

Monocyte-derived Dendritic Cells Perform Hemophagocytosis to Fine-tune Excessive Immune Responses

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SUMMARY

Since immune responses simultaneously defend and injure the host, the immune system must be finely regulated to insure the host's survival. Here, we show that when injected with high TLR ligand doses or infected with LCMV clone 13, which has a high viral turnover, inflammatory monocyte-derived dendritic cells (Mo-DCs) are induced to engulf apoptotic erythroid cells. In this process, called hemophagocytosis, extracellular ATP and phosphatidylserine (PS) serve as “find-me” and “eat-me” signals, respectively. Type I IFNs are necessary for both PS exposure on erythroid cells and the expression of PS receptors in the Mo-DCs. Importantly, hemophagocytosis is required for IL-10 production from Mo-DCs. Blocking hemophagocytosis or Mo-DC-derived IL-10 significantly increases CTL activity, tissue damage, and mortality in virus-infected hosts, suggesting that hemophagocytosis moderates immune responses to assure the host’s survival *in vivo*. This sheds light on the induction mechanisms and physiological relevance of hemophagocytosis in severe inflammatory and infectious diseases.

INTRODUCTION

An immune response is a two-edged sword that, while directly attacking a pathogen, simultaneously damages the body (Schneider and Ayres,2008). In an infected host, DCs and other innate immune cells sense pathogens through receptors that detect pathogen-associated molecular patterns (PAMPs) or damage-associated molecular patterns (DAMPs). These danger signals trigger DC maturation and induce immunogenic DCs that produce inflammatory cytokines and activate both the innate and adaptive immune systems. The adaptive immune system eliminates the pathogen in an Ag-specific manner and protects the host from re-infection (Medzhitov and Janeway,

1997; Banchereau et al., 2000; Cooper and Alder, 2006). However, infection-induced immune responses, particularly virus-specific CTLs, cytokines, and chemical mediators, simultaneously damage tissue (Clark, 2007; Lambth, 2007; Rouse and Sehrawa, 2010). Therefore, immune responses require regulatory machinery to maintain the fine balance between defense and self-injury during infections; the more severe the infection, the greater the regulatory control must be. However, little attention has been paid to the nature of the machinery balancing these immune-response effects. This study focuses on hemophagocytosis, which is conventionally viewed as an indicator of severe inflammation, and newly identifies DC-mediated hemophagocytosis as the machinery that fine-tunes immune responses and assures host survival under severe inflammatory and infectious conditions.

RESULTS

TLR Ligands Induce Hemophagocytosis

Pattern recognition receptors (PRRs), including Toll-like receptors (TLRs) and NOD-like receptors (NLRs), sense microbial infection and mediate immune and inflammatory responses (Kawai and Akira, 2010). In this study, we found that high doses (~200 µg) of various TLR and NLR ligands injected into wild-type (WT) mice induced hemophagocytosis, an indicator of severe inflammatory conditions such as hemophagocytic syndrome (Larroche and Mouthon, 2003; Janka, 2007; Maakaroun et al., 2010). Hemophagocytosis, typically defined as the engulfment of TER119⁺ erythroid cells by CD11c⁺ cells, was detected by the presence of CD11c⁺TER119⁺ cells without permeabilization in the peripheral blood (PB). Of the ligands used, unmethylated CpG DNA (CpG) and poly I:C induced hemophagocytosis in the

peripheral blood (PB) most efficiently (**Figure 1A**). After injecting CpG, the number of CD11c⁺TER119⁺ cells in the PB increased according to the CpG dosage, peaking 18 h later in the bone marrow (BM) and PB and 3 days afterwards in the spleen (**Figures S1A-S1D**). This suggests that hemophagocytosis in the PB correlated well with and reflected that in the BM. Sorted CD11c⁺TER119⁺ cells displayed typical hemophagocytosis with both attached and internalized erythroid cells by Diff-Quick stain and electron microscopic analysis (**Figures 1B and 1C**). CpG also induced other criteria of hemophagocytic syndrome, including fever, splenomegaly, cytopenia (evaluated by the reduction of hemoglobin and platelets), and hypertriglyceridemia (**Figures S1E-S1I**). We used a recently reported protocol (Zoller et al., 2011) to further confirm the intracellular localization of erythroid cells inside CD11c⁺ cells. In brief, TER119 exposed on CD11c⁺ cell surfaces was blocked with saturating amounts of an unlabeled Ab against TER119, and then stained with an anti-TER119 Ab. As expected, no TER119⁺ cells were detected without permeabilization (**Figure S2A**), while intracellular TER119 was stained after permeabilization in a CpG dose-dependent manner (**Figures S2B and S2C**). These results clearly demonstrated the intracellular localization of erythroid cells and correlated well with those of CD11c⁺TER119⁺ staining without permeabilization (**Figures S1D and S2C**). This is reasonable, since hemophagocytosis is a sequential event: apoptotic erythroid cells expressing phosphatidylserine (PS) are initially attached to PS receptors on the CD11c⁺ cells, and then are gradually phagocytosed into the cells.

Hemophagocytosing DCs Are of Monocyte Origin

The phenotype of the hemophagocytosing CD11c⁺ cells was MHC class II^{low}CD11b^{high}F4/80⁺Gr-1^{high}CD40⁻CD80^{high}CD86⁻ (**Figure 1D**). This is characteristic of inflammatory monocyte-derived DCs (Mo-DCs), which are induced under inflammatory conditions (Randolph et al., 1999; Leon et al., 2007; Serbina et al., 2008; Auffray et al., 2009; Geissmann et al., 2010). To confirm the Mo-DCs' origin, we sorted inflammatory monocytes, defined as Ly6c^{high}CX₃CR1^{int}CD11b⁺CD11c⁻ cells, from the BM of *Cx₃cr1*^{GFP/+} mice (CD45.1⁻CD45.2⁺), which express EGFP under the endogenous *Cx₃cr1* locus (Jung et al., 2000) (**Figure S3A**). We injected these cells into WT mice (CD45.1⁺CD45.2⁻) and then injected CpG. After 18 h, the transferred inflammatory monocytes had become CD11c⁺, and approximately 40% were engulfing TER119⁺ cells (**Figures 1E and 1F**). In *Cx₃cr1*^{GFP/+} mice injected directly with CpG, the hemophagocytes consisted mostly of Ly6c^{high}CX₃CR1^{int} inflammatory monocyte-derived DCs (**Figure S3B**), indicating that inflammatory monocytes were the major hemophagocyte precursor. During infection and inflammation, inflammatory monocytes mobilize from the BM to the periphery CCR2-dependently (Boring et al., 1997; Serbina et al., 2006). Since CpG-induced hemophagocytosis is mediated by Mo-DCs, we conducted experiments with *Ccr2*^{-/-} mice, in which circulating monocytes are able to leave the BM but inflammatory monocytes cannot. In WT mice, serum levels of the CCR2 ligands CCL2, CCL7, and CCL12 peaked 3-6 h after CpG injection (**Figures 2A-2C**); CCR2 was predominantly expressed on Ly6c^{high} inflammatory monocytes and to a lesser extent on CD11c⁺TER119⁺ cells (**Figure 2D**). Importantly, the percentage of PB CD11c⁺TER119⁺ cells after CpG injection was greatly reduced in the *Ccr2*^{-/-} mice (Boring et al., 1997), partially reduced in *Ccl2*^{-/-} mice (Lu et al., 1998), probably due to CCL7 and CCL12, which are also CCR2 ligands, and unaffected in *Cx₃cr1*^{GFP/GFP} mice (**Figure 2E**). In contrast, the BM CD11c⁺TER119⁺ cell levels after

CpG injection were comparable between the WT and *Ccr2*^{-/-} mice (**Figure 2F**). Collectively, these results demonstrate that CpG injection caused inflammatory monocytes to mobilize to the periphery, differentiate into Mo-DCs, and engulf mainly erythroid cells.

Involvement of “Find Me” and “Eat Me” Signals in TLR Ligand-Induced Hemophagocytosis

Cells undergoing apoptosis release “find me” signals to recruit phagocytes, and expose PS, which is recognized by phagocytes as an “eat me” signal, on their surface (Nagata et al., 2010; Martin et al., 1996). PS is recognized directly by Tim-1 and -4 and indirectly by the $\alpha_v\beta_3$ and $\alpha_v\beta_5$ integrins through milk fat globule EGF factor 8 (MFG-E8), and by TAM family members through Gas6 and protein S (Martin et al., 1996; Hanayama et al., 2002; Kobayashi et al., 2007; Miyanishi et al., 2007; Nakayama et al., 2009). We next examined the roles of “find me” and “eat me” signals in CpG-induced hemophagocytosis. Serum levels of ATP, recently identified as a “find me” signal (Elliott et al., 2009), increased rapidly after CpG injection, peaking 6 h later (**Figure 3A**). The ATP receptor P2Y2 was predominantly expressed on Ly6c^{high} inflammatory monocytes, on BMDCs, and on Mo-DCs performing hemophagocytosis (CD11c⁺TER119⁺), but not on T or B cells (**Figure 3B**), suggesting that ATP is an important “find me” signal in hemophagocytosis. Indeed, injecting suramin, a competitive inhibitor of ATP-P2Y2 binding, effectively blocked the CpG-induced hemophagocytosis (**Figure 3C**).

Next, we examined the role of the “eat me” signal in CpG-induced hemophagocytosis. The number of apoptotic erythroid cells expressing PS, defined as annexin V⁺TER119⁺

cells, peaked 2-4 h after CpG injection (**Figure 3D**). In this context, most of the apoptotic cells were TER119⁺ erythroid cells, and some apoptotic Gr-1^{high}Ly6c⁺ granulocytes were also detected (**Figures S4A-S4C**). CpG-induced Mo-DCs were the major phagocytes for TER119⁺ erythroid cells (**Figures S5A-S5D**). As “eat me” signal receptors, the Mo-DCs expressed Tim1, Tim4, and α_v and β_3 integrin (**Figure 3E**) and an injection of neutralizing Abs against “eat me” signal receptors significantly blocked hemophagocytosis (**Figure 3F**). CpG injection induced the production of the representative regulatory cytokine IL-10 in the PB at rates that increased with CpG dosage (**Figure 3G**), and this correlated well with hemophagocytosis (**Figures S1D and S2C**). Consistent with these results, excessive CpG stimulation triggers DC IL-10 production (Waibler et al., 2008). Of note, injecting PS receptor-neutralizing Abs, which prevents hemophagocytosis, significantly reduced the IL-10 production (**Figures 3G**). Collectively, these results suggest that “find me” and “eat me” signals are critical for CpG-induced hemophagocytosis, and that high CpG doses induce IL-10 production.

Injecting a high dose of poly I:C into WT mice produced similar results, with Mo-DCs (**Figure S6A**) and “find me” and “eat me” signals (**Figures S6B and S6C**) playing critical roles in the poly I:C-induced hemophagocytosis and accompanying IL-10 production (**Figures S6D and S6E**).

Type I IFN Signaling Is Essential in Virus-Induced Hemophagocytosis

Based on these findings, we examined the induction mechanism and physiological relevance of hemophagocytosis in virus infection. We used the lymphocytic choriomeningitis virus (LCMV) variant clone 13 (C13), which elicits a chronic infection (Matloubian et al., 1990), for two reasons. First, compared with the original LCMV

Armstrong (Arm), C13 binds and infects DCs with high affinity, and it proliferates vigorously owing to two amino acid changes in its glycoprotein and polymerase, resulting in remarkably high viral nucleic acid levels *in vivo* (Sevilla et al., 2010), possibly mimicking the effect of high-dose CpG or poly I:C injection. In this context, we expected that the more severe the infection, the more closely regulated the response. Second, most viruses that induce hemophagocytosis produce a chronic infection in humans (Clark 2007; Lambth 2007; Rouse and Sehrawa, 2010), although the role of hemophagocytosis in chronic infection is unknown. Indeed, the mRNA levels for viral glycoprotein (GP) and nucleoprotein (NP) in C13-infected cells were remarkably higher than in Arm-infected cells, and we newly found that inflammatory monocytes, which are Mo-DC precursors, were a major C13 infection target (**Figure S7A**). As expected from these results, hemophagocytosis was induced more efficiently in C13-infected WT mice than in Arm-infected WT mice (**Figures S7C and S7D**); the intracellular localization of erythroid cells was again confirmed using the Diff-Quick staining and recently reported protocol (Zoller et al., 2011) (**Figures S7B, S7E, and S7F**). As with CpG injection, most of the apoptotic cells were TER119⁺ erythroid cells and some were apoptotic Gr-1^{high}Ly6c⁺ granulocytes, suggesting that the contribution of apoptotic granulocytes was not negligible but relatively low (**Figures S8A-S8C**). Indeed, we sometimes observed the phagocytosis of granulocytes by Mo-DCs isolated from the PB of C13-infected WT mice (**Figure S8D**). In contrast, lymphocyte populations, including T, B, and NK cells, rarely became apoptotic at the same time point after C13 infection.

C13 infection induced type I IFN production (**Figures 4A and 4B**). Interestingly, C13 infection-induced hemophagocytosis was significantly reduced in mice deficient for IFNAR1, a component of the type I IFN-receptor (**Figures 4C and 4D**). Consistent with these results, C13 infection in *Ifnar1*^{-/-} mice severely impaired the PS exposure on

erythroid cells and the Tim-1, Tim-4, and α_V and β_3 integrin expression on Mo-DCs (**Figures 4E-4G**), suggesting that type I IFNs are essential both for inducing PS on erythroid cells and for the development of Mo-DCs that express PS receptors. Indeed, *ex vivo* IFN- α stimulation directly induced the apoptosis of TER119⁺ erythroid cells and, to a lesser extent, Gr-1^{high}Ly6c⁺ granulocytes (**Figure S9A**). In addition, IFN- α stimulation directly induced expression of PS receptors, i.e. Tim-1, Tim-4, and α_V and β_3 integrin, on inflammatory monocytes, precursors of Mo-DCs (**Figures S9B and S9C**).

Hemophagocytes are the major source of IL-10

Finally, we examined the physiological relevance of hemophagocytosis in C13 infection. As expected from our findings thus far, injecting neutralizing Abs against Tim-1, Tim-4, α_V , and β_3 into C13-infected WT mice significantly suppressed hemophagocytosis (**Figure 5A**). Milk fat globule-epidermal growth factor 8 (MFG-E8) contains a PS-binding domain and an arginine-glycine-aspartate (RGD) motif, which binds to both $\alpha_V\beta_3$ and $\alpha_V\beta_5$ integrin on phagocytic cells, and it mediates the integrin-dependent engulfment of apoptotic cells and subsequent signaling (Hanayama et al., 2002; Akakura et al., 2003). As expected, *Mfge8* was clearly expressed in both inflammatory monocytes, a precursor of Mo-DCs, and Mo-DCs performing hemophagocytosis, whereas expression of other PS receptors including *Bai1*, *Stab2*, *Tyro3*, *Axl*, *Mertk*, *Gas6* detected by qPCR (**Figure S10A**) and Stabilin-2, Tyro-3, Axl, MerTK by specific Ab staining (**Figure S10B**) was minimal, suggesting that these molecules are unlikely involved in the uptake of apoptotic erythroid cells by Mo-DCs. Serum levels of IL-10 and TGF- β 1, both of which suppress immune responses, increased after C13 infection,

peaking at 24 h (**Figure 5B** and data not shown). In line with our findings, C13 infection induces IL-10, contributing to the virus persistence *in vivo* (Brooks et al., 2006), although the mechanism of IL-10 production remains unknown. Importantly, injecting Abs against the Mo-DC PS receptor, which block hemophagocytosis, effectively reduced the IL-10 and TGF- β 1 production (**Figure 5B**), whereas injecting IL-10-neutralizing Abs into WT mice did not suppress hemophagocytosis (**Figure 5A**). This suggests that the IL-10 production from Mo-DCs in response to C13 infection is hemophagocytosis-dependent. Of note, ERK and p38 phosphorylations, which are prerequisite for IL-10 production, were detected in hemophagocytosing Mo-DCs (**Figures S11A and S11B**). In this context, blocking IL-10 activity substantially reduced the TGF- β 1 serum levels, implying that IL-10 somehow regulates TGF- β 1 production in C13-infected WT mice (**Figure 5B**). In the PB of C13-infected *Il10*^{Venus} reporter mice (Atarashi et al., 2011), IL-10 was expressed predominantly in CD11c⁺TER119⁺ cells and not in T cells (**Figure 5C**), and the absolute number of IL-10^{Venus+} CD11c⁺TER119⁺ cells was much greater than that of IL-10^{Venus+} cells belonging to other populations, including CD11c⁺TER119⁻ cells (**Figure 5D**). This clearly demonstrates that hemophagocytes are the major source of IL-10 at an early phase of infection *in vivo*.

