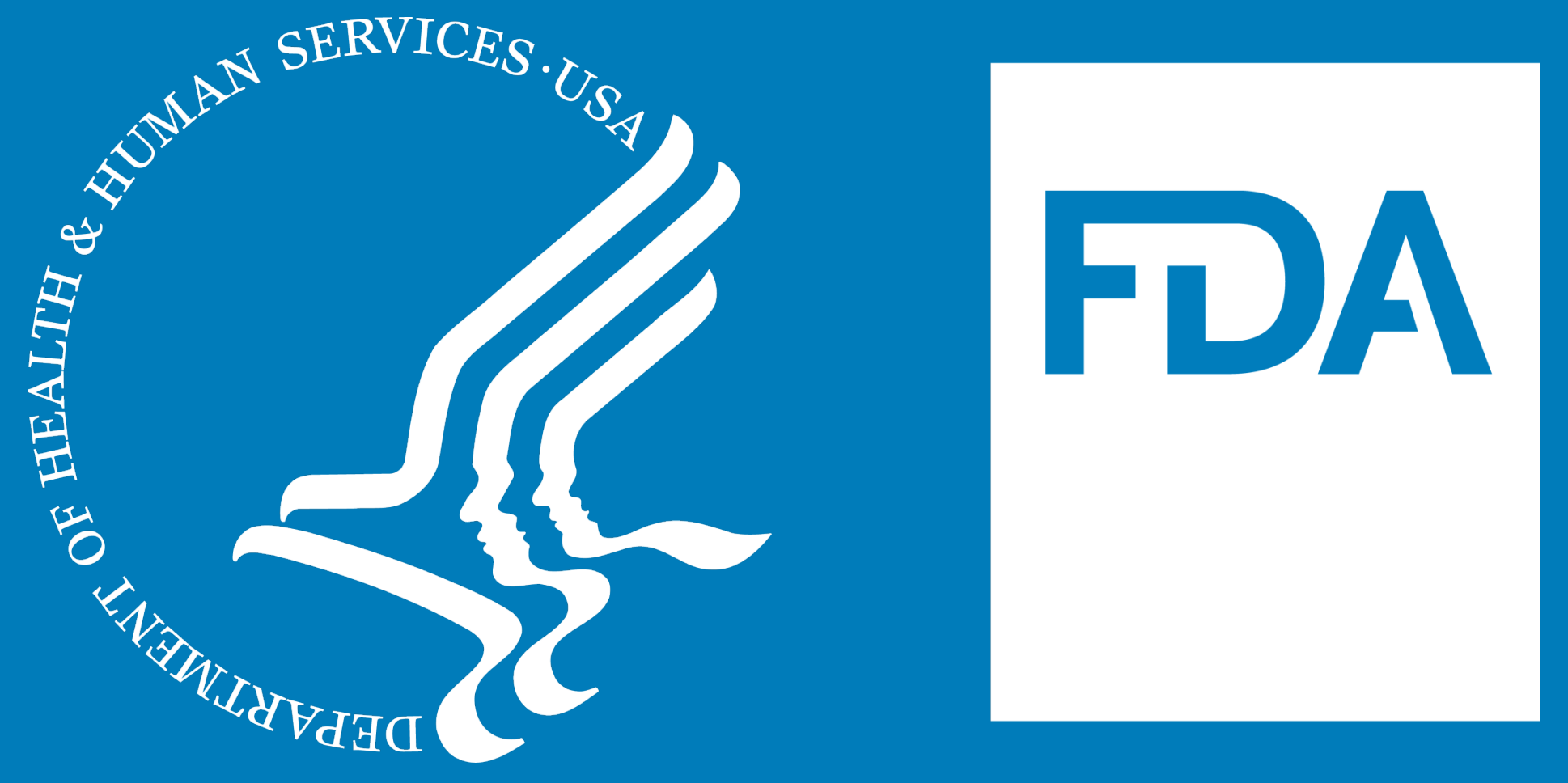


# Development of an immunocompetent mouse model of Dengue virus infection that can be used to test the efficacy of therapeutics and understand Antibody Dependent Enhancement of disease.



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## Abstract

Dengue virus (DENV) infects over 400 million people worldwide every year. While most 1<sup>st</sup> infections are mild, second infections with a heterologous DENV can result in severe disease, encephalitis and death through antibody dependent enhancement (ADE). There are no approved effective therapeutics and no licensed vaccines for dengue virus disease. There are no immunocompetent mouse models available to understand host-pathogen interactions, establish determinants of disease, or explore risk factors for ADE. To address this unmet need, we developed an immunocompetent neonatal mouse model in C57BL/6 mice using DENV serotype 2, New Guinea C strain (DENV2). The DENV2 infected mice fail to thrive, and develop lethargy, ataxia, and tremors around 6-9 dpi and approximately 50% of the infected mice succumbed to disease by 10-12 dpi. The mice have a viremia followed by high levels of virus in the CNS and eyes starting 3 days post infection (DPI). The infection in brains and eyes is associated with significant increase in mRNA expression for interferon-inducible genes (BST2, STAT1&2, IRGM1, IFIT2, IRF7, IFI35), RIG-I-Like receptors (DDX58 and IFIH1), chemokines (CXCL10, CCL5, CCL12, CCL2) antigen presentation and activation (H2-K1, CTSS, TAP1, TAPBP, PTPRC) complement (C4A, C1QB, C2, C1QA) and FC receptors (FCGR1, FCGR4, FCGR2B) as well as a corresponding reduction in genes related to neuronal function (PCP2, PAX2). This model will be used to test the efficacy of anti-DENV therapeutics and understand the determinants of ADE using heterologous infections and antibody transfer studies.

