

1 **Supplementary Material**

2 **High-efficiency magnetophoretic labelling of adoptively-transferred** 3 **T cells for longitudinal in vivo Magnetic Particle Imaging**

4 Rong En Tay^{1*†}, Lokamitra P^{1,2}, Shun Toll Pang^{2^}, Kay En Low³, Hui Chien Tay¹, Charmaine
5 Min Ho¹, Benoit Malleret^{3,4}, Olaf Röttschke^{1,4,5}, Malini Olivo⁶ & Zhi Wei Tay^{2,7*†‡}

6

7 ¹Singapore Immunology Network (SIgN), Agency for Science, Technology and Research
8 (A*STAR), 8A Biomedical Grove, #04-06 Immunos, Singapore 138648, Republic of Singapore.

9 ²Institute of Bioengineering and Bioimaging (IBB), Agency for Science, Technology and
10 Research (A*STAR), 11 Biopolis Way, #02-02 Helios Building, Singapore 138667, Republic of
11 Singapore.

12 ³Electron Microscopy Unit, Yong Loo Lin School of Medicine, National University of
13 Singapore, Singapore 117597, Republic of Singapore.

14 ⁴Department of Microbiology and Immunology, Immunology Translational Research
15 Programme, Yong Loo Lin School of Medicine, National University of Singapore, Republic of
16 Singapore.

17 ⁵School of Biological Sciences, Nanyang Technological University, 50 Nanyang Avenue,
18 Singapore 639798. Republic of Singapore.

19 ⁶A*STAR Skin Research Labs (A*SRL), Agency for Science, Technology and Research
20 (A*STAR), 31 Biopolis Way, #07-01 Nanos, Singapore 138669, Republic of Singapore.

21 ⁷National Institute of Advanced Industrial Science and Technology (AIST), Health and Medical
22 Research Institute (HMRI), 1-2-1 Namiki, Tsukuba, Ibaraki 305-8564, Japan.

23

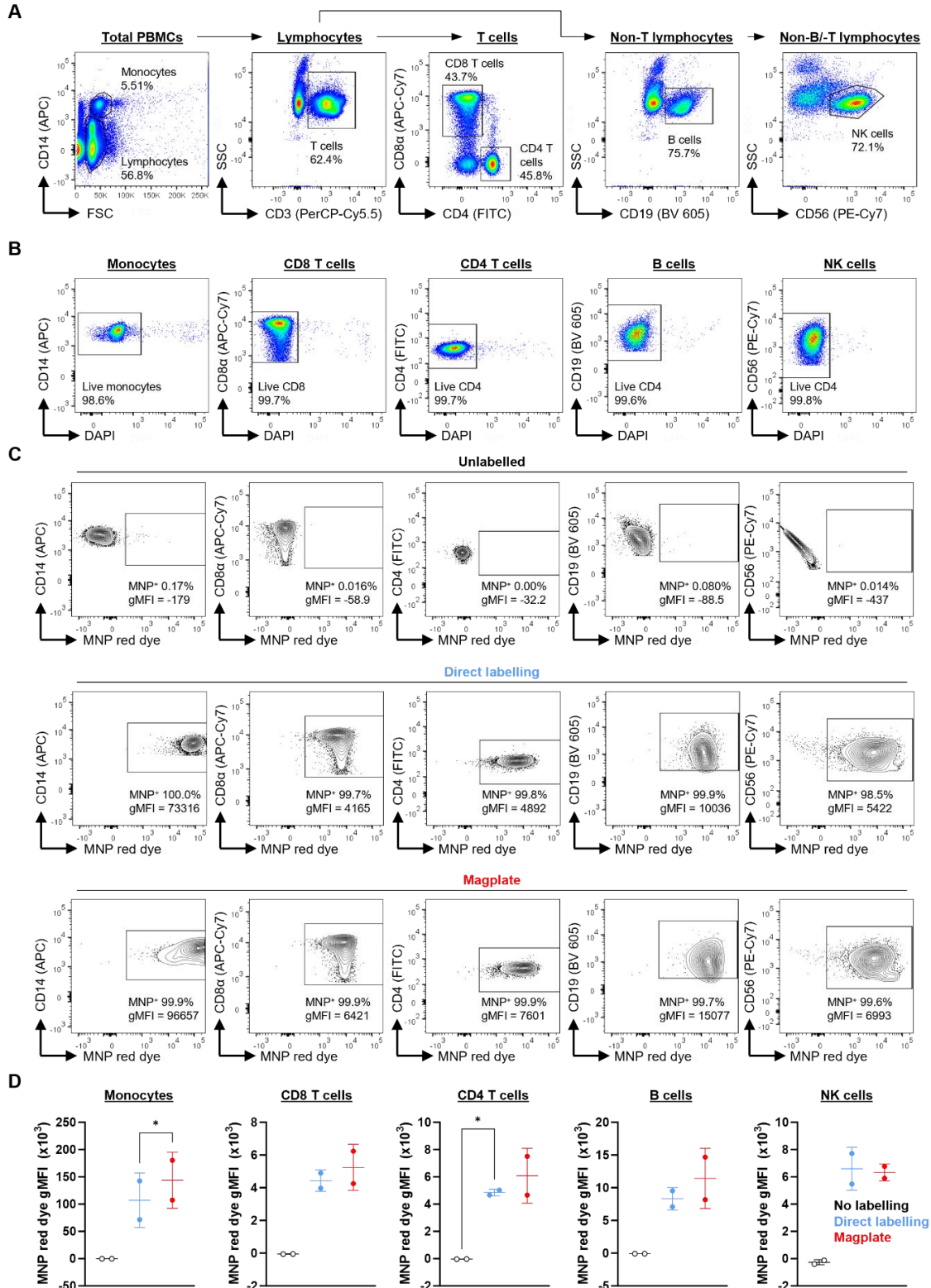
24 *Correspondence: zhiwei.tay@aist.go.jp and tay_rong_en@immunol.a-star.edu.sg

