specificity</i>	<i>clone</i>	<i>fluorochrome</i>	<i>vendor</i>
viability dye		7AAD	Biolegend
fixable viability dye		Zombie Red	Biolegend
fixable viability dye		Zombie Yellow	Biolegend
CD3	UCHT1	AF700	Biolegend
CD4	SK3	Pacific Blue	Biolegend
CD8a	RPA-T8	FITC	Biolegend
CD8a	RPA-T8	APC	Biolegend
CD8a	RPA-T8	BV510	Biolegend
CD14	M5E2	APC/Fire750	Biolegend
CD19	SJ25C1	APC/Fire750	Biolegend
CD20	2H7	PerCP/Cy5.5	Biolegend
CD45	J33	Krome Orange	Beckman Coulter
CD56	5.1H11	APC/Fire750	Biolegend
CD137	4B4-1	PE	Biolegend
TCR $\alpha\beta$	IP26	AF700	Biolegend
murine CD3	17A2	PE/Cy7	Biolegend
murine CD3	17A2	BV421	Biolegend
murine TCR $\beta$	H57-597	PE	Biolegend
HPV-specific HLA-B*35:01 tetramer		PE	NIH TCF
YPL-specific HLA-B*35:01 tetramer		APC	NIH TCF
EPL-specific HLA-B*35:01 tetramer		APC	NIH TCF
GLC-specific HLA-A*02:01 tetramer		PE	NIH TCF
CLG-specific HLA-A*02:01 tetramer		APC	NIH TCF
FLY-specific HLA-A*02:01 tetramer		PE	NIH TCF
YVL-specific HLA-A*02:01 tetramer		APC	NIH TCF

**Supplementary Table 2 Flow cytometry staining reagents**

All reagents were reactive against human species-reactive unless otherwise stated. AF: Alexa Fluor, FITC: fluorescein isothiocyanate, APC: allophycocyanin, BV: Brilliant Violet, PerCP: peridinin chlorophyll, Cy: cyanine, TCF: Tetramer Core Facility

**Supplementary Table 3**

<i>G</i>	<i>peptide</i>	<i>sorted cells (total number)</i>	<i>Successfully TCR<math>\alpha\beta</math>-sequenced cells (total number)</i>	<i>clonal T cells (total number)</i>	<i>number of T cell clones</i>
G1	HPV	92	87	81	6
	EPL	184	176	156	13
G2	HPV	184	144	104	11
	YPL	184	126	98	7
	EPL	184	124	81	17
G3	HPV	184	99	44	9
	EPL	184	128	85	21
G4	EPL	92	92	90	2
G5	EPL	184	158	107	24
G6	GLC	184	49	24	8
	CLG	184	36	32	1
	FLY	184	70	47	11
	YVL	184	27	14	6

**Supplementary Table 3 Frequencies of TCR-recombinant CD8<sup>+</sup> human T cells**

G: stem cell graft, clonal T cells: cells were determined clonal if we detected at least two cells with identical TCR $\alpha\beta$  CDR3 amino acid sequences.

**Supplementary Table 4**

<i>peptide</i>	<i>TRAV</i>	<i>CDR3 <math>\alpha</math> AA sequence</i>	<i>TRAJ</i>	<i>TRBV</i>	<i>CDR3 <math>\beta</math> AA sequence</i>	<i>TRBJ</i>	<i>cf (%)</i>				
							<i>G1</i>	<i>G2</i>	<i>G3</i>	<i>G4</i>	<i>G5</i>
HPV	5*01	CAESYTGGFKTIF	9*01	6-1*01	CASGSEAFF	1-1*01	6		nc		
HPV	19*01	CALSEAGGFGNEKLTF	48*01	10-3*01	CAISDPRDSYEQYF	2-7*01		nc	nc		
EPL	1-2*01	CAVRGSGGSYIPTF	6*01	10-3*01	CATGTGDSNQPQHF	1-5*01		2	11		
EPL	25*01	CAGRFMFSGGYNKLIF	4*01	28*01	CASSLPGANVLTF	2-6*01			4	2	
EPL	2*01	CAVEDMNSGGYQKVTF	13*01	28*01	CASKRTATYEQYF	2-7*01			nc		8