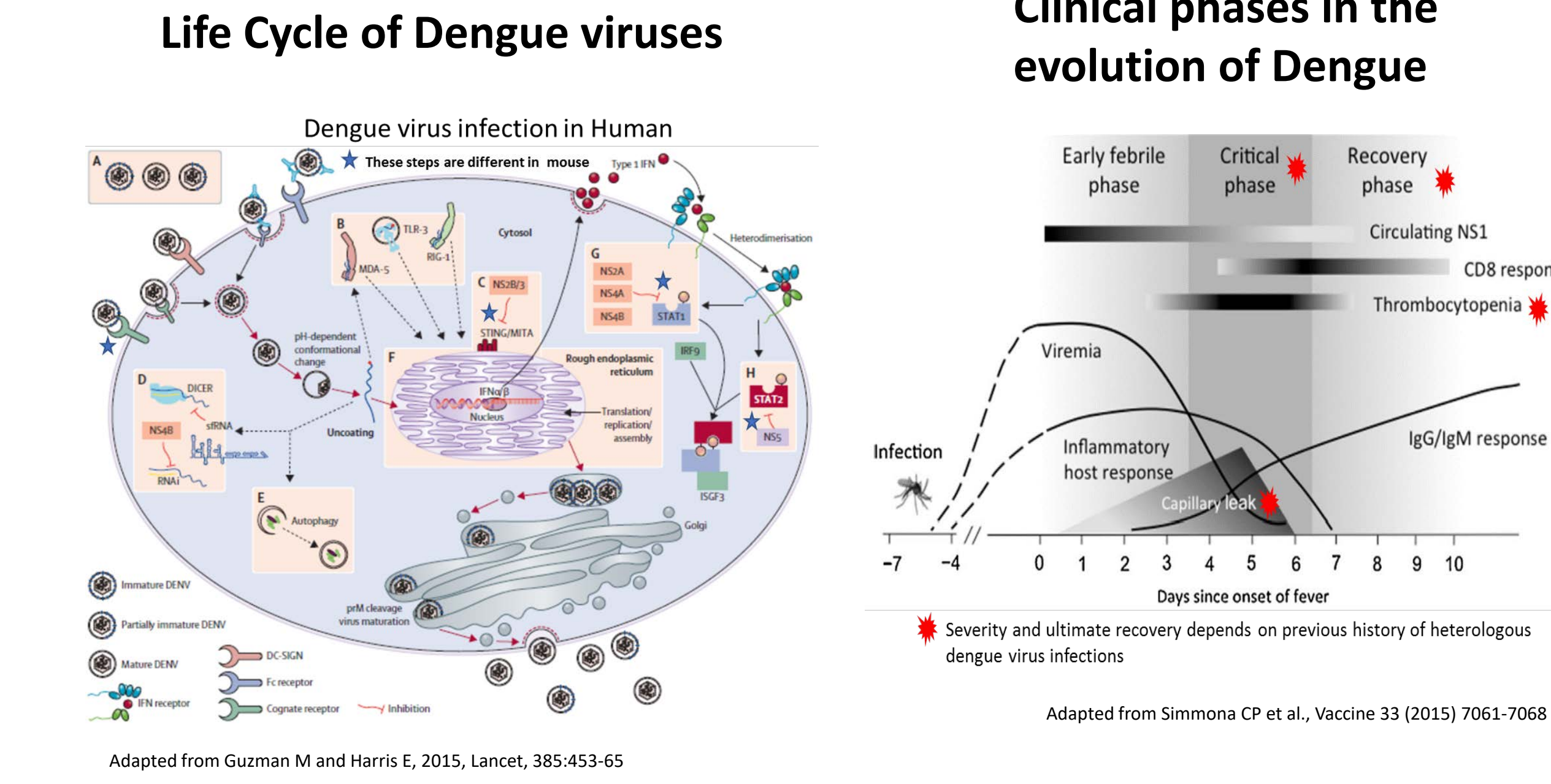
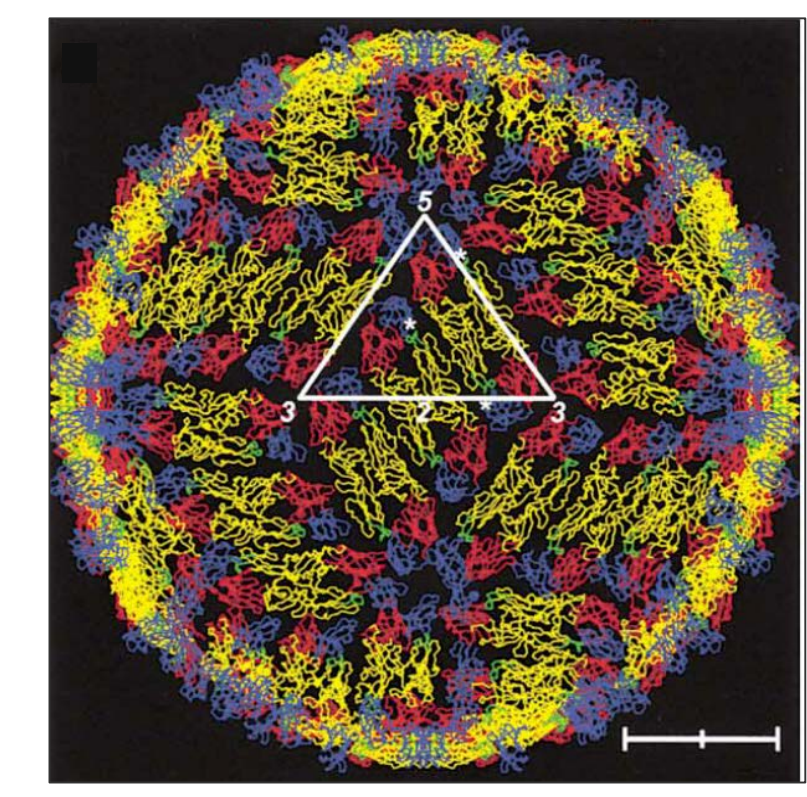
## Introduction

