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A plasmodium falciparum sporozoite's journey: through organs and across CD8+ T-cell challenges

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Chapter 6

Intradermal immunization with *Plasmodium berghei* late-arresting genetically attenuated sporozoites induces PD-L1 expression on regulatory macrophages and dendritic cells

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Intravenous

Intradermal

Abstract

The development of effective and long-lasting malaria vaccines remains a key goal. Late-arresting genetically attenuated sporozoite (LA-GAP SPZ) vaccines, such as PfΔmei2 (GA2), have shown strong protective potential. While GA2 delivered via mosquito bites has demonstrated up to 90% protection in humans, practical vaccine deployment will require alternative administration routes. Intravenous (IV) delivery of whole SPZ vaccines has been highly effective, but intradermal (ID) administration though easier, offers significantly reduced protection, with unclear underlying mechanisms.

In this study, we used a *Plasmodium berghei* GA2 SPZ rodent model to compare immune responses following ID and IV immunization across multiple organs. ID immunization resulted in lower frequencies of CD8⁺ tissue-resident memory T cells (Trm) in the spleen (1.3%, $p = 0.3$), lungs (8.1%, $p = 0.005$) and liver (2.5%, $p = 0.07$), along with reduced activation markers (Granzyme A, Ki67, and KLRG1). In the liver, Granzyme B ($p = <0.0001$) and perforin ($p = <0.0001$) were significantly decreased after ID immunization, indicating diminished cytotoxic potential. Importantly, ID immunization induced a regulatory myeloid phenotype in the skin and skin-draining lymph nodes, marked by low CD86 and high PD-L1 expression, potentially impairing T cell priming. A similar regulatory profile in the liver suggests systemic immunosuppression.

These findings highlight key immunological differences between ID and IV GA2 SPZ delivery and suggest that optimizing ID administration, such as adjusting injection volume to enhance sporozoite migration, may improve efficacy. Understanding how myeloid regulation and T-cell activation vary across organs is essential for enhancing malaria vaccine strategies.

Introduction

The approval of two groundbreaking malaria vaccines, RTS,S and R21, which are based on the circumsporozoite protein (CSP) found on the surface of sporozoites (SPZ, the infective form of malaria) marks a significant milestone in malaria vaccine research (WHO). However, to eliminate malaria, vaccines with higher efficacy and durability are desperately needed. Late-arresting genetically attenuated SPZ vaccination (LA-GAP) presents a promising solution by inducing both cellular immune responses and CSP-targeting antibodies, resulting in high-level protection even after a single immunization in malaria-naïve individuals [1]. This LA-GAP vaccine, known as Pf Δ mei2 (GA2), lacks the mei2-like RNA gene, allowing sporozoites to travel through the bloodstream, infect hepatocytes and develop into liver schizonts similar to wild-type SPZ [2]. However, during the late liver stage, the GA2 liver schizont arrests its development, in contrast to the wild-type liver schizont, which bursts and releases merozoites that infect red blood cells [3, 4]. Previous comparisons between early (GA1) and late (GA2) genetically arresting parasites in humans have demonstrated that development until the late liver stage is essential for achieving high-level protection [5]. Moreover, this protection was associated with T cell activation in peripheral blood mononuclear cells of these individuals, particularly CD4⁺ pluripotent T-cells producing multiple Th1-type cytokines [1, 5].

Whereas in these human trials CD8⁺ T cell activation could not be detected in the peripheral blood circulation, animal models have shown that liver tissue-resident CD8⁺ T cells (CD8⁺ Trm cells) are associated with protection from liver-stage malaria [6-8]. It is thought that these cells cannot be detected in the peripheral circulation due to their low frequency, whereas they are increased in livers of protected animals. It is hypothesized that CD8⁺ Trm cells can recognize and kill malaria-infected hepatocytes, as indicated by their increased expression of granzyme B, perforin, and the cytokine IFN γ [9, 10]. However, these liver CD8⁺ Trm cells require priming before they can specifically recognize and kill malaria-infected hepatocytes [11, 12]. Naïve CD8⁺ T cells can be primed in secondary lymphoid organs by antigen-presenting cells (APC) like macrophages and dendritic cells [13-15]. After priming, naïve CD8⁺ T cells differentiate into CD8⁺ Trm cells and migrate to the liver [6, 7, 16]. Priming of naïve CD8⁺ T cells can occur in different organs before SPZ infect hepatocytes [9].

After a single mosquito bite, roughly 100 SPZ are deposited in the skin [17], where antigens can be processed by APC such as macrophages and dendritic cells. Subsequently, around 15-20% of SPZ migrate to the skin-draining lymph node (skin dLN)

while $\pm 20\%$ enter the bloodstream [17-19], traveling through the lungs and spleen before reaching the liver to infect hepatocytes [20]. In the meantime, skin-resident dendritic cells that have encountered SPZ process the antigens and migrate to the skin dLN [14]. Previous studies have demonstrated that through these routes CD8⁺ T cells, can be primed by APC in the skin dLN and migrate to the liver [11, 13, 21].

Although the administration of GA2 SPZ in humans through mosquito bites has proven effective [1, 5], alternative routes of administration will be required to develop this concept into a vaccine. Interestingly, for other whole SPZ vaccines such as the irradiated SPZ, it has been shown that intradermal (ID) administration results in much lower protection compared to intravenous (IV) administration in humans [22, 23]. This reduced efficacy was initially thought to be related to the number of SPZ entering the liver. Using rodent bioluminescent parasites in animal models, it was indeed shown that IV-injected SPZ reach the liver more efficiently as compared to the ID-administered SPZ. However, when we corrected for the lower parasitic liver load by increasing the ID-administered parasites 5-fold, we found that GA2 *Plasmodium yoelii* (Py) SPZ immunized mice are after ID administration less protected as compared to IV administration [24]. The mechanism behind this observation remained unclear. These findings suggest that the altered biodistribution of SPZ after ID administration in organs other than the liver negatively impacts protective responses in the liver. Therefore, we aimed to investigate whether GA2 SPZ following ID or IV administration in organs other than the liver could negatively impact T-cell responses in the liver.

Methods

Mosquito production

Mosquitoes from a colony of *Anopheles stephensi* (line Nijmegen SDA500) were used to obtain sporozoites. Larval stages were reared in water trays at a temperature of $28 \pm 1^\circ\text{C}$ and a relative humidity of 80%. Adult females were transferred to incubators with a temperature of $26 \pm 0.2^\circ\text{C}$ and a relative humidity of 80%. For all the experiments, 3- to 5-day-old mosquitoes were used. The *Plasmodium berghei* (Pb) infected mosquitoes were maintained at 21°C at 80% relative humidity.

Experimental animals, *P. berghei* parasite line

Female OF1 mice and female and male C57BL/6J mice (Charles River Laboratories, France) of 4-6 weeks old were acclimatized for one week prior to the experiment. Mice were housed between 4-5 mice per cage in ventilated cages with autoclaved aspen woodchip, fun tunnel, wood chew block and nestlets (12:12 hour light-dark cycle; $21 \pm 2^\circ\text{C}$; relative humidity of $55 \pm 10\%$). During the experiment mice were fed with

commercially prepared autoclaved dry rodent diet pellets and water, both available ad libitum. All animal experiments were approved with license 11600202216547 by the Competent Authority after advice on ethical evaluation by the Animal Experiments Committee Leiden and were performed in accordance with the Experiments on Animals Act (Wod, 2014) the applicable legislation in the Netherlands in accordance with the European guidelines (EU directive no 2010/63/EU). The study was executed in a licensed establishment for experimental animals, and conducted in accordance with the ARRIVE guidelines.

Humane endpoints: the animals/body condition was thoroughly examined daily. Animals were humanely sacrificed in case the following defined endpoints are reached: visible pain (abnormal posture and/or movement), abnormal behaviour (isolation, abnormal reaction to stimuli, no food and water intake). If distress of the animals was observed by the animal caretakers, this was reported to the investigators and according to the therefore mentioned criteria, the animals were taken out of the experiment and euthanized. In all experiments no mice were euthanized before termination of the experiment and no mice died before meeting criteria for euthanasia.

The LA-GAP *Pb*Δ*mei2*Δ*lisp2* parasite (2900cl3, mutant RMgm-4937; www.pberghei.eu), which is genetically attenuated by the deletion of the meiosis inhibited 2 (*mei2*) and liver-specific protein 2 (*lisp2*) genes was used [2]. Feeding of *Anopheles stephensi* mosquitoes was performed as described previously [42].

***In vivo* immunization of mice with *Pb* SPZ**

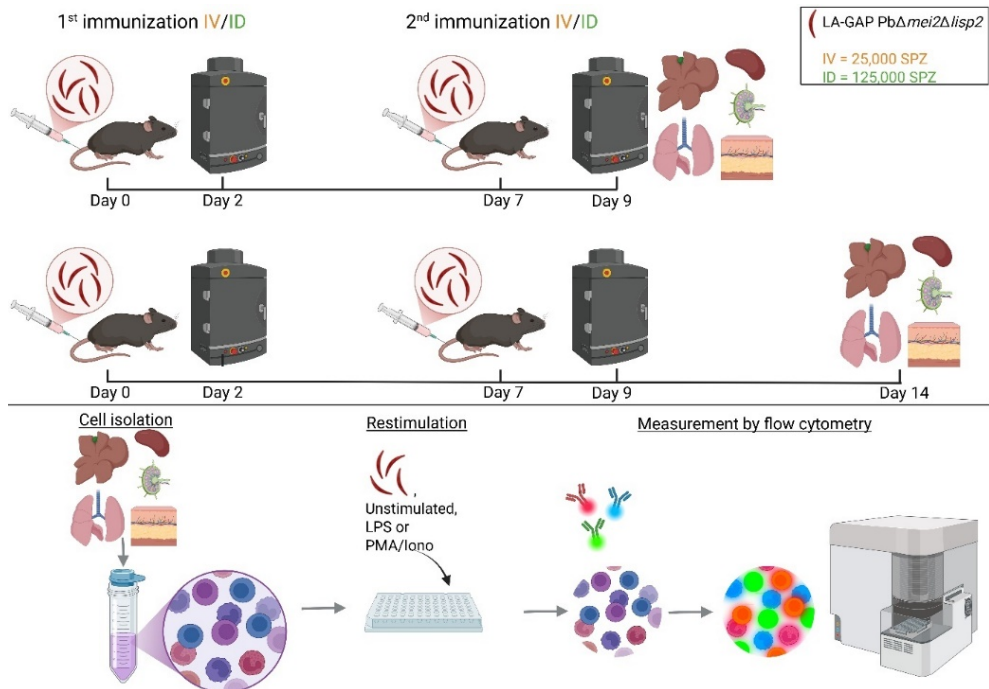
The study including data acquisition was performed blinded. The study was deblinded after analysis of the data.

