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# Characterisation of a Japanese Encephalitis virus genotype 4 isolate from the 2022 Australian outbreak

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Wilson Nguyen<sup>1</sup>, Narayan Gyawali<sup>1</sup>, Romal Stewart<sup>1</sup>, Bing Tang<sup>1</sup>, Abigail L. Cox<sup>1</sup>, Kexin Yan<sup>1</sup>, Thibaut Larcher<sup>2</sup>, Cameron R. Bishop<sup>1</sup>, Nicholas Wood<sup>3</sup>, Gregor J. Devine<sup>1,4</sup>, Andreas Suhrbier<sup>1,4,5</sup> & Daniel J. Rawle<sup>1,5</sup>✉

Human infections with the Japanese encephalitis virus (JEV) are a leading cause of viral encephalitis. An unprecedented outbreak of JEV genotype 4 was recently reported in Australia, with an isolate (JEV<sub>NSW/22</sub>) obtained from a stillborn piglet brain. Herein we conduct a thorough characterization of JEV<sub>NSW/22</sub> in three different mouse strains and in human cortical brain organoids (hBOs), and determined the ability of JEV<sub>NSW/22</sub> to be neutralized by sera from humans vaccinated with IMOJEV. JEV<sub>NSW/22</sub> was less virulent than JEV<sub>FU</sub> (genotype 2) and JEV<sub>Nakayama</sub> (genotype 3) in C57BL/6J mice and in interferon regulatory factor 7 deficient (*Irf7*<sup>-/-</sup>) mice, with infection of wild-type and knockout murine embryonic fibroblasts indicating JEV<sub>NSW/22</sub> is more sensitive to type I interferon responses. *Irf7*<sup>-/-</sup> mice provide a new model for JEV<sub>NSW/22</sub>, showing higher viremia levels compared to C57BL/6J mice, and allowing for lethal neuroinvasive infection. All JEV strains were universally lethal in *Ifnar*<sup>-/-</sup> mice by day 3, with histological signs of brain hemorrhage, but no other lesions. There were no indications of brain infection in *Ifnar*<sup>-/-</sup> mice, with viral protein detected in blood vessels, but not neurons. All JEV isolates showed robust cytopathic infection of human cortical brain organoids, albeit lower for JEV<sub>NSW/22</sub>. IMOJEV vaccination in humans induced antibodies capable of neutralizing JEV<sub>NSW/22</sub>, although, for all JEV strains, cross-neutralization titers declined with increasing divergence from IMOJEV in the envelope amino acid sequences. Overall, our study establishes JEV<sub>NSW/22</sub> mouse and hBO models of infection, allowing for possible lethal neuroinvasive infection in mice that was rarer than for other JEV genotypes. JEV vaccination regimens may afford protection against this newly emerged JEV genotype 4 strain, although neutralizing antibody responses are sub-optimal.

Japanese encephalitis virus (JEV) is a single-stranded positive-sense RNA virus from the *Flaviviridae* family that is transmitted from amplifying hosts (primarily pigs and wading birds) via mosquitoes to humans<sup>1</sup>. While the infection is usually asymptomatic, encephalitis can develop in ≈1 in 250 people, with ~30% of encephalitic cases becoming fatal, and 30–50% of non-fatal encephalitic cases retaining persistent neurological symptoms including seizures, speech impediments, and paralysis<sup>2,3</sup>. JEV is the leading cause of viral encephalitis in Asia, with ~70,000 cases and ~20,000 deaths per annum<sup>4</sup>. After a bite from an infected mosquito, JEV replicates in peripheral blood monocytes producing a viremia that, in some cases, leads to the virus

crossing the blood-brain barrier<sup>5</sup>. JEV primarily infects neurons in the brain<sup>6</sup> leading to uncontrolled inflammation (encephalitis) and neuronal cell death<sup>2</sup>. There are several available and effective vaccines against JEV<sup>7</sup>, but there are no specific licensed treatments.