25 †These authors contributed equally

26 Current affiliations: ^Institute of Technical Education, 2 Ang Mo Kio Drive, Singapore 567720,
27 Republic of Singapore

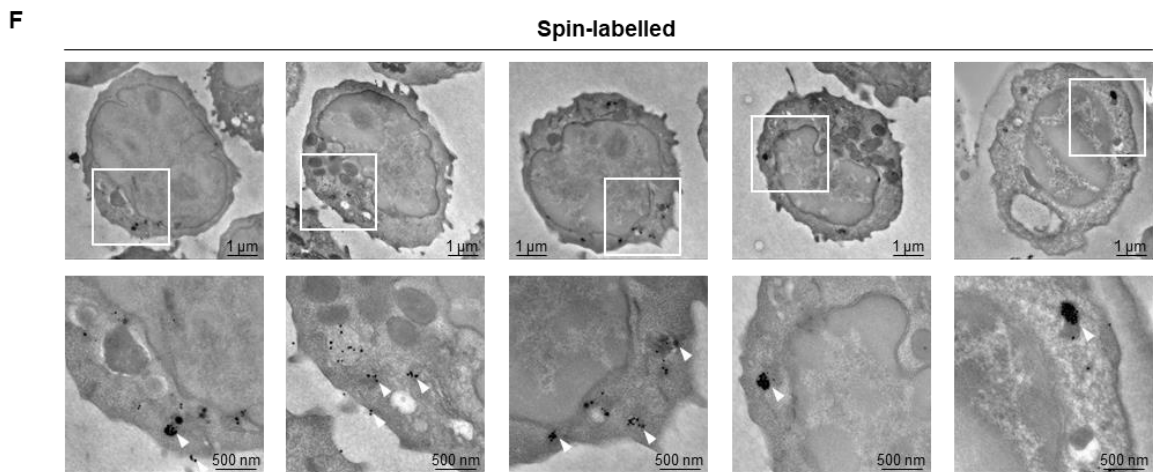
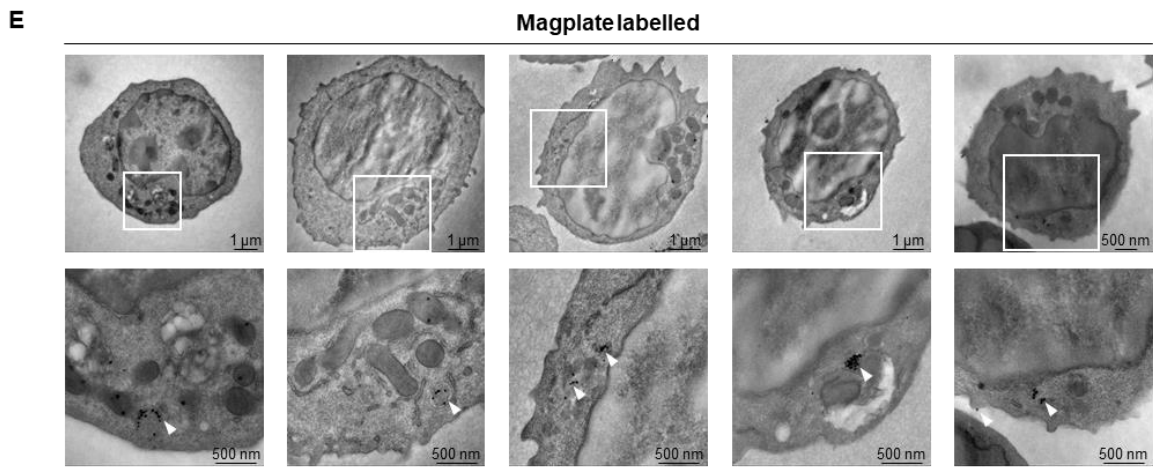