To further examine the IL-10 production machinery from Mo-DCs in response to C13 infection, we newly established *ex vivo* hemophagocytosis assay system, in which TER119⁻ Mo-DCs and TER119⁺ erythroid cells, separately sorted from the BM of C13 infected WT mice, were co-cultured (**Figure 6A**). Of note, the erythroid cells were pHrodo-labelled prior to the co-culture, which allowed the erythroid cells transported into lysosomes upon hemophagocytosis to be detected by their increased light emission under the acidic conditions. Consistent with the *in vivo* results, hemophagocytosis was clearly induced *ex vivo* (**Figures 6A, left panel, 6B, and 6C**). Most importantly, *ex vivo*

blockage of hemophagocytosis with Abs against the Mo-DC's PS receptor impaired Mo-DC-derived IL-10 production (**Figures 6A, right panel, 6D, and 6E**). In this context, adding blocking Abs against Tim-1 and Tim-4 did not affect IL-10 production, whereas blocking Abs against integrin α_v , β_3 , or $\alpha_v\beta_5$ significantly reduced the level of IL-10 production (**Figure 6E**). In addition, neither blocking internalization of apoptotic erythroid cells with cytochalasin D nor opsonization of them induced Mo-DC-derived IL-10 production (**Figure 6E**). These results collectively demonstrated that IL-10 production from Mo-DCs is largely dependent on the integrin-mediated signals. Similar results were obtained from the co-culture in which Mo-DCs and granulocytes, the other apoptotic population in C13-infected mice (**Figure S8A and S8B**), were used in the same experiment (**Figure S12**). Notably, the amounts of IL-10 secreted from the Mo-DCs that engulfed erythroid cells were 60 pg/ml, and from those that engulfed granulocytes were 40 pg/ml in the same experiment (**Figure 6E and S12**). These results suggested that granulocytes are slightly less potent on a per cell basis than erythroid cells for inducing IL-10 secretion from Mo-DCs. To definitively show the contribution of these integrins to the IL-10 secretion from Mo-DCs, we performed *ex vivo* hemophagocytosis assay, in which Mo-DCs and erythroid cells, sorted separately from the BM of WT and *Mfge8*^{-/-} mice after C13 infection, were co-cultured in all combinations (**Figure 6F**). Importantly, the IL-10 secretion was selectively impaired when *Mfge8*^{-/-} Mo-DCs were used for the co-culture, demonstrating that the IL-10 secretion from Mo-DCs is MFG-E8-mediated, and thus dependent on both integrin $\alpha_v\beta_3$ and $\alpha_v\beta_5$ (**Figure 6F**).

We also examined the apoptotic status of these cells during later time points, i.e. 8, 16, 24 days after the infection. The numbers of apoptotic erythroid cells were much reduced and comparable to those of granulocytes (**Figure S13A**), implying the equal

contribution of erythroid cells and granulocytes to phagocytosis-mediated IL-10 production, although the numbers of IL-10 producing cells were substantially lower, compared with that at an early phase of infection (**Figure S13B**). As observed at an early phase of infection, lymphocyte populations, i.e. T and B cells, hardly became apoptotic during later time points (**Figure S13A**).

Hemophagocytosis Prevents Immune Responses-Mediated Damage and Assure the Host's Survival

What is the physiological relevance of the hemophagocyte-derived IL-10 in viral infection? Interestingly, blocking hemophagocytosis with PS receptor Abs or IL-10 activity with IL-10 neutralizing Ab enhanced the virus-specific CTL induction (**Figures S14A and S14B**), CTL-mediated liver-damage (Zinkernagel et al., 1986) (serum AST and ALT) (**Figures S14C, and S15A-S15D**), and mortality (**Figure S14D**). In the context of CTL activity, we also found that blocking hemophagocytosis or IL-10 activity diminished the CTL expression of PD-1 (**Figure S14E**), which causes CTL exhaustion in C13 infection (Barber et al., 2006). As Tim-1/Tim-4-signaling may alter immune responses in some cases (Xiao et al., 2007, 2011; Rodriguez-Manzanet et al., 2008), we examined whether anti-Tim-1 and -4 Abs could modulate immune responses in C13-infected WT mice. However, injecting anti-Tim-1 and -4 Abs did not alter the frequency of hemophagocytosis in the PB, IL-10 level and C13 virus titers in the sera, and percentage and absolute number of IL-10^{venus+} CD11c⁺TER119⁺ cells (**Figures S16A-S16E**), suggesting that these Abs do not significantly affect virus-specific immune responses *in vivo*, at least in this experimental setting.

To further confirm the importance of the hemophagocyte-derived IL-10 in fine-tuning excessive immune responses *in vivo*, *Cd11c-Cre/Il10^{fl/fl}* (CKO) mice, in which the hemophagocytes cannot produce IL-10, were infected with C13. While the hemophagocytosis levels in C13-infected CKO mice were comparable to those seen in control WT mice (**Figure 7A**), the serum IL-10 production was greatly reduced (**Figure 7B**), confirming that hemophagocytes are the major source of IL-10 *in vivo* under these infectious conditions. As with WT mice in which hemophagocytosis was blocked, the CKO mice showed excessive CTL activity, tissue damage, and mortality when infected with C13 (**Figures 7C-F**). Taken together, these results point to hemophagocytosis as a novel machinery for preventing excessive immune response-mediated damage, thus assuring the host's survival in the face of severe viral infection.

DISCUSSION

Our findings indicate that hemophagocytosis induced by TLR ligands or viruses occurs in sequential steps (**Figure S17**). We found that hemophagocytosis could be detected by staining with and without permeabilization. In this context, we would like to stress that hemophagocytosis is a sequential event; that is, apoptotic erythroid cells exposing PS are initially attached to PS receptors expressed on the Mo-DCs, and are then gradually phagocytosed into the cells. Therefore, even without permeabilization, these cells can be detected as CD11c⁺TER119⁺ cells. Supporting the specific recognition of apoptotic erythroid cells by PS receptors on Mo-DCs, hemophagocytosis detected without permeabilization was significantly reduced in WT mice in which hemophagocytosis was blocked with anti-PS receptor Abs or with suramin, a competitive inhibitor of ATP-P2Y2 binding. In C13-infected *Ifnar1^{-/-}* mice, both PS exposure on erythroid cells

and PS receptor induction on Mo-DCs were impaired, suggesting that type I IFNs are critically involved in these processes. As it is well known that CpG injection induces type I IFN production, it is likely that CpG-induced type I IFNs induce apoptosis of erythroid cells. Indeed, *ex vivo* IFN- α stimulation directly induced apoptosis of erythroid cells (**Figure S9A**). F4/80⁺ macrophages are reported to be the hemophagocytosis-performing cells in the spleen of mice infused with IFN- γ (Zoller et al., 2011). In this context, we found that the hemophagocytosis-performing Mo-DCs clearly expressed F4/80 (**Figure 1D**), suggesting that the previously reported F4/80⁺ macrophages and the Mo-DCs are presumably overlapping populations (Domínguez and Ardavín, 2010).

The uptake of experimentally (UV-irradiation, staurosporine, etc.) induced apoptotic peripheral blood lymphocytes, neutrophils, and peripheral blood T cells or Jurkat T cells by phagocytes induces IL-10 production, which suppresses proinflammatory cytokine production (Voll et al., 1997; Byrne and Reen, 2002; Kim et al., 2004). However, it remains unknown whether such *ex vivo* observations are consistent with and represent the *in vivo* situation under severe inflammatory conditions. Remarkably, we showed that, upon CpG injection or LCMV C13 infection, most of the apoptotic cells were erythroid cells, and a small number were granulocytes (**Figure S4 and S8**). As a result, Mo-DCs preferentially engulfed apoptotic erythroid cells, leading to IL-10 production. Given these findings, we do not exclude the contribution of granulocytes, although it is relatively minor, to the IL-10 secretion from Mo-DCs. Consistent with our findings, the type I IFN-dependent apoptosis of erythroid cells, but not of neutrophils or monocytes, is observed in DNase II deficient mice (Kawane et al., 2001; Yoshida et al., 2005). These results suggested that type I IFN signaling is required for the recruitment of nucleated erythroid cells to the PB and that, compared with other hematopoietic cells,

erythroid cells are relatively sensitive to type I IFN signaling and therefore become apoptotic. Indeed, IFN- α induces apoptosis in human erythroid cells (Tarumi et al., 1995). Notably, these phenotypes observed in DNase II deficient mice are consistent with those in WT mice injected with CpG or infected with C13.

Although hemophagocytosis is viewed as an indicator of hemophagocytic syndrome and other severe inflammatory conditions (Clark 2007; Lambth 2007; Rouse and Sehrawa, 2010), our findings suggest that it rather represents a possible biomarker of anti-inflammatory responses in the host. Very recently, it was reported that repeatedly injecting CpG induces hemophagocytic syndrome-like or macrophage activation syndrome-like disease in mice (Behrens et al., 2011). When IL-10 signaling is blocked in this model, a more severe disease with larger numbers of hemophagocytosis develops, raising two possibilities; first, hemophagocytosis contributes to the severity of the disease, and second, it is induced, as a negative feed-back machinery, to suppress severe inflammatory responses. We propose here that hemophagocytosis is prerequisite for IL-10 production, and that IL-10 is secreted by Mo-DCs in a manner dependent on hemophagocytosis, leading to anti-inflammatory effects of IL-10 to fine-tune immune responses. In humans, hemoglobin uptake by CD163⁺ macrophages induces the expression of the anti-inflammatory heme oxygenase (HO-1) *ex vivo* (Schaer et al., 2006), and HO-1 expression is detected in hemophagocytic CD163⁺ macrophages in the BM of patients with fatal sepsis (Schaer et al., 2006). Interestingly, CD163 expression is inducible by IL-10, implying that this anti-inflammatory pathway might also contribute to the prevention of immune responses *in vivo*.

In acute infections, pathogen growth correlates well with indicators of deteriorating health, and inducing immune responses usually restores health by reducing the pathogen

load. However, our findings show that, during severe viral infections, immune responses unaccompanied by efficient hemophagocytosis-mediated fine-tuning lead to a further deterioration in health and eventually to severe immune response-mediated self-damage, despite the virus being effectively eradicated. Of interest, aberrant immune activation with impaired IFN- α expression, a possible cause of limited hemophagocytosis, was evident in macaques infected with the 1918 Spanish pandemic virus (Kobasa et al., 2007).

The more severe the infection, the stronger the immune response must be. The stronger the immune response, the greater the regulatory control must be. Given that most viruses that cause hemophagocytosis, including Epstein-Barr virus, cytomegalovirus, and HIV (Larroche and Mouthon, 2003; Janka, 2007; Maakaroun et al., 2010), establish chronic infections in humans (Clerici et al., 1996; Nordoy et al., 2000; Budiani et al., 2002), the induction of hemophagocytosis may allow the virus to persist in the host, assuring host survival by preventing immune response-mediated damage. Considering this novel trade-off between host and pathogen, our findings may be valuable not only in extending our understanding of immune biology, but also in identifying therapeutic targets for severe inflammatory diseases.

EXPERIMENTAL PROCEDURES

Mice

C57BL/6J (B6, Clea), B6.*Cx3cr1*^{GFP/+} (Jackson) (Jung et al., 2000), B6.*Ccr2*^{-/-} (Jackson) (Boring et al., 1997), B6.*Ccl2*^{-/-} (Lu et al., 1998), B6.*Ifnar1*^{-/-} (B&K Universal), B6.*Il-10*^{Venus} (Atarashi et al., 2011), B6.*Cd11c*-Cre (Jackson) (Strabges et al., 2007), B6.*Mfge8*^{-/-} (Hanayama et al., 2004), and B6.*Il10*^{fl/fl} (Roers et al., 2004) mice were maintained in our SPF facility. All animal experiments were approved by the Institutional Animal Care Committees of Akita University and Tokyo Medical and Dental University.

Reagents and *In Vivo* Injection

LPS, MDP, and cytochalasin D were purchased from Sigma-Aldrich. Other reagents were obtained as follows: poly I:C (GE Healthcare), R848 (Invivogen), CpG (ODN-1668 [5'-TCCATGACGTTTCCTGATGCT-3']; Hokkaido System Science), iE-DAP (Invivogen), and recombinant mouse interferon- α (Dainion Sumitomo Pharma Co. Ltd.). Mice were intravenously (i.v.) injected with TLR ligands (200 μ g poly I:C, 100 μ g LPS, 100 μ g R848, or 200 μ g CpG) or Nod ligands (100 μ g iE-DAP or 100 μ g MDP). Mice were infected by i.v. injection of viruses (2 x 10⁶ pfu LCMV Armstrong or Clone 13).

Diff-Quick Stain

CD11c⁺TER119⁺ cells were sorted from the BM of CpG injected or LCMV C13 infected mice, cytopun at 500 g for 5 minutes onto a slide glass, and stained with Diff-Quick (Sysmex).

Electron Microscopy

CD11c⁺TER119⁺ cells were sorted from the PB of CpG injected mice, fixed with 2.5 % glutaraldehyde in 0.1 M phosphate buffer pH 7.2, washed overnight at 4 °C in the same buffer, and postfixed with 1% OsO₄ at 4 °C for 2 h. Then, the cells were dehydrated in a graded in a series of ethanol and embedded in Epon 812. Ultrathin (1µm) sections were collected on copper grids and with uranyl acetate and lead citrate and then examined by transmission electron microscope, H-7100 (Hitachi).

Hemophagocytosis Blocking

To block ATP activity, 6 mg suramin was intraperitoneally (i.p.) injected into WT mice 30 min after CpG or poly I:C injection, or 3 and 6 days after C13 infection. To block “eat me” signal receptors, WT mice received either 500 µg rat IgG (control) or 500 µg anti-Tim-1 mAb (RMT1-10) (Nakayama et al, 2009), 500 µg anti-Tim-4 mAb (RMT4-54) (Nakayama et al, 2009), 500 µg anti-α_v mAb (RMV-7) (Takahashi et al, 1990), or 500 µg anti-β₃ mAb (HMβ3-1) (Yashuda et al., 1995), or isotype control Ab cocktails including rat IgG2a, rat IgG2c, and hamster IgG (500 µg of each) by i.v., singly or in combination as indicated, 60 min and 120 min after CpG injection or viral infection. To block cytokines, 500 µg rat IgG (control) or 500 µg anti-IL-10 (JES5-2A5) was injected i.p. into WT mice 1 day and 3 days after C13 infection.

ELISAs

Cytokine and chemokine serum levels were determined by ELISA kits according to the manufacturer’s instructions (CCL2, IL-10, and TGF-β1, R&D; CCL7 and CCL12, Bender Med Systems). ATP and AST/ALT serum levels were determined by a luciferin-luciferase assay using an ATP assay kit (Toyo Ink) and a Fuji DRI-CHEM 3500 V analyzer (Fuji Medical Systems), respectively, according to the manufacturer’s instructions.

Flow Cytometric Analysis and Cell Sorting

Antibodies against the following molecules were used for flow cytometric analysis and cell sorting: CD3 ϵ (145-2C11), CD11b (M1/70), CD11c (N418), TER119 (TER119), CD19 (MB19-1), CD40 (1C10), CD80 (16-10A1), CD86 (GL19), NK1.1 (PK136), MHC class II (I-A/I-E; M-15/114.15.2), α_v (RMV-7), β_3 (2C9.G3), PD-1 (RMP1-30) (all from eBioscience), F4/80 (BM8) (Biolegend), and Ly6c (AL-219) (BD), Stabilin-2 (#34-2) (MBL), Tryo-3 (109646), Axl (175128) and MerTK (108921) (all from R&D). Antibodies against Tim-1 (RMT1-17), Tim-3 (RMT3-23), and Tim-4 (RMT4-53) were as described previously (Nakayama et al, 2009). To analyze hemophagocytosis, two protocols were used. First, peripheral blood (PB) samples were collected into heparin-containing tubes and immediately maintained on ice. Red blood cells were lysed by adding 10 ml ACK buffer to the samples and keeping them on ice for 10 min. After washing with cold PBS, the samples were stained with APC-anti-CD11c, PE-anti-CD11b, and FITC-anti-TER119. This protocol detected apoptotic erythroid cells exposed on Mo-DC surfaces, including cells attached to Mo-DC PS receptors or undergoing phagocytosis. Second, a recently reported protocol was used to detect the intracellular localization of apoptotic erythroid cells (Zoller et al, 2011). In detail, TER119 exposed on MO-DC surfaces was first blocked with saturating amounts of an unlabeled Ab against TER119 (10 μ g/10 μ l), then permeabilized with a BD Cytotfix/Cytoperm kit (BD) and stained with FITC-anti-TER119 and APC-anti-CD11c Abs. Apoptotic cells were detected using PE-annexin V (BD) according to the manufacturer's instructions. To purify inflammatory monocytes, defined as CX3CR1⁺Ly6c^{high}CD11b⁺CD11c⁻ cells, BM CD3 ϵ ⁻CD19⁻NK1.1⁻ cells were pre-enriched from *Cx3cr1*^{GFP/+} mice using a PE-Cy5-conjugated antibody mixture (CD3 ϵ , CD19, NK1.1), Cy5 microbeads, and an AutoMACSpro separation system

(Miltenyi Biotec). The cells were then stained with biotin-anti-Ly6c, PE-anti-CD11b, and APC-anti-CD11c antibodies, secondarily labeled with streptavidin-PE-Cy7, sorted on a MoFlo instrument (Beckman Coulter), and analyzed on a FACSCanto II (BD) in conjunction with FlowJo software (TreeStar).

