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# IL-18 and Subcapsular Lymph Node Macrophages are Essential for Enhanced B Cell Responses with TLR4 Agonist Adjuvants

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Designing modern vaccine adjuvants depends on understanding the cellular and molecular events that connect innate and adaptive immune responses. The synthetic TLR4 agonist glycopyranosyl lipid adjuvant (GLA) formulated in a squalene-in-water emulsion (GLA-SE) augments both cellular and humoral immune responses to vaccine Ags. This adjuvant is currently included in several vaccines undergoing clinical evaluation including those for tuberculosis, leishmaniasis, and influenza. Delineation of the mechanisms of adjuvant activity will enable more informative evaluation of clinical trials. Early after injection, GLA-SE induces substantially more Ag-specific B cells, higher serum Ab titers, and greater numbers of T follicular helper (T<sub>FH</sub>) and Th1 cells than alum, the SE alone, or GLA without SE. GLA-SE augments Ag-specific B cell differentiation into germinal center and memory precursor B cells as well as preplasmablasts that rapidly secrete Abs. CD169<sup>+</sup> SIGNRI<sup>+</sup> subcapsular medullary macrophages are the primary cells to take up GLA-SE after immunization and are critical for the innate immune responses, including rapid IL-18 production, induced by GLA-SE. Depletion of subcapsular macrophages (SCM $\phi$ ) or abrogation of IL-18 signaling dramatically impairs the Ag-specific B cell and Ab responses augmented by GLA-SE. Depletion of SCM $\phi$  also drastically reduces the Th1 but not the T<sub>FH</sub> response. Thus the GLA-SE adjuvant operates through interaction with IL-18-producing SCM $\phi$  for the rapid induction of B cell expansion and differentiation, Ab secretion, and Th1 responses, whereas augmentation of T<sub>FH</sub> numbers by GLA-SE is independent of SCM $\phi$ . *The Journal of Immunology*, 2016, 197: 4351–4359.

Augmentation of Ag-specific B cell responses and subsequent Ab production is central to the development of effective vaccines. In settings such as the emergence of new pandemics, intentional release of bioterror agents, and on-demand travelers' vaccines, the rapid initiation of humoral immunity with practical vaccine approaches is highly desirable. The last decade has seen the licensure of several vaccine adjuvants that augment humoral immunity, including the squalene-in-water emulsion (SE) adjuvants MF59 and AS03, and the TLR4 agonist-containing

adjuvants AS01 and AS04. The study of B cell responses following immunization with clinically relevant adjuvants has been primarily limited to serum Ab analyses and the identification of Ab-secreting cells. Compared with the wealth of information regarding T cell and Ab responses to vaccine adjuvants, much less is known regarding the early events following immunization that induce Ag-specific B cell and T follicular helper cell (T<sub>FH</sub>) responses and the mechanisms by which these adjuvants determine the course of these humoral responses.

The initial wave of secreted IgM and class-switched Abs is produced by CD138<sup>+</sup> preplasmablasts residing in extrafollicular spaces. Concurrently, CD95<sup>+</sup> GL7<sup>+</sup> germinal center B cells differentiate within follicles, resulting in affinity-matured Abs, memory cells, and long-lived plasmablasts. Memory B cells expressing IgM and CD38 are generated early in the response — independently of germinal centers — and are distinguished from the bulk population of naive B cells by their Ag specificity (1). The production of a B cell response is coordinated by multiple innate and adaptive responses. The earliest encounter of foreign material in the draining lymph node (LN) can occur within minutes via cell-free transport from the infection or injection site to the draining LNs (2). This material is captured by CD169<sup>+</sup> subcapsular macrophages (SCM $\phi$ ) that reside in LN sinusoidal spaces, allowing delivery of material to B cell follicles (3). Subsequent T dependent B cell differentiation is regulated by multiple subsets of CD4 T helper cells. PD1<sup>+</sup> CXCR5<sup>+</sup> T<sub>FH</sub> expressing the transcription factor Bcl-6 and localized within B cell follicles are particularly important for germinal center responses and the generation of long-lived memory responses (4). T<sub>FH</sub> cells, or perhaps their progenitors, have also been shown to be necessary for extrafollicular responses (5).

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Abbreviations used in this article: CLL, clodronate-loaded liposome; DC, dendritic cell; GC, germinal center; GLA, glycopyranosyl lipid adjuvant; GLA-SE, GLA formulated in SE; LN, lymph node; MPL, monophosphoryl lipid A; T<sub>FH</sub>, T follicular helper cell; SE, squalene-in-water emulsion; SCM $\phi$ , subcapsular macrophage.

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The prototypical TLR4 agonist, LPS, is a powerful stimulant of the innate immune system, but it needs to be detoxified for use in human vaccine preparations. Acid and base treatment of LPS results in deacylation and removal of the polysaccharides and one of the phosphates yielding the biological product monophosphoryl lipid A (MPL), which is a safe and effective immune stimulant, and is present in commercial vaccines produced by GlaxoSmithKline. Glycopyranosyl lipid adjuvant (GLA) is a synthetic monophosphorylated lipid A analog with six acyl chains and, like MPL, is a safe and effective TLR4 agonist for use in adjuvant preparations (6). GLA is an effective vaccine adjuvant when formulated in an SE. It generates a robust Th1 response and augmented Ab production skewed toward IgG2 class-switching against numerous infectious disease- and cancer-associated Ags (7–14). In humans, GLA formulated in SE (GLA-SE) promotes production of Ag-specific Th1 cells making IFN- $\gamma$ , TNF, and IL-2, augmentation of serum Ab titers, preferential switching to IgG1 and IgG3 subtypes, and increases in neutralizing Ab titers when paired with protein Ags (15, 16). The adjuvant activity of GLA is critically dependent on its formulation. When GLA-SE is compared with aqueous formulations of GLA, unique innate and adaptive features emerge (7, 17, 18). We have previously demonstrated that the Th1 potentiating activity of GLA-SE is mediated in part by MyD88 and TRIF signaling, inflammatory caspases and IL-18 as well as by type I and II IFNs, IL-12, and the transcription factor T-bet (18–20). In the present paper we evaluate the cellular and molecular events necessary for augmentation of B cell, Ab, and T<sub>FH</sub> responses by the GLA-SE adjuvant. Defining the cellular and molecular mechanisms of adjuvant activity of GLA-SE will enable more informative clinical evaluation of vaccines containing this adjuvant, and provide a rational pathway for the development of next-generation vaccine adjuvants by identifying the key parameters to optimize during early-stage development.