One day prior to immunizations, mice were randomly divided into three or four different groups (4-5 mice per group, maximum of 20 mice per experiment). On the day of immunizations, salivary glands (21-23 days post blood meal) of *P. berghei* LA-GAP *Pb*Δ*mei2*Δ*lisp2* (*Pb* GA2 SPZ) infected mosquitoes were dissected in cold RPMI 1640 glutamax (Thermo Fisher). As a control, salivary glands from the same batch of uninfected mosquitoes, hence referred to as salivary gland extract (SGE), were dissected in cold RPMI 1640 glutamax (Thermo Fisher). Immediately after dissection the glands were crushed and homogenized and the total number of *Pb* GA2 SPZ was counted. Intravenous immunization in the tail vein was performed after warming the mice under a heat lamp set at 35°C to dilate the veins. For intravenous immunization, 25,000 *Pb* GA2 SPZ were inoculated in 200 μL RPMI 1640 glutamax (Thermo Fisher). For

intradermal immunizations, the mice were anesthetized with isoflurane and 125,000 live *Pb* GA2 SPZ were inoculated in 20 μ L RPMI 1640 glutamax (Thermo Fisher) in the left shaved upper thigh. For the SGE control, an equal amount of SGE was injected in 200 μ L IV or 20 μ L ID RPMI 1640 glutamax (Thermo Fisher) and for the medium control, 200 μ L IV or 20 μ L ID RPMI 1640 glutamax (Thermo Fisher). Second immunizations were given 7 days after the first. All injections were given between 1 pm and 3 pm (methods Figure 6).

Determination of parasite liver load after 1st and 2nd immunization by real-time *in vivo* imaging

After 44 hours post-immunization the parasitic liver load was determined through bioluminescent imaging. Imaging was performed using the IVIS Lumina II Imaging System (Perkin Elmer Life Sciences, Waltham, USA) [43] 8 minutes after a subcutaneous injection with D-luciferin dissolved in PBS (100 mg/kg; Caliper Life Sciences, USA). Quantitative analysis of the bioluminescence of whole bodies was performed by measuring the luminescence signal intensity using the ROI (region of interest) settings of the Living Image[®] 4.4 software. Blood-stage breakthroughs were checked at 5-6 days post 1st immunization and 2nd immunization. If any parasites were discovered in the blood the mice were sacrificed and eliminated from the experiment (methods Figure 6).



Methods Figure 6. Experimental set-up. Overview of the timeline of mice immunizations via IV or ID. 2 days after immunizations, the parasitic liver-load was determined by IVIS. For the early immune response (myeloid activation), organs were harvested 2 days post 2nd immunization ID/IV. For the late immune response (T cell activation), cells were harvested 7 days post 2nd immunization ID/IV. Cells were isolated from the organs and restimulated with GA2 SPZ, LPS, PMA/Iono or unstimulated. After 4 hours of incubation, cells were stained with fluorescent antibodies and measured by flow cytometry.

Organ harvesting and processing

Two days or seven days after the final immunizations, mice were anesthetized via intraperitoneal injection of 10% ketamine (Dechra Pharmaceuticals, Northwich, UK) with 20 mg/mL xylazine (Alfasan, Woerden, The Netherlands). Blood was collected via retro-orbital puncture using a glass capillary tube and transferred into 1.5 mL Eppendorf tubes pre-coated with heparin, and put on ice. The blood was spun down at 4°C, plasma was collected and stored at -80°C. The liver and lungs were perfused by slowly injecting 20 mL cold PBS via the heart, followed by dissection of the liver, lungs and spleen after IV immunizations, and liver, lungs, spleen, skin and skin dLN after ID immunizations, and placed in sterile RPMI 1640 glutamax (Thermo Fisher) on ice. The mice were killed by exsanguination.

Livers were processed via mincing with a blade and placed in a 50 mL tube with 20 mL RPMI 1640 glutamax containing 1 mg/mL Collagenase IV (Sigma-Aldrich) and 2,000 U/mL DNase I (Sigma-Aldrich) and incubated for 45 minutes at 37°C, mixing once during incubation. After incubation, tubes were placed on ice and poured through a 100-micron filter (BD) and washed with 20 mL PBS supplemented with 1% Fetal Calf Serum (FCS) and 2.5 mM ethylenediamine tetra-acetic acid (EDTA, Sigma-Aldrich). The tubes were spun down at 1,500 rpm for 10 minutes at 4°C and supernatants were gently taken off, after which PBS supplemented with 1% FCS and 2.5 mM EDTA was added to the pellets and tubes were spun down at 50 g for 3 minutes at 4°C. To separate the immune cells from the hepatocytes, the supernatant, which contains immune cells, was gently removed. The immune cells were spun down for 10 minutes at 1,600 rpm at 4°C. The supernatant was discarded, and 3 mL sterile PBS supplemented with 0.15M NH₄Cl; 1mM KHCO₃; 0.1 mM Na₂EDTA was added for 2 minutes to lyse red blood cells, followed by the addition of 7 mL PBS supplemented with 1% FCS and 2.5 mM EDTA (Sigma-Aldrich). The tubes were spun down for 10 minutes at 1,600 rpm at 4°C. The supernatant was discarded, and the pellet was resuspended in 10 mL PBS supplemented with 0.5% bovine serum albumin (Fraction V, Roche) and 10 mM EDTA (Sigma-Aldrich) and spun down for 10 minutes at 1,200 rpm at 4°C. The supernatant was removed, and 35 µL CD45 MicroBeads (Miltenyi Biotec) were added and incubated for 15 minutes in the fridge. After incubation, the left-over beads were washed off by adding 10 mL PBS supplemented with 0.5% bovine serum albumin (Fraction V, Roche) and 10 mM EDTA and spun down for 10 minutes at 1,600 rpm at 4°C. The pellets were resuspended in 5 mL PBS supplemented with 0.5% BSA (Fraction V, Roche) and 10 mM EDTA and run through a pre-wetted LS magnetic column according to protocol (Miltenyi Biotec). The columns were washed with RPMI 1640 glutamax, 5% FCS, 0.1% β-mercaptoethanol, 100U/mL penicillin, and 100 µg/mL streptomycin (culture medium) to collect the CD45⁺ cells. The tubes were spun down for 10 minutes at 1,600 rpm at 4°C, supernatants were taken off and resuspended in 5 mL washing medium, and viable cells were counted using trypan blue and a Bürker counting chamber.

To process lungs, the lungs were chopped into small pieces using a scalpel and transferred to a 15-mL tube. To digest the tissue 5 mL RPMI 1640 glutamax supplemented with 1 mg/mL Collagenase IV (Sigma-Aldrich), 2 µL/mL DNase (Sigma-Aldrich), and 2 µL/mL of 1M CaCl₂ (Sigma-Aldrich) and incubated for 30 minutes at 37°C. After incubation, 10 mL cold RPMI 1640 glutamax with 5% FCS was added and poured through a 100-micron filter (BD). A 1 mL syringe was used to mash the digested tissue and washed with 20 mL cold RPMI 1640 glutamax with 5% FCS. The tubes were

spun down for 10 minutes at 1,600 rpm at 4°C, and 2 mL sterile PBS supplemented with 0.15M NH₄Cl; 1mM KHCO₃; 0.1 mM Na₂EDTA was added for 2 minutes to lyse the red blood cells. After 2 minutes 8 mL of washing medium was added, and tubes were spun down for 10 minutes at 1,600 rpm at 4°C. Pellets were resuspended in 5 mL of washing medium, and viable cells were counted using trypan blue and a Bürker counting chamber.

To process the spleen and skin-draining lymph nodes, the organs were crushed with the back of a syringe. A two-fold concentrated enzyme mix in RPMI was added, end concentration of 1 mg/mL Collagenase D (Roche) and 2000 U/mL DNase (Sigma), was added and samples were incubated for 20 minutes at 37°C. After incubation, the cells were poured through a 100-micron filter (BD) and washed with 10 mL cold RPMI medium. The cells were spun down for 10 minutes at 1600 rpm at 4°C. The supernatant was discarded, and 2 mL cold sterile PBS supplemented with 0.15M NH₄Cl; 1mM KHCO₃; 0.1 mM Na₂EDTA was added for 2 minutes to lyse red blood cells. Lysis was stopped by the addition of 8 mL cold RPMI, and the pellet was resuspended in 10-20 mL cold RPMI 1640 glutamax (Thermo fisher) supplemented with 100U/mL penicillin and 100 µg/mL streptomycin (Sigma-Aldrich), 0,1 % β-mercaptoethanol, and 5% FCS and counted using trypan blue and a Bürker counting chamber.

To process the skin, the skin was cut into small pieces and incubated with enzymes P, D and A, diluted in buffer L according to the manufacturer's instructions for the whole skin dissociation kit (Miltenyi). Samples were incubated for 3 hours at 37°C. After incubation, 500 µL of cold RPMI 1640 glutamax (Thermo fisher) supplemented with 100U/mL penicillin and 100 µg/mL streptomycin (Sigma-Aldrich) and 5% FCS was added. The tubes were spun according to the tumor program with the MACS dissociator (Miltenyi) and centrifuged for 10 minutes at 1200 rpm at 4°C. The cells were poured through a 100-micron filter (BD) and washed with 10 mL cold RPMI medium. The cells were spun down for 10 minutes at 1200 rpm at 4°C, and the pellet was resuspended in 1 mL cold RPMI 1640 glutamax (Thermo fisher) supplemented with 100U/mL penicillin and 100 µg/mL streptomycin (Sigma-Aldrich), 0,1 % β-mercaptoethanol, and 5% FCS. Cell counts and viability were assessed using trypan blue exclusion and a Bürker counting chamber.

Cell stimulation and analysis

A total of 200,000 cells per well were added. For specific restimulation, live *Plasmodium berghei* GA2 SPZ in a concentration of 25,000/mL (5,000 per 200,000 cells) in culture

medium: RPMI 1640 glutamax (Thermo fisher) supplemented with 100U/mL penicillin and 100 µg/mL streptomycin (Sigma-Aldrich), 0,1 % β-mercaptoethanol and 5% FCS, were added to each well and spun down for 4 minutes at 1,200 rpm at 4°C. For the specific restimulation, 0.1 µg/mL Phorbol 12-myristate 13-acetate (PMA) + 1 µg/mL ionomycin (Iono) (Sigma-Aldrich) in culture medium was added. To measure intracellular cytokine expression, 10 µg/mL Brefeldin A (Sigma-Aldrich) was directly added. All plates for flow cytometry were incubated for 4 hours at 37°C with 5% CO₂.

