

# A Role for *Ifit2* in Restricting West Nile Virus Infection in the Brain

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**Previous studies have demonstrated that type I interferon (IFN-I) restricts West Nile virus (WNV) replication and pathogenesis in peripheral and central nervous system (CNS) tissues. However, the *in vivo* role of specific antiviral genes that are induced by IFN-I against WNV infection remains less well characterized. Here, using *Ifit2*<sup>-/-</sup> mice, we defined the antiviral function of the interferon-stimulated gene (ISG) *Ifit2* in limiting infection and disease *in vivo* by a virulent North American strain of WNV. Compared to congenic wild-type controls, *Ifit2*<sup>-/-</sup> mice showed enhanced WNV infection in a tissue-restricted manner, with preferential replication in the CNS of animals lacking *Ifit2*. Virological analysis of cultured macrophages, dendritic cells, fibroblasts, cerebellar granule cell neurons, and cortical neurons revealed cell type-specific antiviral functions of *Ifit2* against WNV. In comparison, small effects of *Ifit2* were observed on the induction or magnitude of innate or adaptive immune responses. Our results suggest that *Ifit2* restricts WNV infection and pathogenesis in different tissues in a cell type-specific manner.**

West Nile virus (WNV) is a mosquito-transmitted, neurotropic, positive-stranded RNA virus in the *Flaviviridae* family, which includes other pathogens of global health concern, such as dengue, yellow fever, and Japanese encephalitis viruses. WNV is maintained in an enzootic cycle between *Culex* species mosquitoes and several avian hosts but can cause disease in vertebrate animals, including horses and humans (1). Before 1999, the distribution of WNV was localized largely to Africa, central and southern Asia, the Middle East, southern Europe, and Oceania (2). Since its entry into North America in New York City in 1999, WNV has spread across the continental United States into Canada, Mexico, and parts of South America, likely due to bird migration and the presence of competent mosquito hosts (3, 4). While human transmission usually occurs through mosquito inoculation, other routes (blood transfusion, organ transplantation, and intrauterine transmission) have been reported (1). Although the majority (~50 to 70%) of human infections are asymptomatic, WNV infection in the elderly or immunocompromised can cause severe neuroinvasive disease, including meningitis, encephalitis, and acute flaccid paralysis (5–7). In the United States alone between 1999 and 2012, ~36,000 cases and ~1,500 deaths have been confirmed (<http://www.cdc.gov/ncidod/dvbid/westnile/surv&control.htm>). Despite this, no approved vaccines or therapeutics are available to treat or prevent human WNV infection.

RNA intermediates of viral replication are recognized by cytosolic and endosomal pattern recognition receptors (PRR), such as RIG-I-like receptors (RLR) or Toll-like receptors (TLR), which signal specific transcription factors (e.g., Irf3 or Irf7) to induce type I interferon (IFN-I) expression and secretion. Autocrine and paracrine binding of IFN-I to its receptor (Ifnar) results in a signaling cascade that includes Stat1, Stat2, and Irf9 and results in the induction of many interferon-stimulated genes (ISGs), some of which inhibit virus infections. Although IFN-I responses control the cell and tissue tropism of several families of RNA and DNA viruses, the specific molecules that mediate this remain poorly characterized. While earlier studies defined *Mx1*, *Pkr*, and *RNaseL* as broad-spectrum antiviral ISGs against several families of RNA viruses, more recent investigations have identified other ISGs (e.g., *Ifit1*, *Ifitm* genes, *Bst2*, *Apobec3g*, and *Adar*) that restrict infection of a more limited range of viruses (8–10). Several individ-

ual ISGs have been suggested to have antiviral activity against WNV. *C6orf150*, *Hpse*, *Nampt*, *Phf15*, *Ifitm2*, *Ifitm3*, and *Isg20* inhibit WNV infection *in vitro* (10, 11), and *viperin* (*Rsad2*), *Pkr*, and *RNaseL* restrict WNV pathogenesis *in vivo* and *in vitro* in a tissue- and cell type-specific manner (8, 12).

*Ifit2* (also known as ISG54) is a member of the IFN-induced proteins with tetratricopeptide repeats (IFIT) family. IFIT proteins contain multiple tetratricopeptide repeats, which are domains implicated in the regulation of cell cycle, transcription, protein transport, and protein folding processes (13). Initial experiments indicated that *Ifit2* could restrict infection of several viruses by binding to and inhibiting subunits of eIF3, a key protein involved in initiation of host translation (14). More recent structural and functional studies suggest that IFIT family members also inhibit infection of cytoplasmic RNA and DNA viruses by directly binding nonself RNA, including moieties displaying 5'-ppp RNA (15, 16) and possibly 5' cap 0 structures (7mGpppN) lacking 2'-O methylation (17, 18). *Ifit2*, in particular, has been demonstrated to have antiviral activity in cell culture against WNV (19). While *in vivo* studies have established an antiviral effect of *Ifit2* against vesicular stomatitis virus (VSV) infection (20), analogous experiments have not been conducted with WNV or any other flavivirus.

Beyond its antiviral function, *Ifit2* has been implicated in the regulation of host cell immune responses. One study showed that human IFIT1 and IFIT2 bind to and inhibit MITA (also known as STING), which functions as a mitochondrial adaptor protein that recruits TANK-binding kinase 1 (TBK1) and IRF-3 to a complex with MAVS (also known as IPS-1, CARDIF, and VISA), resulting in the downstream induction of IFN- $\beta$  expression in response to viral RNA or DNA (21). However, these data conflict with the results from mouse fibroblasts, macrophages, and dendritic cells in which silencing of IFIT2 expression did not alter IFN-I re-

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sponses (22). Finally, expression of human IFIT2 also has been suggested to promote apoptosis via a mitochondrial pathway (23).

Here, we examined in detail the function of *Ifit2* *in vivo* against WNV infection and pathogenesis using *Ifit2*<sup>-/-</sup> mice. Our data indicate that *Ifit2* restricts WNV infection in a cell- and tissue-specific manner yet had small effects on the induction and magnitude of the innate and adaptive antiviral immune responses. Thus, *Ifit2* is a key antiviral effector molecule that functions downstream of host pathogen recognition signaling to inhibit infection by WNV and possibly other viruses.

