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ELUCIDATING THE ROLE OF INTERLEUKIN-17A IN

WEST NILE VIRUS INFECTION

by

Dhiraj Acharya

A Dissertation Submitted to the Graduate School and the Department of Biological Sciences at The University of Southern Mississippi in Partial Fulfillment of the Requirements for the Degree of Doctor of Philosophy

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ABSTRACT

ELUCIDATING THE ROLE OF INTERLEUKIN-17A IN

WEST NILE VIRUS INFECTION

by Dhiraj Acharya

May 2017

West Nile virus (WNV) is a neurotropic flavivirus of significant public health importance for which no therapeutics and vaccine are currently available. Interleukin-17A (IL-17A) is an inflammatory cytokine that regulates diverse immune functions, while its role is unclear in host's immune response to WNV. Furthermore, CD8⁺ T cells are crucial components of immunity and play a vital role in recovery from WNV infection. Here, we report a previously unrecognized function of IL-17A in regulating CD8⁺ T cell cytotoxicity. We show that WNV induces the expression of IL-17A in both mouse splenocytes and human peripheral blood mononuclear cells cultured in vitro, and in plasma of WNVinfected mice and humans. In a mouse model of WNV infection, we demonstrate that IL-17A deficient mice ($II17a^{-/-}$) are more susceptible to WNV and develop a higher viral burden compared to wild-type (WT) mice. Interestingly, the CD8⁺ T cells isolated from WNV-infected *II17a^{-/-}* mice are less cytotoxic and express lower levels of cytotoxic mediator genes, which can be restored by supplying recombinant IL-17A in vitro and in vivo. Moreover, treatment of WNV-infected mice with recombinant IL-17A, as late as day 6 post-infection, significantly reduces viral burden and increases survival, suggesting a therapeutic potential of IL-17A. In conclusion, we demonstrate a novel function of IL-17A in promoting

CD8⁺T cell cytotoxicity against WNV infection, which may have broad implications in other microbial infections and cancers.

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DEDICATION

I dedicate this dissertation to my father Durga Datta Sharma Acharya and my mother Sharada Sharma Acharya, who are endless source of inspiration and encouragement throughout my life. I like to extend my gratitude to my beloved wife Ranjana Baral, and my sisters Rasmi and Roshani for their love and care. Finally, I like to thank all my friends for their motivation and encouragement.

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LIST OF ABBREVIATIONS

- ACT-1 NF-κB activator 1
- ANOVA Analysis of variance
- ATCC American type culture collection
- *BBB* Blood brain barrier
- BSL3 Biosafety level 3
- *cDNA* Complementary DNA
- CHIKV Chikungnya virus
- CNS Central nervous system
- *d.p.i.* Days post infection
- *DMEM* Dulbecco's modified Eagle medium
- DNA Deoxyribonucleic acid
- *ELISA* Enzyme linked immunosorbent assay
- *ERK* Extracellular signal-regulated kinase
- FBS Fetal bovine serum
- Fig Figure
- *FCB* Flow cytometry buffer
- *hPBMC* Human peripheral blood mononuclear cells
- *i.p.* Intraperitoneal
- IFN Interferon
- *IL* Interleukin
- *IL-17A* Interleukin-17A
- *IL-17R* Interleukin-17 receptor

- IRB Institutional review board
- *IACUC* Institutional animal care and use committee
- *LCMV* Lymphocytic choriomeningitis virus
- MOI Multiplicity of infection
- *mRNA* Messenger RNA
- NK Natural killer
- *NKT* Natural killer T
- NS Nonstructural
- PBS Phosphate buffer saline
- PFA Paraformaldehyde
- *PFU* Plaque-forming unit
- *qPCR* Quantitative polymerase chain reaction
- *RANTES* <u>R</u>egulated on <u>activation</u>, <u>n</u>ormal <u>T</u> cell <u>expressed</u> and <u>secreted</u>
- *RFC* Relative fold change
- rIL-17A Recombinant IL-17A
- *RNA* Ribonucleic acid
- SEFIR Similar expression to fibroblast growth factor and <u>IL-17R</u>
- SEM Standard error of mean
- TLR Toll-like receptor
- TNF Tumor necrosis factor
- USM The University of Southern Mississippi
- WNV West Nile virus

- WNV-E WNV envelop
- WT Wild-type

CHAPTER I - INTRODUCTION

1.1 West Nile Virus

West Nile virus (WNV), a neurotropic single-stranded RNA virus belonging to *flaviviridae* family, is generally transmitted to human by an infected mosquito bite, primarily of *Culex* species (Colpitts et al., 2012; Kuno and Chang, 2005). However, WNV can also be transmitted through other less frequent transmission routes, including blood transfusion (Pealer et al., 2003; Stramer et al., 2005), organ transplantation (Centers for Disease and Prevention, 2009), breastfeeding (Blazquez and Saiz, 2010), and congenital infections (Alpert et al., 2003). WNV infection in human can cause fever and may result in injury and death of neurons, the latter can lead to various neurological manifestations, such as encephalitis, flaccid paralysis, meningitis, chronic neurologic sequelae and possibly death, particularly in the elderly and immunocompromised individuals (Colpitts et al., 2012; Hayes et al., 2005; Samuel and Diamond, 2006).

Structurally, WNV is an enveloped virus with 50 nm diameter, which comprises an icosahedral nucleocapsid surrounded by a host-derived lipid envelope (Mukhopadhyay et al., 2003). The WNV genome contains a capped, single-stranded, and plus-sensed RNA genome of approximately 11 Kb in size. The WNV genome translates to a polyprotein precursor, which undergoes posttranslational processing by viral and cellular proteases to generate three structural (capsid [C], premembrane [PrM], and envelope [E]), and seven nonstructural proteins (NS1, NS2A, NS2B, NS3, NS4A, NS4B, and NS5). Among these proteins, the structural proteins form the structure of virion, whereas the

non-structural proteins are essential in genome replication, virion assembly, and viral pathogenesis (Chung et al., 2006; Liu et al., 2006; Mukherjee et al., 2011a).

1.1.1 WNV Is a Major Public Health Problem

WNV was first discovered in Uganda about 80 year ago and primarily maintained in a mosquito-bird-mosquito transmission cycle (Colpitts et al., 2012). Until 1990's it caused several sporadic outbreaks of minor public health importance in Africa, Asia, Europe and the Middle East with a limited capacity to cause neuroinvasive diseases (Draganescu, 1979; Kramer et al., 2008). The first report of neuroinvasive cases of WNV in Algeria in 1994 (Le Guenno et al., 1996), and a huge report of neurological impairments during subsequent WNV outbreaks in South-East of Romania (Cernescu et al., 1997), and other parts of southern Europe (Hubalek and Halouzka, 1999) brought this virus into a real public health concern. In northern hemisphere, WNV was first identified in New York in 1999 (Hayes, 2001; Mostashari et al., 2001), which subsequently spread throughout the USA within a few years (Benjelloun et al., 2015).

Currently, WNV is causing endemic diseases throughout the North America and has a wide geographical distribution in all continents of globe except Antarctica (Benjelloun et al., 2015). In the US alone, there have been over 40,000 cases of WNV reported between 1999 and 2014, out of which about 45% were classified as neuroinvasive cases (CDC, 2015). However, the actual burden of WNV is likely much higher than previously thought because only a proportion (about 20%) of infected individual develop a clinical disease and a majority of infected individuals remain asymptomatic (Mostashari et al., 2001). It has been

recently estimated that over 3 million individuals have been infected with WNV in the USA alone, out of which about 780,000 had a symptomatic illness (Petersen et al., 2013). Considering the worldwide circulation of WNV and supportive evidence of its capacity to change pathogenicity and transmission potential (Davis et al., 2005; Ebel et al., 2004; Moudy et al., 2007; Prow et al., 2014; van den Hurk et al., 2014), there is an urgent need to develop a safe and effective antiviral drug or vaccine against WNV infection (Martina et al., 2010). However, no licensed antiviral drug or vaccine is currently available for WNV infection in humans.

1.1.2 Pathogenesis of WNV Infection Is Not Well Understood

After a bite of WNV-infected mosquito inoculates infectious viruses into human skin, WNV first infects skin-resident cells such as dendritic cells (Langerhans cells) and keratinocytes. Infected dendritic cells can carry the viruses to draining lymph nodes, after which WNV circulates through lymphatic systems and enters the blood stream (Lim et al., 2011; Ye et al., 2011). WNV can infect various cells including macrophages, neutrophils, and other leukocytes leading to development of a transient viremia. Subsequently, WNV can disseminate to peripheral organs, such as liver and spleen, and then to the brain and spinal cord. In most of the immune-competent individuals, WNV symptoms range from unapparent infection to mild febrile illness (Colpitts et al., 2012). In a few symptomatic cases, particularly in the elderly and immunocompromised individuals, WNV can cause injury and apoptosis of neuron, which can potentially cause various neurological impairments and death (Colpitts et al., 2012). Viral

entry into the brain is considered a hallmark of WNV pathogenesis and disease severity. Although the mechanism of neurotropism and how WNV enters the central nervous system (CNS) is not clearly understood, hematogenous entry via disrupted blood-brain barrier (BBB) and transneural entry via retrograde axonal transport have been suggested to play a role (Suen et al., 2014; Suthar et al., 2013).

1.1.3 Host's Immunity Is Crucial in WNV Recovery

WNV can infects various type of immune cells and strongly activates host immune responses, which play important roles in controlling viremia, reducing viral dissemination to the CNS, and recovery from the disease (Samuel and Diamond, 2006). Clinical reports of WNV-infected human cases along with research using mouse models or *in vitro* approaches have identified critical roles of several components of immune system in WNV infections (Arjona et al., 2011; Suthar et al., 2013). In brief, type-I interferon (IFNs) (Lazear et al., 2011; Samuel and Diamond, 2005), the complement system (Mehlhop and Diamond, 2006), and humoral immunity (Diamond et al., 2003a; Diamond et al., 2003b) limit viremia and control WNV dissemination to CNS. Similarly, components of cellmediated immunity, including CD4⁺ (Sitati and Diamond, 2006) and CD8⁺ (Shrestha and Diamond, 2004) T cells, have been shown to clear WNV from the CNS, limit viral persistence, and facilitate recovery. In contrast, the roles of neutrophils, NK cells, and $\gamma\delta$ -T cells are still unclear (Bai et al., 2010; Shrestha et al., 2006a; Welte et al., 2008). Cytokine signaling of interleukin (IL)-23 (Town et al., 2009), interferon- γ (IFN- γ)(Shrestha et al., 2006b), and IL-1 β (Ramos et al.,

2012) have been shown to protect against WNV infection, while cytokine such as IL-10 (Bai et al., 2009) and IL-22 (Wang et al., 2012) have been shown to favor WNV pathogenicity. The role of tumor necrosis factor- α (TNF- α) still remains elusive (Shrestha et al., 2008b; Wang et al., 2004a). However, the role of several other components of immune system yet remains unknown and the mechanism of WNV pathogenesis, including its tropism to neurons and CNS invasion, remains unclear. In addition, we have limited knowledge on viral or host factors that contribute to imbalance between viral pathology and host to WNV infection.

1.2 Interleukin-17A and Its Signaling

IL-17A, a major cytokine of IL-17 family (IL-17A, B, C, D, E, and F), was discovered in 1993 as the cytotoxic T-lymphocyte antigen-8 (Rouvier et al., 1993). CD4⁺ Th17 (Th-17) cells are one of the most common sources of IL-17A. Besides Th-17 cells, a wide variety of immune cells, including CD8⁺ T cells, γδ-T cells, and natural killer T (NKT) cells can also produce IL-17A under various pathological conditions (Gu et al., 2013). IL-17A signals through IL-17 receptor (IL-17R) complex composed of IL-17RA and IL-17RC subunits (Ho and Gaffen, 2010). Both subunit of IL-17R complex encode a SEFIR (Similar Expression to Eibroblast growth factor genes and IL-17 Receptor) domain that mediates various downstream signaling events through the adaptor protein NF-κB activator-1 (ACT1) (Gaffen, 2009; Ho and Gaffen, 2010). Specifically, the ACT1 recruits to IL-17R through interaction with SEFIR domain and activates TNF receptor-associated factor-6 (TRAF6) and TRAF3, which are essential upstream activators of the nuclear factor-κB (NF-κB) signaling pathway. In addition, ACT1

is also required for IL-17A mediated stabilization of mRNAs that encode several chemokines and cytokines. Moreover, IL-17A also signals through ACT1 independent pathway by activation of extracellular signal-regulated kinase (ERK) (Ho and Gaffen, 2010).