JEV exists as five genotypes (1–5) which are phylogenetically, antigenically, and geographically distinct<sup>8–10</sup>. Genotype 3 was the main genotype endemic in Asia until 1990, after which genotype 1 have dominated since<sup>11</sup>. Genotype 2 has been identified in Malaysia, Indonesia, Papua New Guinea and Australian outbreaks from 1970–2000. Genotype 5 was originally isolated in Malaysia in 1952, and has since been identified in China, and is now

<sup>1</sup>QIMR Berghofer Medical Research Institute, Brisbane, QLD 4029, Australia. <sup>2</sup>INRAE, Oniris, PANter, APEX, Nantes, France. <sup>3</sup>National Centre for Immunisation Research and Surveillance, Westmead, NSW, Australia. <sup>4</sup>GVN Center of Excellence, Australian Infectious Disease Research Centre, Brisbane, QLD 4029 and 4072, Australia. <sup>5</sup>These authors jointly supervised this work: Andreas Suhrbier, Daniel J. Rawle. ✉e-mail: [Daniel.Rawle@qimrberghofer.edu.au](mailto:Daniel.Rawle@qimrberghofer.edu.au)

dominant in South Korea<sup>12</sup>. Genotype 4 was the least common genotype worldwide, having only been identified in mosquitoes from Indonesia and Papua New Guinea<sup>13</sup>. In February 2021, a fatal JEV infection occurred on the Tiwi Islands, 80 km off the coast of Darwin, Australia<sup>14</sup>. Sequencing identified that the virus belonged to the historically rare JEV genotype 4<sup>14</sup>. In 2022, a geographically widespread outbreak throughout most Australian states was attributed to JEV genotype 4, which caused 44 confirmed human cases and 7 deaths<sup>15</sup>. Outbreaks occurred in several piggeries, causing high abortion and stillbirth rates in sows<sup>16</sup>. Based on proximity to piggeries, estimates indicate that ~740,000 people may be at risk of being infected by JEV in Australia<sup>16</sup>. Given that *Culex annulirostris* mosquitoes, considered the primary vector for JEV in Australia<sup>15,16</sup>, and amplifying vertebrate hosts, including wading birds and pigs, are widespread, it is possible that JEV may become, or is already, endemic in Australia<sup>15</sup>. Murray Valley Encephalitis virus (MVEV) and Kunjin virus are phylogenetically closely related to JEV, have similar vector and reservoir hosts, and are endemic to Australia<sup>17</sup>.

Two JEV vaccines are licensed for use in Australia; IMOJEV (live attenuated) and JESpect (formalin inactivated). Before the 2022 outbreak, JEV vaccination primarily served as a travel vaccine for those traveling to Southeast Asian countries<sup>18</sup>. In response to the outbreak, a priority vaccination program was promptly instituted for people at higher risk of exposure. This included piggery workers, laboratory scientists handling JEV, mosquito surveillance scientists, and those residing in or frequently visiting high-risk locations<sup>18</sup>. Both IMOJEV and JESpect vaccines are designed against a JEV genotype 3 antigen, and there have been no studies assessing their efficacy against the genotype 4 strain responsible for the Australian outbreak.

Efforts to develop treatments for JEV neuropathology are hindered by the difficulties in diagnosing early infection<sup>13</sup> before the virus has infected central nervous system (CNS) neurons; and once the virus has infected CNS neurons, treatments must also be able to effectively enter the brain. While a range of JEV mouse models have been reported (Supplementary Table 1)<sup>19</sup>, there are no studies of genotype 4 peripheral inoculation in adult mice, with the only studies on genotype 4 JEV using 3-week-old mice<sup>20</sup> or intracranial inoculation<sup>21</sup>. Well-characterized, adult mouse models of recent genotype 4 isolates would represent useful tools to study mechanisms of infection and disease and evaluate potential new interventions. Herein, we compare and contrast a JEV genotype 4 isolate from the recent Australian outbreak (JEV<sub>NSW/22</sub>) with historical JEV isolates (genotypes 2 and 3) in three mouse strains, human cortical brain organoids and *in vitro* neutralization assays with human serum after JEV vaccination.

## Materials and methods

### Ethics statement and regulatory compliance

All mouse work was conducted in accordance with the “Australian code for the care and use of animals for scientific purposes” as defined by the National Health and Medical Research Council of Australia. Mouse work was approved by the QIMR Berghofer Medical Research Institute (MRI) Animal Ethics Committee (P3746, A2108-612). Mice were euthanized using CO<sub>2</sub>.

Breeding and use of GM mice were approved under a Notifiable Low Risk Dealing (NLRD) Identifier: NLRD\_Suhrbier\_Oct2020: NLRD 1.1(a). The use of IMOJEV was approved under NLRD\_Suhrbier\_Oct2019: NLRD2.1(d).

