




Malaria abrogates O'nyong-nyong virus pathologies by restricting virus infection in nonimmune cells

Anthony Torres-Ruesta^{1,2}, Teck-Hui Teo¹, Yi-Hao Chan¹, Siti Naqiah Amrun¹ , Nicholas Kim-Wah Yeo¹ , Cheryl Yi-Pin Lee¹, Samantha Yee-Teng Nguetee¹ , Matthew Zirui Tay¹, Francois Nosten^{3,4} , Siew-Wai Fong¹, Fok-Moon Lum¹ , Guillaume Carissimo¹ , Laurent Renia^{1,5,6,7} , Lisa FP Ng^{1,2,8,9} 

O'nyongnyong virus (ONNV) is a re-emerging alphavirus previously known to be transmitted by main malaria vectors, thus suggesting the possibility of coinfections with arboviruses in endemic areas. However, the pathological outcomes of such infections remain unknown. Using murine coinfection models, we demonstrated that a preexisting blood-stage *Plasmodium* infection suppresses ONNV-induced pathologies. We further showed that suppression of viremia and virus dissemination are dependent on *Plasmodium*-induced IFN γ and are associated with reduced infection of CD45⁺ cells at the site of virus inoculation. We further proved that treatment with IFN γ or plasma samples from *Plasmodium vivax*-infected patients containing IFN γ are able to restrict ONNV infection in human fibroblast, synoviocyte, skeletal muscle, and endothelial cell lines. Mechanistically, the role of IFN γ in restricting ONNV infection was confirmed in *in vitro* infection assays through the generation of an IFN γ receptor 1 α chain (IFN γ R1)-deficient cell line.

DOI [10.26508/lsa.202101272](https://doi.org/10.26508/lsa.202101272) | Received 23 October 2021 | Revised 4 January 2022 | Accepted 4 January 2022 | Published online 17 January 2022

Introduction

O'nyongnyong virus (ONNV) is an enveloped, positive-sense, single-stranded RNA virus that belongs to the *Alphavirus* genus of the *Togaviridae* family (1). It is closely related to other arthritogenic alphaviruses from the Semliki Forest antigen complex, such as chikungunya virus (CHIKV), Ross River virus (RRV), and Mayaro virus (MAYV) (2). ONNV pathology in humans is characterized by fever, maculopapular skin rash, myalgia, incapacitating polyarthralgia and to a lesser extent lymphadenopathy (1). The disease is generally self-limiting and resolves within some days, but symptoms

such as polyarthralgia and myalgia may persist in a small proportion of the cases (3). ONNV was first isolated in Gulu, Uganda (4), during an outbreak that lasted 3 yr (1959–1962) and involved more than two million cases (5). After this outbreak, ONNV caused two other major epidemics: one in south-central Uganda in 1996 (3, 6) and another in Liberia and Ivory Coast in 2003 involving thousands of cases (7). More recently, epidemiological surveys have reported high seroprevalence of ONNV in Coastal Kenya (8) and Uganda (9), suggesting an underestimated burden of ONNV infections in sub-Saharan Africa.

The re-emergence and expansion of alphavirus infections in the tropics during the last decade introduce a new risk of coinfections with other highly prevalent endemic mosquito-borne diseases such as malaria. Limited serological studies in malaria-endemic countries in Africa have reported evidence of coinfections between *Plasmodium* parasites and alphaviruses. Specifically, anti-CHIKV antibodies were detected in a Nigerian cohort of *Plasmodium falciparum*-infected patients (10). In another study in Tanzania (11), a considerable proportion of febrile malaria patients were seropositive for CHIKV, suggesting pre-exposure or active CHIKV infection. However, these reports should be interpreted cautiously because antibodies against CHIKV and ONNV are highly cross-reactive (9).

ONNV and *Plasmodium* parasites share common anopheline vectors such as *Anopheles gambiae* and *Anopheles funestus* (1), thus increasing the likelihood of co-transmission. However, reports on ONNV and *Plasmodium* coinfections in humans are lacking despite the increasing rates of ONNV transmission and the overwhelming presence of malaria in Sub-Saharan Africa. This, together with the highly inflammatory signature of both infections (12, 13, 14, 15, 16), the detrimental role of T cell-mediated immunity in the pathologies induced by both infections (17, 18, 19, 20, 21), and the immunosuppressive nature of malaria (22, 23, 24, 25), strongly suggest that immune modulation could happen upon coinfection.

¹A*STAR Infectious Diseases Labs (A*STAR ID Labs), Agency for Science, Technology and Research (A*STAR), Singapore, Singapore ²Department of Biochemistry, Yong Loo Lin School of Medicine, National University of Singapore, Singapore, Singapore ³Shoklo Malaria Research Unit, Mahidol-Oxford Tropical Medicine Research Unit (MORU), Faculty of Tropical Medicine, Mahidol University, Mae Sot, Thailand ⁴Nuffield Department of Medicine, Centre for Tropical Medicine and Global Health, University of Oxford, Oxford, UK ⁵Singapore Immunology Network, Agency for Science, Technology and Research (A*STAR), Singapore, Singapore ⁶Lee Kong Chian School of Medicine, Nanyang Technological University, Singapore, Singapore ⁷School of Biological Sciences, Nanyang Technological University, Singapore, Singapore ⁸National Institute of Health Research, Health Protection Research Unit in Emerging and Zoonotic Infections, University of Liverpool, Liverpool, UK ⁹Institute of Infection, Veterinary and Ecological Sciences, University of Liverpool, Liverpool, UK

Correspondence: lisa_ng@idlabs.a-star.edu.sg; renia_laurent@idlabs.a-star.edu.sg

In this study, we describe the interactions between ONNV and rodent *Plasmodium* parasites in a mammalian host. Using mouse models of coinfections, we demonstrated that preexisting murine malaria restricts ONNV-associated pathologies and this protective effect is driven mainly by *Plasmodium*-induced IFN γ by limiting ONNV infection in the CD45⁺-cell compartment. In vitro experiments using human cell lines and plasma from *Plasmodium vivax*-infected patients confirmed the antiviral role of IFN γ in restricting ONNV infection. Our findings have potential implications in arbovirus and malaria control programs in endemic regions where *Plasmodium* parasites and arboviruses co-circulate.

Results

Murine malaria suppresses ONNV-induced joint swelling and viremia

An immunocompetent mouse model was previously established to recapitulate ONNV-induced joint pathologies (inflammation, edema, muscle necrosis, synovitis, and tenosynovitis) and acute viremia (26). Using this model, we first assessed whether a preexisting acute blood-stage *Plasmodium* infection could alter the development of ONNV pathologies. To do so, 3-wk-old C57BL/6J mice were inoculated with 1E6-infected red blood cells (iRBCs) from either *Plasmodium berghei* ANKA clone 231c11 (PbA), which induces lethal neuropathology known as experimental cerebral malaria (ECM), or the nonlethal self-resolving strain *Plasmodium yoelii* 17XNL clone 1.1 (Py17x). When patent parasitemia was detected at 4 d post iRBC injection, 1E6 ONNV PFU were injected subcutaneously in the right paw. Viremia and joint swelling were measured for 12 and 14 days postinfection, respectively (dpi) (Fig 1A). The patent blood-stage Py17x infection protected coinfecting animals from the development of ONNV-induced joint swelling (Figs 1B and S1A) and significantly reduced viremia levels (Fig 1C). Similarly, a preexisting blood-stage PbA infection was able to abolish virus-induced footpad swelling (Fig 1B). Of note, viremia in animals preinfected with PbA was undetectable during the entire follow-up suggesting a stronger protective effect by PbA than by Py17x against ONNV (Fig 1C).

To assess whether the suppression of viremia and ONNV-induced joint swelling by murine malaria was dependent on the timing of *Plasmodium* inoculation, two additional coinfection conditions were explored: concurrent coinfection and postviral coinfection. In concurrent coinfection, mice were coinfecting with 1E6 ONNV PFU and 1E6 PbA or Py17x iRBC at the same time (Fig 1D). Concurrent coinfection did not affect the development of viremia in coinfecting animals (Fig 1F) but had strain-specific effects on inflammation. Mice coinfecting with Py17x did not display significant reduction in joint swelling, whereas mice coinfecting with PbA displayed a significant reduction in joint swelling from 5 dpi onwards with a major suppression at the peak of swelling at 6 dpi and an earlier resolution of the pathology at 10 dpi compared with ONNV-infected controls (Figs 1E and S1B). Finally, we assessed postviral *Plasmodium* infection, where mice were infected with 1E6 PbA or Py17x iRBC 4 d after 1E6 ONNV PFU injection (Fig 1G). In this

setting, no effect on ONNV-induced joint swelling (Figs 1H and S1C) or viremia (Fig 1I) was observed, indicating that neither PbA nor Py17x infection could alter the course of a preexisting ONNV infection.

The effects of ONNV infection on the dynamics of malarial growth and survival were also assessed. The inoculation of ONNV 4 d post Py17x or PbA infection did not alter parasitemia levels (Fig S2A and B) or PbA-induced mortality (Fig S2B). Concurrent coinfection with nonlethal Py17x and ONNV resulted in increased parasitemia levels (Fig S2C). Py17x parasitemia resolution was delayed as coinfecting animals took 26–28 d to clear blood-stage parasites compared with 20–22 d in controls. On the other hand, simultaneous inoculation with PbA and ONNV did not affect the development of parasitemia or ECM mortality in coinfecting mice (Fig S2D). Finally, the infection with *Plasmodium* parasites 4 d post ONNV inoculation resulted in aggravated Py17x and PbA parasitemia (Fig S2E and F) but did not impact ECM mortality (Fig S2F).

Early stages of ONNV replication in footpad tissues are impaired in animals preinfected by *Plasmodium* parasites

As shown in Fig 1C, a preexisting *Plasmodium* infection, either by PbA or Py17x, significantly reduced viremia in coinfecting animals. To assess any possible differences in the kinetics of virus replication at the site of inoculation (right hind limb footpad), viral dissemination in vivo was assessed using a luciferase-tagged ONNV clone that mimics wild type ONNV infection in mice. Bioluminescence measurements from infected footpads correlate with viral burden in these tissues (26). ONNV-infected and coinfecting animals were monitored during the first 24 hours postvirus inoculation (hpi) given that ONNV dissemination peaks at 12 hpi (31). Significant differences in whole body and footpad radiance were detected as early as 1 hpi in mice preinfected with PbA or Py17x and were maintained at 3, 6, 12, and 24 hpi (Fig 2A–C and Video 1).

The absence of ONNV bioluminescence signals into adjacent tissues such as the tail (Fig 2C) in the coinfecting mice prompted us to further assess ONNV dissemination into other major mouse tissues. Appendages (hind limb footpads and tail), internal organs (spleen, liver, and pLN) and muscle tissue (gastrocnemius and quadriceps) of ONNV-infected and coinfecting animals were harvested at 1 dpi and viral RNA was quantified. Lower viral loads were observed in most of the tissues assessed, with major differences occurring in organs distant from the site of inoculation such as liver and spleen where viral burden in coinfecting mice were on average \approx 10,000-fold and \approx 1,000-fold lower, respectively (Fig 2D).

Preexisting *Plasmodium* infection renders CD45⁺ and CD45⁻ footpad cells less susceptible to ONNV infection

Our data strongly suggested that a preexisting blood-stage *Plasmodium* infection renders footpad cells less susceptible to ONNV. To identify the cellular subsets involved in the suppression of ONNV infection, we defined the differences in ONNV infection of non-immune (CD45⁻) and immune (CD45⁺) cells in footpads of coinfecting mice at 12 hpi (peak of footpad viral load in Fig 2B). For this purpose, a ZsGreen-tagged ONNV infectious clone detectable by flow cytometry under the FITC channel was used.