## MATERIALS AND METHODS

**Virus propagation and titration.** The WNV strain (3000.0259) was isolated in New York in 2000 and passaged once in C6/36 *Aedes albopictus* cells to generate an insect cell- derived stock virus as described previously (24). Mammalian cell-derived WNV was generated from an infectious cDNA clone of the New York 1999 strain, propagated once in C6/36 *Aedes albopictus* cells, and passaged a second time in Vero cells (25). WNV-NS5-E218A was propagated in BHK-21-15 cells as described previously (26). BHK-21-15 and Vero cells were used for titration of virus in tissue and cells by plaque or focus-forming assay as described previously (27, 28). Levels of virus in serum and lymph nodes were determined by quantitative reverse transcription (RT)-PCR using WNV-specific primers and probe as described previously (8).

**Mouse experiments and tissue preparation.** C57BL/6 wild-type mice were obtained commercially (Jackson Laboratory), and the generation of congenic *Ifit2*<sup>-/-</sup> mice was described previously (20). Mice were bred in the animal facility of Washington University School of Medicine, and experiments were performed according to the guidelines and with approval of the Washington University Animal Studies Committee. For infection, 8- to 10-week-old age-matched mice were used. Mice were infected subcutaneously in the footpad (10<sup>2</sup> PFU in 50  $\mu$ l) or intracranially (10<sup>1</sup> PFU in 10  $\mu$ l) with virus diluted in Hanks balanced salt solution (HBSS) supplemented with 1% heat-inactivated fetal bovine serum (FBS). On specific days after infection, mice were perfused extensively with phosphate-buffered saline (PBS), tissues were harvested and weighed, and virus titers were determined by plaque assay on BHK-21-15 cells (27). For survival studies, 8-week-old mice were infected via footpad inoculation and monitored for 21 days.

**IFN bioassay.** Levels of biologically active IFN-I in serum of mice infected with WNV were determined using an encephalomyocarditis virus L929 cell cytopathic effect bioassay as described previously (28, 29).

**Cytokine analysis.** Mice were infected subcutaneously with 10<sup>2</sup> PFU of WNV in the footpad, and serum was collected at 2 days after infection. Cytokine levels in serum were measured using a Bio-Plex Pro 8-plex custom cytokine kit (Bio-Rad) and Bio-Plex 200 (Bio-Rad).

**Measurement of WNV-specific antibodies.** The levels of WNV-specific IgM and IgG were measured using an enzyme-linked immunosorbent assay (ELISA) with purified WNV E protein as described previously (30). The titer of neutralizing antibody in serum was quantitated by a focus reduction neutralization assay in Vero cells as described previously (31).

**Analysis of splenic CD4<sup>+</sup> and CD8<sup>+</sup> T cells.** Splenocytes were harvested from wild-type or *Ifit2*<sup>-/-</sup> mice on day 8 after WNV infection. Intracellular IFN- $\gamma$  or tumor necrosis factor alpha (TNF- $\alpha$ ) staining of CD8<sup>+</sup> T cells was performed after *ex vivo* restimulation with an immunodominant D<sup>p</sup>-restricted NS4B peptide as described previously (32). Samples were processed, stained with antibodies against IFN- $\gamma$  and TNF- $\alpha$  (BD Pharmingen), and analyzed using an LSR II flow cytometer (Becton, Dickinson) and FlowJo software (Treestar).

**CNS leukocyte isolation and phenotyping.** Brains were harvested from wild-type and *Ifit2*<sup>-/-</sup> mice on day 8 after WNV infection, and CNS leukocytes were isolated after Percoll gradient centrifugation as described previously (33). Cells were stained for CD3, CD4, CD8, CD45, and CD11b with directly conjugated antibodies (BD Pharmingen). Intracellular stain-

ing for IFN- $\gamma$  and TNF- $\alpha$  was performed after NS4B peptide restimulation as described before (32).

**Primary cell isolation and infection.** Primary macrophages, dendritic cells, embryonic fibroblasts, cortical neurons, and cerebellar granule cell neurons were prepared from embryonic, neonatal, and adult wild-type and *Ifit2*<sup>-/-</sup> mice as detailed in prior studies (34, 35). For multistep growth curves, cells were infected at a low multiplicity of infection (MOI of 0.001) with or without IFN- $\beta$  pretreatment (12 h, 10 IU/ml; PBL Interferon Source) or IFN- $\zeta$  pretreatment (4 h, 0.4 ng/ml; PBL Interferon Source). Virus was harvested from supernatants at specific times and titrated by focus-forming assay on Vero cells.

**Statistical analysis.** Kaplan-Meier survival curves were analyzed by the log rank test. A two-tailed Student *t* test was used to determine statistically significant differences for *in vitro* experiments. The Mann-Whitney test was used to analyze differences in viral burden. All data were analyzed by using Prism software (GraphPad Software).