All infectious JEV work was conducted in a dedicated suite in a bio-safety level-3 (PC3) facility at the QIMR Berghofer MRI (Australian Department of Agriculture, Water and the Environment certification Q2326 and Office of the Gene Technology Regulator certification 3445). All work was approved by the QIMR Berghofer MRI Safety Committee (P3746).

Human serum samples before and after IMOJEV vaccination were collected from 9 participants with human research ethics approval from Sydney Children’s Hospitals Network Human Research Ethics Committee (2022/ETH02471 HCHNHREC) for an ongoing project led by Dr. Nicholas

Wood; “Japanese encephalitis vaccine via the intradermal route in children and adults (JEVID-2): Critical policy-relevant research for Australia”.

Human serum samples after IMOJEV vaccination were also collected from 10 participants with approval from QIMR Berghofer MRI Human Research Ethics Committee (P3476) provided for neutralization assays (including MVEV cross-neutralization).

Research with JEV was approved under the Queensland Biosecurity Act, Scientific Research Permit (restricted matter) – Permit number PRID000916.

### Cell lines and culture

Vero E6 (C1008, ECACC, Wiltshire, England; obtained via Sigma-Aldrich, St. Louis, MO, USA), BHK-21 (ATCC# CCL-10), and C6/36 cells (ATCC# CRL-1660) were cultured in medium comprising RPMI 1640 (Gibco), supplemented with 10% fetal bovine serum (FBS), penicillin (100 IU/ml)/streptomycin (100 µg/ml) (Gibco/Life Technologies) and L-glutamine (2 mM) (Life Technologies). Vero E6 and BHK cells were cultured at 37 °C and 5% CO<sub>2</sub>, and C6/36 cells were cultured at 27 °C and 5% CO<sub>2</sub>. Cells were routinely checked for mycoplasma (MycoAlert Mycoplasma Detection Kit MycoAlert, Lonza), and FBS was assayed for endotoxin contamination before purchase<sup>22</sup>.

Mouse embryonic fibroblasts (MEFs) were either wild-type or *Irf3*<sup>-/-</sup> and have been described previously<sup>23</sup>. MEFs were cultured in Dulbecco’s Modified Eagle Medium (DMEM) (Gibco), supplemented with 50 µg/ml Penicillin/50 µg/ml Streptomycin and 10% Fetal Bovine Serum (FBS). Cells were cultured at 37 °C and 5% CO<sub>2</sub>. For virus growth kinetics, MEFs were seeded at 2 × 10<sup>5</sup> cells/ml in 12 or 24 well plates one day prior to infection at multiplicity of infection (MOI) 0.1 of the indicated JEV or MVEV. After 1 hr incubation, MEFs were washed twice with 1 ml PBS, and culture medium was added and sampled daily for virus titrations by CCID<sub>50</sub> (see below).

RENcell VM Human Neural Progenitor Cell Line (Sigma-Aldrich) were cultured in medium comprising DMEM F-12 (Thermo Fisher Scientific), penicillin (100 IU/ml)/streptomycin (100 µg/ml) (Gibco/Life Technologies), 20 ng/ml FGF (STEMCELL Technologies), 20 ng/ml EGF (STEMCELL Technologies), and 2% B27 supplements (Thermo Fisher Scientific). Cells were detached using StemPro Accutase Cell Dissociation Reagent (Thermo Fisher Scientific), and were cultured in Matrigel (Sigma). For crystal violet staining of remaining cells after infection, formaldehyde (7.5% w/v)/crystal violet (0.05% w/v) was added to wells overnight, plates washed twice in water, and plates dried overnight.

### Virus isolates and culture

JEV<sub>Nakayama</sub> (GenBank: EF571853), JEV<sub>FU</sub> (GenBank: AF217620), and IMOJEV were obtained from A/Prof. Gregor Devine (QIMR Berghofer MRI). JEV<sub>NSW/22</sub> (GenBank: OP904182) was obtained from an infected porcine neonate by Dr. Peter Kirkland (Elizabeth Macarthur Agriculture Institute, New South Wales, Australia). MVEV<sub>TC123130</sub> (GenBank: JN119814.1) was obtained from the “Doherty Virus Collection” currently held at QIMR Berghofer MRI. Virus stocks were generated by infection of C6/36 cells (for JEV and MVEV) or Vero E6 cells (for IMOJEV and YFV 17D) at MOI ≈ 0.1, with supernatant collected after ~5 days, cell debris removed by centrifugation at 3000 × g for 15 min at 4 °C, and virus aliquoted and stored at –80 °C. Virus stocks used in these experiments underwent less than three passages in our laboratory, with prior passage history in Supplementary Fig. 1. Virus titers were determined using standard CCID<sub>50</sub> assays (see below).