Intracellular p-ERK and p-p38 staining

PBMCs were obtained from WT mice 4 h after LCMV C13 infection. Red blood cells were lysed, and the PBMCs were fixed with formaldehyde 10 min, permeabilized with ice-cold methanol on ice 20 min, and stained with FITC-anti-TER119, APC-anti-CD11c, and PE-pERK (clone 20A) or p-p38 (clone 30) (all from BD).

CTL Response and Virus Titer Detection

Splenocytes were harvested 9 days after infection. To detect LCMV-specific CTL, splenocytes were stained with FITC-anti-CD8 (KT15) (MBL), APC-anti-CD3 ϵ , and PE-H-2D^b-LCMV gp₃₃₋₄₁ tetramer (MBL). IFN- γ induction was analyzed by stimulating splenocytes with an LCMV H-2D^b-binding peptide (gp₃₃₋₄₁, KAVYNFATM; Peptide Institute), 50 U/ml murine IL-2, and Golgi Stop (BD) prior to incubation with CD8 α antibodies. Cells were then fixed and stained with IFN- γ -PE antibodies in PermWash solution (BD) according to the manufacturer's recommendation. For virus titration, MC57G cells were incubated with 10-fold serial dilutions of LCMV-containing serum in a 24-well plate at 37°C for 48 h. Cells were then fixed with 4% formalin, permeabilized with 0.5% Triton X and incubated with 10% fetal bovine serum. After incubation with murine LCMV immunoglobulin G1 (IgG1; Progen Biotechnik) and anti-mouse IgG-horseradish peroxidase (GE Healthcare), the plates were stained using an AEC peroxidase substrate kit (Vector Laboratories), and virus plaques were counted for each well.

Quantitative RT-PCR

Total RNA was extracted using an RNeasy Mini Kit (Qiagen), and cDNA was synthesized using random hexamers and SuperScript III reverse transcriptase. For real-time PCR, cDNA products equivalent to RNA from 2000 cells were amplified using a LightCycler®480 and SYBR Green I Master (Roche Diagnostics). The data were normalized to the levels of *gapdh* RNA expression in each sample. The primers used for real-time PCR were as follows: *Ccr2*, 5'-ATGAGTAACTGTGTGATTGACAAGCA-3' (sense) and 5'-GCAGCAGTGTGTCATTCCAAGA-3' (anti-sense); *P2y2*, 5'-CCGAGAGCTCTTTAGCCATTT-3' (sense) and 5'-GCCATAAGCACGTAACAGACC-3' (anti-sense); *Gapdh*, 5'-TCCACCACCCTGTTGCTGTA-3' (sense) and 5'-ACCACAGTCCATGCCATCAC-3' (anti-sense); LCMV *GP*, 5'-CATTACCTGGACTTTGTCAGACTC-3' (sense) and 5'-GCAACTGCTGTGTTCCCGAAAC-3' (anti-sense); LCMV *NP*, 5'-CAATGGACGCAAGCATTGAG-3' (sense) and 5'-GTTCTTCTGCACTGAGCCTCC-3' (anti-sense); *Mfge8*, 5'-GATCTTTCCAAC AACCT GCCTCC-3' (sense) and 5'-ACCGCTTTCATCCTGGATGAACTC-3' (anti-sense); *Bai1*, 5'-GCTGGCAGAAGCTGACGAT-3' (sense) and 5'-CACGGAGAT GACCTTAGAGTTG-3' (anti-sense); *Stab2*, 5'-GCTGTCGTCCTGGTTACTG-3' (sense) and 5'-CCAAGGCATCAATGTCATC-3' (antisense); *Tyro3*, 5'-TGCCTGCTTCGGA ACTTG-3' (sense) and 5'-GCCTGAGTCGGTACGAATG-3' (antisense); *Axl*, 5'-CCAGTCACAGGACACAGCTC-3' (sense) and 5'-GTGACTCCCTTGGCATTG-3' (anti-sense); *Mertk*, 5'-GGAAAGCGCAGGGACTTAC-3' (sense) and 5'-CTGTGCAGGTGGCATTGTG-3'

(anti-sense); *Gas6*, 5'-GGCATGTGGCAAACCT ATCT-3' (sense) and 5'-CGAGCTCACTCTCCTTGAA3' (antisense). Primers were synthesized for *Ccr2*, *P2y2*, LCMV *GP* (Ejrnaes et al, 2006), LCMV *NP* (Djavani et al., 2001), *Mfge8*, *Bai1*, *Stab2*, *Tyro3*, *Axl*, *Mertk*, and *Gas6* by Operon Biotechnology, and for *Gapdh* by Toyobo.

Ex vivo Hemophagocytosis Assay

TER119⁻ Mo-DCs and TER119⁺ erythroid cells or Gr1^{high}Ly6c⁺ granulocytes were separately sorted from the BM of either WT or *Mfge8*^{-/-} mice at 12 h after LCMV C13 infection. The Mo-DCs were precultured in 10 % FCS-RPMI supplemented with 20 ng/ml GM-CSF, washed, then incubated with either isotype Ab cocktails (50 µg/ml of each) or blocking antibodies against Tim1, Tim4, α_v , β_3 , or $\alpha_v\beta_5$ (Su et al., 2007) (50 µg/ml of each). TER119⁺ erythroid cells were labeled with pHrodo-SE (0.5 µg/ml, Life Technologies) at room temperature for 30 min, then washed with 10 % FCS PBS after stopping the reaction with 1 ml 100% FCS. To opsonize TER119⁺ erythroid cells, the cells were incubated with goat serum at 4°C for 1 h, washed, then incubated with anti-goat IgG antibody at 4°C for 1 h. The Mo-DCs (4×10^4) and pHrodo-labeled erythroid cells (4×10^5), opsonized erythroid cells (4×10^5), or granulocytes (2×10^5) were co-cultured in 100 µl of 10 % FCS-RPMI supplemented with 20 ng/ml GM-CSF in 96 well U-bottom Ultra-Low Attached plate (Corning) at 37°C for 2 h for Diff-Quick staining and flow cytometry or 24 h for ELISA. To block internalization of apoptotic erythroid cells into Mo-DCs, control DMSO (0.01%) or cytochalasin D (5µM) was added in culture for 2 h, washed out, then 24 h supernatants were subjected to IL-10 ELISA.

Ex vivo Induction of PS and PS receptors by IFN- α

Erythroid cells and granulocytes, sorted from the BM of WT mice, and T cells, B cells, NK cells, sorted from the spleen of WT mice, were stimulated *ex vivo* with different doses of IFN- α for 18 h, then subjected to annexin V staining. In addition, inflammatory monocytes, sorted from the BM of WT mice, were stimulated *ex vivo* with IFN- α (1×10^4 U/ml) for 24 h, then, subjected to the analysis of PS receptor expression.

Liver Histology

Livers were isolated 9 days after C13 infection, then fixed in 4% paraformaldehyde, embedded in paraffin, sectioned at 5- μ m thickness, and stained with hematoxylin/eosin.

Statistical Analysis

We evaluated the statistical significance of the obtained values by Student's *t*-test. We considered a *P* value <0.05 as significant.

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study, performed experiments, analyzed data, and wrote the manuscript. T.S., S.Y., K.K., and M.H. assisted N.O. with experiments. H.A., H.Y., K.A., K.H., N.T., M.H., and K.S. provided reagents, mice, discussion, and advice on some parts of this study. T.O. supervised the overall project and wrote the manuscript. The authors declare no competing financial interests.

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Figure legends

Figure 1. TLR Ligands Induce Hemophagocytosis

(A) Hemophagocytosis (%) 18 h after an injection of PBS (vehicle) or the indicated TLR or NLR ligand. (B) Sorted CD11c⁺TER119⁺ cells from the BM of CpG-injected mice were stained with Diff-Quick. Red and black arrows indicate hemophagocytosed and attached erythroid cells, respectively. Original magnification, x100. (C) Sorted CD11c⁺TER119⁺ cells from the PB of CpG injected mice were analyzed by electron microscope. Scale bar, 1 μ m. Arrows indicate hemophagocytosed erythroid cells. (D) Cell-surface staining of CD11c⁺TER119⁺ cells in the peripheral blood (PB) of 200-mg CpG-injected WT mice after blocking with an unlabeled anti-TER119 Ab. (E, F) Representative flow cytometry (FCM) results (E) and percentage (F) of intracellular CD11c⁺TER119⁺ cells in the PB of WT mice 18 h after injection of the indicated doses of CpG. Data represent the mean \pm s.d. of three independent experiments. * $P < 0.01$ and ** $P < 0.05$.

Figure 2. CCR2 Is Involved in Developing CpG-Induced Hemophagocytosis

(A-C) Kinetics of serum CCL2 (A), CCL7 (B), and CCL12 (C) production in WT mice after 200- μ g CpG injection. (D) Relative *Ccr2* mRNA expression in the indicated cell populations from WT mice 18 h after 200- μ g CpG injection. BMDC, bone marrow-derived DCs. (E, F) Percentage of CD11c⁺TER119⁺ cells in the PB (E) and BM (F) of the indicated mice 18 h after 200- μ g CpG injection. Data represent the mean \pm s.d. of three independent experiments. * $P < 0.01$, ** $P < 0.05$.

Figure 3. “Find Me” and “Eat Me” Signals in CpG-Induced Hemophagocytosis

(A) Serum ATP level over time in WT mice after 200- μ g CpG injection. (B) Relative *P2y2* mRNA expression in CD11c⁺TER119⁺ cells from the PB and the indicated BM-derived cell populations 18 h after 200- μ g CpG injection. (C) Percentage of CD11c⁺TER119⁺ cells in the PB of WT mice 18 h after 200- μ g CpG injection with 6 mg suramin. (D) Percentage of annexin V⁺ cells among TER119⁺ cells in the PB of WT mice over time after PBS (vehicle) or 200- μ g CpG injection. (E) Expression of “eat me” signal receptors on CpG-induced CD11c⁺TER119⁺ cells. Shaded histograms represent cells stained with mAbs for the indicated molecules, and open histograms show labeling with isotype controls. (F) Impaired CpG-induced hemophagocytosis in WT mice treated with blocking antibodies against Tim1, Tim4, α_v , and β_3 . The proportion of CD11c⁺TER119⁺ cells 18 h after 200- μ g CpG injection with isotype control Ab cocktails (500 μ g each) or the indicated antibody mixtures (500 μ g each). Ab mix: mixture of antibodies against Tim1, Tim4, α_v , and β_3 . (G) IL-10 serum levels 24 hr after the indicated treatments. Data represent the mean \pm s.d. of three independent experiments. **P* < 0.01.

Figure 4. Essential Role of Type I IFNs in C13-Induced Hemophagocytosis

(A, B) Kinetics of serum IFN- α (A) and IFN- β (B) production in WT mice after 2×10^6 pfu C13 infection. (C) Representative flow cytometry (FCM) profiles of CD11c⁺TER119⁺ cells in the PB of WT and *Ifnar1*^{-/-} mice 24 h after 2×10^6 pfu C13 infection. (D, E) The percentage of CD11c⁺TER119⁺ cells (D) and annexin V⁺ cells among TER119⁺ cells (E) in the PB of WT mice 4 h and 24 h after injection of 2×10^6 pfu C13, respectively. (F, G) Expression of “eat me” signal receptors on

CD11c⁺TER119⁺ cells in the PB of WT (F) and *Ifnar1*^{-/-} (G) mice 24 h after 2 x 10⁶ pfu C13 infection. Shaded histograms represent cells stained with mAbs specific for the indicated molecules, and open histograms show labeling with isotype controls.

Figure 5. Hemophagocytosis-Dependent IL-10 Production from MO-DCs upon C13 Infection

(A) C13-induced hemophagocytosis in WT mice treated with either isotype control Ab cocktails (500 µg each) or blocking antibodies against PS receptors (500 µg each). (B) IL-10 and TGF-β1 serum levels 24 h after C13 infection with the injection of isotype control Ab cocktails (500 µg each), Abs to block PS receptors or neutralize IL-10. (C) A representative FCM profile of IL-10 Venus expression in the indicated cell populations from the PB of *Il10*^{Venus} reporter mice 24 h after C13 infection. (D) The absolute number of cells expressing IL-10 Venus in the indicated cell populations from the PB of *Il10*^{Venus} reporter and WT mice 1 and 9 days after C13 infection. Data represent the mean ± s.d. from three independent experiments. **P* < 0.01.

Figure 6. Critical role of MFG-E8-integrin signal in IL-10 production from Mo-DCs

(A-F) *Ex vivo* hemophagocytosis assay. TER119⁻ Mo-DCs and TER119⁺ erythroid cells were separately sorted from the BM of LCMV C13 infected WT or *Mfge8*^{-/-} mice. The Mo-DCs were pretreated with control rat Ig (A, left panel, and C) or Abs to block PS receptors (B, right panel, and D), and co-cultured with pHrodo-labeled TER119⁺ erythroid cells for 2 h. The cells were stained with Diff-Quick. Red arrows indicate hemophagocytes. Original magnification, x20 (A). Representative FCM profiles of Mo-DCs co-cultured with unlabeled- (B), pHrodo-labeled-erythroid cells (C, D). IL-10

level in the indicated co-culture supernatants (E, F). Data represent the mean \pm s.d. of three independent experiments. * $P < 0.01$.

Figure 7. Physiological Significance of C13 Infection-Induced Hemophagocytosis

(A-F) Hemophagocytosis (%) (A) and IL-10 serum levels (B) 24 h after C13 infection. The percentage of LCMV GP₃₃₋₄₁-specific CD8⁺ T cells in the spleen (C), titer of LCMV C13 in the serum (D), and level of AST/ALT in the serum (E) of WT and *Cd11c-Cre Il10^{fl/fl}* (CKO) mice 9 days after C13 infection, and survival (%) of WT or CKO mice after LCMV C13 infection (F). Data represent the mean \pm s.d. of three independent experiments. * $P < 0.01$.

Figure 1. Ohyagi, et al.

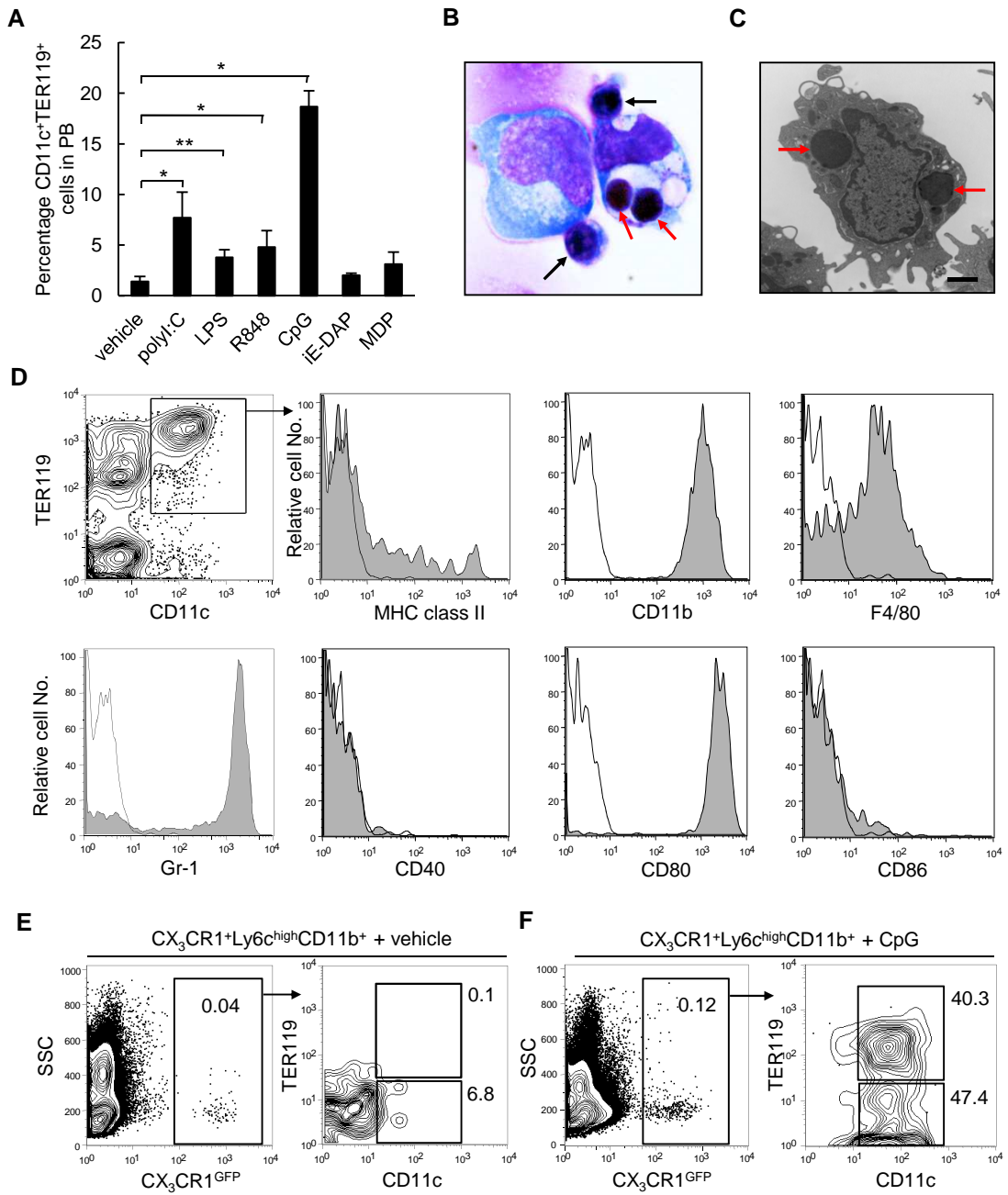


Figure 2. Ohyagi, et al.

