1 **Peli1 facilitates virus replication and promotes neuroinflammation during West Nile virus**

2 **infection**

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Abstract:

 The E3 ubiquitin ligase - Pellino (Peli)1 is a microglia-specific mediator of autoimmune encephalomyelitis. Its role in neurotropic flavivirus infection is largely unknown. Here, we report that mice deficient of Peli1 (*Peli1−/−*) were more resistant to lethal West Nile virus (WNV) infection and exhibited reduced viral loads in tissues and attenuated brain inflammation. Peli1 mediates chemokine and proinflammatory cytokine production in microglia and promotes T cells and macrophages infiltration into the central nervous system. Unexpectedly, Peli1 was required for WNV entry and replication in mouse macrophages, and mouse and human neurons and microglia. It was also highly expressed on WNV-infected neurons and adjacent inflammatory cells in acute post-mortem WNV encephalitis patients. WNV- passaged in *Peli1−/−* macrophages or neurons induced a lower viral load and impaired activation in wild-type microglia thereby a reduced lethality in mice. Smaducin-6, which blocks interactions between 39 Peli1 to IRAK1, RIP1, and IKK_& did not inhibit WNV-triggered microglia activation. Collectively, our findings suggest a non- immune regulator role of Peli1 in promoting microglia activation during WNV infection and identify a potential novel host factor for flavivirus cell entry and replication.

INTRODUCTION

 Pellino (Peli)1, an E3 ubiquitin ligase, is an important regulator in innate and adaptive 48 immunity. It is essential for nuclear factor (NF)- k B activation induced by TIR-domain-containing, adapter-inducing interferon-β (TRIF)/toll-like receptor (TLR)-mediated signaling in innate immune cells such as macrophages and dendritic cells (DCs) [\(1\)](#page-20-0). In microglial cells, where Peli1 is predominantly expressed in the brain, it regulates TLR/MyD88 signaling by promoting degradation of TNF receptor-associated factor 3 (Traf3), a process which leads to microglia activation during the course of induction of experimental autoimmune encephalomyelitis (EAE) [\(2\)](#page-20-1). Peli1 is also implicated in regulation of adaptive immune cell functions. For example, it is a critical factor in the maintenance of peripheral T cell tolerance by regulating c-Rel via its K48 ubiquitination [\(3\)](#page-20-2). Peli1 upregulation plays a role in B cell lymphomas tumorigenesis [\(4\)](#page-20-3).

 West Nile virus (WNV), a mosquito-borne, single-stranded flavivirus that caused outbreaks in Asia, Europe, and Australia, has been the leading cause of viral encephalitis in the United States (US) for more than one decade [\(5,](#page-20-4) [6\)](#page-20-5). The features of acute illness range from WN fever to neuroinvasive conditions, including meningitis, encephalitis, acute flaccid paralysis, and death [\(7\)](#page-20-6). In addition, up to 50% of WNV convalescent patients have been reported to have long-term neurological sequelae or chronic kidney disease [\(8-16\)](#page-20-7). Currently, there is no specific therapeutic agent for treatment of WNV infection, and an approved vaccine is not available for humans. Studies in animal models suggest that WNV-induced central nervous system (CNS) disease is caused by neuronal degeneration, a direct result of viral infection, and/or by bystander damage from the immune response to the pathogen in the CNS [\(17,](#page-21-0) [18\)](#page-21-1) that is induced by infiltrating inflammatory cells and CNS resident cells, including microglia/macrophages, neutrophils, and lymphocytes [\(19-23\)](#page-21-2). Microglial cells, the resident macrophages of the CNS that express various pattern recognition receptors (PRRs), become activated and produce innate proinflammatory molecules upon encountering microbial infection. Microglia activation in the CNS is the hallmark of acute WNV- infection [\(24,](#page-22-0) [25\)](#page-22-1). It was also

72 shown to drive neuronal and synaptic loss which together contributes to memory impairment in

WNV-induced chronic cognitive sequelae [\(26\)](#page-22-2). The underlying immune mechanisms of

microglia activation are not clearly understood. Here, we hypothesize that Peli1 mediates the

activation of microglia and promotes neuroinflammation during lethal WNV infection.

- Unexpectedly, we found that Peli1 is required for WNV entry and replication in both peripheral
- 77 myeloid cells and CNS resident cells. In particular, a defective WNV replication in Peli1-

deficient microglia and neurons directly contributes to attenuated neuroinflammation in the CNS

- and ultimately decreases host susceptibility to lethal encephalitis.
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RESULTS

83 *Peli1^{-/−}* mice were more resistant to systemic WNV infection.

 Peli1 promotes microglia-mediated brain inflammation in the course of EAE induction [\(2\)](#page-20-1). To investigate the role of Peli1 in WNV encephalitis, we infected WT and *Peli1 −/−* mice i.p. 86 with 100 FFU WNV 385-99 strain and monitored daily for survival (Figure 1A). *Peli1^{-/-}* mice (39% survival) were more resistant to WNV infection than WT controls (9.5% survival). To further understand viral pathogenesis, we measured viral burden in the peripheral organs and brain. *Peli1 −/−* mice had lower viremia at days 2 and 3 post infection (pi) and decreased splenic viral load at day 6 compared to WT mice (Figure 1, B and C). WNV crosses the blood brain barrier (BBB) and infects the CNS around day 3 in mice [\(25\)](#page-22-1). Viral RNA levels in *Peli1−/−* mouse brains were more than 15–fold lower than that of WT mice at day 6 pi (Figure 1D). This trend continued but became insignificant at day 9 when both groups of mice started to succumb to lethal WNV infection. On day 6 pi, meningitis (inflammation of the leptomeninge) was noted in 95 WT mice, but not in *Peli1^{-/-}* mice (Figure 1E). Inflammation further spread to the brain parenchyma (encephalitis) as seen in striatum, hippocampus, and cerebellum in both groups of mice at day 9. Encephalitis, particularly perivascular cuffing and microglia activation (cells with elongated nuclei), was much more extensive in WT mice. Thus, CNS inflammation started 99 earlier and was more severe in WNV-infected WT mice compared to *Peli1^{-/−}* mice. No histopathology differences were noted in naïve mouse brains between WT and *Peli1^{-/−}* groups (Supplemental Figure 1).

103 *Peli1^{-/-}* mice exhibited impaired innate cytokine production, but modestly enhanced **adaptive immune responses in the periphery.**

 Peli1 is known to facilitate TRIF-dependent TLR signaling and proinflammatory cytokine production [\(1\)](#page-20-0). Following WNV infection, Peli1 expression was increased in the blood of WT mice (Supplemental Figure 2A). The RNA levels of interferon (*Ifn*)*a* and *Ifnb* at day 6,

 interleukin (*Il)6*, tumor necrosis factor (*Tnf*)*a* at days 3 and 6, and *Il12* at day 3 were all diminished in *Peli1 −/−* mice (Supplemental Figure 2B). The blood plasma protein levels of IL-1β 110 and IL-10 were also reduced in *Peli1^{-/−}* mice (Supplemental Table 1), though no differences 111 were noted in plasma IFN- γ and IL-17 levels between the two groups of mice. To study 112 adaptive immune responses in the periphery, we first measured antibody production in the blood. WNV -specific IgM responses were modestly enhanced in *Peli1 −/−* mice at days 3 and 9 pi (Supplemental Figure 2C). WNV-specific IgG responses were similar between the two groups of mice (Supplemental Figure 2D). Peli1 is also known to negatively regulate T cell signaling [\(3\)](#page-20-2). We next collected the spleen tissues from naive and WNV infected WT and 117 *Peli1^{-/-}* mice. No necrosis was observed in any of the spleen sections examined. There was a trend of the white pulp expansion resulting from germinal center proliferation in WT group at days 3 and 6 pi, but returned to similar levels as naive mice at day 9; in *Peli1 −/−* mice the expansion was continuously enhanced (Supplemental Figure 2, E and F). On day 7, both 121 splenic CD4⁺ and CD8⁺ T cells of *Peli1^{-/−}* mice produced more IFN-γ than WT mice upon *ex vivo* re-stimulation with WNV-specific peptides. CD4⁺ T cells of *Peli1 −/−* mice also induced higher IL-6 and IL-10 production (Supplemental Figure 2, G and H).