## Materials and Methods

### Animals and immunizations

Female C57BL/6 mice and IL-18R1<sup>-/-</sup> mice aged 6–10 wk were purchased from The Jackson Laboratory. CD169-DTR mice were provided by the RIKEN BRC through the National Bio-Resource Project of the Ministry of Education, Culture, Sports, Science and Technology, Japan (21, 22). All animal experiments and protocols used in this study were approved by the Infectious Disease Research Institute's Institutional Animal Care and Use Committee. For the adaptive response assessments mice were immunized s.c. via foot pad or base of tail injection with 1.25 or 5  $\mu$ g of GLA respectively, in combination with 1  $\mu$ g of R-PE (Prozyme, Hayward, CA) or 1  $\mu$ g of the *Mycobacterium tuberculosis* Ag ID87 (23) or 2.5  $\mu$ g of the *M. tuberculosis* Ag ID97 (24). Squalene oil formulations were emulsified with egg phosphatidylcholine or synthetic dimyristoyl phosphatidylcholine as described (25) and used after dilution in saline for injection to 2% oil. For the innate response assessment mice were immunized via an i.m. injection in the quadriceps with 5  $\mu$ g of DiD-labeled GLA-SE. To label GLA-SE, DiD Oil, DiC18 (5) (Molecular Probes) was solubilized in DMSO at 25 mg/ml. DiD was added to GLA-SE at 25  $\mu$ g/ml, mixed well, and incubated at room temperature for 10 min. Excess DiD was removed using a PD-10 desalting column (GE).

### SCM $\phi$ depletion

Mice were treated with 30  $\mu$ l of clodronate-loaded liposomes (5 mg/ml clodronate disodium salt) (Encapsula Nano Sciences, Brentwood, TN) via intradermal hock injection. Mice were immunized 6 d later via footpad, hock or i.m. injection in the quadriceps.

### Selective deletion of CD169<sup>+</sup> macrophages

CD169-DTR mice were injected i.p. with diphtheria toxin (Sigma-Aldrich), at 20 ng/g body weight, 2 d prior to subsequent immunization.

### Serum endpoint titer determination

Serum endpoint titer ELISAs were performed on Corning high bind 384 well plates. Plates were coated overnight at 4° with 1  $\mu$ g/ml PE. Detection

Abs were Ig (H+L) HRP and IgG2c HRP (Southern Biotech). Endpoints were set as the minimum dilution at which values were lesser than or equal to the mean + 3 SD of naive controls.

### LN single-cell suspension preparation

Single-cell suspensions of LN were generated by mechanical homogenization in PBS, 0.5% BSA in the presence of Halt protease inhibitor (Thermo Scientific), and 10  $\mu$ g/ml Brefeldin A (BD Bioscience). For Fig. 2, LN were homogenized and digested at 37°C for 30 min in RPMI 1640 in the presence of 8  $\mu$ g/ml Collagenase IV (Worthington), and 20  $\mu$ g/ml DNase I (Roche).

### Flow cytometry

Samples were prepared by Fc receptor blocking (clone 2.4G2). B cells were stained in 1% FBS, 2 mM EDTA in PBS with the following: CD95-Bv421 (clone JO2), CD138-Bv605 (clone 16A8), B220-Bv785 (clone RA3-6B2) or CD19-Bv785 (clone 1D3), IgM-PerCP-eFluor710 (clone IL/41), IgD-APC-Cy7 (clone 1126C.2A), CD38-AF700 (clone 90), lineage mixture: Ly6G-FITC (clone 1A8), CD11b-FITC (clone M1/70), CD11c-FITC (clone N418), F4/80-FITC (clone BM8), Ter119-FITC (clone TER119), and CD90.2-FITC (clone OX-7). Cells were fixed and permeabilized with Fix/Perm buffer (BD Biosciences) and stained for intracellular IgG2-Bv510 (clone 5.7). PE-specific B cells were identified using 1  $\mu$ g/ml PE for surface staining, and 0.1–0.3  $\mu$ g/ml PE for intracellular staining.

For peptide-MHCII tetramer staining, the Ag85B p25 tetramer (NIH Tetramer Core Facility) was incubated in 100  $\mu$ l with a maximum of 4  $\times$  10<sup>6</sup> cells at 37°C for 1.5 h at 13  $\mu$ g/ml with anti-CD16/32 Ab. Cells were washed and stained for CD4-APC-H7 (clone RM4-5), CD44-AF700 (clone IM7), PD-1-Bv605 (clone 29F1A12), CXCR5-PerCP-710 (clone SPRCL5), and lineage mixture: CD8a-FITC (clone 53-6.7), CD11b-FITC, CD11c-FITC, Ly6G-FITC, Ter-119-FITC, F4/80-FITC, and CD19-FITC at room temperature for 30 min. Cells were then fixed and permeabilized as above and stained for Bcl6-AF647 (clone BCL-DWN), and Tbet-Bv-421 (clone 4B10) overnight at 4°C.

Identification of DiD<sup>+</sup> cells was performed by surface staining draining LN cells by blocking CD16/32 and staining for CD11b-eF450, CD90.2-Bv510 (clone 53-2.1), CD19-Bv785 (clone 6D5), purified SIGNR1 (clone eBio22D1, Armenian hamster host), Ly6C-PerCP-Cy5.5 (clone HK1.4), CD169-PE (clone 3D6.112), Ly6G-PE-CF594, and CD11c-PE-Cy7 at 4°C for 30 min. Cells were then washed and stained with a secondary Ab against Armenian hamster coupled to AF488.

Quantitation of cells was carried out by applying the frequencies of cell populations to cell counts obtained with a GUAVA EasyCytometer (Millipore, Billerica, MA). Data were collected on LSRII or Fortessa flow cytometers (BD Bioscience). Data were analyzed using FlowJo (Tree Star, Ashland, OR). Statistical analysis was performed using Prism software (GraphPad Software, La Jolla, CA).