### Biology of Dengue Virus

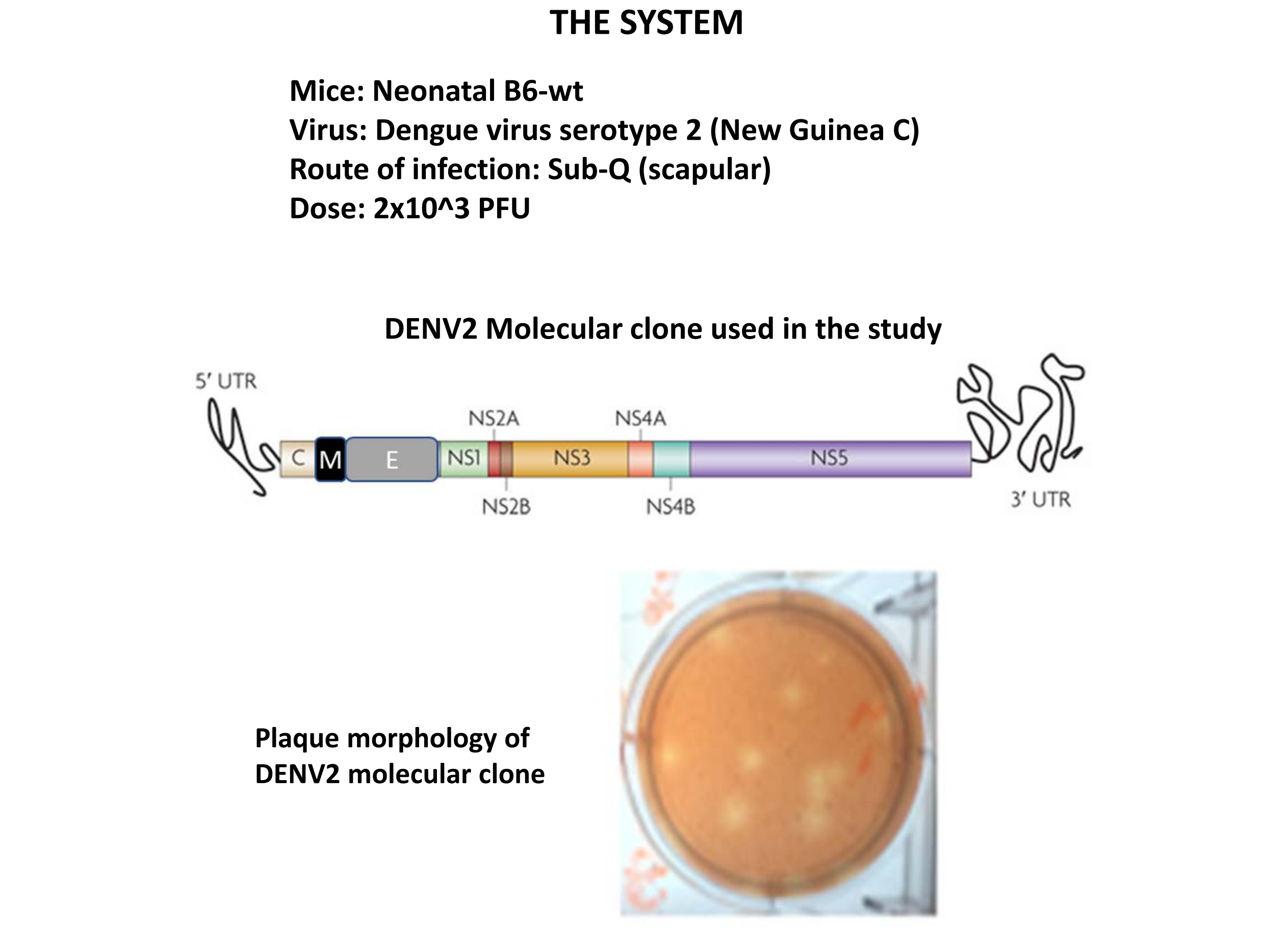
Family: Flaviviridae  
 Genus: Flavivirus  
 Species: Dengue

Dengue virus is presently the most common cause of arboviral disease globally. All four serotypes can be found worldwide. WHO estimates an annual incidence of ~100 million infections, with approximately 500,000 people with Dengue Hemorrhagic fever (DHF).