After 4 hours (flow cytometry measurement), the cells were transferred into a V-bottom plate and washed with cold PBS. Cells were stained with live/dead marker Zombie NIR (Thermo Fisher), fixed with eBioscience FOXP3/Transcription Factor Fixation/Permeabilization kit, and stained with different markers for 3 different panels: (1) Myeloid panel (Table 1), (2) T-cell panel (Table 2), or (3) Advanced T-cell panel (Table 3, implemented halfway through the experiments). To all panels, Fc-block (BD bioscience), True-Stain Monocyte Blocker (BioLegend), and Brilliant Violet buffer (Thermo Fisher) was added. The cells were measured by flow cytometry using Aurora 5 laser (Cytek Bioscience B.V., Amsterdam) and analyzed using Spectroflow (Cytek Bioscience B.V.), FlowJo version 10.8 (FlowJo LLC), and R-studio version 1.4.1717. For gating strategy see supplementary figure 2. Of the CD4⁺ and CD8⁺ T-cells were gated into effector memory T cells (Tem, CD44^{hi} CD62L⁻ CD69⁻), resident memory -cells (Trm, CD44^{hi} CD62L⁻ CD69⁺) and DN cells into DN1, DN2, DN3 and DN4 based on expression of CD44 and CD25 expression.

Table 1. Myeloid panel

Target	Antibody clone	Fluorochrome
MHCII	2G9	BUV395
PD-L1	B7-H1	BUV737
CD11c	N418	BV421
Siglec F	E50-2440	BV480
CD40	3-23	BV510
CCR7	4B12	BV605
XCR1	ZET	BV650
F4/80	T45-2342	BV711
CD45	30-f11	BV785
CD70	FR70	FITC
Ly6C	HK1.4	Percp-cy5.5
CD80	16-10A1	PE

CD64	X54-5/7.1	PE-Dazzle
B220	RA3-6B2	PE-Cy5
CD11b	M1/70	PE-Cy7
CD206	C068C2	APC
TIM4	F31-5G3	AF647
CD86	GL1	AF700
NK1.1	PK135	APC-Cy7

Table 2. T-cell panel

Target	Antibody clone	Fluorochrome
CD3	17A2	BUV661
CD45	30-F11	BUV805
CXCR3	CXCR3-173	Super Bright 436
Ki67	SoIA15	eFluor 506
$\gamma\delta$ T-cells	GL3	BV605
CD4	GK1.5	BV650
PD-L1	10F.9G2	BV711
PD1	29f.1A12	BV785
Perforin	eBio0mak	FITC
CD44	1M7	AF532
CD8	53-6.7	PerCP
Granzyme A	GrA-368.5	PerCP-ef710
CD137	4-1BB	PE
CD69	H1.2F3	PE-cf594
IFN γ	XMG1.2	PE-Cy5
KLRG1	2F1	PE-Cy5.5
TNF α	MPG-XT22	PE-Cy7
FOXP3	FjK-16s	APC
Granzyme B	NGZB	efluor 660
CD25	PC61.5	AF700

Table 3. Advanced T-cell panel

Target	Antibody clone	Fluorochrome
MHCII	2G9	BUV395
CD25	PC 61.5	BUV563
CD3	17A2	BUV661
CD62L	Mel-14	BUV737

CD45	30-f11	BUV805
CXCR3	CXCR3-173	Super bright 436
$\gamma\delta$ T-cells	GL3	BV605
CD4	GK1.5	BV650
CD11c	N418	BV711
PD1	29f.1A12	BV785
CD40L	SA047C3	FITC
CD44	IM7	AF532
CD8	53-6.7	PerCP
CD64	X54-5/7.1	PerCP-eFluor 710
CD69	H1.2F3	PE-cf594
IFN γ	XMG1.2	PE-Cy5
KLRG1	2F1	PE-Cy5.5
TNF α	MPG-XT22	PE-Cy7
CD86	GL1	AF700
NK1.1	PK135	APC-Cy7

Statistical analysis

Data was analysed using Spectroflow (Cytex Bioscience B.V., Amsterdam) and FlowJo version 10.8 (FlowJo LLC, Ashland, OR, USA), R studio version 4.3.1 and Adobe Illustrator 2023. Sample size estimation was performed with a power analysis using an alpha of 0.05 and a power of 90%. Statistical analyses were performed using one-way ANOVA with multiple comparisons with Bonferroni correction or an independent t-test with R studio version 4.3.1. Significance was defined as a p-value of less than 0.05. Data subjected to parametric statistical analyses had its normality confirmed beforehand.

Results

Distribution of immune cells in the spleen, lung and liver after intradermal or intravenous sporozoite GA2 immunization

First, we observed the parasitic liver load after IV or ID immunization. Here we found a negative parasitic liver load after the 2nd IV GA2 SPZ immunization but a comparable parasitic liver load between the 1st and 2nd ID GA2 SPZ immunization (Sup Fig. 1). To compare the immune response after ID immunization or IV immunization with GA2 SPZ, we analyzed shifts in the number of immune cells in the spleen, lung and liver two days after the 2nd and last immunization. Here, we observed that IV GA2 SPZ immunization leads to marked increases in total numbers of immune cells in the liver, and small increases in the spleen but no changes in the lungs (Fig. 1A). After ID immunization,

these increases were not detected, resulting in a statistically significant lower number of cells in the ID vs IV condition in the liver ($p = 0.0003$). Furthermore, only in the liver after IV immunization did we observe a significant difference between SGE and LA-GAP ($p = <0.0001$, Fig 1A). This is in line with the absolute number of various immune cell populations, where we primarily observe a significant increase in the liver following IV immunization compared to ID immunization (Sup Fig. 3). Next, after identifying immune cell subsets by antibody staining and flow cytometry, we investigated the overall kinetics of immune cell distribution in the spleen, lungs and liver after ID versus IV GA2 SPZ immunization. Looking at cell immune subsets 2 days after the 2nd immunization, we observed slightly lower total numbers of $\gamma\delta$ T cells (5.5%, $p = 0.05$) and $CD3^+$ T cells (8.5%, $p = 0.0003$) in the lungs and $CD3^+$ T cells in the liver (9.3%, $p = 0.0001$) after ID GA2 SPZ immunization compared with IV GA2 SPZ immunization (Fig. 1B). However, 7 days after the 2nd immunization of the total number of $CD3^+$ T cells, the $CD4^+$ (2.2%, $p = 0.01$) and double negative (DN) T cells (3.2%, $p = 0.04$) in the spleen, and DN (7.4%, $p = 0.01$) and $CD8^+$ T cells (6.1%, $p = 0.06$) in the lungs were most affected with a lower number after ID GA2 SPZ immunization compared with IV GA2 SPZ immunization. In contrast, these differences were not observed in the liver. Interestingly, $CD4^+$ T cells in the lungs exhibited the opposite trend, showing an increase after ID immunization (8.4%, $p = 0.02$, Fig. 1C). Last, we focused on the differentiation markers within different lineages, e.g. within the $CD4^+$, $CD8^+$ and DN T cells 7 days after the 2nd GA2 SPZ immunization. Here we observed a lower percentage of $CD4^+$ Trm in the spleen (4.2%, $p = 0.01$), lungs (10.7%, $p = 0.0006$) and liver (4.2%, $p = 0.2$) after ID GA2 SPZ immunization compared with IV GA2 SPZ immunization (Fig. 1D). For the $CD4^+$ Tem we also found lower numbers in the liver (3.3%, $p = 0.06$) after ID immunization compared with IV immunization, but not in the spleen and lungs (Fig. 1D). For the $CD8^+$ T cells we similarly observed a lower percentage Trm in the spleen (1.3%, $p = 0.3$), lungs (8.1%, $p = 0.005$) and liver (2.5%, $p = 0.07$) and also Tem in the spleen (27.3%, $p = 0.006$), lungs (13.2%, $p = 0.002$) and liver (19.6%, $p = 0.0001$) after ID GA2 SPZ immunization compared with IV GA2 SPZ immunization (Fig. 1E). Interestingly, for the DN T cells we found a higher percentage of DN1 T cells in the spleen (13%, $p = 0.004$), lungs (12%, $p = 0.01$) and liver (16%, $p = 0.01$) after ID GA2 SPZ immunization compared with IV GA2 SPZ immunization (Fig. 1F). This indicates that the differentiation of DN T cells after ID SPZ immunization is less progressed at 7 days after 2nd immunization compared with IV SPZ immunization. Overall, these findings show a lower amount of $\gamma\delta$ T cells, $CD3^+$ T cells, $CD4^+$ Tem, $CD8^+$ Tem and Trm and more DN1 T cells after ID SPZ immunization compared with IV SPZ immunization in the spleen, lungs and liver.