### Immunofluorescence microscopy

LN cells were collected and embedded in OCT Compound (Tissue-Tek). Cryostat sections (6  $\mu$ m thick) were mounted on poly-lysine microscope slides (Thermo Scientific). Brightfield and DiD images of cryostat sections were acquired using Nikon Eclipse Ti-5 before acetone fixation. Sections were subsequently dried and fixed in cold acetone for 5 min at -20°C, blocked with 3% BSA in PBS-blocking buffer at room temperature for 30 min, and incubated with anti-CD169 (Thermo Scientific) and SIGNR1 (eBioSciences) Abs diluted in the blocking buffer for 1 h. Primary Abs were detected with Goat-anti-Rat-AF488 (against CD169; BioLegend) and Goat-anti-Hamster IgG-568 (against SIGNR1; Abcam). Images were acquired with the Nikon Eclipse Ti-5 microscope, using a 4  $\times$  objective lens and NIS-Elements D3.2 Software, and processed with Fiji software.

### Cytokine and chemokine protein levels quantitation

The concentration of cytokines and chemokines were measured using a microbead-based ELISA system (ProcartaPlex Mouse; Affymetrix eBioscience), according to the manufacturer's directions.

## Results

### GLA-SE induces rapid B cell responses

To better understand the B cell responses induced by GLA-SE, we analyzed Ag-specific B cells via immunization with PE either alone or adjuvanted with GLA-SE or its constituent components, SE or GLA, a strategy previously described to identify Ag-specific B cells

by flow cytometry (26, 27). We also included alum as a well-established adjuvant with which to benchmark the adjuvanticity of GLA-SE. Seven days after immunization with PE alone, very few Ag-specific B cells were detected in the draining LN (Fig. 1A). These rare PE<sup>+</sup> B cells elicited by immunization with unadjuvanted Ag were predominately IgD<sup>+</sup>, CD38<sup>+</sup>, IgM<sup>+</sup> putative memory B cells (Fig. 1B). Adjuvanting PE with alum, GLA or SE elicited a minor but not statistically significant increase in PE<sup>+</sup> B cells with an increase in the proportion of CD95<sup>+</sup>, GL7<sup>+</sup>, IgD<sup>-</sup> germinal center B cells (Fig. 1A, 1B). In contrast, GLA and SE synergized to elicit substantially more PE-specific B cells (Fig. 1A). In addition to inducing germinal center and putative memory B cells, GLA-SE uniquely produced isotype switched preplasmablasts (B220<sup>lo</sup>, CD138<sup>+</sup>, IgD<sup>-</sup>, IgM<sup>-</sup>), which are an early source for circulating Ab (Fig. 1B). Accordingly, early after immunization PE-specific serum Ab titers were dramatically increased with GLA-SE compared with the other immunizations including alum (Fig. 1A). Furthermore, compared with alum, GLA or SE, GLA-SE uniquely augmented the frequency of Ag-specific B cells expressing intracellular IgG2c and the IgG2c serum Ab titers (Fig. 1C).

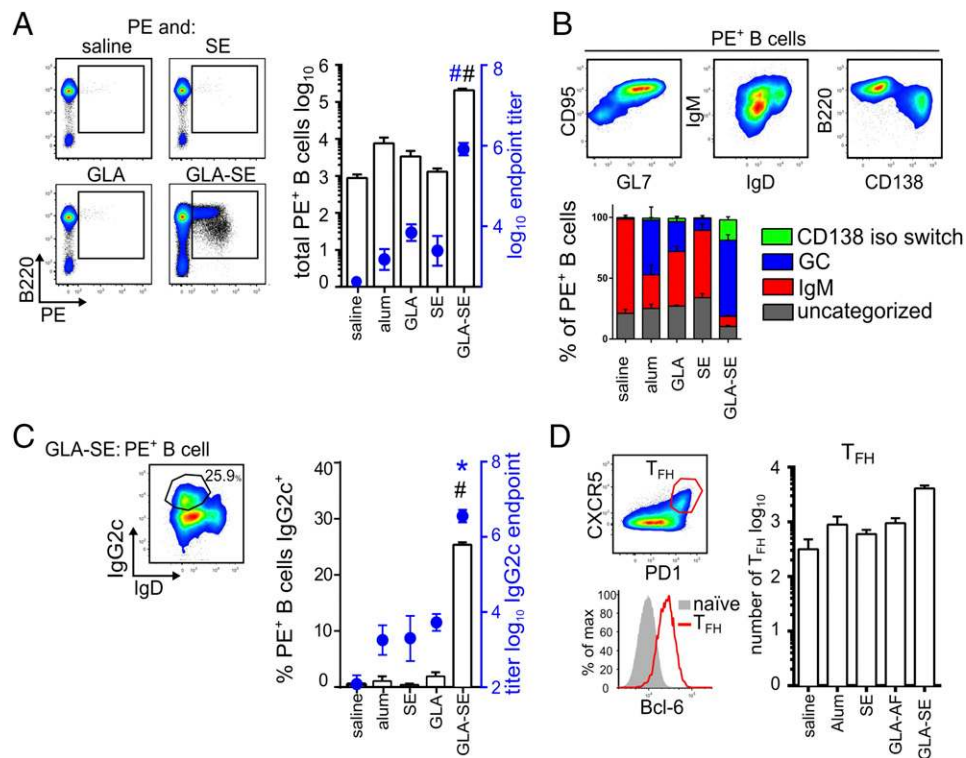
Given the importance of T<sub>FH</sub> cells to humoral immunity (4), we also enumerated T<sub>FH</sub> cells based on surface expression of CXCR5 and PD1 and expression of the transcription factor Bcl-6. GLA-SE induced more T<sub>FH</sub> cells in the draining LNs when compared with Ag alone or adjuvanted with alum, GLA, or SE (Fig. 1D). Taken together, these results suggest that the rapid and dramatic expansion of

early B cell responses, especially preplasmablasts, Ab production, and T<sub>FH</sub> cells is unique to the combination GLA-SE adjuvant formulation.