 Peli1 was required for WNV entry and viral replication in macrophages and dendritic cell (DC)s which further activated innate cytokine responses.

 Macrophages and DCs are innate immune cells expressing many PRRs and both cell types are permissive to WNV infection in the peripheral organs [\(27\)](#page-22-3). To study Peli1-mediated innate immune responses upon WNV infection, we first measured viral loads in these two cell types by quantitative (q)PCR and focus-forming assays (FFA) and found that both viral RNA levels and viral titers were significantly diminished in primary myeloid *Peli1 −/−* macrophages and DCs compared to WT controls (Figure 2, A and B and Supplemental Figure 3, A and B). We

 next performed virus attachment and entry assays to determine Peli1 involvement in the WNV replication cycle. Macrophages of both groups were incubated with WNV (MOI of 3 or 10) at 4°C for 1 h, allowing the virus to attach to the cell surface. After 1 h incubation, the cells were washed to remove unattached virus, and the amounts of viruses that had attached to the cell 137 surface were measured by qPCR. It was noted that *Peli1^{-/-}* macrophages attached 28% less virus than the WT cells [\(Figure](https://www.sciencedirect.com/science/article/pii/S2211124717314481?via%3Dihub#fig2) 2C). Cells were further incubated at 37°C to initiate viral entry. At 139 1.5 h and 5 h pi, the infected cells were stringently washed to remove free virus as well as cell surface-associated virus, and the intracellular viral RNA was quantified. As shown in Figure 2D, levels of viral RNA in *Peli1 −/−* macrophages were 27% to 54% lower than those of WT cells. These results indicated that Peli1 is involved in WNV attachment and entry. To confirm these results, we performed ultrastructural analysis of macrophages of both groups at 0 min, 5 min, and 10 h after exposure to WNV (Figure 2, E and F). At 0 h, we observed 16 WNV particles ranging between 41 to 46 nm in diameter associated with the plasma membrane in WT macrophages; whereas a total of 7 WNV particles were found in association with the plasma 147 membrane of *Peli1^{-/-}* macrophages. At 5 m pi, 38 viral particles with a diameter of about 42 nm were noted in small uncoated vesicles of WT macrophages**.** However, only 11 WNV particles 149 vith the same size were identified in *Peli1^{-/-}* macrophages. By 10 h, there were 55 WNV viral particles with a diameter of 42 nm inside the double membrane vesicles of WT macrophages. Interestingly, 18 viral particles with much smaller size (about 25 nm) were found inside the double membrane vesicles of *Peli1 −/−* macrophages (Figure 2, E-F). At day 4 pi, virus particles detected in the supernatants of WT macrophages ranged in diameter from 41 nm to 47 nm, which are in similar dimensions to those previously reported [\(28\)](#page-22-4); whereas virions detected in the supernatant of the *Peli1 −/−* macrophages showed significant variations in size ranging from 23.5 nm to 39.5 nm (Figure 2G). These observations suggest that Peli1 not only plays a role in the initial cell attachment and entry, but also in other aspects of WNV life cycle within the host cells. We next determined the infectivity of WNV that was passaged in WT or *Peli1−/−*

159 macrophages. WNV passaged in *Peli1^{-/−}* macrophages had a lower replication rate in WT macrophages versus WNV passage in WT cells (Figure 2H). Furthermore, we challenged WT 161 and *Peli1^{-/−}* mice with 100 FFU of WNV passaged in WT or *Peli1^{-/−}* macrophages. WT mice 162 infected with WNV passaged in *Peli1^{-/-}* macrophages showed an increased survival rate (28.5%, Figure 2I) compared to WT mice infected with WNV passaged in WT cells (0%). *Peli1−/−* mice infected with WNV passaged either in WT macrophages (43%) or *Peli1−/−* cells (56%) also showed similar resistance to lethal WNV infection.

 We next determined if Peli1 mediated immune responses during WNV infection in macrophages and DCs.Following WNV infection, *Peli1* expression was upregulated in macrophages at day 4. In contrast, the other two Pellino family members, including *Peli2* and *Peli3*, remained at low levels (Figure 3A). Compared to WT cells, *Peli1^{-/−}* macrophages had reduced *Ifna* and *Ifnb* RNA levels at days 1 and 4 after WNV infection (Figure 3, B and C). Production of inflammatory cytokines, including IL-6, TNF-α, and IL-12, was also impaired in *Peli1^{-/−}* macrophages at day 4 (Figure 3, D-F). WNV infection in *Peli1^{-/-}* DCs also resulted in diminished levels of *Ifnb*, *Il1b*, and *Il12* compared to WT DCs (Supplemental Figure 3C). PRRs, 174 including TLR3, TLR7, and retinoid acid-inducible gene-I (RIG-I) –like receptor (RLR)s, such as RIG-I and melanoma differentiation antigen 5 (MDA-5), are involved in WNV recognition and 176 trigger the signaling cascade leading to the production of type 1 IFNs and pro-inflammatory cytokines [\(25,](#page-22-1) [29-31\)](#page-23-0). To understand the role of Peli1 in PRR-mediated signaling pathways, we 178 treated WT and *Peli1^{-/-}* macrophages with TLR agonists (Poly I:C for TLR3, CL097 for TLR7, and Poly I:C LyoVec for RLRs). Both Poly I:C or Poly I:C/LyoVec triggered lower *Ifna*, *Ifnb*, *Il6*, 180 and *II12* levels in *Peli1^{-/-}* macrophages. In contrast, stimulation with TLR7 agonist induced similar antiviral cytokine responses in both groups of macrophages (Supplemental Figure 3, D- G). Peli1 is known to be dispensable for MyD88-dependent TLRs, but is required for TRIF- dependent TLR signaling in macrophages and DCs [\(1\)](#page-20-0). Consistent with the previous findings, we demonstrate that Peli1 positively regulates TLR3 and RLR but not TLR7 -mediated innate

 cytokine responses in macrophages. Finally, to understand the effects of defective viral 186 replication on antiviral immunity, we infected WT macrophages with the same dose of WNV passaged once in WT or *Peli1−/−* macrophages. Interestingly, WNV passaged in *Peli1−/−* cells triggered diminished *Ifnb* RNA and IL-6 protein levels in WT macrophages compared to those isolated from WT cells. The levels of reduction induced by WNV passaged in *Peli1−/−* cells were similar to WT virus induced in *Peli1−/−* cells (Figure 3G and Supplemental Figure 3H). Collectively, these data show that Peli1 positively mediates antiviral cytokine responses mainly 192 by a direct involvement in WNV replication life cycle in macrophages and DCs. **Peli1 was predominantly involved in CNS inflammation during WNV infection** Peli1 was previously reported to be highly expressed by neural tissues [\(2\)](#page-20-1). Here, we noted *Peli1* expression was induced in the brain on days 6 and 9 pi (Figure 4A). The mRNA levels of inflammatory cytokines including *Il1b*, *Il6*, and *Tnfa* and chemokines such as *Ccl2*, *Ccl7*, and *Cxcl10* were decreased in *Peli1 −/−* mice (Figure 4B-G). *Ifnb* but not *Ifna* RNA levels 199 were also reduced in *Peli1^{-/−}* mouse brains (Supplemental Figure 4, A and B). To study brain 200 leukocyte phenotype, we performed flow cytometry analysis of these cells isolated at day 9 and 201 found that the number and the percentage of infiltrating CD4⁺ T cells, activated microglial cells 202 (CD11bhiCD45^{lo}), and macrophages (CD11bhiCD45hi) were decreased up to 60% in *Peli1^{-/-}* 203 mice (Figure 4, H-I & Supplemental Figure 4C). To exclude the effects of Peli1 on WNV infection in the periphery tissues, we inoculated WNV intracranially (i.c.) into both groups of 205 mice (Figure 4J). *Peli1^{-/-}* mice showed a similar resistance as in systemic WNV infection (44% 206 versus 11%, *Peli1^{-/−}* versus WT). Collectively, these results indicate that Peli1 is involved in

WNV encephalitis predominantly by mediating CNS infection and induction of

neuroinflammation.