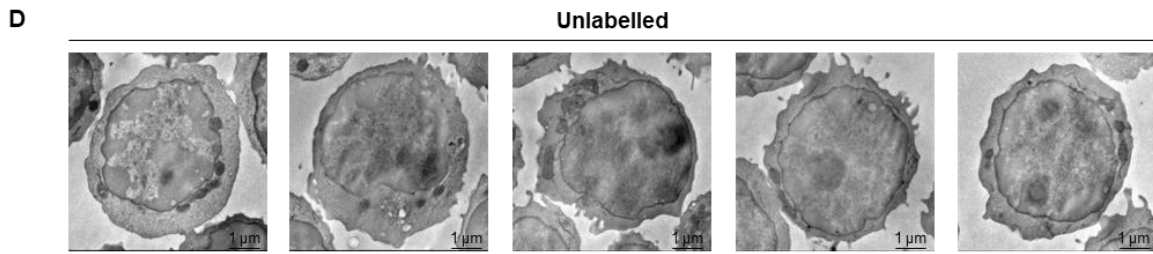
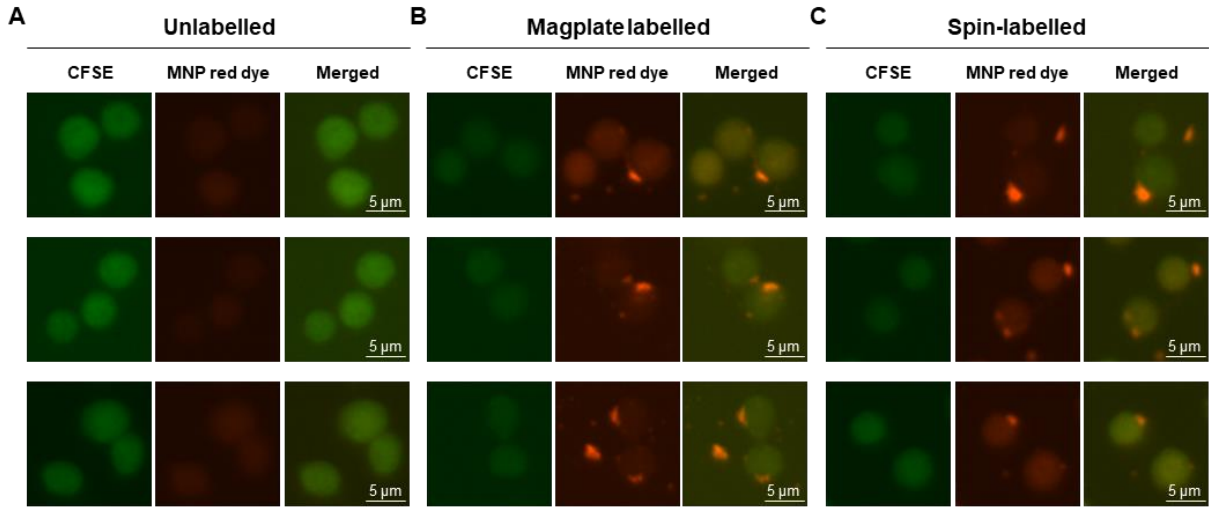
28 ‡Health and Medical Research Institute (HMRI), National Institute of Advanced Industrial
29 Science and Technology (AIST), 1-2-1 Namiki, Tsukuba, Ibaraki 305-8564, Japan