1.2.1 IL-17A Regulates Diverse Immune Functions

The finding that IL-23 drives expression of IL-17 in CD4⁺ T cells led to the realization of a new paradigm that many functions formerly attributed to CD4⁺ Th-1 cells were in fact mediated by a distinct, IL-17-producing T cell subset known as Th-17 cells (Aggarwal et al., 2003; Langrish et al., 2005). It is now well established that IL-1 β , IL-6, TGF- β , and IL-21 drive development, while IL-23 stabilizes Th-17 cells (Korn et al., 2007; Nurieva et al., 2007; Sutton et al., 2006; Zhou et al., 2007).

IL-17A signaling regulates diverse immune functions including the expression of various inflammatory cytokines, chemokines and antimicrobial peptide, activation and recruitment of leukocytes, and production of antibodies (Gu et al., 2013; Song and Qian, 2013) during infection and immunity. For example, IL-17A has often been described as a mediator of inflammation (Witowski et al., 2004) with prominent roles in allergic and autoimmune diseases including multiple sclerosis (McFarland and Martin, 2007; Zepp et al., 2011), rheumatoid arthritis (van den Berg and Miossec, 2009), psoriasis (Raychaudhuri, 2013), asthma (Newcomb and Peebles, 2013), and Crohn's disease (Siakavellas and Bamias, 2012). Similarly, IL-17A also plays protective roles against some bacterial and fungal infections. For instance, IL-17A may enhance neutrophil

recruitment and protect against bacterial and fungal pathogens such as *Klebsiella pneumonia* and *Escherichia coli* (Happel et al., 2005; Shibata et al., 2007; Ye et al., 2001), *Listeria monocytogenes* (Hamada et al., 2008), *Mycobacterium tuberculosis* (Witowski et al., 2004), *Francisella tularensis* (Lin et al., 2009), *Chlamydia muridarum* (Zhang et al., 2009), *Candida albicans* (Conti et al., 2009; Huang et al., 2004) and *Pneumocystis carinii* (Rudner et al., 2007). Conversely, IL-17A signaling has been suggested to facilitate toxoplasmosis (Guiton et al., 2010) and certain fungal infections (Zelante et al., 2007).

1.2.2 Role of IL-17A in Viral Infections Is Unclear

The IL-17A has been originally identified as a homolog of virally encoded ORF13 gene of *herpesvirus saimiri* (Yao et al., 1995). This discovery generated interest to study the role of this cytokine in viral infection. It has been demonstrated that genetically constructed vaccinia virus (VV) expressing IL-17A (VV^{IL-17A}) caused more severe disease in mice (Patera et al., 2002), but VV^{IL-17A} was also reported less virulent and IL-17 deficient (*II17a^{-/-}*) mice were more susceptible to VV infection (Kohyama et al., 2007). Thus, the role of IL-17A during VV infection remains inconclusive.

Interestingly, it has been reported that type I IFNs potently suppress IL-17A expression (Curtis et al., 2009; Tilg et al., 2009), which further makes the functional role of IL-17A during viral infections more complex to understand. In most of the viral infections, type I IFN-mediated antiviral response primarily occurs during early phase of infection (Lazear et al., 2011; Pinto et al., 2011). It can be speculated that the IL-17A functions may be more dominant in the later

phase of infection when type I IFN expression levels become low. It has been shown that viral infections can induce expression of IL-17A (Mukherjee et al., 2011b; Town et al., 2009; Wang et al., 2013), which has been implicated in priming T cell responses during lymphocytic choriomeningitis virus (LCMV) hepatitis (Jie et al., 2014) and mediating immunopathogenicity of viral infections such as influenza virus (Crowe et al., 2009), respiratory syncytial virus (de Almeida Nagata et al., 2014; Mukherjee et al., 2011b), murine encephalomyelitis virus (Hou et al., 2009) and hepatitis B virus (Wang et al., 2013). However, the role of IL-17A in WNV infection in not clearly understood.

1.3 Possible Implication of IL-17A in WNV Infection

It has been previously reported that WNV infection induces Toll-like receptor-7 (TLR7) dependent production of IL-23 in mice (Town et al., 2009). IL-23 is known as a prime regulator for stabilization and maintenance of CD4⁺ Th-17 cells, which are the major cell type secreting IL-17A (Aggarwal et al., 2003; McGeachy et al., 2009; Stritesky et al., 2008). Thus, it is likely that WNV infection induces IL-17A expression, which may play a role in WNV immunity. The following literature provides strong evidence for possible involvement of IL-17A signaling during WNV infection.

1.3.1 IL-17A Is Crucial in the CNS Inflammation

WNV primarily targets neurons and causes inflammation in the CNS, which play important role in its pathogenesis. Astrocytes and microglia upregulate the expression of functional IL-17A receptor and respond to this cytokine during brain inflammatory conditions (Das Sarma et al., 2009). In addition, IL-17A is also described as a mediator of CNS inflammation (McFarland and Martin, 2007; Zepp et al., 2011) and suggested as a blood-brain-barrier permeability factor (Kebir et al., 2007). Therefore, the WNV pathogenesis and host's immune response in the CNS are likely impacted by IL-17A.

1.3.2 IL-17A May Control Host's Immunity to WNV Infection

Considering the role of IL-17A in regulating diverse immune function, it can be speculated that this cytokine may play an important role in WNV immunity. For instance, IL-17A has been shown to regulate the expression of other cytokines including IL-1 β , IFN- γ and TNF- α (Maione et al., 2009; Song and Qian, 2013), infiltration of leukocytes, and production of antibodies (Yao et al., 1995; Yuan et al., 2010). As these cytokines are previously described to modulate immunity to WNV (Ramos et al., 2012; Shrestha et al., 2006b; Shrestha et al., 2008b; Wang et al., 2004a), one can expect possible involvement of IL-17A in WNV immunity. Moreover, brain-infiltrating leukocytes are crucial in clearance of WNV (Bai et al., 2010; Shrestha and Diamond, 2004), which can be potentially controlled by IL-17A signaling. Furthermore, the humoral immune response, which plays an important role in controlling viremia and WNV dissemination to CNS (Diamond et al., 2003a), can be potentially regulated by IL-17A (Yuan et al., 2010).

Among the CNS-infiltrating leukocytes, CD8⁺ T cells play crucial role in WNV immunity (Shrestha and Diamond, 2004, 2007; Shrestha et al., 2012; Shrestha et al., 2006a). Although the relation between IL-17A signaling and CD8⁺ T cell cytotoxicity is not yet studied, several literatures support this novel axis may play a role in WNV and other viral infections. First, both Th-17 cells and $\gamma\delta$ -T cells, the major cells that produce IL-17A, have been previously shown to promote CD8⁺ T cell cytotoxicity during intracellular infections (Hamada et al., 2009; Xu et al., 2010). Second, the Th-17 cells and IL-17 have also been shown to control CD8⁺ T cell function in autoimmune diseases (Ankathatti Munegowda et al., 2011a). Third, the CD8⁺ T cells have a crucial role in clearance of tumor cells (Benchetrit et al., 2002; Martin-Orozco and Dong, 2009). Several studies also demonstrated the expression of IL-17A in tumor cells (Benchetrit et al., 2002; Martin-Orozco and Dong, 2009). Interestingly, it has been shown that Th-17 cell and IL-17A induce cytotoxic function of CD8⁺ T cells and control tumor progression (Benchetrit et al., 2002; Martin-Orozco and Dong, 2009), however, the mechanism for this IL-17A-CD8⁺ T cells in WNV immunity, WNV is a suitable model to study IL-17A-CD8⁺ T cell axis.

CHAPTER II – SIGNIFICANCE, HYPOTHESIS, AND INNOVATION

2.1 Significance

WNV infection is a significant public health problem, for which no therapeutics or vaccine are currently available. In addition, IL-17A and CD8⁺ T cells regulate diverse immune functions during microbial infections, malignancies, and autoimmune diseases. In this study, we studied the function of IL-17A in WNV immunity and identified a potential therapeutic role of IL-17A in facilitating WNV clearance by promoting CD8⁺ T cells cytotoxicity. Further understanding of the interplay between CD8⁺ T cells and IL-17A axis may lead to identification of novel therapeutic strategies. Moreover, this novel function of IL-17A may have broad implications in deciphering the immunopathology of other viral infections and malignancies, where CD8⁺ T cells functions are crucial.

2.2 Hypothesis

Despite intensive research over the past 15 years, WNV pathogenesis and host's immune response still remains unclear. IL-17A, a major cytokine in the IL-17 family regulates diverse immune functions including the expression of various inflammatory cytokines and chemokines, activation and recruitment of leukocytes, and production of antibodies (Gu et al., 2013; Song and Qian, 2013). Moreover, IL-17A has often been described as a mediator of inflammation (Witowski et al., 2004) with an important role in allergic and autoimmune diseases (McFarland and Martin, 2007; Newcomb and Peebles, 2013; Raychaudhuri, 2013; van den Berg and Miossec, 2009; Zepp et al., 2011), malignancies (Martin-Orozco and Dong, 2009; Murugaiyan and Saha, 2009), and

infections (Conti et al., 2009; Hamada et al., 2008; Happel et al., 2005; Huang et al., 2004; Lin et al., 2009; Rudner et al., 2007; Shibata et al., 2007; Ye et al., 2001; Zhang et al., 2009). However, the role of IL-17A signaling is not clear during viral infections and has been previously studied in WNV infection. As we described in the introduction (section 1.7), there are several evidences suggesting a possible role of IL-17A in controlling host's immune response to WNV. Therefore, *we hypothesized that IL-17A may play a crucial role in WNV immunity*. In this study, we tested this hypothesis under following four specific aims.

Specific Aim 1: Study the expression of IL-17A during WNV infection in vitro and in vivo. It has been previously reported that TLR7 mediates IL-23dependent protective immune responses against WNV infection in mice (Town et al., 2009). IL-23 is known as a prime regulator for stabilization and maintenance of CD4⁺ Th-17 cells, which are the major cell type secreting IL-17A (Aggarwal et al., 2003; McGeachy et al., 2009; Stritesky et al., 2008). Since IL-17A is also described as a mediator of CNS inflammation (McFarland and Martin, 2007; Zepp et al., 2011), and a factor contributing to blood-brain-barrier permeability (Kebir et al., 2007), we asked if WNV induces IL-17A production and whether it involves IL-23 signaling.

Specific Aim 2: Study the role IL-17A in WNV immunity in a mouse model. Based on the literature, it is likely that IL-17A may play either protective or pathogenic role during WNV infection. To dissect this question, we used a mouse

model of WNV infection and compared survival and viral burden of WNV infected IL-17A deficient (*II17a^{-/-}*) and WT control mice.

Specific Aim 3: Study the immune functions of IL-17A during WNV infection. IL-17A has been shown to recruit leukocytes (Happel et al., 2005; Ye et al., 2001), control cytokine and chemokine expression (Maione et al., 2009; Song and Qian, 2013), and regulate humoral immune response (Grund et al., 2012; Tarlinton, 2008; Yuan et al., 2010). Under this specific aim, we asked how various immune functions were controlled by IL-17A during WNV infection.

Specific Aim 4: Study the mechanism of IL-17A mediated regulation of WNV immunity. From Aim 3, we identified a novel function of IL-17A in regulation of CD8⁺ T cell cytotoxicity. Under Aim 4, we studied the mechanism of IL-17A mediated control of CD8⁺ T cells cytotoxicity and its potential therapeutic role against WNV infection.