**Supplementary Table 4 Overlap of epitope-specific TCRs between different stem cell grafts**

TRAV: TCR $\alpha$  V-gene and allele, TRAJ: TCR $\alpha$  J-gene and allele, TRBV: TCR $\beta$  V-gene and allele, TRBJ: TCR $\beta$  J-gene and allele, AA: amino acid, cf: clone frequency among clonally expanded cells specific for the respective epitope, G: stem cell graft, nc: detectable but not clonally expanded

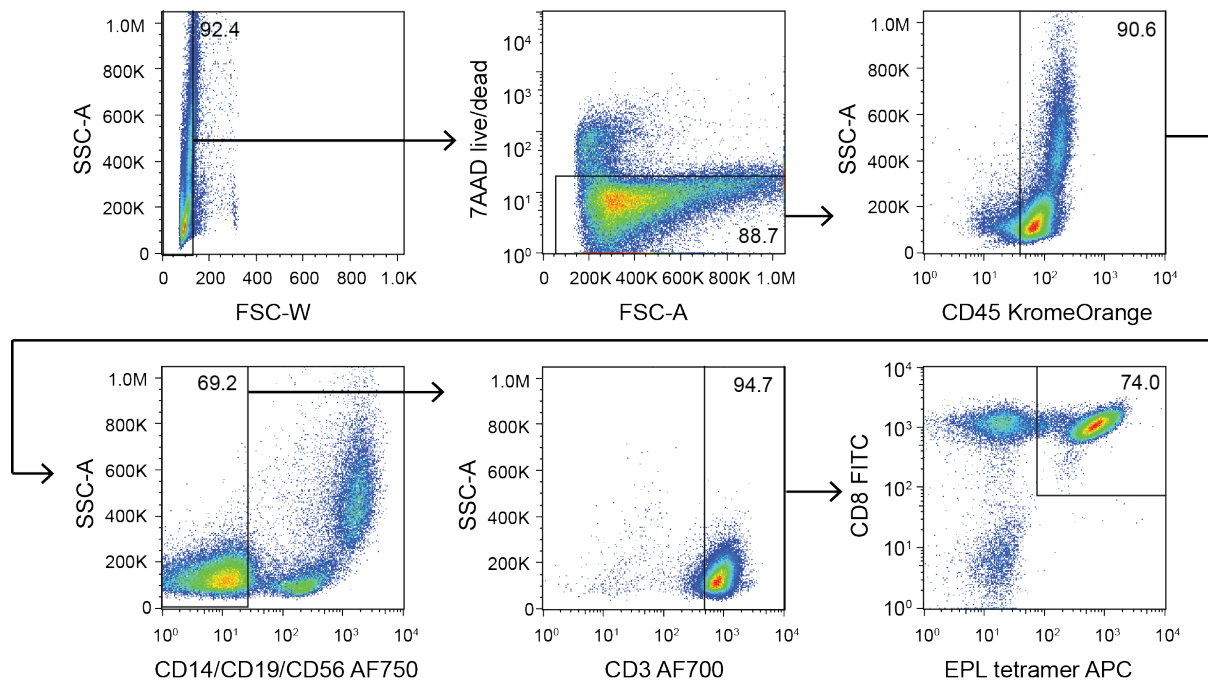
**Supplementary Table 5**

<i>cell type and name</i>	<i>HLA-A</i>		<i>HLA-B</i>		<i>HLA-C</i>	
miniLCL	01:01	26:01	35:01	57:01	04:01	06:02
miniLCL	02:01	29:02	44:02	45:01	06:02	-
LCL B01	03:01	24:02	15:01	35:01	03:03	04:01
LCL B03	02:01	23:01	15:01	58:01	03:04	07:01
LCL DJS	02:01	03:01	35:01	37:02	04:01	06:02
LCL JY	02:01	-	07:02	-	07:02	-