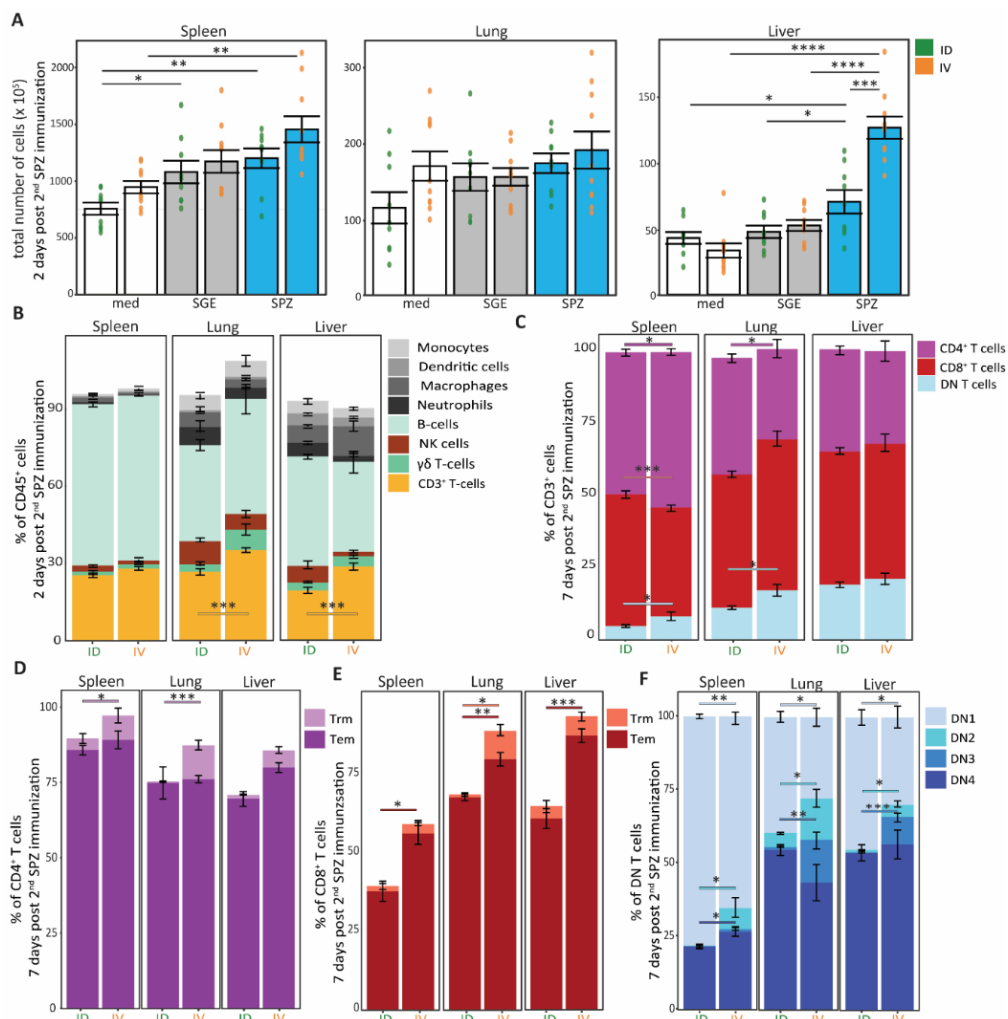


Figure 1. Distribution of immune cells in the spleen, lungs and liver after intradermal (ID) or intravenous (IV) SPZ GA2 immunization. (A) Total number of immune cells in the spleen, lungs or liver, 2 days after 2nd ID (green) or IV (orange) immunization *ex vivo* with medium (white), salivary gland extract (SGE) (grey) or SPZ (blue). **(B)** Kinetics of immune cells in the spleen, lungs and liver 2 days after 2nd SPZ ID or IV immunization followed by *in vitro* restimulation with LA-GAP SPZ. **(C)** CD3⁺ T cell distribution in the spleen, lungs and liver, 7 days after 2nd SPZ ID or IV immunization followed by *in vitro* restimulation with LA-GAP SPZ. **(D)** Distribution of Trm and Tem CD4⁺ T cells in the spleen, lungs and liver, 7 days after 2nd SPZ ID or IV immunization. **(E)** Distribution of tissue-resident memory (Trm) and effector memory (Tem) CD8⁺ T cells in the spleen, lungs and liver, 7 days after 2nd SPZ ID or IV immunization followed by *in vitro* restimulation with LA-GAP SPZ. **(F)** Distribution of double negative 1 (DN1), DN2, DN3 and DN4 T cells in the spleen, lungs and liver, 7 days after 2nd SPZ ID or IV immunization followed by *in vitro* restimulation with LA-GAP SPZ.

SPZ ID or IV immunization followed by *in vitro* restimulation with LA-GAP SPZ. ID immunization n=9 per group, IV immunization n=10 per group divided over 2 independent experiments. Statistical significance between groups was assessed by one-way ANOVA with multiple comparisons or an independent t-test. *p<0.05, **p<0.005, ***p<0.0005, and ****p<0.0001.

Activation of T cells in the spleen, lungs and liver after intravenous or intradermal GA2 SPZ immunization

Having found a difference in CD4⁺ Tem, CD8⁺ Tem and Trm and DN1 T cells after ID immunization, we next investigated the activation of T cells in the spleen, lungs and liver after ID and IV immunization at the later timepoint: 7 days post 2nd GA2 SPZ immunization. An optimized t-Distributed Stochastic Neighbor Embedding (opt-SNE) analysis based on different T cell markers revealed activation of two cell clusters after IV GA2 SPZ immunization, which were not activated after ID GA2 SPZ immunization. We identified that these clusters consisted of CD8⁺ T cells and DN T cells which highly expressed Granzyme A, Ki67 and KLRG1 (Fig. 2A+B+C). Therefore, we analyzed the percentage of Granzyme A⁺, Ki67⁺ and KLRG1⁺ CD8⁺ T cells and DN T cells. Here we observed an overall lower percentage of expression on DN T cells after ID GA2 SPZ immunization compared with IV GA2 SPZ immunization in markers for Granzyme A⁺ (spleen 6.1%, lung 19.3%, liver 13.8%), Ki67⁺ (spleen 5.6%, lung 8.8%, liver 11.6%) and KLRG1⁺ (spleen 6.3%, lung 13.6%, liver 8.9%) CD8⁺ T cells and Granzyme A⁺ (spleen 11.9%, lung 26%, liver 20.1%), Ki67⁺ (spleen 2.4%, lung 8.6%, liver 15.2%) and KLRG1⁺ (spleen 9.1%, lung 12.9%, liver 12.4%) respectively. Interestingly, after SGE immunization, we observed minimal differences in the percentages of Granzyme A⁺, Ki67⁺ and KLRG1⁺ CD8⁺ T cells and DN T cells between ID and IV immunization, which indicates that the immune differences are GA2 SPZ specific (Fig. 2D+E). In parallel, $\gamma\delta$ T cells exhibited significantly lower expression of Ki67 (p = 0.02) and KLRG1 (p = 0.04) in the lungs as well as a reduced Ki67 expression in the liver (p = 0.02) after ID GA2 SPZ immunization compared with IV (Sup Fig. 4B), indicating that these cells are phenotypically also less activated after ID immunization. To further investigate the killing pathway of T cells we looked into the expression of Granzyme B and perforin. In the spleen, no differences were observed between ID and IV GA2 SPZ immunization. However, in the lung expression of Granzyme B on both CD8⁺ T cells (p = 0.0003) and DN T cells (p = 0.07) and in the liver expression of both Granzyme B and perforin on CD8⁺ T cells (p = <0.0001, p = <0.0001) and DN T cells (p = 0.06, p = <0.0001) was lower after ID GA2 SPZ immunization compared with IV GA2 SPZ immunization (Fig. 2F). This is in line with the decreased expression of Granzyme B on CD4⁺ T cells in the liver after

ID immunization compared with IV immunization ($p = 0.001$, Sup Fig. 4A). Last, we investigated the expression of cytokines IFN γ and TNF on CD8 $^+$ T cells and DN T cells. Here we did not observe any differences in cytokine expression between ID and IV GA2 SPZ immunization in any of the organs (Fig. 2G).

In conclusion, ID GA2 SPZ immunization compared to IV delivery results in suppressed activation of the CD8 $^+$ T cell and DN T cell compartments, primarily in the lungs and liver, visible by the number of Granzyme A $^+$, Ki67 $^+$ and KLRG1 $^+$ CD8 $^+$ T cells and DN T cells. Furthermore, in the liver, ID GA2 SPZ immunization leads to decreased perforin and Granzyme B expression compared to IV GA2 SPZ immunization.

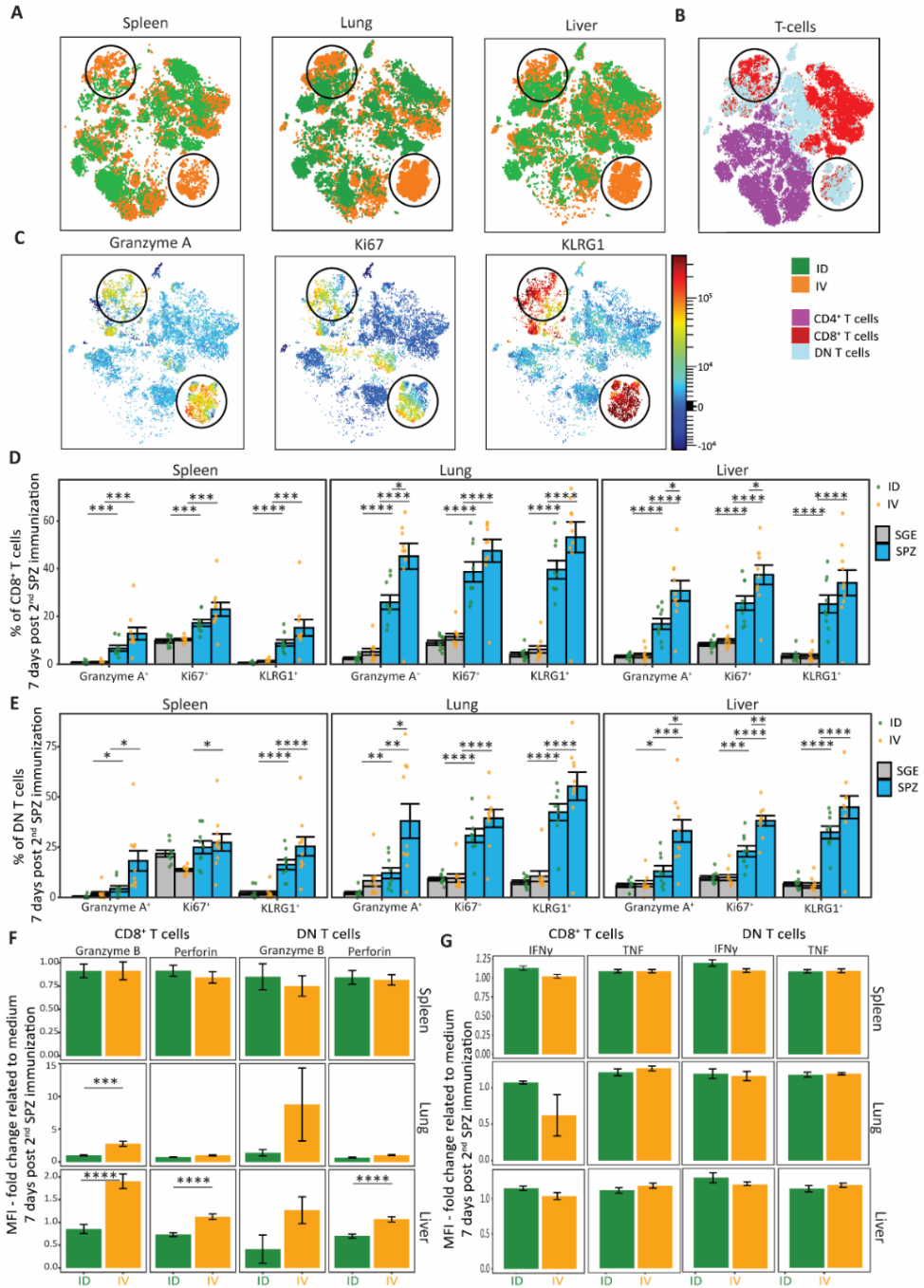


Figure 2. T cell activation in the spleen, lung and liver after intradermal (ID) or intravenous (IV) GA2 immunization. (A) Opt-SNE of different activation markers on T cells after ID

