



Depletion of Regulatory T Cells Augments a Vaccine-Induced T Effector Cell Response against the Liver-Stage of Malaria but Fails to Increase Memory

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Abstract

Regulatory T cells (T_{reg}) have been shown to restrict vaccine-induced T cell responses in different experimental models. In these studies $CD4^+CD25^+ T_{reg}$ were depleted using monoclonal antibodies against CD25, which might also interfere with CD25 on non-regulatory T cell populations and would have no effect on $Foxp3^+CD25^- T_{reg}$. To obtain more insights in the specific function of T_{reg} during vaccination we used mice that are transgenic for a bacterial artificial chromosome expressing a diphtheria toxin (DT) receptor-eGFP fusion protein under the control of the *foxp3* gene locus (depletion of regulatory T cell mice; Dereg). As an experimental vaccine-carrier recombinant *Bordetella* adenylate cyclase toxoid fused with a MHC-class I-restricted epitope of the circumsporozoite protein (ACT-CSP) of *Plasmodium berghei* (*Pb*) was used. ACT-CSP was shown by us previously to introduce the $CD8^+$ epitope of *Pb*-CSP into the MHC class I presentation pathway of professional antigen-presenting cells (APC). Using this system we demonstrate here that the number of CSP-specific T cells increases when T_{reg} are depleted during prime but also during boost immunization. Importantly, despite this increase of T effector cells no difference in the number of antigen-specific memory cells was observed.

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Introduction

The liver stage of malaria is an attractive target for vaccination. During the liver stage sporozoites invade hepatocytes where they replicate without provoking clinical illness. After the infected hepatocytes burst, merozoites are released into the blood stream and infect red blood cells, where they undergo a massive replication leading to the symptoms of malaria. Infected hepatocytes present plasmodial antigens on MHC class I molecules where they can be recognized by antigen-specific $CD8^+$ T cells [1,2]. A high level of protection was induced using irradiated sporozoites [3] IFN γ production by these cells is implicated in protection [4,5]. More recently genetically modified sporozoites were used as vaccines which induce antigen-specific T cells but are not capable of establishing the infectious cycle [6,7]. *P. berghei* is a worldwide employed infection model for malaria in mice. The efficacy of experimental vaccines can be tested employing infection of BALB/c mice with *P. berghei* sporozoites [8]. Here a peptide of the circumsporozoite protein (CSP) was identified that is presented on H-2K^d. Thus in this model the number and function of antigen-specific T cells [9] can be monitored. Up to now several different methods were used to induce CSP-specific T cells [10]. Some of these strategies indeed induce promising numbers of CSP-specific $CD8^+$ T cells but the degree of protection often varies. Up to now

the most promising strategies rely on heterologous prime/boost immunization [11].

The adenylate cyclase toxoid (ACT) of *Bordetella pertussis* is capable of delivering its catalytic domain and inserted cargo $CD8^+$ T cell epitopes into the cytosol of CD11b-expressing professional antigen-presenting cells. Thus recombinant and detoxified ACT containing different epitopes was repeatedly used for delivery into the MHC class I presentation pathway to generate $CD8^+$ T cells against model antigens [12], which demonstrates the versatility of this tool as antigen-delivery system. We used recombinant detoxified ACT containing an epitope of *Plasmodium berghei* CSP (ACT-CSP) in other studies to induce high numbers of IFN γ secreting $CD8^+$ T cells, which confer sterile immunity against sporozoite challenge when combined with a blockade of CTLA-4 or using a heterologous prime/boost approach with CSP-expressing *Salmonella* [13].

Regulatory T cells (T_{reg}) were shown to inhibit immune responses and thereby play a central role in peripheral tolerance. T_{reg} also attenuate the immune response during infections and vaccination. In naïve mice $CD4^+Foxp3^+$ T cells express CD25. Thus antibodies against CD25 have been used to deplete $CD4^+CD25^+$ T_{reg} cells [14,15]. Recent studies have shown that a depletion of $CD4^+CD25^+$ T_{reg} by antibodies led to an increased number of vaccine-induced T cells [16]. Understanding the exact role of T_{reg} during vaccination is hampered due to the multiple

effects anti-CD25 might have also on non-regulatory T cells that express CD25. Until recently no tools were available to selectively deplete T_{reg} . Thus we employed transgenic mice in which $Foxp3^{+} T_{reg}$ can be specifically depleted to study the specific function of T_{reg} during vaccination against the liver-stage of *P. berghei* [17]. These mice are transgenic for a bacterial artificial chromosome expressing a diphtheria toxin (DT) receptor-eGFP fusion protein under the control of the *foxp3* gene locus. This allows a selective depletion of $Foxp3^{+} T_{reg}$ by DT injection.

Collectively our data demonstrate that $CD4^{+}Foxp3^{+} T_{reg}$ dampen the magnitude of a CSP-specific $CD8^{+}$ T cell response that was induced by vaccination during priming and/or boosting. In addition, we demonstrated that an enhanced number of T effector cells are not necessarily accompanied by an increased number of memory T cells.

Materials and Methods

Mice and parasites

Male DEREg-mice [17] were crossed with female Balb/c mice in the animal facility of the Bernhard Nocht Institute for Tropical Medicine. The resulting offsprings were typed by PCR and flow cytometry. In each experiment hemizygotic DEREg mice were used with sex and age-matched wildtype littermates as controls. For some experiments we used the F1 generation of male DEREg mice on BALB/c background and female C57BL/6 wild type mice (H-2Kb). The resulting F1 generation expresses H-2Kd and H-2Kb, which allows the simultaneous analysis of T cell responses against the malaria CSP-epitope (SYIPSAEKI corresponding to CSP 245–253) and the SIINFEKL epitope of ovalbumin. Animal maintenance and experimental procedures were all approved by the the relevant regulatory agency in Germany, the “Amt fuer Gesundheit und Verbraucherschutz of the Hansestadt Hamburg”. The procedures followed were in accordance with the ethical standards of the review board of the Bernhard-Nocht-Institute for Tropical Medicine and the Ethic Commission of the state of Hamburg as well as the related recommendations of the EU.

For all experiments 6–8 weeks old mice were used. *Plasmodium berghei* ANKA was maintained by alternating cyclic passage of the parasite in *Anopheles stephensi* mosquitoes and BALB/c mice at the mosquito colony of the Bernhard Nocht Institute for Tropical Medicine. Sporozoites were collected by manual dissection of infected mosquito salivary glands in minimal essential medium (MEM) 18–21 days after the mosquito had taken an infectious blood meal.

Depletion of Treg cells

For depletion of T_{reg} cells, DEREg and control mice were injected i.p. with 1 μ g diphtheria toxin (Merck) diluted in endotoxin-free PBS for three consecutive days, starting on day 1 after prime or boost immunization.

ACT-CSP toxoid construction and purification

The construction of ACT-CSP was described in a previous study [9]. The amino acid sequence VRVRKNNDDSYIPSAEKILEFVKQ, which comprises the MHC I epitope SYIPSAEKI corresponding to CSP 245–253, was inserted at position 336 into the catalytic domain of the adenylate cyclase of *B. pertussis*. The resulting ACT-CSP also includes the OVA-derived SIINFEKL epitope, presented in the context of H-2K^b (corresponding to residues 257 to 264 from OVA). The detoxified ACT-CSP was produced in the *E. coli* strain XL1-Blue (Stratagene) transformed with the appropriate plasmid construct.