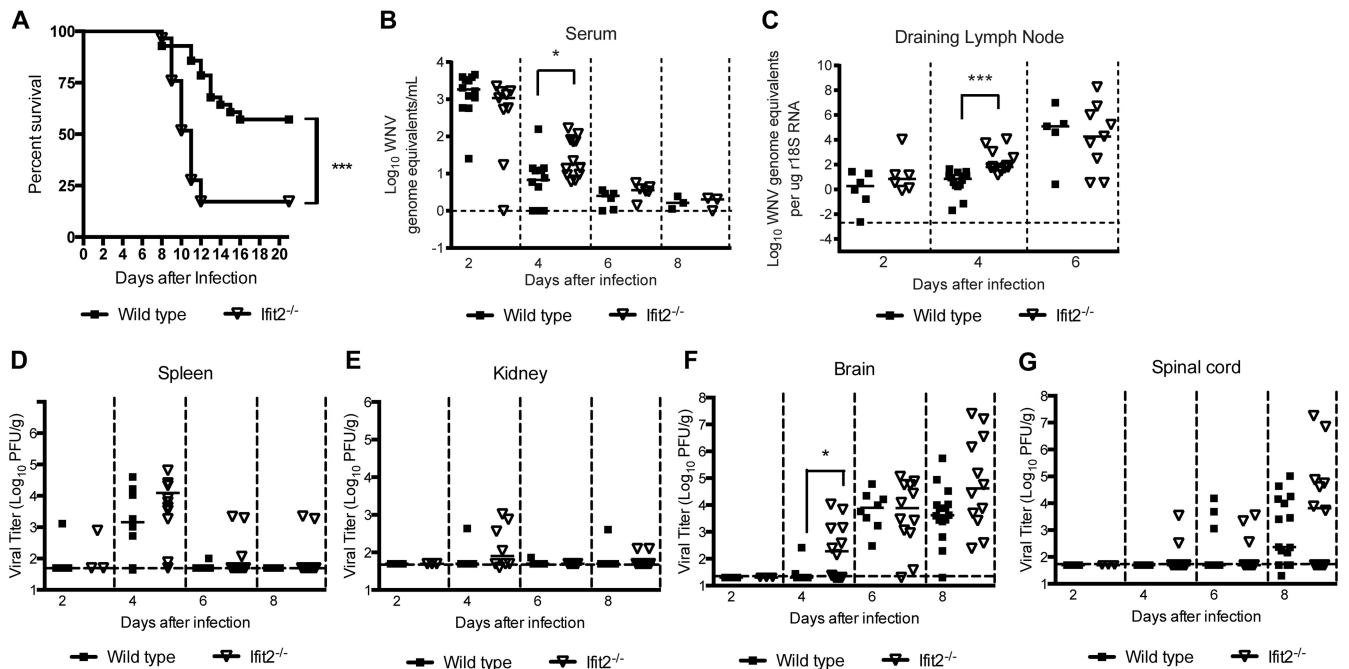
## RESULTS

**Ifit2 is required for control of WNV infection *in vivo*.** To determine whether a deficiency of *Ifit2* resulted in increased susceptibility to lethal WNV infection, we infected wild-type and congenic *Ifit2*<sup>-/-</sup> mice via a subcutaneous route with 10<sup>2</sup> PFU of WNV, and survival was monitored (Fig. 1A). A higher percentage of *Ifit2*<sup>-/-</sup> mice succumbed to lethal WNV infection than wild-type mice (92% compared to 38%; *P* < 0.001). Moreover, we observed a significant decrease in the mean time to death (10.0 compared to 12.2 days; *P* < 0.01) in *Ifit2*<sup>-/-</sup> compared to wild-type mice. As IFIT genes are implicated in the recognition and control of flaviviruses, coronaviruses, and poxviruses lacking 2'-O methylation of the 5' viral RNA cap structures (17, 18, 36), we also tested whether an absence of *Ifit2* affected pathogenicity of WNV-NS5-E218A. Previous studies established that this WNV mutant virus lacked 2'-O methyltransferase activity and was attenuated in wild-type mice (26) yet more virulent in *Ifit1*<sup>-/-</sup> mice (17, 18). Unexpectedly, WNV-NS5-E218A remained attenuated in *Ifit2*<sup>-/-</sup> adult mice, as no illness or mortality was observed after subcutaneous or intracranial challenge (data not shown). Thus, while *Ifit2* had a protective effect against a virulent strain of WNV *in vivo*, it was not required for restricting viruses lacking 2'-O methylation.

**Ifit2 restricts WNV replication in different tissues.** To begin to understand how *Ifit2* restricts WNV infection *in vivo*, we measured viral burdens in serum, peripheral organs (spleen, kidney, and draining lymph node), and CNS tissues (brain and spinal cord) at different days after subcutaneous infection.

In the serum and draining lymph nodes, we observed a small (3- to 140-fold) yet statistically significant (*P* < 0.02) increase in WNV replication in *Ifit2*<sup>-/-</sup> mice, but only at day 4 after infection (Fig. 1B and C). However, we failed to detect differences in the kinetics or magnitude of WNV infection in the spleen or kidney (Fig. 1D and E) compared to that in mice deficient in IFN-I signaling (34, 37). Overall, compared to mice lacking other innate immune signaling molecules (e.g., *Mavs*, *Irf3*, *Irf7*, or *Ifnar*) (37–40), a loss of *Ifit2* expression had a limited effect on WNV infection in peripheral organs.

We next assessed the effects of an absence of *Ifit2* on WNV infection in CNS tissues (Fig. 1F and G). At day 4 after subcutaneous infection, we observed higher levels (*P* = 0.02) of infection in the brains of *Ifit2*<sup>-/-</sup> mice than in wild-type mice. At day 8 after infection, there was a trend toward increased viral burden in the brain and spinal cord of *Ifit2*<sup>-/-</sup> mice, although this failed to attain statistical significance.

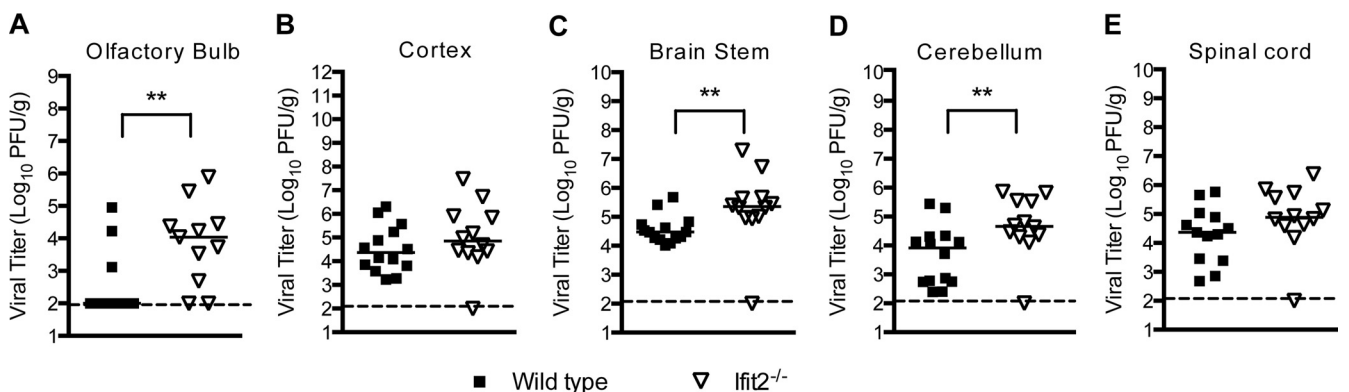


**FIG 1** Survival and viral burden analysis of wild-type and *Ifit2*<sup>-/-</sup> mice. (A) Eight-week-old age-matched wild-type ( $n = 28$ ) and *Ifit2*<sup>-/-</sup> ( $n = 29$ ) mice were infected via the subcutaneous route with  $10^3$  PFU of WNV and monitored for mortality for 21 days. Survival differences were judged by the log rank test and were statistically significant ( $P < 0.0001$ ). (B to G) WNV tissue burden and spread in mice after subcutaneous infection. WNV levels in serum (B), draining lymph node (C), spleen (D), kidney (E), brain (F), and spinal cord (G) of wild-type and *Ifit2*<sup>-/-</sup> mice were measured by quantitative RT-PCR (qRT-PCR) (B and C) or by infectious plaque assay (D to G) of samples harvested at the indicated time points. Data are shown as WNV genome equivalents per microgram of 18S rRNA (r18S) or PFU per gram of tissue for 3 to 15 mice per time point. Solid lines represent the median viral titers, and dotted lines denote the limits of detection of the respective assays. Asterisks indicate values that are statistically significant (\*,  $P < 0.05$ ; \*\*\*,  $P < 0.0001$ ).