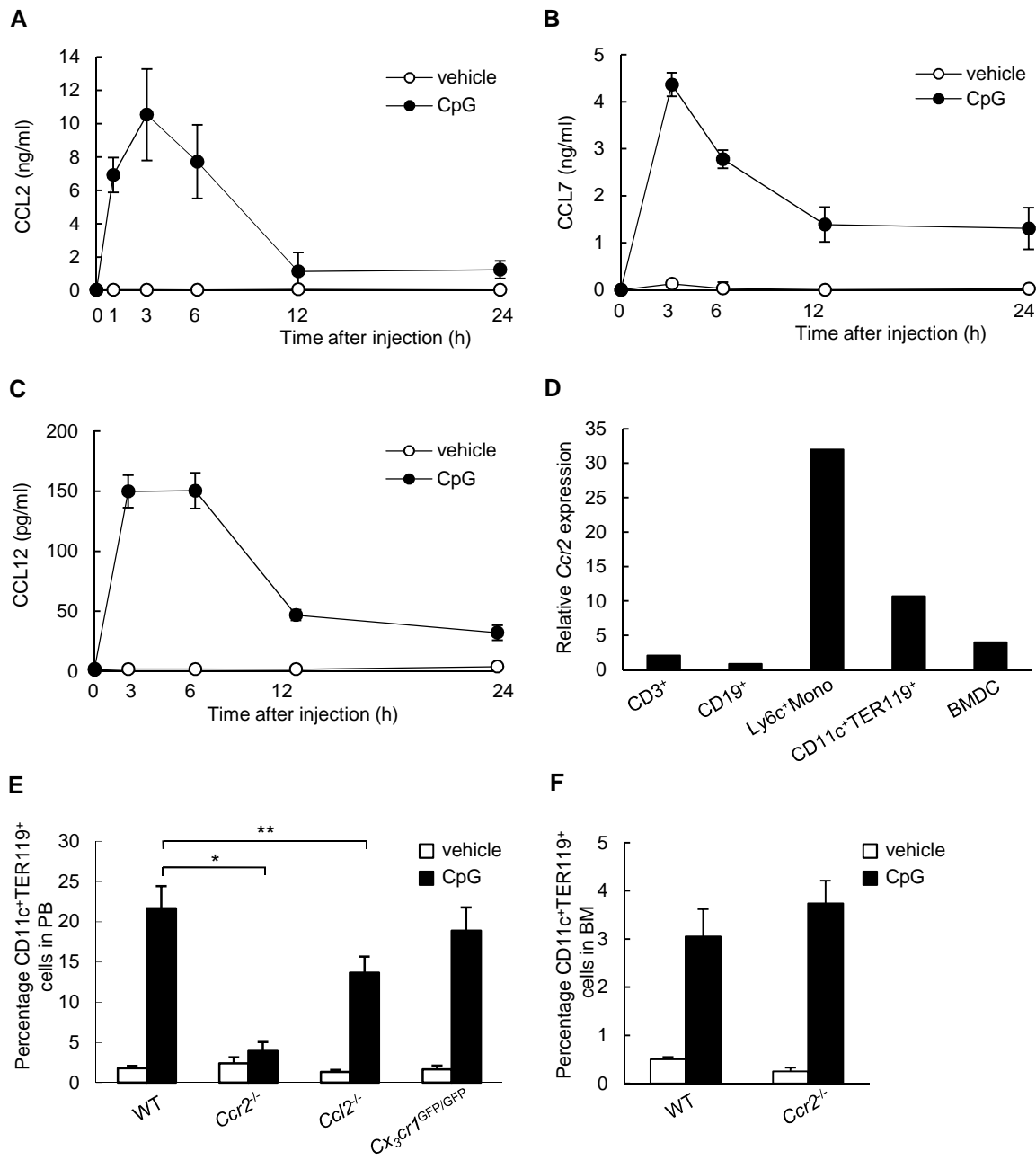


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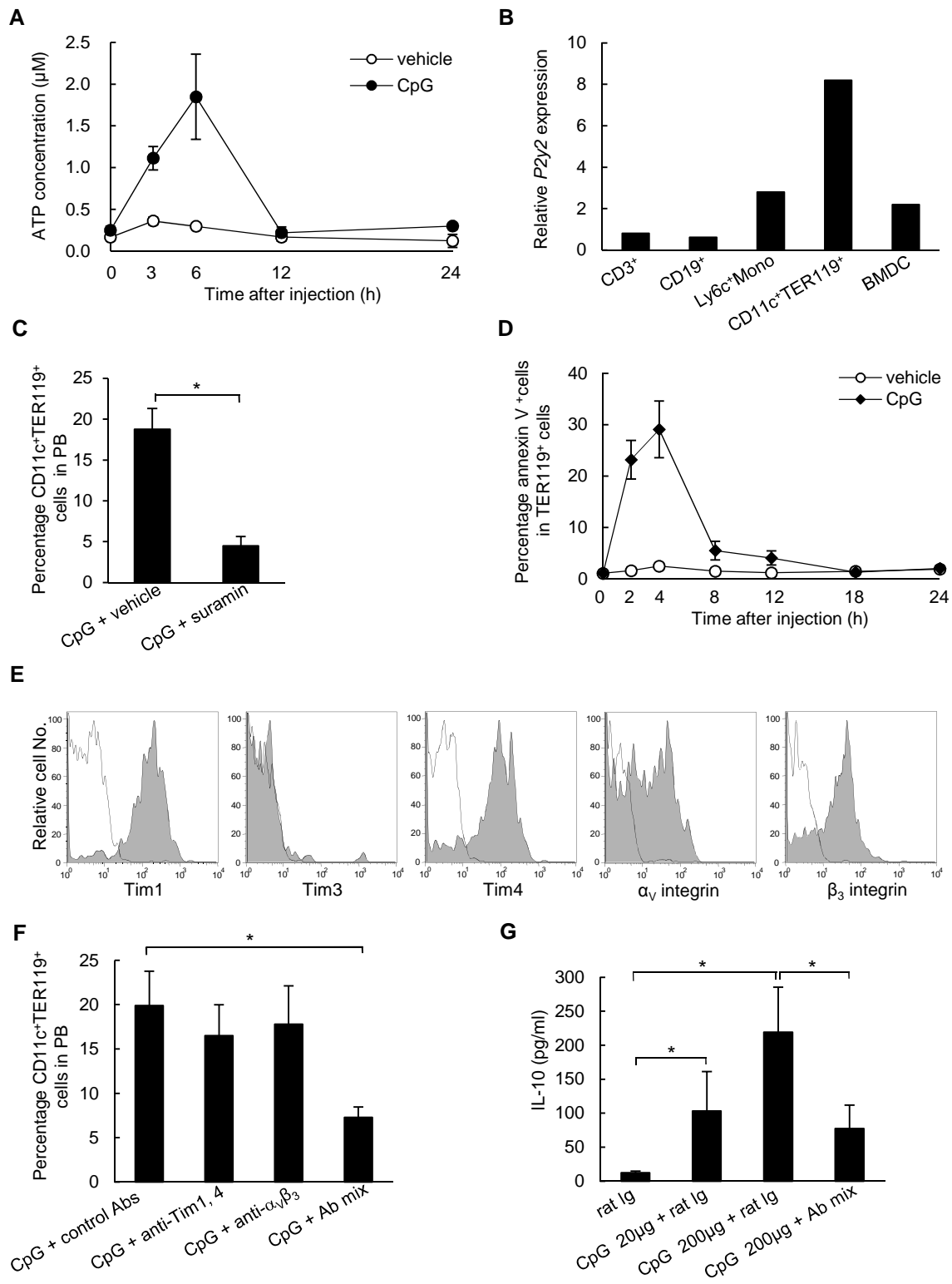


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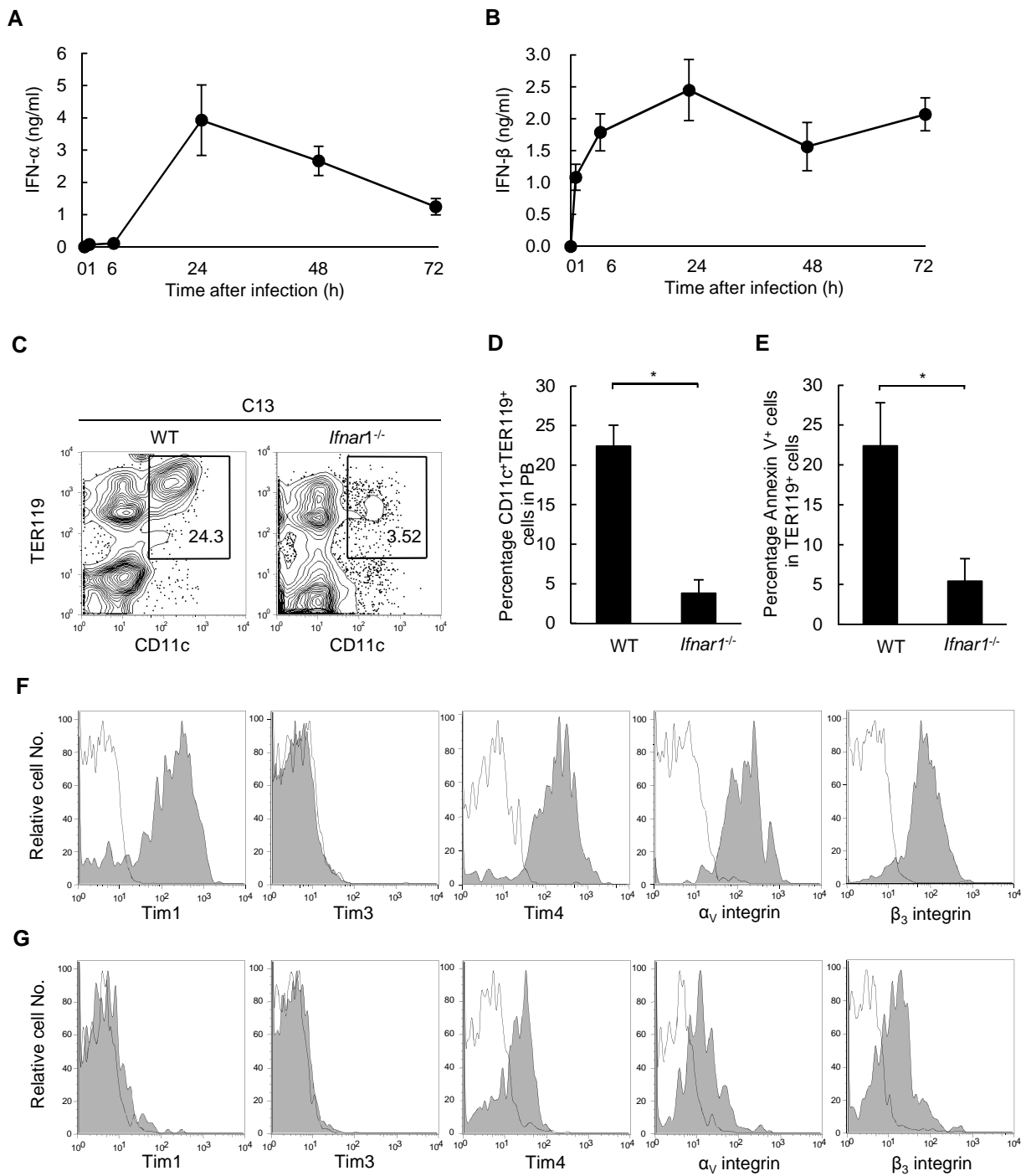


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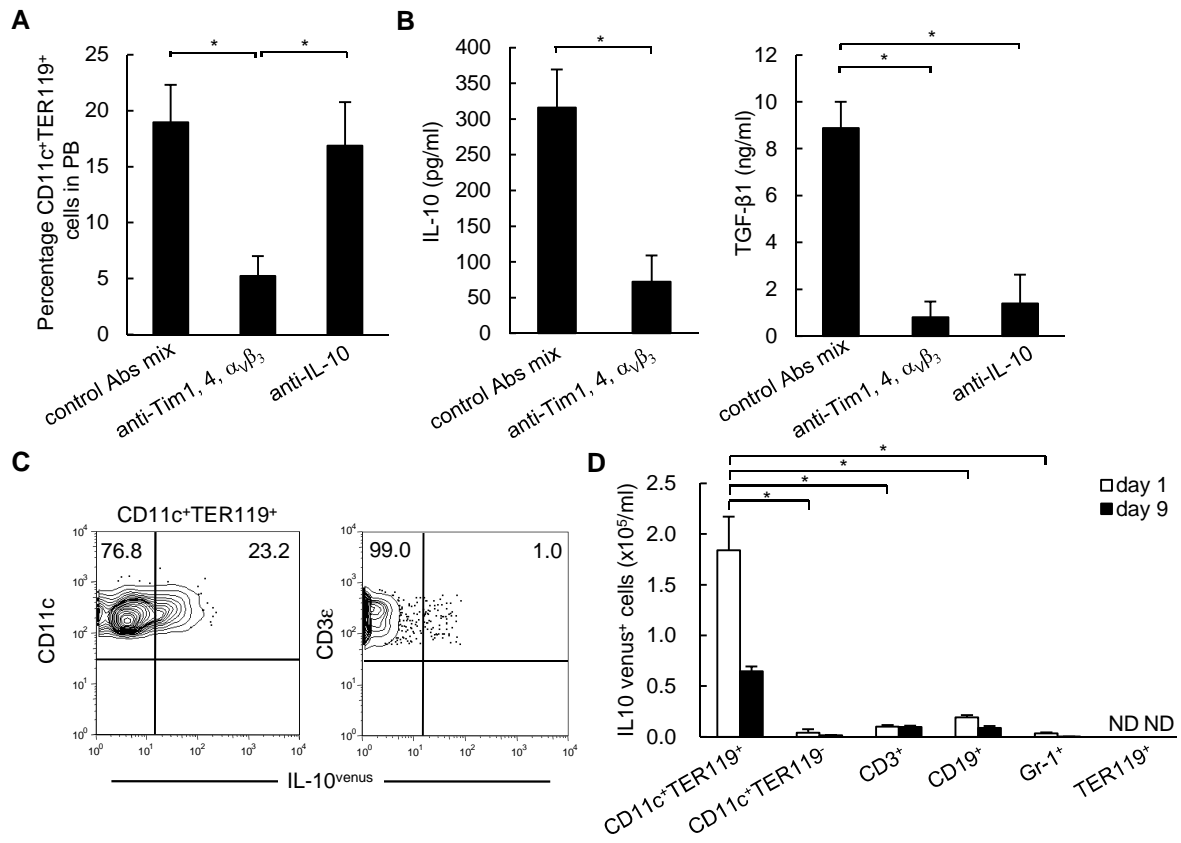