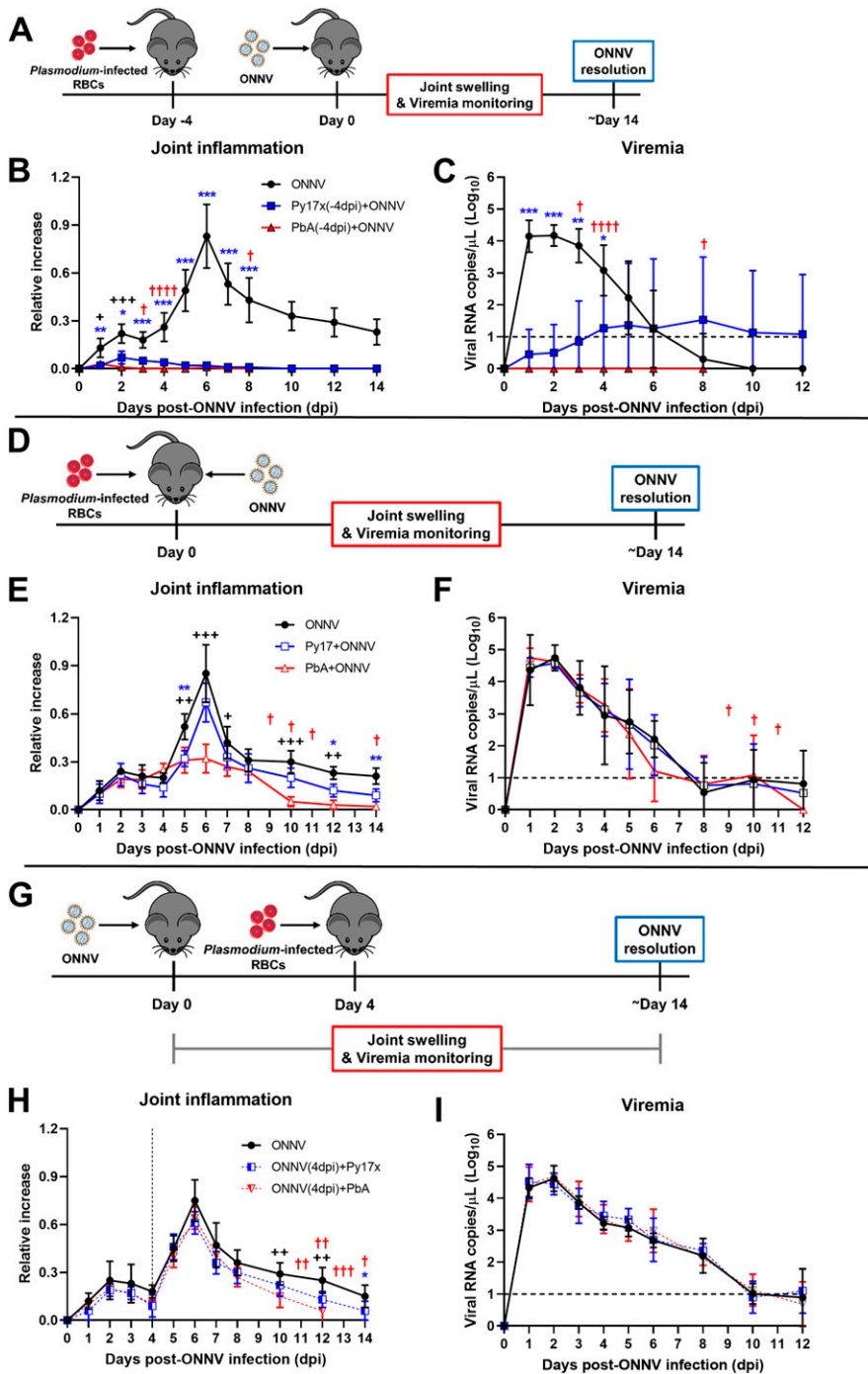


Figure 1. Preexisting murine malaria protects mice from ONNV-induced pathologies.

(A) Preval *Plasmodium* infection: mice were infected with PbA or Py17x 4 d prior ONNV inoculation according to the schematic in (A). **(B, C)** Joint swelling and (C) viremia measurements in ONNV, Py17x(-4dpi)+ONNV, and PbA(-4dpi)+ONNV groups. **(D)** Concurrent coinfection: animals were simultaneously infected with ONNV and PbA or Py17x on the same day according to the schematic in (D). **(E, F)** Joint swelling and (F) viremia measurements in ONNV, Py17x+ONNV, and PbA+ONNV groups. **(G)** Postviral *Plasmodium* infection: mice were infected with ONNV 4 d prior PbA or Py17x inoculation according to the schematic in (G). **(H, I)** Joint swelling and (I) viremia measurements in ONNV, ONNV(-4dpi)+Py17x, and ONNV(-4dpi)+PbA groups. Data are presented as mean \pm SD of at least five animals per experimental group and are representative of two independent experiments. Differences between ONNV controls and coinfecting mice with PbA ($++P < 0.01$, $+++P < 0.01$) or Py17x ($*P < 0.05$, $**P < 0.01$, $***P < 0.001$) were calculated using two-tailed Kruskal–Wallis and post hoc Dunn’s tests. When PbA-infected animals succumbed to ECM, differences between ONNV singly infected controls and coinfecting mice with Py17x were computed using two-tailed Mann–Whitney U test instead. “+” represents one mouse that succumbed of PbA-induced ECM on the respective day. Horizontal dashed lines in (C), (F) and (I) represent the qRT-PCR detection limit. Vertical dashed line in (H) represents the day on which *Plasmodium* parasites were inoculated.

To understand the individual contribution of CD45⁺ and CD45⁻ compartments to the total ONNV-infected cells at 12 hpi, the infectivity profile of footpads was analyzed in ONNV-infected mice. It was observed that ~80% of the ONNV-infected footpad cells at 12 hpi were part of the CD45⁻ nonimmune compartment (Fig S3A). When the analysis was extended to median fluorescence intensity values, CD45⁻ cells appeared to harbour on average a higher number of ONNV-ZsGreen copies than CD45⁺ cells (Fig S3B). These

observations highlight a major role of CD45⁻ cells as an early target for ONNV replication.

The impact of coinfections on the infectivity rates of CD45⁺ and CD45⁻ cells was then assessed. Animals preinfected with either PbA or Py17x displayed a marked reduction in the percentage and total counts of CD45⁺ZsGreen⁺ and CD45⁻ZsGreen⁺ cells compared with ONNV controls (Fig S3C and D). Given the importance of nonimmune cells as alphavirus targets during the early stages of infection

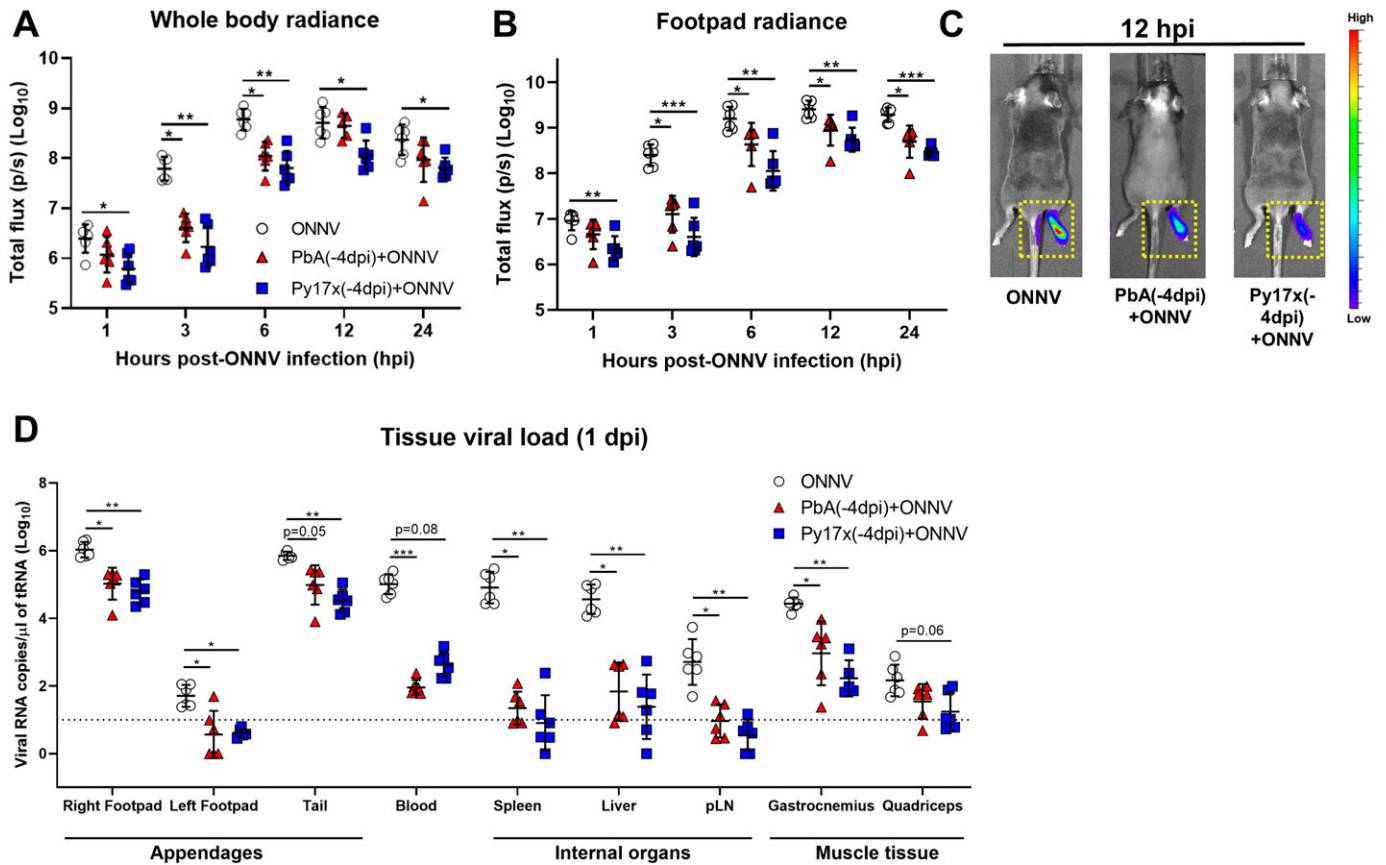


Figure 2. Early stages of ONNV replication and dissemination are suppressed by preexisting *Plasmodium* infection.

Mice were infected with Py17x or PbA and 4 d postinfection were inoculated with a firefly luciferase-tagged ONNV clone in the right hind limb footpad. Before data acquisition, mice were injected with 100 μ l of D-Luciferin (5 mg/ml) subcutaneously. (A, B) Whole body radiance and (B) footpad radiance of ONNV, Py17x(-4dpi)+ONNV, and PbA(-4dpi)+ONNV groups at 1, 3, 6, 12, and 24 hpi. (C) Representative pseudo-coloured images of bioluminescence readings depicting reduction of tissue viral load and restriction of viral dissemination in coinfecting animals at 12 hpi (yellow-dotted boxes). (D) Tissue viral load in mouse appendages, internal organs and muscle detected by qRT-PCR at 24 hpi. Data are presented as mean \pm SD of at least five animals per experimental group. Differences between ONNV controls and coinfecting mice with PbA or Py17x were calculated using two-tailed Kruskal–Wallis and post hoc Dunn’s tests (* $P < 0.05$ ** $P < 0.01$, *** $P < 0.001$).

(27, 28, 29), the differences in infectivity within the CD45⁺ compartment were further characterized using Uniform Manifold Approximation and Projection (UMAP) on the flow cytometry data. Manually gated CD45⁺ populations based on the expression of signature surface markers (30, 31, 32, 33, 34) were overlapped in the UMAP plot and allowed the identification of four major cell lineages: endothelial cells (CD45⁺CD29⁺CD31⁺), myoblasts (CD45⁺CD31⁺Sca-1⁺), fibroblasts (CD45⁺CD9⁺CD29⁺CD31⁺Sca-1⁻), and mesenchymal stem cells (CD45⁺CD9⁺CD29⁺CD31⁻Sca-1⁻) (Fig 3A). UMAP plots displaying the distribution of ZsGreen⁺ events in ONNV-infected and coinfecting mice are also shown in Fig 3A.

As observed from the UMAP plots, ONNV infection was globally suppressed in endothelial cells, myoblasts, and fibroblasts from coinfecting animals at 12 hpi (Fig 3A). A similar trend was observed when ZsGreen⁺ cells were quantified revealing a reduced number of infected endothelial cells, myoblasts, and fibroblasts (Fig 3B). To explore whether the reduced infectivity in CD45⁺ cells was a transient or maintained effect beyond 12 hpi, footpads from coinfecting mice with Py17x were harvested at 48 and 72 hpi and analyzed by flow cytometry. Coinfecting mice with PbA were not

included in these sets of experiments because of the high mortality rate induced by ECM. Interestingly, suppression of ONNV infection in coinfecting animals was observed across all the different time points assessed, suggesting a sustained suppression of viral load and not a delay in viral load appearance (Fig 3C).

Murine malaria up-regulates proinflammatory immune mediators in footpad tissues

Acute *Plasmodium* infections are known to induce strong proinflammatory responses characterized by the production and release of cytokines/chemokines and other mediators into the bloodstream to control parasite burden (15, 16). Thus, we hypothesized that some of these soluble factors could be responsible for the decreased susceptibility to ONNV infection observed in footpad tissues.