Immunization and challenge

Mice were immunized i.p. with a single dose of 20 μ g ACT-CSP diluted in 200 μ l of phosphate buffered saline (PBS) on day 0. Boost immunization was performed 14 days after prime immunization. Challenge was performed i.v. seven days after prime or boost immunization using 1000 *Pb* sporozoites. For experiments concerning induction of memory responses the challenge was performed at later time points as indicated. Mice were examined every day and parasitemia was determined every two days by light microscopy of blood smears with Wright's stain (Sigma, Taufkirchen, Germany).

Quantification of liver-stage burden

Quantification of liver-stage parasite burden was performed as described previously (Mol. Biochem. Parasitol. 2001, 118, p233–245). Briefly, at 30 h post-challenge, livers were perfused with PBS and removed. Total RNA was extracted with Trizol (Invitrogen, Darmstadt, Germany) according to the manufacturers instructions. RNA was transcribed using random hexamer primer and RevertAid H minus reverse transcriptase (Thermo Scientific, St. Leon-Rot, Germany) according to the manufacturers instructions. The resulting cDNA was amplified using the following primers: 5'-GGATGTATTTCGCTTTATTTAATGCTT-3' and 5'-CACGCGTGCAGCCTAGTAT-3' for the detection of 18S rRNA of *PbA*. As reference gene mouse GAPDH was amplified with the primers 5'-GGGTGTGAACCACGAGAAAT-3' and 5'-CCTTCCACAATGCCAAAGTT-3'. Cycling conditions were as following: 15 min 95°C, 40 cycles at 95°C for 15 s, 50°C for 20 s and 68°C for 20 s. For each cycle a melting curve analysis was performed with a ramp from 67° to 95°C. The relative concentration of *P. berghei* 18S rRNA was determined with the comparative C_t method (delta delta C_t).

Isolation of splenocytes and PBL

Spleens were removed at the indicated time points and RBCs were lysed by addition of ammonium chloride. Blood was collected from the tail vein and RBCs were lysed by addition of ammonium chloride.

IFN γ -ELISPOT

Single cell suspensions (2×10^5 /well) were cultivated in Millipore HTS HA plates coated with anti-mouse IFN γ . Cells were stimulated with 1 μ g/ml CSP_{245–253} peptide and in some experiments with 0.1 μ g OVA 257–264. Supernatants were removed after 18 hours and the number of IFN γ -producing cells was determined using ELISPOT, as described elsewhere [18]. Antibodies and Avidin-HRP were purchased from BD Pharmingen (Heidelberg, Germany). The synthetic peptides SYIPSAEKI and SIINFEKL, corresponding to CD8⁺ T-cell epitopes CSP 245–253 and OVA 257–264 respectively, were obtained from MWG (Ebersberg, Germany).

Analysis of cytokine production and proliferation

Single-cell suspensions (10^5 cells/well) were cultivated in 96-well plates. Cells were stimulated with 0.01 μ g/ml SIINFEKL, 0.1 μ g/ml CSP, 3 μ g/ml anti-CD3 or medium RPMI as control. After 48 h, supernatants were removed and the IFN γ concentration was determined by indirect sandwich enzyme linked immunosorbent assay (ELISA). Antibodies and cytokine standard were purchased from R&D Systems (Wiesbaden, Germany). Proliferation was analyzed after 24 h by incorporation of [³H] thymidine. Therefore, cells were pulsed with 0.5 mCi of [³H] thymidine for 6 h.

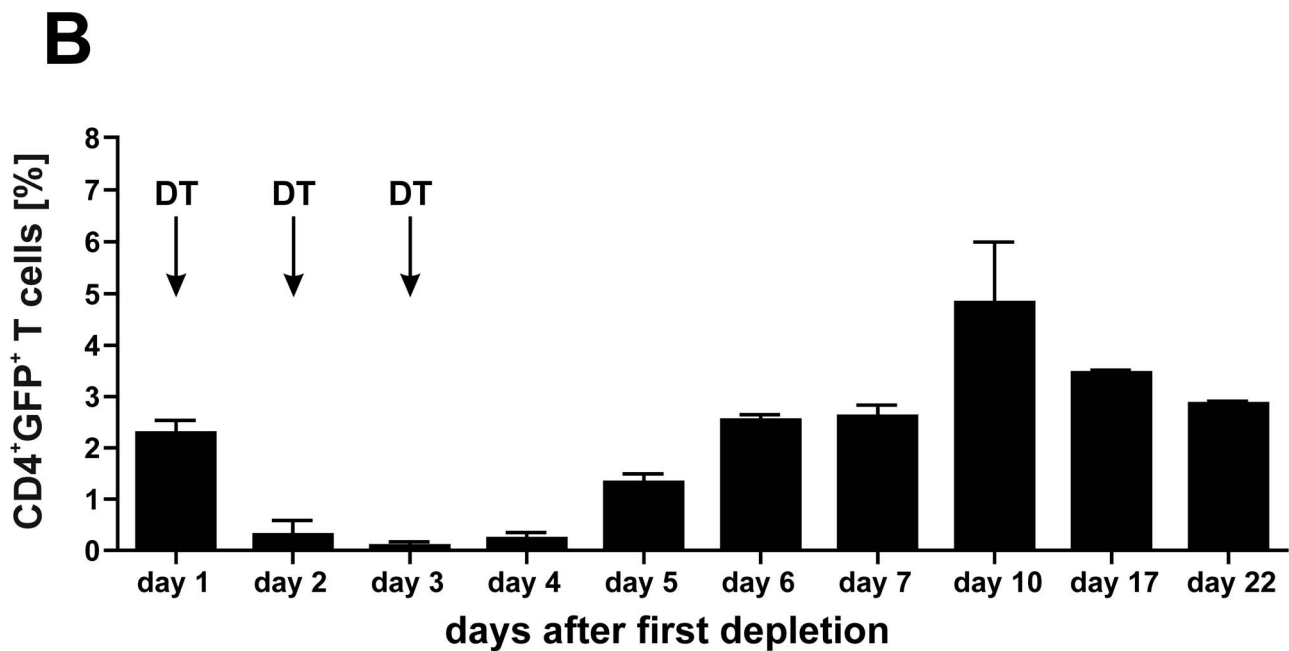
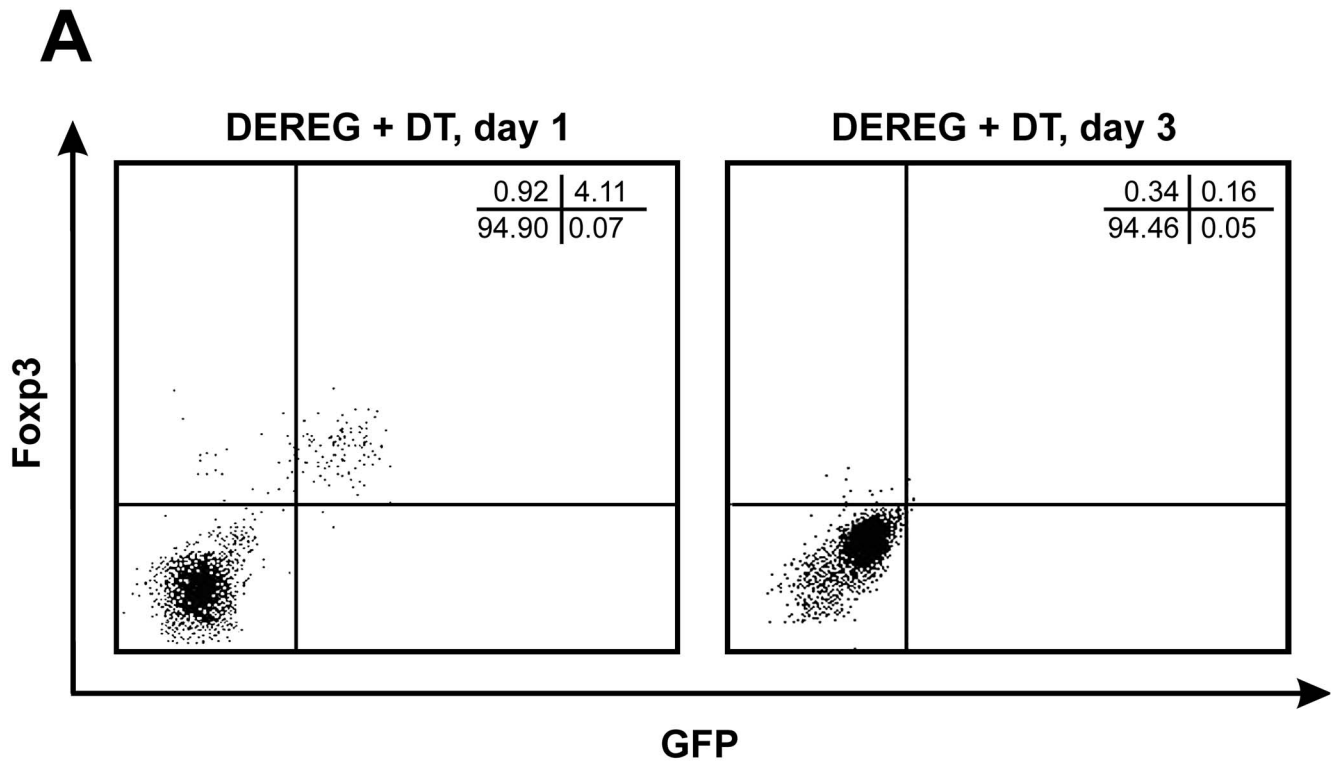