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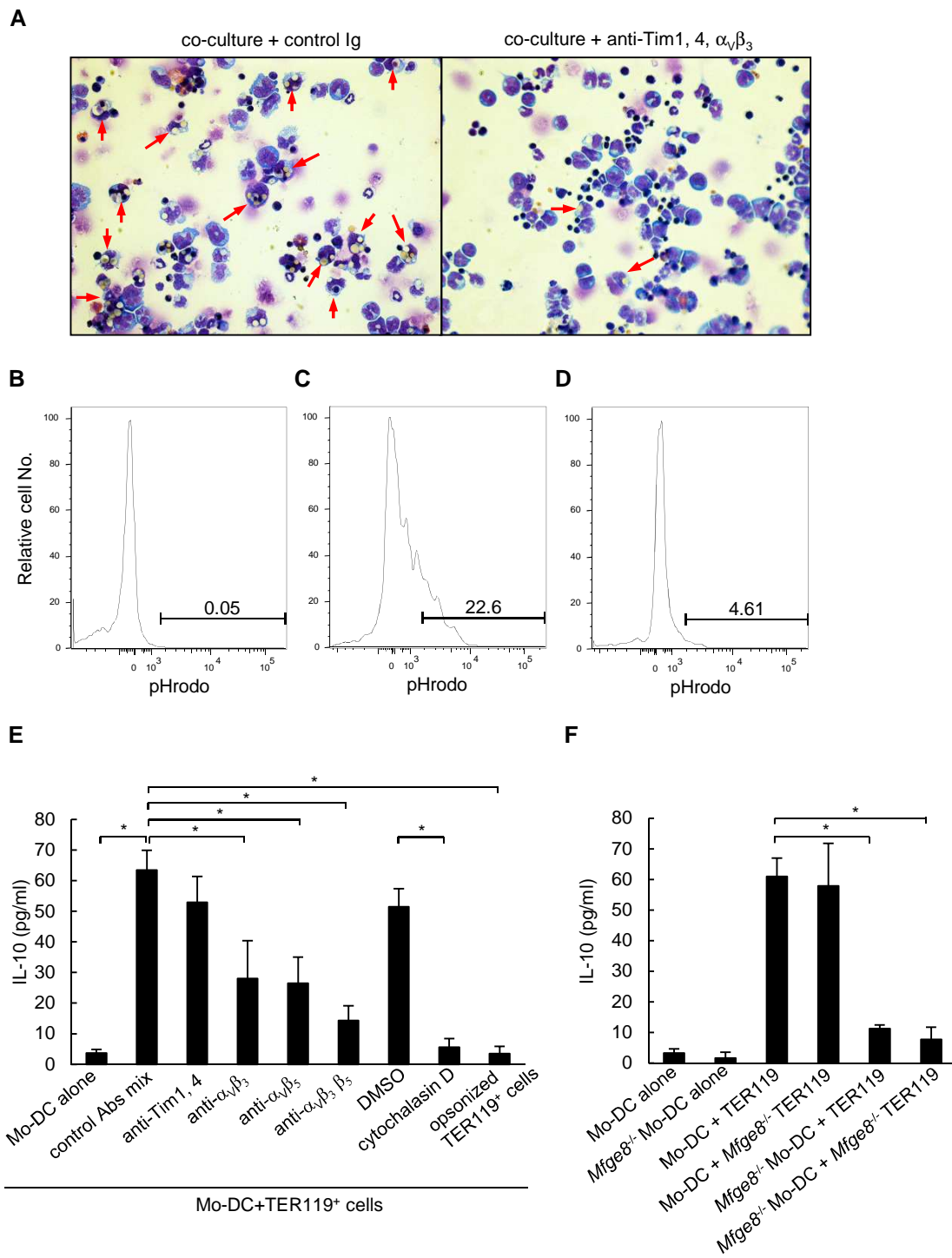
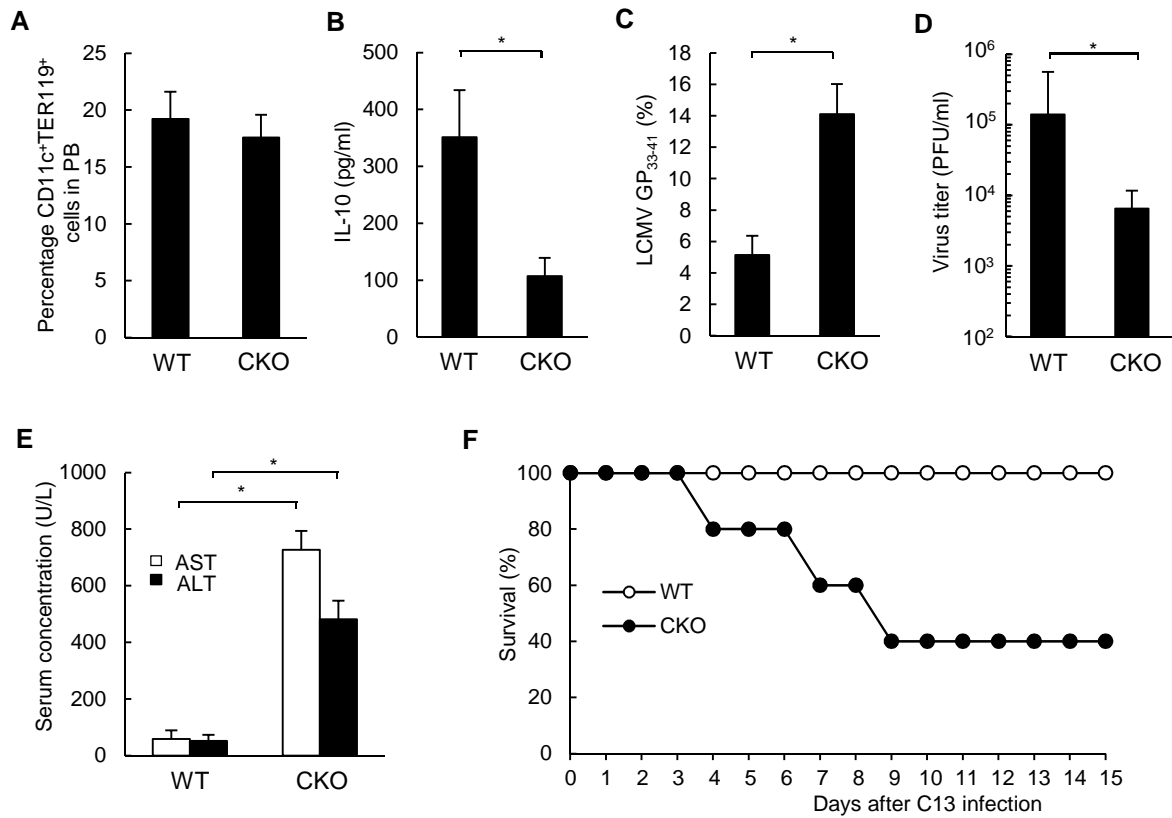


Figure 7. Ohyagi, et al.



Supplementary Information

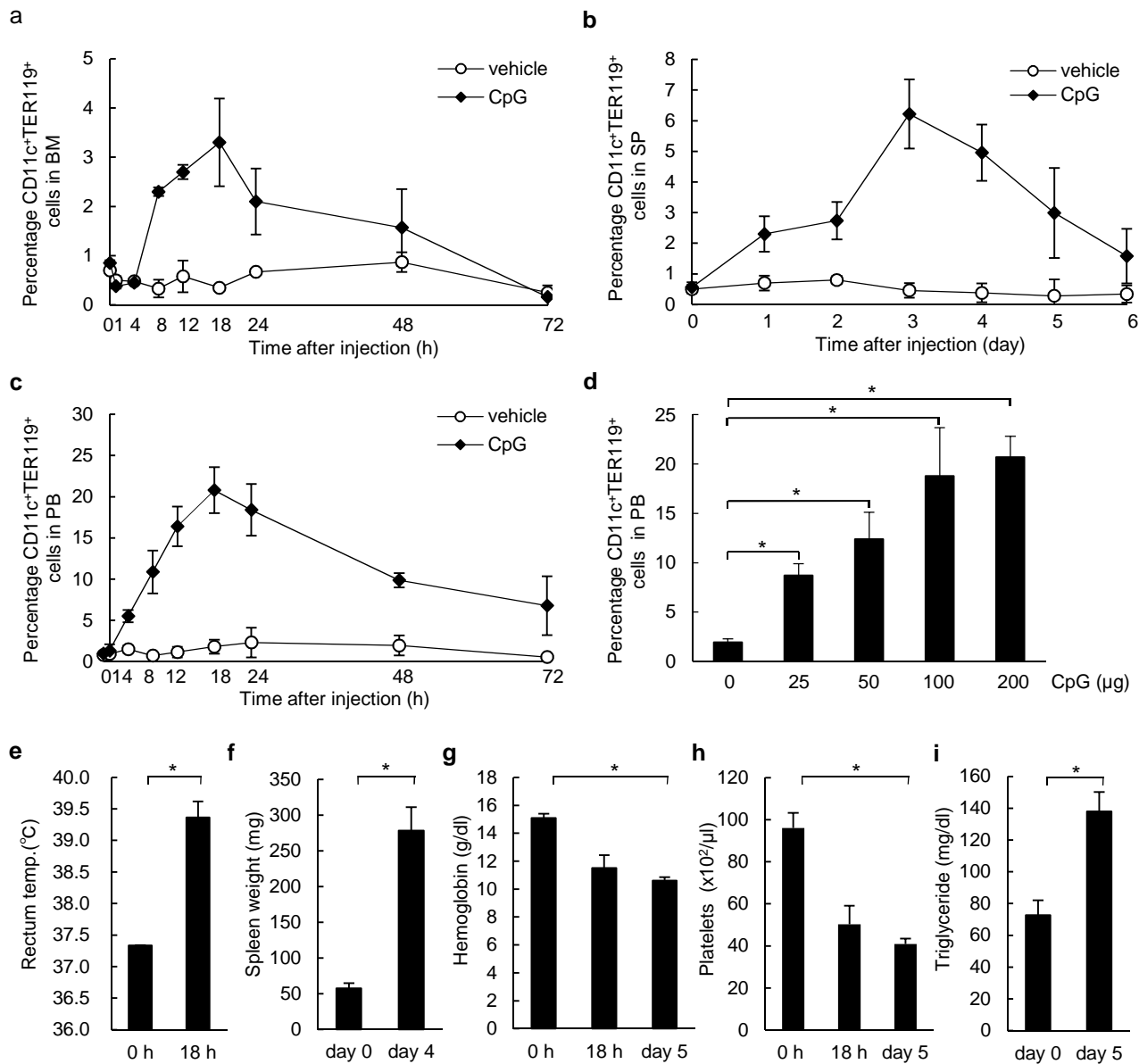


Figure S1. Hemophagocytosis in CpG-injected mice. **(a-c)** Time course of the percentage of CD11c⁺TER119⁺ cells in the bone marrow (BM) **(a)**, spleen (SP) **(b)**, and PB **(c)** of WT mice after vehicle (PBS) or 200- μ g CpG injection. **(d)** Percentage of CD11c⁺TER119⁺ cells in the PB of WT mice 18 h after injection of the indicated doses of CpG. **(e-i)** Clinical parameters for HPS, i.e., rectal temperature ($^{\circ}$ C) **(e)**, spleen weight (mg) **(f)**, hemoglobin (g/dl) **(g)**, platelets ($\times 10^2/\mu$ l) **(h)**, and triglycerides (mg/dl) **(i)** were evaluated at 18 h **(e)** and at the indicated time points **(e-i)**, after an injection of PBS (vehicle) or 200- μ g CpG. Data represent the mean \pm s.d. of three independent experiments. * $P < 0.01$.

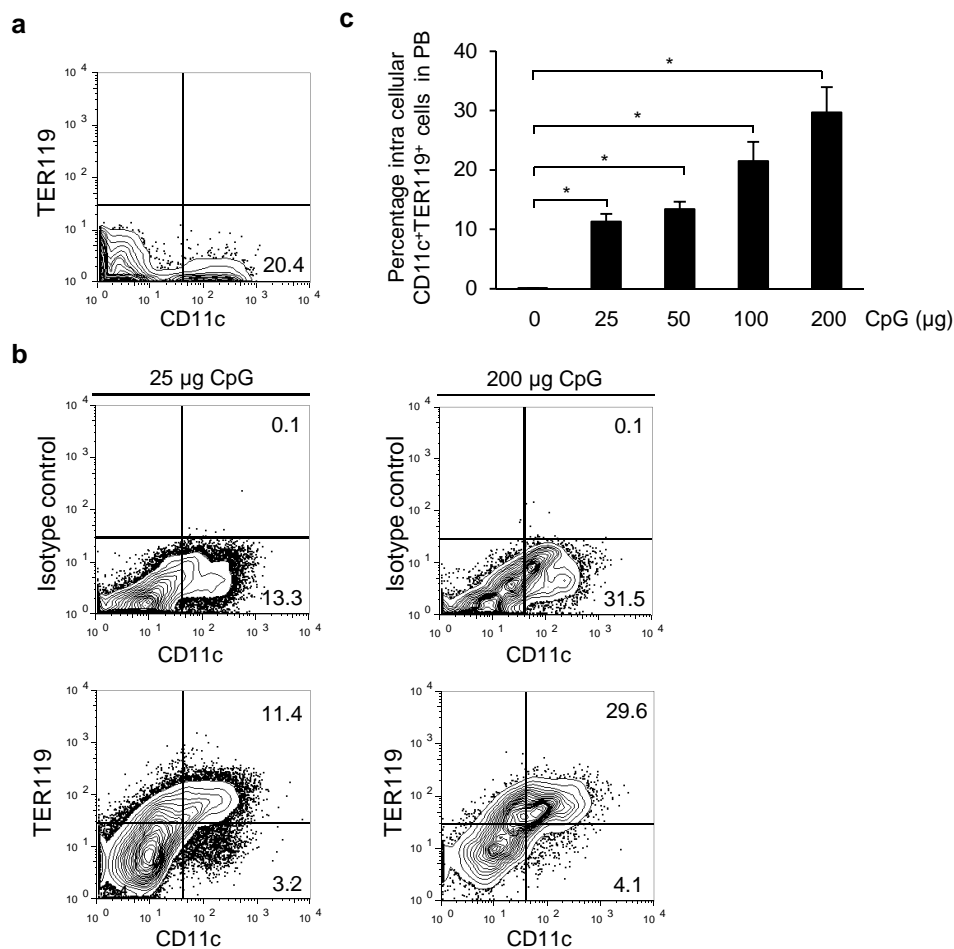


Figure S2. Intracellular detection of TER119⁺ cells inside CD11c⁺ cells. **(a)** Cell-surface staining of CD11c⁺TER119⁺ cells in the peripheral blood (PB) of 200-µg CpG-injected WT mice after blocking with an unlabeled anti-TER119 Ab. **(b, c)** Representative flow cytometry (FCM) results **(b)** and percentage **(c)** of intracellular CD11c⁺TER119⁺ cells in the PB of WT mice 18 h after injection of the indicated doses of CpG. Data represent the mean \pm s.d. of three independent experiments. * $P < 0.01$.

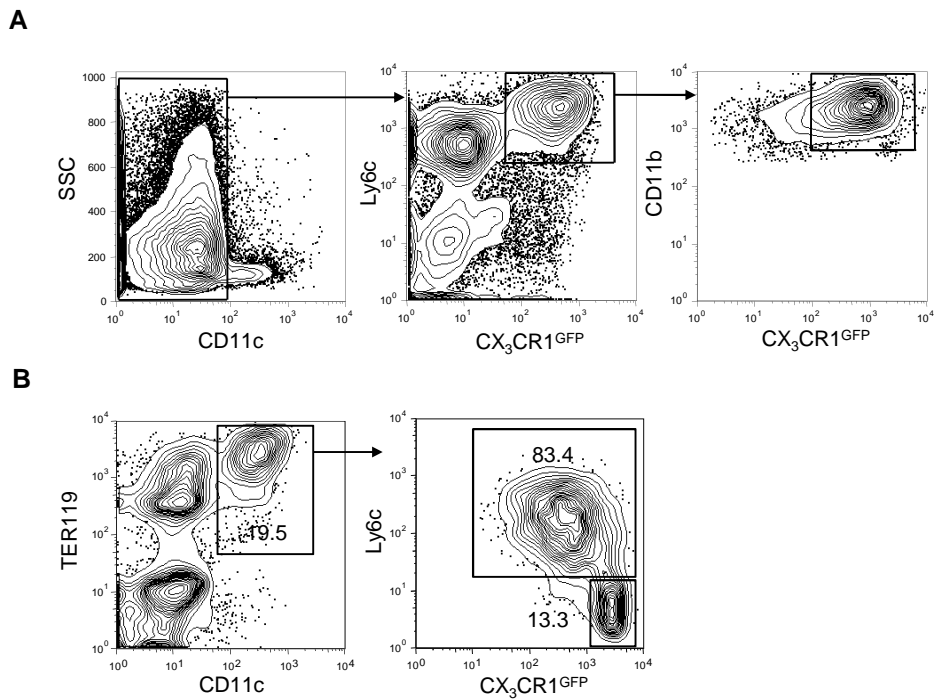


Figure S3. Inflammatory monocyte-derived DCs are the major hemophagocytes in CpG-injected mice. (A) Purification of inflammatory monocytes, defined as Ly6c⁺CX₃CR1^{GFP/+}CD11b⁺CD11c⁻ cells, from the BM of *Cx₃cr1^{GFP/+}* mice. **(B)** Representative FCM profiles of CD11c⁺ TER119⁺ cells in the PB of *Cx₃cr1^{GFP/+}* mice 18 h after CpG injection.