2.3 Innovation

Cytotoxic CD8⁺ T cells are important component of immunity, particularly in clearance of intracellular infections and tumors. One of the effector mechanisms used by these cells is cytotoxic killing of target cells. However, regulators of CD8⁺ T cell cytotoxicity, in particular, the expression of cytotoxicity mediators in these cells is poorly understood. In addition, functions of several molecules expressed in CD8⁺ T cells remain uncharacterized. One such molecule is the IL-17A receptor (IL-17R), which is expressed on the surface of CD8⁺ cells (Lindemann et al., 2008; Yao et al., 1995). In additions, subsets of CD8⁺ cells can produce IL-17A (Hamada et al., 2009). Considering the role of IL- 17A in regulation of diverse immune functions during microbial infections, malignancies, and autoimmune diseases, it is likely that IL-17A signaling may play a role in biology of CD8⁺ T cells. CD8⁺ T cells are particularly important in controlling viral infections and have been shown to play a crucial role in clearance of WNV infection (Shrestha and Diamond, 2004). In this study, we studied the role of IL-17A in a mouse model of WNV infections and discovered a novel, previously unrecognized, function of IL-17A in promoting CD8⁺ T cells cytotoxicity.

CHAPTER III – EXPERIMENTAL APPROACHES

3.1 Ethics Statement and Biosafety

Written informed consent was obtained from all human volunteers and human WNV cases prior to their enrollment in this study. The protocol for human subject has been reviewed and approved by the University of Southern Mississippi (USM) Institutional Review Board (IRB, protocol # CH-R11120601, Appendix A). All animal experimental procedures were reviewed and approved by the Institutional Animal Care and Use Committee at USM (protocol #12041201, Appendix A). All the *in vitro* experiments and animal studies involving infectious WNV were performed by the certified personnel in biosafety level 3 (BSL3) or animal biosafety level 3 (ABSL3) laboratories following standard biosafety protocols approved by USM Institutional Biosafety Committee.

3.2 Virus Stock and Animal Studies

Low-passaged WNV isolate CT2741 (Anderson et al., 1999) was provided by Dr. John F. Anderson at the Connecticut Agricultural Experiment Station. Lowpassaged CHIKV LR OPY1 2006 strain was provided by Dr. Robert B. Tesh at the University of Texas Medical Branch. WNV and CHIKV stocks used in this study were prepared by propagating the viruses in Vero cells through a single passage and titrated in Vero cells by a plaque-forming assay as previously described (Bai et al., 2005). Vero cells (ATCC CCL-81) were cultured in a 37°C incubator with 5% CO₂ in Dulbecco's modified Eagle medium (DMEM, Life Technologies) supplemented with 10% fetal bovine serum (FBS). Interleukin-17A deficient (*II17a-*) mouse breeding pairs (C57BL/6J background) were provided

by Dr. Richard A. Flavell at the Yale University School of Medicine and wild-type (WT) control mice (C57BL/6J) were purchased from The Jackson Laboratory (Bar Harbor, ME). IL-23 deficient (*II23p19^{-/-}*) breeding pairs on a mixed C57BL/6 × 129 background were obtained from the Mutant Mouse Regional Resource Center (MMRRC). Mice were housed under standard conditions in the animal facility at USM. Gender-matched and 7 to 9 weeks old *II17a^{-/-}* and WT control mice were infected with 1,000 plaque forming units (PFUs) of WNV by i.p. injection in 100 µl of phosphate buffer saline (PBS) containing 5% gelatin (Town et al., 2009). For footpad inoculation, 100 PFU of WNV in 50 µl PBS containing 1% FBS was injected into the mouse footpad after isofluorane anesthesia (Bai et al., 2009). Infected animals were observed twice daily for up to 21 days for morbidity and mortality.

3.3 Cell Culture and In Vitro Infection

Human peripheral blood mononuclear cells (hPBMCs) were isolated from blood of healthy human volunteers using Ficoll-paqueTM PLUS (GE healthcare). To isolate murine splenocytes, healthy C57BL/6J mice (7-week old) were euthanized and spleens were collected to make single cell suspension. After red blood cell lysis, both mouse splenocytes and hPBMCs were purified and cultured in the Dulbecco's modified Eagle medium (DMEM, Life Technologies) supplemented with 10% FBS, 2 mM L-glutamine, and 1% non-essential amino acids. Cells were infected with WNV (MOI = 0.1, 1 or 5) and collected in TRIreagent (Molecular Research Center) at 24 h and 48 h for total RNA extraction.

3.4 Quantitative Polymerase Chain Reaction (qPCR)

Total RNA was extracted from cells or animal tissues (i.e. blood, spleen, and brain) using the TRIreagent or RNeasy kit with on-column DNA digestion (Qiagen). First-stranded complementary DNA (cDNA) was synthesized using the iSCRIPTTM cDNA synthesis kit (Bio-Rad). WNV-envelope (*WNVE*) RNA copy numbers were quantified using probe-based qPCR and normalized to cellular β -*actin* gene, as previously described (Bai et al., 2005). qPCR assays for cytokine, chemokine, and other immunological marker genes were performed using SYBR Green supermix (Bio-Rad) and data are presented either as the relative fold change (RFC) by $\Delta\Delta$ CT method using β -*actin* as a housekeeping gene or as a copy number ratio of target gene to cellular β -*actin*. Primer sequences for mice (Bai et al., 2005) and human (Kozaci et al., 2007) β -*actin* are previously described. Primers sequences for human *II17a* (F, 5'-

TGTGATCTGGGAGGCAAAGT-3'; R, 5'-GATCTCTTGCTGGATGGGGA-3'), and mouse *II17a* (F, 5'-TCTCCACCGCAATGAAGACC-3', R, 5'-

TTTCCCTCCGCATTGACACA-3'); perforin-1 (F, 5'-

TGTTCCTCCTGGGCCTTTTC-3'; R, 5'-CCATACACCTGGCACGAACT-3'),

granzyme-A (F, 5'-CACGTGAGGGGGGATCTACAAC-3'; R, 5'-

TCTCCCCCATCCTGCTACTC-3'), granzyme-B (F, 5'-

TGCTACTGCTGACCTTGTCTC-3'; R, 5'-CCATGTAGGGTCGAGAGTGG-3'),

fasL (F, 5'-GAACTGGCAGAACTCCGTGA-3'; R, 5'-

TGAGTGGGGGTTCCCTGTTA-3'), *lfn-*α (F, 5'-TTCCCCTGACCCAGGAAGAT-3'; R, 5'-CTTCTGCTCTGACCACCTCC-3'), *lfn-*β (F, 5'-

TGTCCTCAACTGCTCTCCAC-3'; R, 5'-ATCTCTGCTCGGACCACCAT-3'), *Act-1* (F, 5'-GAGGACGAGCATGGCTTACA-3'; R, 5'-TGGCATTTGGGAAGAGCACA -3') were designed using NCBI's primer designing tool and synthesized by Integrated DNA Technologies.

3.5 Enzyme Linked Immunosorbent Assay (ELISA)

IL-17A, IFN- β , IL-1 β , IFN γ , TNF α , IL-6, IL-10, IL-12p40, and anti-WNV-E IgM antibody in plasma of WNV-infected mice (1,000 PFU, i.p.) were measured using ELISA kit (R&D Systems) following manufacturer's instructions. Level of IL-17A in culture media, and sera of human WNV cases and healthy controls was measured by ELISA kit from Enzo Life Sciences.

3.6 Confocal Microscopy

Brains were collected from WNV-infected mice (1,000 PFU, i.p.) after intracardial PBS perfusion, fixed overnight in 4% PFA at 4°C, and cryoprotected in sucrose. Para-median sagittal sections (25 µm) of brain were pre-blocked for 30 min at room temperature and then probed overnight at 4°C with a combination of primary antibodies against CD11b, CD45 and WNV antigen (anti-WNV antibody provided by John F. Anderson, other antibodies purchased from BD Biosciences). After a PBS wash, sections were probed with appropriate fluorescent labeled secondary antibodies for 1 h at room temperature, counterstained with DAPI (Invitrogen), and mounted in fluorescence mounting medium (ProLong Gold). Images were acquired in independent channels using a Nikon A1R confocal microscope.

3.7 Flow Cytometry

Brains were collected from WNV-infected (1,000 PFU, i.p.) mice after intracardial PBS perfusion and processed into a single cell suspension. Brain leukocytes were isolated using a discontinuous Percoll gradient (GE Healthcare) and probed with CD45, CD4, CD8, and CD11b (BD Biosciences or eBioscience). After staining, cells were washed two times in flow cytometry buffer (FCB, PBS with 2% FBS) and fixed in 4% paraformaldehyde (PFA) for 15 minutes. For intracellular staining, cells were permeabilized and probed with antibodies against perforin, and granzyme A (eBioscience). Cells were then washed and resuspended in FCB. Data were acquired on a flow cytometer (BD LSRFortessa) and analyzed with FlowJo or FACSDiva[™] software (BD Biosciences).

3.8 CD8⁺ T cell Isolation and Cytotoxicity Assay

Spleens were collected from WNV-infected (100 PFU, i.p.) mice at 10 d.p.i.. Splenic CD8⁺ T cells were isolated by negative antibody selection with magnetic beads using the mouse CD8⁺ T Lymphocyte Enrichment Set-DM (BD Biosciences). Purity of CD8⁺ T cells was examined by flow cytometry after staining with fluorescent labeled anti-CD3 and anti-CD8 antibodies (eBioscience). Cytotoxicity of CD8⁺ T cells was measured as described previously (Shrestha and Diamond, 2004), with some modifications. Briefly, purified CD8⁺ T cells (~ 80-90% purity) were co-cultured for 4 h in 96-well plates with target cells expressing the ectodomain of *WNV-E* (MC57GL_{WNV-E}) or control cells containing the expression vector (MC57GL_{vector}) (gifted by Dr. Michael S. Diamond) with effector to target cell ratios of 50:1. CD8⁺ T cell cytotoxicity was measured using a LDH Cytotoxicity Detection Kit (Thermo Scientific). Transcripts of *perforin-1*, *granzyme A*, *granzyme B* and *Fas-ligand* (*FasL*) genes in purified CD8⁺ T and CD8⁻ cells were measured by gPCR, as described above.

3.9 Ex Vivo and In Vivo IL-17A Treatment Assays

For *ex vivo* studies, splenocytes were isolated from WNV-infected (100 PFU, i.p.) mice at 8 d.p.i. and splenic CD8⁺ T cells were purified as described above. In some experiments, CD4⁺ T cells were depleted from splenocytes using CD4 Magnetic Particles - DM (BD Biosciences). Splenocytes, purified CD8⁺ T cells, or CD4⁻ splenocytes were cultured for 24 h in the presence of mouse recombinant IL-17A (50 ng/ml, eBioscience). In some experiments, splenocytes were cultured for 24 h in the presence of mouse recombinant IL-17A (50 ng/ml) and subjected to CD8⁺ T cell purification. Expression of *perforin-1, granzyme A*, *granzyme B* and *FasL* in splenocytes, CD8⁺ T cells, and splenic CD8⁻ cells were measured by qPCR, as described above.

For *in vivo* IL-17A treatment and survival studies, WT female mice (8week old) were inoculated with WNV (100 PFU) via i.p. route. At day 6 p.i, mice were treated (2.5 µg/mouse) with i.p. administration of carrier-free mouse recombinant IL-17A (eBioscience) or PBS, and monitored daily for mortality and morbidity for up to 21 days. Randomly selected mice were euthanized at day 8 p.i. and WNV burden in brain and the expression of the cytotoxic mediators in splenic CD8⁺ and CD8⁻ T cells were measured by qPCR as described above.