(green) or IV (orange) immunization in the spleen, lung and liver. **(B)** Identification of CD4⁺ (purple), CD8⁺ (red) and DN T cells (light blue) in the opt-SNE. **(C)** Expression of Granzyme A, Ki67 and KLRG1 in the opt-SNE. **(D + E)** Percentage of Granzyme A⁺, Ki67⁺ and KLRG1⁺ CD8⁺ **(D)** or DN T cells **(E)** in the spleen, lung and liver after SGE (grey) or GA2 SPZ (blue) ID (green dots) or IV (orange dots) immunization. **(F + G)** Fold change relative to medium expression of Granzyme B, Perforin **(F)** and IFN γ and TNF **(G)** on CD8⁺ T cells and DN T cells in the spleen, lung and liver. ID GA2 SPZ immunization in green, IV GA2 SPZ immunization in orange. ID immunization n=9 per group, IV immunization n=10 per group divided over 2 independent experiments. All conditions were *in vitro* restimulated with LA-GAP SPZ. Statistical significance between groups was assessed by one-way ANOVA with multiple comparisons or an independent t-test. *p<0.05, **p<0.005, ***p<0.0005, and ****p<0.0001.

Activation of myeloid cells in the spleen, lungs and liver after intravenous or intradermal GA2 SPZ immunization

To investigate the underlying mechanism behind the reduced number of activated T cells after ID GA2 SPZ immunization, we analyzed the impact of ID GA2 SPZ immunization on the activation of myeloid cells in the spleen, lungs and liver compared with IV GA2 SPZ immunization at the early timepoint 2 days post 2nd immunization. Principal Component Analysis (PCA) revealed distinct immunological profiles between the two routes of administration across the spleen, lungs and liver (Fig. 3A). To identify the markers and cell types driving these differences, we focused on the four markers with the highest PCA loadings, which contribute most strongly to the observed variation.

In the spleen, we found that monocytes expressing PD-L1 (p= 0.02), CD40 (p= 0.01), and CD70 (p= < 0.0001), as well as neutrophils expressing TIM-4 (p= 0.01), showed the greatest contribution to the separation between groups. In the lungs, macrophages expressing CD40 (p= 0.002) and CD64 (p= 0.04), along with neutrophils expressing CD80 (p= 0.005), were most distinct. In the liver, macrophages expressing CD40 (p= 0.001) and neutrophils expressing CD70 (p= 0.01) were the primary contributors (Fig. 3B).

Notably, the spleen displayed a more regulatory profile following IV immunization, characterized by decreased expression of activation markers CD40 and CD70, as well as the immune checkpoint marker PD-L1 (Fig. 3B). In contrast, in the lungs, IV immunization led to increased expression of CD64 on lung macrophages compared to ID immunization, suggesting enhanced activation and infiltration of monocyte-derived macrophages (Fig. 3B). In the liver, IV immunization led to upregulation of pro-

inflammatory markers CD70, CD40, and CD80, suggesting a shift toward a more inflammatory phenotype relative to ID immunization (Fig. 3B).

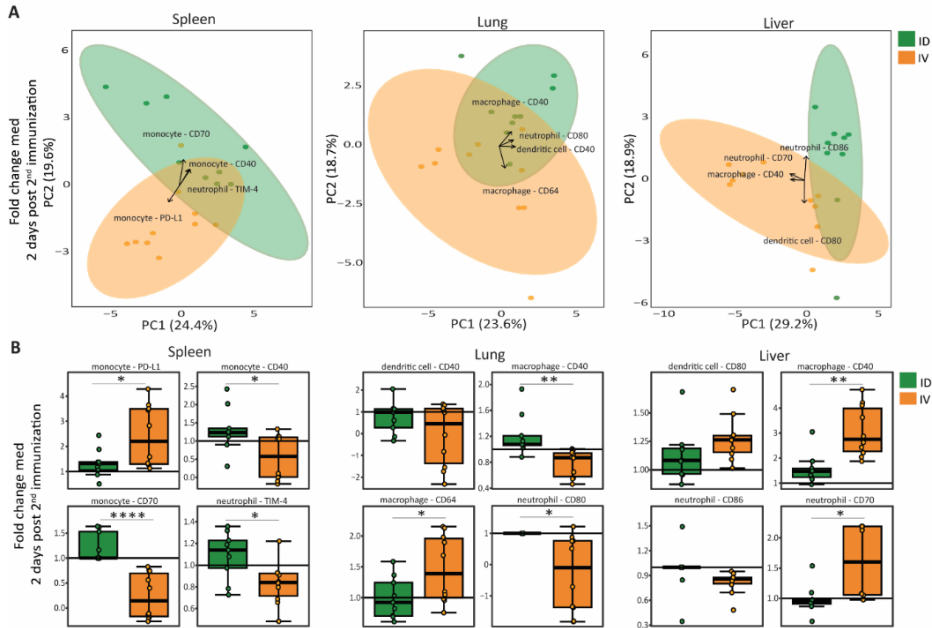


Figure 3. Activation of myeloid cells after intradermal (ID) and intravenous (IV) GA2 SPZ immunization. (A) Principal Component Analysis (PCA) plot of different markers on B-cells, neutrophils, macrophages, dendritic cells and monocytes after ID (green) of IV (orange) GA2 SPZ immunization in the spleen, lung and liver. MFI fold change relative to medium immunization, 2 days after the 2nd immunization. Markers and cell types responsible for the four highest loadings shown in the figure **(B)** Box plot of the markers and cell types responsible for the four highest loadings per organ between IV (orange) and ID (green) GA2 SPZ immunization. MFI fold change relative to medium, 2 days post 2nd immunization. ID immunization n=9 per group, IV immunization n=10 per group divided over 2 independent experiments. All conditions were *in vitro* restimulated with LA-GAP SPZ. Statistical significance between groups was assessed by an independent t-test. *p<0.05, **p<0.005, ***p<0.0005, and ****p<0.0001.

Distribution of immune cells in the skin and skin-draining lymph node

In an attempt to trace the source of the observed myeloid alteration, we examined the distribution of immune cells in the skin and skin-draining lymph node (skin dLN) after ID immunization. Two days after the second ID immunization, we observed an early increase in total number of immune cells in the skin after both SGE (p = 0.0006) and GA2 SPZ (p = 0.003) immunization, which normalized rapidly at 7 days after 2nd immunization (Fig. 4A). In the skin dLN we observed an increase in the total number of

immune cells at the early time point of 2 days after 2nd immunization with SGE ($p = 0.0002$) or GA2 SPZ ($p = <0.0001$), which remained stable at the later 7 days after 2nd immunization time point (Fig. 4B). This suggests that APC cells from the skin along with circulating B and T cells, migrate to the skin dLN within 2 days where antigen presentation and subsequent B and T cell activation occurs.

When we investigated the overall kinetics of immune cell distribution in the skin and skin dLN 2 days after 2nd ID immunization, we observed an increase in the percentage of eosinophils in SGE compared with medium ($p = 0.02$) and in SGE compared with GA2 SPZ ($p = 0.009$) immunization. At 7 days after 2nd immunization, we found an increase in the percentage of eosinophils after SGE ($p = 0.06$) and GA2 SPZ ($p = 0.04$) immunization. Additionally, we observed an increase in the percentage CD3⁺ T cells after GA2 SPZ ($p = 0.04$) immunization (Fig. 4C). Interestingly, in the skin dLN we observed a striking increase in B-cells after GA2 SPZ administration ($p = <0.0001$) and SGE ($p = 0.001$) 2 days after 2nd immunization, but a lower percentage CD3⁺ T cells after SGE ($p = <0.0001$) and GA2 SPZ ($p = <0.0001$) immunization (Fig. 4D). However, the total number of CD3⁺ T cells after SGE (mean 408.10^4) and GA2 SPZ (366.10^4) immunization was slightly higher than medium (232.10^4) immunization (Sup Fig. 5A). Overall we found in the skin and skin dLN an increase in total number of monocytes, dendritic cells, macrophages, CD3⁺ T cells and B-cells 2 days after 2nd GA2 SPZ immunization and SGE compared with medium (Sup Fig. 5A).

Next, we investigated the expression of activation and regulatory markers on myeloid cells in the skin and skin dLN 2 days after 2nd GA2 SPZ immunization compared with 2 days after 2nd medium immunization. Here we observed, next to an increase in activation marker CD64 and CD70 on monocytes and neutrophils, a striking decrease in expression of activation marker CD86 on dendritic cells, macrophages, monocytes and neutrophils (Fig. 4E+G). Compared to SGE immunization, GA2 SPZ immunization led to the upregulation of regulatory marker PD-L1 on dendritic cells ($p = 0.09$) and macrophages ($p = 0.06$) in the skin dLN (Fig. 4F+H). Last, in the skin, T cell activation appeared overall comparable between GA2 SPZ and SGE but in the skin dLN the activation of T cells after GA2 SPZ immunization seems slightly downregulated compared with SGE immunization for most activation markers (Sup Fig. 6A). In line with previous findings, we observed a decrease over time in the percentage of IFN γ ⁺ CD8⁺ T cells in the skin dLN (3.3%) [13], suggesting migration of these cells to other organs. This effect appeared to be specific to SPZ restimulation and was not observed following PMA/Iono restimulation (Sup Fig. 6B+C+D).