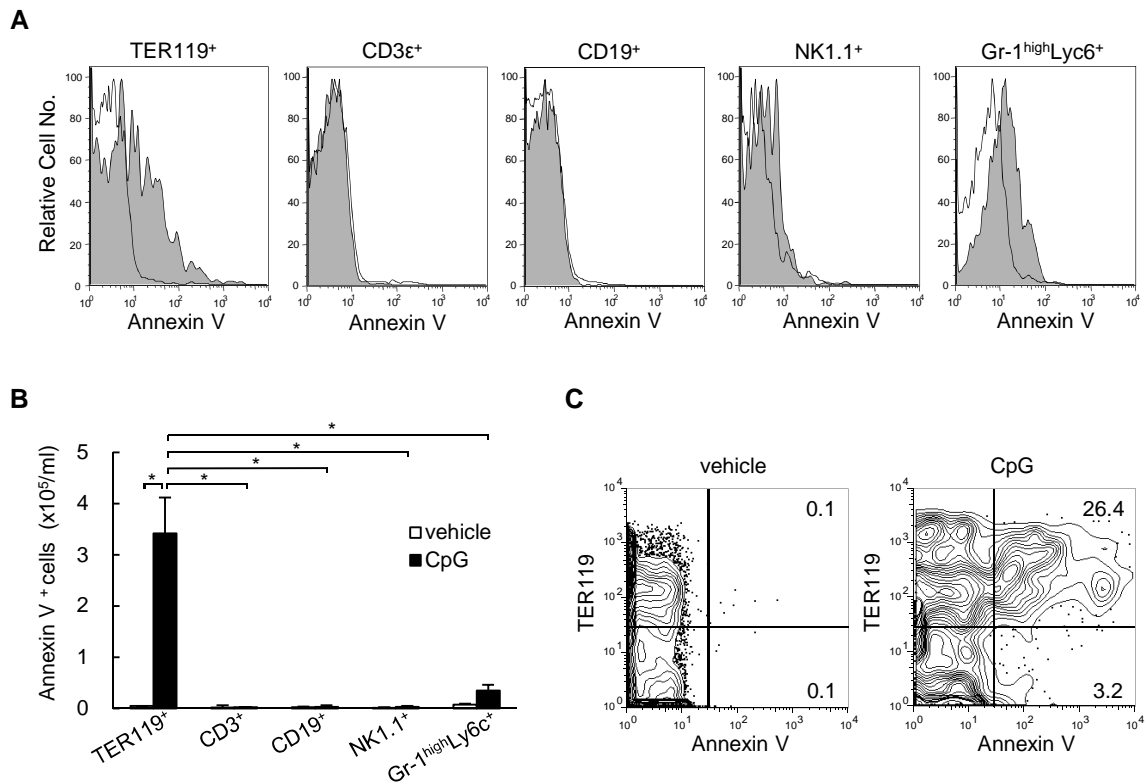


Figure S4. Apoptosis in CpG-injected WT mice.

(A) Representative FCM analysis of Annexin V staining in the different cell subset of the PB form WT mice 4h after PBS (open histogram) or CpG injection (shad histogram). (B,C) Absolute number of Annexin V⁺ cells (B) and representative FCM analysis of Annexin V and TER119 staining (C) in the PB form WT mice 4h after PBS (vehicle) or CpG injection. Data represent the mean \pm s.d. of three independent experiments. * $P < 0.01$.

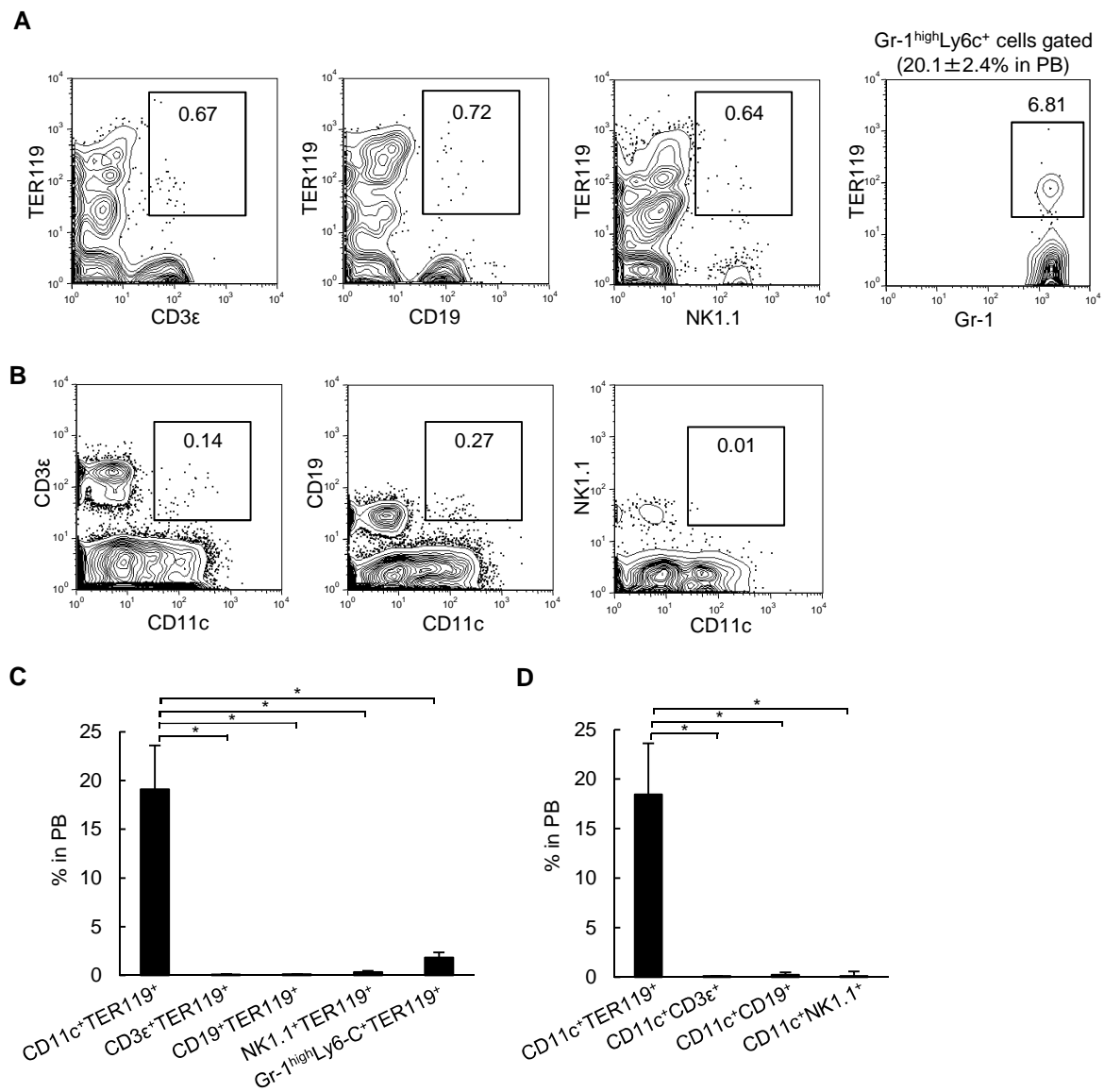


Figure S5. CD11c⁺ cells are the major phagocytes for TER119⁺ cells in the PB of Hemophagocytosis-induced mice. (A,C) Representative FCM analysis of the PB (A), and percentage of CD11c⁺TER119⁺, CD3 ϵ ⁺TER119⁺, CD19⁺TER119⁺, NK1.1⁺TER119⁺, and Gr-1^{high}Ly6c^{int}TER119⁺ cells in CpG-injected mice (C). (B,D) Representative FCM analysis of the PB (B), and percentage of CD11c⁺TER119⁺, CD11c⁺CD3 ϵ ⁺, CD11c⁺CD19⁺, and CD11c⁺NK1.1⁺ cells in CpG-injected mice (D). Data represent the mean \pm s.d. of three independent experiments. * P < 0.01.

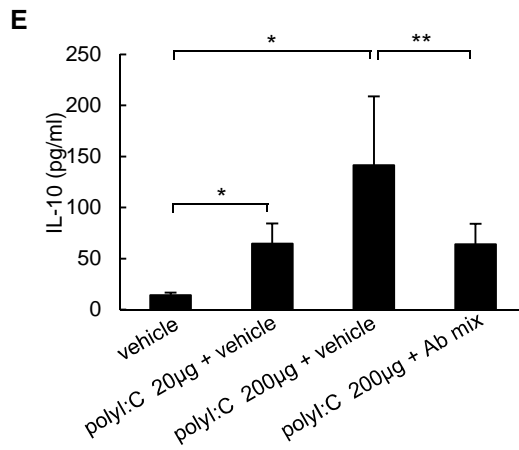
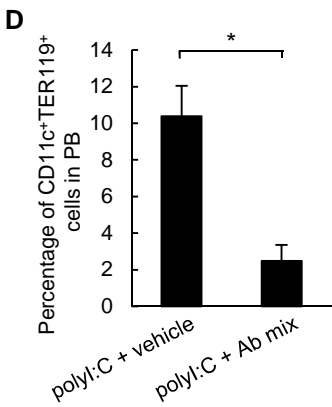
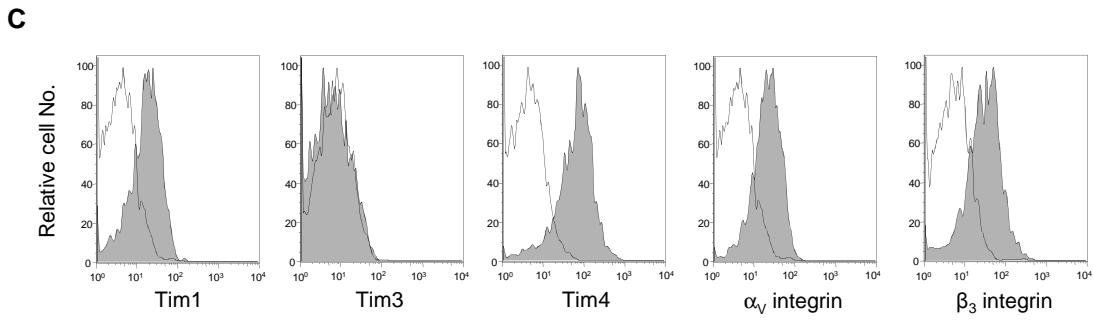
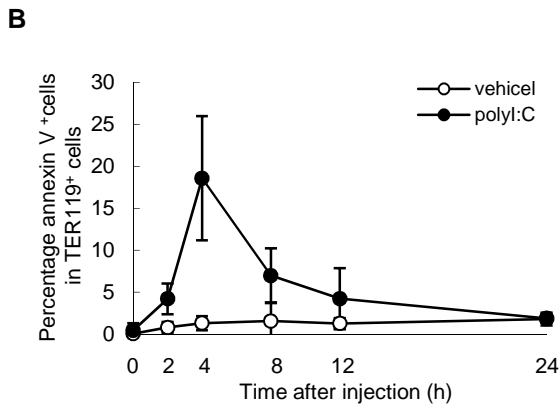
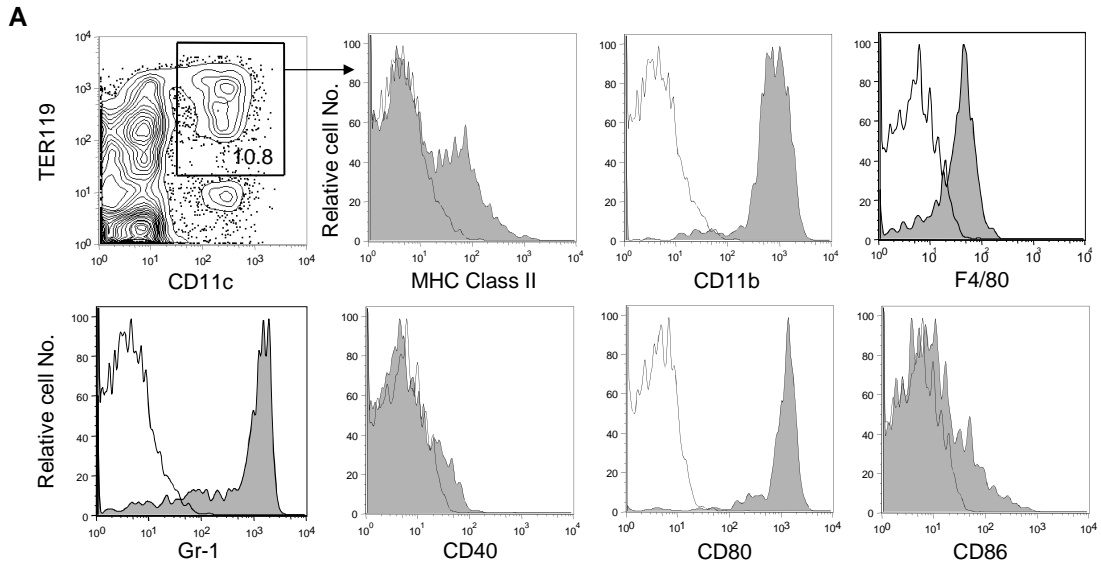


Figure S6. Mechanisms of poly I:C-induced hemophagocytosis.

(A) Phenotypic analysis of CD11c⁺TER119⁺ cells. CD11c⁺TER119⁺ cells gated on the contour plot were stained with mAbs to the indicated markers (shaded histograms) and isotype controls (open histograms). (B) Time course of the percentage of annexin V⁺ cells among the TER119⁺ cells in the PB of WT mice after PBS (vehicle) or 200- μ g poly I:C injection. (C) Expression of “eat me” signal receptors on poly I:C-induced CD11c⁺TER119⁺ cells. Shaded histograms represent cells stained with mAbs specific for the indicated molecules, and open histograms show labeling with isotype controls. (D) Impaired poly I:C-induced hemophagocytosis in WT mice treated with neutralizing antibodies against Tim1, Tim4, α_v , and β_3 . The proportion of CD11c⁺TER119⁺ cells 18 h after a 200- μ g poly I:C injection together with a control rat IgG or with the indicated antibody mixtures (500 μ g each). Ab mix, a mixture of antibodies against Tim1, Tim4, α_v , and β_3 . (E) Serum IL-10 levels 24 hr after the indicated treatments. Data represent the mean \pm s.d. of three independent experiments. * $P < 0.01$. ** $P < 0.05$.

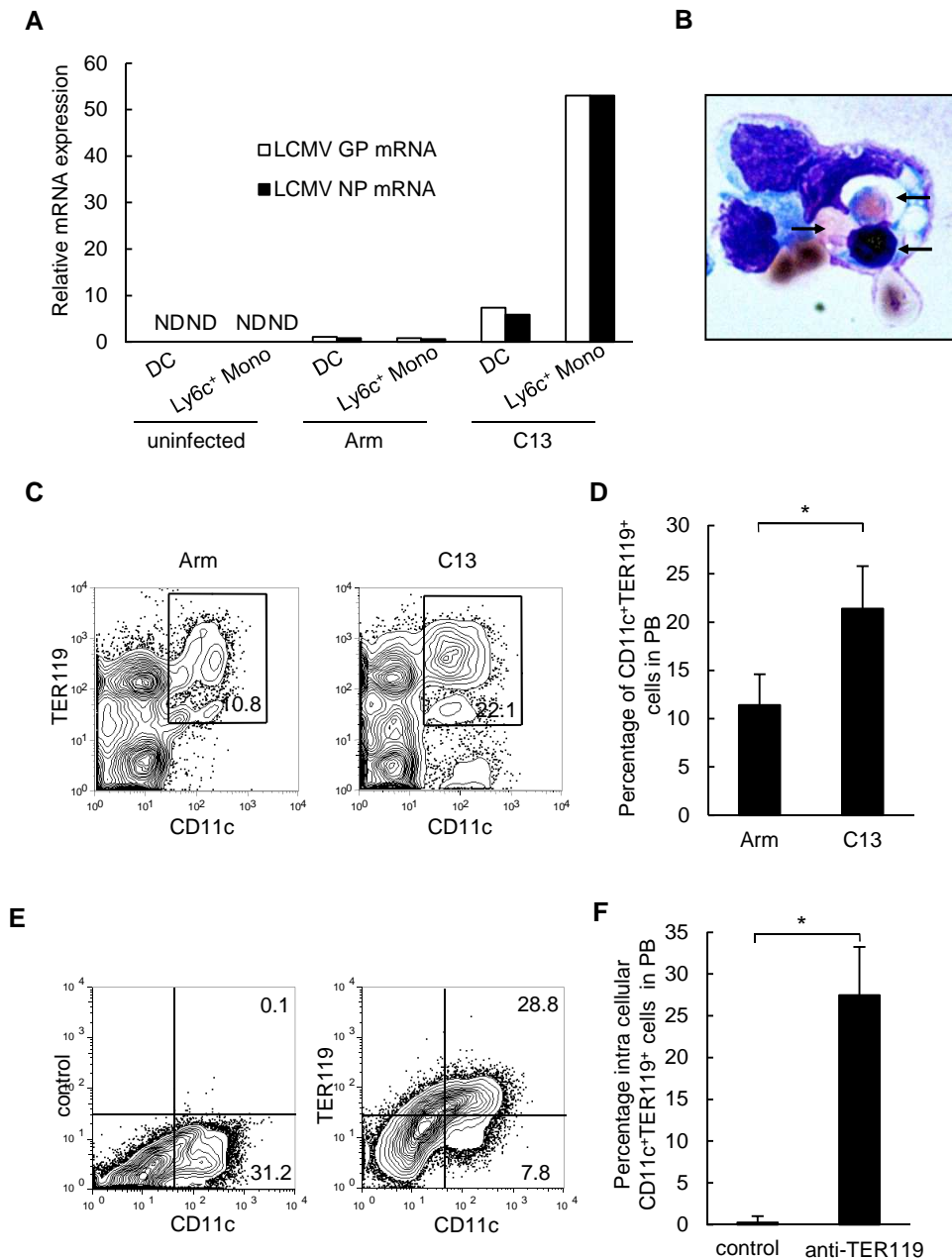


Figure S7. LCMV C13 induces efficient hemophagocytosis.

