

# Supplemental Figures

## Mapping immune processes in intact tissues at cellular resolution

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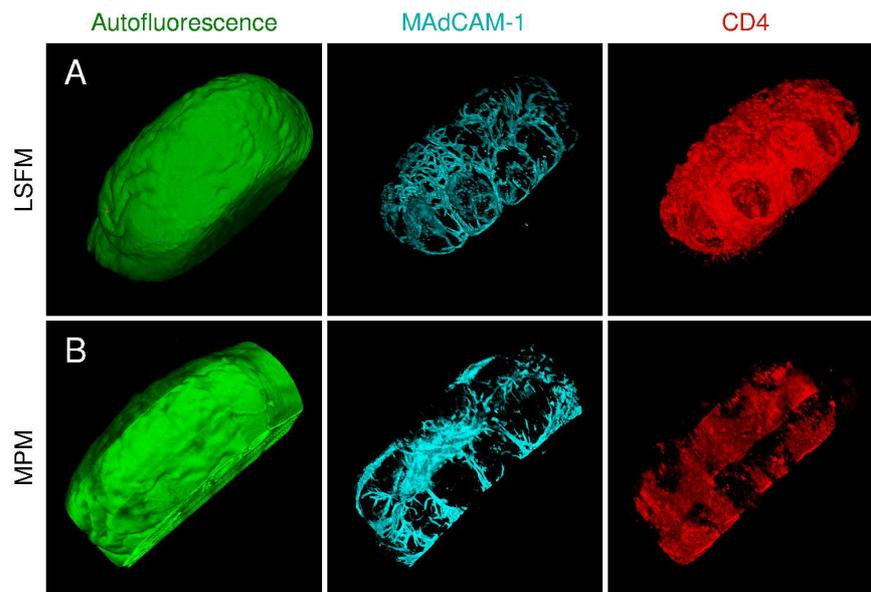
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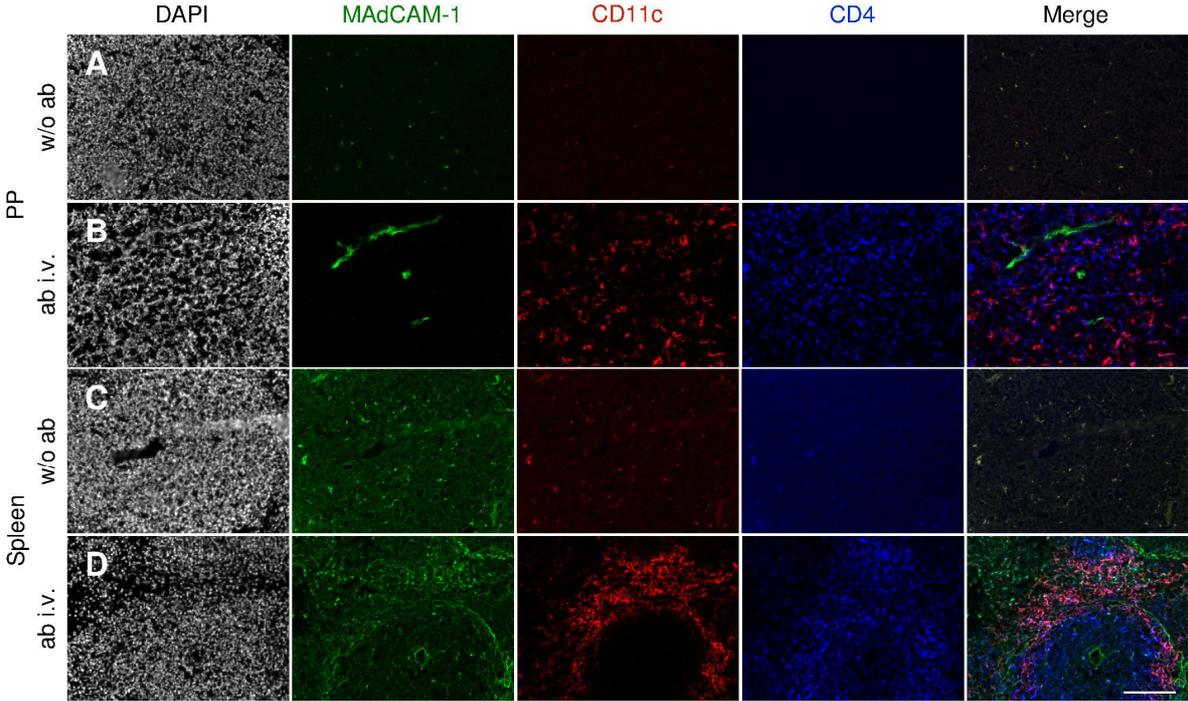
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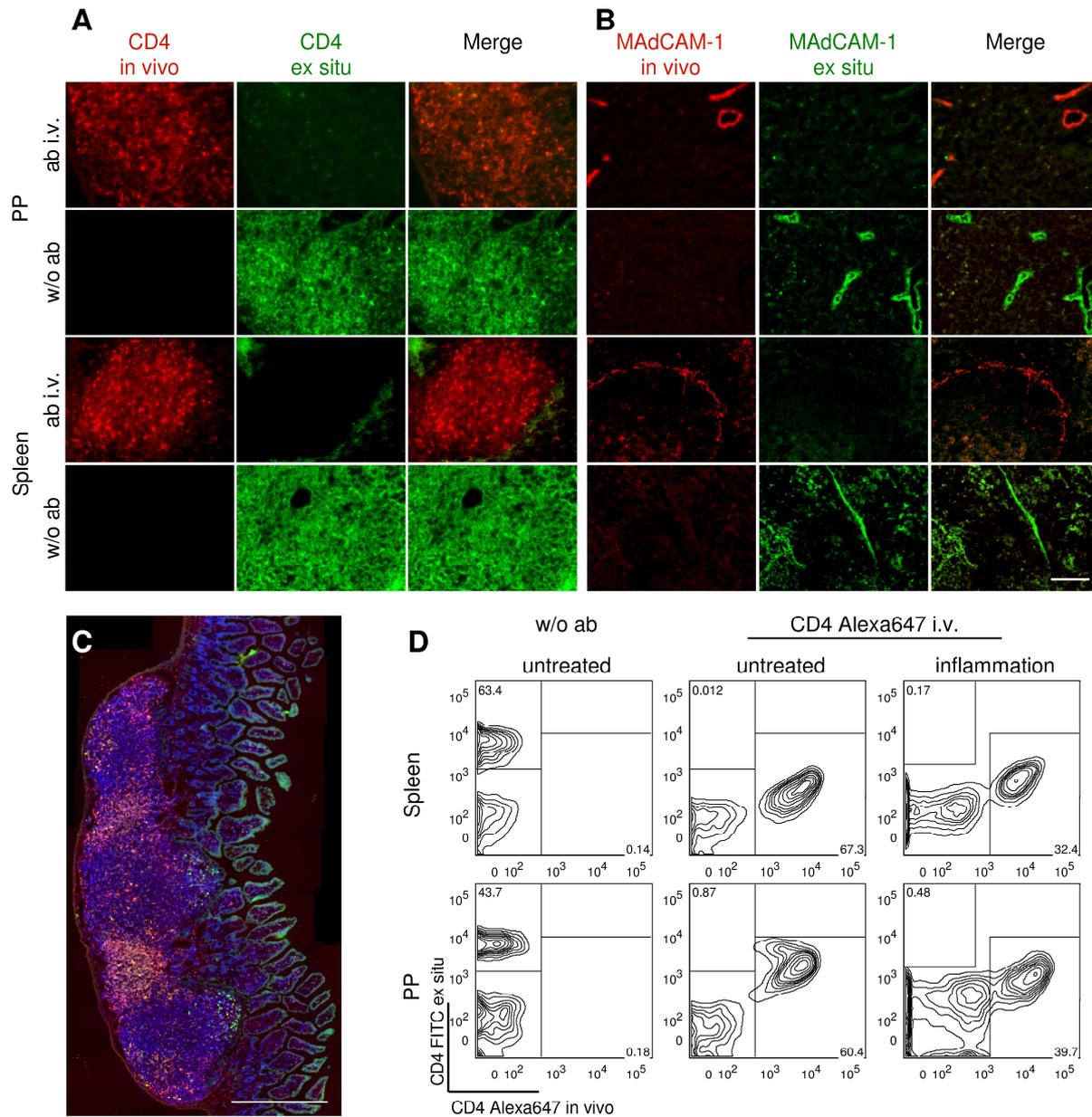
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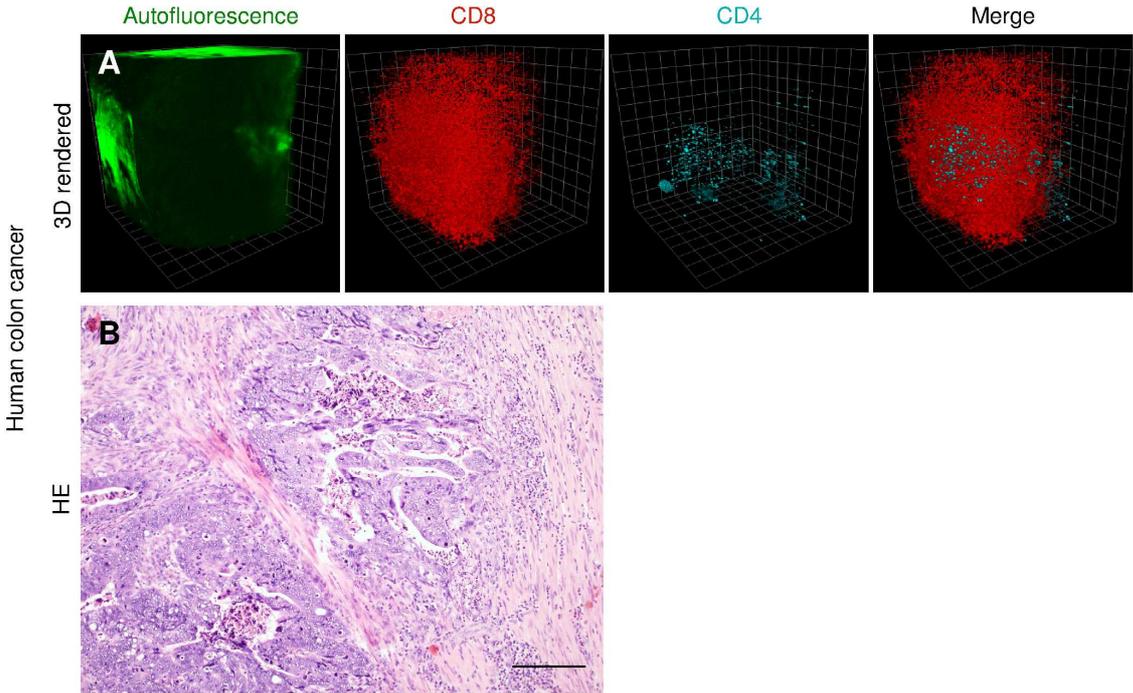
**Supplemental Figure 1: Comparison of LSFM and MPM for whole PP imaging.** (A) LSFM allowed imaging of whole PP using a x5 objective with 0.66 adapter (5 $\mu$ m increment, 300 optical sections / detection channel) within 12 minutes. (B) Due to low photon efficiency, MPM imaging of the same PP required a x20 objective (5 $\mu$ m increment, 243 optical sections / detection channel) and acquisition time took approximately 6 hours.