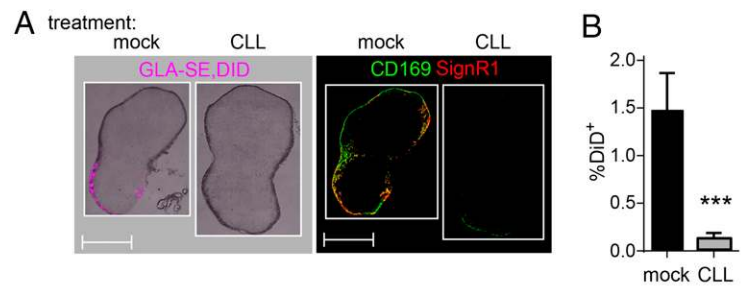
*SCMφ are essential for GLA-SE uptake*

To better understand how GLA-SE augments rapid B cell responses, we identified the cells interacting with GLA-SE immediately after immunization. The cells interacting with the adjuvant were visualized by labeling the adjuvant with the lipophilic fluorescent dye DiD. Fifteen minutes after injection GLA-SE colocalized with the CD169<sup>+</sup> SIGNR1<sup>+</sup> medullary macrophage population in the draining LN (28) (Fig. 2A, 2C). Other populations including monocytes (CD11b<sup>+</sup> Ly6C<sup>+</sup>) also took up the adjuvant, but in lower numbers and with less DiD signal intensity (Fig. 2C, 2D). Of note, DiD<sup>dim</sup> B cells were the most abundant DiD<sup>+</sup> population by absolute numbers, although these cells represented only a small fraction of the total LN B cell population (~3%) (Fig. 2D).

To determine the significance of GLA-SE capture by SCMφ, including the medullary and sinusoidal macrophage subsets, we depleted these cells by injection of clodronate-loaded liposomes (CLL), and waited for 6 d to allow the repopulation of resident phagocytes (monocytes, dendritic cells, and neutrophils) but not the SCMφ (3), as confirmed by analysis of the cellular content of the lymph nodes of CLL-treated animals (Supplemental Fig. 1). SCMφ depletion dramatically affected GLA-SE uptake by LN cells with <0.2% of the LN cells being DiD<sup>+</sup> 15 mins postinjection compared with an average of 1.5% in the mock-treated mice (Fig. 2B). Surprisingly,



**FIGURE 1.** Rapid B cell responses elicited by different formulations of the TLR4 adjuvant GLA. B cell responses in the injection site draining inguinal and axillary LN of mice immunized with the indicated adjuvant and PE were analyzed 7 d after immunization. (A) Ag specific B cells were enumerated by PE staining and cell populations were determined by cell surface marker staining. Total numbers of PE staining B cells in draining LNs were determined (bars). Overlaid are the total Ig (H+L) serum Ab endpoint titers (circles). (B) FACS plots of GLA-SE induced PE<sup>+</sup> B cells are shown. Subsets of B cells were identified by surface marker expression as CD38<sup>+</sup> IgM<sup>+</sup> memory B cells (IgM), IgD<sup>-</sup> CD95<sup>+</sup> GL7<sup>+</sup> GC cells, CD138<sup>+</sup> B220<sup>lo</sup> CD38<sup>-</sup> IgM<sup>-</sup> IgD<sup>-</sup> preplasmablasts (CD138 iso switch), or uncategorized. The stacked histograms represent the draining LN PE-specific B cell composition for each immunization. (C) Representative FACS plot for the identification of PE<sup>+</sup> IgG2c B cells in draining LNs are shown. Frequency of IgG2c<sup>+</sup> PE-specific B cells (bars) are overlaid with serum IgG2c endpoint titers (circles). (D) Representative FACS plot for Bcl-6 staining of T<sub>FH</sub> cells (CXCR5<sup>+</sup>PD1<sup>+</sup>) versus naive cells. Following immunization the numbers of T<sub>FH</sub> cells per LN were determined. Bars and circles are drawn to the mean values + SEM. Data are representative of two independent experiments with three or four animals per group. The *p* values were determined by one-way ANOVA/Dunnett's test, #*p* < 0.001 compared with all other groups, \**p* < 0.05 compared with saline.



**FIGURE 2.** GLA-SE-DiD cellular uptake after SCM $\phi$  depletion. Mice were treated with clodronate-loaded liposomes (CLL) or mock treated with saline 6 d prior to injection with DiD-labeled GLA-SE or saline. Draining inguinal LN were harvested 15 min after immunization. **(A)** Immunofluorescence microscopic images of representative LN sections were taken before and after staining with CD169 and SIGNR1 to assess GLA-SE distribution. **(B)** FACS analysis of total DiD staining cells per LN were determined. **(C)** A representative histogram of DiD-GLA-SE staining per cellular population in a mock treated DiD-GLA-SE immunized mouse is shown and DiD-GLA-SE geometric means for each cellular population were plotted. **(D)** The number of DiD<sup>+</sup> cells per population and the percentage of DiD<sup>+</sup> cells within each population was determined for mock- and CLL-treated groups. Data are representative of two independent experiments with four or five animals per group. Bars are drawn to the mean + SEM. The *p* values were determined by Student *t* test, \*\*\**p* < 0.005.

adjuvant uptake was abrogated in cell populations that were not depleted by CLL treatment including monocytes, macrophages, dendritic cells, T cells, and B cells (Fig. 2D). These results indicate that SCM $\phi$ , and most probably the medullary macrophages, are critical to early GLA-SE capture and subsequent distribution in the LN.