Peli1 promoted microglia activation via facilitating WNV replication in neurons and

microglia in mice and humans.

 Microglial cells are involved in CNS neuroinflammation [\(24,](#page-22-0) [32\)](#page-23-1) and are permissive to WNV infection (Supplemental Figure 5A). *Peli1* is highly expressed on microglial cells [\(2\)](#page-20-1) and was enhanced following WNV infection, however, *Peli2* and *Peli3* remained at low levels (Figure 5A and Supplemental Figure 5B). WNV triggered higher mRNA levels of inflammatory cytokines (*Tnfa* and *Il12*) and chemokines (*Ccl2*, *Ccl4*, and *Cxcl10*) in microglial cells (Supplemental Figure 5C). Interestingly, WNV infection was nearly abolished in both *Peli1 −/−* 218 primary microglial cells (Figure 5, B and C) and Peli1-depleted microglial cells (Supplemental Figure 5D), which was accompanied by diminished levels of inflammatory cytokines and chemokines (Figure 5D and Supplemental Figure 5E). Neurons are most permissive to WNV infection in the CNS [\(18,](#page-21-1) [33\)](#page-23-2). WNV infection upregulates *Peli1* expression on neurons up to 222 200% (Figure 5E). WNV replication was also significantly reduced in *Peli1^{-/−}* neurons compared to WT neurons (Figure 5, F and G). The mRNA levels of chemokines (*Ccl2*, *Ccl7*, *Cxcl10*), inflammatory cytokine (*Il6*), and *Ifnb* were all diminished in WNV-infected *Peli1−/−* neurons at day 4 (Figure 5H, and Supplemental Figure 5F). Western blot analysis showed that the phosphorylation levels of p38MAPK and p65 were both reduced in WNV- infected *Peli1−/−* 227 neurons at day 3 compared to WT controls (Supplemental Figure 5, G and H) which suggests a role of Peli1 in positive regulation of NF-κB and p38MAPK activation. Immunostaining of acute post-mortem WNV encephalitis patient hippocampal tissues show membrane Peli1 on WNV- positive neurons and adjacent inflammatory cells, but not in the same region of the age-match patient controls (Figure 6A). *Peli1* expression was also upregulated following in vitro WNV infection in SHSY-5Y- differentiated neurons and HMC3 cells (human microglial cell line (Figure 6, B and C) and WNV- infected PBMCs (Supplemental Figure 6A). Nevertheless, its expression was not changed in WNV-infected neural stem cell- derived neurons nor THP1- derived macrophages (Supplemental Figure 6A). Knockdown of Peli1

 expression on human microglial cells or SHSY-5Y- differentiated neurons decreased viral loads by 20- 35% at day 4 pi (Figure 6, D and E and Supplemental Figure 6B). Peli1 deficiency also decreased IL-6, CCL2 and CCL5 production in human microglial cells (Figure 6, F-H). WNV infection did not induce inflammatory cytokine or chemokine production in human neurons (data not shown). It was noted that *Peli1* knockdown (36% reduction, Supplemental Figure 6C) in human fetal cortical neural stem cells (hNSCs)- derived neurons led to a 30% decrease on *Ifnb* levels at day 4 pi compared to control-siRNA- treated neurons (Supplemental Figure 6D). These 243 data suggests that Peli1 is involved in WNV replication in neural cells and induction of inflammatory cytokines and chemokines in human microglial cells.

 To determine if defective replication contributes to attenuated inflammatory responses in 146 the CNS, we infected WT microglial cells with WNV passaged in WT or *Peli1^{-/−}* neurons or 247 macrophages. WT microglia- infected with WNV grown in *Peli1^{-/−}* cells had lower viral titers (Figure 7, A and B) and reduced mRNA levels of *Il6*, *Ccl2*, and *Ccl7*, but not *Tnfa* and *Cxcl10* (Figure 7, C -E**)** compared to cells infected with WNV grown in WT cells.It is known that microglia respond to viral infection via activation of p38MAPK [\(24\)](#page-22-0). Microglia infected with WNV 251 passaged in *Peli1^{-/−}* cells also showed lower phosphorylation levels of p38MAPK than infected with WNV passaged in WT neurons (Figure 7F). Smaducin-6, a membrane-tethered palmitic acid-conjugated peptide composed of amino acids 422-441 of Smad6 was reported to interact 254 with Peli1 and disrupt the formation of IRAK1, RIP-1, and IKK ϵ , but not MAPK mediated signaling complexes [\(34\)](#page-23-3). Smaducin-6 treatment did not block WNV infection nor induction of inflammatory cytokine/chemokine production in microglial cells or macrophages, though it decreased levels of type 1 IFNs in macrophages (Figure 7G-H & Supplemental Figure 7). Overall, our results suggest that Peli1 is required for WNV replication in the neural cells which promotes p38MAPK activation in microglia and induction of inflammatory immune responses in 260 the CNS.

DISCUSSION

 Peli1 has been reported to be an important mediator in activation of NF- κ B or p38MAPK in TLR- dependent signaling pathways [\(2\)](#page-20-1). In this study, we have provided evidence demonstrating that Peli1 facilitates WNV replication and mediates innate immunity in the 266 periphery and CNS (Figure 8). In particular, we conclude that Peli1 is predominantly involved in CNS neuroinflammation during WNV infection based on the following facts: First, although Peli1 promotes WNV replication in myeloid cells (macrophages and DCs), it also positively regulates antiviral innate immune responses in these cells, which compromise its overall pathogenic effects in the periphery. Second, compared to myeloid cells, Peli1 expression is highly enriched 271 in CNS resident cells and is even upregulated following WNV infection. Third, Peli1 facilitates WNV replication in microglia and neurons, especially in the latter which are the major cells 273 infected during in vivo challenge. It also positively mediates NF-KB and /or p38MAPK activation in these cells, and boosts a robust inflammatory cytokine and chemokine production, which 275 attracts more inflammatory cells infiltrated from the periphery and ultimately contributes to lethal WNV encephalitis. Thus, Peli1 synergistically promotes virus dissemination and inflammation in 277 the CNS. Lastly, *Peli1^{-/-}* mice displayed similar levels of resistance compared to WT mice following systemic and direct intracranial WNV infection. This further indicates that Peli1 promotes WNV- induced pathology primarily in the CNS.

 WNV-induced CNS disease is partially caused by bystander damage from both the immune response induced in the CNS residential cells [\(17,](#page-21-0) [18\)](#page-21-1) and by infiltrating inflammatory 282 cells [\(19-23\)](#page-21-2). *Peli1^{-/−}* mouse brains had a significantly decreased number of infiltrating CD4⁺ T cells, activated microglia, and macrophages. Both CCL2 and CCL7 are involved in the monocytosis and monocyte accumulation in the brain [\(35\)](#page-23-4), whereas CXCL10 helps to recruit the antigen specific T cells [\(36\)](#page-23-5). Microglial cells are the major producers of inflammatory cytokines and chemokines in the CNS following WNV infection [\(25,](#page-22-1) [37\)](#page-24-0). In vitro WNV infection in *Peli1−/−* or Peli1- depleted mouse and human microglia resulted in a lower viral load and induced an

 impaired production of IL-6, CCL2, CCL7, and CXCL10. Neurons are the primary targets during 189 in vivo WNV replication in the CNS [\(18,](#page-21-1) [38\)](#page-24-1). WNV passaged in *Peli1^{-/−}* neurons induced similarly lower levels of p38MAPK phosphorylation in WT microglia, and this was accompanied 291 by impaired chemokine CCL2 and CCL7, but not CXCL10 production in these cells. Collectively, 292 our results suggest that Peli1 mediates proinflammatory cytokine and chemokine production predominantly via facilitating WNV replication in neural cells (microglia and neurons), which ultimately leads to macrophages/monocytes and T cell infiltration into the CNS.