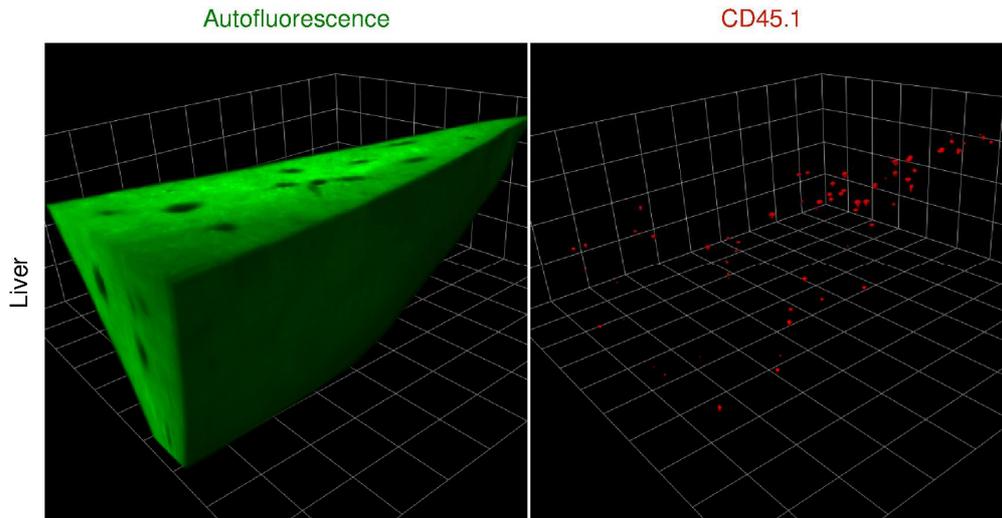
**Supplemental Figure 2: Antibody in vivo staining.** Antibodies were injected i.v. 2.5 h (CD4, CD11c) or 0.5 h (MAdCAM-1) before perfusion. Specific antibody distribution was confirmed by conventional immunofluorescence microscopy. PP of (A) an untreated mouse and (B) of a mouse intravenously injected with fluorescent labeled antibodies (MAdCAM-1 (green), CD11c (red) and CD4 (blue)). Spleen of (C) an untreated mouse (w/o antibody) and (D) a mouse injected with antibodies. Sections were stained with DAPI ex situ (Scale bar: 100  $\mu$ m). Abbreviations: ab = antibody, i.v. = intravenously, w/o = without.