**Supplementary Table 5 HLA class-I data of all LCL and miniLCL**

## Supplementary Figures

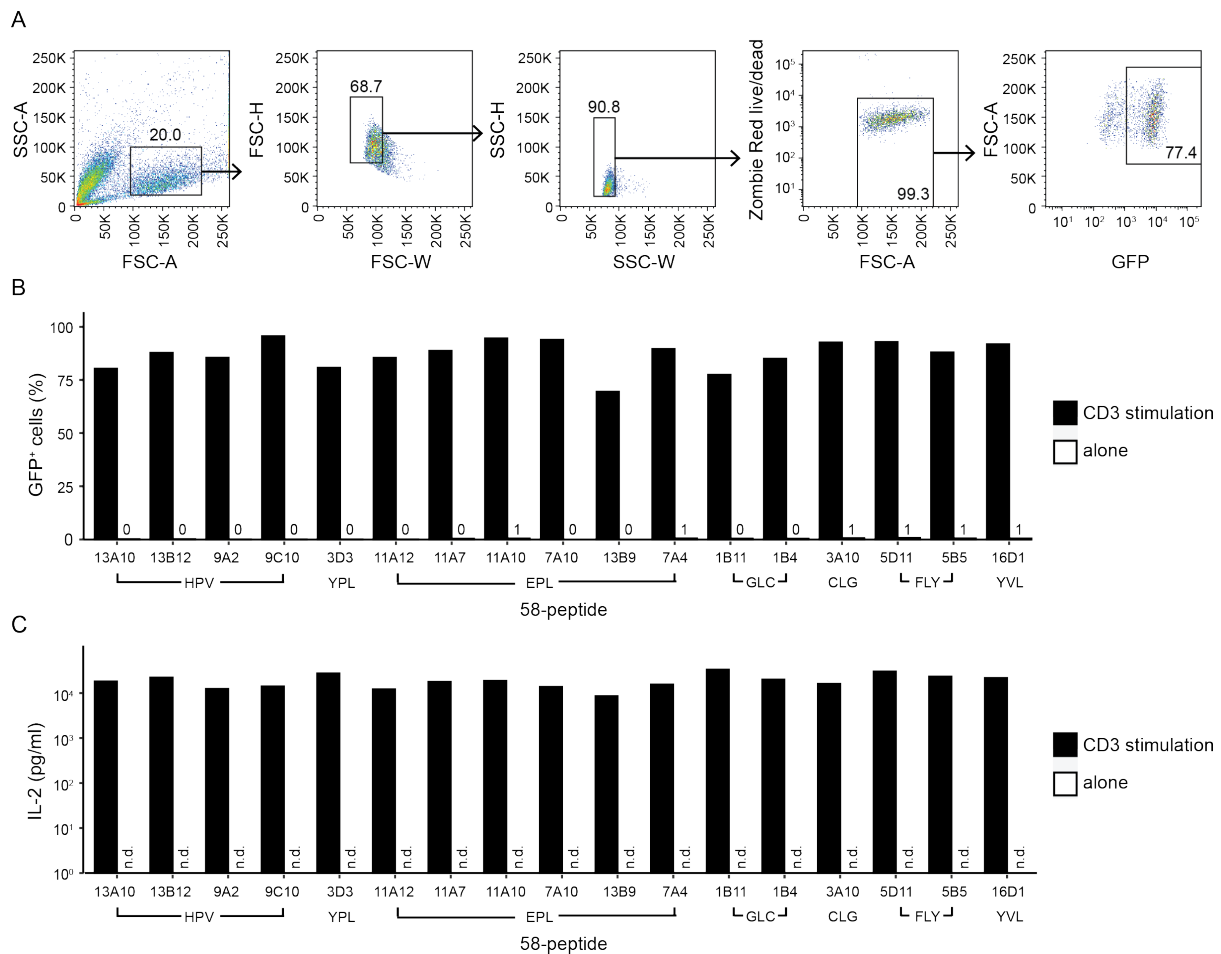
## Supplementary Figure 1



### Supplementary Figure 1 Identification of epitope-specific T cells by pMHC tetramer staining

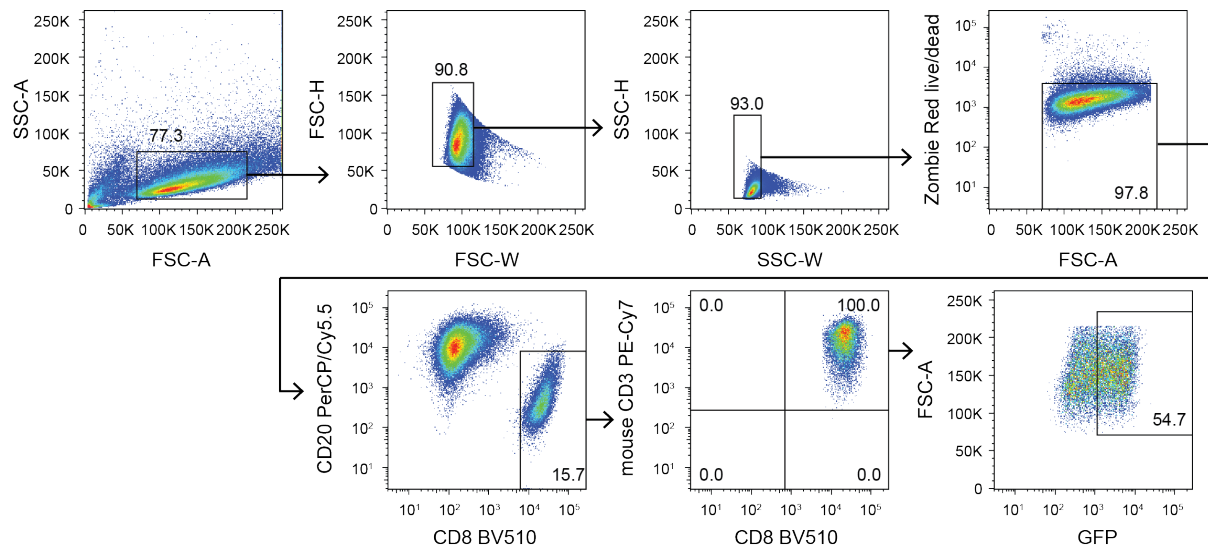
Gating strategy for peptide-specific T cells before (day 0) and after (day 9) peptide stimulation. Gating on single, live, CD45<sup>+</sup>, CD14<sup>-</sup>CD19<sup>-</sup>CD56<sup>-</sup> cells. After selection of CD3<sup>+</sup> T cells, the tetramer gate was set based on staining of cells that were expanded in presence of a different peptide. Numbers within or adjacent to gates indicate percentages.

## Supplementary Figure 2



### Supplementary Figure 2 Stimulation of TCR-recombinant 58 $\alpha$ $\beta$ <sup>-</sup> cells with plate-bound anti-CD3