3.10 Statistical Analyses

Data were analyzed using two-tailed student's *t*-test or analysis of variance (ANOVA) in GraphPad Prism (GraphPad Software, version 6), with p < 0.05 considered statistically significant.
CHAPTER IV – RESULTS

4.1 WNV Induces Expression of *II17a* and *II17ra* in Humans

We previously reported that WNV induces IL-23 production in mice in a TLR7-dependent manner (Town et al., 2009). Considering the role of IL-23 in Th17 cell stabilization and IL-17A production (Aggarwal et al., 2003), we hypothesized that IL-17A may play a role in WNV infection. To test this, we measured the expression of *II17a* in human cells infected with WNV *in vitro*. Human peripheral blood mononuclear cells (hPBMCs) isolated from healthy human volunteers without a history of WNV infection were infected with WNV (MOI = 0.1 and 5) for 24 h and 48 h *in vitro*. WNV-infected hPBMCs were collected, total RNA isolated, and cDNA synthesis and quantitative real-time PCR (qPCR) was performed to measure transcripts of *II17a* and cellular *β-actin* as a housekeeping gene. The qPCR results showed that *II17a* gene expression was up-regulated in WNV-infected hPBMCs (Fig. 4.1A), which was further confirmed by measuring IL-17A production in hPBMC culture supernatants (Fig. 4.1B) by an enzyme-linked immunosorbent assay (ELISA).

To relate these *in vitro* results to WNV infection in humans, we used ELISA to measure the production of IL-17A in the sera of human cases with active WNV infection (fever or neuroinvasive disease), with a history of recovered neuroinvasive WNV diseases, and healthy controls who have no history of WNV infection. The cases with active and long-standing history of neuroinvasive WNV diseases showed a trend of higher levels of IL-17A in sera compared to WNV fever cases and healthy controls (Fig. 4.1C), with no difference between the latter

two. These results demonstrate that WNV infection induces a production of IL-17A in humans and suggest that this cytokine may play a role in WNV infection.



Figure 4.1 WNV induces expression of *II17a* and *II17ra* in humans

(A) *II17a* transcripts were measured by qPCR and expressed as relative fold change (RFC) after normalization to cellular β -actin in human PBMCs infected with WNV for 24 h or 48 h. (B) IL-17A production in culture supernatant of WNVinfected hPBMC was measured by ELISA. (C) Levels of IL-17A in sera of human WNV patients and healthy controls were measured by ELISA. Error bars represent standard error of mean (mean ± SEM). Data in A and B represent two independent experiments performed in triplicates and analyzed by a one-way ANOVA. "*ns*" denotes no significant difference (p > 0.05).

4.2 WNV Induces Expression of II17a and II17ra in Mice

To expand upon the findings from human samples, we used a mouse model of WNV infection because it reflects various aspects of human WNV disease (Bai et al., 2009; Bai et al., 2005; Town et al., 2009). Splenocytes isolated from C57BL/6J mice were infected with WNV (MOI = 0.1) *in vitro* for 24 h and 48 h, and the expression of *II17a* gene was measured by qPCR. Similar to hPBMC, *II17a* transcript levels were up-regulated at both 24 and 48 h.p.i. in mouse splenocytes infected with WNV *in vitro* (Fig. 4.2A).



Figure 4.2 WNV induces expression of II17a and II17ra in mice

(A) RFC of *II17a* transcripts after normalization to cellular β -actin in mouse splenocytes infected with WNV (MOI = 0.1) in vitro. (B to C) WT (C57BL/6) mice were infected with WNV (1,000 PFU, i.p.) and expression of *II17a* (B) and *II17ra* (C) transcripts were measured in brain tissue by qPCR. Error bars represent standard error of mean (mean ± SEM). Data in A represent two independent experiments performed in triplicates and analyzed by a one-way ANOVA. Data in B and C represent two independent experiments (n = 5 mice/group) and analyzed by two-tailed student's t-test; "*ns*" denotes no significant difference (*p* > 0.05).

Since astrocytes, microglia, and brain infiltrating immune cells express functional interleukin-17 receptor A (IL-17RA) in brain inflammatory conditions (Das Sarma et al., 2009), we measured the expression of *II17a* and *II17ra* genes in brain of WNV-infected mice. For this, we infected a group of WT mice with WNV (1,000 PFU, i.p.), sacrificed them at various time points to collect brains, and measured levels of *II17a* and *II17ra* transcripts by qPCR. Indeed, there was a significantly up-regulated expression of both *II17a* (Fig. 4.2B) and *II17ra* (Fig. 4.2C) genes in brain of WNV-infected mice compared to uninfected controls. Collectively, these results indicate that WNV infection elevates the expression of both *II17a* and *II17ra* in mice, suggesting a possible role of IL-17A in WNV infection.

4.3 WNV-mediated IL-17A Production Depends on IL-23 Signaling

To further measure *II17a* expression in mice and to test whether its production was IL-23 dependent, we intraperitoneally (i.p.) infected a group of wild-type (WT) littermates and IL-23 deficient (*II23p19^{/-}*) mice (both are in mixed C57BL/6×129 background) with 1,000 plaque-forming units (PFUs) of WNV and measured IL-17A protein in plasma by ELISA. The results showed that WNV infection in mice induced IL-17A production (Fig. 4.3), but the level of this cytokine was undetectable in serum samples from mock-infected control mice (data not shown).



Figure 4.3 WNV-mediate IL-17A response in a IL-23 dependent manner

IL-17A production was measured in plasma of $ll23p19^{-/2}$ and their littermate WT control mice (7-9 week old) infected with WNV (1,000 PFU, i.p.) by ELISA. Error bars represent standard error of mean (mean ± SEM). Data represent two independent experiments (n = 3 to 5 mice/group) and analyzed by two-tailed student's t-test; "*ns*" denotes no significant difference (p > 0.05).

Moreover, there was approximately 80% reduction in *II17a* expression in *II23p19^{-/-}* mice at 3 days post infection (d.p.i), suggesting that IL-17A production during WNV infection in mice largely depends on IL-23 signaling (Fig. 4.3).

4.4 IL-17A Protects Mice from Lethal WNV Infection

To further investigate the role of IL-17A in WNV immunity, we used a mouse model of WNV encephalitis (Town et al., 2009). IL-17A deficient (*II17a^{-/-}*) and WT control mice (7-8 weeks old, sex-matched, strain C57BL/6J) were challenged *via* i.p. injection with 1,000 PFUs of WNV (Town et al., 2009), a dose that kills approximately 40 ~ 50% of WT animals. Morbidity and mortality was monitored twice daily for 21 days. We found strikingly greater susceptibility of *II17a^{-/-}* mice (20% survival) vs. WT control mice (60% survival) to lethal WNV infection (Fig. 4.4A).



Figure 4.4 *II17a^{-/-}* mice are more susceptible to WNV infection

Seven to nine weeks old WT (C57BL/6J) and *II17a^{-/-}* mice were infected with WNV via i.p. (1,000 PFU) or footpad (100 PFU) routes and monitored for mortality twice daily for 21 days; percentage (%) survival was compared using the Kaplan-Meier survival and log-rank test. (A) Survival curves after i.p inoculation, or (B) footpad inoculation.

To exclude the possibility of an inoculation route specific response, we also challenged *II17a^{-/-}* and WT mice with 100 PFUs of WNV *via* footpad (Bai et al., 2009), and performed survival analysis. Similar to the i.p. route, footpad inoculation showed that *II17a^{-/-}* mice were more susceptible to WNV infection (Fig. 4.4B). Together, these data suggest that IL-17A protects mice from lethal WNV infection.

4.5 Mice Deficient in IL-17A Develop Higher Viral Burden

To further expand the protective function of IL-17A in controlling WNV infection, we compared the virological profiles of WNV-infected *II17a^{-/-}* vs. WT mice in various tissues such as blood, spleen, liver and brain. Measurement of WNV viremia by qPCR revealed no difference at 2 d.p.i, however, a 3-fold increase in the transcript level of WNV envelope gene (*WNVE*) was observed at 4 d.p.i. in WNV-infected *II17a^{-/-}* mice compared to WT controls (Fig. 4.5A).

To assess the viral burden in peripheral organs, we sacrificed WNVinfected mice, collected liver, spleen and brain samples at the selected time points, and performed a qPCR analysis. When compared to WT controls, we found approximately 2-fold increase in *WNV-E* transcripts in the liver of *II17a^{-/-}* mice (Fig. 4.5B). Consistent with the survival results, there were about 6-fold (at 7 d.p.i.) and 30-fold (at 8 d.p.i.) increases in *WNVE* transcripts in the brain of WNV-infected *II17a^{-/-}* mice compared to WT controls (Fig. 4.5C). Although there was no difference in viral burden in spleens of WT vs. *II17a^{-/-}* mice at 4 d.p.i, *II17a^{-/-}* mice had significantly higher level (about 3-fold) of *WNVE* transcripts on 8 d.p.i (Fig. 4.5D).



Figure 4.5 IL-17A deficiency results in deficient WNV clearance in mice

Seven to nine weeks old WT (C57BL/6J) and $ll17a^{-/-}$ mice were infected with WNV via i.p. (1,000 PFU). qPCR was performed to measure *WNV-E* RNA in blood (A), liver (B), brain (C), and spleen (D) with viral burden expressed as the ratio of *WNV-E* RNA copies to cellular β -actin transcripts. The ratios of viral load between WT and $ll17a^{-/-}$ mice (mean ± SEM) were compared by two-tailed student's t-test; "*ns*" denotes no significant difference (*p* > 0.05).

These data demonstrate that mice deficient in IL-17A develop higher viral burden in blood and liver at day 4, and have deficient clearance of WNV from brain and spleen at day 8 p.i., leading to greater WNV susceptibility. Collectively, these results indicate that IL-17A plays a protective role during WNV infection.

4.6 CNS Leukocyte Slightly Elevates in WNV-infected *II17a^{-/-}* mice

IL-17A has been shown to recruit leukocytes including neutrophils during microbial infections (Happel et al., 2005; Ye et al., 2001), and inflammatory conditions (Griffin et al., 2012). Since we observed a higher WNV burden in brain

of *II17a^{-/-}* mice, we asked if this was related to IL-17A-mediated control of leukocyte infiltration into brain during WNV infection. For this, we performed confocal microscopy to detect WNV-E antigen, CD45 (pan-leukocyte marker), and CD11b (microglial and macrophage marker) in brain sections of WNVinfected *II17a^{-/-}* and WT mice sacrificed at 6 d.p.i. We focused on the olfactory bulb because we have shown that this brain region is most sensitive to WNV infection (Bai et al., 2009; Town et al., 2009; Wang et al., 2004a). Consistent with the qPCR measurement of *WNVE* RNA in brain tissue (Fig. 4.5C), the confocal imaging revealed more WNVE antigens in the brains of WNV-infected *II17a^{-/-}* mice compared to WT controls (Fig. 4.6A and B). Similar results were also obtained in other brain regions, including the cerebral cortex, brainstem, cerebellum, and striatum (data not shown). Unexpectedly, confocal imaging results showed more CD45⁺ (Fig. 4.6A) and CD11b⁺ leukocytes (Fig. 4.6B) in the brain of WNV-infected *II17a^{-/-}* mice than WT controls.

To confirm these data and further quantify brain infiltrating immune cells, we performed flow cytometric analysis of brain leukocytes isolated from WT and *II17a^{-/-}* mice infected with 1,000 PFU (i.p.) of WNV for 6 days. We characterized CD45⁺, CD11b⁺, CD3⁺CD4⁺ and CD3⁺CD8⁺ cell populations, as previously described (Town et al., 2009). Consistent with the confocal imaging results, we observed a trend towards elevation of all leukocyte population in the brain of WNV-infected *II17a^{-/-}* mice compared to WT controls (Fig. 4.6C).



Figure 4.6 CNS leukocyte profile in WNV-infected II17a-/- mice

Seven to nine weeks old WT (C57BL/6J) and *II17a^{-/-}* mice were challenged with WNV (1,000 PFU) through i.p.. (A to B) PBS perfused brains were isolated on 6 d.p.i, and WNV antigen (green signal) and CD45 (leukocyte common antigen, red, panel A) or CD11b (macrophage and microglia marker, red, panel B) were

detected by Nikon A1R confocal microscope (original magnification = 20X). DAPI (blue signal) was used as a nuclear counterstain; representative images are shown. (C) Brain leukocytes isolated on day 6 d.p.i. were characterized by flow cytometry after probing with antibodies against WNVE, CD45, CD4, CD8, and CD11b. Signal color used in dot plots and numbers representing the percentages of positive cells within gated populations are shown (right). Data represent two independent experiments (n = 5 mice per group for each experiment).