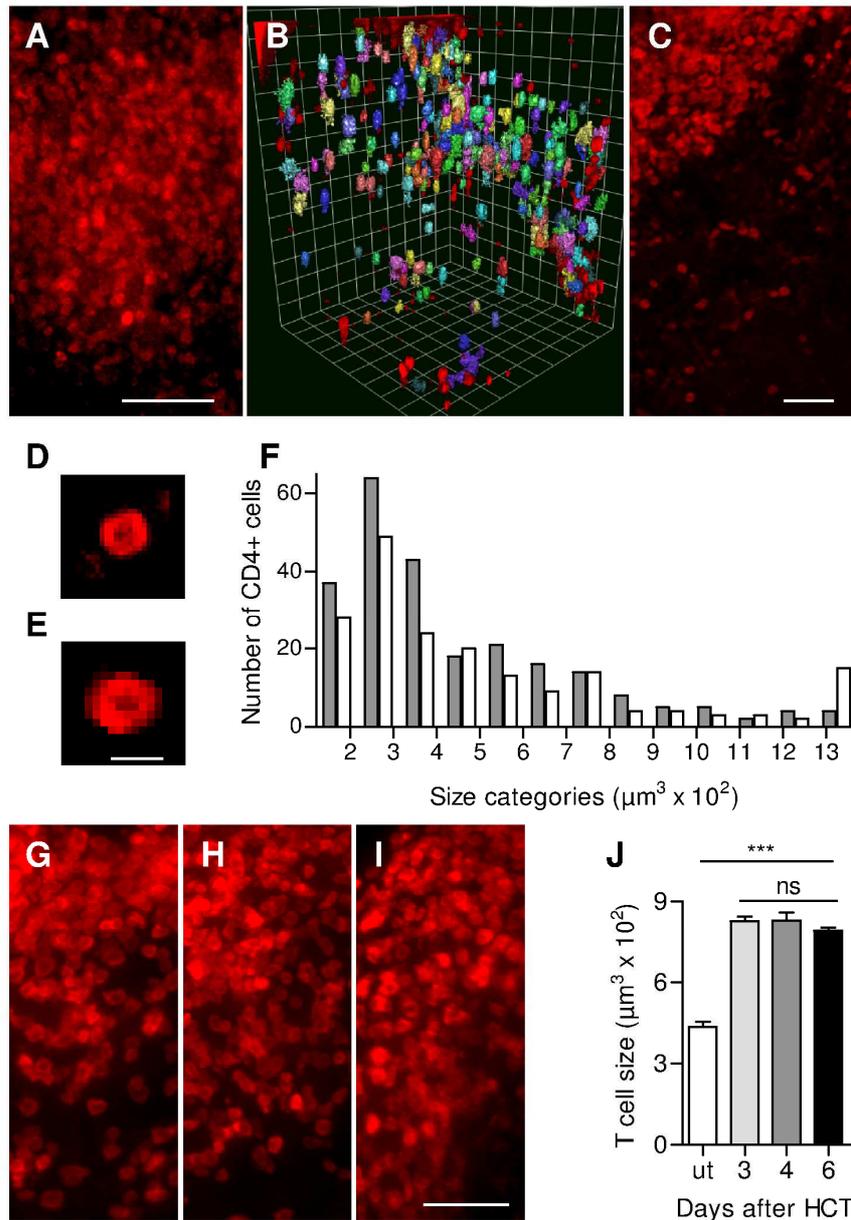
**Supplemental Figure 3: Distribution and specificity of i.v. antibody staining.** To confirm antibody i.v. staining in an untreated BALB/c mouse (A) CD4 (red) was injected 2.5 h and (B) MAdCAM-1 (red) was injected 0.5 h before perfusion and post-section staining was performed with DAPI and either with CD4 (green) or MAdCAM-1 (green), respectively (x20, scale bar: 100  $\mu$ m). Sections from mice injected with antibody (ab i.v.) were compared to sections from mice that did not receive an antibody injection (w/o antibody). (C) Tiling of a whole PP cross-section with surrounding mucosal tissue indicates specific whole tissue distribution of i.v. injected CD4 (red) antibody (x20, stitched from 30 images, scale bar: 500  $\mu$ m). (D) Confirmation by flow cytometry of i.v. antibody staining in inflamed and untreated tissues. CD4-Alexa647 antibody was injected in untreated mice or in mice at day+6 after allo-HCT (inflammation). 2.5 h after injection spleen and PPs were digested in presence of CD4-FITC antibody and subsequently analyzed with flow cytometry. Cell frequencies demonstrate efficiency of i.v. staining (> 98 %) of cellular events. Abbreviations: ab = antibody, i.v. = intravenously, w/o = without.