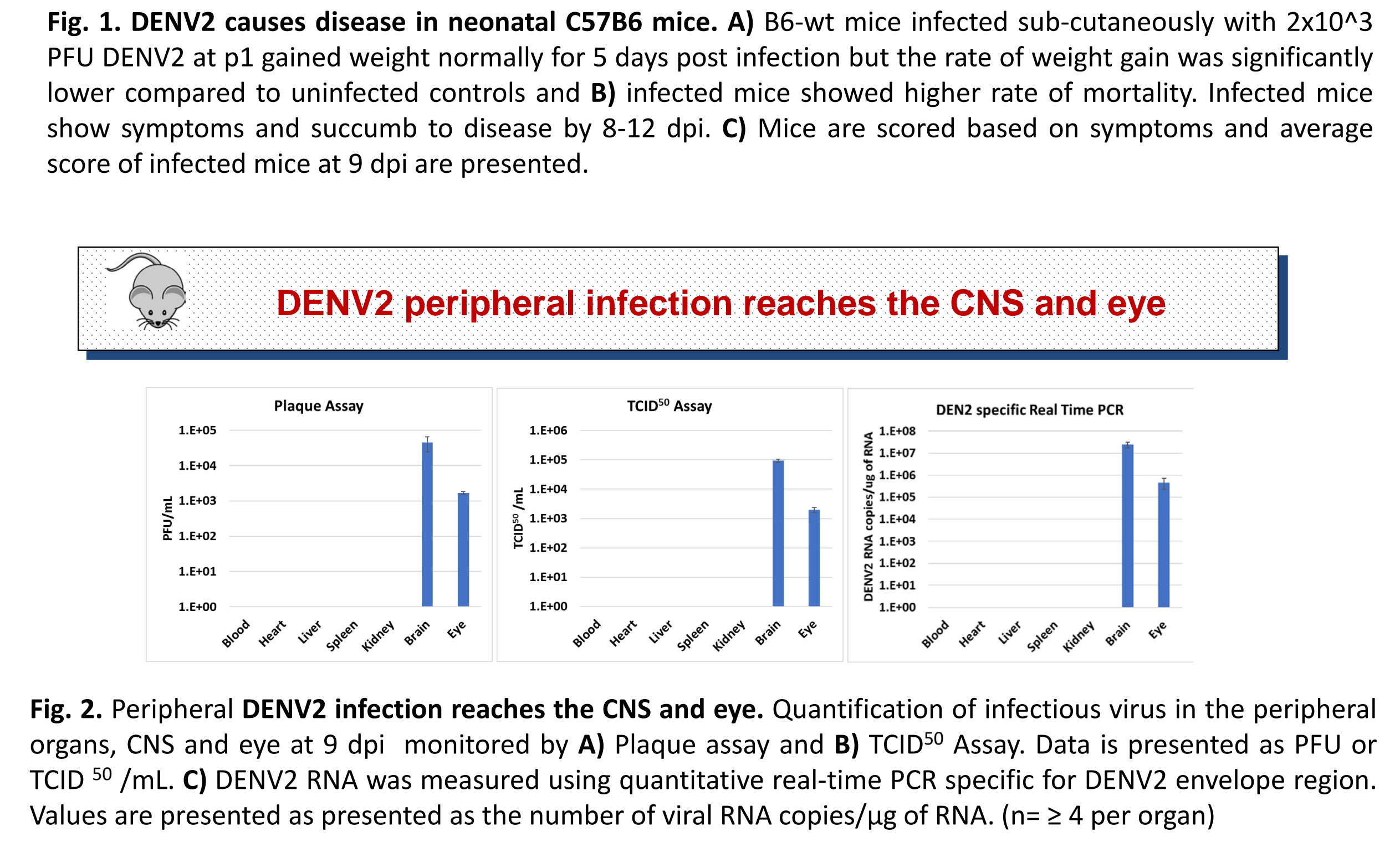
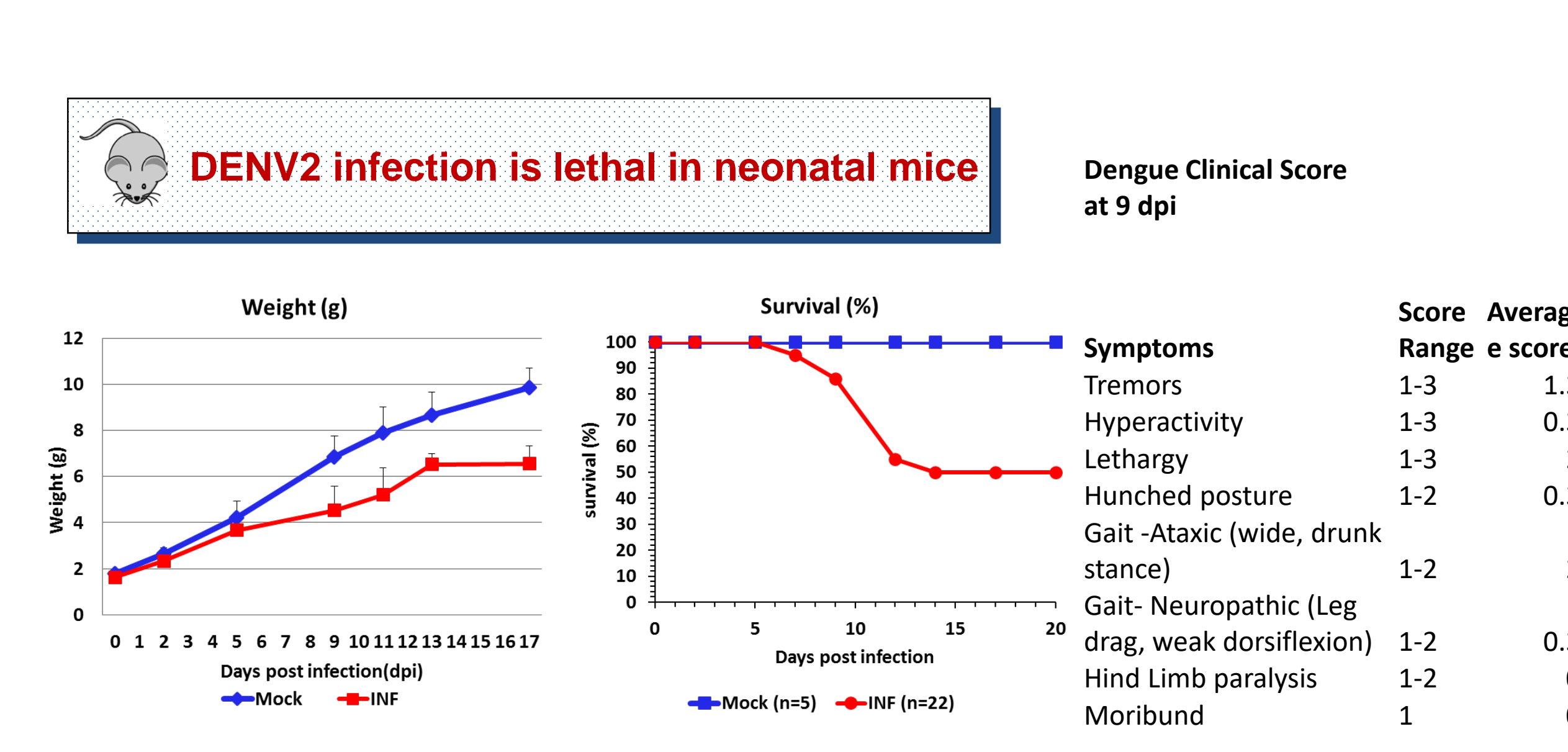
Enveloped virus with single positive sense RNA ~ 10 kb. Four serologically and genetically distinct serotypes Dengue 1-4.