36 different immune mediators were assessed in sera (Fig 4A) and footpad tissue lysates (Fig 4B) of mice singly infected with lethal PbA or self-resolving Py17X at 4 d postparasite infection (time of ONNV inoculation in the coinfection model). Interestingly, results

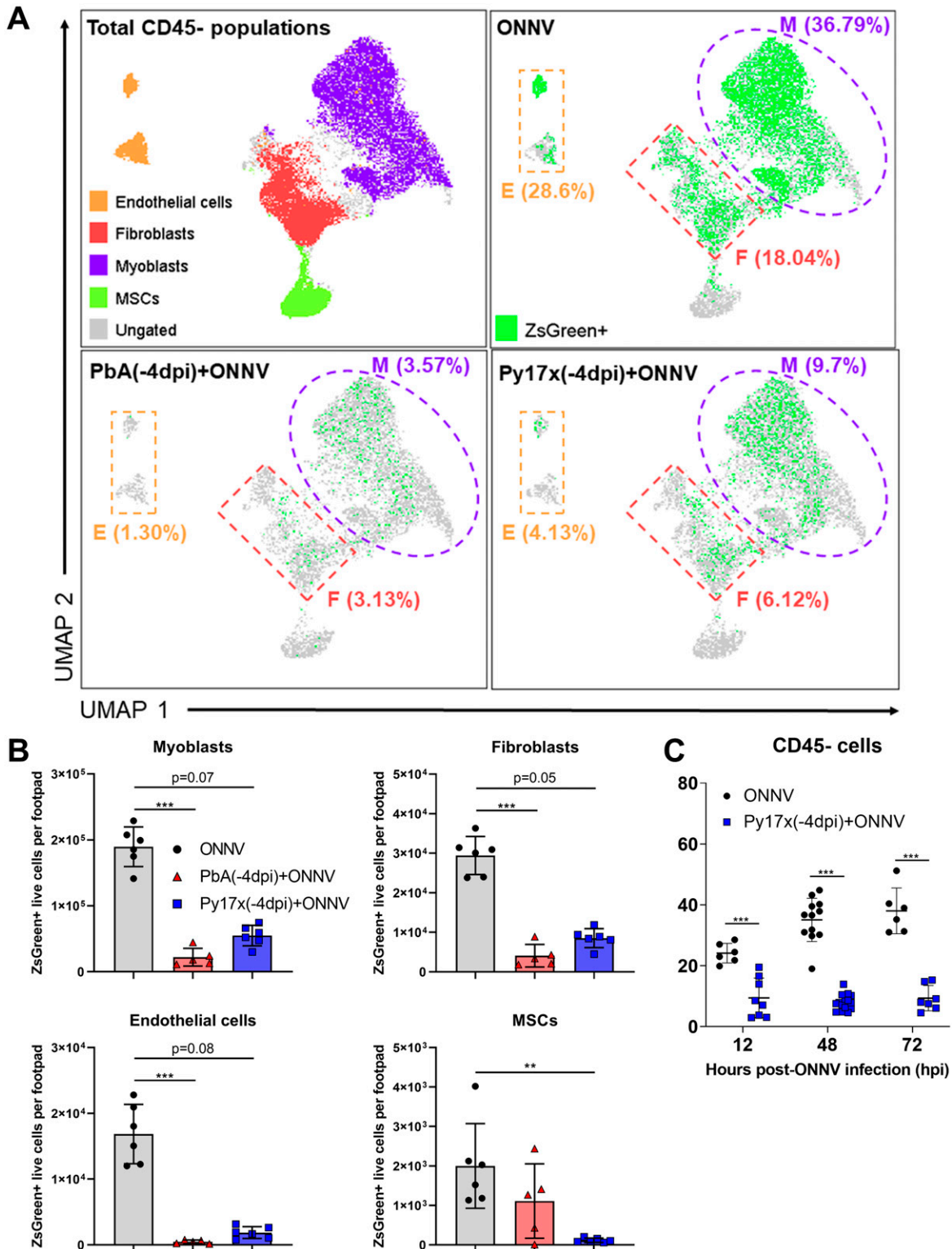


Figure 3. Prior *Plasmodium* infection restricts ONNV replication in CD45⁻ footpad cells.

(A) UMAP analysis of 105,000 live CD45⁻ footpad cells from naive, ONNV, PbA(-4dpi)+ONNV, and Py17x(-4dpi)+ONNV groups at 12 hpi. The UMAP plot was generated by concatenation of samples containing 5,000 randomly selected live CD45⁻ cells from each sample. ONNV, PbA(-4dpi)+ONNV, and Py17x(-4dpi)+ONNV UMAP plots show the global distribution of ZsGreen⁺ events (ONNV-infected cells). Colored dashed boxes highlight myoblasts (M), fibroblasts (F), and endothelial cells (E) and median ONNV infection rates per population. (B) Total counts of ONNV-ZsGreen⁺ cells in endothelial cells, myoblasts, fibroblasts and mesenchymal stem cells (MSCs) in ONNV, PbA(-4dpi)+ONNV, and Py17x(-4dpi)+ONNV groups at 12 hpi. (C) Percentages of infected CD45⁻ cells in ONNV and Py17x(-4dpi)+ONNV groups at 12, 48, and 72 hpi. Data are

showed that PbA- and Py17x-infected mice generated distinct pro-inflammatory cytokine profiles. Using principal component analysis on these cytokine profiles, a distinct separation was revealed between samples from mock (green), PbA-infected (red), and Py17x-infected mice (green) (Fig 4C).

Levels of pro-inflammatory IFN γ were increased by ~292- and ~28-folds in sera from PbA and Py17x-infected mice, respectively (Fig 4D), in line with previous reports (35, 36, 37). Up-regulation of chemokines such as CXCL10 (IFN γ -induced protein 10: IP-10), CXCL1, CCL2, CCL3, CCL5, and CCL7 was also observed in the sera (Fig 4D). Locally at the footpad, changes in immune mediator profiles were characterized by increased chemokine production, particularly IFN γ -induced CXCL10, which was found to be elevated by ~20- and ~11-folds in tissue lysates of PbA and Py17x-infected animals, respectively (Fig 4D). Other up-regulated chemokines shared between PbA and Py17x-infected mice were CCL2, CCL3, CCL4, CCL5, CCL7, and CCL11. In addition, IL-18 (IFN γ inducing factor: IGIF) and IFN γ were found to be slightly elevated in footpad tissues upon PbA and Py17x infection (Fig 4D).

Differentially regulated cytokines/chemokines in the footpads were subjected to STRING analysis (performed with a high confidence threshold of 0.9), revealing key interactions between eight of these immune mediators (Fig S4). The predicted interactions are linked to cellular responses to IFN γ , host-negative regulation of viral transcription and leukocyte recruitment and activation. Collectively, these data suggest that *Plasmodium* infection (either by PbA or Py17x) triggers a systemic immune response characterized by the up-regulation of pro-inflammatory immune mediators not only in the blood but also locally in footpad tissues. Importantly, the increased levels of immune mediators linked to IFN γ signaling, a known antiviral cytokine (38), in footpad lysates of *Plasmodium*-infected mice suggested a possible involvement of this cytokine in the restriction of ONNV infection.

***Plasmodium*-induced IFN γ mediates the suppression of ONNV replication and dissemination in coinfecting animals**

To assess the potential role of IFN γ in suppressing ONNV pathologies, IFN γ -deficient animals were infected with nonlethal Py17x. 4 d postinfection, luciferase-tagged ONNV was inoculated subcutaneously in the right footpad. Lack of IFN γ in coinfecting animals abolished the antiviral effects exerted by Py17x infection in mice. Bioluminescence readings at 3, 6, 12, and 24 hpi were comparable with those from IFN γ -deficient animals singly infected with ONNV (Fig 5A and Video 2). Similarly, ONNV viremia in coinfecting IFN γ -deficient mice was also restored (Fig 5B) despite a suppression of joint swelling at 6 dpi (Fig S5). This suggests that coinfection might modulate other immune responses leading to joint swelling suppression in the absence of IFN γ .

Because nonimmune cells express IFN γ receptors (38) and can respond to IFN γ signaling, the infection profile of these subsets was assessed in coinfecting mice deficient in IFN γ . Likewise, ONNV infectivity profiles of CD45⁺ cells were restored in myoblasts, endothelial

cells, fibroblasts, and mesenchymal stem cells (Fig 5C and D). To further validate these findings, in vivo IFN γ neutralization was performed using anti-mouse IFN γ antibodies. Coinfecting animals treated with anti-mouse IFN γ displayed comparable ONNV tissue viral loads at 3, 6, 12, and 24 hpi (Fig 5E and Video 3) than isotype-control treated mice, suggesting a major role of IFN γ in the antiviral effects exerted by preexisting *Plasmodium* infections.

Blood-stage *Plasmodium* infections in humans and mice are known to induce the production of type I IFN responses (39, 40, 41). Type I IFN is a major regulator of susceptibility to alphavirus infection (13, 42, 43) and evidence has suggested crosstalk mechanisms with IFN γ signaling (44, 45, 46). To evaluate any possible contribution of type I IFN responses in the reduced susceptibility to ONNV upon coinfection, the effect of preexisting murine malaria on ONNV replication was assessed in IFN α / β receptor-deficient mice (deficient of IFN- α / β receptor). Viremia measurements at 12, 24, and 48 hpi in coinfecting IFN α / β receptor-deficient mice (Fig S6) revealed that murine malaria was still able to restrict ONNV infection, ruling out the involvement of type I IFN responses in the antiviral effects exerted by *Plasmodium*-induced IFN γ .

IFN γ inhibits ONNV infection in human fibroblast, synoviocyte, skeletal muscle, and endothelial cell lines

The antiviral role of IFN γ in nonimmune cell lineages upon ONNV infection remains poorly defined. To extrapolate our findings in the human context, the antiviral effect of IFN γ was assessed in human cell lines representing skin fibroblasts (BJ), synoviocytes (SW982), endothelial cells (HPMECs), and skeletal muscle cells (RD) in an in vitro infection system. Interestingly, each cell type displayed different susceptibility to ONNV infection. Particularly, skin fibroblast and synoviocyte cell lines are highly susceptible to ONNV, whereas endothelial cells and skeletal muscle cells are poorly infected. Nevertheless, regardless of cell type, IFN γ pretreatment successfully reduced ONNV infection in BJ, SW982, HPMEC, and RD cells at 24, 48, and 72 hpi in a dose-dependent manner (Fig 6A).

Finally, we characterized the ability of plasma samples from acute *P. vivax*-infected patients to render human cell lines less susceptible to ONNV infection. Plasma IFN γ levels were quantified, and samples were categorized and pooled into low (n = 13, median IFN γ concentration = 69.53 pg/ml) and high IFN γ producers (n = 14, median IFN γ concentration = 293.675 pg/ml). 10 plasma samples from healthy individuals were pooled and included in the experiments as controls (IFN γ concentration under the quantification limit) (Fig 6B). Incubation of skin fibroblasts (BJ) with plasma from either low or high IFN γ producers reduced ONNV infection compared with those treated with healthy control plasma (Fig 6C). To prove that IFN γ present in the plasma samples from acute *P. vivax*-infected patients was responsible for the antiviral effects exerted in vitro, we generated a HEK293T cell line with impaired IFN γ signaling (Fig S7A and B) by knocking down the expression of the IFN γ receptor 1 α chain (IFN γ R1). Upon treatment with plasma from malaria patients, HEK293T cells with intact IFN γ R1 expression

presented as mean \pm SD of at least five animals per experimental group. Differences between ONNV controls and coinfecting mice with PbA or Py17x were calculated using two-tailed Kruskal–Wallis and post hoc Dunn's tests (** P < 0.01, *** P < 0.001).