**Validation of virus stock sequences.** Viral RNA was extracted from virus stock culture supernatants using NucleoSpin RNA Virus kit (Machery Nagel) as per manufacturer’s instructions. cDNA was synthesized using ProtoScript First Strand cDNA Synthesis Kit (New England Biolabs) as per manufacturer’s instructions. PCR was performed using Q5 High-Fidelity 2X Master Mix (New England Biolabs) as per manufacturer’s instructions with the following primers; JEV envelope Forward 5’ GGAAGCATTGACACATGTGC 3’ and Reverse 5’

TCTGTGCACATACCATAGGTTGTG 3', and MVEV envelope Forward 5' GAGCATTGACACATGCGCAAAG and Reverse 5' TGTGCACATCCCATAAGTGGTTC 3'. PCR products were run on a 1% agarose gel, and DNA was extracted using Monarch DNA Gel Extraction Kit (New England Biolabs). DNA was sequenced by Sanger sequencing using either the forward or reverse primer. Sequences of our virus stocks matched the sequences on GenBank (Supplementary Fig. 1).

### Cell culture infectious dose 50% (CCID<sub>50</sub>) assays

CCID<sub>50</sub> assays were performed as previously described<sup>24–26</sup>. C6/36 cells were plated in 96 well flat bottom plates at  $2 \times 10^4$  cells per well in 100  $\mu$ l of medium. For tissue titrations, tissues were homogenized in tubes each containing 4 ceramic beads twice at  $6000 \times g$  for 15 seconds, followed by centrifugation twice at  $21,000 \times g$  for 5 min. Samples underwent 10-fold serial dilutions in 100  $\mu$ l RPMI 1640 supplemented with 2% FBS, performed in quadruplicate for tissue homogenate and in duplicate for serum and cell culture supernatant. For mouse tissues and serum, a volume of 100  $\mu$ l of serially diluted samples was added to each well of 96 well plates containing C6/36 cells, and the plates were cultured for 5 days at 37 °C and 5% CO<sub>2</sub>. 25  $\mu$ l of supernatant from infected C6/36 cells were then passaged on to Vero E6 cells plated the day before at  $2 \times 10^4$  cells per well in 96 well flat bottom plates. Supernatants from in vitro cultures were titered directly onto Vero E6 cells. Vero E6 cells were cultured for 5 days cytopathic effect was scored, and the virus titer was calculated using the method of Spearman and Karber<sup>27</sup> (a convenient Excel CCID50 calculator is available at <https://www.klinikum.uni-heidelberg.de/zentrum-fuer-infektiologie/molekulare-virology/welcome/downloads>).

### Mouse infections

C57BL/6J mice were purchased from the Animal Resources Center, Canning Vale WA, Australia. *Ifnar*<sup>-/-</sup> were kindly provided by Dr P. Hertzog (Monash University, Melbourne, Australia). *Irf7*<sup>-/-</sup> mice were kindly provided by T. Taniguchi (University of Tokyo)<sup>28–30</sup>. C57BL/6N mice were purchased from The Jackson Laboratory (stock no. 005304). C57BL/6NJ<sup>ANnt8-12</sup> were generated by The Australian Phenomics Network, Monash University, Melbourne, Australia, as described<sup>31</sup>. Mice used in this study were female, except *Irf7*<sup>-/-</sup> mice where both males and females were used. The age/age range at infection is indicated in the figure legends. Mice were sorted into groups so that each group had a similar mean age and age distribution, and in the case of *Irf7*<sup>-/-</sup> mice, had equal numbers of males and females in each group. All mice strains except C57BL/6J were bred in-house at QIMR Berghofer MRI, and mice housing conditions were described previously<sup>32</sup>. Mice were infected subcutaneously (s.c.) at the base of the tail with 100  $\mu$ l of virus inoculum (doses ranging from  $5 \times 10^2$  to  $5 \times 10^5$  as indicated in the figure legends). Serum was collected via tail nicks into Microvette Serum-Gel 500  $\mu$ l blood collection tubes (Sarstedt, Numbrecht, Germany). Mice were weighed and monitored for disease manifestations and were humanely euthanized using CO<sub>2</sub> based on a scorecard system (Supplementary Fig. 2B). At necropsy, brain, and spleens were collected for virus titrations by CCID<sub>50</sub> assays and/or for histology.