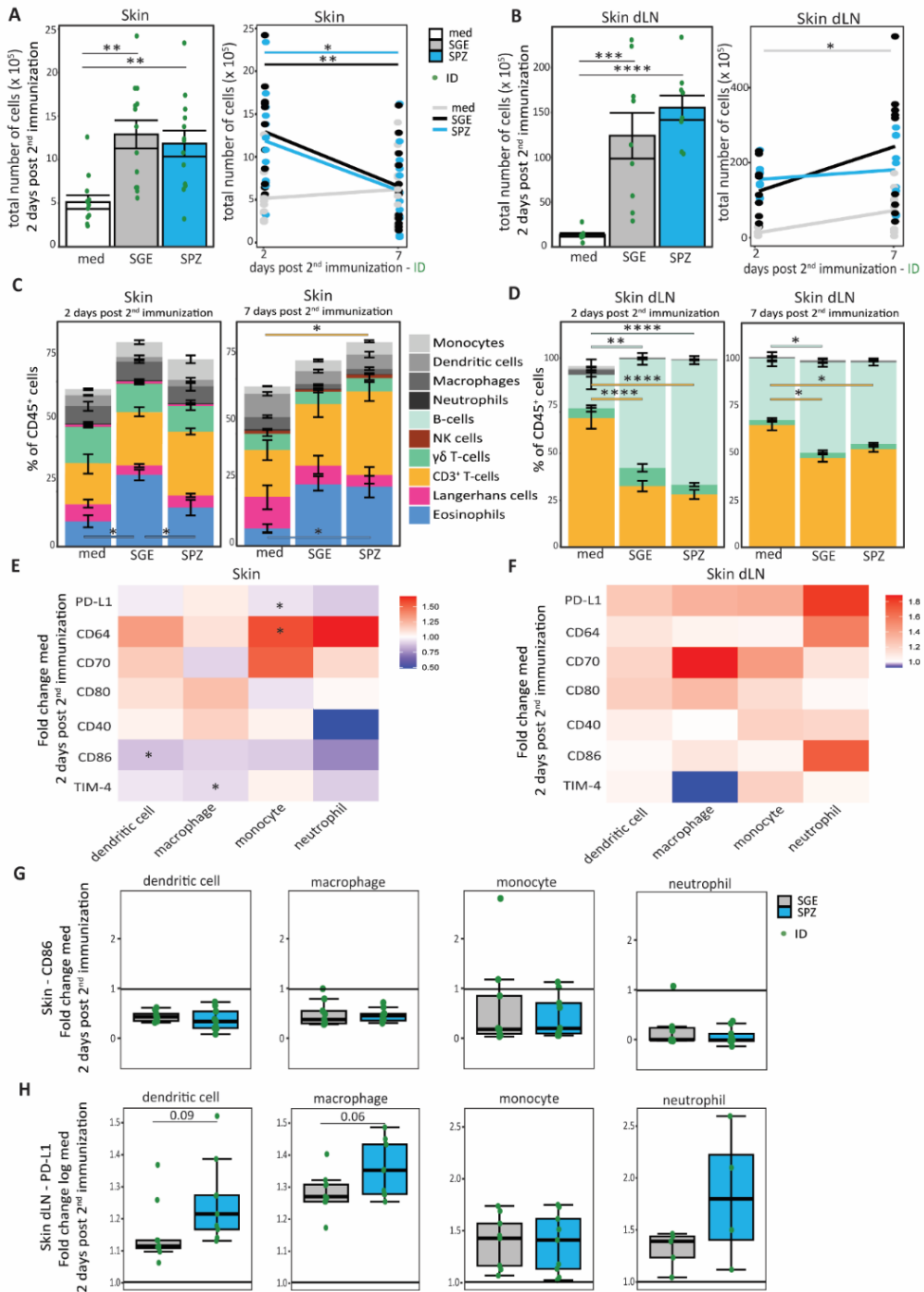


Figure 4. Cell distribution and myeloid activation in the skin and skin-draining lymph node (dLN). (A+B) The total number of immune cells in the skin (A) or skin dLN (B) *ex vivo*, after

medium (white), SGE (grey) or GA2 SPZ (blue) ID (green dots) immunization at 2 days or 7 days after 2nd immunization. **(C+D)** cell distribution in the skin **(C)** and skin dLN **(D)** after medium, SGE or GA2 SPZ immunization at 2 days or 7 days after 2nd immunization followed by *in vitro* restimulation with LA-GAP SPZ. **(E+F)** Heatmap of fold change relative to medium expression of different markers on myeloid cells for the skin **(E)** and skin dLN **(F)** 2 days after 2nd immunization followed by *in vitro* restimulation with LA-GAP SPZ. **(G)** Expression of CD86 in the skin on B-cells, dendritic cells, macrophages, monocytes and neutrophils after SGE (grey) or GA2 SPZ (blue) ID (green dots) immunization after *in vitro* restimulation with LA-GAP SPZ. **(H)** Expression of PD-L1 in the skin dLN on B-cells, dendritic cells, macrophages, monocytes and neutrophils after SGE (grey) or GA2 SPZ (blue) ID (green dots) immunization after *in vitro* restimulation with LA-GAP SPZ. Skin n=9-13 mice per group, skin dLN n=9 mice per group divided over 2 or 3 independent experiments. Statistical significance between groups was assessed by one-way ANOVA with multiple comparisons or an independent t-test. *p<0.05, **p<0.005, ***p<0.0005, and ****p<0.0001.

Overall, we observed an increase in the number of immune cells in both the skin and skin dLN, but an impressive decrease of activation marker CD86 on myeloid cells in the skin after GA2 SPZ immunization which also occurs after SGE immunization. However, GA2 SPZ immunization specifically led to the upregulation of regulatory marker PD-L1 on dendritic cells and macrophages in the skin dLN, suggesting the induction of a regulatory phenotype after ID GA2 SPZ immunization. This regulatory profile was not seen after SGE immunization and may impair the antigen-presenting capacity of dendritic cells and macrophages as they migrate from the skin dLN to other malaria-infected organs, potentially influencing the quality and magnitude of the systemic T-cell response (Fig. 5).

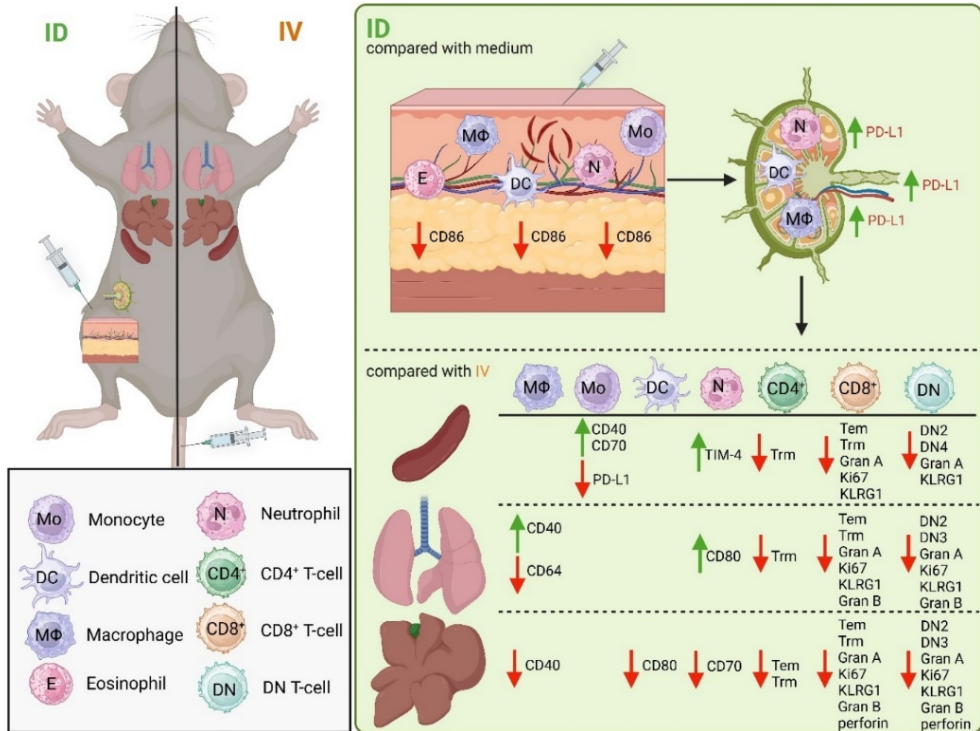


Figure 5. Summary. Immune responses after two immunizations with late arresting genetically attenuated parasite sporozoite (GA2 SPZ) in the skin, skin-draining lymph node, spleen, lungs and liver. Skin and skin draining lymph node GA2 SPZ intradermal immunization (ID, green) compared with medium ID immunization and spleen, lungs and liver GA2 SPZ ID compared with GA2 SPZ intravenous immunization (IV, orange).

Discussion

In summary, we compared the immune responses in the spleen, lung and liver after GA2 SPZ ID or IV immunization in a *Plasmodium berghei* (*Pb*) rodent malaria model to unravel the potential mechanisms underlying decreased protection of ID-immunized mice. Overall, our findings revealed a lower number of CD8⁺ Trm cells and reduced activation of CD8⁺ T cells and DN T cells after ID immunization compared with IV immunization, corroborating prior findings in the *Py* rodent malaria model [24]. In the skin, skin dLN and the liver we identified a large population of myeloid cells with a predominantly regulatory phenotype following ID GA2 SPZ immunization. These cells exhibited decreased CD86 expression in the skin and increased PD-L1 expression in the skin dLN. These results suggest that myeloid cells in the skin and skin dLN orchestrate a regulatory immune response resulting in a lower number of CD8⁺ Trm cells and

diminished T cell activation in the spleen, lungs and liver after ID GA2 SPZ immunization as compared to IV immunization.

Our findings in mouse models may provide insight into why ID-administered whole SPZ vaccines are less effective at inducing protective immunity in humans. First, it has been shown that SPZ injected intradermally by needle are 3-fold less efficient in migrating and infecting mouse livers compared to mosquito-inoculated SPZ. Previously we suggested that this is a consequence of the changes in the dermal tissue morphology caused by the injection of larger volumes via needles, in contrast to the small volume saliva inoculation by mosquitoes [25]. Second, recent findings demonstrate that some SPZ remaining in the skin after ID immunization migrate to the skin dLN, where they may contribute to the induction of regulatory responses, characterized by PD-L1 expression on myeloid cells. These myeloid cells may, in turn, orchestrate the suppression of T cell activation, specifically to a reduction of CD8⁺ Trm cells as well as decreased expression of Granzyme B and perforin. Previously, we have investigated the responses of human monocyte-derived macrophages when stimulated with SPZ and observed a similar regulatory phenotype. Similarly, we have shown that *in vitro* human dermal myeloid cells showed an increase in PD-L1 expression following exposure to SPZ [26]. Altogether these findings support our belief that the mechanisms observed in the *Pb* rodent model may be translatable to humans.