Figure 1. Kinetics of T_{reg} depletion in DEREG mice. In DEREG mice on BALB/c background Foxp3⁺ T_{reg} can be identified by their eGFP expression using flow cytometry. The consecutive application of DT i.p. over three days leads to a depletion of eGFP⁺Foxp3⁺ T_{reg} within 72 h (A). For immunization experiments we used an optimized protocol where mice were treated with DT on days +1, +2 and +3. The percentage of CD4⁺GFP⁺ T_{reg} in the blood was measured in two F1 DEREG C57BL/6 x BALB/c mice by flow cytometry at the indicated time points (B). Data is expressed as mean +/– SEM in relation to all cells in the lymphocyte gate as defined by the respective forward- and sideward scatter. doi:10.1371/journal.pone.0104627.g001

Antibodies and flow cytometry

The following antibodies were purchased from BD Pharmingen (Heidelberg, Germany): APC-, FITC- or PE-labeled rat anti-mouse IFN γ , IL-10, CD4, CD8a, CD25, CD62L, and FITC-labeled hamster anti-mouse CD69. APC-labeled rat anti-mouse

TNF α , PE-labeled rat anti-mouse IL-2, PE- and FITC-labeled anti-mouse Foxp3, as well as the appropriate isotype controls were from eBioscience (San Diego, CA). The frequency of CSP-specific T cells was determined using an APC-labeled H2-Kd pentamer loaded with the CSP 245–253 peptide (Proimmune, Oxford, UK).