### Histopathology and immunohistochemistry

Brains and spleens were fixed in 10% formalin and embedded in paraffin. Human cortical brain organoids were embedded in agarose by adding 4% melted agarose and incubating on ice to solidify, prior to standard paraffin embedding. Sections were stained with H&E (Sigma-Aldrich), and slides were scanned using Aperio AT Turbo (Aperio, Vista, CA USA) and analyzed using Aperio ImageScope software (LeicaBiosystems, Mt Waverley, Australia) (v10). Leukocyte infiltrates were quantified by measuring nuclear (strong purple staining)/cytoplasmic (total red staining) pixel ratios in scanned H&E stained images, and was undertaken using Aperio Positive Pixel Count Algorithm (Leica Biosystems)<sup>33</sup>.

For anti-flavivirus non-structural protein 1 (NS1) immunohistochemistry using 4G4, sections were affixed to positively charged adhesive slides and air-dried overnight at 37 °C. Sections were dewaxed and

rehydrated through xylol and descending graded alcohols to water. Sections were transferred to Dako Epitope Retrieval Solution (pH 9.0) and subjected to heat antigen retrieval (100 °C for 20 min) using the Biocare Medical de-cloaking chamber, and slides allowed to cool for 20 minutes before transferring to TBS plus 0.025% Tween-20 (TBS-T). Endogenous mouse Ig was blocked by incubating sections with donkey anti-mouse Fab fragments (Jackson ImmunoResearch) diluted 1:50 in Biocare Medical Rodent block M for 60 minutes. Sections were washed three times in TBS-T, then incubated with anti-mouse Fc for 15 minutes, before a further three washes in TBS-T. Nonspecific antibody binding was inhibited by incubation with Biocare Medical Background Sniper with 1% BSA 20% donkey serum, and 20% goat serum for 15 minutes. Primary antibody 4G4 (mouse anti-flavivirus NS1<sup>25,34,35</sup>) was diluted 1 in 4 with Da Vinci Green diluent and applied to the sections overnight at room temperature. Sections were washed three times in TBS-T, and endogenous peroxidase was blocked by incubating slides in Biocare Medical Peroxidase 1 for 5 minutes. Sections were washed three times in TBS-T, and Perkin Elmer Opal HRP Polymer or Perkin Elmer Goat anti-mouse HRP diluted 1:500 in TBS-T was applied for 60 minutes. Sections were washed three times in TBS-T, and signal developed in Vector Nova Red for 5 minutes, after which they were washed three times in water. Sections were lightly counterstained in Haematoxylin (program 7 Leica Autostainer), washed in water, dehydrated through ascending graded alcohols, cleared in xylene, and mounted using DePeX or similar.

Apoptosis was detected using the ApopTag Peroxidase In Situ Apoptosis Detection Kit (Merck Catalog No. S7100) as per the manufacturer's instructions.

For anti-GFAP IHC, antigen retrieval was performed in 0.1 M citric acid buffer (pH 6.0) at 100 °C for 20 min. Endogenous peroxidase activity was blocked using 1.0% H<sub>2</sub>O<sub>2</sub> and 0.1% sodium azide in TBS for 10 min. Endogenous mouse Ig was blocked by incubating sections with goat anti-mouse Fab fragments (Jackson ImmunoResearch) diluted 1:50 in Biocare Medical Mouse block M for 60 minutes. Nonspecific antibody binding was inhibited by incubation with Biocare Medical Background Sniper + 2% BSA for 30 min. Mouse anti-GFAP clone GA-5 (Biocare Medical, CM065C), was diluted 1:250 in the above buffer and incubated in sections for 1 hr at room temperature. After washes, Perkin Elmer Goat anti-mouse HRP diluted 1:500 in TBS-T was applied for 60 minutes. Nova Red development and counter staining was performed as for 4G4 above.