Given that a deficiency of *Ifit2* resulted in a trend toward increased viral burden in the CNS at day 8 after infection, we hypothesized that *Ifit2* might restrict infection in some but not all regions, which could obscure small differences. To test this, we infected wild-type and *Ifit2*<sup>-/-</sup> mice and at day 9 harvested tissues from different regions of the CNS and analyzed their viral burden. Notably, viral infection in *Ifit2*<sup>-/-</sup> mice was higher in the olfactory bulb ( $10^{5.0}$  versus  $10^{3.9}$  PFU/g;  $P = 0.004$ ), brain stem ( $10^{6.4}$  versus  $10^{4.9}$  PFU/g;  $P = 0.004$ ), and cerebellum ( $10^{5.3}$  versus  $10^{4.6}$  PFU/g;

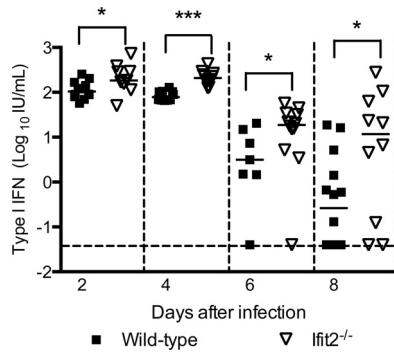
$P = 0.009$ ) than that in wild-type mice (Fig. 2A, C, and D). In comparison, no difference in viral burden was observed in the cerebral cortex or the spinal cord ( $P > 0.1$ ; Fig. 2B and E). Overall, these data indicate that *Ifit2* has a role in restricting WNV infection in specific regions of the brain.

**Cytokine levels in circulation of *Ifit2*<sup>-/-</sup> mice.** Because *Ifit2* has been suggested to inhibit STING, a mitochondrial adaptor protein that recruits TBK1 and IRF-3 to a complex with MAVS to induce IFN- $\beta$  expression (21), we assessed whether a deficiency of



**FIG 2** WNV replication in regions of the CNS of *Ifit2*<sup>-/-</sup> mice after subcutaneous infection. Wild-type and *Ifit2*<sup>-/-</sup> mice were infected with  $10^2$  PFU of WNV via the subcutaneous route. Different regions of the brain were harvested at day 9: olfactory bulb (A), cerebral cortex (B), brain stem (C), cerebellum (D), and spinal cord (E). Tissue homogenates were analyzed for viral burden by plaque assay. Data are shown as PFU per gram of tissue from 11 to 14 mice per time point. Solid lines represent the median viral titers, and dotted lines indicate the limits of detection of the respective assays. Asterisks indicate values that are statistically significant (\*\*,  $P < 0.001$ ).



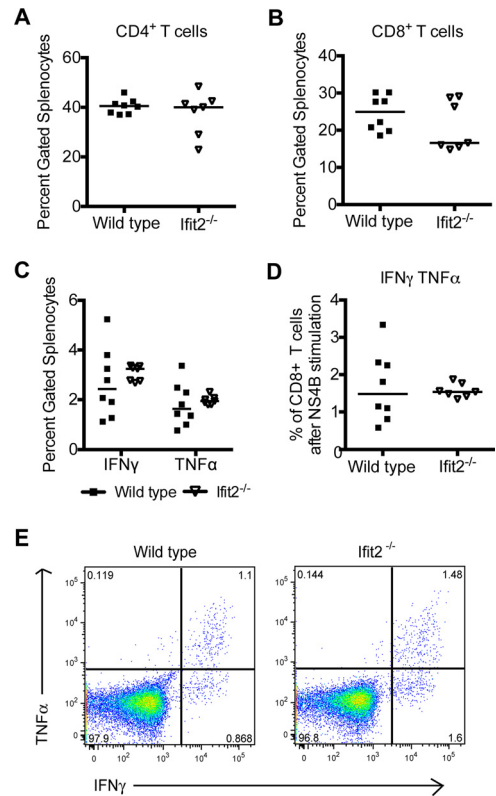


**FIG 3** Cytokine levels in serum of wild-type and *Ifit2*<sup>-/-</sup> mice after infection with WNV. Wild-type and *Ifit2*<sup>-/-</sup> mice were inoculated subcutaneously with 10<sup>2</sup> PFU of WNV, and serum was collected on days 2 to 8 after infection. IFN-I activity was determined in a bioassay. Solid lines represent the median concentrations, and dotted lines indicate the limits of detection of the respective assays. Asterisks indicate differences that are statistically significant by the Mann-Whitney test (\*, *P* < 0.05; \*\*\*, *P* < 0.0001).

*Ifit2* affected IFN-I production. Wild-type and *Ifit2*<sup>-/-</sup> mice were infected with WNV, and the level of biologically active IFN-I in serum was monitored in a validated encephalomyocarditis virus (EMCV) L929 cell bioassay (29). Notably, in *Ifit2*<sup>-/-</sup> mice we observed IFN-I activity levels that were moderately higher (2- to 15-fold; *P* < 0.05) than those in the wild type at every time point tested (Fig. 3). This may be due to increased viral replication in cells of *Ifit2*<sup>-/-</sup> mice or to the proposed loss of negative regulatory effects on cytokine production (41). Because of the elevated IFN-I levels and prior reports suggesting that *Ifit2* might regulate inflammatory responses (41), we also measured cytokine levels in serum from wild-type and *Ifit2*<sup>-/-</sup> mice at day 2 after infection. No differences in interleukin 1β (IL-1β), IL-6, IL-10, IL-12 (p40 subunit), IFN-γ, TNF-α, CXCL1, and CCL4 cytokines and chemokines were observed (data not shown).

**Effect of *Ifit2* on adaptive immune responses to WNV infection.** We next investigated whether an absence of *Ifit2* influenced the development of an effective adaptive immune response during infection, as prior studies suggest that depressed T and B cell responses can result in enhanced WNV replication in the CNS (27, 42–44). Initially, we evaluated T cell responses in the spleen. Equivalent percentages of CD4<sup>+</sup> and CD8<sup>+</sup> T cells were isolated from spleens of wild-type and *Ifit2*<sup>-/-</sup> mice (Fig. 4A and B). Moreover, we observed no difference in the percentage of WNV-specific CD8<sup>+</sup> T cells that expressed IFN-γ, TNF-α, or IFN-γ and TNF-α (double positive) after *ex vivo* NS4B peptide restimulation in wild-type and *Ifit2*<sup>-/-</sup> mice (Fig. 4C to E).