(A) Relative mRNA levels of LCMV glycoprotein (GP) and nucleoprotein (NP) in DCs (DC) and inflammatory monocytes (Ly6c⁺ Mono) from WT mice 2 days after 2×10^6 pfu Arm or C13 infection. Cells were isolated from the BM of infected WT mice. Data are representative of three independent experiments. (B) Sorted CD11c⁺TER119⁺ cells from the BM of LCMV C13 infected mice were stained with Diff-Quick. Arrows indicate hemophagocytosed erythroid cells. Original magnification, x40. (C, D) Representative FCM profiles (C) and percentage (D) of CD11c⁺TER119⁺ cells in the PB of WT mice 24 h after 2×10^6 pfu Arm or C13 infection. Intracellular staining of TER119⁺ cells inside CD11c⁺ cells in mice infected with C13. (E, F) Representative FCM results (E), and percentage (F) of intracellular CD11c⁺TER119⁺ cells in the PB of WT mice 24 h after C13 infection. Data represent the mean \pm s.d. of three independent experiments. * $P < 0.01$.

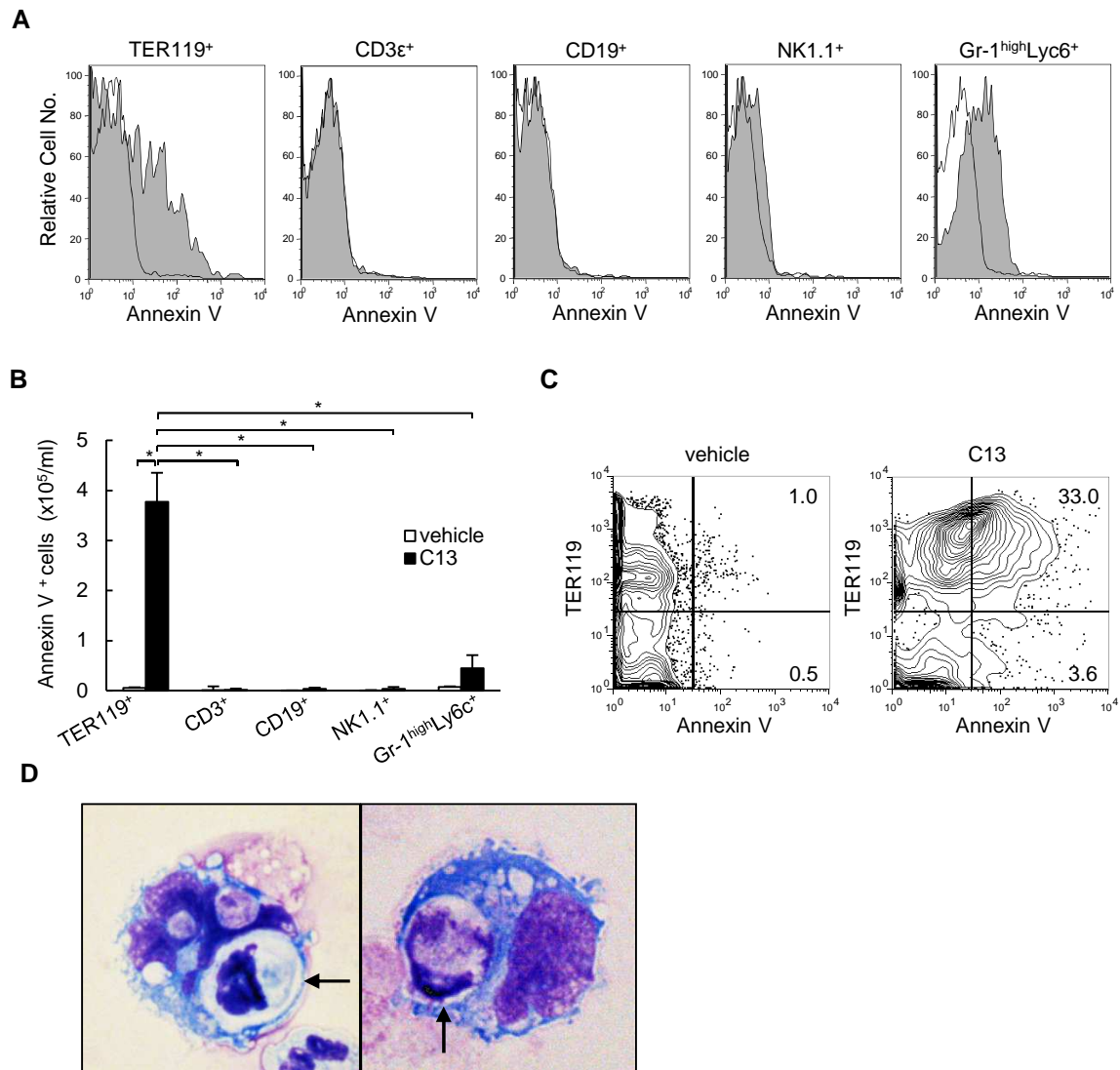


Figure S8. Apoptosis in LCMV C13-infected WT mice.

(A) Representative FCM analysis of Annexin V staining in the PB cell subsets from WT mice 4h after PBS (open histogram) or LCMV C13 infection (shad histogram). (B,C) Absolute number of Annexin V⁺ cells (B) and representative FCM analysis of Annexin V and TER119 staining (C) in the PB from WT mice 4h after PBS injection (vehicle) or LCMV C13 (C13) infection. (D) Sorted hemophagocytes from PB of LCMV C13 infected WT mice, were stained with Diff-Quick. Arrows indicate hemophagocytosed granulocytes. Original magnification, x40. Data represent the mean \pm s.d. of three independent experiments. * $P < 0.01$.

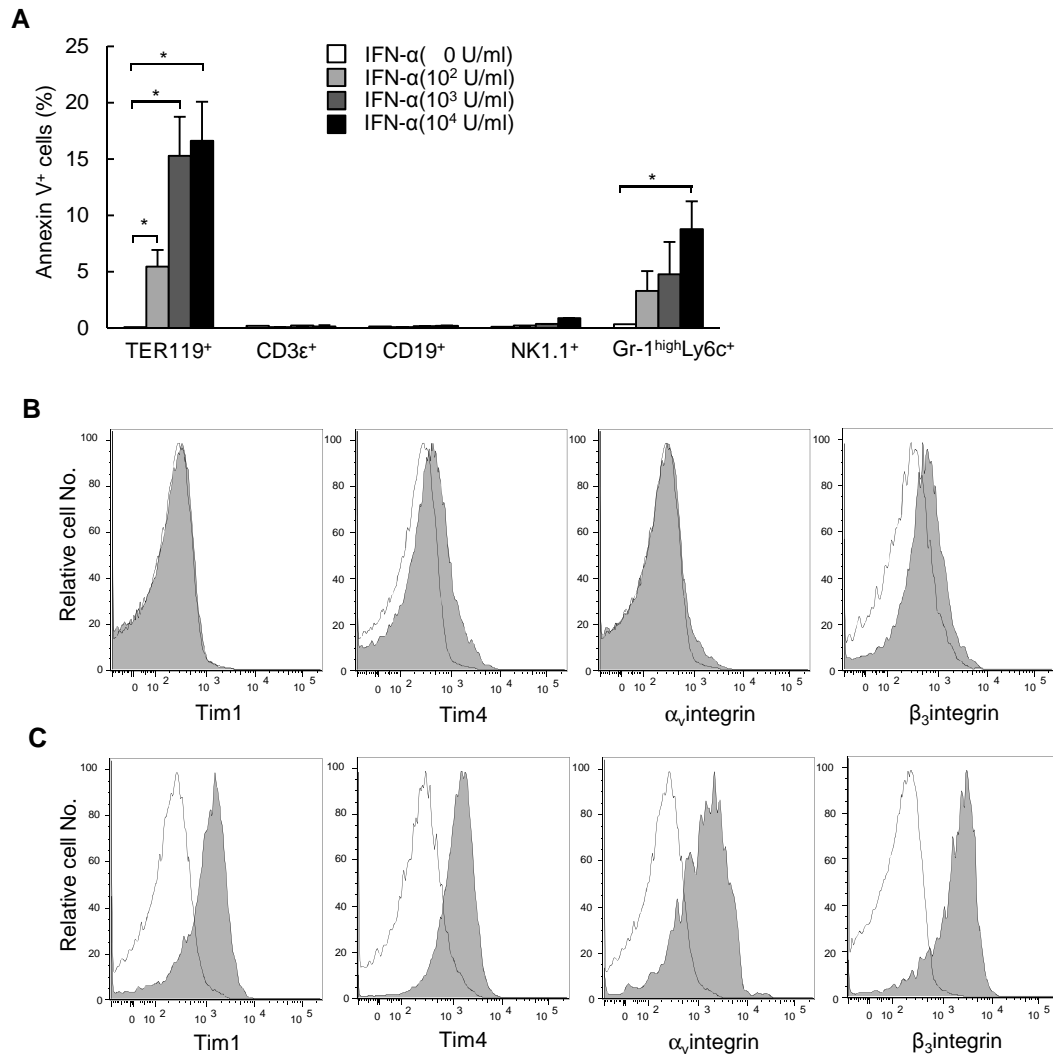


Figure S9. Interferon- α induces apoptosis and PS receptor expression in inflammatory monocytes.

(A) Percentage of annexin V⁺ cells in different cell type. Different cells from spleen of WT mice were stimulated with different concentration of interferon- α for 18h.

Data represent the mean \pm s.d. of three independent experiments. * $P < 0.01$.

(B,C) Representative FCM analysis of “eat me” signal receptors on inflammatory monocytes from BM of WT mice. Ly6c⁺ inflammatory monocytes were stimulated without (B) or with (C) interferon- α (10⁴ U/ml) for 24h. One of representative data from two independent experiments.

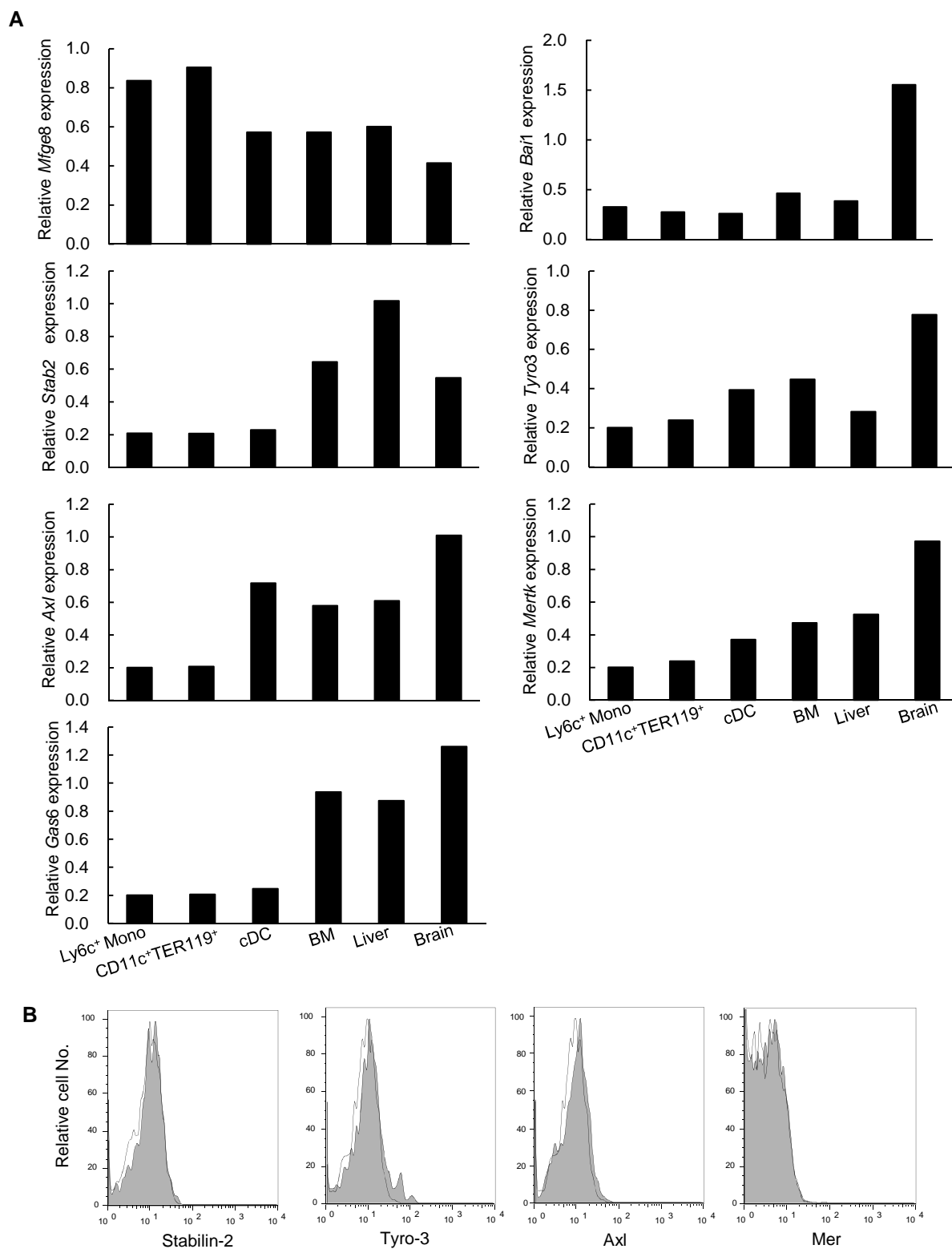


Figure S10. Expression of phosphatidylserine (PS) receptors and its recognition molecule in WT mice.

(A) Relative PS receptors and recognition molecule mRNA expression in the indicated cell populations and tissue from WT mice. cDC, conventional DC. (B) Expression of PS receptors on CD11c⁺TER119⁺ cells in the PB of WT mice 24 h after 2 x 10⁶ pfu C13 infection. Shaded histograms represent cells stained with Abs specific for the indicated molecules, and open histograms show labeling with isotype controls.

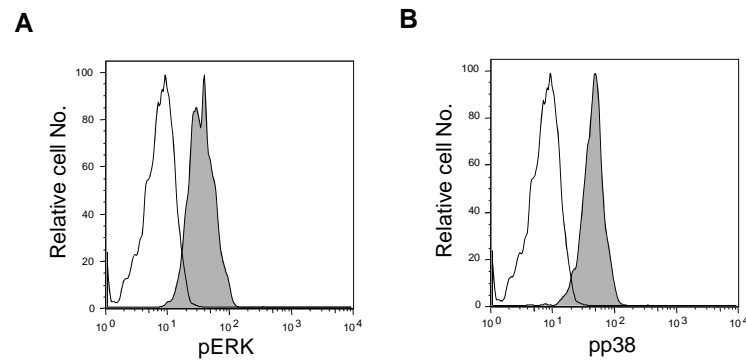


Figure S11. Phosphorylation of ERK and p38 in the hemophagocytosis-performing Mo-DCs of LCMV clone 13-infected WT mice. (A, B) PBMCs from WT mice 4 h after LCMV C13 infection were fixed with formaldehyde, permeabilized with methanol, and stained with cell-surface makers (CD11c and TER119), and pERK (A) or pp38 (B) (shaded histogram) and isotype control (open histogram). The data were gated on the CD11c⁺TER119⁺ cells. Data are representative of three independent experiments.

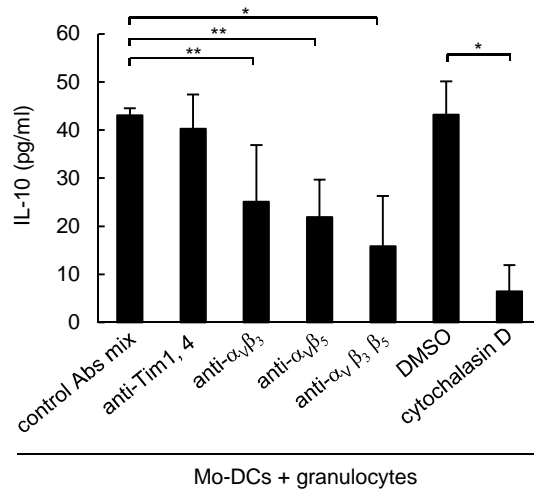


Figure S12. IL-10 production from Mo-DCs is dependent on hemophagocytosis. IL-10 level in the co-culture supernatants from the *ex vivo* hemophagocytosis assay. TER119⁺ Mo-DCs and granulocytes from the BM of C13-infected mice were co-cultured with the indicated reagent(s) for 24 h. Data represent the mean \pm s.d. of three independent experiments. * $P < 0.01$. ** $P < 0.005$.