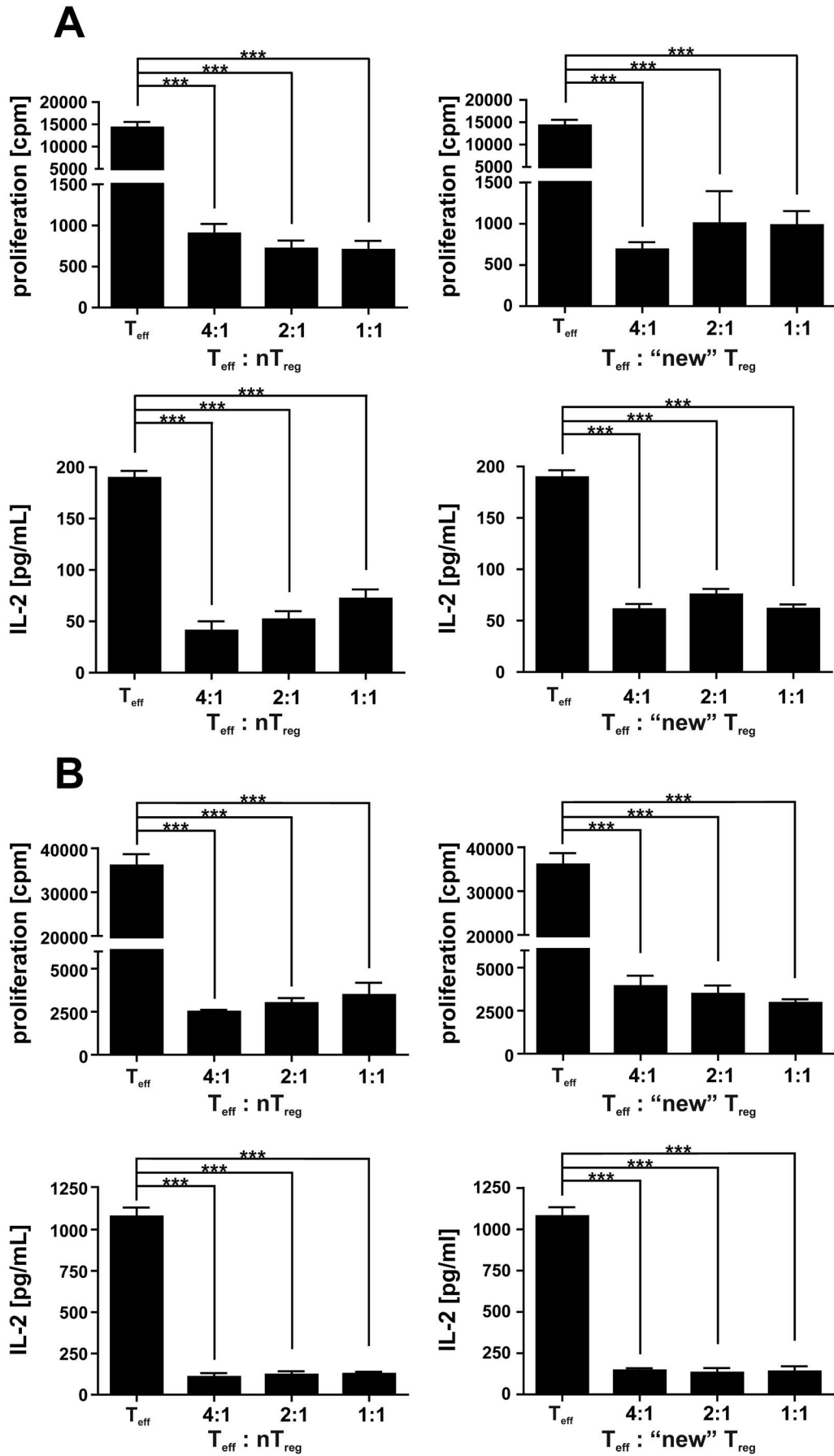


Figure 2. T_{reg} that repopulated depleted DEREg-mice are functional suppressors. (A) A F1 DEREg C57BL/6 x BALB/c-mouse were depleted by three consecutive DT injections and 30 days after depletion the T_{reg} that repopulated the animals were sorted by flow cytometry due to their eGFP-expression ("new" T_{reg}). Similarly GFP⁺ T_{reg} were sorted from an untreated F1 DEREg mice. Purity of both T_{reg} preparation was >95%. Naïve spleen cells were stimulated with anti-CD3 and incubated with different amounts of the initial GFP⁺ T_{reg} (natural GFP⁺ T_{reg} , n T_{reg}) before depletion and the GFP⁺ "new" T_{reg} that reappear after depletion. After 48 h the supernatants were harvested and IL-2 production was measured by ELISA. The proliferation of cells was measured by ³H-Thymidin incorporation. Data are expressed as mean \pm SD. The experiment was repeated twice with similar results. (B) A similar assay was performed using either the initial T_{reg} purified from F1 DEREg C57BL/6 x BALB/c mice or those "new" T_{reg} that reappear in these mice after depletion. Initial T_{reg} or "new" T_{reg} were incubated with different amounts of spleen cells from OT-1 T cell receptor transgenic mice, which were subsequently stimulated with SIINFEKL-peptide. After 48 h IL-2 production and proliferation of cells was measured. Data are expressed as mean \pm SD. The experiment was repeated once with similar results. *** p <0.0001. doi:10.1371/journal.pone.0104627.g002

Intracellular Foxp3 staining was performed with the Foxp3 staining set (eBioscience, San Diego CA), according to the manufacturer's instructions. Dead cells were excluded by propidium iodide staining. Data were acquired on a FACSCalibur flow cytometer (Becton Dickinson, Mountain View, CA) and analyzed with the CellQuest program (Becton Dickinson). Facs plots are minimally adjusted in brightness and contrast using Adobe Photoshop CS6 (Adobe Systems, San Jose, CA).

Statistical analysis

Statistical analysis was done with GraphPad PRISM (GraphPad Software Inc., San Diego, USA). The Student's t-test was used for measuring the significance of difference between two distributions. statistical significance. Multiple groups were compared using a one-way or two-way analysis of variance (ANOVA). The Mann-Whitney test and Bonferroni Multiple Comparison test were used as a post hoc comparison of the ANOVA. For survival analysis we employed Kaplan-Meier estimator function for lifetime data. If not otherwise stated, data are expressed as means \pm SD. Statistical significance was indicated as followed: **** p <0.0001; *** p <0.001; ** p <0.01; * p <0.05.

Results

Kinetics of T_{reg} depletion in DEREg mice

GFP⁺Foxp3⁺ T_{reg} can be rapidly depleted in DEREg mice by i.p. injection of DT. The percentage of CD4⁺Foxp3⁺ T_{reg} cells dropped from approximately 3% to under 1.5% in blood and spleen when analyzed 12 h after a single i.p. application of DT (data not shown). We tested several protocols for the depletion of T_{reg} . Most efficient was a treatment on three consecutive days [18]. Using this protocol the T_{reg} numbers were significantly reduced for almost 5 days. Thus we decided to define a protocol where we inject DT on days +1, +2, +3 (Fig. 1 A, Fig. S1). Using this schedule T_{reg} were strongly reduced from day +1 to +5 when the strongest expansion of vaccine-induced T cells could be expected (Fig. 1 B). Modified schedules i.e. starting from -1 and/or prolonged depletion do not lead to statistical significant enhanced T cell responses (data not shown). The number of eGFP⁺/Foxp3⁺ T cells started to increase from days +5 to +6 (Fig. 1 B). On later time points depleted mice have similar numbers of eGFP⁺/Foxp3⁺ T_{reg} compared to DT treated non transgenic littermates control mice.

Functional analysis of reappearing T_{reg}

After transient depletion of T_{reg} by DT injection GFP⁺Foxp3⁺ T_{reg} cells repopulate depleted mice rapidly (Fig. 1B). Currently it is not known if this is due to proliferation of remaining T_{reg} , *de novo* repopulation by thymic derived T_{reg} or due to conversion of conventional T cells. However, we were interested if T_{reg} that repopulate depleted DEREg mice ("new" T_{reg}) have the same regulatory capacity as the initial T_{reg} population. To this end BALB/c DEREg mice were depleted by three consecutive DT

injections and 30 days after depletion the T_{reg} that repopulated the animals were sorted by flow cytometry due to their eGFP-expression. Similarly GFP⁺ T_{reg} were sorted from untreated DEREg mice. Spleen cells of untreated mice were stimulated with anti-CD3 and incubated with different amounts of the initial GFP⁺ T_{reg} before depletion and the "new" T_{reg} that reappear after depletion. Both types of T_{reg} exhibited similar suppressive capacity and inhibited IL-2 production and proliferation (Fig. 2 A). Similar assays were performed using either the initial T_{reg} purified from F1 DEREg C57BL/6 x BALB/c mice or those that reappear in these mice after depletion. Initial T_{reg} or "new" T_{reg} were incubated with different amounts of spleen cells from OT-1 T cell receptor transgenic mice, which were subsequently stimulated with SIINFEKL-peptide. Again, both types of T_{reg} exhibit a strong suppressive capacity (Fig. 2 B). Thus T_{reg} that repopulate mice after transient depletion were functional.

Effects of T_{reg} depletion during prime immunization

To analyze the effect of T_{reg} depletion on the efficacy of immunization, we used an adenylate cyclase toxoid (ACT-CSP) that carries an insert of the *P. berghei* CSP-derived CD8⁺ T cell epitope SYIPSAEKI presented on H-2K^d (corresponding to residues 245 to 253 from CSP) and in addition the OVA-derived SIINFEKL epitope, presented in the context of H-2K^b (corresponding to residues 257 to 264 from OVA) [9,13]. For initial immunization experiments we used the F1 generation from male DEREg-mice on C57BL/6 background and female wildtype BALB/c mice (F1 DEREg). Cells from these mice express the MHC class I molecule H-2K^b and H-2K^d. This allows the simultaneous characterization of T cell responses against *P. berghei*-CSP (H-2K^d) and the SIINFEKL (H-2K^b) peptide used as model antigen in various studies. We have previously shown that a single application of ACT-CSP can induce IFN γ producing CSP-specific CD8⁺ T cells. However, the number of CSP-specific CD8⁺ T cells remained under a threshold needed for protection unless it was combined with a blockade of CTLA-4 [9]. Accordingly, when F1 wildtype mice were immunized i.p. with ACT-CSP and treated with DT, an induction of CSP- and SIINFEKL-specific T cells in spleens was found to produce IFN γ upon restimulation with the respective peptide. However, when F1 DEREg littermates were immunized and treated with DT a strong increase in IFN γ production was observed (Fig. 3 A). To further evaluate if the number of CSP-specific CD8⁺ T cells also increased, we specifically stained these cells using the MHC-class I multimer technology and analyzed the CSP-specific T cells subsequently by flow cytometry. Here we also found that upon depletion of T_{reg} an increased number of CSP-specific T cells after immunization with ACT-CSP was induced (Fig. 3 B). Not immunized control mice shown no increased specific response upon restimulation. (Fig. 3 C and D). These data show that a transient depletion of T_{reg} augments the priming of antigen-specific CD8⁺ T cells.