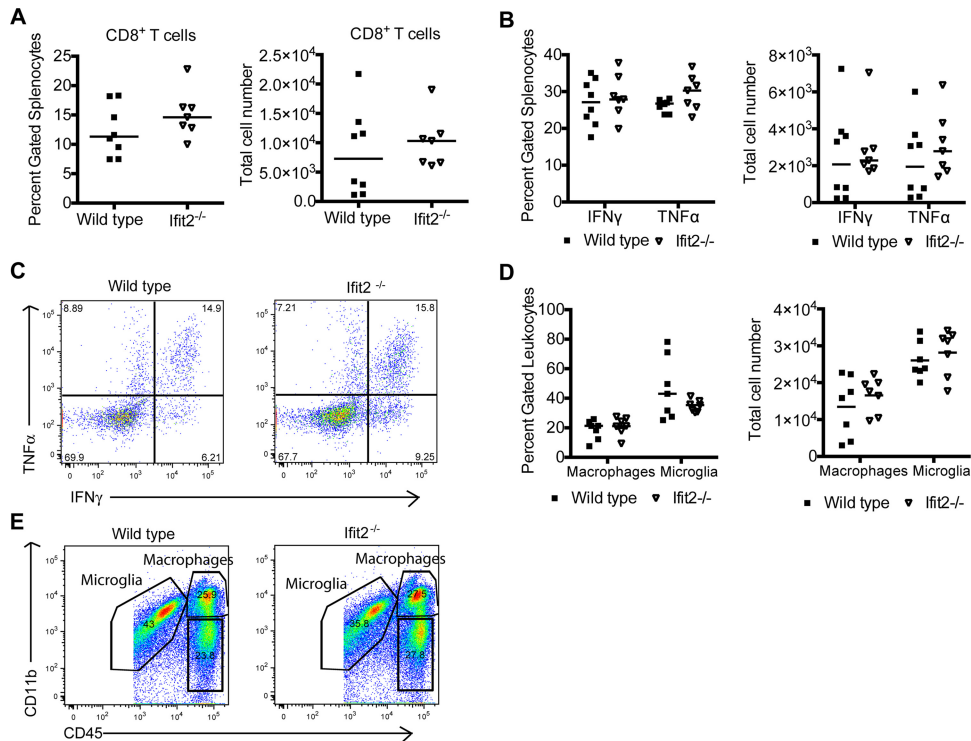
Although peripheral T cell priming was equivalent, we assessed whether leukocyte migration into the CNS might be altered in *Ifit2*<sup>-/-</sup> mice, as infiltrating leukocytes into the brain clears WNV infection (43–45). In the brain, at day 8 after infection, we observed similar percentages and numbers of CD8<sup>+</sup> T cells (*P* > 0.6) (Fig. 5A). Antigen specificity was inferred after intracellular staining of IFN-γ or TNF-α in brain CD8<sup>+</sup> T cells that were restimulated with the NS4B peptide. No statistical differences were observed in the percentage or number of antigen-specific IFN-γ<sup>+</sup> CD8<sup>+</sup> or TNF-α<sup>+</sup> CD8<sup>+</sup> T cells in the brain (*P* > 0.2) (Fig. 5B and C). We also observed similar numbers of CD11b<sup>high</sup> CD45<sup>high</sup> macrophages (*P* = 0.4) and CD11b<sup>high</sup> CD45<sup>low</sup> microglia (*P* > 0.7) (Fig. 5D and E).



**FIG 4** Peripheral T cell responses after WNV infection in *Ifit2*<sup>-/-</sup> mice. Wild-type and *Ifit2*<sup>-/-</sup> mice were inoculated subcutaneously with 10<sup>2</sup> PFU of WNV, and spleens were harvested at day 8 after infection. (A and B) Percentage of CD3<sup>+</sup> CD4<sup>+</sup> (A) or CD3<sup>+</sup> CD8<sup>+</sup> (B) cells in the spleen after being gated on live cells; (C and D) data are shown as the percentage of CD3<sup>+</sup> CD8<sup>+</sup> T cells that expressed intracellular IFN-γ or TNF-α after D<sup>b</sup>-restricted NS4B peptide restimulation *ex vivo*. The differences were not statistically significant, and the data were pooled from two independent experiments with a total of 7 or 8 mice. (E) Flow cytometry dot plots showing intracellular IFN-γ and TNF-α staining after NS4B peptide restimulation and gating on CD8<sup>+</sup> T cells.

To assess the effect of *Ifit2* on WNV-specific humoral responses, we analyzed serum from *Ifit2*<sup>-/-</sup> and wild-type mice for binding to the WNV E protein. At days 6 and 8 after infection, no appreciable differences in levels of anti-WNV E protein IgM were observed between *Ifit2*<sup>-/-</sup> and wild-type mice (*P* > 0.08) (Fig. 6A). Similar levels of WNV-specific IgG also were detected in *Ifit2*<sup>-/-</sup> and wild-type mice at day 8 after infection (*P* > 0.1) (Fig. 6B). Consistent with this, no defect in the levels of neutralizing antibodies was observed in *Ifit2*<sup>-/-</sup> mice (*P* > 0.3) (Fig. 6C). Together, these results suggest that the virological phenotype observed in *Ifit2*<sup>-/-</sup> mice was not due to major defects in T or B cell function.

***Ifit2* controls WNV replication in subsets of primary cells.** Given the variation in tissue-specific virological phenotypes in *Ifit2*<sup>-/-</sup> mice, we speculated that there might be cell-specific functions of *Ifit2* in restricting WNV replication. To assess this, we compared multistep growth kinetics in several different wild-type and *Ifit2*<sup>-/-</sup> primary cells, including macrophages, myeloid dendritic cells, embryonic fibroblasts (MEFs), cortical neurons, and cerebellar granule cell neurons after WNV infection at a low MOI (0.001 to 0.01). In macrophages, cortical neurons, and MEFs, we observed no difference in WNV infection between wild-type and