30

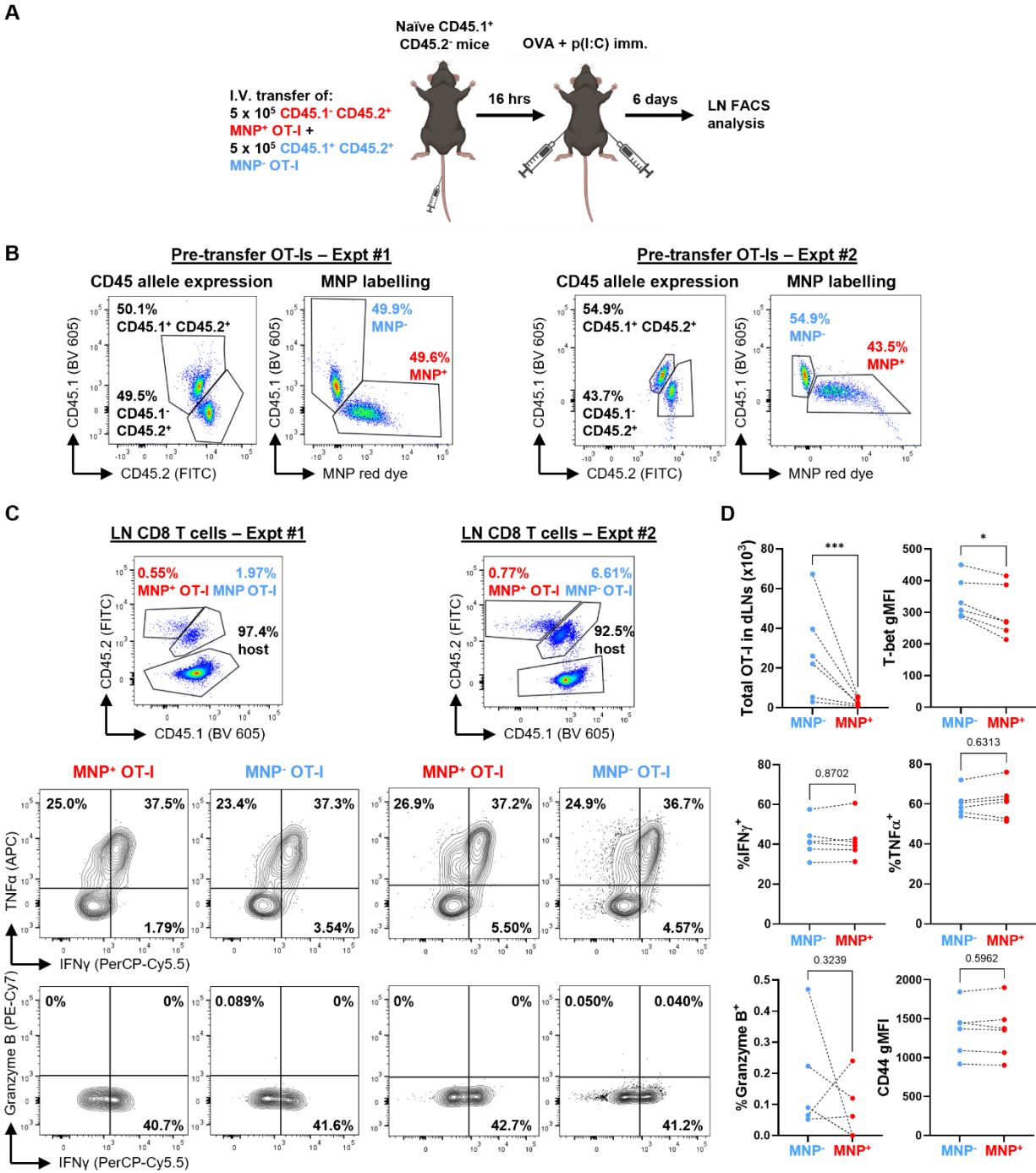


31
 32 **Figure S1. Magplate labelling increases MNP label uptake across five major subsets of healthy human**
 33 **PBMCs, related to Figure 1. (A-D)** PBMCs from two healthy human donors were labelled with MNPs
 34 either by direct labelling or magplate labelling for 3 h. As a control, parallel samples of PBMCs were
 35 incubated for 3 h without MNP labels. Label uptake within five major subsets of immune cells in the

36 PBMCs was then assessed by flow cytometry **(A)**. Representative flow cytometry plots showing gating
37 strategy to identify (left to right) monocytes, CD8 and CD4 T cells, B cells, and NK cells in PBMCs. Data
38 are representative of labelled and unlabelled PBMCs from two separate donors. **(B)** Live/Dead exclusion
39 by DAPI staining for the five immune cell subsets, gated as in **(A)**. **(C)** Representative flow cytometry plots
40 showing gating and MNP label uptake in the five immune cell subsets, gated as in **(A)**, in PBMC samples
41 that were (top to bottom rows) left unlabelled, directly labelled, and magplate-labelled. **(D)** Quantification
42 of label uptake across immune cell subsets from PBMCs from two healthy donors. Data are mean \pm SD,
43 statistical analysis using two-tailed one-way ANOVA with Tukey's test for multiple comparisons. * $p <$
44 0.05.
45