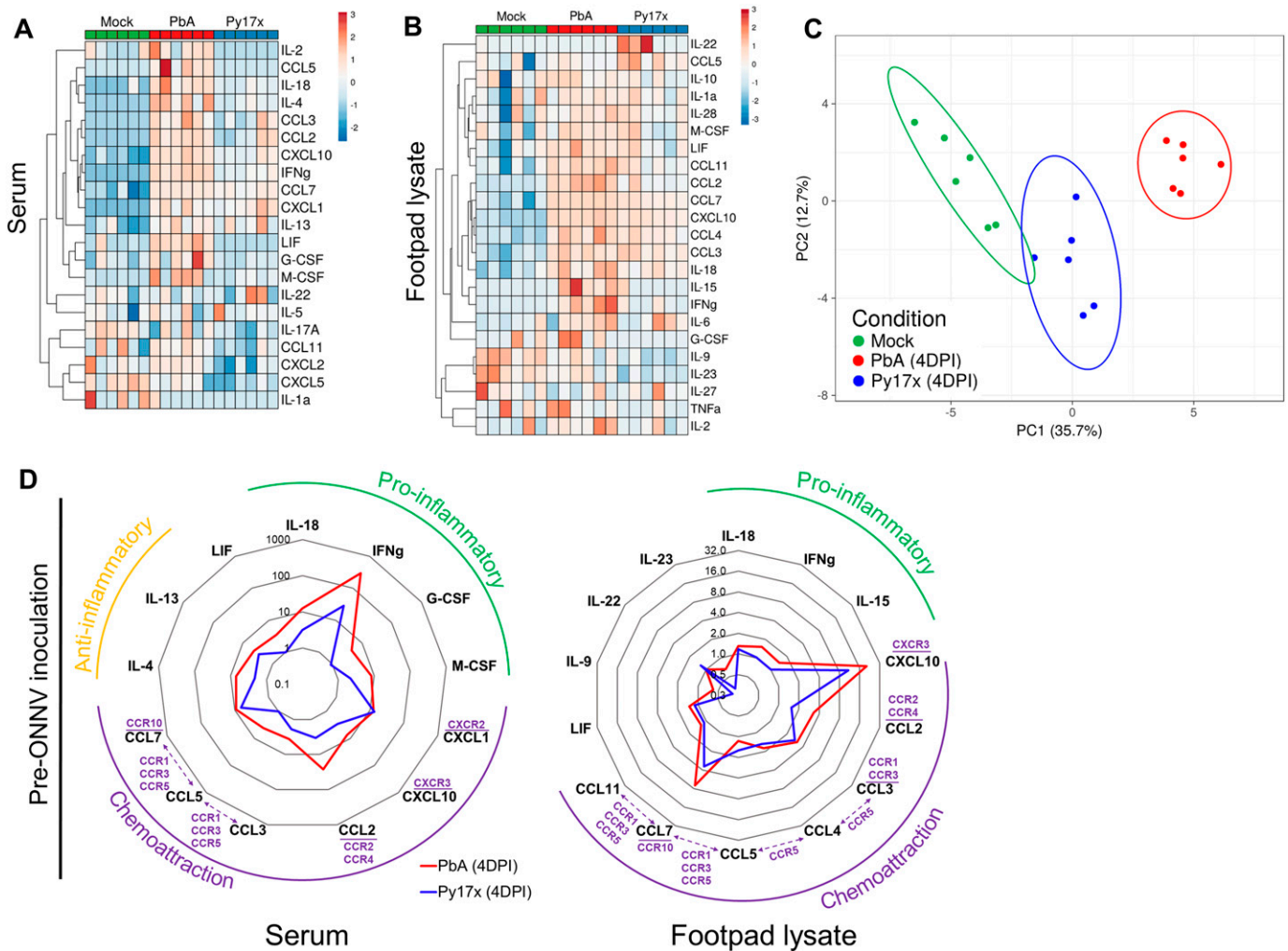


Figure 4. Footpads of *Plasmodium*-infected mice display a pro-inflammatory milieu.

(A, B) Heat map plots showing the detected cytokines/chemokines in (A) serum and (B) footpad lysates of mock-infected (green), PbA-infected (red), and Py17x-infected (blue) mice at 4 dpi. Analyte concentrations (pg/ml + 1) were logarithmically transformed (\log_{10}) and Z-scores were calculated for representation purposes. Principal component analysis (PCA) and heat map plots were constructed using ClustVis. (C) PCA using differentially expressed analytes in footpad lysates and sera of mock, PbA-infected (4 dpi), and Py17x-infected (4 dpi) groups. PCA plot shows that PC1 (responsible for 35.7% of the variation) and PC2 (responsible for 12.7% of the variation) segregate the populations in three clusters: mock (green), PbA-infected (red), and Py17x-infected (blue). Colored ellipses were calculated with 95% confidence levels. (D) Radar plots showing median fold changes of differentially expressed cytokines/chemokines in serum and footpad lysates of PbA-infected (4 dpi) and Py17x-infected (4 dpi) groups relative to mock animals. Each cytokine/chemokine is grouped according to its immunological function (green: pro-inflammatory, yellow: anti-inflammatory) or homing receptors (purple) as indicated. Shared chemokine receptors are shown in dashed lines. Data correspond six animals per experimental group. Differences between naive, PbA, or Py17x-infected mice calculated using two-tailed Kruskal-Wallis and post hoc Dunn's tests.

displayed lower ONNV infection compared with untreated controls. This antiviral effect was lost in cells with impaired IFN γ R1 expression (Δ IFN γ R1) (Fig 6D), in agreement with an IFN γ -specific effect.

Discussion

O'nyongnyong virus and *Plasmodium* parasites share common anopheline vectors and co-circulate in sub-Saharan Africa with risk of human coinfection. This is the first study investigating the pathological outcomes of coinfection by *Plasmodium* parasites and alphavirus ONNV in a mammalian host. Here, we showed that a preexisting murine *Plasmodium* infection is able to suppress the

development of ONNV pathologies by restricting viral infection at the site of inoculation and dissemination to distant organs. We demonstrated that *Plasmodium*-induced IFN γ is the main cytokine driving the antiviral effects observed.

IFN γ is a pleiotropic cytokine known for its ability to regulate immune responses by promoting macrophage activation, enhancing antigen presentation, modulating helper T cell development and mediating viral and bacterial immunity, among others (47). In ONNV-infected animals, it was observed that nearly 80% of the total virus-infected cells at 12 hpi (Fig S3A) belonged to the CD45⁺ compartment corroborating previous observations from other closely related alphaviruses, whereby non-immune cells support the early stages of viral replication (13, 27, 28, 29, 48). In contrast, animals harboring a *Plasmodium* infection displayed

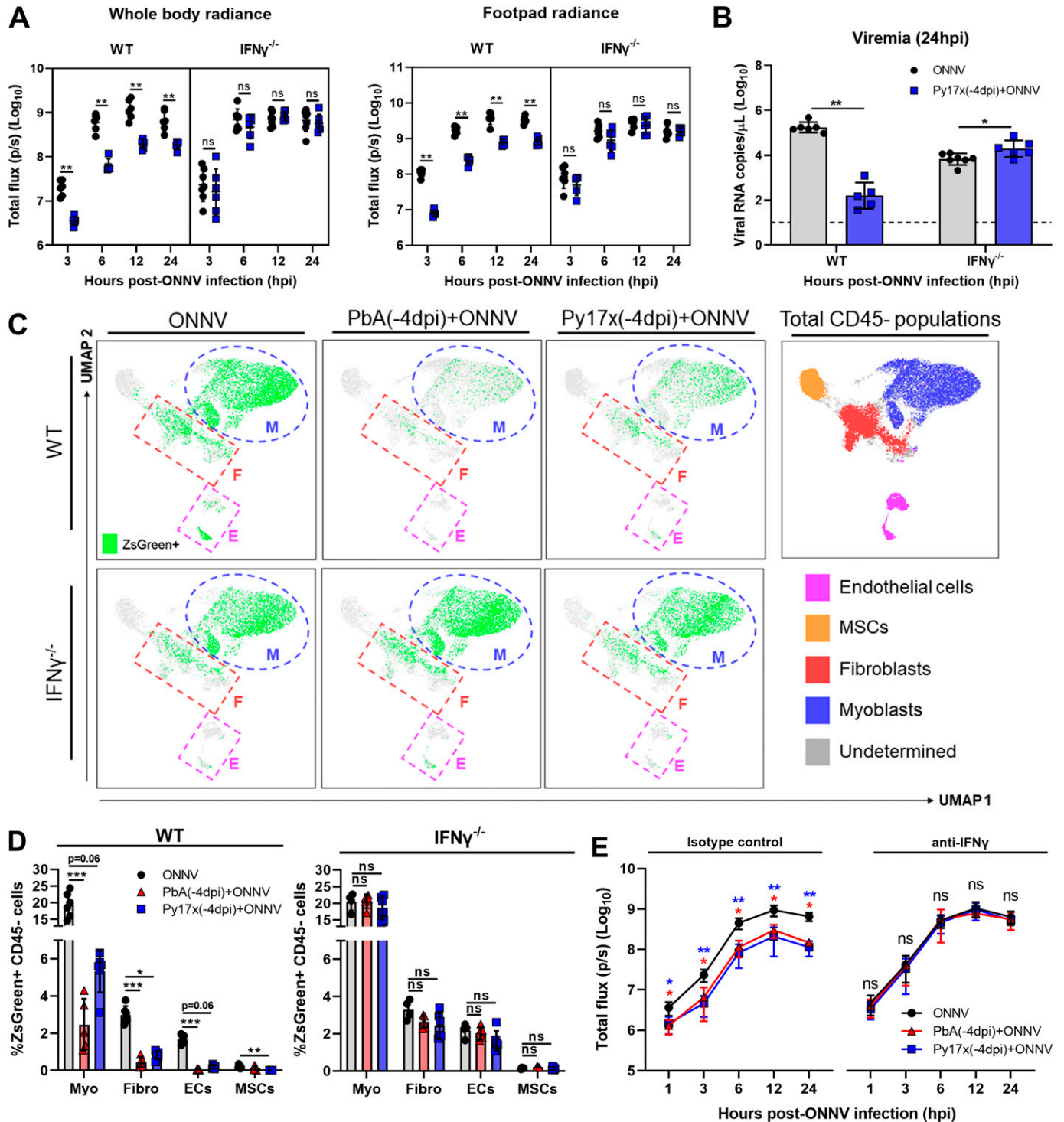


Figure 5. Plasmodium-induced IFN γ mediates the suppression of ONNV replication and dissemination in coinfecting animals.

(A, B) In vivo luminescence readings of (A) whole body and footpad radiance at 3, 6, 12, and 24 hpi and (B) viremia at 24 hpi of ONNV and Py17x(-4dpi)+ONNV wild-type (WT), or IFN γ -deficient (IFN γ ^{-/-}) animals. (C) UMAP analysis of 160,000 live CD45⁺ footpad cells from WT or IFN γ ^{-/-} ONNV, PbA(-4dpi)+ONNV, and Py17x(-4dpi)+ONNV groups at 12 hpi. The UMAP plot was generated by concatenation of samples containing 5,000 randomly selected live CD45⁺ cells from each sample. ONNV, PbA(-4dpi)+ONNV, and Py17x(-4dpi)+ONNV UMAP plots show the global distribution of ZsGreen⁺ events (ONNV-infected cells). Colored dashed boxes highlight ONNV infection in myoblasts, M; fibroblasts, F and endothelial cells, E. (D) Frequency of CD45⁺ ZsGreen⁺ footpad cells of WT or IFN γ ^{-/-} ONNV, PbA(-4dpi)+ONNV, and Py17x(-4dpi)+ONNV groups at 12 hpi. Myo, myoblasts; Fibro, fibroblasts; ECs, endothelial cells and MSCs, mesenchymal stem cells. (E) Footpad radiance at 1, 3, 6, 12, and 24 hpi of ONNV, PbA(-4dpi)+ONNV, and Py17x(-4dpi)+ONNV groups in animals treated with mouse anti-IFN γ or isotype control. Data are presented as mean \pm SD of at least five animals per experimental group. (A, B) Two-tailed Mann-Whitney U test was used to compute differences between ONNV and Py17x(-4dpi)+ONNV groups in (A) and (B) (* P < 0.05 ** P < 0.01). Differences

reduced numbers of ONNV-infected myoblasts, fibroblasts and endothelial cells and this protective effect was reverted in IFN γ -deficient mice or upon in vivo IFN γ neutralization. We therefore hypothesize that production of IFN γ in response to acute blood-stage *Plasmodium* infection could stimulate cells from the CD45⁻ compartment activating antiviral processes (Fig S4) which in turn restrict a subsequent ONNV infection. In line with this, the antiviral effects of IFN γ have been observed in other alphaviruses such as Sindbis virus (SINV). In vitro studies suggested that IFN γ affects SINV replication in mature neurons by interfering with the synthesis of genomic and sub-genomic viral RNA (49) and that this effect is dependent on JAK/STAT signaling (50). The contribution of *Plasmodium*-induced type I IFN (40, 41) to the reduced susceptibility to ONNV infection was also assessed in IFN α R^{-/-} mice. Considerable viremia differences were observed between ONNV-infected wild type and IFN α R^{-/-} controls (~4–5 Log₁₀ at 48 hpi) highlighting the importance of IFN- α/β signaling in the control of ONNV infection as observed in other alphavirus animal models (42, 43, 51, 52). Nonetheless, type I IFN-induced upon *Plasmodium* infection seems to be negligible for the establishment of protective effects by murine malaria as coinfecting IFN α R^{-/-} mice still displayed reduced ONNV infection to a comparable level than coinfecting wild-type mice (Fig S6).