 In line with findings in a previous report [\(37\)](#page-24-0), we did not note significant induction of inflammatory cytokine responses in human neurons following WNV infection. Interestingly, WNV induced higher *Peli1* expression on mouse neurons than on microglia. WNV-infected *Peli1−/−* 298 mouse neurons had reduced levels of NF-_KB and p38MAPK activation accompanied by an impaired production of inflammatory cytokines and chemokines. It was also noted that noninfected *Peli1 −/−* mouse neurons also had lower levels p38MAPK activation despite of normal basal levels of cytokine production compared to WT controls (data not shown). Thus, it is likely Peli1-positively regulates inflammatory cytokine and chemokine responses in WNV-infected mouse neurons via activation of NF- κ B. Peli1 is known to activate NF-KB signaling via interaction with RIPK1. RIPK3, another RIPK family member was recently shown to mediate neuronal chemokine induction and recruit T lymphocytes and inflammatory myeloid cells in the CNS [\(39\)](#page-24-2). Whether Peli1 interacts with RIPK3 and regulates inflammatory responses remains to be investigated.

 Smad6 and Smad7 are critical mediators for effective TGF-β1-mediated suppression of IL-1R/TLR signaling, by simultaneous binding to discrete regions of Peli1 [\(40\)](#page-24-3). Smaducin-6, which is composed of amino acids 422-441 of Smad6, has been reported to disrupt the formation of IRAK1-, RIP1-, and IKK ε -, mediated TRIF signaling complexes, but not phosphorylation of p38MAPK in macrophages [\(34\)](#page-23-3). Smaducin6 did not inhibit WNV replication

 in microglia or macrophages. Further, Smaducin-6 did not block the production of inflammatory cytokines and chemokines in WNV-infected microglia and macrophages. These results support a role of Peli1 in induction of inflammatory responses in both cell types during WNV infection via promoting p38MAPK activation.

 Consistent with the findings in myeloid and CNS resident cells, the production of type I IFNs, inflammatory cytokines, and chemokines in the blood and CNS tissues were all impaired in WNV-infected *Peli1 −/−* mice. Our results suggest a role of Peli1 as a positive regulator of innate immunity. There are conflicting reports on the role of Peli1 in induction of type I IFN response during viral infection. For example, Xiao *et al*. found that Peli1 negatively regulates IFN signaling in microglia and macrophages during vascular stomatitis virus infection in the CNS [\(41\)](#page-24-4). Another group has shown that Peli1 interacts with DEAF1 and positively regulates IFNβ production following Sendai virus infection [\(42\)](#page-24-5). While our results are in line with the latter, the main difference between ours and the previous findings is that Peli1 positively mediates immune induction predominantly via facilitating WNV entry and replication.

 WNV life cycle includes attachment/entry, translation, RNA replication, and egress of viral particles. Although the virus relies heavily on host proteins during its life cycle [\(43,](#page-24-6) [44\)](#page-24-7), our understanding of the molecular interactions of virus and mammalian host cells and their impacts on viral pathogenesis is currently limited. Previous RNA interference screening of human genes responsible for interaction with WNV proteins identified the Ub ligase CBLL1 as being critical for WNV internalization during in vitro infection [\(45\)](#page-24-8). Several host factors, including AXL and TIM1 have been shown to be involved in flavivirus replication from in vitro cell culture studies. However, their roles remain to be confirmed in vivo [\(46-48\)](#page-24-9). In this study, by using both in vivo and in vitro models, we have identified Peli1 as a novel host factor required for WNV initial cell attachment, and entry, a process which further promotes TLR-mediated inflammatory responses in mouse and human neural cells. Interestingly, the size of the virions generated after

passaging WT WNV once in *Peli1−/−* cells was reduced significantly. No genetic changes were

 noted in the passaged viruses (data not shown). However, the reduction of virion size is likely 340 due to an altered protein composition or changes on the ratio of viral proteins in the virions 341 during virus assembly [\(49\)](#page-25-0). In particular, the smaller viral particles produced in *Peli1⁻* cells are consistent with the size of recombinant subviral particles assembled in the endoplasmic reticulum during flavivirus infection in mammalian cells reported previously [\(50\)](#page-25-1). The subviral particles consist of the pre-membrane (PrM)-E structural proteins, retain functional properties and are transported from the endoplasmic reticulum through the secretory pathways to undergo cleavage maturation by the cellular protease furin. Mutation of the furin recognition site of PrM of tick-borne encephalitis virus resulted in secretion of the smaller subviral particles. Altogether, our data suggest that Peli1 is involved in WNV attachment, entry and assembly. The smaller 349 virions generated *Peli1^{-/-}* cells had reduced replication rates in WT cells and triggered attenuated inflammatory responses in these cells which contribute to a higher resistance in WT mice. These results further support our hypothesis that Peli1- mediates inflammatory responses and promotes WNV encephalitis via facilitating virus replication. Consistent with our findings in the murine model, we also demonstrate Peli1 expression is associated with WNV infection and inflammatory cell activation in acute post-mortem WNV encephalitis patient hippocampal tissues. Results from this study provide us with a better understanding of the mechanisms by which WNV induce lethal encephalitis. They will ultimately help to identify therapeutic targets for intervention, such as Peli1 as a new strategy for development of the inhibitors of virus 358 replication to prevent and treat WNV-induced encephalitis. Lastly, Peli1 is expressed on many cell types and is highly enriched in the CNS tissues. Future investigation will be needed to determine the role of Peli1 in other virus models, in particular the neurotropic flaviviruses.

METHODS

 Mice: 5-8-week-old C57BL/6 (B6) mice were purchased from the Jackson Laboratory. 365 Peli^{1-'-} mice (on a B6 background) [\(2,](#page-20-1) [51\)](#page-25-2) were bred at the University of Texas Medical Branch (UTMB). Both female and male mice were used in this study and were age- and sex- matched. Mice were inoculated intraperitoneally (i.p.) with100 FFU of WNV 385-99 [\(52,](#page-25-3) [53\)](#page-25-4) or WNV 385- 368 09 passage once in WT and *Peli1^{-|-}* macrophages. In some experiments, mice were challenged intracranially (i.c.) with 5 FFU of WNV 385-99.

 Cells: Bone marrow (BM)-derived dendritic cell (DC)s, macrophages, and primary microglia cultures were isolated as described previously [\(2,](#page-20-1) [54\)](#page-25-5). Neurons were generated according to [\(55\)](#page-25-6) with slight modifications. After the dissection and dissociation of cortices of mouse embryos (E18.5), cells were enriched using a mouse neuron isolation kit (Miltenyi Biotec) and cultured for 5 days in neurobasal medium containing B-27 supplement (Invitrogen). BV2 cells were kindly provided by Dr. A Cardona (University of Texas San Antonio). BM-DCs or macrophages, microglia, and BV2 cells were infected with WNV at a MOI of 0.1, or 0.02 and neurons were infected at a MOI of 0.003. SH-SY5Y cells were cultured in F12K medium and EMEM (Invitrogen) and seeded in the 6-well plate for 1 day, then replaced with fresh medium with 30 uM Retinoic acid 1% (Sigma) and B27 Supplements (Gibco,17504) for 5 days, the cells were differentiated to neurons. Smaducin-6 or Pal-Scram peptides [\(34\)](#page-23-3) were purchased from Sigma-Aldrich and used at 100 nm 1 h after infection. Supernatants and cells were harvested at 382 24 h and 96 h pi to measure viral load and cytokine production. In some experiments, WT and *Peli1^{-/-}* macrophages or neurons were infected with WNV 385-99. Culture supernatants were then harvested at day 4 for virus titration by FFA. Equal titers of viruses from WT and *Peli1*−/− culture were subsequently used for in vivo and in vitro infection studies. **FFA:** Vero cells were incubated with sample dilutions for 1 h. A semi-solid overlay containing 0.8% methylcellulose (Sigma-Aldrich), 3% fetal bovine serum, 1% Penicillin-

Streptomycin, and 1% L-glutamine was added. At 48 h, the semisolid overlay was removed,

 cells were washed, and fixed with 1:1 of acetone: methanol solution for at least 30 min at -20ºC. Cells were next subjected to immunohistochemical staining with a rabbit WNV polyclonal antibody (the World Reference Center for Emerging Viruses and Arboviruses (WRCEVA), T35502) followed by goat anti-rabbit HRP-conjugated IgG (KPL, 474-1516) for 1 h. Cells were next incubated with a peroxidase substrate (Vector Laboratories) until color developed. The number of foci was used to calculate viral titers.

 qPCR: Samples were re-suspended in Trizol (Invitrogen) for RNA extraction. cDNA was synthesized by using a qScript cDNA synthesis kit (Bio-Rad). The sequences of the primer sets for WNV envelope (*Wnve*), *Peli1*, *Peli2*, *Peli3*, and cytokines cDNA and PCR reaction conditions were described previously [\(2,](#page-20-1) [25,](#page-22-1) [36,](#page-23-5) [53,](#page-25-4) 56). The assay was performed in the CFX96 real-time PCR system (Bio-Rad). Gene expression was calculated based on C_t values by using the formula 2^ \cdot ^{[C}t^{(target gene)-C}t^{(GAPDH} or β-actin)] [\(53\)](#page-25-4).