### Infection of human cortical brain organoids

hBOs were reprogrammed from adult dermal fibroblasts (HDFa, Gibco, C0135C) using the CytoTune-iPS 2.0 Sendai Reprogramming Kit (Invitrogen, A16518)<sup>36</sup>, and were grown using the CelVivo Clinostar incubator (Invitro Technologies) as described<sup>37</sup>. On the day of infection, ~30-day-old hBOs were transferred from each Clinoreactor into an ultra-low-binding 24-well plate (one hBO per well), and each hBO was infected with 10<sup>5</sup> CCID<sub>50</sub> of either JEV<sub>Nakayama</sub>, JEV<sub>FU</sub>, JEV<sub>NSW/22</sub>, MVEV<sub>TC123130</sub>, IMOJEV<sup>38</sup>, or YFV 17D<sup>25</sup> for ~4 h. For shorter-term culture (up to 4 days for viral growth kinetics), virus inoculum was removed, and hBOs were washed twice with media in the well, before 1 ml differentiation media was added to each well, and the 24-well plate was placed within a humidified tissue culture incubator at 37 °C, 5% CO<sub>2</sub> for up to 4 days. For culture up to 11 dpi, hBOs were washed twice with media by transferring them to an ultra-low-binding six-well plate containing 5 ml of media, and then hBOs were transferred into 50 mm LUMOX gas exchange dishes (SARSTEDT) (4 organoids per dish) containing 7 ml of differentiation media, and placed within a humidified tissue culture incubator at 37 °C, 5% CO<sub>2</sub> for up to 11 days. hBOs were imaged using an EVOS FL (Advanced Microscopy Group), and organoid 2D image circumference was determined by drawing around the edge of the organoid using Image J v1.53<sup>39</sup>.

### Human serum neutralization assays

Cohort 1 is human serum samples from 9 participants (age >50 years), and were collected ~28 days ( $\pm 2$  days) after vaccination (administered via either subcutaneous or intradermal injection) with the IMOJEV vaccine (Sanofi-

Aventis Australia). Serum was also collected pre-IMOJEV vaccination for these same participants for baseline analysis. Cohort 2 is human serum samples from 10 participants (age range 24 to 60 years), collected at variable times post-vaccination (~2 months to 1 year). Pre-vaccination serum was not collected for Cohort 2, and the vaccine was administered subcutaneously. Neutralizing antibodies against IMOJEV, JEV<sub>Nakayama</sub>, JEV<sub>FU</sub>, JEV<sub>NSW/22</sub>, and MVEV<sub>TC123130</sub> were measured by plaque reduction neutralization (PRNT) assay<sup>40,41</sup>. BHK-21 cells were seeded at  $1.6 \times 10^5$  cells per well in 24 well plates overnight at 37 °C. Serum samples were heat-inactivated (56 °C, 30 min) and serially diluted fourfold from 1:5 to 1:160 in BHK-21 cell culture media. The serum was then incubated with 100–110 pfu of IMOJEV, JEV<sub>Nakayama</sub>, JEV<sub>FU</sub>, JEV<sub>NSW/22</sub>, and MVEV<sub>TC123130</sub> for 1 h at 37 °C. Serum plus virus mixtures were then added to BHK-21 cell monolayers and incubated for 1 hr at 37 °C to enable non-neutralized virus to adsorb to cells. Thereafter, 1 ml of 0.375% w/v carboxymethyl cellulose (CMC, Sigma-Aldrich)/RPMI 1640 was added, and the plates were incubated at 37 °C in a CO<sub>2</sub> incubator for 3 days for JEV<sub>Nakayama</sub> and 4 days for JEV<sub>NSW/22</sub>. The CMC medium was then removed, and the cell monolayers were fixed and stained with 0.1% w/v crystal violet (Sigma-Aldrich) in formaldehyde (1% v/v) and methanol (1% v/v). Plate wells were washed with tap water, dried and the plaques were counted. The PRNT<sub>50</sub> titer was interpolated from plaque count compared to the average plaque count for the naive or no serum control.

### Envelope protein structure visualizations

JEV envelope protein structure was downloaded from the Protein Data Bank (PDB: 5WSN)<sup>42</sup>. Envelope structure visualizations were generated using PyMol Molecular Graphics System (version 2.3.3; Schrodinger, NY, USA). Virus sequences were downloaded from GenBank (see Supplementary Fig. 1 for accession numbers) and aligned using Mega-X (Molecular Evolutionary Genetics Analysis 10, Penn State University, State College, PA, USA) and the ClustalW plugin with default parameters. Differences in amino acids compared to IMOJEV were colored according to structural conservation as described<sup>43</sup>.