To test whether the higher leukocyte infiltration into WNV-infected *II17a^{-/-}* mouse brains was affected by leukocyte expansion or differentiation in the periphery, we compared leukocyte populations in spleen of *II17a^{-/-}* and WT mice infected with WNV. There was no difference in leukocyte populations in spleens of WNV-infected *II17a^{-/-}* vs. WT mice (data not shown), suggesting that more leukocytes in brain may not be due to the possible effects of IL-17A on leukocyte expansion in the periphery.

4.7 WNV-infected *II17a^{-/-}* Mice Upregulate CCL5

To further dissect the mechanism by which more leukocytes migrate into the brains of WNV infected *II17a*-/- mice, we performed qPCR to measure the expression of selected chemokines genes (*Cxcl1*, *Cxcl10*, *Ccl5*) known to mediate recruitment of leukocytes. There was significantly elevated expression of *Ccl5* (also known as RANTES) (Fig. 4.7A) and its receptor *Ccr5* (Fig. 4.7B) in blood of WNV infected *II17a*-/- mice at 4 d.p.i., but no difference in expression of other chemokines, such as *Cxcl1* and *Cxcl10* (data not shown). In addition, there was a significant increase of *Ccl5* expression (Fig. 4.7C) in the brains of WNVinfected *II17a*-/- mice at 8 d.p.i., but no difference in expression of other leukocyte *Cxcr3* (Fig. 4.7E). These results may imply a link between deficient IL-17A and higher *Ccl5* expression that may contribute to more leukocyte homing to the brain of $II17a^{-/-}$ mice during the course of WNV infection.





Expressions of *Ccl5* (A) and *Ccr5* (B) gene at the mRNA levels were measured by qPCR in the blood of WNV-infected mice at 2 and 4 d.p.i.. mRNA levels of *Ccl5* (C), *Cxcl10* (D) and *Cxcr3* (E) genes were measured by qPCR in brain of WNV-infected mice at 8 d.p.i. Gene expression data were normalized to cellular β -actin mRNA and compared by two-tailed student's t-test; "*ns*" denotes no significant difference (*p* > 0.05).

4.8 IL-17A Does Not Affect Interferon Response and Inflammation

WNV infection induces potent type I IFN responses in mice, which plays a critical role in controlling both viremia and encephalitis (Samuel and Diamond, 2005). We tested whether deficient IL-17A alters type I IFN expression during

WNV infection by qPCR and ELISA. The qPCR results showed no difference in *lfn-* α expression in blood of *ll17a*- 2^{-} vs. WT control mice at 4 d.p.i. (Fig. 4.8A). Similarly, no difference in expression of *lfn-* β gene was observed in blood (Fig. 4.8B), spleen (Fig. 4.8C), liver (Fig. 4.8D), and brain (Fig. 4.8E) samples from WNV-infected *ll17a*- 2^{-} vs. WT control mice measured at various time points. To further confirm these results, we also measured IFN- β protein in plasma of WNV-infected WT and *ll17a*- 2^{-} mice at 3 d.p.i. by ELISA and found no difference in IFN- β expression (Fig. 4.8F). These results suggest that the type I IFNs response remains unaltered in *ll17a*- 2^{-} mice during WNV infection.

IL-17A cause potent inflammation and regulate expression of several cytokines including IL-1β, IFNγ and TNFα (Maione et al., 2009; Song and Qian, 2013). We assessed the possible role of IL-17A in inflammatory responses during WNV infection by measuring inflammatory cytokine expression in plasma by ELISA. Again, there was no difference in levels of IL-1β, IL-6, IL-10, IFN-γ, IL-12 p40, and TNF-α in plasma from WNV-infected *II17a*-/- vs. WT control mice at both 1 and 3 d.p.i. (Fig. 4.8G-L). In addition, no significant difference in expression of these cytokines was detected by qPCR in brain of WNV-infected *II17a*-/- vs. WT control mice at 8 d.p.i. (data not shown). Collectively, these results demonstrate that higher viral load in *II17a*-/- mice is likely not due to altered production of interferon and other inflammatory cytokines.



Figure 4.8 Antiviral and inflammatory responses of WNV-infected II17a-/- mice

Seven to nine weeks old WT (C57BL/6J) and $ll17a^{-/-}$ were infected with 1,000 PFU (i.p.) of WNV. Blood, plasma, and tissue samples were collected at the selected time points (d.p.i.) for cytokine and anti-WNV-E IgM measurement. Expression of interferon- \Box (*lfn*- \Box) gene transcripts in blood (A), and expression of interferon- β (*lfn*- β) gene transcripts in blood (B), spleen (C), liver (D), and brain (E) were measured by qPCR (normalized to cellular β -actin mRNA). (F to L)

Protein level of IFN- β IL-1 β IL-6, IL-10, IFN- γ , IL-12p40, and TNF- α in plasma were measured by ELISA. Data (mean ± SEM) represent at least two independent experiments performed in triplicates and analyzed by two-tailed student's t-test; "*ns*" denotes no significant difference (*p* > 0.05).

4.9 IL-17A Does Not Have a Direct Anti-WNV Effect

We next asked if IL-17A has a direct antiviral activity against WNV infection, which has been shown for some other cytokines, such as TNF-α (Ruggiero et al., 1989; Shrestha et al., 2008b) and IL-6 (Moore et al., 2012). However, no effect in replication of WNV was observed in Raw 264.7 (mouse macrophage, Fig. 4.9A) and Neuro 2a (mouse neuroblast, Fig. 4.9B) cells that were pretreated with mouse recombinant IL-17A (1 to 100 ng/ml), suggesting that IL-17A may not have a direct antiviral effect against WNV replication.



Figure 4.9 IL-17A does not directly control WNV replication

Replication of WNV was analyzed by qPCR in Raw 264.7 (A) and Neuro 2a (B) cells that were pretreated with mouse recombinant IL-17A (1 to 100 ng/ml) for 6 h followed by infection with WNV (1 MOI) for 24 h. Data (mean \pm SEM) represent at least two independent experiments performed in triplicates and analyzed by one-way ANOVA; "*ns*" denotes no significant difference (*p* > 0.05).

4.10 IL-17A Does Not Control Humoral Immune Response

Besides type I IFN, humoral immune response also plays an important role in clearance of WNV from the blood and peripheral organs and limits viral dissemination to the CNS (Diamond et al., 2003a). Although the role of IL-17A in humoral immunity is not well understood, it has been shown that B cells express IL-17RA (Yao et al., 1995), whereas Th17 cells (major IL-17A producers) promote B cells to produce antibodies (Yuan et al., 2010). To test the possible effect of IL-17A in humoral immune responses during WNV infection, we compared WNV-E specific IgM antibody production in WNV-infected *II17a^{-/-}* and WT mice by ELISA. Both *II17a^{-/-}* and WT mice produced similar levels of anti-WNV-E IgM when measured at 2, 3, and 5 d.p.i. (Fig. 4.10). These results demonstrate that higher viral load in *II17a^{-/-}* mice is likely, not due to altered antibody responses.



Figure 4.10 WNV-mediated IL-17A does not affect antibody production

Seven to nine weeks old WT (C57BL/6J) and $II17a^{-/-}$ were infected with 1,000 PFU (i.p.) of WNV. Anti-WNV-E IgM antibody levels in plasma were measured by ELISA. Data (mean ± SEM) represent at least two independent experiments performed in triplicates and analyzed by two-tailed student's t-test; "*ns*" denotes no significant difference (*p* > 0.05).

4.11 IL-17A Deficiency Causes the Reduced CD8⁺ T Cell Cytotoxicity

Brain-infiltrating leukocytes play a vital role in clearing WNV from the CNS during WNV infection (Shrestha and Diamond, 2004; Sitati and Diamond, 2006; Town et al., 2009). In particular, CD8⁺ T cells are crucial for clearance of WNV

from the CNS and spleens (Shrestha and Diamond, 2004, 2007; Shrestha et al., 2006a; Wang et al., 2004b). Despite a modest elevation trend of brain infiltrating CD8⁺ T cells in *II17a^{-/-}* mice, viral burden in the brains of these mice was higher than in WT controls (Fig. 4.5C). In addition, $II17a^{-/-}$ mice were also deficient in clearing WNV from spleen (Fig. 4.5D). Therefore, we hypothesized that CD8⁺ T cells in *II17a^{-/-}* mice may be functionally defective in their ability to clear WNVinfected target cells. To test this, we infected WT and *II17a^{-/-}* mice through i.p. with a sub-lethal dose of WNV (100 PFU) to prolong the course of WNV infection. This is important because CD8⁺ T cells play a major role in clearing WNVinfected cells during the later phase (day 8 to 12) of infection (Shrestha and Diamond, 2004; Wang et al., 2004b), and most of *II17a^{-/-}* mice infected with a higher dose (e.g. 1,000 PFU or more) develop severe diseases and die during this time period. At 10 d.p.i., mice were sacrificed and splenic CD8⁺T cells were purified using a negative antibody selection method. The purified effector CD8⁺T cells were then co-cultured with the target cells (MC57GL_{WNV-E}) or control cells (MC57GL_{vector}) for 4 h. The target cells express the ectodomain of the WNVE in a pcDNA3.1 vector, while the control cells only express the parent vector (Shrestha and Diamond, 2004). The cytotoxicity of effector CD8⁺ T cells to WNV specific target cells was assessed by measuring the quantity of intracellular lactate dehydrogenase released into culture supernatants from the lysed target cells. Strikingly, the cytotoxicity assay showed about 2-fold reduction in cytotoxicity of CD8⁺ T cells isolated from WNV-infected *II17a^{-/-}* mice in comparison to WNVinfected WT mice (Fig. 4.11A). These results demonstrate that CD8⁺ T cells from

II17a^{-/-} mice failed to mount an effective target cell-specific cytotoxic response during WNV infection.



Figure 4.11 CD8⁺ T cells from *II17a^{-/-}* mice have reduced cytotoxicity

Seven to nine weeks old WT (C57BL/6J) and *ll17a^{-/-}* mice were infected with WNV at 100 PFU (i.p.) for 10 days. (A) Purified splenic CD8⁺T cells were co-cultured with the target (MC57GL_{WNV-E}) or control (MC57GL_{vector}) cells at a 50:1 effector/target ratio for 4 h and cytotoxicity was assayed by measuring the release of intracellular lactate dehydrogenase in culture supernatants. (B to E) RFC in transcripts of *perforin-1* (B), *granzyme A* (C), *granzyme B* (D), and *FasL* (E) in indicated tissues of WT or *ll17a^{-/-}* mice were measured by qPCR (normalized to cellular β -actin mRNA). Data (mean ± SEM) represent three independent experiments (n = 3 mice/group). "*ns*" denotes no significant difference (*p* > 0.05).

Cytotoxicity of CD8⁺ T cells employ granule (e.g., perforin and granzyme) exocytosis and Fas-Fas ligand (FasL) dependent mechanisms to kill target cells (Shresta et al., 1998). To test whether IL-17A regulates the expression of cytotoxicity mediator genes, we performed qPCR assay in blood, spleen, liver, and brain of WT and $II17a^{-/-}$ mice that were infected with WNV (100 PFU) for 10 days. Interestingly, the expression levels of *perforin-1* (Fig. 4.11B), *granzyme A* (Fig. 4.11C), *granzyme B* (Fig. 4.11D) and *FasL* (Fig. 4.11E) were significantly lower in most of these tissues collected from $II17a^{-/-}$ mice compared to WT controls.

4.12 II17a^{-/-} CD8⁺ T Cells Express Less Cytotoxicity Mediators

Although CD8⁺ cells are the major cells that express these cytotoxic mediators, other cells, such as NK cells, may also express these genes and contribute to cytotoxic effector function by the mechanism similar to CD8⁺ T cells. To specifically test if the attenuated cytotoxicity of CD8⁺ T cells in *ll17a^{-/-}* mice was due to the lower expression of these cytotoxic mediator genes, we performed qPCR to measure the expression of *perforin-1*, *granzyme A*, *granzyme B* and *FasL* in CD8⁺ T cells purified from spleens of WNV-infected WT and *ll17a^{-/-}* mice at 10 d.p.i.. Consistent with the cytotoxicity assay results, the qPCR showed that CD8⁺ T cells isolated from the *ll17a^{-/-}* mice have significantly reduced expression of *perforin-1* (Fig. 4.12A), *granzyme A* (Fig. 4.12B), *granzyme B* (Fig. 4.12C) and *FasL* (Fig. 4.12D) compared to WT controls.