 Viral attachment and entry assays: BM-macrophages were incubated with WNV (MOI of 3 or 10) at 4°C for 1 h, allowing the virus to attach to the cell surface. After 1 h incubation, infected cells were washed three times with cold PBS to remove unbound virions. Cell surface- associated viruses were removed by washing with cold alkaline-high-salt solution (1 M NaCl and 50 mM sodium bicarbonate, pH 9.5). After twice cold-PBS washes, the cells were harvested, and suspended in 3 ml DMEM medium containing 2% FBS. Total cells were collected by 407 centrifugation at 1,000 \times g for 5 min. The cell pellets were resuspended in Trizol for RNA extraction to measure viral titer by qPCR. Some cells were further incubated at 37°C to initiate viral entry. At 1.5 h and 5 h pi, the infected cells were stringently washed to remove free virus as well as cell surface-associated virus, and the intracellular viral RNA was quantified by qPCR.

 Transmission electron microscopy (TEM): Cells were washed with ice cold PBS and incubated on ice for 15 min before exposure to WNV (MOI= 10) for 1 h. Cells were then rinsed with ice cold PBS, resuspended in pre-warmed media and incubated at 37°C. At 0 h, 5 min, and

 10 h pi, cells were pelleted and fixed for at least 1 h in a mixture of 2.5% formaldehyde and 0.1% glutaraldehyde in 0.05 M cacodylate buffer (pH 7.3) to which 0.01% picric acid and 0.03% CaCl² were added (EM fixative). The pellets were washed in 0.1 M cacodylate buffer followed by post-fixation in 1% OsO4 in 0.1M cacodylate buffer for 1 h, washed and *en bloc* stained with 2% aqueous uranyl acetate for 20 min at 60°C. The pellets were next dehydrated in ethanol, processed through propylene oxide, and embedded in Poly/Bed 812 (Polysciences). Sections were cut on Leica EM UC7 ultramicrotome (Leica Microsystems), stained with lead citrate, and examined in a Philips CM-100 transmission electron microscope at 60 kV. Images were acquired with a Gatan Orius SC200 digital camera. In some experiments, supernatants of WNV-infected cells were concentrated at day 4 pi using a 3 kD spin columns (Sartoris, 424 Germany). The concentrated supernatants were centrifuged for 10 min at 3000 xg to remove debris. Next, nickel grids were incubated with clarified supernatants for 10 min followed by glutaraldehyde fixation and 2% uranyl acetate staining. Micrographs were taken using a CM100 transmission electron microscope (Philips).

 Flow cytometry: Brain leukocytes were isolated as described before [\(20\)](#page-22-5) and were stained with antibodies for cell surface markers, including CD3 (eBioscience, clone 145-2C11), CD4 (eBioscience, clone GK1.5), CD8 (eBioscience, clone 53-6.7), CD11b (eBioscience, clone M1/70, Gr-1-(eBioscience, clone RB6-8C5) and CD45 (BD Biosciences, clone 30-F11), fixed with 1% paraformaldehyde in PBS and examined with a C6 Flow Cytometer (BD Biosciences). Dead cells were excluded on the basis of forward and side light scatter.

 Western blot: Protein was dissolved in 1X SDS loading buffer, separated on SDS- PAGE gels, electroblotted onto nitrocellulose membranes, probed with primary and secondary antibodies, and detected using the enhanced chemiluminescence (ECL) system (Pierce). The primary antibodies against phospho-p38 (Thr180/Tyr182, #9211s), p38 (#9212), Phospho-NF- κB RelA (Ser 468, #3039) and NF-κB (#3034) were from Cell Signaling Technology (Beverly) and mouse monoclonal anti-β-actin (Sigma-Aldrich, clone AC-15) was used as a loading control.

 Histology: Mice were transcardially perfused with PBS. Brains and spleens were 441 removed and placed in 4% paraformaldehyde (PFA) for 3 days at 4°C, followed by 70% ethanol 442 before embedding in optimal cutting temperature compound. H & E staining was performed at 443 the Histopathology Laboratory Core at Baylor College of Medicine.

 Immunohistochemistry: Human paraffin-embedded hippocampal tissues were obtained from fatal WNV encephalitis cases (generously provided by Dr. Beth Levy at St. Louis University, St. Louis, MO) and control post-mortem cases from patients without neurologic diseases (St. Louis, MO). Sections were deparaffinized in Xylene, rehydrated in serial dilutions of ethanol and boiled in citrate buffer (pH 6.0) for 30 min for antigen retrieval. Next, they were incubated for 2 h at room temperature in blocking solution (10% serum and .2% Tween 20 in PBS) to permeabilize and block nonspecific binding. This was followed by incubating with primary antibodies for Peli1 (Santa Cruz Biotechnology, sc-271065, 1:50), WNV antigen (UTMB WRCEVA, T35502, 1:500) and NeuN (Millipore, clone A60, 1:20) or isotype matched IgG in 453 PBS supplemented with10%serum O/N at 4°C. After washing, sections were incubated for 1 h at room temperature with fluorescently conjugated secondary antibodies (Life Technologies, 1:500). Autofluorescence was darkened using .5% Sudan Black B (diluted in 70% Ethanol) incubated for 10 min. After washing, nuclei were counterstained with Dapi and coverslips were applied with ProLong™ Gold Antifade Mountant (ThermoFisher Scientific). Immunofluorescence was captured using a Zeiss LSM 510 laser-scanning confocal microscope.

 Statistics:Survival curve comparisons were performed using Prism software (GraphPad) statistical analysis, which uses the log rank test. Values for viral burden, cytokine 461 production, and antibody and T cell responses experiments were presented as means \pm SEM.

P values of these experiments were calculated with a non-paired 2-tailed Student's t test.

Statistical significance was accepted at *P* < 0.05.

 Study approval: All experiments were performed in compliance with and under the approval of the Animal Care and Use Committee at UTMB.