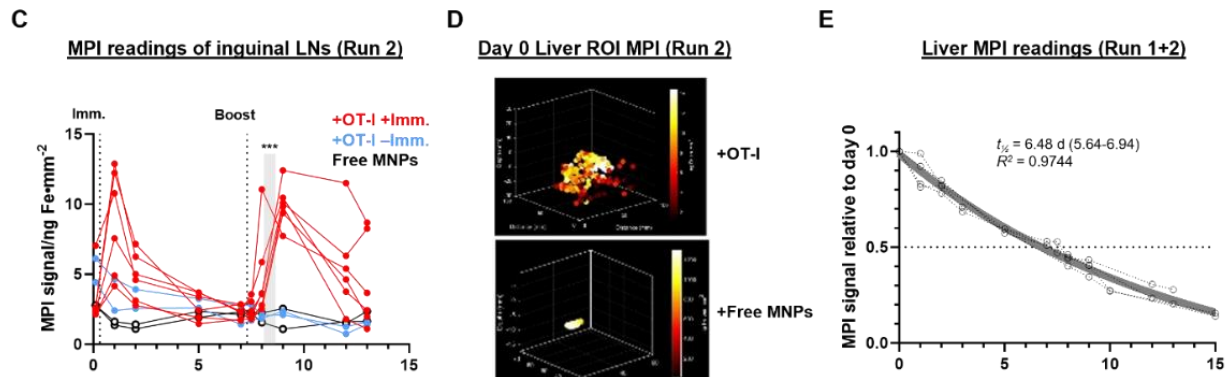
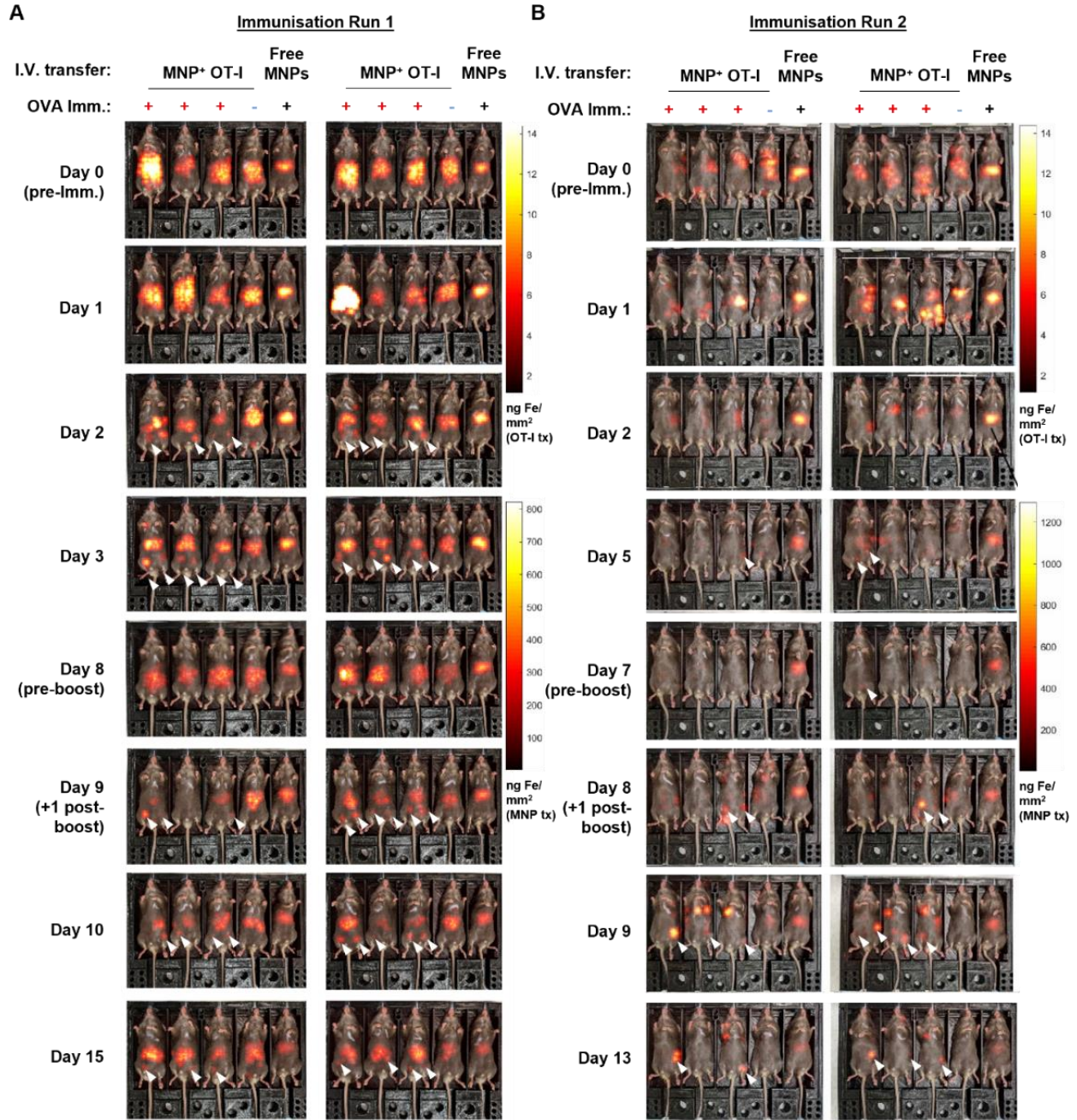
47 **Figure S2. Extended sets of live cell microscopy and transmission electron microscopy images of**
48 **MNP-labelled primary mouse CD8 T cells, related to Figure 2 (A-C)** Representative live cell fluorescent
49 images of primary mouse OT-I CD8 T cells labelled with cytosolic CFSE (green channel) prior to indicated
50 labelling procedures. **(D-F)** Representative transmission electron microscope images of primary mouse OT-
51 I CD8 T cells treated with indicated labelling procedures. White squares indicate regions of interest
52 displayed at higher magnification below each whole cell image **(E and F)**. White arrowheads indicate
53 MNPs in high-magnification images **(E and F)**.
54