Figure 4.12 *II17a^{-/-}* CD8⁺ T cells express reduced cytotoxicity mediators

Seven to nine weeks old WT (C57BL/6J) and *ll17a*^{-/-} mice were infected with WNV at 100 PFU (i.p.) for 10 days and euthanized to obtain splenic CD8⁺T cells. (A to D) RFC in expression of transcripts of *perforin-1* (A), *granzyme A* (B), *granzyme B* (C), and *FasL* (D) in splenic CD8⁺ T cells or CD8⁻ cells from WT or *ll17a*^{-/-} mice were measured by qPCR (normalized to cellular β -*actin* mRNA). (E to F) RFC in the expressions of *perforin-1* (E), and *granzyme A* (F) in the spleens of CHIKV-infected (10⁵ PFU, i.p.) WT and *ll17a*^{-/-} mice were measured on 12 d.p.i. by qPCR. Data (mean ± SEM) shown in A to D represent three independent experiments (n = 3 mice/group); and data in E to F represent two independent experiments (n = 3 mice/group). "*ns*" denotes no significant difference (*p* > 0.05).

To test if these cytotoxicity mediators were also less expressed in other immune cells in *II17a^{-/-}* mice, we performed qPCR assay on CD8⁻ splenocytes (mixed immune cells including NK cells) isolated from WNV-infected WT and *II17a^{-/-}* mice at 10 d.p.i. In contrast to the CD8⁺ cells, we found no difference in the expression of *perforin-1*, *granzyme A*, *granzyme B* and *FasL* genes in CD8⁻ splenocytes from *II17a^{-/-}* mice (Fig. 4.12A-D). Whereas in WNV-infected WT mice, the expression of these cytotoxicity marker genes was significantly higher in CD8⁺ than CD8⁻ cells, no such differences were detected between CD8⁺ and CD8⁻ cells isolated from *II17a^{-/-}* mice (Fig. 4.12A-D). In addition, there was no significant difference in the expression of these cytotoxicity marker genes between uninfected *II17a^{-/-}* and WT mice (data not shown).

To further test whether the IL-17A-mediated cytotoxicity mediator expression is specific to WNV, or common across viral infections, we measured the expression of cytotoxicity mediator genes in WT and *II17a^{-/-}* mice infected with chikungunya virus (CHIKV), a single-stranded RNA virus from genus *Alphavirus* of the family *Togaviridae*. Similar to WNV, we found significantly reduced expressions of *perforin*-1 (Fig. 4.12E) and *granzyme A* (Fig. 4.12F) in the splenocytes isolated from CHIKV-infected *II17a^{-/-}* mice compared to WT controls. Collectively, these results suggest that IL-17A promotes the expression of the cytotoxicity mediators and facilitate CD8⁺ T cells cytotoxicity during WNV and other viral infections, such as CHIKV.

4.13 IL-17A Induces Cytotoxic Mediators Independent of CD4⁺ T Cells

The role of IL-17A in regulating the expression of cytotoxic mediators in CD8⁺ T cells has not been previously reported. To further investigate and confirm this, we isolated splenocytes from WNV-infected WT mice at 8 d.p.i., cultured them with recombinant IL-17A (50 ng/ml) *ex vivo* for 24 h, and measured the expression of *perforin-1*, *granzyme A*, *granzyme B* and *FasL* genes by qPCR assay. The treatment with recombinant IL-17A significantly up-regulated the expression of *perforin-1*, *granzyme A*, and *granzyme B*, but not *FasL* (Fig. 4.13A-D), in the splenocytes isolated from WNV-infected WT mice. Similar results were

obtained when splenocytes were isolated from WNV-infected *II17a^{-/-}* mice and treated with recombinant IL-17A *ex vivo* (Fig. 4.13A-D).



Figure 4.13 IL-17A promotes cytotoxic mediators independently of CD4⁺ cells

Seven to nine weeks old WT (C57BL/6J) and $ll17a^{-/-}$ mice (n = 6 mice/group) were infected with 100 PFU (i.p.) of WNV and sacrificed on 8 d.p.i. to isolate splenocytes and CD8⁺ T cells. Splenocytes or CD4⁺ T cells depleted splenocytes (CD4⁻ splenocytes) were cultured with or without recombinant mouse IL-17A (50 ng/ml) for 24 h *ex vivo* and RFC of transcripts of *perforin-1* (A), *granzyme A* (B), *granzyme B* (C), and *FasL* (D) were measured by qPCR (normalized to cellular β -*actin* mRNA). All data (mean ± SEM) represent two independent experiments (n = 3 mice/group) and compared by two-tailed student's t-test. All data from WT and $ll17a^{-/-}$ mice were normalized to respective mock treated controls.

To assess the possible role of CD4⁺ T cells in IL-17A-mediated expression of cytotoxic mediators, we depleted CD4⁺ T cells from the splenoocytes isolated from WNV-infected *II17a^{-/-}* and WT mice on 8 d.p.i., and cultured CD4⁻ splenocytes with recombinant IL-17A (50 ng/ml) *ex vivo* for 24 h. The supply of recombinant IL-17A induced expression of the cytotoxic mediators even in the absence of CD4⁺T cells (Fig. 4.13A-D), suggesting that IL-17A-mediated induction of cytotoxic mediators expression in CD8⁺T cell is independent of CD4⁺ T cells.

4.14 IL-17A-induced Cytotoxicity Mediators May Involve ACT-1 Signaling

IL-17A exerts its function through IL-17A receptor (IL-17R), which is a heterodimeric receptor complex of IL-17RA and IL-17RC. Although the signaling pathway downstream of IL-17R has not been completely understood, one common pathway that has been characterized is the activation of the classical NF-κB pathway (Gaffen, 2009; Gu et al., 2013). Upon binding to IL-17A, SEFIR domain of IL-17R recruits the signaling adaptor ACT-1, which further recruits an essential upstream activator of the classical NF-κB pathway called TRAF6 (Gaffen, 2009). To test if IL-17A signaling in WNV-infected cells occurs *via* ACT-1, we measured the expression of *Act-1* in splenocytes and CD8⁺ T cells isolated from WNV-infected mice and treated *ex vivo* with recombinant IL-17A, as described above. The qPCR results showed that IL-17A treatment up-regulated the expression of *Act-1* gene in splenocytes (Fig. 4.14A) and CD8⁺ T cells (Fig. 4.14B) isolated from WNV-infected mice.

To further test if IL-17A-mediated expression of cytotoxicity mediators involves NF-κB activation, we treated splenocytes isolated from WNV-infected WT mice with recombinant IL-17A in the presence of Bay-11-7082 (an inhibitor of NF-κB) and measured the expression of the cytotoxicity mediators, as described above. Consistent with previous reports (Huang et al., 2007; Ribaux et al., 2007), Bay-11-7082 inhibited the IL-17A mediated *Cxcl1* expression that involves NF-κB pathway. However, inhibition of NF-κB did not inhibit, but interestingly further up-

regulated the IL-17A-mediated expression of *perforin-1*, *granzyme A*, and *granzyme B* (Fig. 4.14C). These data indicate that IL-17A-mediated expression of cytotoxicity mediators may involve ACT-1, but may occur independently of NF ĸB pathway, which requires further investigation.



Figure 4.14 IL-17A induces the expression of ACT-1

Seven to nine weeks old WT (C57BL/6J) and *ll17a*^{-/-} mice (n = 6 mice/group) were infected with 100 PFU (i.p.) of WNV and sacrificed on 8 d.p.i. to isolate splenocytes and CD8⁺ T cells. Splenocytes (A) or purified CD8⁺ T cells (B) were cultured *ex vivo* with or without recombinant IL-17A (rlL17A, 50 ng/ml) for 24 h, and expression (RFC) of *Act*-1 was by measured qPCR. (C) RFC of *Act*-1 in splenocytes cultured *ex vivo* for 24 h with or without rlL17A (50 ng/ml), in the presence of Bay-11-7082 (4 μ M) or DMSO as vehicle control (< 0.05%). All data (mean ± SEM) represent two independent experiments (n = 3 mice/group) and compared by two-tailed student's t-test. All data from WT and *ll17a*^{-/-} mice were normalized to respective mock treated controls. * denotes *p* < 0.05; ** denotes *p* < 0.005; and "*ns*" denotes not significant (*p* > 0.05).

4.15 IL-17A Directly Induces Cytotoxic mediators in CD8⁺ T Cells

IL-17R is expressed by virtually all cell types and tissues examined (Moseley et al., 2003; Yao et al., 1995; Yao et al., 1997). CD8⁺ T cells also express IL-17R (Lindemann et al., 2008; Yao et al., 1995), however, the functional role of this receptor in CD8⁺ T cell biology has not yet been recognized. To test if IL-17A may directly promote cytotoxic mediators expression in CD8⁺ T cells, we treated CD8⁺ T cells purified (~ 80 to 90% pure) from spleens of WNV-infected WT mice with the recombinant IL-17A (50 ng/ml) *ex vivo* for 24 h, then measured expression of cytotoxicity mediators by qPCR. IL-17A treatment significantly induced the expression of *perforin-1* (Fig. 4.15A), *granzyme A* (Fig. 4.15B), and *granzyme B* (Fig. 4.15C), but not *FasL* (Fig. 4.15D) in CD8⁺ T cells purified from WNV-infected WT mice. Similar results were also obtained in CD8⁺ T cell purified from WNV-infected *II17a^{-/-}* mice (Fig. 4.15A-D). In a separate experiment, we cultured splenocytes from WNV-infected *II17a^{-/-}* and WT mice with recombinant IL-17A (50 ng/ml) *ex vivo* for 24 h, separated CD8⁺ T cells from CD8⁻ cells, and then measured expression of cytotoxic mediators by qPCR. Consistently, IL-17A treatment induced the expression of cytotoxic mediators in CD8⁺ T cells (data not shown), but not in CD8⁻ cells (Fig. 4.15A-D).

To further confirm the role of IL-17A in promoting the expression of the cytotoxic mediators in brain CD8⁺ T cells *in vivo*, we infected *II17a^{-/-}* mice with WNV (100 PFU), treated them with recombinant IL-17A (2.5 µg per mouse at day 6 p.i.), and performed flow cytometric analysis of brain leukocytes. The results showed that treatment of WNV-infected *II17a^{-/-}* mice with recombinant IL-17A significantly induced the production of perforin and granzyme A in brain infiltrating CD8⁺ T cells (Fig. 4.15E). Taken together, these results suggest that IL-17A promotes the expression of cytotoxic mediator genes in CD8⁺ T cells during WNV infection in mice.



Figure 4.15 IL-17A directly induces cytotoxicity mediators in CD8⁺ T cells

(A-D) Seven to nine weeks old WT (C57BL/6J) and *ll17a^{-/-}* mice (n = 6) were infected with 100 PFU (i.p.) of WNV and sacrificed at 8 d.p.i. to collect spleens, followed by magnetic separation of CD8⁺ T cells. Relative fold change (RFC) of the transcripts of *perforin-1* (A), *granzyme A* (B), *granzyme B* (C), and *FasL* (D) were measured in CD8⁺ or CD8⁻ T cells cultured *ex vivo* for 24 h with or without recombinant mouse IL-17A (50 pg/ml) by qPCR. (E) Eight week old *ll17a^{-/-}* mice (n = 4 per group) infected with WNV (100 PFU, i.p.) were

treated with recombinant IL-17A (rIL17A, 2.5 µg/mouse) or PBS (control) via i.p. at day 6 p.i and sacrificed at 8 p.i. to characterize brain leukocytes by flow cytometry. Mean fluorescence intensities of perforin and granzyme A within the gated CD45^{hi} CD8⁺ cells (green) are shown in bar (right). All data (mean ± SEM) represent two independent experiments and compared by two-tailed student's t-test. All data from WT and $II17a^{-/-}$ mice (shown in figures A to D) were normalized to respective mock treated controls. "*ns*" denotes no significant difference (*p* > 0.05).