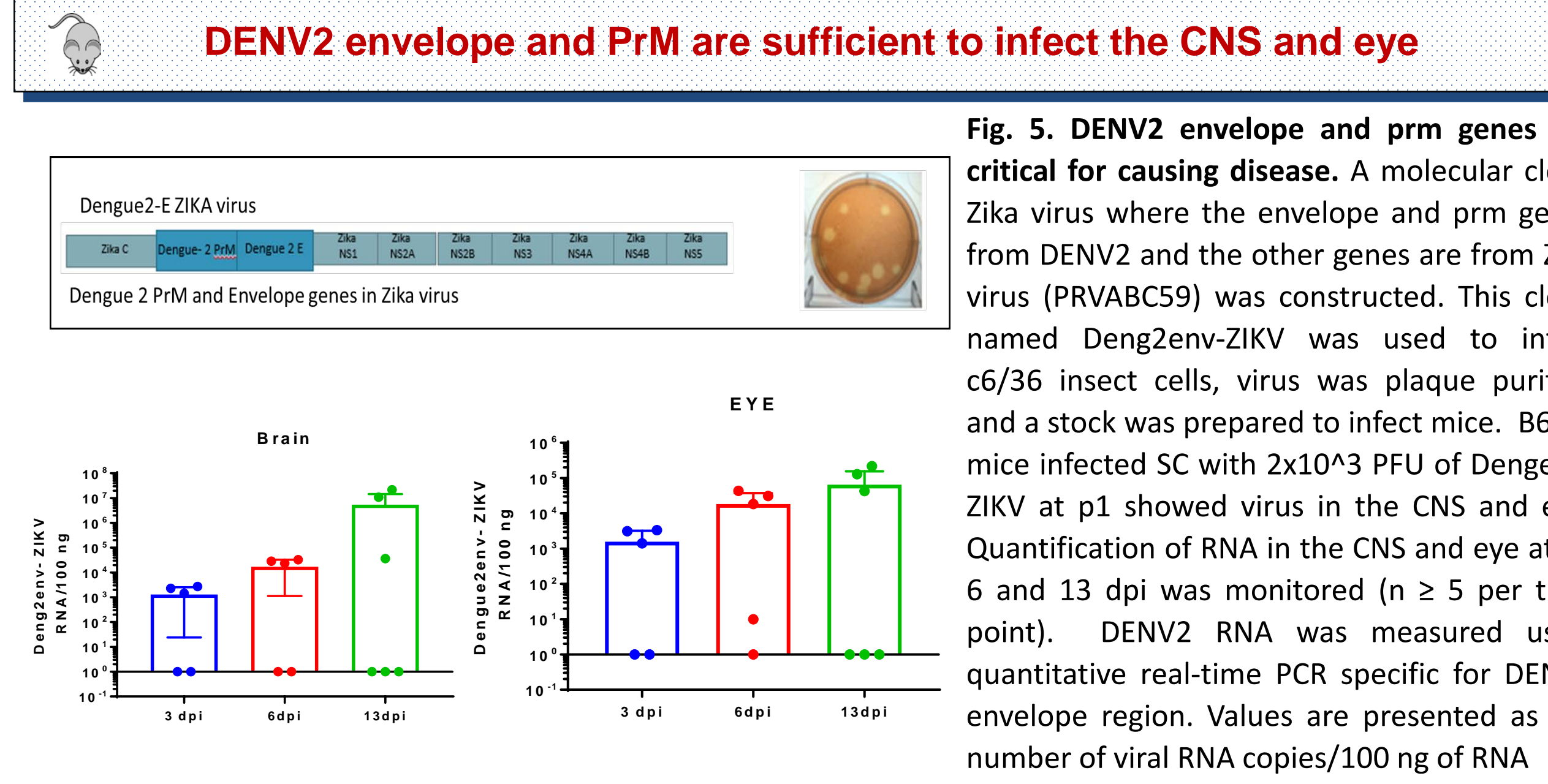
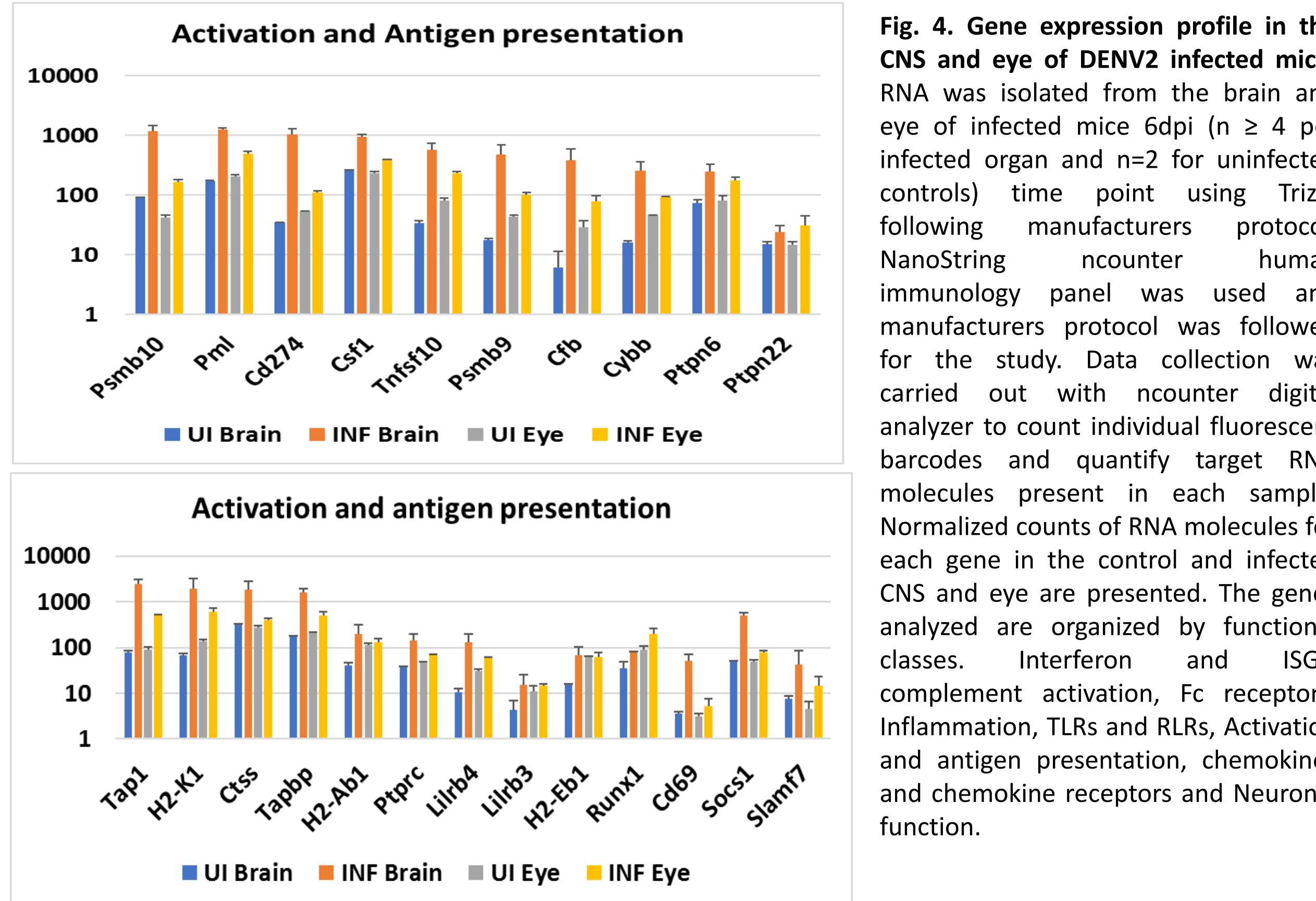