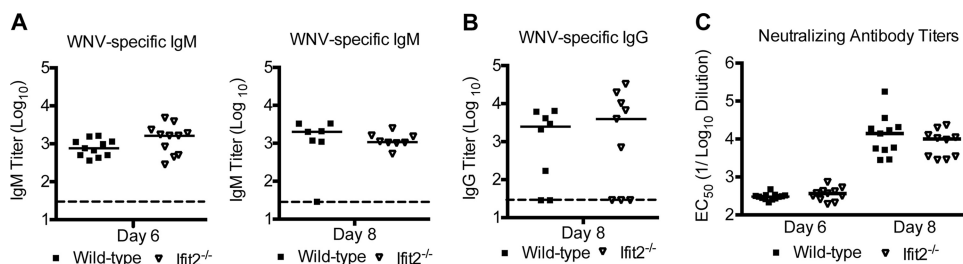


**FIG 5** Leukocyte accumulation in the CNS of *Ifit2*<sup>-/-</sup> mice after WNV infection. Wild-type and *Ifit2*<sup>-/-</sup> mice were infected with 10<sup>2</sup> PFU of WNV by a subcutaneous route, and 8 days later brains were harvested and leukocytes were isolated by Percoll gradient centrifugation. (A) Total percentages and numbers of CD3<sup>+</sup> CD8<sup>+</sup> T cells; (B) total percentages and numbers of CD3<sup>+</sup> CD8<sup>+</sup> T cells that expressed intracellular IFN- $\gamma$  or TNF- $\alpha$  after restimulation with a D<sup>p</sup>-restricted NS4B peptide; (C) flow cytometry dot plots showing intracellular IFN- $\gamma$  and TNF- $\alpha$  cells after peptide restimulation and gating on CD8<sup>+</sup> T cells; (D) total percentages and numbers of activated microglia (CD11b<sup>high</sup> CD45<sup>low</sup>) and macrophages (CD11b<sup>high</sup> CD45<sup>high</sup>). The differences were not statistically significant, and the data were pooled from two independent experiments with a total of 7 or 8 mice. (E) Flow cytometry dot plots indicating gating strategy for defining activated microglia and macrophages.

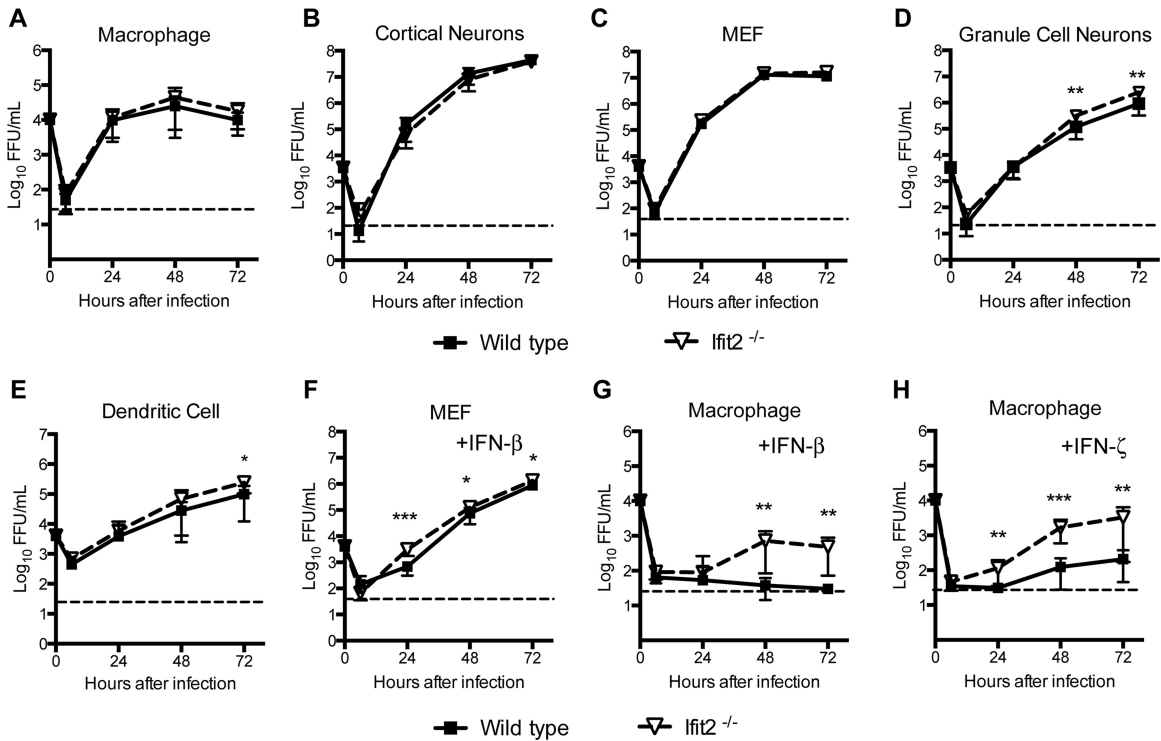
*Ifit2*<sup>-/-</sup> cells (Fig. 7A to C). In comparison, small (2- to 3-fold) yet statistically significant ( $P < 0.05$ ) increases in WNV infection were observed at the 48- and 72-h time points in *Ifit2*<sup>-/-</sup> cerebellar granule cells and at the 72-h time point in *Ifit2*<sup>-/-</sup> dendritic cells (Fig. 7D and E). The antiviral effect of Ifit2 was magnified when cells were primed with different subtypes of IFN-I. The level of WNV replication was higher in IFN- $\beta$ -pretreated *Ifit2*<sup>-/-</sup> MEFs (2- to 5-fold;  $P < 0.05$ ) and macrophages (20-fold;  $P < 0.01$ ) than in wild-type cells (Fig. 7F and G). Similar results were observed after treatment with another mouse IFN-I, IFN- $\zeta$  (also known as limitin) (Fig. 7H). Overall, our results establish a cell

type-specific effect of Ifit2 on WNV replication, which can be magnified in the context of an IFN-I response.

**Ifit2 has a direct role in restricting WNV replication in the CNS.** Previous studies suggested a direct role of Ifit2 in controlling virus replication in the CNS (20, 46). Because our survival and virological experiments through a peripheral route suggested that Ifit2 contributed to the control of WNV infection in selected CNS tissues, we directly introduced virus into the brain after intracranial injection and monitored spread to the contralateral brain regions (olfactory bulb, cerebral cortex, brain stem, spinal cord, cerebellum) on days 2, 4, and 6 after inoculation (Fig. 8A to E).



**FIG 6** WNV-specific IgM and IgG responses in wild-type and *Ifit2*<sup>-/-</sup> mice. Wild-type and *Ifit2*<sup>-/-</sup> mice were infected with WNV via the subcutaneous route, and serum was collected at the indicated time points. The development of WNV-specific IgM (A) or IgG (B) was determined by ELISA using purified WNV E protein. (C) Sera were harvested from animals at the indicated times postinfection and tested for neutralization activity using a focus reduction assay (see Materials and Methods). Data represent the effective reciprocal dilution of sera that produced 50% neutralization of WNV infection (EC<sub>50</sub>). Data are from 7 to 11 mice per group. Differences were not statistically significant ( $P > 0.05$ ) as judged by the Mann-Whitney test.



