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Nucleotide at position 66 of NS2A in Japanese encephalitis virus is associated with the virulence and proliferation of virus

Original Paper | Published: 08 November 2023

(2023) [Cite this article](#)



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Abstract

Most wild strains of Japanese encephalitis virus (JEV) produce NS1' protein, which plays an important role in viral infection and immune escape. The G66A nucleotide mutation in NS2A gene of the wild strain SA14 prevented the ribosomal frameshift that prevented the production of NS1' protein, thus reduced the virulence. In this study, the 66th nucleotide of the NS2A gene of SA14 was mutated into A, U or C, respectively. Both the G66U and G66C mutations cause the E22D mutation of the NS2A protein. Subsequently, the expression of NS1' protein, plaque size, replication ability, and virulence to mice of the three mutant strains were examined. The results showed that the three mutant viruses could not express NS1' protein, and their proliferation ability in nerve cells and virulence to mice were significantly reduced. In addition, the SA14(G66C)

was less virulent than the other two mutated viruses. Our results indicate that only when G is the 66th nucleotide of NS2A, the JEV can produce NS1' protein, which affects the virulence.

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Data availability

The authors confirm that the data supporting the findings of this study are available within the article.

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Funding

This work was supported by the Research fund of North Sichuan Medical College (No. CBY22-ZDA06).

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Contributions

LY and NT designed the study, analyzed the data and wrote the manuscript. CC, YR and RH cultured viruses, collected and tested the samples and analyzed the data. ZZ, KX and XY carried out the animal experiment. JY and LY reviewed and edited the manuscript. All authors approved the final manuscript.

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Ethics declarations

Conflicts of interest

The authors declared that they have no conflicts of interest to this work.

Ethical approval

All procedures performed in this study involving animals were approved by the Institutional Animal Care and Use Committee of North Sichuan Medical College (approval number AE2022-068), Nanchong city, China and followed National Institutes of Health guidelines.

Additional information

Edited by Sassan Asgari.

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Cite this article

Tan, N., Chen, C., Ren, Y. *et al.* Nucleotide at position 66 of NS2A in Japanese encephalitis virus is associated with the virulence and proliferation of virus. *Virus Genes* (2023).

<https://doi.org/10.1007/s11262-023-02036-5>

Received

09 August 2023

Accepted

10 October 2023

Published

08 November 2023

DOI

<https://doi.org/10.1007/s11262-023-02036-5>

Keywords

[Japanese encephalitis virus](#)

[Ribosomal frameshift](#)

[NS2A gene](#)

[NS1' protein](#)

[Virulence](#)