It is important to note that in our experiments, mice only experienced suppression of ONNV viremia and virus dissemination after 4 d post-*Plasmodium* inoculation (Figs 1B and 2A–D) and not upon concurrent or sequential (postviral) coinfection. These observations strongly suggested that the timing of parasite inoculation and induction of IFN γ are critical for the protective effects to happen. Interestingly, although the main suppression of ONNV infection occurred in joint footpad cells, we observed lower concentrations of IFN γ in joint footpad tissues compared with serum samples at 4 d post-*Plasmodium* inoculation. Thus, it is likely that IFN γ levels in joint footpads could have increased at an earlier time point. In support of this, we observed high concentrations of IFN γ -induced immune mediators in joint footpad tissues, particularly CXCL10 (53) and CCL7, known to be produced by fibroblasts and mononuclear cells upon IFN γ stimulation (54). The development of T-cell responses, major IFN γ -producing subsets during malaria (ref), could also influence the outcome of a *Plasmodium*–ONNV coinfection in murine models. Early in a blood-stage infection, a large number of IFN γ -secreting Th1 cells are produced, whereas Th2-like responses govern during the chronic phase of infection (17). Because ONNV inoculation occurs in the early stages of murine malaria (4 dpi), it is likely that the antiviral effects of IFN γ are associated to the establishment of Th1 immunity against the parasite. It can be speculated that the degree of virus suppression might differ if ONNV is inoculated during the chronic stage of the murine malaria, particularly when Th1 responses are waning.

We explored the relevance of our findings in the context of ONNV human infection by treating four human cell lines from different nonimmune lineages (fibroblast, synovial cell, endothelial, and skeletal muscle cells) with IFN γ before ONNV infection. Mechanistically, our results demonstrated that IFN γ is able of restricting ONNV

infection in human cell lines in a dose dependent manner. Similarly, we also observed that stimulation of skin fibroblasts with plasma from acute *P. vivax*-infected patients containing IFN γ -reduced cell susceptibility to ONNV infection. Of note, the biological effects of IFN γ are mediated conventionally through the activation of the JAK/STAT pathway (55). We postulate that the attenuation of ONNV infection by *Plasmodium*-induced IFN- γ observed in mouse models could be translated to a real coinfection scenario in endemic populations given that the JAK/STAT signaling pathway between humans and mice is highly conserved (56, 57). This suggests that similar downstream effector proteins could be involved in the IFN γ -mediated restriction of alphavirus infection.

A recent study (58) showed that *Plasmodium* infection protected mice from Ebola virus (EBOV)-induced mortality via up-regulation of IFN γ , supporting field reports where coinfecting patients by EBOV and *P. falciparum* displayed increased survival rates (59). Conversely, two other murine coinfection models with respiratory viral pathogens such as murine pneumonia virus (PVM) and murine gammaherpesvirus 68 (MHV68) using nonlethal *Plasmodium chabaudi* and *P. yoelii* 17XNL have reported detrimental outcomes for the host such as increased viral loads in the lungs (60) and mortality due to severe anemia (61). These observations were linked to altered type I IFN production (60) and antiviral humoral responses (61) upon coinfection. Thus, the protective or detrimental effects of murine malaria on viral pathogens are likely associated to the modulation of distinct immune responses governing the control of different viral infections. Conversely, other protozoan parasites highly prevalent in the tropics and known to induce the up-regulation of IFN γ in response to infection such as *Leishmania* spp. (62), *Toxoplasma* spp. (63), or *Trypanosoma* spp. (64) could potentially display similar protective effects in coinfection settings. In line with this, mice infected with *Trypanosoma brucei* were shown to be resistant to cutaneous leishmaniasis through the induction of IFN γ generating a hostile pro-inflammatory environment impairing *L. major* colonization of the skin (65).

Results in Fig 1B and E suggest the impairment of ONNV-induced joint pathologies upon *Plasmodium* infection. Interestingly, concurrent coinfection with PbA parasites did not affect viremia levels but significantly reduced the major peak of footpad swelling at 6 dpi. This can be the result of two different mechanisms. First, the development of viremia in concurrently coinfecting mice can be attributed to the absence of *Plasmodium*-induced IFN γ in the early stages of ONNV infection in the footpads. It has been shown that the earliest IFN γ production during blood-stage murine malaria only occurs after 24 h postparasite injection (35). During this period, footpad cells in concurrently coinfecting mice are still susceptible to ONNV infection which results in viremia levels similar to control mice (Fig 1F). On the other hand, the suppression of joint swelling upon coinfection could be linked to the dysregulation of virus-specific CD4 T-cell responses, main drivers of joint inflammation at 6 dpi (26) by malaria. It has been reported that murine *Plasmodium* infections impair the development of CD4 T-cell responses against heterologous antigens (66, 67). Thus, PbA infections could alter the establishment of virus-specific CD4 T-cell immunity resulting in decreased footpad swelling.

between ONNV, PbA(-4dpi)+ONNV, and Py17x(-4dpi)+ONNV groups were calculated using two-tailed Kruskal–Wallis and post hoc Dunn's tests (* $P < 0.05$ ** $P < 0.01$, *** $P < 0.001$).

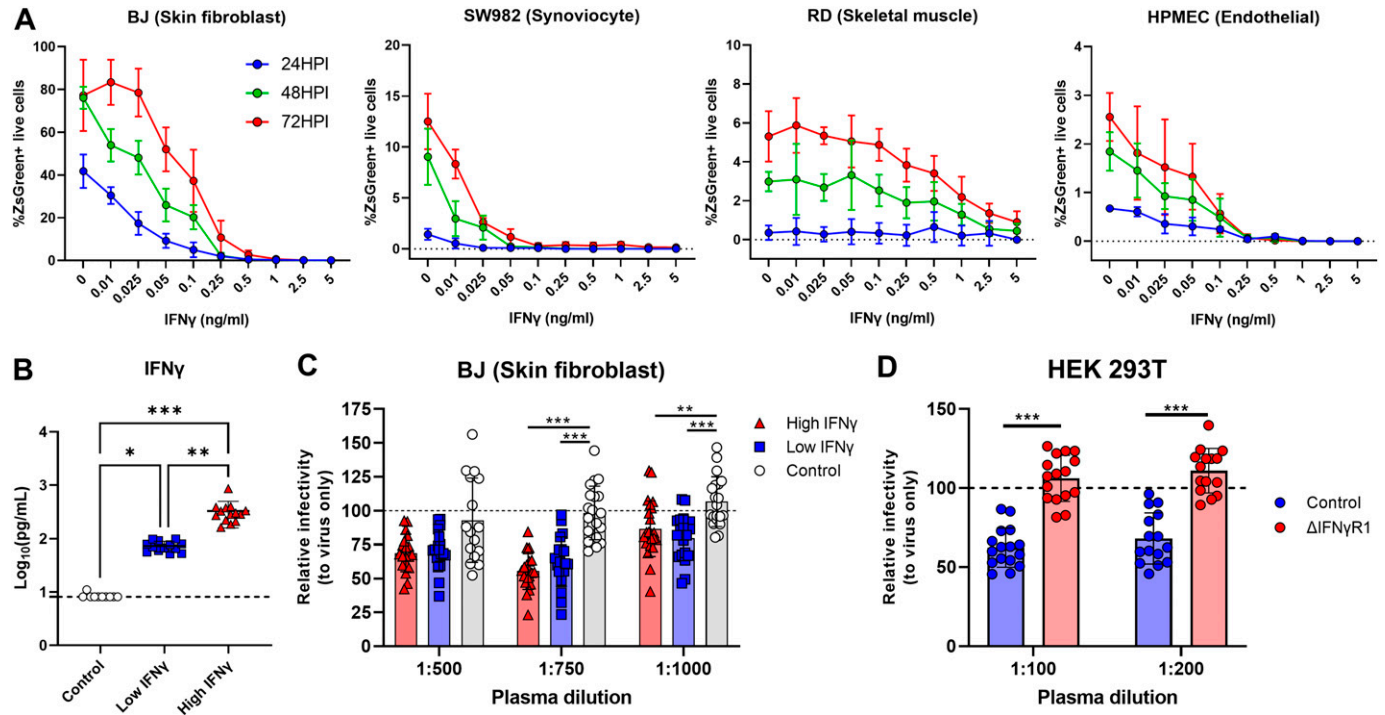


Figure 6. In vitro stimulation with human IFN γ or plasma from malaria patients reduces susceptibility to ONNV infection. For (A), cells were treated with recombinant human IFN γ for 24 h before ONNV infection at MOI 10. (A) ONNV infection rates in BJ, SW982, HPMEC, and RD at 24, 48, and 72 hpi. (B) IFN γ levels in plasma of healthy controls (HC, n = 10), low (n = 13), and high (n = 14) IFN γ responders. For (C), BJ cells were treated with pooled plasma dilutions from *P. vivax*-infected patients or healthy controls for 12 h before ONNV infection at MOI 1. For (D), control or Δ IFN γ R1 HEK293T cells were treated with pooled plasma (1:100 or 1:200) from *P. vivax*-infected patients for 12 h before ONNV infection at MOI 1. Differences between three groups were calculated using two-tailed Kruskal–Wallis and post hoc Dunn’s tests, differences between two groups were calculated using two-tailed Mann–Whitney U test (* $P < 0.05$ ** $P < 0.01$, *** $P < 0.001$). Data are presented as mean \pm SD values and representative of two independent experiments.

The great success of malaria control programs during the last decade has been accompanied by a sharp rise in the number of arbovirus infections worldwide (68). Our data highlight a possible causative link: that this phenomenon could be due in part to the loss of protective effects exerted by *Plasmodium* infections on alphavirus-induced pathologies that had hitherto masked the real burden of ONNV and other arbovirus infections. This may occur via two mechanisms: first, higher arbovirus case reporting because of more severe symptoms of arbovirus infections in the absence of malaria; second, and worse, increased arbovirus cases might reflect increased arbovirus transmissibility and/or fitness because of increased viral titers in the absence of malaria. Our data in this report will be of value in the fight against *Plasmodium* and ONNV infections in areas where both pathogens co-circulate, particularly highlighting the need for screening and clinical studies of underlying alphavirus infections in malaria intervention programs.

Materials and Methods

Mice

3- to 4-wk-old gender-matched wild-type (JAX #000664), IFN γ ^{-/-} (JAX #002287), and IFN α 1^{-/-} (JAX #028288) mice in C57BL/6J background were used in this study. Animals were bred and maintained under

specific pathogen-free conditions at the Biological Resource Centre of the Agency for Science, Technology, and Research, Singapore (A*STAR).

Viruses

The IMTSSA/5163 ONNV isolate used in this study was obtained from an acute patient in Chad in 2004 (kindly provided by Marc Grandadam from the Unité de Virologie Tropicale, IMTSSA) (69). Full-length infectious cDNA clones of the IMTSSA/5163 isolate were used to generate ONNV variants expressing the firefly luciferase gene (ONNV-Fluc) and ZsGreen protein (ONNV-ZsGreen) (70, 71). Viruses were propagated in *Aedes albopictus* C6/36 cell line (ATCC CRL-1660) and purified by sucrose-gradient ultracentrifugation. Viral stock titers were determined by standard plaque assay using Vero E6 cells (ATCC CCL-81).

Parasites

P. yoelii 17XNL clone 1.1 (referred as Py17x) was used to induce self-resolving infections in mice (72). Lethal infections were induced by inoculation of *P. berghei* ANKA (PbA) (231c11) expressing luciferase and GFP under the control of the e1- α promoter (73). iRBCs were obtained by in vivo serial passage in C57BL/6J mice and were stored in Alsever’s solution in liquid nitrogen.

Human cell lines

BJ (ATCC CRL-2522), SW-982 (ATCC HTB-93), and RD (ATCC CCL-136) cells were grown in DMEM medium with 10% heat-inactivated FBS. HPMEC (#3000; ScienCell) were grown in supplemented EC medium (#1001; ScienCell). Cells were grown at 37°C, relative humidity of 95%, and 5% CO₂.

Generation of Δ IFN γ R1 HEK293T cell line

Δ IFN γ R1 HEK293T cell line was generated by phosphorylating and annealing primers (5'-CACCGACATGAACCTATCGTATAT-3') and (5'-AAACATATACGATA GGGTTCATGTC-3') (NG_007394.1) using T4 Polynucleotide Kinase (NEB M0201S) in provided buffer supplemented with 1 mM ATP (A2383; Sigma-Aldrich) at 37°C for 30 min followed by 5 min at 95°C and ramped down to 25°C. Annealed primers were then ligated using Instant Sticky-end Ligase Master Mix (M0370; NEB) in pSB-CRISPR (kindly gifted by Dr Gao and Dr Hu (74)) previously linearized using Esp3I (R0734S; NEB) in CutSmart Buffer (NEB) and purified using Nucleospin gel and PCR clean up kit (740609.50; Macherey-Nagel). HEK293T cells were co-transfected with pCMV(CAT)T7-SB100 (a gift from Zsuzsanna Izsvak, plasmid # 34879; Addgene; RRID: Addgene_34879 (75)) and pSB-CRISPR (at 1:1 ratio) using Lipofectamine 2000 (11668019; Thermo Fisher Scientific) in Opti-MEM medium (31985070; Gibco) following the manufacturer's recommendations. 3 d post co-transfection, cells were passaged and cultured in complete media containing 1 μ g/ml puromycin (P8833; Sigma-Aldrich). When control cells fully succumbed to puromycin selection, co-transfected cells were cultured in complete media and IFN γ R1 expression was assessed by flow cytometry.