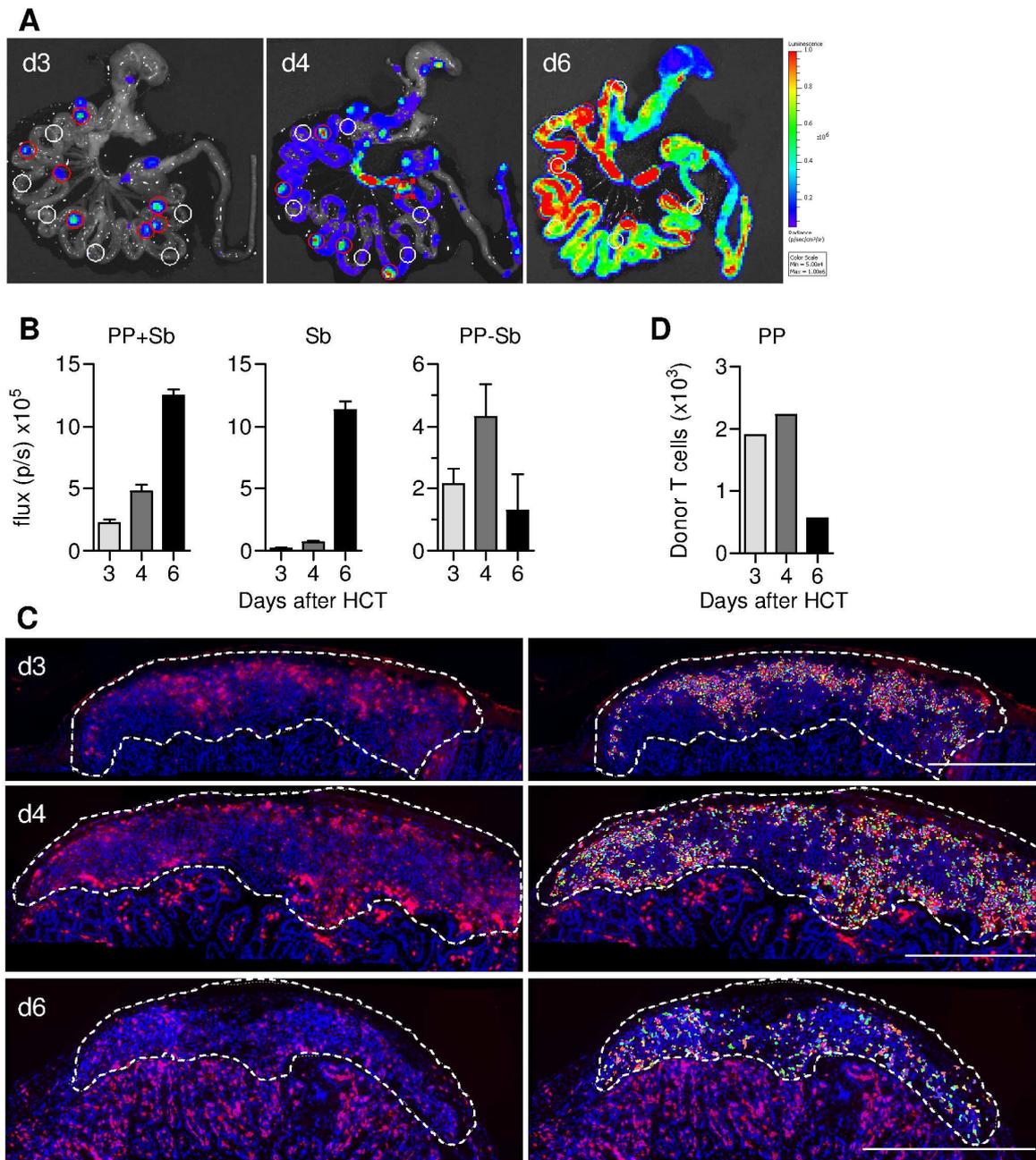
**Supplemental Figure 4: LSFM imaging of human colon cancer tissue.** (A) 3D reconstruction of a moderate differentiated human colon adenocarcinoma specimen with tumor infiltrating CD8+ T cells (red) and CD4+ T cells (cyan). (B) Histopathologic image of a sample from the same specimen stained with hematoxylin and eosin (HE) after formalin fixation and paraffin embedding confirmed the lymphocytic infiltration of the cancer tissue (scale bar: 500µm).