#### SCM $\phi$ are critical for the innate response to GLA-SE

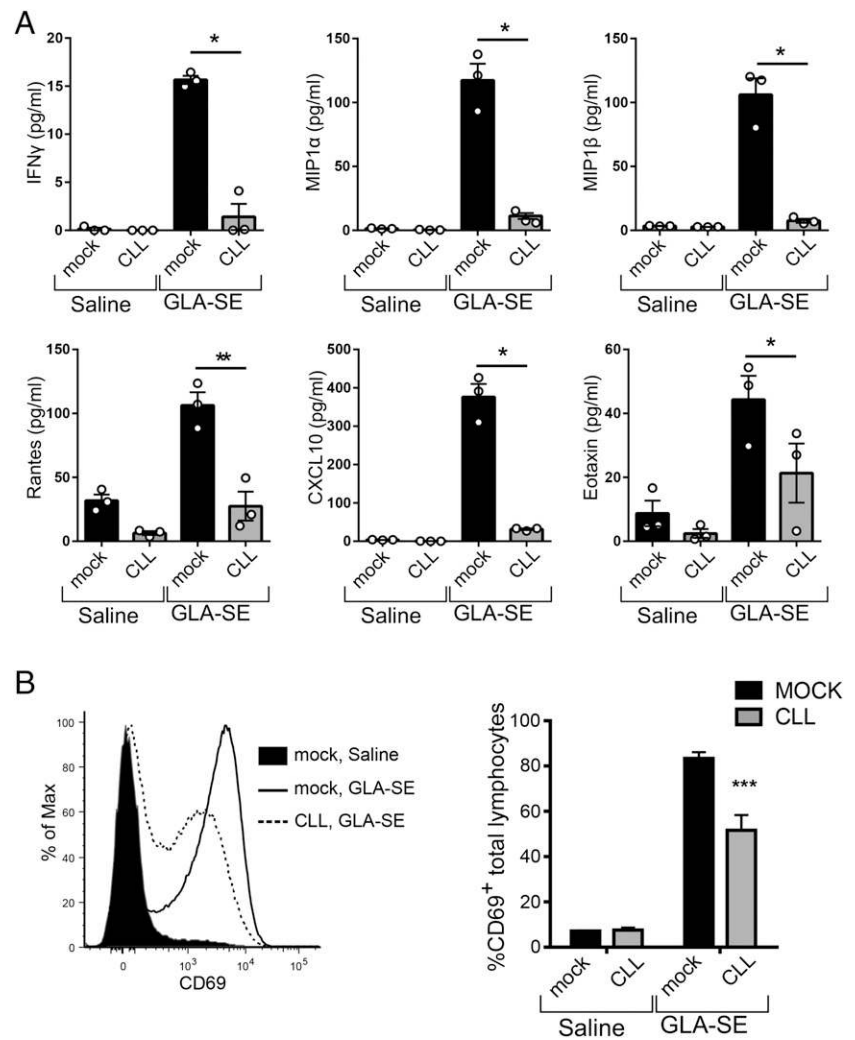
Immunization with GLA-SE initiates a complex innate immune response including production of IFN- $\gamma$ , which is essential for the adjuvanticity of GLA-SE (18) and a number of chemoattractants including RANTES, MIP1 $\alpha$  and  $\beta$ , Eotaxin, and CXCL10 (Fig. 3A). Depletion of SCM $\phi$  by CLL treatment severely compromised the innate response to GLA-SE (Fig. 3A). CD69 expression increases upon GLA-SE immunization in a type I and II IFN-dependent manner and is thus a sensitive readout for the response to GLA-SE (18, 19). Treatment with CLL resulted in a substantial decrease in the induction of CD69 on total lymphocytes 18 h after injection with GLA-SE (Fig. 3B). The decrease in these early immune responses to GLA-SE after SCM $\phi$  depletion indicated that these cells were critical for the innate immune response to GLA-SE.

#### SCM $\phi$ are necessary for B cell and Th1 but not T<sub>FH</sub> induction by GLA-SE

To assess the contribution of the SCM $\phi$  to the adjuvanticity of GLA-SE, we examined the adaptive immune response to PE+GLA-SE immunization in CLL-treated mice. Seven days after immuniza-

tion, the number of PE-specific B cells were severely diminished by CLL treatment (Fig. 4A). PE-specific Ig (H+L) and IgG2c serum titers were also reduced (Fig. 4B). This demonstrated that SCM $\phi$  play a central role in the B cell expansion and Ab responses elicited by immunization with GLA-SE. Surprisingly, we observed no difference in the total numbers of T<sub>FH</sub> cells in the draining LNs between the mock-treated and CLL-treated mice in response to PE+GLA-SE immunization (Fig. 4C).

To determine whether Ag-specific T<sub>FH</sub> numbers are specifically affected by CLL depletion we immunized mock- or CLL-treated mice with the recombinant tuberculosis Ag ID87, which contains the Ag85B P25 epitope (23), and analyzed peptide-MHCII (pMHCII) tetramer staining. The number of Ag specific T<sub>FH</sub> cells (pMHCII<sup>+</sup>, CD44<sup>+</sup>, CXCR5<sup>+</sup>, PD1<sup>+</sup>, Bcl-6<sup>+</sup>) was virtually unaffected by CLL treatment. In contrast, significantly fewer Th1 cells (pMHCII<sup>+</sup>, CD44<sup>+</sup>, CXCR5<sup>-</sup>, PD1<sup>-</sup>, Tbet<sup>+</sup>) were present in LNs from CLL-treated mice compared with mock-treated mice (Fig. 5). To confirm that the observed altered adaptive responses in CLL-treated mice were due to the absence of SCM $\phi$  and not unexpected off-target effects of CLL treatment, we also examined responses in CD169-DTR mice (21, 22). Similar to CLL treatment, depletion of CD169<sup>+</sup> SCM $\phi$  via diphtheria treatment prior to immunization with GLA-SE reduced the B cell and Th1 responses but not the T<sub>FH</sub> responses to immunization (Supplemental Fig. 2). These data suggest that Ag-specific Th1 cell numbers, not T<sub>FH</sub> cell numbers, may be important for the early B cell response in GLA-SE immunized animals.



**FIGURE 3.** Innate responses to GLA-SE after SCM $\phi$  depletion. Mice were treated with clodronate-loaded liposomes (CLL) or saline (mock) 6 d prior to injection with GLA-SE or saline and draining LNs were harvested either 4 or 18 h after immunization. **(A)** The concentration of cytokines and chemokines 4 h after injection are plotted. **(B)** A representative histogram of CD69 staining on total lymphocytes 18 h after injection is shown and the percentage of CD69<sup>+</sup> lymphocytes is plotted. Data are representative of two independent experiments with three animals per group. Bars are drawn to the mean + SEM. The *p* values were determined by one-way ANOVA/Dunnett's test, \*\*\**p* < 0.001, \*\**p* < 0.01, \**p* < 0.05.