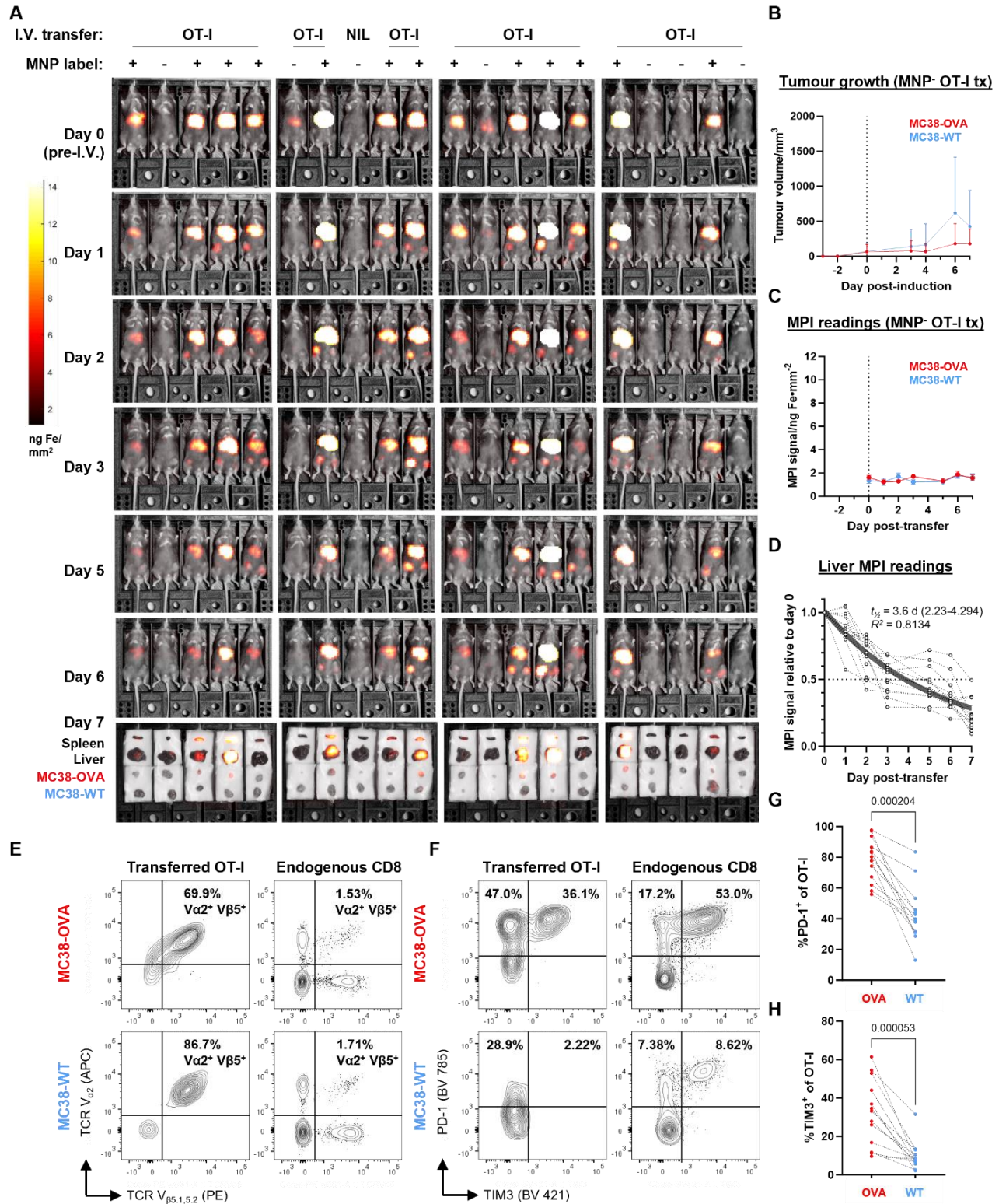
55

56 **Figure S3. In vivo competition experiment between MNP-labelled and –unlabelled mouse**
 57 **primary CD8 T cells, related to Figure 3.** (A) OT-I CD8 T cells harvested from CD45.1⁺ CD45.2⁻ mice
 58 were labelled with MNPs for 3 h with the magplate (red). In parallel, OT-I CD8 T cells harvested from
 59 CD45.1⁺ CD45.2⁻ mice were left unlabelled for 3 h (blue). 5 × 10⁵ MNP⁺ and MNP⁻ OT-I cells were then
 60 adoptively transferred into CD45.1⁺ CD45.2⁻ congenic C57BL/6J mice by intravenous injection and the
 61 mice were immunised with OVA + p(I:C) the following day by subcutaneous injection into both flanks.
 62 After 6 days, cells suspensions prepared from draining inguinal lymph nodes were analysed by flow
 63 cytometry following restimulation with 5 µg/mL OVA SIINFEKL peptide + protein transport inhibitors.

64 **(B)** Pre-transfer flow cytometry analysis of the 1:1 mix of CD45.1⁻ CD45.2⁺ MNP⁺ OT-I (red) + CD45.1⁺
65 CD45.2⁺ MNP⁻ OT-I (blue). Gated on live CD8 α ⁺ events. **(C and D)** Representative flow cytometry plots
66 (C) and quantification (D) of parameters of OT-I CD8 T cell effector function in inguinal lymph nodes of
67 mice. Gated on live TCR β ⁺ CD8 α ⁺ CD4⁻ events. Data are means \pm SD and are from two independent
68 experiments with two and four recipient immunised mice, respectively. Numbers indicate *p*-values of
69 comparisons using two-tailed ratio paired t-test (all parameters except % Granzyme B⁺) or two-tailed paired
70 t-test (% Granzyme B⁺).
71