While investigating the myeloid activation in the skin we further observed a decrease in CD86 and TIM-4 expression, which suggests reduced T cell activation [27]. However, the increased expression of CD64 and CD70 points to a different, non-traditional pathway of T cell stimulation, possibly involving CD70 on myeloid cells and CD27 on T cells, which needs further study [28, 29]. In the skin dLN, we primarily observed increased PD-L1 expression, indicating a regulatory phenotype [30]. In the liver, we found decreased expression of activation markers CD40, CD70 and CD80 on myeloid cells after ID GA2 SPZ immunization compared with IV immunization. This suggests that regulatory immune activity in the skin dLN may modulate hepatic immune responses. Notably, dendritic cells, key players in T cell activation [31, 32], along with the expression of perforin on CD8⁺ and DN T cells, were only affected in the liver after ID GA2 SPZ immunization. Together, these findings indicate that myeloid cell activity and T cell priming in the skin and dLN can shape downstream immune responses in the liver.

By passing through the skin, SPZ undergoes a “filtering” process allowing only viable and motile SPZ to enter the bloodstream. This filtering effect does not occur during IV

administration, where SPZ bypass the skin and its filtering mechanism, allowing also less motile or less viable SPZ to enter the bloodstream [25, 33]. By administering IV instead of ID, the skin and skin dLN will be passed, allowing “unfiltered” SPZ to enter the lungs. Consequently, a larger number of less motile and viable SPZ may become trapped in the lungs following IV immunization which could lead to more localized immune activation in the lungs [20]. Interestingly, although SPZ from ID immunization still travel through the lungs, we did not observe the presence of CD8⁺ Trm cells in this organ. Furthermore, we saw increased Granzyme B expression in CD8⁺ T cells and DN T cells following IV immunization but not after ID immunization, indicating lower activation in the lungs after ID immunization compared with IV immunization.

ID immunization is considered technically easier to perform than IV immunization [34]. Currently, several vaccines, including those for tuberculosis, influenza, and rabies, are administered through ID injection. These vaccines have been shown to be more effective when delivered at reduced doses [35-39]. For humans, the recommended volume for ID immunization typically ranges between 50 and 100 μ L (NIH). In mice, the standard volume ranges from 10 to 50 μ L (IACUC guideline), but recent studies suggest that whole SPZ ultra-low volumes as small as of 2.5 μ L are more effective as they facilitate migration of SPZ out of the skin injection site [40, 41]. Reducing the volume of the vaccine administered may thus enhance the efficacy of ID GA2 SPZ vaccinations, changing the ratio of trapped versus migratory SPZ. This may result in a lower regulatory immune response and improve T cell activation. Additionally, if more SPZ exit the skin, a larger number will likely infect hepatocytes and develop into late liver schizonts, which is essential for generating effective protection after GA2 immunization [5]. Consequently, a lower dose for GA2 SPZ ID administration could lead to a more effective, easier-to-administer, and cost-effective vaccine.

In conclusion, we compared the immune response in the spleen, lungs and liver after GA2 SPZ ID and IV immunization and our findings revealed decreased T cell differentiation and activation after ID immunization. Additionally, upon examining the skin and skin dLN we observed a regulatory myeloid phenotype. These observations suggest that the regulatory myeloid cells in the skin and skin dLN influence the T cell differentiation and activation in the spleen, lungs and liver. These findings warrant further investigation to examine their impact on vaccine efficacy and to assess whether protection following GA2 SPZ ID immunization can be strengthened.

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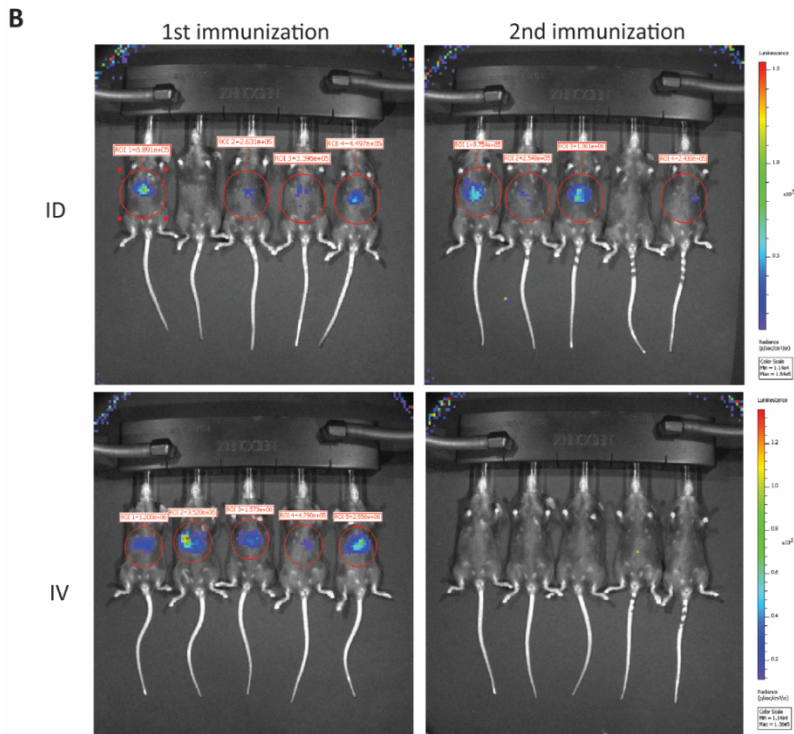
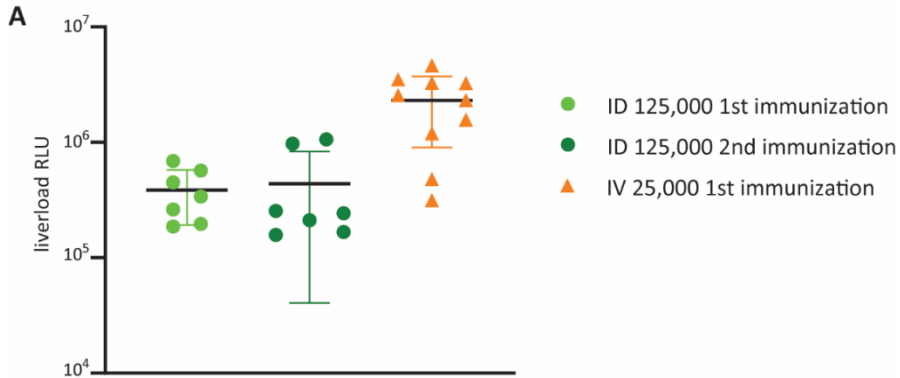
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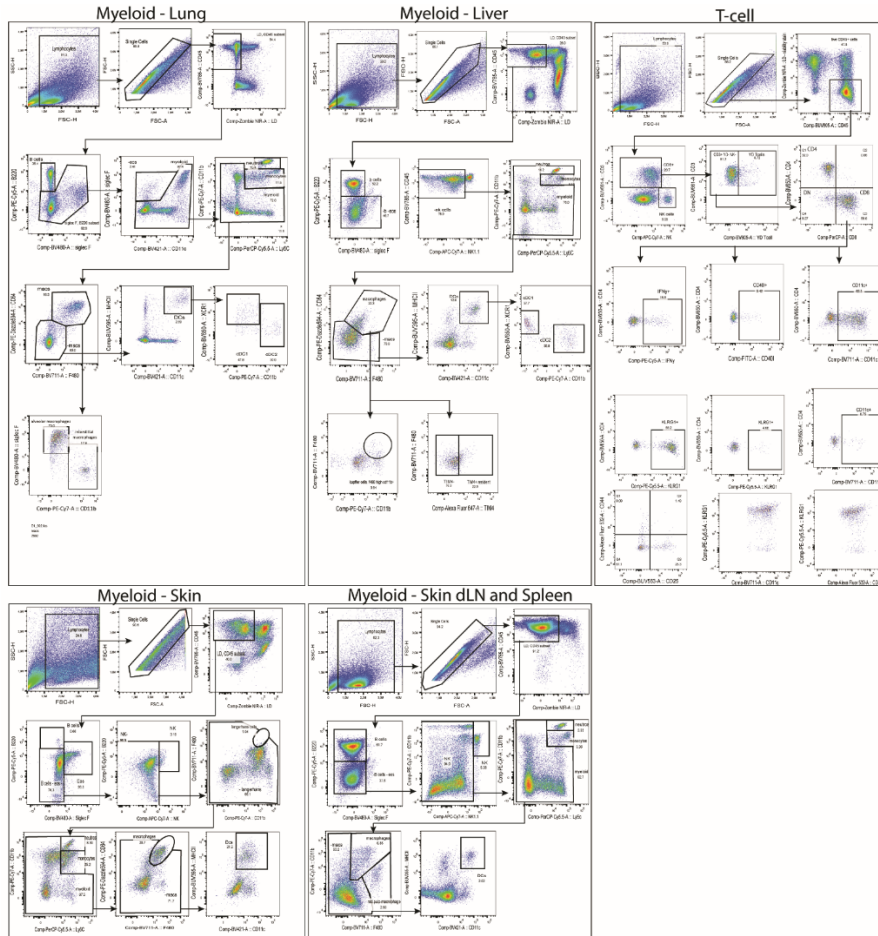
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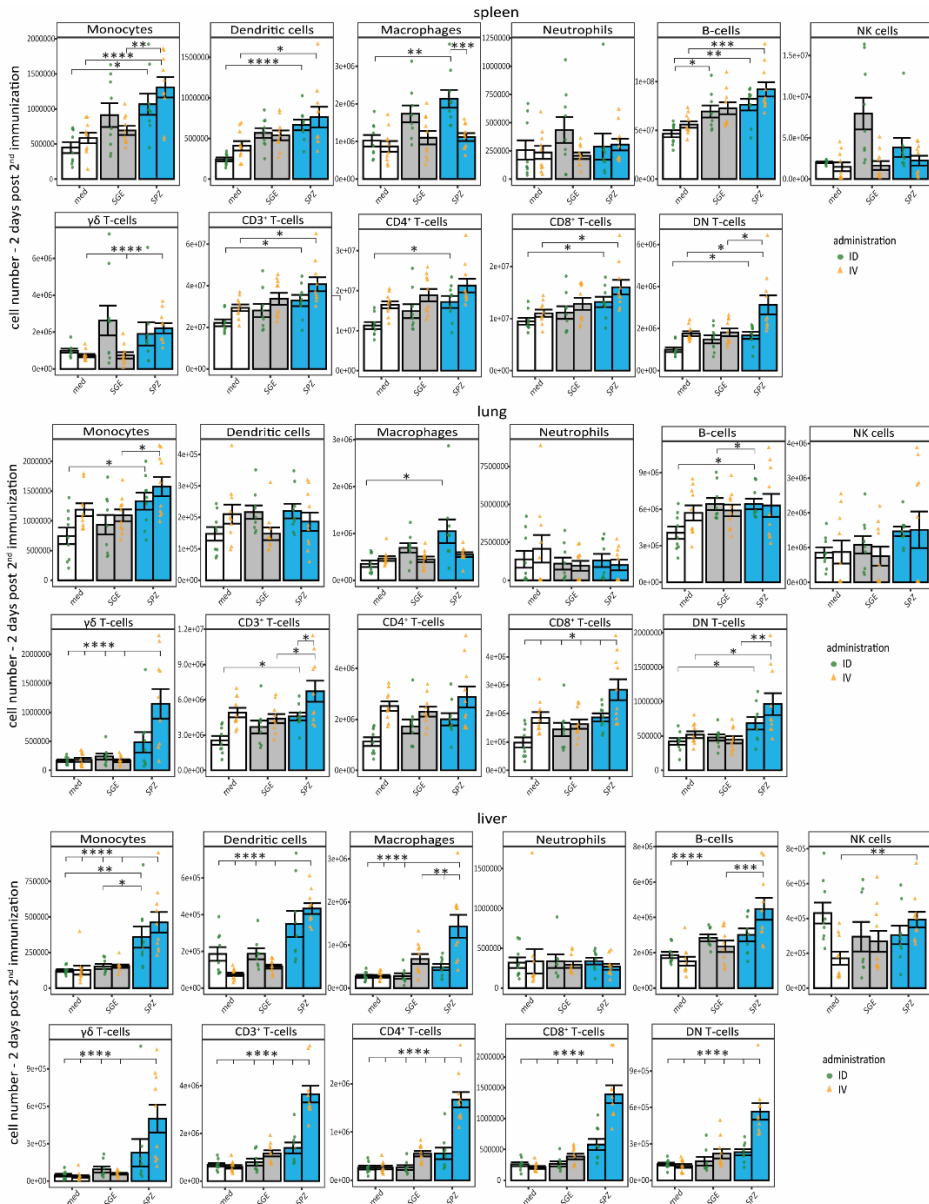
Supplementary