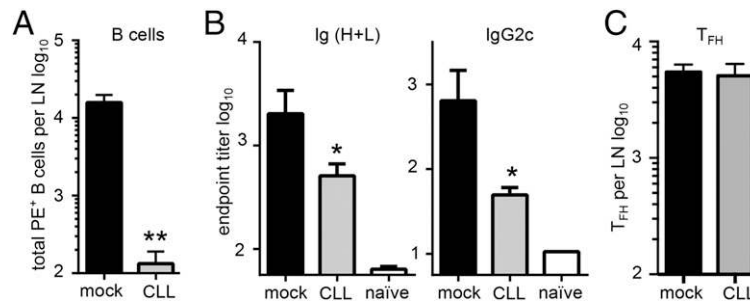
#### *IL-18R1* contributes to the induction of early B cell responses elicited by GLA-SE

Th1 induction with GLA-SE is dependent on the caspase-1 mediated inflammasome and rapid IL-18 production which is essential for IFN- $\gamma$  production by memory CD8 T cells and neutrophils upon GLA-SE immunization (18). SCM $\phi$  are an important source of IL-18 in draining LN (29). Thus, we hypothesized that SCM $\phi$  played a role in the innate and adaptive responses elicited by immunization with GLA-SE in part by producing IL-18. CLL depletion prior to GLA-SE immunization significantly diminished the amount of secreted IL-18 in the draining LN 1 h after immunization indicating that the SCM $\phi$  are the main source of IL-18 early after immunization (Fig. 6A). To determine whether IL-18 was important for the induction of Ag-specific B cells and Ab production, we compared responses to immunization in wild-type or IL18R1 deficient (IL-18R1<sup>-/-</sup>) mice. Compared with wild type, total Ig (H+L) and IgG2c PE-specific serum titers were diminished in IL-18R1<sup>-/-</sup> mice (Fig. 6B). Concordantly, in the immunized IL-18R1<sup>-/-</sup> mice significantly fewer PE-specific B cells were present (Fig. 6C) and IgG2c<sup>+</sup> class-switching was impaired (Fig. 6D). These data indicate that, similar to CLL or DTR depletion of SCM $\phi$ , ablation of IL-18 signaling impaired the humoral responses elicited by immunization with GLA-SE. Of note DTR- or CLL-depletion had a more dramatic impact on adaptive immune responses than IL-18R1 deficiency indicating SCM $\phi$  also control adjuvant-driven responses in an IL-18R1-independent manner.

#### Discussion

Rapid Ab production after immunization could mean the difference between protection and infection against a new strain of pathogen such as a pandemic avian influenza virus or biological weapons. As evident in the data presented here, different adjuvants can alter the kinetics and quality of the humoral immune response. Few studies have examined changes in Ag-specific B cell populations elicited by adjuvants, particularly those destined for clinical use. Of note, Ag-staining was recently used to identify germinal center B cell responses during a prime-boost regimen with the emulsion adjuvant MF59 (30). Overall, those results were consistent with our studies of the similar adjuvant SE. Alum, GLA alone or SE alone only weakly elicited early B cell responses — primarily germinal center (GC) and memory B cells — whereas the combination GLA-SE drove a rapid and robust B cell and Ab response that also included preplasmablasts important for early Ab secretion. We previously found that a rapid humoral response induced with GLA-SE correlated with early-onset protection against the highly pathogenic avian influenza virus H5N1 as early as 4–6 d after immunization, underlining the importance that such an adjuvant could have in a rapidly emerging epidemic (13).

After injection, GLA-SE was predominantly associated with the SIGNR1<sup>+</sup> medullary subset of SCM $\phi$ , which correlated with the ability to induce strong B cell responses. This is consistent with microscopic analysis of cells taking up DiD-labeled MF59 (31). Strikingly, when the SCM $\phi$  population was disrupted by CLL



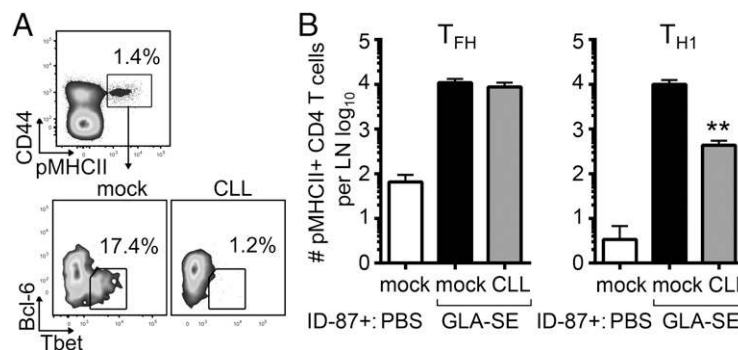
**FIGURE 4.** B cell and T<sub>FH</sub> induction via GLA-SE after SCM $\phi$  depletion. Mice were treated with clodronate-loaded liposomes (CLL) or saline (mock) 6 d prior to immunization with PE+GLA-SE and responses were analyzed 7 d after immunization. **(A)** Shown are the number of PE<sup>+</sup> B cells in draining inguinal lymph nodes, **(B)** Shown are the total Ig (H+L) and IgG2c endpoint titers. **(C)** The total numbers of T<sub>FH</sub> cells (CD4<sup>+</sup>, CD44<sup>+</sup>, CXCR5<sup>+</sup>, PD1<sup>+</sup>, Bcl-6<sup>+</sup>) in draining LNs were enumerated. Data are representative of two independent experiments with five animals per group. Bars are drawn to the mean + SEM. The *p* values were determined by one-way ANOVA/Dunnett's test, \*\**p* < 0.01, \**p* < 0.05.

injection, the number of B cells that were associated with labeled GLA-SE fell by nearly 10-fold, suggesting that SCM $\phi$  may be necessary to capture and transfer the GLA-SE adjuvant to B cells. In support of this possibility sinusoidal SCM $\phi$  facilitate transport of virus particles to B cell follicles (3). Presumably this process facilitates the induction of germinal center responses as the disruption of sinusoidal SCM $\phi$  lattice architecture inhibits the induction of germinal center responses (32). Using an i.p. immunization method Nikbakht and colleagues (33) found that CLL depletion of marginal-zone macrophages from the spleen also impaired B cell responses to alum adjuvanted immunization. In contrast, we do not detect substantial Ag-specific B cell responses with alum in our system, likely due to the different route of delivery (i.m. versus i.p.). Nevertheless, these findings largely support those of Nikbakht that CLL-dependent macrophages are important for the humoral response to adjuvanted immunization. Our studies do not discount the contribution of other cell types to the adjuvanticity of GLA-SE, however, it is intriguing to note that LN-resident macrophages but not migratory macrophages that traffic to the immunization site were crucial for B cell responses to a rabies virus vector immunization (34). The importance of these migratory macrophages in the adjuvanticity of GLA-SE remains to be determined.