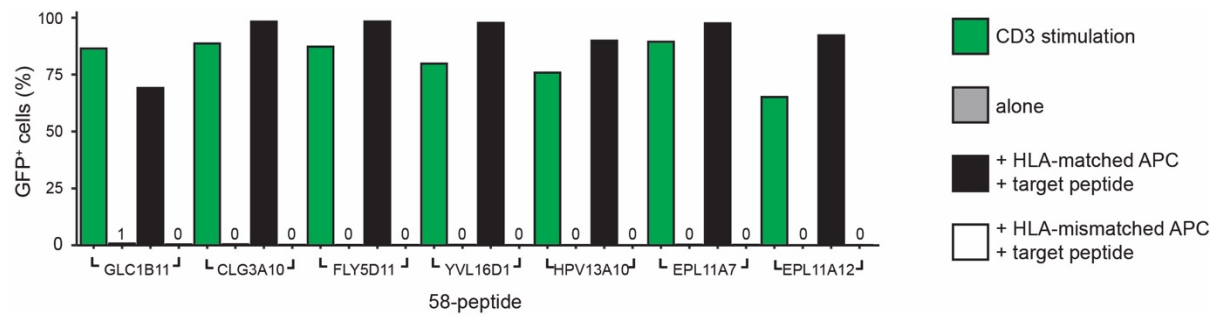
(A) Gating strategy for the identification of GFP<sup>+</sup> TCR-recombinant cells. 58-GLC1B11 stimulated with CD3 is shown as an example. GFP gates were set based on non-stimulated controls. (B) Summarizes GFP expression of all TCR-recombinant 58 $\alpha$  $\beta$ <sup>-</sup> cell lines upon CD3 stimulation following the gating strategy presented in figure part A. “Alone” refers to TCR-recombinant 58 $\alpha$  $\beta$ <sup>-</sup> reporter cell lines alone. Numbers in places of bars indicate percentages. (C) IL-2 production measured by ELISA in cell culture supernatants corresponding to data in figure part B. n.d. = not detectable.

**Supplementary Figure 3**

**Supplementary Figure 3 Gating strategy for identification of GFP expressing TCR-recombinant 58 $\alpha$  $\beta$ - cells**

The figure shows GFP expression in 58-GLC1B11 in response to stimulation with GLC peptide-loaded antigen-presenting cells as an example. Numbers within or adjacent to gates indicate percentages.