The observation that a single immunization with ACT-CSP in combination with the depletion of T_{reg} already induces a high

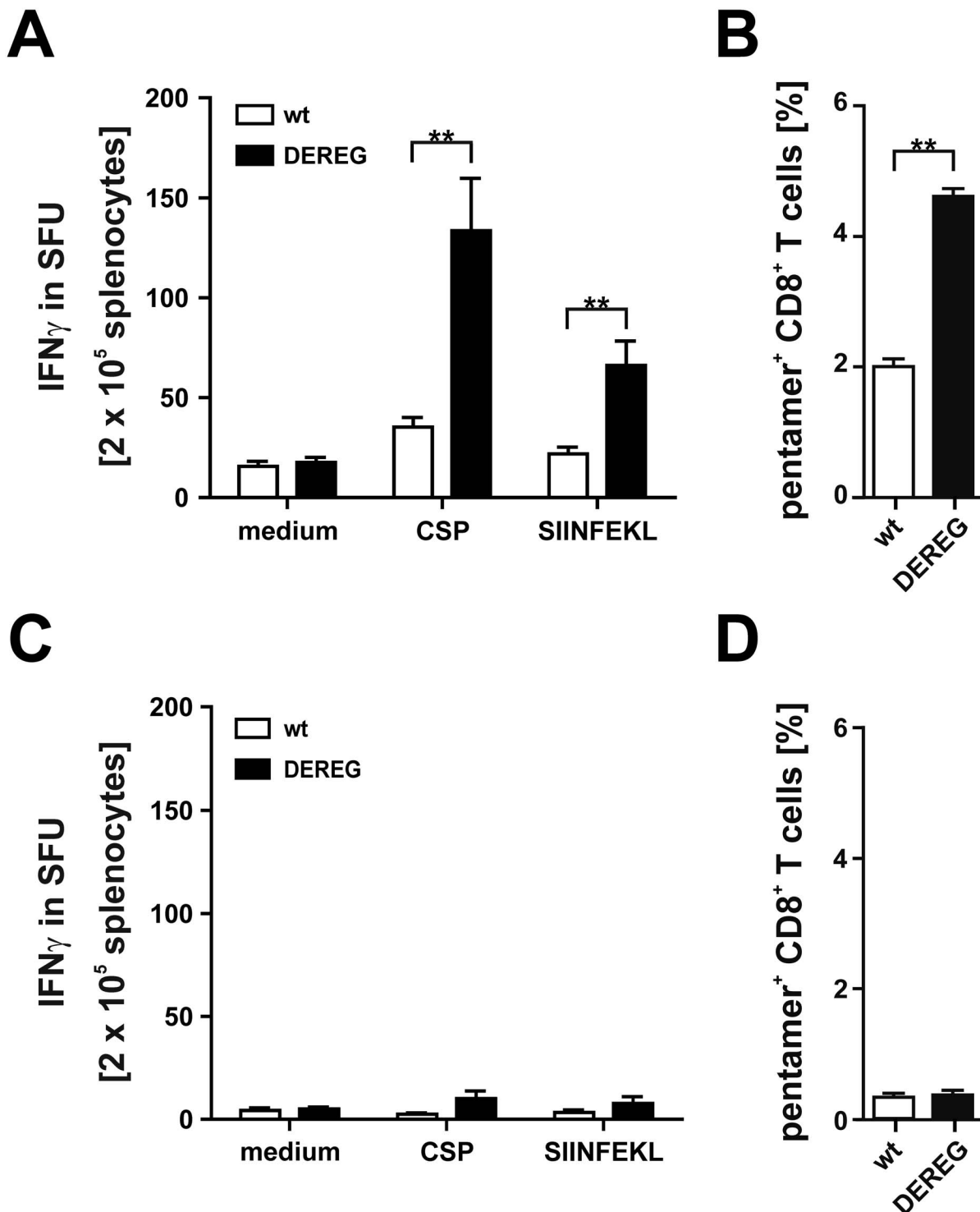


Figure 3. Effect of T_{reg} depletion on the efficacy of a primary immunization. (A) To analyze the effect of T_{reg} depletion on the efficacy of immunization, we immunized the F1 generation from male DERE mice on C57BL/6 background and female wildtype BALB/c mice (F1 DERE) on day 0 using i.p. application of 20 μ g of an adenylate cyclase toxoid (ACT-CSP) that carries an insert of the *P. berghei* CSP-derived CD8⁺ T cell epitope SYIPSAEKI presented on H-2K^d (corresponding to residues 245 to 253 from CSP) and the OVA-derived SIINFEKL epitope, presented in the context of H-2K^b (corresponding to residues 257 to 264 from OVA). On day +1, +2, and +3 immunized F1 DERE and F1 non-transgenic littermates were treated with DT. On day +7, spleen cells were restimulated with CSP-peptide or OVA-peptide and IFN γ production in spot forming units (SFU) was analyzed using ELISPOT. Data are expressed as mean \pm SD of 5 mice. This experiment was repeated twice with similar results. (B) Percentage of CD8⁺CSP⁺ spleen cells stained using PE-labeled anti-CD8 and APC-labeled MHC-I-pentamer loaded with CSP-peptide to detect CSP-specific CD8⁺ T cells. The mean of five immunized F1 DERE mouse or a F1 non-transgenic littermates is shown. The percentage of CD8⁺CSP⁺ T cells is expressed as mean \pm SD of whole lymphocytes. (C) A similar experiment was performed using again F1 DERE mice and F1 non-transgenic littermates that were treated on days +1, +2, and +3 with DT but without previous immunization. Data are expressed as mean \pm SD of 5 mice. This experiment was repeated once with similar results. (D) Percentage of CD8⁺CSP⁺ spleen cells stained using PE-labeled anti-CD8 and APC-labeled MHC-I-pentamer loaded with CSP-

peptide to detect CSP-specific CD8⁺ T cells. The mean of five immunized F1 DEREg mouse or a F1 non-transgenic littermates is shown. The percentage of CD8⁺CSP⁺ T cells is expressed as mean \pm SD of whole lymphocytes. **, $p < 0.01$. doi:10.1371/journal.pone.0104627.g003

number of CSP-specific CD8⁺ T cells, prompted us to consider whether this regimen would be capable to induce protection. To this end BALB/c DEREg mice and their non-transgenic littermates were immunized with ACT-CSP and treated as described with DT to deplete T_{reg}. At day 7 post immunization both groups of mice were challenged with *P. berghei* sporozoites (Fig. 4). The course of the infection was analyzed at the indicated time points. In the DEREg group, a lower increase in the levels of parasitemia after challenging was observed (Fig. 4 A). No significant differences in the survival time of the two groups could

be observed. It is noteworthy that a transient depletion of T_{reg} in DEREg mice without immunization did not influence the parasitemia (Fig. 4 B).