GLA-SE also uniquely induced preplasmablasts, which have been shown to localize to medullary regions of the LN (35). This result suggested that initial priming could occur at sites distal to the medullary sinus, but it is conceivable that medullary macrophages may promote the activation of B cells and their differentiation into preplasmablasts in situ. Furthermore, Ab responses in

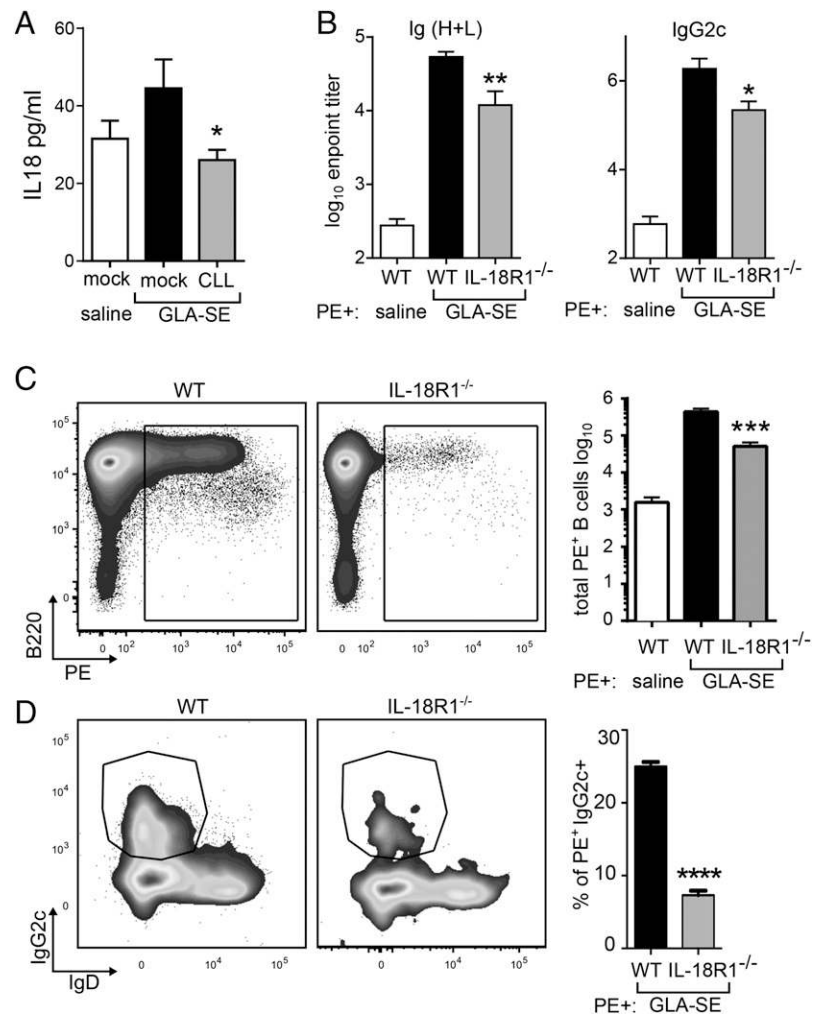
transiently dendritic cell (DC) depleted mice (CD11c-DTR) were mostly unaffected, suggesting that DC are not required for B cell priming (36). Thus we suggest that SCM $\phi$ , and not DC, are critical for adjuvant enhancement of B cell priming upon immunization, by creating a microenvironment conducive to B cell expansion and preplasmablast generation (e.g., increases in local cytokines such as IL-18, IFN- $\gamma$ , and others reduced by SCM $\phi$  ablation) and/or facilitating B cell-Ag interactions. It remains to be determined whether SCM $\phi$  or another cell population such as follicular dendritic cells or neutrophils are also critical for capturing and transporting unprocessed Ag to the B cells during the priming event or if the B cells are directly capturing the Ag (18, 37). In support of a direct role for Ag presentation to B cells by SCM $\phi$ , direct targeting of Ags to these cells by coupling OVA to anti-CD169 increased the OVA-specific germinal centers and humoral response (38).

SCM $\phi$  were also important for the production of IL-18 upon immunization. We previously demonstrated that IL-18 production mediated by caspase 1 activation was important for the production of IFN- $\gamma$  by memory CD8 T cells and neutrophils (18). This pathway was also important for the generation of Ag-specific CD4 T cells with GLA-SE. This IL-18 axis was also at least partially responsible for the SCM $\phi$ -dependent expansion of Ag-specific B cells. It remains to be determined whether IL-18 is acting directly on B cells to drive their expansion, or via IL-18 mediated production of IFN- $\gamma$  production. Another possibility is that the IL-18 axis is primarily affecting the ability of CD4 T cells to provide sufficient help to B cells for optimal expansion after immunization, as suggested by our previous findings that IL-18 is important



**FIGURE 5.** T<sub>FH</sub> and Th1 dependency upon SCM $\phi$ . Peptide-MHCII tetramer staining was used to identify T cell subsets in the context of CLL treatment. Mice were treated with clodronate-loaded liposomes (CLL) or saline (mock) 6 d prior to immunization with GLA-SE and the recombinant protein ID87 and responses were analyzed 7 d after immunization. **(A)** FACS plots of Tbet and Bcl-6 staining of CD4<sup>+</sup>pMHCII<sup>+</sup> cells from draining lymph nodes. **(B)** The total numbers of T<sub>FH</sub> (pMHCII<sup>+</sup>, CD4<sup>+</sup>, CD44<sup>+</sup>, CXCR5<sup>+</sup>, PD1<sup>+</sup>, Bcl-6<sup>+</sup>) and Th1 (pMHCII<sup>+</sup>, CD4<sup>+</sup>, CD44<sup>+</sup>, CXCR5<sup>+</sup>, PD1<sup>+</sup>, Tbet<sup>+</sup>) per LN were enumerated. Data are representative of two independent experiments with three or four animals per group. Bars are drawn to the mean + SEM. The *p* values were determined by one-way ANOVA/Dunnett's test, \*\**p* < 0.01.