4.16 IL-17A Treatment Increases Survival of WNV-infected Mice

Since CD8⁺T cells are essential to clear WNV from the CNS at a later phase of infection, we hypothesized that recombinant IL-17A treatment *in vivo* may offer protection from WNV infection by promoting cytotoxicity of CD8⁺T cells. To test if IL-17A may serve as a therapeutic reagent to treat WNV infection in mice, we infected WT female mice with WNV (100 PFU) via i.p. route. At day 6 p.i, mice were injected with the carrier-free mouse recombinant IL-17A (eBioscience) or PBS as control *via* i.p. route for survival analysis.



Figure 4.16 Recombinant IL-17A increases survival of WNV-infected mice WT (C57BL/6J) mice (8-week old, female) were infected with 100 PFU of WNV via i.p. route. At day 6 p.i, mice were injected with i.p. administration of carrier-free mouse recombinant IL-17A (rIL17A, 2.5 μ g/mouse) or PBS. (A) Percent (%) of survival was compared using the Kaplan-Meier survival and log-rank test. Viral burden in brain (B) was measured by qPCR and compared by two-tailed student's t-test; "*ns*" denotes no significant difference (*p* > 0.05).

Mice that received recombinant IL-17A showed significantly increased survival rate compared to the PBS-treated control mice (Fig. 4.16A). Consistent with the survival results, mice treated with recombinant IL-17A also showed lower viral burden in the brain compared to the PBS-treated control mice at day 8 p.i. (Fig. 4.16B).

4.17 IL-17A Induces Cytotoxicity Mediators in WNV-infected Mice

To test if the effect of the recombinant IL-17A treatment involves a promotion of cytotoxicity of CD8⁺ T cells, we also measured the expression of the cytotoxicity mediators in splenic CD8⁺ and CD8⁻ T cells



Figure 4.17 Recombinant IL-17A induces cytotoxicity in WNV-infected mice.

WT (C57BL/6J) mice (8-week old, female) were infected with 100 PFU of WNV via i.p. route. At day 6 p.i, mice were injected with i.p. administration of carrier-free mouse recombinant IL-17A (rIL17A, 2.5 μ g/mouse) or PBS. (A to E) The transcripts of *perforin-1*, *granzyme A*, *granzyme B*, and *FasL* in splenic CD8⁺ and CD8⁻ T cells (A-D), and brain (E) were measured by qPCR and compared by two-tailed student's t-test; "*ns*" denotes no significant difference (*p* > 0.05).

. Consistent with reduced viral burden and increased survival,

administration of recombinant IL-17A in WNV-infected mice induced the expression of the cytotoxic mediators in CD8⁺ T cells, but not in CD8⁻ T cells (Fig. 4.17A-D), which agreed with our *in vitro* results. In addition, we also detected increased expression of *granzyme A*, *granzyme B*, and *FasL* in brain tissues of WNV-infected mice after IL-17A treatment (Fig. 17E). Collectively, these results suggest a novel and promising therapeutic role of IL-17A in facilitating WNV clearance by promoting CD8⁺ T cell cytotoxicity.

CHAPTER V – DISCUSSION

This study reveals a novel role of IL-17A in facilitating WNV clearance by inducing the expression of cytotoxic mediator genes and promoting CD8⁺T cell cytotoxicity. Specifically, we report here that 1) WNV induces IL-17A expression in both mice and humans; 2) *II17a^{-/-}* mice, generate a higher viral burden and are more susceptible to WNV infection; 3) CD8⁺T cells purified from *II17a^{-/-}* mice are less cytotoxic and express lower levels of cytotoxic mediators, i.e., *perforin-1*, *granzyme A*, *granzyme B* and *FasL*; and most importantly; and 4) *ex vivo* supply of recombinant IL-17A as late as day 6 post-infection significantly reduces viral burden in the brain and increases survival rate of WNV-infected mice, suggesting a therapeutic potential of IL-17A.

5.1 IL-17A Has a Protective Function in WNV Infection

It has been previously reported that WNV mediates production of IL-23, a cytokine that regulate IL-17A expression (Town et al., 2009). As expected, we found that WNV induced IL-17A production in human and mouse in an IL-23 dependent manner. We further found a protective function of IL-17A against WNV and its crucial role in clearance of WNV from brain, which was also consistent with function of IL-23 (Town et al., 2009). In support of our finding, it has been previously reported that mice depleted of IL-17A function by administration of anti-IL-17A antibody showed about 20% reduced survival rate compared to control mice after a lethal WNV challenge (Welte et al., 2011). However, no difference in WNV viral burden was observed between anti-IL-17A antibody treated and control mice (Welte et al., 2011). In contrast, our results

clearly demonstrate that *II17a*^{-/-} mice are more susceptible to WNV infection. The discrepancy between these two studies may be due to the transient effects of anti-IL-17A treatment when compared to complete genetic deficiency of IL-17A in *II17a*^{-/-} mice. More to this point, $\gamma \delta$ -T cell produce IL-17A during early phage of WNV infection, whereas CD4⁺ Th-17 cells become the major IL-17A producer during the late phase of infection (Hamada et al., 2008). Thus, the administration of anti-IL-17A antibody at an early phase of WNV infection (0 and 5 d.p.i.) (Welte et al., 2011), may not have sufficiently blocked IL-17A production and its function during the later phase of infection. Indeed, we found that IL-17A was not essential for controlling the viremia on 2 d.p.i, but became critical for reducing viremia on 4 d.p.i and clearing the viruses from spleen and brain on 8 d.p.i.. Thus, it appears that IL-17A-mediated protective immunity against WNV infection predominantly occurs during the late phase of infection, which requires further investigation.

5.2 Role of IL-17A in IFN, Cytokines and Antibody Response

IL-17A is a pleiotropic cytokine that plays key roles during infection (Iwakura et al., 2008), inflammation (Park et al., 2005) and autoimmune diseases (Onishi and Gaffen, 2010). Further, it regulates the expression of a number of cytokines, including IL-1β, IFN- γ , and TNF- α (Maione et al., 2009; Song and Qian, 2013). Considering their critical roles in WNV pathogenesis (Ramos et al., 2012; Shrestha et al., 2006b; Shrestha et al., 2008b; Wang et al., 2004a), it is plausible that IL-17A may regulate the expression of these cytokines during WNV infection. However, we did not detect significant differences in the expression of IL-1β, IFN-γ, and TNF-α in WNV-infected *II17a*^{-/-} mice when compared to WT controls. These results suggest the likelihood that IL-17A-mediated protective immunity against WNV infection may not be associated with functions controlled by these cytokines. Previous studies have suggested that type I IFNs (antiviral cytokines), which can potently suppress IL-17A expression (Curtis et al., 2009; Tilg et al., 2009), play a prominent role in controlling WNV infection (Lazear et al., 2011; Samuel and Diamond, 2005). Again, we did not detect any change in the expression of type I IFNs in *II17a*^{-/-} vs. WT control mice during WNV infection. Moreover, IL-17A has been suggested to regulate the humoral immune response (Grund et al., 2012; Tarlinton, 2008; Yuan et al., 2010), however, we did not see such effect during WNV infection in mice. Despite unaltered inflammatory response, type I IFN expression and humoral immune response, *II17a*^{-/-} mice generated higher viral burden, suggesting that IL-17A mediated protective immunity during WNV infection is independent of these immune responses.

5.3 Role of IL-17A in Leukocyte Homing to CNS

WNV invades the CNS and infects neurons and CNS-resident cells, such as microglia and astrocytes, which leads to production of cytokines and chemokines that recruit peripheral leukocytes into the brain (Glass et al., 2005; Hussmann and Fredericksen, 2014; Shrestha and Diamond, 2004; Town et al., 2009). We previously reported that IL-23 has an important role in recruiting CD11b⁺ monocytes and macrophages into the CNS to control WNV infection (Town et al., 2009). Considering the role of IL-23 in producing IL-17A, and the role of both of these cytokines in inducing and sustaining leukocyte recruitment to infected sites (Roussel et al., 2010; Shahrara et al., 2009), we hypothesized that leukocyte migration into the CNS would be reduced in *II17a^{-/-}* mice. Surprisingly, we detected a modestly elevated leukocyte influx into WNV-infected *II17a^{-/-}* mouse brains by both flow cytometry and confocal microscopy.

In dissecting the mechanism, we found that the expression of CCL5 was elevated in both the CNS and the peripheral tissues of WNV-infected *II17a^{-/-}* mice compared to WT controls. CCL5 plays a protective role in WNV, influenza and parainfluenza infections by promoting leukocyte trafficking to infected tissues (Glass et al., 2005; Kohlmeier et al., 2008). Also, Ccl5 expression sustains CD8+ T cell responses during influenza (Kohlmeier et al., 2008), parainfluenza (Kohlmeier et al., 2008) and chronic lymphocytic choriomeningitis virus (LCMV) (Crawford et al., 2011) infections. The up-regulation of Ccl5 in $II17a^{-/-}$ mice may be due to several reasons. First, the higher viral load in *II17a^{-/-}* mice could trigger a stronger Ccl5 expression (Matsukura et al., 1998; Saito et al., 1997). Secondly, the presence of IL-17A in WT mice may suppress Ccl5 expression, since IL-17A can down-regulate Ccl5 expression (Andoh et al., 2002; Shen et al., 2005). Although increased *Ccl5* expression may account for modest elevation in brain infiltrating leukocytes in *II17a^{-/-}* mice, this did not result in protection of these mice from WNV infection, implying a possible functional defect in brain infiltrating effector leukocytes in *II17a^{-/-}* mice.

5.4 IL-17A Regulates CD8⁺ T Cell Cytotoxicity

CD8⁺ T cells play a critical role in clearance of viruses from the CNS (Binder and Griffin, 2001; Griffin, 2010; Shrestha and Diamond, 2004). CD8⁺ T

cells can control viral infection directly by inducing apoptosis of virus-infected cells via perforin, granzyme, or Fas-FasL interactions (Harty and Badovinac, 2002; Shresta et al., 1998), or indirectly by immune-mediated non-cytolytic clearance of viruses from neurons by producing cytokines, such as TNF- α and IFN-y (Binder and Griffin, 2001; Griffin, 2010). Mice deficient in the expression of cytotoxic mediators such as *perforin*, *FasL*, or cytokines, such as TNF- α and IFN- γ , exhibit increased mortality and viral burden in the CNS and peripheral organs following WNV infection (Shrestha and Diamond, 2007; Shrestha et al., 2012; Shrestha et al., 2006a; Shrestha et al., 2006b; Shrestha et al., 2008b; Wang et al., 2004b). However, we did not detect any significant difference in the expression of the cytokines including TNF- α and IFN-y between WNV-infected WT and *II17a^{-/-}* mice, suggesting that IL-17A-mediated control of WNV by CD8⁺T cells may occur independently of the effects of these cytokines. We found that CD8⁺ T cells isolated from *ll17a^{-/-}* mice have significantly reduced cytotoxicity compared to WT controls, which may explain higher viral burden and lower survival rates of *II17a^{-/-}* mice, despite elevated leukocyte infiltration (including CD8⁺ T cells) into the CNS. While we are not certain whether CD8⁺ T cells predominantly contribute to WNV clearance by causing cytolysis of WNV-infected neurons due to lower MHC I expression in these cells (Kimura and Griffin, 2000; Mucke and Oldstone, 1992), it is likely that CD8⁺ T cells can cause cytolysis of WNV-infected non-neural cells, such as infiltrating macrophages and neutrophils, and microglia in the CNS, thus facilitating the virus clearance (Bai et al., 2010; Shrestha and Diamond, 2004, 2007; Shrestha et al., 2006a). In line with reduced

cytotoxicity, CD8⁺ T cells from *II17a^{-/-}* mice had significantly reduced expression of *perforin-1*, *granzyme A*, *granzyme B* and *FasL* in most of the tissue examined during WNV infection. Although NK cells (type of CD8⁻ cells) also express these genes and mediate apoptosis of target cell, we did not detect difference in expression of cytotoxic mediators in CD8 negative (CD8⁻) cell populations isolated from WNV-infected WT and *II17a^{-/-}* mouse spleens. This indicates that NK cells (type of CD8⁻ cells) may not play a prominent role in IL-17A-mediated cytotoxicity during WNV infection. This notion is consistent with the previous reports that NK cells have a little or no role in controlling WNV infection (Shrestha et al., 2006a).