Due to the very high number of merozoites released by a single infected hepatocyte, a direct correlation between liver-stage burden and parasitemia is unlikely. In order to directly compare the protection induced by vaccination of T_{reg} depleted and not depleted mice, parasite burden in the liver after challenge with *P. berghei* sporozoites was determined using a quantitative PCR (qPCR) specific for *P. berghei* 18S rRNA normalized to a murine

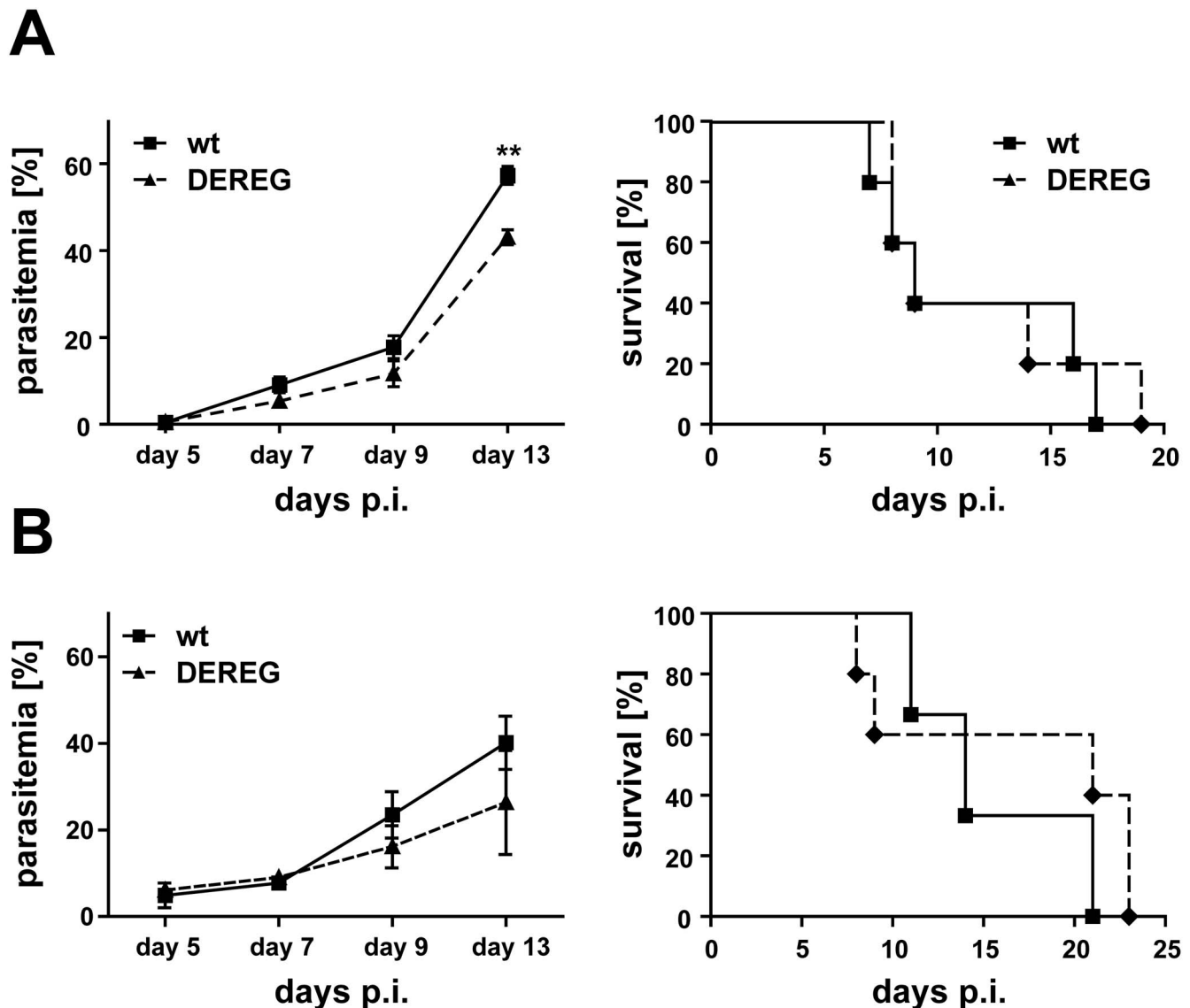


Figure 4. T_{reg}-depletion during primary immunization confers only a partial protection against challenge with *P. berghei* sporozoites. (A) To analyze the effect of T_{reg} depletion on protection DEREg-mice were immunized on day 0 using i.p. application of 20 μ g ACT-CSP. On day +1, +2, and +3 immunized DEREg-mice and non-transgenic littermates were treated with DT. At day 9 post immunization both groups of mice were challenged with 1000 *P. berghei* sporozoites i.v. The resulting parasitemia was monitored at the indicated time points. Parasitemia is expressed as mean \pm SD of five mice per group. This experiment was repeated twice with similar results. A single application of ACT-CSP to DT-treated wild type mice delayed the onset of the blood-stage. (B) The transient depletion of T_{reg} in DEREg mice without immunization did not influence the parasitemia. **, $p < 0.01$. doi:10.1371/journal.pone.0104627.g004