ONNV infection and disease monitoring

Mice were infected with ONNV by subcutaneous inoculation of 10⁶ PFU in 30 μ l of PBS in the ventral side of the right hind footpad. Viremia was monitored daily from 1 to 6 dpi and thereafter on every alternate day until 12 dpi. Briefly, 10 μ l of blood collected from the tail of each mouse was mixed in 120 μ l of PBS supplemented with 10 μ l of citrate-phosphate-dextrose solution (Sigma-Aldrich). RNA isolation was performed using the QIAamp Viral RNA kit (QIAGEN) following the manufacturer's instructions with a final elution volume of 60 μ l. 1 μ l of purified RNA was quantified by qRT-PCR using QuantiTect Probe RT-PCR (QIAGEN) as previously described (26). Joint swelling was measured for 2 wk post-ONNV inoculation as a function of height \times width relative to measurements preinfection (relative increase) (26).

Tissue viral load determination

Ketamine xylazine-anesthetized mice (150 mg/kg of ketamine, 10 mg/kg of xylazine) were intracardially perfused with PBS and organs were collected in tubes containing zirconia beads (TOMY Digital Biology) and 1 ml TRIzol (Invitrogen) and stored at -80°C. To isolate RNA, tissues were thawed on ice and homogenized using the Bead Ruptor Elite (OMNI International) at a speed of 6 m/s (3 cycles of lysis \times 30 s). Tissue lysates were then transferred to 1.5-ml

Eppendorf tubes and mixed with 230 μ l of chloroform and incubated at RT for 2 min. Samples were centrifuged at 12,000g for 10 min at 4°C and recovered supernatants were transferred into clean Eppendorf Tubes and mixed with 70% ethanol (1:1 volume). RNA was purified using the RNeasy Mini Kit (QIAGEN) according to the manufacturer's protocol. Viral RNA copies were quantified by qRT-PCR as described above.

In vivo virus tissue dissemination assay

To quantify tissue viral load and virus dissemination in vivo, mice were inoculated with a firefly luciferase-tagged ONNV infectious clone (ONNV-Fluc) and virus dissemination was tracked using the IVIS Spectrum In Vivo Imaging System (Perkin-Elmer) (26). Animals were kept anesthetized during the experiment using an oxygen flow rate of 1 liter/minute with 2% isoflurane. Full-body shaved mice were subcutaneously injected with 100 μ l of D-luciferin potassium salt (Caliper Life sciences) diluted in PBS (5 mg/ml). Whole body and footpad bioluminescence readings were independently taken 7 min post-D-luciferin injection with a field of view (FOV) of 21.7 cm (ventral position) and 13.1 cm (dorsal position) for whole body (FOV-D) and footpad (FOV-C) measurements, respectively. Two pictures were taken per FOV with exposure times set to "AUTO" and 60 s. Regions of interest were drawn using the software Living Image 3.0 and total flux values (photons/second) were calculated. Readings of naïve mice injected with D-luciferin were used for background subtraction.

Plasmodium infection and disease monitoring

Mice were infected with Py17x or PbA by i.p. injection of 10⁶ iRBC in Alsever's buffer. Parasitemia was monitored by flow cytometry as previously described (76) using a staining mix containing anti-mouse APC-tagged CD45 antibodies, 8 μ M dihydroethidium (Sigma-Aldrich), and 5 μ g/ml Hoechst 33342 (Sigma-Aldrich). Successful infections were confirmed 4 d postinoculation.

Isolation of footpad cells

Homogenous cell suspensions were obtained for immune profiling of footpads of infected and naïve animals. Mice were culled by cervical dislocation and right paws were harvested and immediately placed in 4 ml of digestion medium containing Collagenase IV (20 μ g/ml; Sigma-Aldrich), Dispase I (2 U/ml; Invitrogen), and DNase I (50 μ g/ml; Roche Applied Science) mixed in RPMI medium complemented with 10% FBS. Using forceps, footpads were deskinning and deboned to maximize digestion. Processed samples were placed on a shaker and incubated at 37°C at 100 rpm (Biosan PSU-10i) for 3 h. After digestion, tissues were passed through a 40- μ m cell strainer (Fisherbrand). Any remaining tissue trapped in the strainer was grinded using the top of a 1 ml syringe plunger to maximize cell recovery. 1 \times Flow Cytometry Mouse Lysis Buffer (R&D Systems) was used to lyse contaminating RBCs. Samples were resuspended in 1 ml of complete RPMI, overlaid to 35% vol/vol Percoll (Sigma-Aldrich)/RPMI mixture, and centrifuged at 2,400 rpm for 20 min at 4°C. Footpad cell pellets were washed and resuspended in appropriate volumes for counting using haemocytometers.

Profiling of immune and nonimmune cells by flow cytometry

Cell suspensions were stained for viability using LIVE/DEAD Aqua dye (Life Technologies). Cells were washed and resuspended in 50 μ l of blocking buffer containing TruStain FcX PLUS (anti-mouse CD16/32, clone S17011E) antibody diluted in PBS and incubated in the dark for 15 min on ice. Conjugated anti-mouse antibodies CD45 (30-F11), CD9 (eBioKMC8), CD29 (HM β 1-1), CD31 (390), and Integrin α 7 (334908), Sca-1 (D7) were used to stain cell surface markers for 30 min on ice. Finally, cells were fixed with 50 μ l of eBioscience IC Fixation Buffer (Thermo Fisher Scientific) for 5 min and acquired using a 5-laser LSR II flow cytometer (BD Biosciences) with BD FACSDiva software. Data were analyzed with FlowJo v10.6.2 (Becton, Dickinson and Company).

Dimensionality reduction analysis of flow cytometry data

Live CD45⁺ singlets events were pre-gated and then randomly down-sampled to a fixed number ($n = 5,000$) for each sample using FlowJo v10.7 (Becton, Dickinson and Company). Down-sampled files were concatenated and analyzed using UMAP for Dimension Reduction plug-in v3.1 using default parameters (number of nearest neighbours = 15, minimum distance = 0.1).

In vivo IFN γ neutralization

Anti-mouse IFN γ (0.5 mg per mouse, clone XMG1.2, Bio X Cell) was i.p. injected at 0-, 2-, and 4 d postparasite inoculation. Control groups were given rat IgG1 isotype control (0.5 mg, clone TNP6A7, Bio X Cell) at similar time points as treatment groups.

Cytokine/chemokine quantification by multiplexed bead-based immunoassays

Cytokine and chemokine concentrations (pg/ml) were quantified in footpad and serum samples. For footpad samples, animals were anesthetized with ketamine-xylazine and intracardially perfused with PBS. The right paw was cut at the ankle and placed in a gentleMACS M tube (Miltenyi) filled with 1.5 ml of RIPA buffer (50 mM Tris-HCl, pH 7.4, 1% NP-40, 0.25% sodium deoxycholate, 150 mM NaCl, and 1 mM EDTA) complemented with 1 \times cComplete Protease Inhibitor Cocktail (Roche). Samples were lysed in a Xiril Dispomix Tissue Homogenizer, centrifuged and the supernatants transferred into clean 2-ml microcentrifuge tubes for sonication in a Branson Ultrasonics Sonifier S-450 (70% intensity \times 15 s). For serum samples, blood was collected from the retro-orbicular sinus using a glass Pasteur pipette and allowed to clot for 30 min at room temperature. Clotted blood was centrifuged at 14,000 rpm for serum isolation. Footpad lysates and serum samples were analyzed using the Cytokine & Chemokine 36-Plex Mouse ProcartaPlex Panel 1A (Thermo Fisher Scientific) according to the manufacturer's protocol. Human plasma samples were analyzed using the Cytokine/Chemokine/Growth Factor 45-plex Human ProcartaPlex Panel 1 (Thermo Fisher Scientific). Data were acquired with Luminex FLEXMAP 3D instrument (Millipore) using xPONENT 4.0 software and analyzed with BioPlex Manager 6.1.1 (Bio-Rad Laboratories).

In vitro IFN γ treatment and ONNV infection

Recombinant human IFN γ (PHC4033; Gibco) diluted in supplemented DMEM at various concentrations was used to treat skin fibroblasts (BJ), synoviocytes (SW-982), skeletal muscle cells (RD), and endothelial cells (HPMEC) for 24 h before virus infection. Cells were washed with PBS and then infected with ONNV-ZsGreen virus at MOI of 10. Cells were harvested at 24-, 48-, and 72-h postinfection (hpi) and percentage of infection was quantified by flow cytometry.

Data and statistical analyses

Statistical analyses were performed using GraphPad Prism 8.4.3 (GraphPad Software). Data are presented as mean \pm SD unless otherwise specified. Nonparametric Mann-Whitney U statistical test was used to compute differences between two groups. Differences between three groups were calculated using two-tailed Kruskal-Wallis and post hoc Dunn's tests. Values obtained for viremia, parasitemia, in vivo imaging, and cytokine/chemokines were log-transformed for representation purposes. P -values < 0.05 were considered statistically significant.

Study approval

Animal experiments were approved by the Institutional Animal Care and Use Committee (IACUC #211635) of A*STAR in accordance with the guidelines of the Agri-Food and Veterinary Authority (AVA) and the National Advisory Committee for Laboratory Animal Research of Singapore (NACLAR). Plasma samples from febrile *P. vivax*-infected patients from Mae Sot, Thailand, were collected and tested in accordance with protocols approved by the University of Oxford Tropical Research Ethics Committee (OXTREC 17-11) and the Ethics Committee of the Faculty of Tropical Medicine at Mahidol University (MUTM 2008-215). Written informed consent was received before participation.

Supplementary Information

Supplementary Information is available at <https://doi.org/10.26508/lsa.202101272>.

Acknowledgements

The authors would like to thank Dr Carla Claser for critical discussion and valuable suggestions on the study. We also thank the SigN Flow Cytometry Core and SigN Mouse Core for assistance with cytometry analyses and support in animal breeding, respectively. We thank Wilson How from the SigN Immunomonitoring platform for his support in the multiplexed bead-based immunoassays. We also thank Professor Andres Merits from the University of Tartu for providing the tagged ONNV infectious clones used in this study. The study was supported by a core research grant provided to A*STAR Infectious Diseases Labs and Singapore Immunology Network by the Biomedical Research Council (BMRC) from the Agency for Science, Technology and Research (A*STAR). A Torres-Ruesta is supported by the A*STAR Singapore International Graduate Award (SINGA) scholarship. Flow cytometry platform is supported by the Health and Biomedical Sciences (HBMS) Open Fund Shared Infrastructure Support Grant under the Immunomonitoring Service

Platform project (NRF2017_SISFP09). The funders had no role in the study design, data collection and analysis, decision to publish, or preparation of the manuscript.

Author Contributions

A Torres-Ruesta: data curation, formal analysis, investigation, visualization, methodology, and writing—original draft, review, and editing.

T-H Teo: data curation, formal analysis, investigation, methodology, and writing—review and editing.

Y-H Chan: formal analysis, investigation, methodology, and writing—review and editing.

SN Amrun: formal analysis, investigation, methodology, and writing—review and editing.

NK-W Yeo: formal analysis, investigation, methodology, and writing—review and editing.

CY-P Lee: formal analysis, investigation, methodology, and writing—review and editing.

SY-T Ngeee: formal analysis, investigation, methodology, and writing—review and editing.

MZ Tay: formal analysis, investigation, methodology, and writing—review and editing.

F Nosten: formal analysis, investigation, methodology, and writing—review and editing.

S-W Fong: formal analysis, investigation, methodology, and writing—review and editing.

F-M Lum: formal analysis, investigation, methodology, and writing—review and editing.

G Carissimo: formal analysis, investigation, methodology, and writing—review and editing.

L Renia: conceptualization, supervision, and writing—original draft, review, and editing.

LFP Ng: conceptualization, resources, supervision, funding acquisition, project administration, and writing—original draft, review, and editing.

Conflict of Interest Statement

The authors declare that they have no conflict of interest.