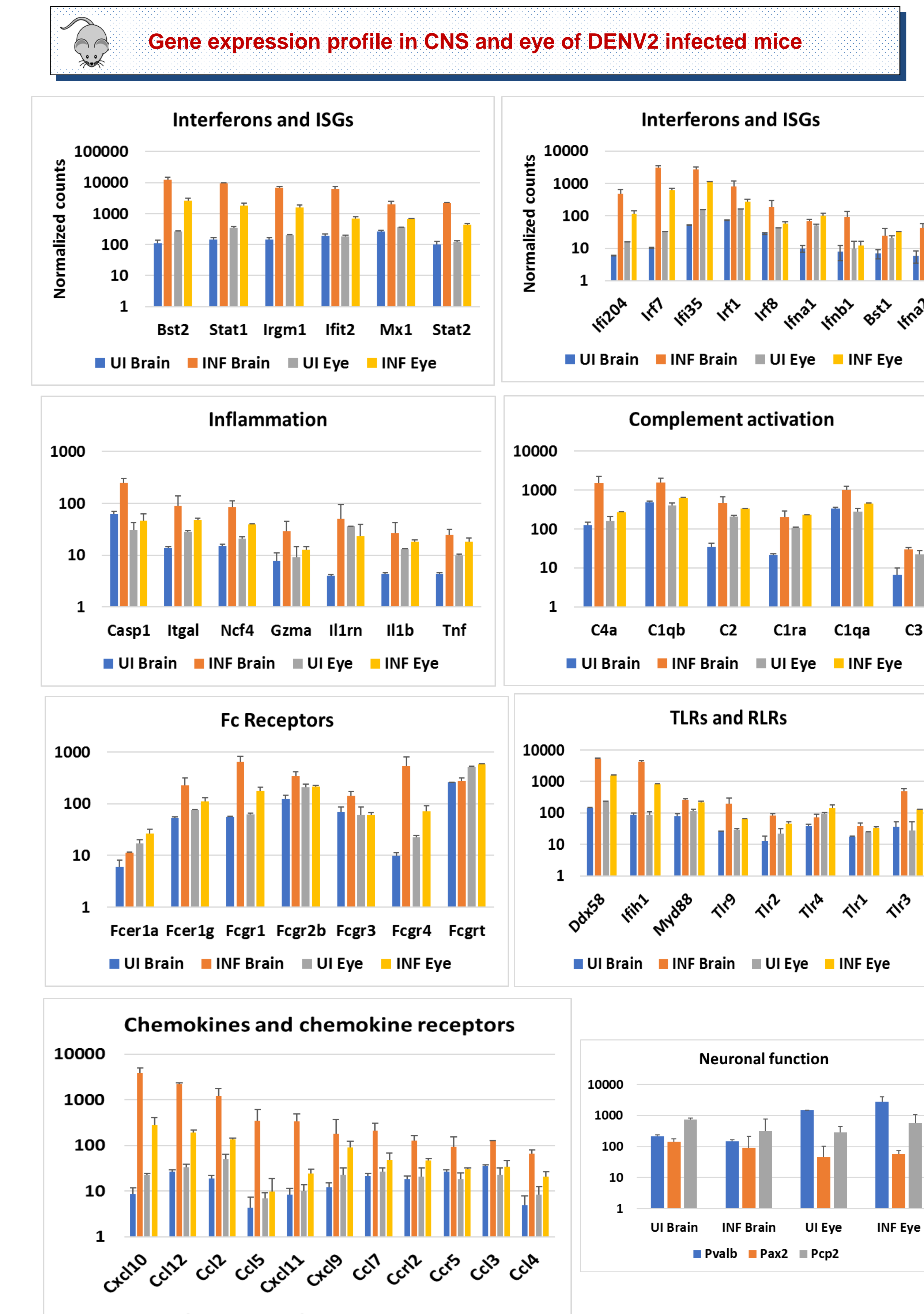