### Statistics

The *t* test was used when the difference in variances was <4-fold, skewness was >−2, and kurtosis was <2 (Excel 2016). Otherwise, the non-parametric Kolmogorov–Smirnov exact test or Mann–Whitney test was used (GraphPad Prism 8). Paired *t* test was used for comparing neutralization of different viruses with the same human serum sample. For matched human serum samples, a paired non-parametric Wilcoxon matched-pairs signed rank test (GraphPad Prism 8) was used since the difference in variance was >4-fold. Kaplan–Meier statistics were determined by log-rank (Mantel–Cox) test. Area under the curve analyses was performed in GraphPad Prism 8, with an area under the curve values then compared by *t* test. Correlation analyses of PRNT<sub>50</sub> with envelope conservation used the non-parametric Spearman's rank-order correlation.

### Results

#### JEV<sub>NSW/22</sub> infection produces a viremia but is not lethal in C57BL/6J mice

The JEV<sub>NSW/22</sub> G4 virus was isolated from the brain of a stillborn piglet in New South Wales (NSW), Australia, in February 2022 (GenBank accession OP904182). The JEV<sub>Nakayama</sub> genotype 3 prototype was isolated in Japan in 1935 (GenBank accession EF571853), and the JEV<sub>FU</sub> genotype 2 was isolated in Australia in 1995<sup>44,45</sup> (GenBank accession AF217620). MVEV was isolated in Australia in 1974<sup>46</sup> (MVEV<sub>TC123130</sub>, GenBank accession JN119814). The latter three viruses were isolated from human patients. Brief descriptions of the viral isolates and their phylogenic relationships with other flaviviruses are provided in Supplementary Fig. 1.

Six-week-old adult C57BL/6J mice were infected subcutaneously (s.c.) with  $5 \times 10^5$  CCID<sub>50</sub> of the aforementioned viruses. Viremia for all four viruses was broadly similar, peaking 1 day post infection (dpi) at 2–3 log<sub>10</sub>CCID<sub>50</sub>/ml of serum, with nearly all mice showing no detectable