**FIGURE 6.** IL-18 is important for GLA-SE augmentation of B cell responses. Mice were treated with clodronate-loaded liposomes (CLL) or saline (mock) prior to immunization. (A) Mice were immunized with GLA-SE and IL-18 production in the draining lymph node was assessed 1 h postinjection. (B–D) Wild-type or IL18R<sup>-/-</sup> mice were immunized with PE<sup>+/+</sup>GLA-SE and responses were analyzed 7 d after immunization. (B) Total Ig (H+L) and IgG2c PE reactive endpoint titers. (C) FACS plots of PE<sup>+</sup> staining are shown and the total numbers of PE-specific B cells in the draining LNs are plotted. (D) FACS plots of IgG2c and IgD staining of PE<sup>+</sup> B cells are shown and frequencies of IgG2c<sup>+</sup> PE<sup>+</sup> B cells are plotted. Data are representative of two independent experiments with four or five animals per group. Bars are drawn to the mean + SEM. The *p* values were determined by one-way ANOVA/Dunnett's test, \*\*\*\**p* < 0.0001, \*\*\**p* < 0.001, \*\**p* < 0.01, \**p* < 0.05.



for Th1 expansion upon immunization with GLA-SE (18). Our recent work with Tbet-deficient mice indicates that the IFN- $\gamma$  axis likely plays a role in humoral responses as loss of Th1 cells in the Tbet-deficient animals correlated with a decrease in Ab responses (19).

Substantial evidence has accumulated describing the relationship between T<sub>FH</sub> cells and the induction of B cell responses (4). Importantly, T<sub>FH</sub> and GC B cell numbers correlate during the induction of an immune response by vaccination (39). In agreement with these observations, GLA-SE drives an expansion of Ag-specific T<sub>FH</sub> cells concordant with the increase in Ag-specific B cells and this is superior in magnitude to the other adjuvants tested. However, the contribution of Ag-specific T<sub>FH</sub> cells to B cell expansion and early Ab titers during immunization with GLA-SE is unclear. Surprisingly Ag-specific T<sub>FH</sub> cells were unaffected by SCM $\phi$  depletion, whereas B cell and Th1 responses were dramatically impaired. Canonically Th1 cells are associated with CD8 T cell responses, not B cell responses, although Th1 cells can facilitate B cell priming in some circumstances (40). This underlines the current models suggesting that separate subsets of Ag presenting cells mediate the induction of Th1 cells and T<sub>FH</sub> cells (41, 42). In support of this we have recently shown that Tbet deficient mice, which have no Th1 responses to GLA-SE, do not induce an IgG2c skewed response, further supporting a role for Th1 cells in B cell responses (20). Although this does not demonstrate that T<sub>FH</sub> cells are irrelevant to the early B cell response, it does indicate that T<sub>FH</sub> cells are insufficient to support all B cell responses.

The unique adjuvant properties of GLA-SE to expand B cell responses, relative to GLA alone or SE alone, are likely due both to its biophysical characteristics, as well as molecular signaling capacities. For example, the innate response to GLA-SE depends not only upon MyD88 and TRIF, which are critical components of TLR4 signaling, but also upon the inflammasome (18, 19), which is only engaged by TLR4 agonists upon cytoplasmic recognition (43, 44). On their own, oil-in-water emulsions such as MF59 or SE promote humoral immunity via activation of canonical and non-canonical inflammasome pathways (45–48). Inflammasome activation could be due to direct recognition of the squalene via distinct, but unknown, receptors. GLA and SE in turn likely collaborate to produce bioactive IL-18 in a two-step process in which TLR4 signaling triggered by GLA increases production of pro-IL-18 and SE engagement of the inflammasome generates active caspase-1 to cleave pro-IL-18 into bioactive IL-18. Additionally, the nanoparticle nature of GLA-SE may result in more efficient delivery to the LN. Nevertheless, the particulate nature cannot fully explain the enhancement mediated by GLA-SE as other oil-in-water/TLR4 agonist compositions, with similar biophysical properties, do not recapitulate the adjuvanticity of GLA-SE (7, 49).

In summary, when formulated as GLA-SE the TLR4 agonist GLA induces rapid B cell expansion and differentiation, augmented Ab production, and T<sub>FH</sub> expansion. These responses were superior in magnitude to the widely used adjuvant alum, as well as GLA in an aqueous base or the SE alone adjuvant. SCM $\phi$  and IL-18 production were central to optimal immunization with GLA-SE as



the innate immune response, B cell expansion, and Th1 induction were impaired upon SCM $\phi$  depletion or abrogation of IL-18R signaling. Interestingly, T<sub>FH</sub> induction after GLA-SE immunization was not affected in the absence of SCM $\phi$ . Thus, GLA-SE may be ideally suited for use in vaccines that require a rapid-onset response with a single immunization.

Our results point to early IL-18 production as a candidate biomarker of immunogenicity for immunization with GLA-SE adjuvanted vaccines, and may indicate that early IL-18 production could be a hallmark of different classes of vaccine adjuvants. Additionally, the role of SCM $\phi$  in the mechanism of action of other adjuvants will help determine whether this subpopulation needs to be targeted by all adjuvants through the activation of TLR or other PRRs expressed by SCM $\phi$  or by receptor-specific delivery mechanisms such as Ag complexed to SIGNR1, similar to the approaches targeting Ags to DCs via anti-DEC205 (38). Defining the key pathways that determine the activity of clinical stage adjuvants including GLA-SE, MF59, and the AS01-04 series is crucial to rational development of next-generation vaccine adjuvants tailored to elicit specific immune responses.

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## Disclosures

The authors have no financial conflicts of interest.

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