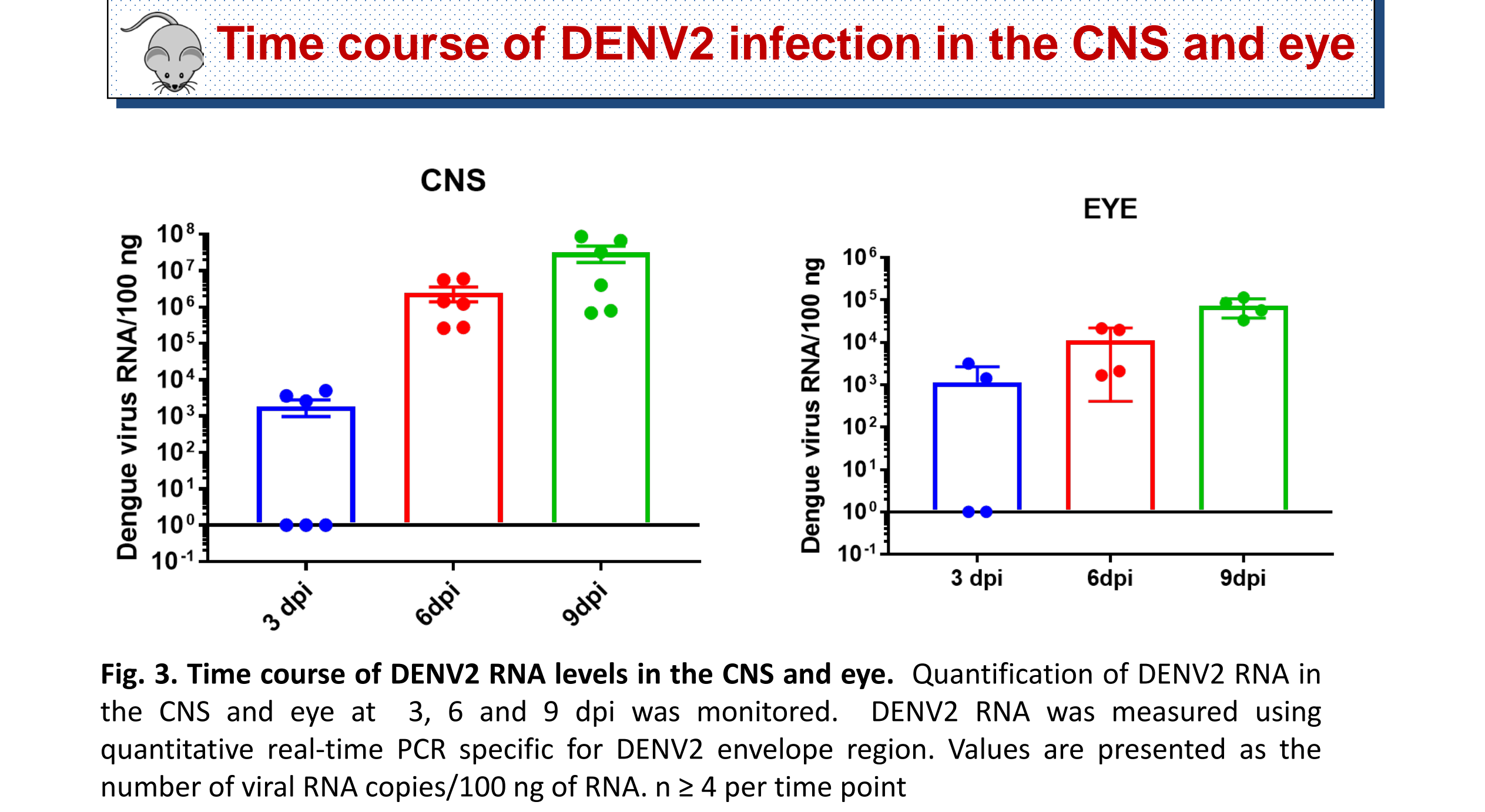
## Materials and Methods