References

- Rezza G, Chen R, Weaver SC (2017) O'nyong-nyong fever: A neglected mosquito-borne viral disease. *Pathog Glob Health* 111: 271–275. doi:[10.1080/20477724.2017.1355431](https://doi.org/10.1080/20477724.2017.1355431)
- Powers AM, Brault AC, Shirako Y, Strauss EG, Kang W, Strauss JH, Weaver SC (2001) Evolutionary relationships and systematics of the alphaviruses. *J Virol* 75: 10118–10131. doi:[10.1128/JVI.75.21.10118-10131.2001](https://doi.org/10.1128/JVI.75.21.10118-10131.2001)
- Sanders EJ, Rwaguma EB, Kawamata J, Kiwanuka N, Lutwama JJ, Ssengooba FP, Lamunu M, Najjemba R, Were WA, Bagambisa G, et al (1999) O'nyong-nyong fever in south-central Uganda, 1996–1997: Description of the epidemic and results of a household-based seroprevalence survey. *J Infect Dis* 180: 1436–1443. doi:[10.1086/315073](https://doi.org/10.1086/315073)
- Williams MC, Woodall JP, Corbet PS, Gillett JD (1965) O'nyong-Nyong fever: An epidemic virus disease in East Africa. 8. Virus isolations from Anopheles mosquitoes. *Trans R Soc Trop Med Hyg* 59: 300–306. doi:[10.1016/0035-9203\(65\)90012-x](https://doi.org/10.1016/0035-9203(65)90012-x)
- Haddow AJ, Davies CW, Walker AJ (1960) O'nyong-nyong fever: An epidemic virus disease in East Africa 1. Introduction. *Trans R Soc Trop Med Hyg* 54: 517–522. doi:[10.1016/0035-9203\(60\)90025-0](https://doi.org/10.1016/0035-9203(60)90025-0)
- Rwaguma EB, Lutwama JJ, Sempala SD, Kiwanuka N, Kamugisha J, Okware S, Bagambisa G, Lanciotti R, Roehrig JT, Gubler DJ (1997) Emergence of epidemic O'nyong-nyong fever in southwestern Uganda, after an absence of 35 years. *Emerg Infect Dis* 3: 77. doi:[10.3201/eid0301.970112](https://doi.org/10.3201/eid0301.970112)
- Posey DL, O'Rourke T, Roehrig JT, Lanciotti RS, Weinberg M, Maloney S (2005) O'Nyong-nyong fever in West Africa. *Am J Trop Med Hyg* 73: 32. doi:[10.4269/ajtmh.2005.73.1.0730032](https://doi.org/10.4269/ajtmh.2005.73.1.0730032)
- LaBeaud AD, Banda T, Brichard J, Muchiri EM, Mungai PL, Mutuku FM, Borland E, Gildengorin G, Pfeil S, Teng CY, et al (2015) High rates of O'nyong nyong and Chikungunya virus transmission in coastal Kenya. *PLoS Negl Trop Dis* 9: e0003436. doi:[10.1371/journal.pntd.0003436](https://doi.org/10.1371/journal.pntd.0003436)
- Clements TL, Rossi CA, Irish AK, Kibuuka H, Eller LA, Robb ML, Kataaha P, Michael NL, Hensley LE, Schoepp RJ (2019) Chikungunya and O'nyong-nyong viruses in Uganda: Implications for diagnostics. *Open Forum Infect Dis* 6: ofz001. doi:[10.1093/ofid/ofz001](https://doi.org/10.1093/ofid/ofz001)
- Baba M, Logue CH, Oderinde B, Abdulmaleek H, Williams J, Lewis J, Laws TR, Hewson R, Marcello A, D'Agaro P (2013) Evidence of arbovirus co-infection in suspected febrile malaria and typhoid patients in Nigeria. *J Infect Dev Ctries* 7: 51–59. doi:[10.3855/jidc.2411](https://doi.org/10.3855/jidc.2411)
- Kinimi E, Patrick BN, Misinzo G (2018) Serological evidence of chikungunya and malaria co-infection among febrile patients seeking health care in district, Tanzania. *Tanzania J Health Res* 20: 1–8. doi:[10.4314/thrb.v20i4.1](https://doi.org/10.4314/thrb.v20i4.1)
- Mostafavi H, Abeyratne E, Zaid A, Taylor A (2019) Arthritogenic alphavirus-induced immunopathology and targeting host inflammation as a therapeutic strategy for alphaviral disease. *Viruses* 11: 290. doi:[10.3390/v11030290](https://doi.org/10.3390/v11030290)
- Assunção-Miranda I, Cruz-Oliveira C, Da Poian AT (2013) Molecular mechanisms involved in the pathogenesis of -induced arthritis. *Biomed Res Int* 2013: 973516. doi:[10.1155/2013/973516](https://doi.org/10.1155/2013/973516)
- Septembre-Malaterre A, Bedoui Y, Giry C, Gasque P, Guiraud P, Sélambarom J (2021) Quercetin can reduce viral RNA level of O'nyong-nyong virus and resulting innate immune cytokine responses in cultured human synovial fibroblasts. *Sci Rep* 11: 6369. doi:[10.1038/s41598-021-85840-z](https://doi.org/10.1038/s41598-021-85840-z)
- Clark IA (2007) The advent of the cytokine storm. *Immunol Cell Biol* 85: 271–273. doi:[10.1038/sj.icb.7100062](https://doi.org/10.1038/sj.icb.7100062)
- Clark IA, Alleva LM, Budd AC, Cowden WB (2008) Understanding the role of inflammatory cytokines in malaria and related diseases. *Trav Med Infect Dis* 6: 67–81. doi:[10.1016/j.tmaid.2007.07.002](https://doi.org/10.1016/j.tmaid.2007.07.002)
- Kurup SP, Butler NS, Harty JT (2019) T cell-mediated immunity to malaria. *Nat Rev Immunol* 19: 457–471. doi:[10.1038/s41577-019-0158-z](https://doi.org/10.1038/s41577-019-0158-z)
- Rénia L, Howland SW, Claser C, Charlotte Gruner A, Suwanarusk R, Hui Teo T, Russell B, Ng LF (2012) Cerebral malaria: Mysteries at the blood-brain barrier. *Virulence* 3: 193–201. doi:[10.4161/viru.19013](https://doi.org/10.4161/viru.19013)
- Howland SW, Claser C, Poh CM, Gun SY, Renia L (2015) Pathogenic CD8+ T cells in experimental cerebral malaria. *Semin Immunopathol* 37: 221–231. doi:[10.1007/s00281-015-0476-6](https://doi.org/10.1007/s00281-015-0476-6)
- Rénia L, Potter SM, Mauduit M, Rosa DS, Kayibanda M, Deschemin JC, Snounou G, Grüner AC (2006) Pathogenic T cells in cerebral malaria. *Int J Parasitol* 36: 547–554. doi:[10.1016/j.ijpara.2006.02.007](https://doi.org/10.1016/j.ijpara.2006.02.007)
- Claser C, Ngeee SYT, Balachander A, Howland SW, Becht E, Gunasegaran B, Hartimath SV, Lee AWQ, Ho JTT, Ong CB, et al (2019) Author Correction: Lung endothelial cell antigen cross-presentation to CD8+ T cells drives malaria-associated lung injury. *Nat Commun* 10: 5066. doi:[10.1038/s41467-019-13025-4](https://doi.org/10.1038/s41467-019-13025-4)

22. Wangoo A, Ganguly NK, Mahajan RC (1990) Immunosuppression in murine malaria: Suppressor role of macrophages and their products during acute and chronic Plasmodium berghei infection. *APMIS* 98: 407–414. doi:[10.1111/j.1699-0463.1990.tb01051.x](https://doi.org/10.1111/j.1699-0463.1990.tb01051.x)
23. Chemtai AK, Okello GB (1989) Suppression of T-cell proliferative response in Plasmodium falciparum malaria patients—preliminary results. *East Afr Med J* 66: 787–791.
24. Theander TG, Bygbjerg IC, Andersen BJ, Jepsen S, Kharazmi A, Odum N (1986) Suppression of parasite-specific response in Plasmodium falciparum malaria. A longitudinal study of blood mononuclear cell proliferation and subset composition. *Scand J Immunol* 24: 73–81. doi:[10.1111/j.1365-3083.1986.tb02071.x](https://doi.org/10.1111/j.1365-3083.1986.tb02071.x)
25. Hviid L, Theander TG, Abu-Zeid YA, Abdulhadi NH, Jakobsen PH, Saeed BO, Jepsen S, Bayoumi RA, Jensen JB (1991) Loss of cellular immune reactivity during acute Plasmodium falciparum malaria. *FEMS Microbiol Immunol* 3: 219–227. doi:[10.1111/j.1574-6968.1991.tb04218.x](https://doi.org/10.1111/j.1574-6968.1991.tb04218.x)
26. Chan YH, Teo TH, Torres-Ruesta A, Hartimath SV, Chee RS, Khanapur S, Yong FF, Ramasamy B, Cheng P, Rajarethinam R, et al (2020) Longitudinal [18F]FB-IL-2 PET imaging to assess the immunopathogenicity of O'nyong-nyong virus infection. *Front Immunol* 11: 894. doi:[10.3389/fimmu.2020.00894](https://doi.org/10.3389/fimmu.2020.00894)
27. Lentscher AJ, McCarthy MK, May NA, Davenport BJ, Montgomery SA, Raghunathan K, McAllister N, Silva LA, Morrison TE, Dermody TS (2020) Chikungunya virus replication in skeletal muscle cells is required for disease development. *J Clin Invest* 130: 1466–1478. doi:[10.1172/JCI129893](https://doi.org/10.1172/JCI129893)
28. Gardner J, Anraku I, Le TT, Larcher T, Major L, Roques P, Schroder WA, Higgs S, Suhrbier A (2010) Chikungunya virus arthritis in adult wild-type mice. *J Virol* 84: 8021–8032. doi:[10.1128/JVI.02603-09](https://doi.org/10.1128/JVI.02603-09)
29. Sourisseau M, Schilte C, Casartelli N, Trouillet C, Guivel-Benhassine F, Rudnicka D, Sol-Foulon N, Le Roux K, Prevost MC, Fsihi H, et al (2007) Characterization of reemerging chikungunya virus. *PLoS Pathog* 3: e89. doi:[10.1371/journal.ppat.0030089](https://doi.org/10.1371/journal.ppat.0030089)
30. Halfon S, Abramov N, Grinblat B, Ginis I (2011) Markers distinguishing mesenchymal stem cells from fibroblasts are downregulated with passaging. *Stem Cells Dev* 20: 53–66. doi:[10.1089/scd.2010.0040](https://doi.org/10.1089/scd.2010.0040)
31. Gabrielis K (2012) Surface markers distinguishing mesenchymal stem cells from fibroblasts. *Acta Med Lituanica* 19: 75–79. doi:[10.6001/actamedica.v19i2.2313](https://doi.org/10.6001/actamedica.v19i2.2313)
32. Kafadar KA, Yi L, Ahmad Y, So L, Rossi F, Pavlath GK (2009) Sca-1 expression is required for efficient remodeling of the extracellular matrix during skeletal muscle regeneration. *Dev Biol* 326: 47–59. doi:[10.1016/j.ydbio.2008.10.036](https://doi.org/10.1016/j.ydbio.2008.10.036)
33. Mierzejewski B, Archacka K, Grabowska I, Florkowska A, Ciemerych MA, Brzoska E (2020) Human and mouse skeletal muscle stem and progenitor cells in health and disease. *Semin Cell Dev Biol* 104: 93–104. doi:[10.1016/j.semcdb.2020.01.004](https://doi.org/10.1016/j.semcdb.2020.01.004)
34. Yao Z, Liu H, Yang M, Bai Y, Zhang B, Wang C, Yan Z, Niu G, Zou Y, Li Y (2020) Bone marrow mesenchymal stem cell-derived endothelial cells increase capillary density and accelerate angiogenesis in mouse hindlimb ischemia model. *Stem Cell Res Ther* 11: 221. doi:[10.1186/s13287-020-01710-x](https://doi.org/10.1186/s13287-020-01710-x)
35. De Souza JB, Williamson KH, Otani T, Playfair JH (1997) Early gamma interferon responses in lethal and nonlethal murine blood-stage malaria. *Infect Immun* 65: 1593–1598. doi:[10.1128/IAI.65.5.1593-1598.1997](https://doi.org/10.1128/IAI.65.5.1593-1598.1997)
36. Mitchell AJ, Hansen AM, Hee L, Ball HJ, Potter SM, Walker JC, Hunt NH (2005) Early cytokine production is associated with protection from murine cerebral malaria. *Infect Immun* 73: 5645–5653. doi:[10.1128/IAI.73.9.5645-5653.2005](https://doi.org/10.1128/IAI.73.9.5645-5653.2005)
37. Miller JL, Sack BK, Baldwin M, Vaughan AM, Kappe SHI (2014) Interferon-mediated innate immune responses against malaria parasite liver stages. *Cell Rep* 7: 436–447. doi:[10.1016/j.celrep.2014.03.018](https://doi.org/10.1016/j.celrep.2014.03.018)
38. Kang S, Brown HM, Hwang S (2018) Direct antiviral mechanisms of interferon-gamma. *Immune Netw* 18: e33. doi:[10.4110/in.2018.18.e33](https://doi.org/10.4110/in.2018.18.e33)
39. He X, Xia L, Tumas KC, Wu J, Su XZ (2020) Type I interferons and malaria: A double-edge sword against a complex parasitic disease. *Front Cell Infect Microbiol* 10: 594621. doi:[10.3389/fcimb.2020.594621](https://doi.org/10.3389/fcimb.2020.594621)
40. Sebina I, Haque A (2018) Effects of type I interferons in malaria. *Immunology* 155: 176–185. doi:[10.1111/imm.12971](https://doi.org/10.1111/imm.12971)
41. Mooney JP, Wassmer SC, Hafalla JC (2017) Type I interferon in malaria: A balancing act. *Trends Parasitol* 33: 257–260. doi:[10.1016/j.pt.2016.12.010](https://doi.org/10.1016/j.pt.2016.12.010)
42. Couderc T, Chrétien F, Schilte C, Disson O, Brigitte M, Guivel-Benhassine F, Touret Y, Barau G, Cayet N, Schuffenecker I, et al (2008) A mouse model for chikungunya: Young age and inefficient type-I interferon signaling are risk factors for severe disease. *PLoS Pathog* 4: e29. doi:[10.1371/journal.ppat.0040029](https://doi.org/10.1371/journal.ppat.0040029)
43. Seymour RL, Rossi SL, Bergren NA, Plante KS, Weaver SC (2013) The role of innate versus adaptive immune responses in a mouse model of O'nyong-nyong virus infection. *Am J Trop Med Hyg* 88: 1170–1179. doi:[10.4269/ajtmh.12-0674](https://doi.org/10.4269/ajtmh.12-0674)
44. Crisler WJ, Lenz LL (2018) Crosstalk between type I and II interferons in regulation of myeloid cell responses during bacterial infection. *Curr Opin Immunol* 54: 35–41. doi:[10.1016/j.coi.2018.05.014](https://doi.org/10.1016/j.coi.2018.05.014)
45. Rayamajhi M, Humann J, Kearney S, Hill KK, Lenz LL (2010) Antagonistic crosstalk between type I and II interferons and increased host susceptibility to bacterial infections. *Virulence* 1: 418–422. doi:[10.4161/viru.1.5.12787](https://doi.org/10.4161/viru.1.5.12787)
46. Gough DJ, Messina NL, Hii L, Gould JA, Sabapathy K, Robertson AP, Trapani JA, Levy DE, Hertzog PJ, Clarke CJ, et al (2010) Functional crosstalk between type I and II interferon through the regulated expression of STAT1. *PLoS Biol* 8: e1000361. doi:[10.1371/journal.pbio.1000361](https://doi.org/10.1371/journal.pbio.1000361)
47. Schroder K, Hertzog PJ, Ravasi T, Hume DA (2004) Interferon-gamma: An overview of signals, mechanisms and functions. *J Leukoc Biol* 75: 163–189. doi:[10.1189/jlb.0603252](https://doi.org/10.1189/jlb.0603252)
48. Young AR, Locke MC, Cook LE, Hiller BE, Zhang R, Hedberg ML, Monte KJ, Veis DJ, Diamond MS, Lenschow DJ (2019) Dermal and muscle fibroblasts and skeletal myofibers survive chikungunya virus infection and harbor persistent RNA. *PLoS Pathog* 15: e1007993. doi:[10.1371/journal.ppat.1007993](https://doi.org/10.1371/journal.ppat.1007993)
49. Burdeinick-Kerr R, Griffin DE (2005) Gamma interferon-dependent, noncytolytic clearance of virus infection from neurons in vitro. *J Virol* 79: 5374–5385. doi:[10.1128/JVI.79.9.5374-5385.2005](https://doi.org/10.1128/JVI.79.9.5374-5385.2005)
50. Burdeinick-Kerr R, Govindarajan D, Griffin DE (2009) Noncytolytic clearance of virus infection from neurons by gamma interferon is dependent on Jak/STAT signaling. *J Virol* 83: 3429–3435. doi:[10.1128/JVI.02381-08](https://doi.org/10.1128/JVI.02381-08)
51. Schilte C, Couderc T, Chrétien F, Sourisseau M, Gangneux N, Guivel-Benhassine F, Kraxner A, Tschopp J, Higgs S, Michault A, et al (2010) Type I IFN controls chikungunya virus via its action on nonhematopoietic cells. *J Exp Med* 207: 429–442. doi:[10.1084/jem.20090851](https://doi.org/10.1084/jem.20090851)
52. Ryman KD, Klimstra WB, Nguyen KB, Biron CA, Johnston RE (2000) Alpha/beta interferon protects adult mice from fatal Sindbis virus infection and is an important determinant of cell and tissue tropism. *J Virol* 74: 3366–3378. doi:[10.1128/jvi.74.7.3366-3378.2000](https://doi.org/10.1128/jvi.74.7.3366-3378.2000)
53. Metzemaekers M, Vanheule V, Janssens R, Struyf S, Proost P (2017) Overview of the mechanisms that may contribute to the non-redundant activities of interferon-inducible CXC chemokine receptor 3 ligands. *Front Immunol* 8: 1970. doi:[10.3389/fimmu.2017.01970](https://doi.org/10.3389/fimmu.2017.01970)
54. Liu Y, Cai Y, Liu L, Wu Y, Xiong X (2018) Crucial biological functions of CCL7 in cancer. *PeerJ* 6: e4928. doi:[10.7717/peerj.4928](https://doi.org/10.7717/peerj.4928)
55. Gough DJ, Levy DE, Johnstone RW, Clarke CJ (2008) IFN-gamma signaling—does it mean JAK-STAT? *Cytokine Growth Factor Rev* 19: 383–394. doi:[10.1016/j.cytogfr.2008.08.004](https://doi.org/10.1016/j.cytogfr.2008.08.004)
56. Durbin JE, Hackenmiller R, Simon MC, Levy DE (1996) Targeted disruption of the mouse Stat1 gene results in compromised innate immunity to viral disease. *Cell* 84: 443–450. doi:[10.1016/s0092-8674\(00\)81289-1](https://doi.org/10.1016/s0092-8674(00)81289-1)