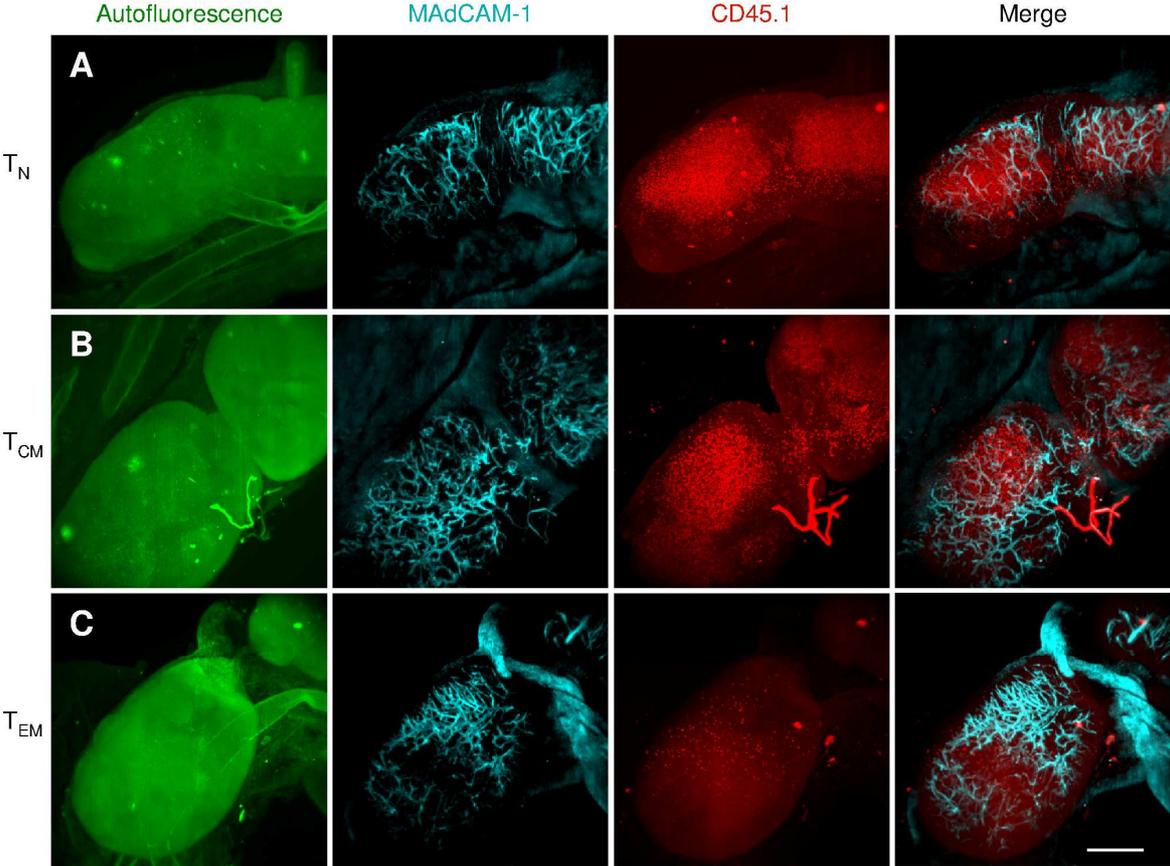
**Supplemental Figure 5: Imaging of single donor T cells in the liver 60h after allo-HCT.** Bone marrow together with luciferase<sup>+</sup> CD45.1<sup>+</sup> transgenic T cells were transplanted into myeloablatively irradiated allogeneic recipients to induce acute GVHD. Detection of single infiltrating T cells (red) in the liver (autofluorescence in green) by LSMF (unit: 246 $\mu$ m).