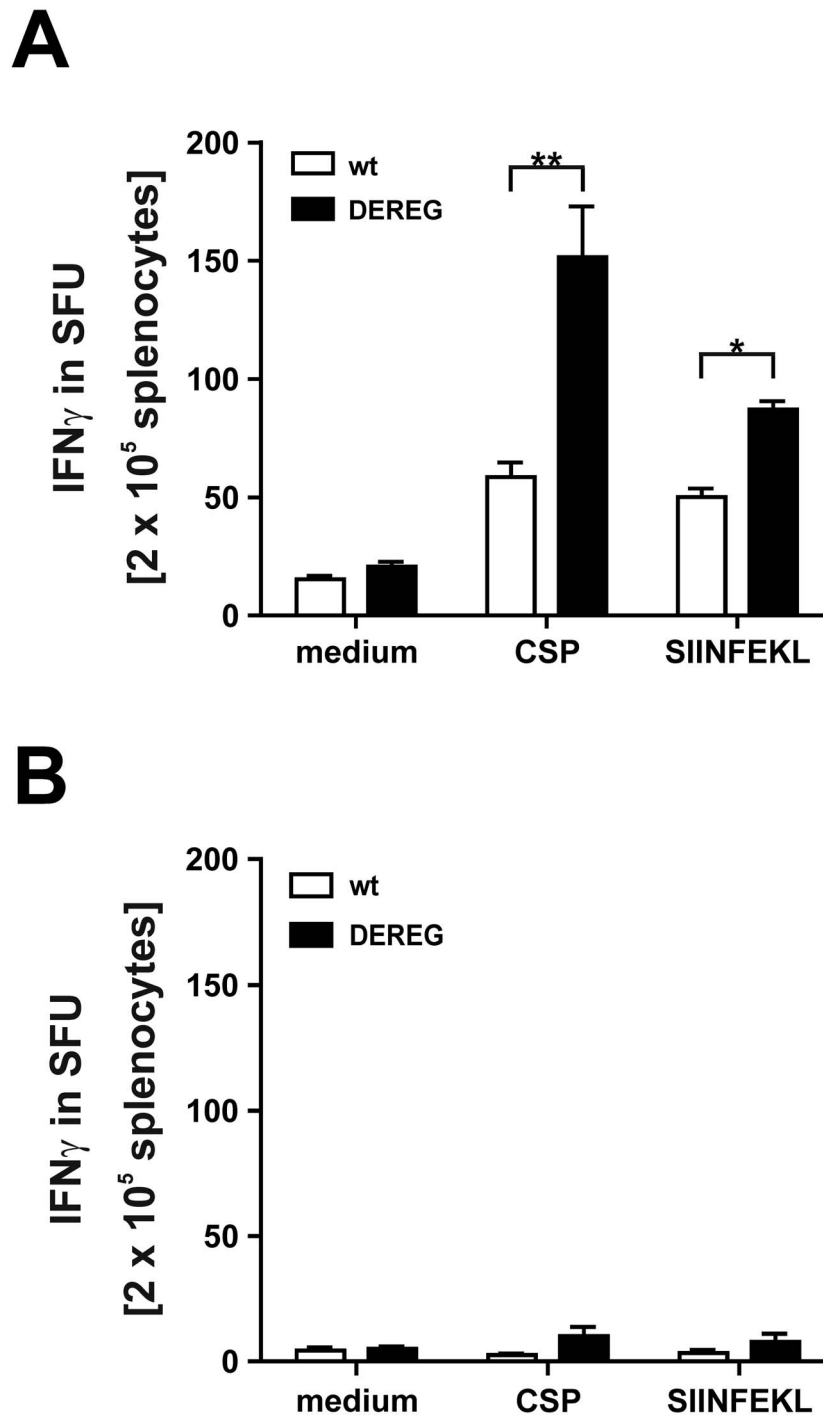


Figure 6. Effect of T_{reg} -depletion on the efficacy of a boost immunization. (A) To analyze the effect of T_{reg} depletion on the efficacy of a boost immunization, we vaccinated F1 DERE mice or their non-transgenic littermates with ACT-CSP on day 0 and day 14. Both groups of mice were treated on days 15, 16 and 17 with DT. On day +25, spleen cells were restimulated with CSP-peptide or OVA-peptide and $IFN\gamma$ production in spot forming units (SFU) was analyzed using ELISPOT. Data are expressed as mean \pm SD of 5 mice. This experiment was repeated twice with similar results. *, $p < 0.05$, $p < 0.01$. (B) A similar experiment was performed using as well F1 DERE mice and F1 non-transgenic littermates that were treated on days 15, 16 and 17 with DT but without previous immunization. doi:10.1371/journal.pone.0104627.g006

Using the previously described DERE mouse it is possible to deplete $Foxp3^+$ T_{reg} at selected time points by DT injection [17]. Upon injection on three consecutive days the vast majority of T_{reg} cells were depleted although they started to reappear soon after stopping DT injection. Using this scheme we generated a time

window of 3 to 5 days were the numbers of T_{reg} were significantly reduced without observing overt pathology. $Foxp3^+$ T_{reg} rapidly reappear after depletion and these reappearing T_{reg} have a similar suppressive potential as the initial T_{reg} population. Thus, DERE-mice are an ideal model to study the influence of T_{reg} on T cell

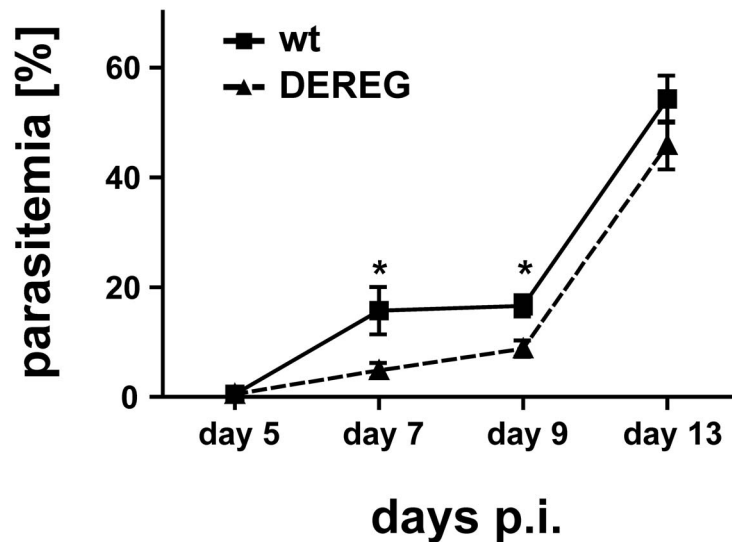


Figure 7. T_{reg} -depletion during boost immunization leads to decreased parasitic burden after challenge with *P. berghei* sporozoites. BALB/c DEREg mice or non-transgenic littermates were immunized with ACT-CSP on day 0 and day 14. Both groups of mice were treated on days 15, 16 and 17 with DT. At day +21 mice were challenged i.v. with 1000 *P. berghei* sporozoites and the resulting parasitemia was analyzed at the indicated time points. Data are expressed as mean \pm SD of 5 mice per group. This experiment was repeated twice with similar results. *, $p < 0.05$. doi:10.1371/journal.pone.0104627.g007

priming during initial antigen encounter elicited by vaccines, whereas all subsequent steps will take place in the presence of functional T_{reg} . In contrast, data on anti-CD25 treatment suggest a long lasting effect of at least 13 to 20 days and therefore not only priming but also expansion and contraction will be influenced by the lack of T_{reg} [31].

We used in the present study ACT-CSP that induces IFN γ producing CD8⁺ T cells against a single epitope of CSP of *Pb* [13]. Although CSP induces a good immune response and confers protection, is also known that in certain models it has a minimal role in sterile protection, because other antigens are responsible for the induction of sterile immunity [32,33].

The number and function of these T cells can be analyzed by MHC-class I multimer technology and ELISPOT, respectively. Upon depletion of T_{reg} during priming we found a strong enhanced number of these CSP-specific T cells producing IFN γ . It is noteworthy that at the time point of *ex vivo* analysis, the T_{reg} population already had reached its original size and was proven to be functional. This suggests that the presence of T_{reg} during initial antigen-encounter limits the number of responding T cells.

Although T_{reg} depletion exhibited a strong effect on the vaccine-induced response upon primary vaccination and boost immunization, the secondary vaccination did not show an additional increase in the magnitude of CSP-specific T-cell expansion. The enhanced number of antigen specific CD8⁺ T cells after T_{reg} depletion was only transient, which indicates that the strong induction of T cells in the absence of T_{reg} is followed by a strong contraction of the T cell pool in the presence of T_{reg} . Ultimately no differences in the number of CSP-specific T cells were found in both groups several weeks after immunization, although in both groups a significant number of CSP-specific T cells remained detectable. A transient depletion of T_{reg} during boost immunization was also accompanied by an increased number of antigen-specific T cells. However, very similar to the situation observed during priming the differences waned when analyzed at later time points. This result shows that a simple correlation between the number of antigen-specific T effector that were primed initially during immunization and the number of remaining memory T

cells does not exist. Hence our data suggest that the efficacy of vaccines, which are intended to induce memory responses, cannot be estimated by analyzing the number of T cells that were primed initially.

T_{reg} were shown to directly interact with DC and to mitigate their maturation process [34,35]. Previously it was shown that depletion of T_{reg} in DEREg-mice is accompanied by an increase of DC with higher costimulatory activity [36]. Thus T_{reg} might limit the signal strength by preventing maturation of DC, which subsequently influence the priming of T cells. In this scenario a depletion of T_{reg} would augment priming of T cells leading to an increase of T effector cells but a decrease of memory T cells. However, T_{reg} were also shown to interact with T cells by either contact-dependent [37,38] or independent mechanisms [39–41]. Hereby they might influence the expansion of T cells. We observed no increased induction of memory T cells upon depletion of T_{reg} . If T effector cells and memory T cells develop in parallel our data would imply that only the size of the T effector cell pool but not the size of the memory T cell pool is regulated by T_{reg} . Further studies are needed to pinpoint the exact function of T_{reg} during priming and expansion on the subsequent development of T cell subsets. Whatever the exact function might be, our data have important implications for vaccine development since they indicate that T_{reg} depletion, although leading to an enhanced number of short-lived T effector cells, might not be associated with an enhanced number of long-living T memory cells, which is the major goal of prophylactic vaccines. Hence, transient inhibition of T_{reg} function might be more beneficial when a strong but short-lived immune response is desired for example in therapeutical vaccinations e.g. against tumours where an immune response against self-antigens should be elicited [42].