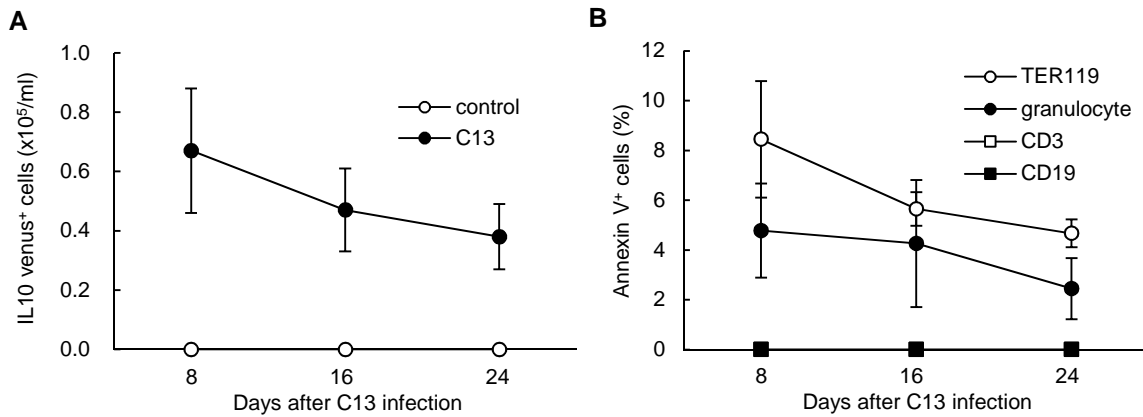


Figure S13. IL-10 production and apoptosis of TER119⁺ cells and granulocytes are detected in the late phase of C13 infection.

(A) The absolute number of cells expressing IL-10 Venus in the Mo-DCs from the PB of *Il10*^{Venus} reporter mice after C13 infection. **(B)** Percentage of annexin V⁺ cells in the indicated cells of PB of WT mice after C13 infection. Data represent the mean \pm s.d. of three independent experiments.

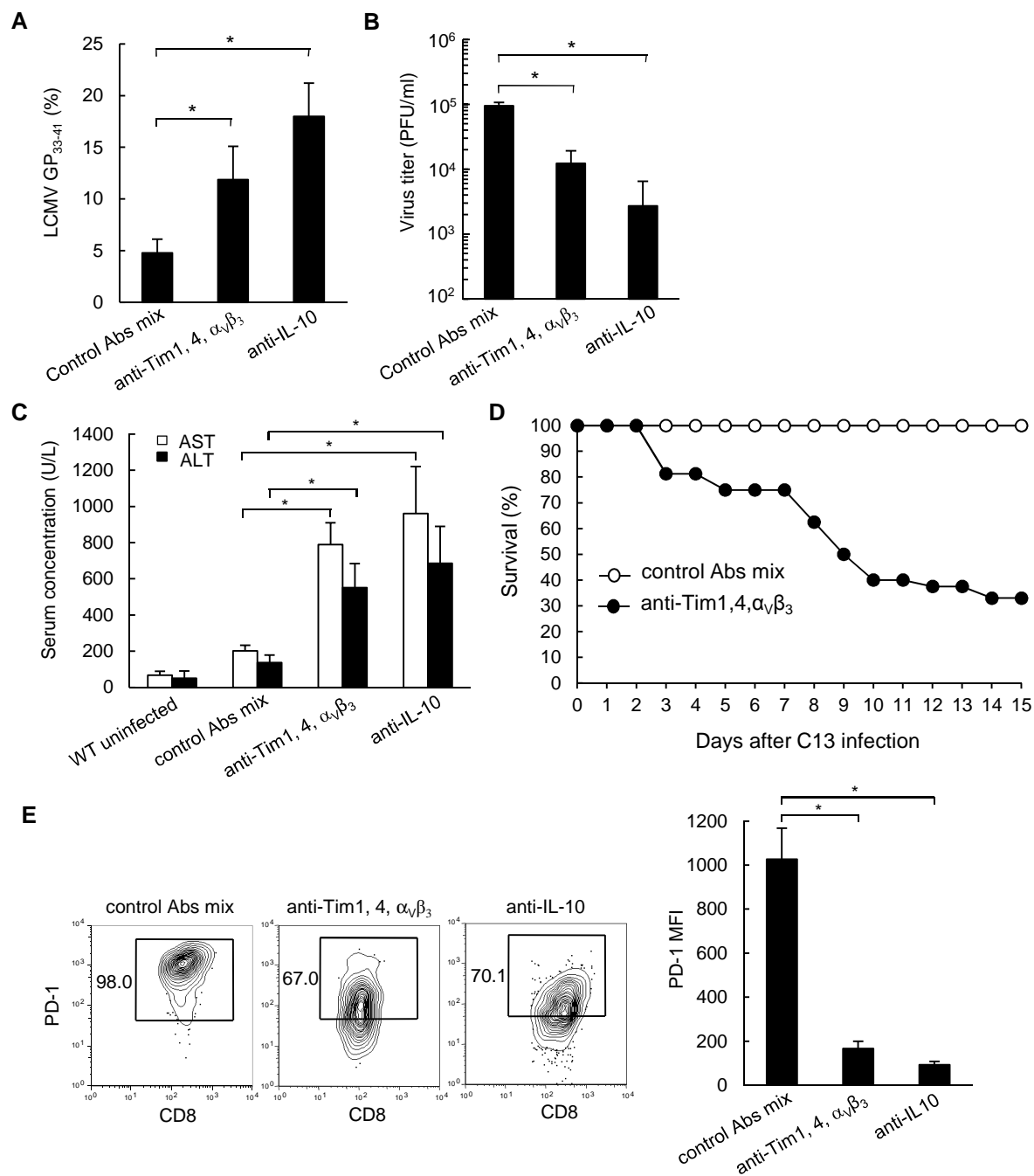


Figure S14. Altered immune responses in mice with limited hemophagocytosis and blocking of IL-10 production. (A-C) Percentage of LCMV GP₃₃₋₄₁-specific CD8⁺ T cells in the spleen (A), titer of LCMV C13 in the serum (B), and level of AST/ALT in the serum (C) of uninfected WT mice and WT mice 9 days after LCMV C13 infection, with or without the injection of Abs to block PS receptors or neutralize IL-10. (D) Survival (%) of WT mice after LCMV C13 infection with or without the injection of Abs to block PS receptors. (E) FCM profile of PD-1 expression on LCMV GP₃₃₋₄₁-specific CD8⁺ T cells in the spleen of WT mice 8 days after C13 infection, with or without injection of Abs to block PS receptors or IL-10-neutralizing Abs. MFI: mean fluorescence intensity. Data represent the mean \pm s.d. from three independent experiments. * $P < 0.01$.

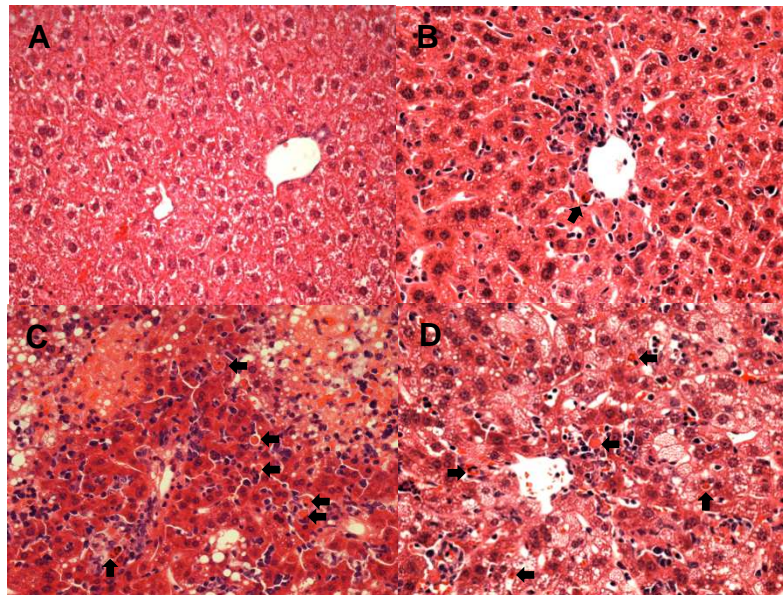


Figure S15. Inhibition of hemophagocytosis results in liver damage.

(A-D), HE staining of the liver from uninfected WT mice (A), WT mice infected with C13 (B), WT mice infected with C13 and treated with a mixture of antibodies against Tim1, Tim4, α_V , and β_3 (C), and WT mice infected with C13 and treated with anti-IL-10 Ab (D). Livers from C13-infected mice were prepared 9 days after infection. Many acidophilic bodies (apoptotic cells), indicated by arrows, are observed in (C) and (D). Data are representative from 5 mice. Original magnification, x 400.

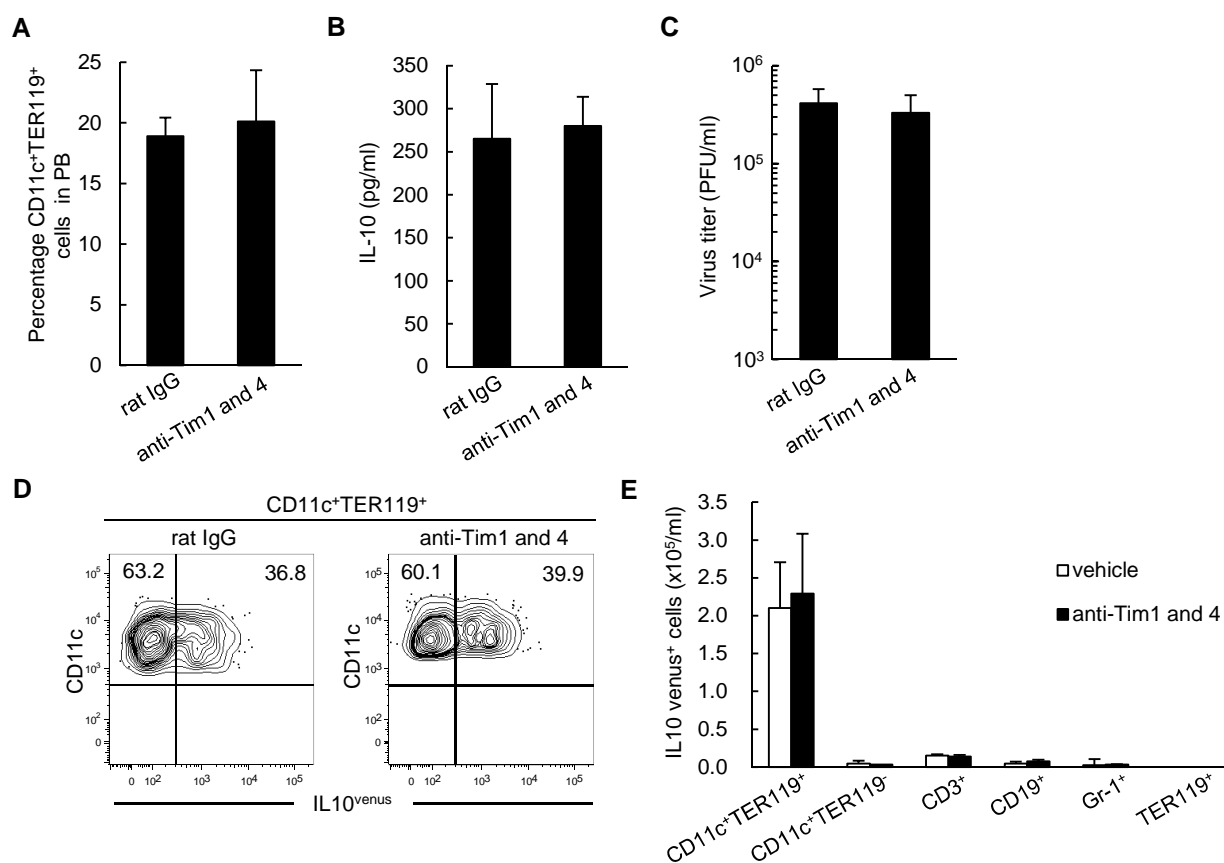


Figure S16. Administration of anti-Tim1 and 4 Abs does not alter immune responses in mice infected with LCMV C13. (A-C) Percentage of CD11c⁺TER119⁺ cells in the PB (A), IL-10 level (B), titer of LCMV C13 (C) in the serum of WT mice 9 days after LCMV C13 infection, with or without the injection of anti-Tim1 and 4 Abs. (D) A representative FCM profile of IL-10 Venus expression in CD11c⁺TER119⁺ cells from the PB of *Il10*^{Venus} reporter mice 24 h after C13 infection. (E) The cell number of IL-10 Venus⁺ in the indicated cell populations from the PB of *Il10*^{Venus} reporter and WT mice 1 and 9 days after C13 infection. Data represent the mean ± s.d. from three independent experiments. **P* < 0.01.

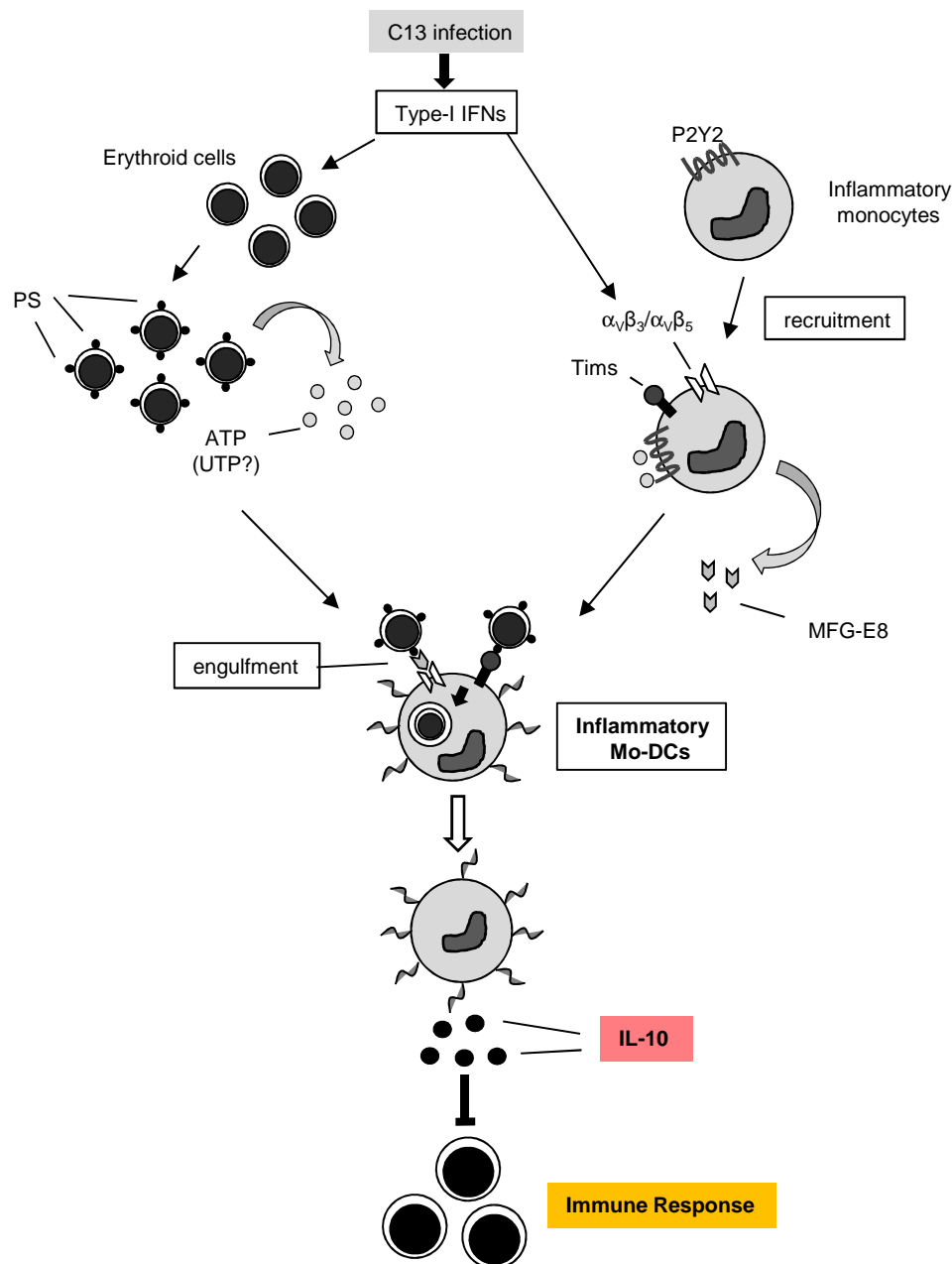


Figure S17. Schema of Hemophagocytosis development and Its Immunological Role.

Type I IFNs are produced immediately after C13 infection. The virus infection triggers both the activation of inflammatory monocytes and their differentiation into monocyte-derived dendritic cells (Mo-DCs), on which the cell-surface expression of phosphatidylserine (PS)-recognizing molecules ($\alpha_v\beta_3/\alpha_v\beta_5$ integrin, MFG-E8, and Tims) are upregulated; meanwhile, TER119⁺ erythroid cells begin to expose PS on their surface as an “eat me” signal. Both events are induced in a type I IFN-dependent manner. Then, Mo-DCs perform hemophagocytosis and release an immunoregulatory cytokine, IL-10, to fine-tune the immune responses, including CTL activity.