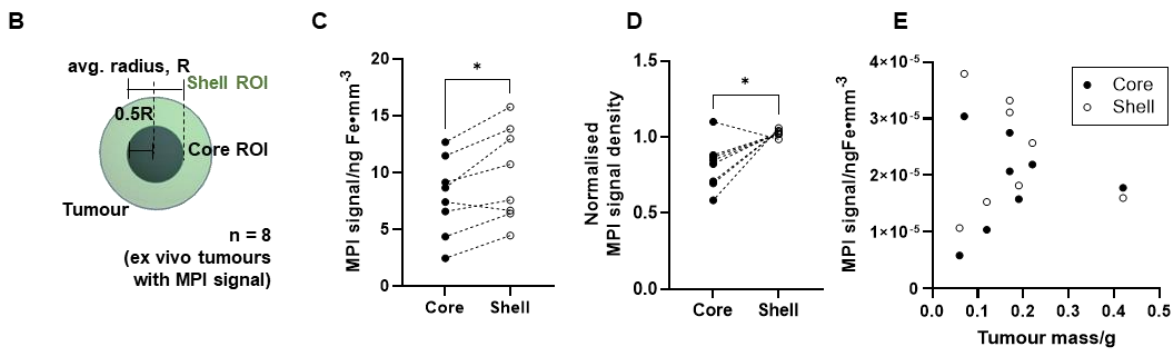
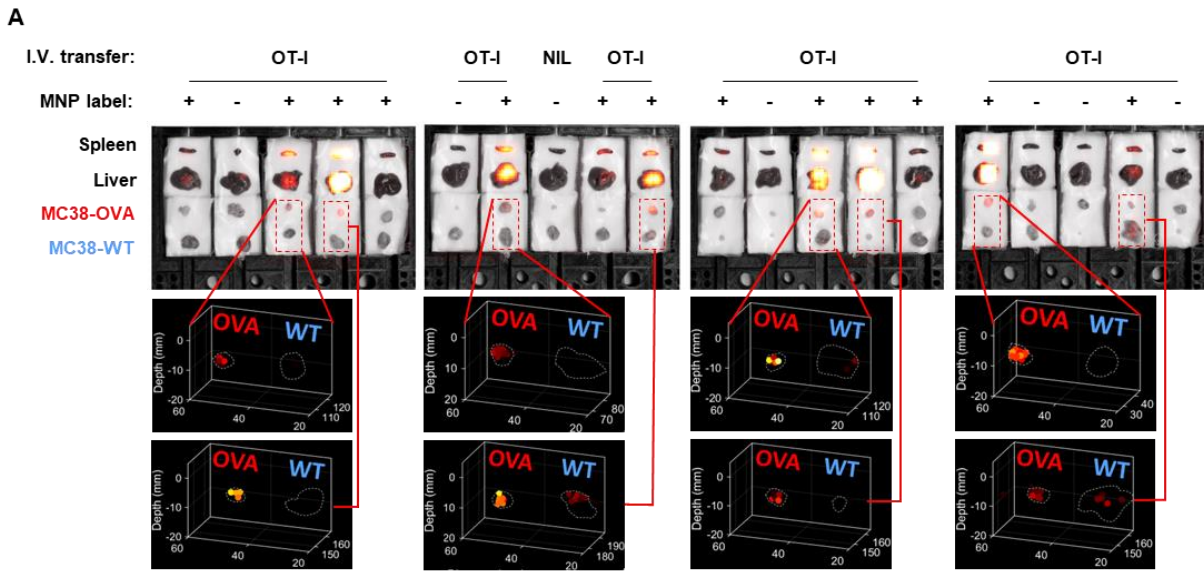
73 **Figure S4. Additional data from MPI tracking of adoptively-transferred antigen-specific CD8 T cells**
74 **in response to immunisation, related to Figure 6. (A and B)** MPI imaging time courses of two
75 independent immunisation experiments with 10 mice each. 2×10^6 CD45.1⁺ CD45.2⁺ OT-I CD8 T cells
76 were labelled with MNPs, then adoptively transferred into C57BL/6J mice (CD45.1⁻ CD45.2⁺) by
77 intravenous injection. As an imaging control, one mouse in each set of five mice was injected with a
78 suspension of free MNPs of an equivalent mass of iron. The following day, the mice received subcutaneous
79 flank injections of OVA + p(I:C) (immunisation, red) or PBS (mock immunisation, blue) and were boosted
80 7 days later as shown in **Figure 6A**. Total body MPI was performed throughout the duration of the
81 experiment. White arrowheads indicate local MPI signal peaks corresponding to the anatomical locations
82 of the inguinal lymph nodes draining the immunisation sites. **(C)** Time course of MPI signal in inguinal
83 lymph nodes of mice from **(B)**, with 6, 2, and 2 mice in the +OT-I+Imm. (red), +OT-I-Imm. (blue), and
84 MNP (black) treatment groups, respectively. Statistical analysis by 2-way ANOVA with Sidak's test for
85 multiple comparisons between the +OT-I+Imm. (red) and +OT-I-Imm. (blue) treatment groups as indicated
86 by the grey bars. *** $p < 0.001$. **(D)** Representative 3D MPI images of abdominal cavities of mice from **(B)**
87 that received OT-I CD8 T cells and free MNPs on day 0. **(E)** Decay of MPI signal relative to day 0 in liver
88 ROIs of mice that received intravenous injections of free MNPs (two per experiment, four total). Grey curve
89 shows the best-fit one-phase decay curve, with the best-fit and (95% confidence interval) of the half-life
90 ($t_{1/2}$) and R^2 goodness-of-fit as indicated.



91
92 **Figure S5. Additional data from MPI tracking of tumour antigen-specific CD8 T cells in tumour-**
93 **bearing hosts, related to Figure 7. (A) MPI imaging time course of all mice in the experiment shown in**
94 **Figure 7. C57BL/6J mice received subcutaneous implantation of 5×10^5 MC38-OVA and -WT tumour**
95 **cells injected into their right and left flanks, respectively. After 10 days, mice with dual palpable tumours**