**FIG 7** WNV replication in wild-type and *Ifit2*<sup>-/-</sup> primary MEFs, myeloid cells, and neurons. Primary cells from wild-type and *Ifit2*<sup>-/-</sup> mice were infected with WNV, and viral replication from 6 to 72 h was measured by focus-forming assay. Shown are data from bone marrow-derived macrophages (MOI, 0.01) (A), cortical neurons (MOI, 0.001) (B), MEFs (MOI, 0.01) (C), cerebellar granule cell neurons (MOI, 0.001) (D), and bone marrow-derived dendritic cells (MOI, 0.001) (E). (F to H) MEFs (F) and bone marrow-derived macrophages (G and H) were pretreated with 10 IU/ml of IFN-β (F and G) or 0.4 ng/ml of IFN-ζ (H) for 4 h and infected with WNV, and viral replication was measured. Asterisks indicate differences that are statistically significant by the Mann-Whitney test (\*, *P* < 0.05; \*\*, *P* < 0.001; \*\*\*, *P* < 0.0001).

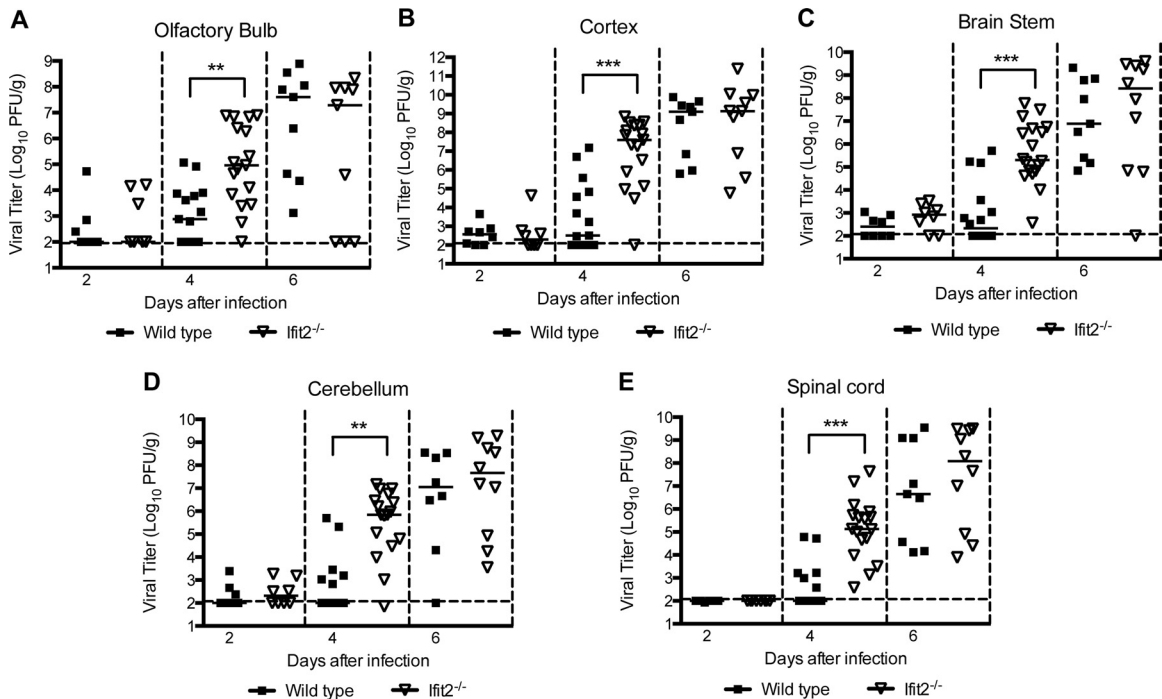
Threshold levels of infection were detected at day 2 after infection in wild-type and *Ifit2*<sup>-/-</sup> mice. By day 4, viral titers of *Ifit2*<sup>-/-</sup> mice were substantially higher in the olfactory bulb ( $10^{5.0}$  versus  $10^{3.1}$  PFU/g; *P* = 0.002), cerebral cortex ( $10^{6.9}$  versus  $10^{3.5}$  PFU/g; *P* = 0.0001), brain stem ( $10^{5.6}$  versus  $10^{2.9}$  PFU/g; *P* < 0.0001), cerebellum ( $10^{5.5}$  versus  $10^{2.8}$  PFU/g; *P* = 0.0002), and spinal cord ( $10^{5.1}$  versus  $10^{2.6}$  PFU/g; *P* < 0.0001) than in wild-type mice. Nonetheless, by day 6, viral titers of wild-type mice became equivalent to those observed in *Ifit2*<sup>-/-</sup> mice. These results suggest that the antiviral action of *Ifit2* restricts WNV spread in the CNS, especially during the early stages of virus spread.

**DISCUSSION**

Subsets of ISGs are responsible for the antiviral effector functions of IFN-I. While several hundred ISGs have been identified by transcriptional profiling and RNAseq studies (47), relatively few have been shown directly to inhibit viral infections (48). IFIT genes encode a family of proteins that are induced after IFN treatment, viral infection, or pathogen-associated molecular pattern (PAMP) recognition (49, 50). Four family members have been characterized in humans (*IFIT1* [ISG56], *IFIT2* [ISG54], *IFIT3* [ISG60], and *IFIT5* [ISG58]), and three members are expressed in mice (*Ifit1* [Isg56], *Ifit2* [Isg54], and *Ifit3* [Isg49]). We initiated these pathogenesis studies because preliminary data suggested that *Ifit2* had antiviral activity against attenuated WNV strains in cell culture (17, 19). Indeed, our analysis here shows a key role for *Ifit2*-dependent restriction of WNV infection *in vivo*. *Ifit2*<sup>-/-</sup> mice

showed enhanced susceptibility to virulent WNV infection, and this was associated with elevated levels of infection in subsets of CNS tissues. A direct effect of *Ifit2* on neuronal cell infection and spread *in vivo* also was suggested, as deficient mice exhibited a higher viral burden in CNS tissues following intracranial inoculation. Finally, a multistep viral growth analysis of primary cells confirmed a cell type-specific antiviral function of *Ifit2*, as *Ifit2*<sup>-/-</sup> myeloid cells, embryonic fibroblasts, and cerebellar granule cell neurons but not cortical neurons showed enhanced infectivity. This phenotype was magnified in the context of priming with IFN-I, which is consistent with the hypothesis that IFN-induced *Ifit2* expression in some cell types restricts WNV replication.