AUTHOR CONTRIBUTION

H.L., E.R.W., S.C., P.W., W.Z., S.J.T, P.V.A., and T.W. designed the experiments. E.R.W., H.L.,

E.M., J.A. S., W.R., L.L.V., S.Z., J.G., N.E.B., C.C., G.X., G.L., R.T., V.L.P., and T.W. performed

the experiments. E.R.W., H.L., E.M., J.A.S., S.Z., B.H.P., C.C., V.L.P., W.Z., R.S.K., P.V.A., and

- T.W. analyzed the data. S.C., S.P.Y., W.Z., and S.C.S. provided key reagents. H.L., E.R.W.,
- J.A.S., P.W., R.S.K., P.V.A. and T.W. wrote the manuscript.
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ACKNOWLEDGEMENTS

- This work was supported in part by the Institute for Human Infections & Immunity at UTMB
- (T.W.), NIH grants R01 AI099123 (T.W.), R01 AI27744 (T.W.), R01NS079166 (S.J.T),
- R01NS095747 (S.J.T), R01DA036165 (S.J.T), U19 AI083019 (R.S.K.) and R01 NS052632
- (R.S.K.), and R01 EY022694 and R01 EY026629 (W.Z.). J.A.S was supported by NIH grant
- F31 AI124662-01. P.V.A and V.P were partially supported by NIH grant R24 AI120942. We
- thank Texas A&M Institute for Genomic Medicine for *Peli1 −/−* mice, Dr. A Cardona from the
- University of Texas San Antonio and Dr. Partha Sarkar from UTMB for BV2 cells and SHSY-5Y
- cells, Dr. Beth Levy for human post-mortem hippocampal tissues, Ms. Lan Pang for technique
- support, and Dr. Linsey Yeager for assisting in manuscript preparation.
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COMPETING FINANCIAL INTERESTS

- The authors declare no competing financial interests.
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Figure 1. *Peli1 −/−* **mice are more resistant to lethal WNV infection***.* **A.** Survival of WT and *Feli1^{-/−}* mice after i.p. injection with WNV 385-99. *n* = 21 and n = 23 for WT and *Peli1^{-/−}* mice respectively. ** *P* < 0.01 compared to WT group (log rank test). **B.** Viremia was determined by 651 using FFA at days 2 and 3 post infection (pi). Data are presented as mean \pm SEM (n= 3 to 6) of samples collected from one representative of three similar experiments. **C-D.** Viral load in the samples collected from one representative of three similar experiments. **C-D.** Viral load in the spleen and brain of infected and non-infected (NF) mice was determined by qPCR assay. Data 654 are presented as mean \pm SEM, $n = 7$ to 12 collected from 3 independent experiments. For 655 Panels **B-D**, ** $P < 0.01$ or * $P < 0.05$ compared to WT group (Unpaired t test). **E**. H & E stair Panels **B-D**, ** *P* < 0.01 or **P* < 0.05 compared to WT group (Unpaired t test). **E**. H &E staining. Representative images (20X) shown are brains collected from 4 to 5 WNV-infected mice per group at indicated time points. Scale bar: 100um.

 Figure 2. Peli1 facilitated WNV replication in macrophages and promoted high mortality in vivo. A-B*.* Viral load of WNV-infected macrophages was measured by qPCR (**A**) and FFA (**B**). Data shown are representative of five similar experiments and are presented as means ± SEM, *n* = 3 to 6. **C-D**. Macrophages were infected with WNV for 1 h at 4ºC, washed and 667 collected to measure intracellular viral RNA by qPCR in the attachment assay (C). For virus 668 entry (D), cells were subsequently resuspended in medium and incubated at 37°C. At indica entry (D), cells were subsequently resuspended in medium and incubated at 37°C. At indicated time points, cells were washed to determine intracellular viral RNAs. n= 6. **E-F**. Thin-section transmission electron micrographs of WNV- infected macrophages. **E**. Viral particles were observed at the plasma membrane at 0 min, in small uncoated vesicles at 5 min and in double membrane vesicles at 10 h. **F***.* Quantitation of 10 fields of view of ultrathin sections. **G.** Negative staining micrograph of WNV. Virions range in size from 41 nm to 47 nm and 23.5 nm to 39.5 nm in the supernatants of WT and *Peli1* −/− macrophages at day 4 pi respectively. **H***.* WT macrophages were infected at MOI of 0.02 with viruses passaged in WT (WTP) and *Peli1*−/− (*Peli1*−/− P) macrophages. Viral load was measured by FFA at day 4 pi. Data shown are 677 representative of two similar experiments and are presented as means \pm SEM, $n = 4$. Data in A- **D** & **H** were analyzed by Unpaired t test. ** *P* < 0.01 or **P* < 0.05 compared to WT group. **I.** Survival rate of mice i.p. injected with 100 FFU of WNV passaged in WT macrophages (WTP), 680 and *Peli1^{-/−}* (*Peli1^{-/−}P*) macrophages. *n*= 7 or 9. **P* < 0.05 compared to WT mice infected with WTP (WT (WTP), Log-rank test).

688 Data shown are representative of three similar experiments and are presented as means \pm 689 SEM, $n = 3$. $^{***} P < 0.01$ compared to non-infected (NF) group (Unpaired t test). **B- F.** Cytok SEM, $n = 3$. $#$ $P < 0.01$ compared to non-infected (NF) group (Unpaired t test). **B- F.** Cytokine

RNA or protein levels were measured at indicated time points by qPCR and Bioplex

691 respectively. Data represent means \pm SEM of 5 to 10 samples collected from 2 independent

experiments. ** *P* < 0.01 or **P* < 0.05 compared to WT group. **G.** WT macrophages were

infected at MOI of 0.02 with WNV passaged in WT (WTP) or *Peli1*−/− (*Peli1*−/− P) macrophages.

694 At day 4 pi, cytokine production was determined by using qPCR. Data are presented as the fold increase compared to mock- infected (means \pm SEM) and are representative of two similar

695 increase compared to mock- infected (means \pm SEM) and are representative of two similar 696 experiments, $n = 3$ to 6 per group. *P < 0.05 compared to WTP group (Unpaired t test).

experiments, $n = 3$ to 6 per group. $*P < 0.05$ compared to WTP group (Unpaired t test).

 Figure 4. *Peli1* **mediated neuroinflammation in the CNS after WNV infection. A.** RNA levels of *Peli1* in the brain of WT mice following WNV infection were determined by qPCR assay. Data are presented as the means \pm SEM of samples pooled from 2 to 3 independent 703 experiments, $n= 8$. $** P < 0.01$ compared to non-infected (NF) group (Unpaired t test). **B-G.** RNA levels of cytokines and chemokines in the brain at indicated time points were determined by 705 qPCR assay. Data are presented as the means \pm SEM of pooled 6 to 10 samples from 2
706 independent experiments. **H-J**. Brain leukocyte infiltration following WNV infection. **H.** The independent experiments. **H-J***.* Brain leukocyte infiltration following WNV infection. **H.** The number of brain leukocytes are presented as the means \pm SEM of 9 to 10 mice from 2 708 independent experiments, including naïve microglia cells (NMG), activated microglial cells
709 (AMG), macrophages (M), CD4⁺ and CD8⁺ T cells on day 9 pi analyzed by flow cytometry. (AMG), macrophages (M), CD4⁺ and CD8⁺ T cells on day 9 pi analyzed by flow cytometry. **I.** Representative flow image is shown. For **B-G** & H, ** *P* < 0.01 or **P* < 0.05 compared to WT The group (Unpaired t test). **J.** Survival of WT and *Peli1^{-/−}* mice after i.c. injection with WNV 385-99.
712 *n* = 9 per group. **P* < 0.05 compared to WT group (Log-rank test). $n = 9$ per group. $*P < 0.05$ compared to WT group (Log-rank test).

 $\frac{716}{717}$ **Figure 5. WNV infection and induction of immune responses in microglia and neurons**.

718 **A.** RNA levels of *Peli1* in WT microglia at day 4 pi were determined by qPCR assay Data are presented as means \pm SEM and are representative of 2 independent experiments, n =5. ^{##} P presented as means \pm SEM and are representative of 2 independent experiments, n =5. $^{\#}P$ <

0.01 compared to non-infected (NF) group (Unpaired t test). **B- D.** Primary microglia were

721 infected with WNV 385-99 and harvested at indicated time points. Data are presented as the means \pm SEM of pooled 4 to 8 samples from 2 independent experiments. **B-C.** Viral load was

means ± SEM of pooled 4 to 8 samples from 2 independent experiments. **B-C.** Viral load was

measured at day 4. **D.** Cytokine RNA levels were measured at indicated time points by qPCR.

 E- H*.* Primary mouse neurons were infected with WNV 385-99 and harvested at indicated time points. Data are presented as means ± SEM, n =6 per group. **E.** RNA levels of *Peli1* in WT

726 neurons at day 4 pi. ^{##} P < 0.01 compared to NF group (Unpaired t test). **F-G**. Viral load was

measured at indicated time points by qPCR (**F**) and FFA (**G***)*. **H**. *Il6, Ccl2, Ccl7*, and *Cxcl10*

RNA levels were measured at indicated time points by qPCR. For **B-D & F-H** panels ** *P* < 0.01

or **P* < 0.05 compared to WT group (Unpaired t test).