It is worthy to note that CD8⁺T cells have also been shown an immunopathological role when mice were infected intravenously with a high dose (10⁸ PFU) of WNV (Sarafend strain), resulting in 100% mortality with a 6-day mean survival time (Wang et al., 2003). However, such an immunopathology role of CD8⁺ T cells was likely not due to their cytotoxic functions because these cells are usually activated after one week post-infection. In support of this, that same study and many other reports also showed a recovery role of CD8⁺ T cells when mice were infected with low doses (10²-10³ PFU) of WNV, which is similar to the current study (Brien et al., 2011; Brien et al., 2007; Shrestha et al., 2008a; Shrestha et al., 2012; Shrestha et al., 2006a; Wang et al., 2003). In addition, Th17 cells and IL-17A have been implicated in inflammation and immunopathology associated with autoimmune diseases (McFarland and Martin, 2007; van den Berg and Miossec, 2009; Zepp et al., 2011) and some virus-

induced chronic CNS diseases (Hou et al., 2014; Hou et al., 2009; Savarin et al., 2012), suggesting that Th-17/IL-17A axis may have different implications in such immunopathological and chronic inflammatory conditions.

5.5 IL-17A Has a Potential Therapeutic Role Against WNV

Our data demonstrates that IL-17A deficiency leads to an attenuated CD8⁺ T cell cytotoxicity and deficient WNV clearance. Both CD4⁺ Th-17 cells and $\gamma \delta \Box T$ cells, the major cells that produce IL-17A, have been previously shown to promote CD8⁺ T cell cytotoxicity during infection (Hamada et al., 2009; Xu et al., 2010), autoimmune disease (Ankathatti Munegowda et al., 2011a), and cancers (Ankathatti Munegowda et al., 2011b; Martin-Orozco et al., 2009). Also, NK cells treated with IL-17A overexpress *perforin* and *granzymes*, and have better cytotoxic functions (Al Omar et al., 2013), which provides additional evidence that IL-17A signaling can induce the expression of cytotoxicity mediators. These reports support our finding and suggest that IL-17A may facilitate cytotoxicity of CD8⁺ T cells in diverse disease conditions.

Various types of immune cells, such as CD4⁺ Th-17 cells, $\gamma \delta$ -T cells, NKT cells, and CD8⁺ T cells can produce IL-17A (Gu et al., 2013; Welte et al., 2011). In addition, the IL-17A receptor (IL-17R) is expressed ubiquitously in virtually all cell types and tissues examined (Andoh et al., 2002; Moseley et al., 2003; Yao et al., 1997). Although CD8⁺ T cells can produce IL-17A and also express its cognate receptor (Lindemann et al., 2008; Yao et al., 1995), the link between IL-17A and CD8⁺ T cell cytotoxic function has not been previously recognized. In theory, it is possible that IL-17A may affect CD8⁺ T cell development and/or

function either by directly acting on these cells or indirectly through other cell types. Since we found similar levels of CD8⁺ T cells and CD4⁺/CD8⁺ ratios in splenocytes isolated from *II17a^{-/-}* and WT mice infected with WNV, it appears that CD8⁺ T cell development remains unaltered in $II17a^{-/-}$ mice and IL-17A may largely control CD8⁺ T cell function during WNV infection. This hypothesis is confirmed by ex vivo treatment of IL-17A to splenocytes isolated from WNVinfected *II17a^{-/-}* and WT mice, which showed an up-regulation of *perforin-1*, granzyme A, and granzyme B expression in CD8⁺ T cells by IL-17A, in a CD4⁺ T cell-independent manner. These results suggest that IL-17A can induce the expression of cytotoxic mediators by directly acting on CD8⁺ T cells. However, the induction of FasL by recombinant IL-17A was differentially affected in our experiments. For instance, in vitro IL-17A treatment did not induce the expression of FasL in CD8⁺T cells, while it was induced after in vivo administration of IL-17A. This discrepancy is likely due to the experimental conditions or may reflect differential role of IL-17A in regulating the expression of these cytotoxic mediators, which requires further investigation. We further demonstrated that the treatment of WNV-infected *II17a^{-/-}* mice with recombinant IL-17A significantly induced the production of perforin and granzyme A in brain infiltrating CD8⁺ T cells, which confirmed the role of IL-17A in promoting the expression of the cytotoxic mediators in brain infiltrating CD8⁺ T cells in vivo.

To extend upon our finding, we tested therapeutic potential of IL-17A in a mouse model of WNV infection. Importantly, administration of a single dose of recombinant IL-17A to WNV-infected mice, even as late as day 6 p.i., induced
the expression of cytotoxic mediators in CD8⁺T cells, dramatically reduced viral burden in the brain, and significantly increased survival rate, suggesting a promising therapeutic role of IL-17A against WNV infection.

CHAPTER VI – CONCLUSION AND FUTURE DIRECTIONS

6.1 Conclusion

IL-17A and CD8⁺ T cells regulate diverse immune functions during microbial infections, malignancies, and autoimmune diseases. IL-17A is a proinflammatory cytokine produced by diverse cell types, while the CD8⁺ T cells (known as cytotoxic T cells) are major cells that provide immunity against intracellular pathogens. Previous studies have demonstrated a crucial role of CD8⁺ T cells in recovery from WNV infection. Here, we demonstrate that IL-17A protects mice from lethal WNV infection by promoting CD8⁺ T-cell-mediated clearance of WNV. In addition, treatment of WNV-infected mice with recombinant IL-17A reduces viral burden and increases survival of mice, suggesting a potential therapeutic. In conclusion, this study uncovers a novel function of IL-17A in promoting CD8⁺ T cell axis may also have broad implications in immunity to other microbial infections and cancers, where CD8⁺ T cell functions are crucial.

6.2 Future Directions

In this study, we found the production of IL-17A in WNV-infected human and mice, which we also confirmed in a mixed culture of human and mouse immune cells infected *in vitro* with WNV. We further demonstrated a novel function of IL-17A in promoting the cytotoxicity of CD8⁺ T cell against WNV infection. In addition, a supply of recombinant IL-17A to a culture of CD8⁺ T cells and WNV-infected mice clearly induced the expression of cytotoxicity mediator genes. The role of IL-17A in promoting the expression of cytotoxic mediators in

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CD8⁺ T cells has not been previously reported and the signaling mechanism by which IL-17A regulates the expression of cytotoxicity mediators is not currently understood. In addition, diverse cell types including immune cells, astrocytes, and microglia can produce IL-17A and express IL-17A receptor. Future studies are required for characterization of IL-17A producing and responding cells in both CNS and periphery during various stage of WNV infection. In addition, a better understanding of the regulation of IL-17A production and dissection of the signaling mechanism by which IL-17A induces expression of cytotoxicity mediators and promotes CD8⁺ T cell cytotoxicity in the context of WNV and other diseases may help to exploit novel IL-17A-based therapeutic strategies.

APPENDIX A IACUC and IRB Approval Letter

THE UNIVERSITY OF SOUTHERN MISSISSIPPI.

INSTITUTIONAL ANIMAL CARE AND USE COMMITTEE

118 College Drive #5116 | Hattiesburg, MS 39406-0001 Phone: 601.266.6791 | Fax: 601.266.4377 | iacuc@usm.edu | www.usm.edu/iacuc

NOTICE OF COMMITTEE ACTION

The proposal noted below was reviewed and approved by The University of Southern Mississippi Institutional Animal Care and Use Committee (IACUC) in accordance with regulations by the United States Department of Agriculture and the Public Health Service Office of Laboratory Animal Welfare. The project expiration date is noted below. If for some reason the project is not completed by the end of the approval period, your protocol must be reactivated (a new protocol must be submitted and approved) before further work involving the use of animals can be done.

Any significant changes should be brought to the attention of the committee at the earliest possible time. If you should have any questions, please contact me.

PROTOCOL NUMBER:15101601 (ReplacesPROJECT TITLE:ImmunotherapeuticPROPOSED PROJECT DATES:10/2015 - 09/2018PROJECT TYPE:RenewalPRINCIPAL INVESTIGATOR(S):Fengwei BaiDEPARTMENT:Biological SciencesFUNDING AGENCY/SPONSOR:Wilson Research ForIACUC COMMITTEE ACTION:Full Committee AppPROTOCOL EXPIRATON DATE:September 30, 2018

15101601 (Replaces 12041201) Immunotherapeutics Against Flavivirus es and Alphaviruses 10/2015 - 09/2018 Renewal Fengwei Bai Biological Sciences Wilson Research Foundation Full Committee Approval September 30, 2018

Date

Frank Moore, PhD IACUC Chair

10/01/2015



The University of Southern Mississippi

Institutional Animal Care and Use Committee

118 College Drive #5147 Hattiesburg, MS 39406-0001 Tel: 601.266.6820 Fax: 601.266.5509 www.usm.edu/spa/policies/animals

INSTITUTIONAL ANIMAL CARE AND USE COMMITTEE NOTICE OF COMMITTEE ACTION

The proposal noted below was reviewed and approved by The University of Southern Mississippi Institutional Animal Care and Use Committee (IACUC) in accordance with regulations by the United States Department of Agriculture and the Public Health Service Office of Laboratory Animal Welfare. The project expiration date is noted below. If for some reason the project is not completed by the end of the three year approval period, your protocol must be reactivated (a new protocol must be submitted and approved) before further work involving the use of animals can be done.

Any significant changes (see attached) should be brought to the attention of the committee at the earliest possible time. If you should have any questions, please contact me.

PROTOCOL NUMBER: PROJECT TITLE: PROPOSED PROJECT DATES: PROJECT TYPE: PRINCIPAL INVESTIGATOR(S): DEPARTMENT: FUNDING AGENCY/SPONSOR: IACUC COMMITTEE ACTION: PROTOCOL EXPIRATON DATE: 7/31/2015

12041201 Immunotherapeutics against flaviviruses 08/01/2012-07/31/2015 New Fengwei Bai

Biological Sciences

ie M. Jawor, Ph.D. CUC Chair

10 September 2012 Date



INSTITUTIONAL REVIEW BOARD

118 College Drive #5147 | Hattiesburg, MS 39406-0001 Phone: 601.266.5997 | Fax: 601.266.4377 | www.usm.edu/research/institutional-review-board

NOTICE OF COMMITTEE ACTION

The project has been reviewed by The University of Southern Mississippi Institutional Review Board in accordance with Federal Drug Administration regulations (21 CFR 26, 111), Department of Health and Human Services (45 CFR Part 46), and university guidelines to ensure adherence to the following criteria:

- The risks to subjects are minimized.
- The risks to subjects are reasonable in relation to the anticipated benefits.
- The selection of subjects is equitable.
- Informed consent is adequate and appropriately documented.
- Where appropriate, the research plan makes adequate provisions for monitoring the data collected to ensure the safety of the subjects.
- Where appropriate, there are adequate provisions to protect the privacy of subjects and to maintain the confidentiality of all data.
- Appropriate additional safeguards have been included to protect vulnerable subjects.
- Any unanticipated, serious, or continuing problems encountered regarding risks to subjects must be reported immediately, but not later than 10 days following the event. This should be reported to the IRB Office via the "Adverse Effect Report Form".
- If approved, the maximum period of approval is limited to twelve months.
 Projects that exceed this period must submit an application for renewal or continuation.

PROTOCOL NUMBER: CH-R11120601 PROJECT TITLE: The Role of Neutrophic Virus Pathogenesis PROJECT TYPE: Change to a Previously Approved Project RESEARCHER(S): Fengwei Bai, Ph.D. COLLEGE/DIVISION: College of Science & Technology DEPARTMENT: Biological Sciences FUNDING AGENCY/SPONSOR: N/A IRB COMMITTEE ACTION: Expedited Review Approval PERIOD OF APPROVAL: 04/21/2014 to 04/20/215

Lawrence A. Hosman, Ph.D. Institutional Review Board

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