The infection with *P. berghei* sporozoites is quite a challenge for the induced immune response. Only very few hepatocytes are infected and within 48 h a single infected cell may release up to 30,000 merozoites which is sufficient to trigger the blood-stage of malaria. Thus CD8⁺ T cells specific for antigen presented on MHC-I of hepatocytes must find very few infected cells within a large excess of uninfected cells and combat them in a very narrow

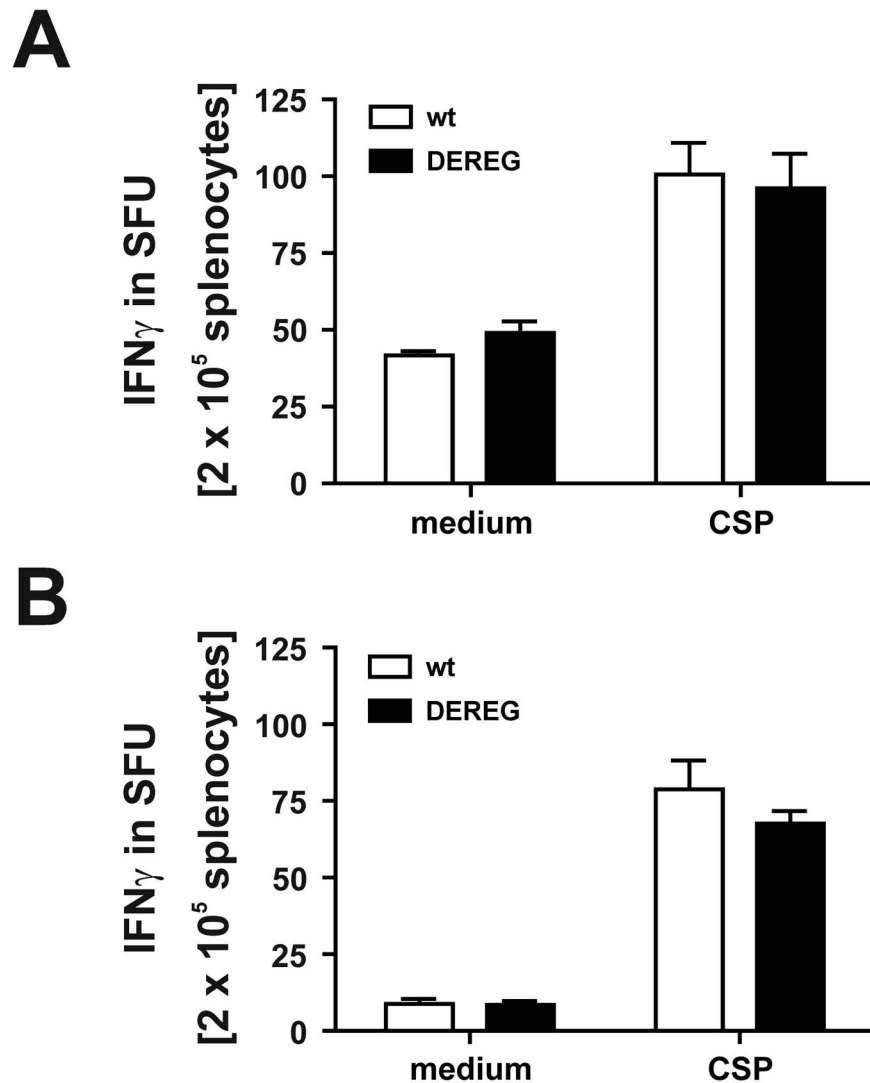


Figure 8. T_{reg} -depletion during boost immunization does not affect memory responses. BALB/c DEREg mice or non-transgenic littermates were immunized with ACT-CSP on day 0 and day 14. Both groups of mice were treated on days 15, 16 and 17 with DT. Spleen cells of mice were restimulated with CSP-peptide after (A) 6 weeks, and (B) 9 weeks. The number of $IFN\gamma$ producing T cells in spot forming units (SFU) was analyzed by ELISPOT. Data are expressed as mean \pm SD of 5 mice per group and time-point. This experiment was repeated once with similar results. doi:10.1371/journal.pone.0104627.g008

time-window. This was demonstrated in a study where different amounts of CSP-specific cells were transferred [43]. Using this approach a threshold of protection was defined. For sterile protection against the liver-stage almost 100 to 1000-fold more T cells were needed in comparison with bacterial or viral infections of the liver. Therefore the requirements on a protective immune response in the *P. berghei* model are very high and probably higher in comparison to human malaria since here the liver-stage lasts much longer [44]. Nevertheless we observed a reduced parasitemia when mice were immunized in the absence of T_{reg} . Using qPCR to detect *P. berghei* 18S RNA in the liver 30 h postchallenge, we found that depletion of T_{reg} during immunization was also accompanied by a significant reduction of parasite burden, which indicates that the increased number of CSP-specific $CD8^+$ T cells are capable to combat infected hepatocytes. However, the difference in the parasitemia between immunized mice that were depleted for T_{reg} or immunized control mice vanishes when mice were challenged at later time-points (data not shown). This corroborated our observation that the enhanced

induction of antigen-specific T cells is not accompanied by an increased level of long-living T cells of a memory subtype.

Mice were challenged at day 7 post-priming or boosting and 3 days after DT injection, a time at which innate immune responses might still interfere with the infection. Since T_{reg} are known to suppress innate immunity [45], it is not certain whether the slight decrease of parasitemia in DEREg mice was due to increased numbers of $CD8^+$ T cells and the subsequent decrease in liver-stage burden or due to a more active innate immune response against the blood-stage. Our data cannot support the notion that the reduced parasitemia was mediated by $CD8^+$ T cells, because the numbers of $IFN\gamma$ -producing $CD8^+$ T cells at day 7 post primary vaccination (Fig. 3) and at day 7 post secondary vaccination (Fig. 5) were similar. To clarify this, it is important for further investigations to include a control group with mice depleted of $CD8^+$ T cells before parasite challenge.

Collectively our results demonstrate that a specific and transient depletion of T_{reg} could enhance a vaccine induced $CD8^+$ T cell response during prime and boost immunization. Importantly, the

enhanced expansion of T effector cells after T_{reg} -depletion was not accompanied by an increased long-lasting memory response, demonstrating that the design of novel vaccine strategies must not only be focussed on the optimisation of a strong short term immune response since this is not necessarily leading to a strong memory response.

Supporting Information

Figure S1 Kinetics of T_{reg} depletion in DEREg mice. The consecutive application of DT i.p. in DEREg mice over three days leads to a depletion of $CD4^{+}Foxp3^{+} T_{reg}$ within 72 h. Mice were treated with DT on days +1, +2 and +3. The percentage of

$CD4^{+}Foxp3^{+} T_{reg}$ in the blood was measured in three F1 DEREg C57BL/6 \times BALB/c mice by flow cytometry at the indicated time points. Data is expressed as mean \pm SEM in relation to all cells in the lymphocyte gate as defined by the respective forward- and sideward scatter.

(TIF)

Author Contributions

Conceived and designed the experiments: MdREM TJ BF. Performed the experiments: MdREM CS. Analyzed the data: MdREM AL TJ. Contributed reagents/materials/analysis tools: VH TS ST. Wrote the paper: MdREM AL TJ.

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