 Figure 7. Peli1 promoted p38MAPK activation in microglia via facilitating WNV replication. **A- F.** BV2 cells were infected at MOI of 0.02 with viruses passaged in WT (WTP) and *Peli1^{-/-}* (*Peli1^{-/-}P*) macrophages or neurons. **A- B.** At day 4 pi, viral load was measured by q PCR (**A**) or FFA (**B**). **C- E.** IL-6 and TNF-α production and Cc/2, Cc/7, and Cxc/10 RNA levels qPCR **(A)** or FFA (**B**). **C- E.** IL-6 and TNF-α production and *Ccl2*, *Ccl7*, and *Cxcl10* RNA levels 753 were measured at day 4 by Bioplex or qPCR respectively. Data are presented as means \pm SEM and are representative of 2 similar experiments, n =4. **A-E**, ** *P* < 0.01 compared to WT group (Unpaired t test). **F**. Western blot assay for p38MAPK activation. One representative of two samples per group was shown. **G**-**H**. BV2 cells were infected at MOI of 0.02 with WNV 385-99 and treated with Smaducin-6 or control peptides at 1 h pi. **G.** Viral load was measured at day 4 758 pi by qPCR. **H.** Cytokine and chemokine levels were measured at day 4 by qPCR. Data are
759 presented as fold increase compared to mock- infected (means ± SEM) and represent 8 759 presented as fold increase compared to mock- infected (means \pm SEM) and represent 8
760 samples pooled from 2 independent experiments. samples pooled from 2 independent experiments.

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Figure 8. Proposed model illustrating Peli1 promotes virus replication and

neuroinflammation during WNV infection.

Supplemental Figure 1. **Histology study of WT and** *Peli1 −/−* **mice.** Representative images 780 (20X) shown are H&E staining of brains from non-infected WT and *Peli1^{-/−}* mice. Samples were collected from 4 mice per group of 2 independent experiments. Scale bar: 100µm.

B 200 Day 3 150 A Ċ Fold change 100 D IgM IgG 50 $80¹$ 2.0 expression WT $\overline{\mathbf{3}}$ 60 P eli t [/] 1.5 $\overline{3}$ $\frac{1}{2}$ 40 \overline{a} 응 1.0 Ifna **Ifnb** $II6$ Tnfa $II12$ Peli1 50 0.5 Fold change Day 6 $\ddot{}$ $40 -$ **WNV** NI 30° $0₀$ $\overline{Day 3}$ $Day 6$ Day 3 Day 9 Day 9 Day 6 $20₁$ 10 **Ifnb** $\overline{116}$ Tnfa II12 .
Ifna E **NF** Day 3 Day 6 Day 9 F 1.00 WT Pe lit \sim 0.75 WT 50.50 0.25 0.00 Peli1-Day 3 Day 6 Day 9 **NF** G н CD4⁺ 20000-20000 $CD8⁺$ WТ 10000 10000 Ξ $\overline{\epsilon}$ P eli1^{-/-} ğq þg/ 200 $100 100₁$ 50 $\overline{0}$ Ω IFN- γ $IL-6$ $IL-10$ $IL-17$ $IL-10$ $IL-17$ IFN-y $IL-6$

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Supplemental Figure 2. *Peli1* 786 *−/−* **mice exhibited impaired cytokine production but** 787 **modestly enhanced adaptive immune responses in the periphery. A***.* RNA levels of *Peli1* in 788 the blood of WT mice at day 3 were determined by qPCR assay. Data represent the means \pm SEM of 6 to 7 samples from 2 independent experiments. $#P < 0.05$ compared to non-infected

790 (NF) group (Unpaired t test). **B.** Blood cytokines levels were determined by gPCR assay. Data 790 (NF) group (Unpaired t test). **B.** Blood cytokines levels were determined by qPCR assay. Data 791 are presented as the fold increase compared to NF group and represent the means \pm SEM of 6 792 to 11 samples from 3 independent experiments. **C- D**. The development of specific IgM (C) or 793 IgG (D) Abs to WNV was determined using purified rWNV-E protein antigen. Data represent the 794 means \pm SEM of 4 to 7 samples from 2 independent experiments. **E-F**: H&E staining. 794 means ± SEM of 4 to 7 samples from 2 independent experiments. **E-F***:* H&E staining. 795 Representative images (20X) shown are spleens from NF and WNV-infected mice. Two 796 photographs of different regions were taken from each spleen section at lowest magnification
797 (2.5X objective) to cover maximal area for comparison. The white pulp area was measured. $(2.5X$ objective) to cover maximal area for comparison. The white pulp area was measured, 798 expressed in pixels, using the ImageJ program (*https://imagej.nih.gov/ij/*). Scale bar: 250µm.
799 Data are presented as the ratios of white pulp-to-total area (W/T) of each group and represer 799 Data are presented as the ratios of white pulp-to-total area (W/T) of each group and represent 800 the means \pm SEM, n= 2 to 3, ** $P < 0.01$ or $\pm P < 0.05$ compared to NF group (Unpaired t test). the means \pm SEM, n= 2 to 3. ** $P < 0.01$ or $P < 0.05$ compared to NF group (Unpaired t test). 801 **G- H**. Splenocytes were harvested at day 7 pi and were cultured ex vivo with WNV peptides for 802 3 days, and cytokine production in culture supernatant was measured. Data represent the 803 means ± SEM of 4 to 7 samples from 2 independent experiments. For **B-D & G-H** panels, ** P < 804 0.01 or *P < 0.05 compared to WT group (Unpaired t test).

Supplemental Figure 3. WNV infection and PRR agonist treatment in *Peli1 −/−* **DCs and**

macrophages. A-C. WT and *Peli1^{-/-}* DCs were infected with WNV. Viral load was measured at 809 indicated time points by qPCR (A) and FFA (B). C. Cytokine RNA levels were measured at day indicated time points by qPCR (A) and FFA (B). C. Cytokine RNA levels were measured at day 3. Data represent the means ± SEM of 3 to 6 samples from 2 independent experiments. **D- G**. **WT** and *Peli1^{-/-}* macrophages were treated with TLR3, TLR7, and RLR agonists for 24 h. *Ifna*, *Ifnb*, *Il12* RNA and IL-6 protein levels were measured by qPCR or Bioplex. Data are presented 813 as fold increase compared to NF or mock-treated cells. Data represent the means \pm SEM of 3 to 4 samples are representative of 2 independent experiments. **A-G**, ** P < 0.01 or *P < 0.05 compared to WT group (Unpaired t test). **H.** WT macrophages were infected at MOI of 0.02 with WNV passaged in WT (WTP) or *Peli1*−/− (*Peli1*−/− P) macrophages. At day 4 pi, IL-6 production 817 was determined by Bioplex. Data represent the means \pm SEM of 4 samples from 2 independent

experiments. ** P < 0.01 compared to WTP group (Unpaired t test).

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825 825 **Supplemental Figure 4**. **Type I IFN induction and leukocyte infiltration in the CNS. A-B***.* 826 RNA levels of type I IFNs in the brain at indicated time points were determined by qPCR assay.
827 Data are presented as fold increase compared to NF group (means \pm SEM) and represent 4 to 827 Data are presented as fold increase compared to NF group *(means ± SEM)* and represent 4 to 828 5 samples pooled from 2 independent experiments. **C**. Brain leukocyte infiltration following WNV 828 5 samples pooled from 2 independent experiments. **C***.* Brain leukocyte infiltration following WNV 829 385-99 infection. The percentage (means \pm SEM) of brain leukocytes collected from 9 to 10
830 mice of 2 independent experiments, including naïve microglia cells (NMG), activated microg 830 mice of 2 independent experiments, including naïve microglia cells (NMG), activated microglial
831 cells (AMG), macrophages (M), CD4⁺ and CD8⁺ T cells on day 9 pi was analyzed by flow 831 cells (AMG), macrophages (M), CD4⁺ and CD8⁺ T cells on day 9 pi was analyzed by flow eyer optionetry. ** $P < 0.01$ or $*P < 0.05$ compared to WT group (Unpaired t test). cytometry. ** $P < 0.01$ or $*P < 0.05$ compared to WT group (Unpaired t test).