**Supplemental Figure 6: Quantification of T cells in intact PPs.** (A) Clustering of T cells within a PP (CD4, red) impede automated cell counting (z-projection scale bar: 50  $\mu\text{m}$ ). (B) To calculate cell volumes, single CD4+ cells are rendered (colored objects) and measured at less dense areas. (C) Comparison of cell imaging with confocal microscopy and LSFM measurements (z-projection; scale bar: 50  $\mu\text{m}$ ). Representative single cells imaged with (D) LSFM and (E) confocal microscopy using a x20 objective (z-projection scale bar: 10  $\mu\text{m}$ ). (F) Size distribution of measured CD4+ cells shown for LSFM (grey bars; mean volume 435.6  $\mu\text{m}^3 \pm 19.7$ , N = 233) and confocal microscopy (white bars; mean volume 561.1  $\mu\text{m}^3 \pm 33.4$ , N = 195). (G) Donor T cells on day +3, (H) day +4 (I) and day +6 after allo-HCT (z-projections scale bar: 50  $\mu\text{m}$ ). (J) Mean T cell volume was increased during activation (day +3 mean volume 826.9  $\mu\text{m}^3 \pm 17.2$  N = 816; day +4 mean volume 830.2  $\mu\text{m}^3 \pm 27.9$  N = 301, day +6 mean volume 788.8  $\mu\text{m}^3 \pm 15.2$  N = 976). Statistics: ANOVA, Bonferroni corrected (ns =  $P > 0.05$ , \*\*\* =  $P < 0.001$ ).