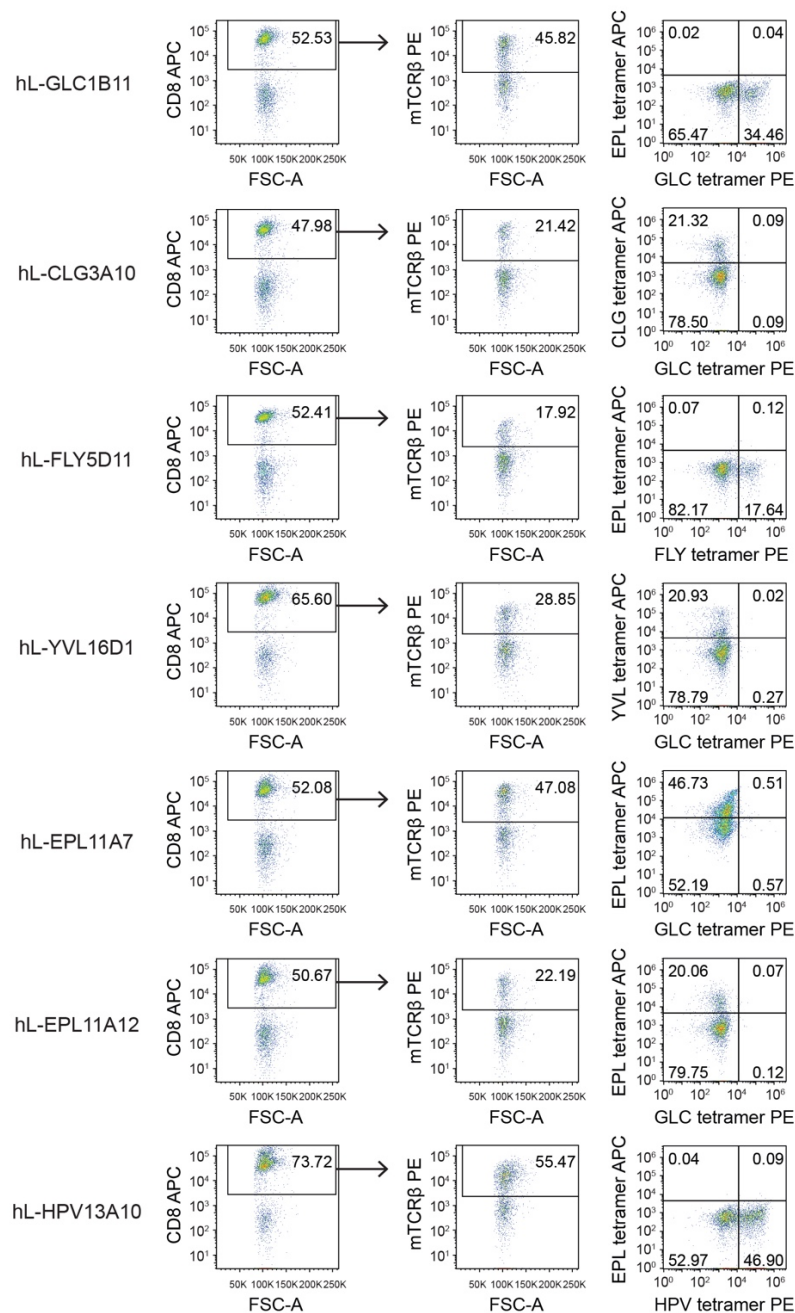


**Supplementary Figure 4**

**Supplementary Figure 4 No cross-reactivity of four HLA-A\*02:01 and three HLA-B\*35:01-restricted TCRs with HLA-mismatched miniLCL**

GFP expression of seven TCR-recombinant  $58\alpha\beta^+$  cell lines upon co-culture with HLA-matched and HLA-mismatched miniLCL loaded with the corresponding target peptides. Detailed HLA class-I data of the used miniLCL can be found in Supplementary Table 5. Numbers in places of bars indicate percentages. All bars represent mean values of two co-cultures within one experiment. APC = antigen presenting cells.

## Supplementary Figure 5

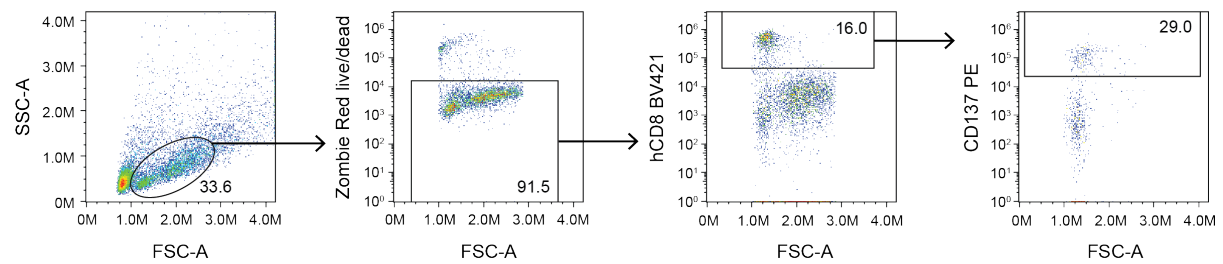


## Supplementary Figure 5 Recombinant TCR expression on human lymphocytes

Plots in the left column were pre-gated on lymphocytes by scatter characteristics. Plots in middle and right columns were pre-gated on CD8<sup>+</sup> lymphocytes. pMHC tetramer staining was done in a separate experiment using T cells of the same transduction and FITC-conjugated CD8 antibody. mTCRβ and pMHC tetramer gates were set based on controls with non-transduced T cells.