A previous study also suggested an antiviral function of *Ifit2* in a subset of CNS tissues against VSV *in vivo* (20). In that report, *Ifit2* inhibited VSV replication in the cortex, brain stem, midbrain, and cerebellum after intranasal infection. Analogously, in our study, *Ifit2* limited WNV infection in multiple regions of the brain after intracranial infection. These results suggest that *Ifit2* contributes to protection against viral pathogenesis in the brain, which is consistent with the observation that *Ifit2* is induced in many regions of the brain upon viral infection (51). Nonetheless, in our subcutaneous infection model, some regions of the CNS (cerebral cortex and spinal cord) failed to show enhanced infection in *Ifit2*<sup>-/-</sup> mice, although this could reflect the time point that was assessed in these experiments. Moreover, *Ifit2*<sup>-/-</sup> mice did not show increased viral infection in visceral organs such as the spleen and kidney, which could reflect the restricted expression pattern



**FIG 8** WNV replication in regions of the CNS of *Ifit2*<sup>-/-</sup> mice after intracranial infection. Wild-type and *Ifit2*<sup>-/-</sup> mice were infected with 10<sup>1</sup> PFU of WNV in 10  $\mu$ l of HBSS supplemented with 1% FBS via the intracranial route. Different regions of the contralateral brain were harvested at the indicated time points: olfactory bulb (A), cerebral cortex (B), brain stem (C), cerebellum (D), and spinal cord (E). Tissue homogenates were analyzed for viral burden by plaque assay. Data are shown as PFU per gram of tissue for 7 to 18 mice per time point. Solid lines represent the median viral titers, and dotted lines indicate the limits of detection of the respective assays. Asterisks indicate values that are statistically significant (\*\*,  $P < 0.001$ ; \*\*\*,  $P < 0.0001$ ).

of *Ifit2* in those tissues (46). The antiviral role of *Ifit2* has been described in the context of other virus infections. Our data showing an antiviral role of *Ifit2* are consistent with recent data with influenza A virus, although the mechanisms of action may differ. *Ifit2* was suggested to inhibit influenza A virus by forming a complex with *Ifit1* and *Ifit3* to recognize the 5'-ppp moiety on genomic viral RNA (22). In comparison, *Ifit2* expression failed to show any antiviral effects on alphavirus infection *in vitro* (52).

The identification of *Ifit2* as an ISG that restricts WNV infection *in vivo* adds to a small number of studies suggesting that the deletion of individual ISGs can impact WNV pathogenesis. While a redundancy of antiviral ISGs against a given virus might preclude phenotypes when a single antiviral ISG is targeted for deletion, recent studies with WNV suggest otherwise. Using genetically deficient mice, several ISGs, including the PKR (protein kinase R, also known as *Eif2ak2*), RNase L, and viperin (*Rsad2*) genes, were shown to inhibit WNV *in vivo*. Activated PKR phosphorylates the  $\alpha$  subunit of eukaryotic translation initiation factor 2 (eIF2- $\alpha$ ), resulting in a block of protein synthesis (26). RNase L is an endoribonuclease, and, once activated, it cleaves viral RNA and mRNA, leading to a decrease in protein synthesis and viral replication (23). *Pkr*<sup>-/-</sup>  $\times$  *RnaseL*<sup>-/-</sup> mice were vulnerable to subcutaneous WNV infection, with increased mortality and viral replication. Viperin (*Rsad2*) is an endoplasmic reticulum (ER)-associated protein that has inhibitory activity against several viruses, possibly because it inhibits bulk protein secretion, lipid raft formation, and virus budding and localizes to ER-derived lipid droplets, which are required for efficient replication by some RNA viruses (53). *viperin*<sup>-/-</sup> mice infected with WNV also showed in-

creased lethality and/or enhanced viral replication in CNS tissues (12).

How does *Ifit2* inhibit WNV infection? Although more detailed mechanism-of-action studies are required, *Ifit2* has been proposed to inhibit viral translation through its binding to and inhibition of subunits of eIF3 (14). eIF3 is a multisubunit complex that functions during the translation initiation step, including assembly of the eIF2-GTP-Met-tRNA ternary complex, formation of the 43S preinitiation complex, and mRNA recruitment to the 43S preinitiation complex (54). Mouse *Ifit2* and human IFIT2 can block the formation of the 48S complex by binding to eIF3c, and human IFIT2 can block eIF3 binding to the ternary complex by interacting with eIF3e (14). Another IFIT family member, human IFIT1, has been suggested to inhibit influenza and Rift Valley fever virus replication by binding to the 5'-ppp moiety and sequestering RNA from replication (22), and this phenotype reportedly required a complex with IFIT2 and IFIT3. While capped WNV genomic RNA lacks a 5'-ppp end, the negative-strand intermediate may have this moiety exposed (55). Thus, it is possible that *Ifit2* inhibits WNV replication by recognizing a 5'-ppp motif on the negative-strand RNA in a complex with *Ifit1*. Against the need for a complex of *Ifit* proteins for inhibiting wild-type WNV infection, no increased replication or lethality was observed in *Ifit1*<sup>-/-</sup> mice (18). In comparison, recent studies showed that flavivirus mutants lacking 2'-O methyltransferase activity (e.g., WNV-NS5-E218A) were attenuated in wild-type mice (26, 56, 57) yet more virulent in *Ifit1*<sup>-/-</sup> mice (17, 18), suggesting that *Ifit1* recognizes RNA cap structures lacking 2'-O methylation to inhibit viral replication. Although our prior studies suggested that ectopic expres-



sion of Ifit2 might inhibit such viruses, WNV-NS5-E218A remained attenuated in *Ifit2*<sup>-/-</sup> mice; this demonstrates that Ifit2 is not absolutely required for inhibition of WNV strains lacking 2'-O methylation.

Our experiments also revealed no appreciable difference in B and T cell responses between *Ifit2*<sup>-/-</sup> and wild-type mice during WNV infection. This suggests that the absence of Ifit2 does not influence the development of effective adaptive immunity during viral infection.

In summary, our results show that Ifit2 contributes to the antiviral response against WNV *in vivo*, as its targeted deletion was associated with increased lethality and selectively enhanced replication in specific tissues, without an appreciable negative effect of the innate or adaptive T cell or antibody responses. Based on the tissue-specific antiviral effects of Ifit2, we speculate that different cell types might differentially require Ifit2 to restrict WNV replication. The importance of Ifit2 as an antiviral molecule in a given cell type may reflect the qualitative and quantitative ISG signature that is induced (10, 58). A more detailed biochemical and cellular analysis is planned to reveal mechanistically how Ifit2 restricts infectivity of WNV in different subsets of cells.

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