Supplementary figure 1. (A) Parasitic liver load 44 hours after ID (green) or IV (orange) administration measured by RLU. **(B)** Images of IVIS measurements of mice injected for the first or second time with GA2 SPZ via ID or IV immunization.



Supplementary figure 2. Gating strategy of myeloid cells for the lung, liver, skin, skin dLN and spleen. For the T cell gating same strategy has been used for all the organs.

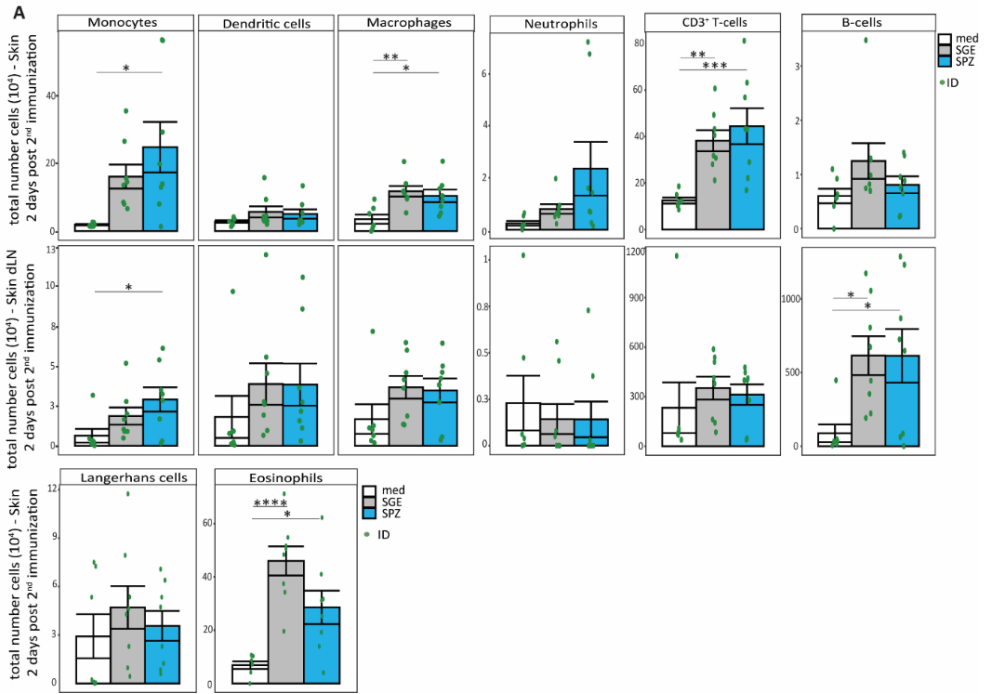


Supplementary figure 3. The absolute cell number of various immune cells in the spleen, lung and liver after medium (white), salivary gland extract (SGE) (grey) or GA2 sporozoite (SPZ) (blue) ID (green) or IV (orange) immunization. All conditions were *in vitro* restimulated with LA-GAP SPZ. ID immunization n=9 per group, IV immunization n=10 per group divided over 2 independent experiments. Statistical significance between groups was assessed by one-way ANOVA with multiple comparisons with Bonferroni correction. *p<0.05, **p<0.005, ***p<0.0005, and ****p<0.0001.

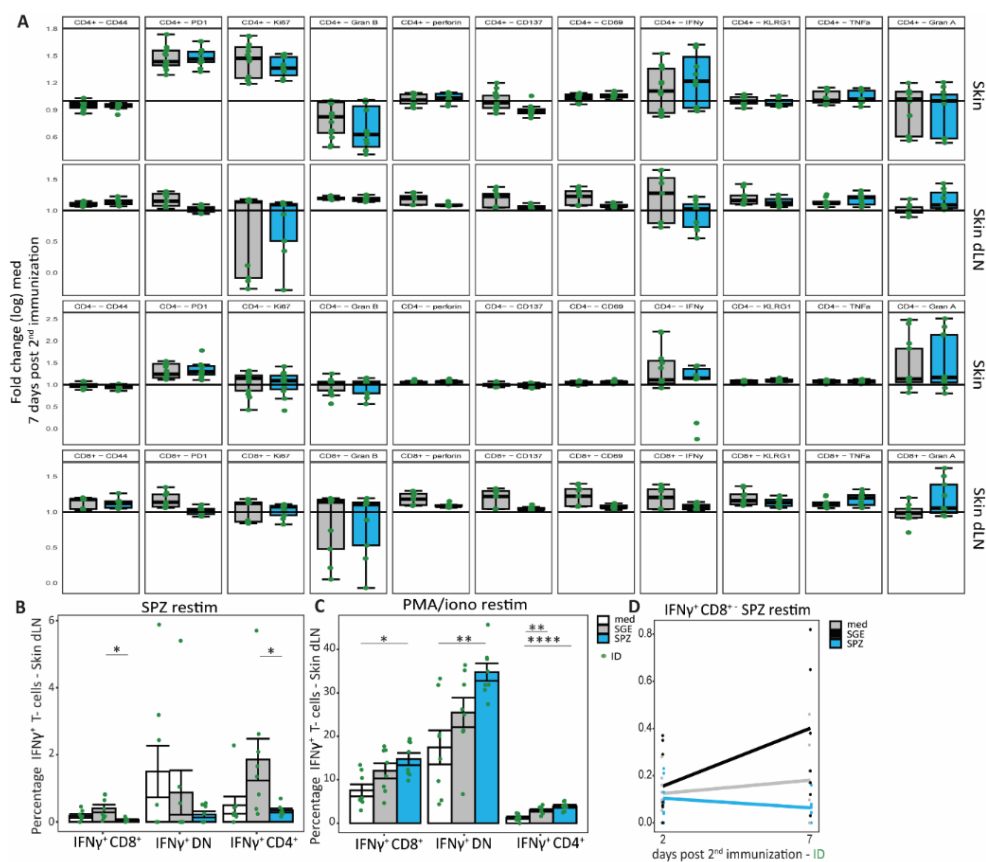
Chapter 6



Supplementary figure 4. (A) Differences between expression of different markers on CD4⁺ T cells in the spleen, lung and liver after ID (green) or IV (orange) GA2 SPZ immunization. Measured by fold change relative to medium. **(B)** Differences between expression of different markers on $\gamma\delta$ T cells in the spleen, lung and liver after ID (green) or IV (orange) GA2 SPZ immunization. Measured by fold change log scale relative to medium. All conditions were *in vitro* restimulated with LA-GAP SPZ. ID immunization n=9 per group, IV immunization n=10 per group divided over 2 independent experiments. Statistical significance between groups was assessed by one-way ANOVA with multiple comparisons with Bonferroni correction. *p<0.05, **p<0.005, ***p<0.0005, and ****p<0.0001.



Supplementary figure 5. (A) The total number of monocytes, dendritic cells, macrophages, neutrophils CD3⁺ T cells and B cells in the skin and skin dLN, and Langerhans cells and eosinophils in the skin. Ex vivo counted 2 days after 2nd ID immunization with medium (white), SGE (grey) or GA2 SPZ (blue). Skin n=9-13 mice per group, skin dLN n=9 mice per group divided over 2 or 3 independent experiments. Statistical significance between groups was assessed by one-way ANOVA with multiple comparisons or an independent t-test. *p<0.05, **p<0.005, ***p<0.0005, and ****p<0.0001.



Supplementary figure 6. (A) Activation of CD4⁺ and CD4⁻ T cells in the skin and CD4⁺ and CD8⁺ T cells in the skin dLN. 7 days post 2nd immunization of SGE (grey) or GA2 SPZ (blue), measured by fold change relative to medium log scale. **(B)** Percentage of IFN γ ⁺ CD8⁺, CD4⁺ or DN T cells after SPZ **(B)** or PMA/Iono **(C)** restimulation 7 days after 2nd ID immunization. **D.** The amount of IFN γ ⁺ CD8⁺ T cells over time measured by 2 days post 2nd GA2 SPZ immunization and 7 days post 2nd GA2 SPZ immunization. medium (grey), SGE (black) or GA2 SPZ (blue). All conditions were *in vitro* restimulated with LA-GAP SPZ. Skin n=9-13 mice per group, skin dLN n=9 mice per group divided over 2 or 3 independent experiments. Statistical significance between groups was assessed by one-way ANOVA with multiple comparisons or an independent t-test. *p<0.05, **p<0.005, ***p<0.0005, and ****p<0.0001.