57. Meraz MA, White JM, Sheehan KC, Bach EA, Rodig SJ, Dighe AS, Kaplan DH, Riley JK, Greenlund AC, Campbell D, et al (1996) Targeted disruption of the Stat1 gene in mice reveals unexpected physiologic specificity in the JAK-STAT signaling pathway. *Cell* 84: 431–442. doi:[10.1016/s0092-8674\(00\)81288-x](https://doi.org/10.1016/s0092-8674(00)81288-x)
58. Rogers KJ, Shtanko O, Vijay R, Mallinger LN, Joyner CJ, Galinski MR, Butler NS, Maury W (2020) Acute Plasmodium infection promotes interferon-gamma-dependent resistance to Ebola virus infection. *Cell Rep* 30: 4041–4051.e4. doi:[10.1016/j.celrep.2020.02.104](https://doi.org/10.1016/j.celrep.2020.02.104)
59. Rosenke K, Adjemian J, Munster VJ, Marzi A, Falzarano D, Onyango CO, Ochieng M, Juma B, Fischer RJ, Prescott JB, et al (2016) Plasmodium parasitemia associated with increased survival in Ebola virus-infected patients. *Clin Infect Dis* 63: 1026–1033. doi:[10.1093/cid/ciw452](https://doi.org/10.1093/cid/ciw452)
60. Edwards CL, Zhang V, Werder RB, Best SE, Sebina I, James KR, Faleiro RJ, de Labastida Rivera F, Amante FH, Engwerda CR, et al (2015) Coinfection with blood-stage Plasmodium promotes systemic type I interferon production during pneumovirus infection but impairs inflammation and viral control in the lung. *Clin Vaccin Immunol* 22: 477–483. doi:[10.1128/CVI.00051-15](https://doi.org/10.1128/CVI.00051-15)
61. Matar CG, Anthony NR, O’Flaherty BM, Jacobs NT, Priyamvada L, Engwerda CR, Speck SH, Lamb TJ (2015) Co-infection with malaria suppresses anti-parasitic humoral immunity. *PLoS Pathog* 11: e1004858. doi:[10.1371/journal.ppat.1004858](https://doi.org/10.1371/journal.ppat.1004858)
62. Kima PE, Soong L (2013) Interferon gamma in leishmaniasis. *Front Immunol* 4: 156. doi:[10.3389/fimmu.2013.00156](https://doi.org/10.3389/fimmu.2013.00156)
63. Sturge CR, Yarovinsky F (2014) Complex immune cell interplay in the gamma interferon response during Toxoplasma gondii infection. *Infect Immun* 82: 3090–3097. doi:[10.1128/IAI.01722-14](https://doi.org/10.1128/IAI.01722-14)
64. Rodrigues AA, Saosa JS, da Silva GK, Martins FA, da Silva AA, Souza Neto CP, Horta CV, Zamboni DS, da Silva JS, Ferro EA, et al (2012) IFN-γ plays a unique role in protection against low virulent Trypanosoma cruzi strain. *PLoS Negl Trop Dis* 6: e1598. doi:[10.1371/journal.pntd.0001598](https://doi.org/10.1371/journal.pntd.0001598)
65. Pereira L, Oliveira F, Townsend S, Metangmo S, Meneses C, Moore IN, Brodskyn CI, Valenzuela JG, Magez S, Kamhawi S (2018) Coinfection with trypanosoma brucei confers protection against cutaneous leishmaniasis. *Front Immunol* 9: 2855. doi:[10.3389/fimmu.2018.02855](https://doi.org/10.3389/fimmu.2018.02855)
66. Millington OR, Di Lorenzo C, Phillips RS, Garside P, Brewer JM (2006) Suppression of adaptive immunity to heterologous antigens during Plasmodium infection through hemozoin-induced failure of dendritic cell function. *J Biol* 5: 5. doi:[10.1186/jbiol34](https://doi.org/10.1186/jbiol34)
67. Millington OR, Gibson VB, Rush CM, Zinselmeyer BH, Phillips RS, Garside P, Brewer JM (2007) Malaria impairs T cell clustering and immune priming despite normal signal 1 from dendritic cells. *PLoS Pathog* 3: 1380–1387. doi:[10.1371/journal.ppat.0030143](https://doi.org/10.1371/journal.ppat.0030143)
68. Ferguson NM (2018) Challenges and opportunities in controlling mosquito-borne infections. *Nature* 559: 490–497. doi:[10.1038/s41586-018-0318-5](https://doi.org/10.1038/s41586-018-0318-5)
69. Bessaud M, Peyrefitte CN, Pastorino BA, Gravier P, Tock F, Boete F, Tolou HJ, Grandadam M (2006) O’nyong-nyong Virus, Chad. *Emerg Infect Dis* 12: 1248–1250. doi:[10.3201/eid1208.060199](https://doi.org/10.3201/eid1208.060199)
70. Carissimo G, Ng LFP (2019) Understanding molecular pathogenesis with chikungunya virus research tools. *Curr Top Microbiol Immunol* 1–21. doi:[10.1007/82_2019_158](https://doi.org/10.1007/82_2019_158)
71. Pohjala L, Utt A, Varjak M, Lulla A, Merits A, Ahola T, Tammela P (2011) Inhibitors of entry and replication identified with a stable Chikungunya replicon cell line and virus-based assays. *PLoS One* 6: e28923. doi:[10.1371/journal.pone.0028923](https://doi.org/10.1371/journal.pone.0028923)
72. Weiss WR, Good MF, Hollingdale MR, Miller LH, Berzofsky JA (1989) Genetic control of immunity to Plasmodium sporozoites. *J Immunol* 143: 4263–4266.
73. van Dijk MR, Waters AP, Janse CJ (1995) Stable transfection of malaria parasite blood stages. *Science* 268: 1358–1362. doi:[10.1126/science.7761856](https://doi.org/10.1126/science.7761856)
74. Hu K, Li Y, Wu W, Chen H, Chen Z, Zhang Y, Guo Y, Dong Y (2018) High-performance gene expression and knockout tools using sleeping beauty transposon system. *Mob DNA* 9: 33. doi:[10.1186/s13100-018-0139-y](https://doi.org/10.1186/s13100-018-0139-y)
75. Mátés L, Chuah MK, Belay E, Jerchow B, Manoj N, Acosta-Sanchez A, Grzela DP, Schmitt A, Becker K, Matrai J, et al (2009) Molecular evolution of a novel hyperactive Sleeping Beauty enables robust stable gene transfer in vertebrates. *Nat Genet* 41: 753–761. doi:[10.1038/ng.343](https://doi.org/10.1038/ng.343)
76. Malleret B, Claser C, Ong AS, Suwanarusk R, Sriprawat K, Howland SW, Russell B, Nosten F, Rénia L (2011) A rapid and robust tri-color flow cytometry assay for monitoring malaria parasite development. *Sci Rep* 1: 118. doi:[10.1038/srep00118](https://doi.org/10.1038/srep00118)



License: This article is available under a Creative Commons License (Attribution 4.0 International, as described at <https://creativecommons.org/licenses/by/4.0/>).