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834 **Supplemental Figure 5**. **WNV infection in microglia and neurons. A- C**. BV2 cells were 835 infected with WNV 385-99. **A.** Viral load was measured at indicated time points by qPCR. Data 836 represent the means \pm SEM of 7 to 10 samples pooled from 2 independent experiments. **B.** represent the means ± SEM of 7 to 10 samples pooled from 2 independent experiments. **B.** *Peli1*, *Peli2* and *Peli3* RNA levels in BV2 cells were measured by qPCR at day 4 pi. Data represent the means ± SEM of 12 to 13 samples pooled from 3 independent experiments. **A-B**, ** $P < 0.01$ or ** $P < 0.05$ compared to non-infected (NF) group (Unpaired t test). **C.** RNA levels of 840 cytokines were measured at indicated time points. Data are presented as means \pm SEM and are representative of 2 independent experiments (n = 2 to 3). **D- E.** BV2 cells were treated with control and Peli1 siRNA (Peli1KD) and infected with WNV 385-99 at 24 h. Data are presented 843 as means \pm SEM and are representative of 2 independent experiments ($n = 5$). ** $P < 0.01$ compared to control group (Unpaired t test). **D.** Viral load was measured at day 4 by qPCR. **E.** *Il6, Ccl2, Ccl7, and Cxl10* RNA levels were measured at day 4. **F.** *Ifnb* RNA levels in mouse neurons at day 4 pi. Data are presented as means ± SEM, n =6 per group. **G-H**. Western blot assay for p38MAPK and NF-κB activation in NF and WNV-infected neurons at day 3 pi. **G.** One representative of three samples per group was shown. **H.** The densitometric analysis of western blot data. Data presented as the ratio of pP65 or pP38 to β-actin compared to NF control. For **F& H** panels, ** *P* < 0.01 or **P* < 0.05 compared to WT group (Unpaired t test).

853 **Supplemental Figure 6**. **WNV infection in human cells**. **A.** RNA levels of *Peli1* in non-854 infected (NF) and WNV-infected hNSCs-derived neuron (SNeuron)s, THP1-derived
855 macrophages (THPM) and peripheral blood mononuclear cell (PBMC)s were detern 855 macrophages (THPM) and peripheral blood mononuclear cell (PBMC)s were determined by 856 gPCR assay. Data represent the means \pm SEM of 4 to 6 samples pooled from 2 independen 856 qPCR assay. Data represent the means \pm SEM of 4 to 6 samples pooled from 2 independent 857 experiments. ${}^{#}P$ < 0.05 compared to non-infected (NF) group (Unpaired t test). **B- D**. S-neuro experiments. ${}^{#}P$ < 0.05 compared to non-infected (NF) group (Unpaired t test). **B- D**. S-neurons 858 were treated with control and Peli1 siRNA knockdown (Peli1KD), infected with WNV 385-99 at 859 48 h and harvested at day 4 pi. Viral load (**B**) and *Ifnb* (**D**) levels were measured by qPCR. **C.** 860 *Peli1* expression on control and Peli1 siRNA knockdown cells at 24 h post-treatment. Data 861 represent the means \pm SEM and are representative from 2 independent experiments, n =4 to 5.
862 $*$ $*$ P < 0.01 compared to control group (Unpaired t test). ** P < 0.01 compared to control group (Unpaired t test). 863

Supplemental Figure 7. **Smaducin-6 treatment in macrophages during WNV infection.**

BM-macrophages were infected at MOI of 0.02 with WNV 385-99 and treated with Smaducin-6

or control peptides at 1 h pi. **A.** Viral load was measured at day 4 pi by qPCR. **B-C***.* Cytokine

871 levels were measured at day 4 by qPCR or Bioplex. Data are presented as fold increase

872 compared to mock- infected or pg/ ml and represent the means \pm SEM of 8 samples pooled from 2 independent experiments. $*P < 0.05$ compared to control group (Unpaired t test). from 2 independent experiments. * $P < 0.05$ compared to control group (Unpaired t test).

Table S1: Serum cytokine levels at day 3 post infection

Supplemental Methods:

 WNV infection in human cells: HMC3 (human microglial cell line), THP-1 cells and PBMCs from healthy donors were purchased from ATCC and Astarte Biologics, Inc respectively. THP-1 cells were treated with phorbol 12-myristate 13-acetate (PMA, Sigma) to 892 differentiate into macrophage cells as described previously [\(57\)](#page-45-0) before infection with WNV at a MOI of 1. HMC3 cells and PBMC were infected with WNV at a MOI of 5 or 10 respectively. Human fetal cortical neural stem cells (hNSCs) were propagated as neurospheres in DMEM/F12 basic media supplemented with 20 ng/mL epidermal growth factor (EGF), 20 ng/mL fibroblast growth factor 2 (FGF2), 10 ng/mL leukemia inhibitor factor (LIF) and N2; and passaged every 10 days [\(58\)](#page-45-1). For neuron differentiation, dissociated hNSCs were plated into 898 T75 flasks. After a 3-day incubation, 1.2×10^6 or 5×10^5 hNSCs (small spheres) were seeded into T25 flasks or 6-well plates pre-coated with 0.01% poly-D-lysine and 1 μg/cm² laminin (Invitrogen), respectively. Cells were incubated for 4 days with the priming media containing 901 EGF (20 ng/mL), LIF (10 ng/mL) and laminin (1 µg/mL), followed by 9-day incubation with a differentiation medium consisting of N2 plus glutathione (1 μg/mL) (Sigma), biotin (0.1 μg/mL) (Sigma), superoxide dismutase (2.5μg/mL) (Worthington), DL-α-tocopherol (1 μg/mL, Sigma), DL--α-tocopherol acetate (1μg/mL, Sigma) and catalase (Sigma). Cells were subsequently infected with WNV at a MOI of 1.

In vitro T cell activation: Splenocytes (0.3 x 10⁶) were stimulated with WNV-specific 907 NS3 and E peptides respectively [\(59\)](#page-45-2) for CD4⁺ T cells, or WNV-specific NS4B and E peptides [\(60,](#page-45-3) [61\)](#page-45-4) for CD8⁺ T cells for 72 h at 37°C. Culture supernatant was collected for cytokine assay. **ELISA***:* Microtiter plates were coated with recombinant WNV-E protein [\(62\)](#page-45-5) overnight at 910 4°C. Sera were diluted 1/30 in PBS with 2% BSA, and incubated for 1 h at room temperature. Alkaline phosphatase-conjugated goat anti-mouse IgG (Sigma-Aldrich, A3562) or IgM (Sigma-Aldrich, A9688) at a dilution of 1/1000 in 1XPBS with 0.05% Tween (Sigma-Aldrich) was then

 added for 1 h. Color was developed with *p*-nitrophenyl phosphate (Sigma-Aldrich) and intensity read at an absorbance of 405 nm.

 siRNA knockdown for Peli1: Mouse cells were transfected with 187.5nM of pooled Peli1 specific siRNA (CUCAUGACAGCAACACUGA, CAACCAUGGGUAUAUCUAA, CUUUACAGCUCGGAUUUAU), or 187.5nM control siRNA (MISSION Universal Negative Control #1) (all from Sigma-Aldrich) by using Superfect (Qiagen) per the manufacturer's 919 instructions. Human cells were transfected with 37.5 pM control siRNA or Peli1 siRNA (Santa Cruz Biotechnology) per the manufacturer's instructions. Transfected cells were grown in RPMI medium containing 10% FBS. QPCR analysis of Peli1mRNA was used to confirm the effects of the siRNA knockdown. At 48 h post-infection, cells were infected with WNV with MOI of 0.1 (mouse cells) or MOI of 1 or 5 (human cells) respectively. Supernatants and cells were harvested at 24 h and 96 h pi to measure viral load and cytokine production. **Cytokine Bioplex:** Analysis of cytokine production by using a Bio-Plex Pro Mouse Cytokine Assay (Bio-Rad). .

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