96 received 10×10^6 MNP-labelled CD45.1⁺ CD45.2⁺ OT-I CD8 T cells by intravenous injection ($n = 13$). As
97 controls, some mice received unlabelled OT-I CD8 T cells ($n = 6$), and one mouse was injected with PBS
98 (mock cell transfer). Total body MPI was performed throughout the duration of the experiment and endpoint
99 *ex vivo* MPI was performed on freshly harvested organs on day 7 post-cell transfer. **(B and C)** Tumour
100 growth data **(B)** and MPI signal data **(C)** from mice that received unlabelled OT-I CD8 T cells ($n = 6$).
101 Statistical analysis using (left) 2-way ANOVA with Sidak's test for multiple comparisons between MC38-
102 OVA and -WT tumour volumes with matching for tumour pairs from the same mouse (no significant
103 differences detected). **(D)** Decay of MPI signal relative to day 0 in liver ROIs of mice that received MNP-
104 labelled OT-I CD8 T cells ($n = 13$). Grey curve shows the best-fit one-phase decay curve, with the best-fit
105 and (95% confidence interval) of the half-life ($t_{1/2}$) and R^2 goodness-of-fit as indicated. **(E)** Representative
106 flow cytometry plots showing positive identification of OT-I cells in tumours from mice that received MNP-
107 labelled OT-I cells by co-expression of TCR V α 2 and V β 5 chains. Gated on TCR β^+ CD8 α^+ CD4 $^-$ CD45.1 $^+$
108 events as shown in **Figure 7G**. **(F-H)** Representative flow cytometry plots **(F)** and quantification **(G and**
109 **H)** of T cell exhaustion marker expression in OT-I cells in tumours from mice that received MNP-labelled
110 OT-I cells ($n = 13$). Gated on TCR β^+ CD8 α^+ CD4 $^-$ CD45.1 $^+$ events as shown in **Figure 7G**. Statistical
111 analysis using ratio paired t-test with the Benjamini-Krieger-Yekutieli FDR approach for multiple
112 comparisons between paired MC38-OVA and -WT tumours from the same mouse (number indicates q -
113 value)

114



115

116 **Figure S6. Quantitative 3D image analysis of MNP biodistribution within excised tumours, related to**
 117 **Figure 7. (A)** MPI imaging in 3D of excised tumours from the same experiment presented in Fig. 7. Only
 118 mice with labelled OT-I CD8 T cells and the excised tumour having detectable MPI signal at Day 7 ($n = 8$)
 119 in the MC38-OVA tumour were selected for zoom-in imaging. A narrow field-of-view (FOV) around the
 120 excised tumours demarcated by the dashed red lines in the 2D photo background was then plotted in 3D
 121 with MC38-OVA and MC38-WT (OVA^-) tumours in the field-of-view with spot size set as 2 mm
 122 corresponding to the spatial resolution limit of 2 mm of the MPI device used. Tumour outlines within each
 123 FOV are marked by dashed white lines. **(B-E)** Quantitative analysis of the average signal density in the
 124 inner (core) versus outer (shell) ROIs of the tumours. **(B)** The core was defined as the inner region of the
 125 tumour demarcated by a spherical surface with half the average radius of the whole tumour, with the shell
 126 defined as the rest of the tumour volume. MPI signal voxels were binned into their respective core or shell
 127 ROIs, and the average signal density for each ROI was calculated. **(C)** Paired core-shell MPI signal density
 128 values for excised OVA^+ tumours ($n=8$). **(D)** Paired core-shell MPI signal density values for excised
 129 MC38-OVA tumours ($n=8$) normalized to the average signal density assessed across entire tumour
 130 inclusive of both core and shell regions. **(E)** Correlation plot of MPI signal density values for the core and
 131 shell ROI versus tumour mass. Data were analysed using two-tailed ratio-paired t-test, $*p < 0.05$ (C and
 132 D).

133

134 **Table S1. Details of the labelling reagents used for flow cytometry in this work.**

Step	Species reactivity	Fluorophore	Epitope	Clone	Dilution buffer	Dilution	Manufacturer			
Live/Dead dye	N/A	Zombie Aqua	Free amines	N/A	PBS	1:500	Biologend			
	N/A	DAPI	dsDNA	N/A	FACS	1:200	Biologend			
Fc receptor blocking	Mouse	N/A	CD16/32	93	FACS	1:50	Biologend			
	Human	N/A	F _c receptors	TruStain FcX	FACS	1:50	Biologend			
Surface epitope staining	Mouse	PE/Dazzle 594	CD4	RM4-5	FACS	1:200	Biologend			
		BV 650	CD8 α	53-6.7						
		AlexaFluor 700	CD44	IM7						
		BV 605	CD45.1	A20		1:100				
		FITC	CD45.2	104						
		BV 785	PD-1	29F.1A1 2				1:200		
		APC-Cy7	TCR β	H57-597		FACS		1:200	Biologend	
		APC	TCR V α 2	B20.1						
		PE	TCR V β 5.1,5.2	MR9-4						
	BV 421	TIM3	RMT3-23							
	Human	APC-Cy7	CD3	OKT3	FACS		1:200			Biologend
		PerCP-Cy5.5	CD3	OKT3						
		FITC	CD4	RPA-T4						
		PE/Dazzle 594	CD4	RPA-T4						
		APC-Cy7	CD8 α	RPA-T8						
		BV 650	CD8 α	RPA-T8						
		APC	CD14	M5E2						
		BV 605	CD19	HIB19						
		PE-Cy7	CD56	MEM-188						

Intracellular epitope staining	Mouse	PE-Cy7	Granzyme B	NGZB	Perm	1:50	Thermo Fisher Scientific
		PerCP-Cy5.5	IFN γ	XMG1.2			Biolegend
		PE	IL-2	JES6-5H4			
		BV 421	T-bet	4B10			
		APC	TNF α	MP6-XT22			
	Human	PE-Cy7	Granzyme B	QA16A02	Perm	1:50	Biolegend
		PerCP-Cy5.5	IFN γ	B27			
		PE	IL-2	MQ1-17H12			
		BV 421	Perforin	B-D48			
		APC	TNF α	W19063 E			

135