**Supplementary Figure 7: Visualizing and quantifying cellular changes within Peyer's Patches after allo-HCT.** Bone marrow together with CD90.1<sup>+</sup> luciferase<sup>+</sup> transgenic T cells were transplanted into irradiated allogeneic recipients to induce acute GVHD. (A) Ex vivo BLI was performed at the transition from GVHD initiation phase to effector phase (day +3 and day +4 after allo-HCT) and at the effector phase (day+6). (B) Emitted photons of donor T cells in PPs (red circles) or intestinal mucosa (Sb, white circles) were quantified respectively. (C) PPs from day +3, day +4 and day +6 were stained for donor T cells with CD90.1 (red) and DAPI (blue) and whole PP sections were imaged in high-resolution with IFM (x20) and stitched together. Donor T cells were automatically counted with the Software Velocity (colored objects) by determining nuclei with a high MFI in the red channel, indicating cellular staining. (D) Quantification of whole PP sections. (scale bar: 500  $\mu$ m)



**Supplemental Figure 8: Homing capacity of individual T cell subsets to mesenteric lymph nodes (mLN).** LSFM imaging of whole mLNs allowed the detection of donor T cell subsets 20 h after transfer. Accordant to the homing capacity to PPs, T cell numbers in the mLNs revealed a better homing capacity of (A) TN and (B) TCM over (C) TEM (z-projection scale bar: 500µm).

## Supplemental Videos

### Mapping immune processes in intact tissues at cellular resolution

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## Supplemental Video Legends

**Supplemental Video 1: Virtual journey through an intact Peyer's patch by multicolor LSFM.** Three-dimensional reconstruction of approximate 1500 optical sections created by LSFM. Tissue autofluorescence (green) revealed microanatomical details of the intestinal mucosa and submucosa such as villi, crypts and a PP with its subepithelial dome regions. Within the PP, CD4<sup>+</sup> T lymphocytes (red) colocalize to high endothelial venules (MAdCAM-1, cyan).

**Supplemental Video 2: Optical sectioning of an intact Peyer's patch by multicolor LSFM.** Whole PPs were imaged by creating approximate 1500 optical sections (x5 Objective, 5 µm increments) in multiple colors (Overlay: tissue autofluorescence, green; MAdCAM-1, cyan; CD4<sup>+</sup> T cells, red). The optical sections were used for 3D tissue reconstruction shown in the **Supplemental Video 1**.

**Supplemental Video 3: Multicolor LSFM of diverse murine organs.** A C57Bl/6 mouse was injected i.v. with fluorescent labeled antibodies CD31 (cyan) and LYVE-1 (red). 3D reconstruction of (a) heart, (b) testicle, (c) skin and (d) mLNs are shown.