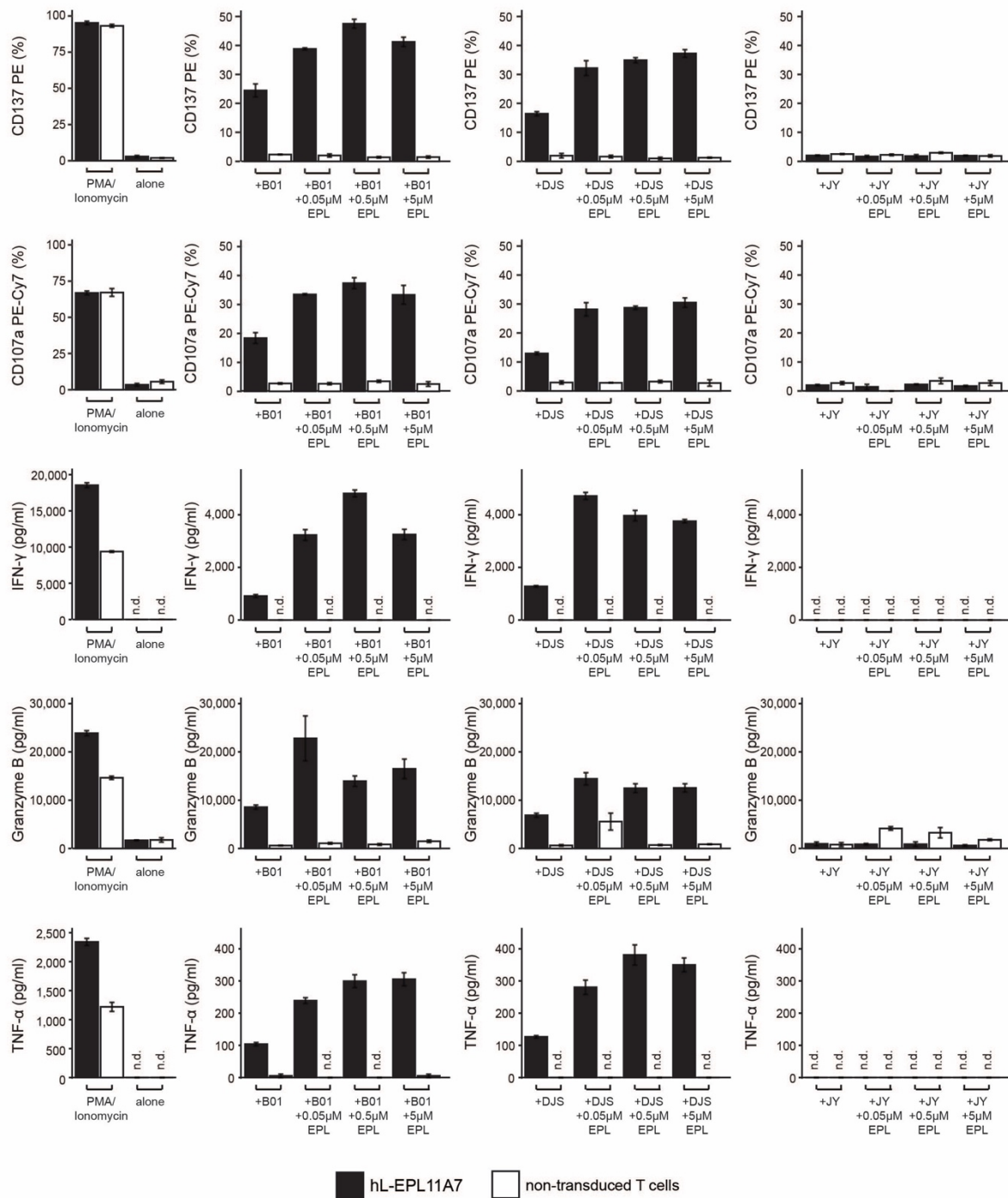
**Supplementary Figure 6**

hl-GLC1B11 + B03

**Supplementary Figure 6 Gating strategy for identification of CD8<sup>+</sup>CD137<sup>+</sup> human T cells**

Data from one co-incubation of hl-GLC1B11 with LCL B03 are shown as a representative example. CD137 gates were set based on non-stimulated controls.

Supplementary Figure 7



Supplementary Figure 7 Detailed activation characteristics of hL-EPL11A7

hL-EPL11A7 were incubated with HLA-B\*35:01-matched LCL (B01 and DJS) or HLA-B\*35:01-mismatched LCL (JY) in presence or absence of increasing target peptide (EPL)

concentrations. CD137 and CD107a expression were determined by flow cytometry after pre-gating on live CD8<sup>+</sup> lymphocytes. IFN- $\gamma$ , granzyme B, and TNF- $\alpha$  were measured in cell culture supernatants by ELISA. All bars represent mean values  $\pm$  standard error of three co-cultures within one experiment. n.d. = not detectable.