## Results and Discussion



## Results and Discussion



## Conclusion

- ✓ Infection of immunocompetent neonatal B6 wt mice with DENV2 (Dengue virus serotype 2, New Guinea C strain) results in systemic infection, reaches the CNS and eye and results in neurological symptoms including tremors and hyperactivity and results in death in 50% of infected mice.
- ✓ 2. DENV2 RNA and infectious virus can be transiently detected in the periphery but persistent in the CNS and eye even after clearance from the periphery.
- ✓ 3. Gene expression in the CNS and eye of infected mice show inflammation and innate immune response that correlates with the level of infection. Some infected mice that do not show DENV2 RNA at 6 and 9dpi still show signs of inflammation indicating past or ongoing infection.
- ✓ 4. Infection of neonatal mice with DengZenv-ZIKV show that envelope and prM genes are critical for the tropism exhibited in this mouse model of DENV2 virus disease.
- ✓ 5. This mouse model will be used to understand Antibody Dependent Enhancement of dengue virus disease.

**FDA Mission Relevance Statement**

Dengue virus mouse model that we have developed will help product developers and reviewers to test the efficacy of immunomodulatory and anti-DENV therapeutics and to improve our understanding of the determinants of increased disease severity following a heterologous DENV infection or vaccine.

**Thanks**

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