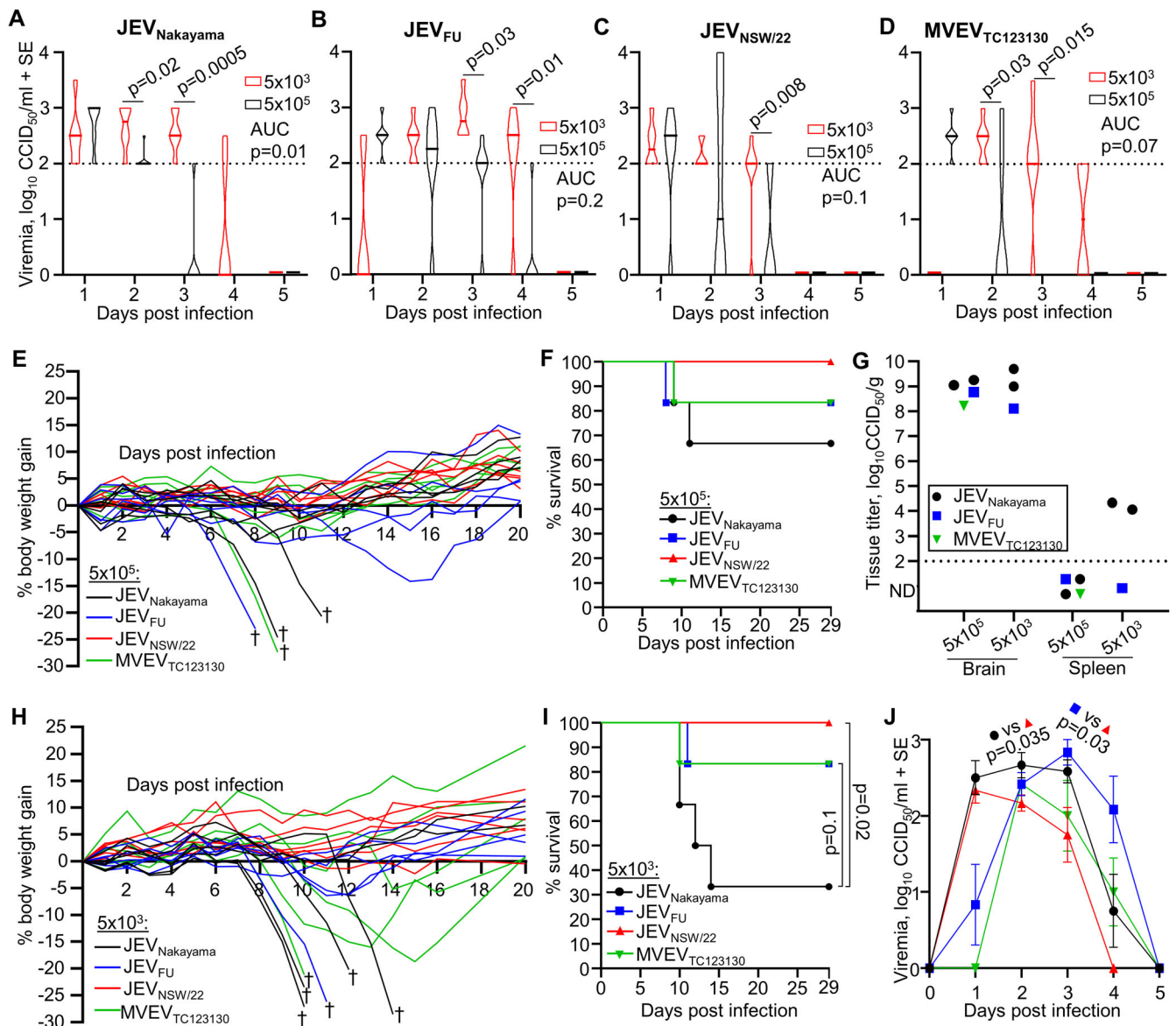
viremia by day 4 (Fig. 1A–D). The infection appeared to stall weight gain for most mice until day ~10 (Fig. 1E). Between 8 and 12 dpi, four mice (2 infected with JEV<sub>Nakayama</sub>, 1 JEV<sub>FU</sub> and 1 MVEV<sub>TC123130</sub>) out of the total of 24 mice showed weight loss ≥20% and were euthanized (Fig. 1E, †). A further four mice (1 JEV<sub>Nakayama</sub>, 2 JEV<sub>FU</sub>, 1 MVEV<sub>TC123130</sub>) lost >5% of their body weight, but subsequently recovered. None of the JEV<sub>NSW/22</sub> infected mice lost more than ~3% of their body weight (Fig. 1E). Kaplan–Meier survival curves provided no significant differences for the different viral isolates (Fig. 1F). Mice that were euthanized also displayed varying levels of abnormal posture (hunching), reduced activity, and fur ruffling, on the day of euthanasia (Supplementary Fig. 2A). For the four mice that were euthanized (Fig. 1E), these mice had very high levels of brain infection (≈8–9 log<sub>10</sub>CCID<sub>50</sub>/mg) (Fig. 1G). At the time of euthanasia, viral titers in the spleen in these mice were below the level of detection (Fig. 1G), consistent with the viremia data. Viral titers in the brain and spleen were assessed exclusively in mice that were humanely euthanized due to disease. Euthanizing healthy mice at corresponding time points was avoided to ensure the generation of a Kaplan–Meier survival curve.

We then infected six-week-old adult C57BL/6J mice with  $5 \times 10^3$  CCID<sub>50</sub> (s.c.) of the aforementioned viruses to determine if reduced inoculation dose can increase viremia and neuropenetrance, as has been reported previously<sup>47</sup>. Compared to mice infected with  $5 \times 10^5$  CCID<sub>50</sub>, viremia in mice infected with  $5 \times 10^3$  CCID<sub>50</sub> of virus was significantly higher at later time points for JEV<sub>Nakayama</sub> (Fig. 1A, day 2–3), JEV<sub>FU</sub> (Fig. 1B, day 3–4), and MVEV<sub>TC123130</sub> (Fig. 1D, day 2–3). For JEV<sub>NSW/22</sub>, viremia at day 3 post infection was higher on average but this did not reach statistical significance (Fig. 1C). The viremia area under the curve was significantly higher for JEV<sub>Nakayama</sub> inoculated with the lower dose (Fig. 1A), but was not significantly different for the other viruses. This coincided with an increase in mortality for JEV<sub>Nakayama</sub> inoculated at the lower dose, but not for the other viruses (Fig. 1H, I). Viral titers in the brains or spleens of mice that succumbed to infection were not significantly different between inoculum doses (Fig. 1G). JEV<sub>NSW/22</sub> had a significantly lower viremia compared to JEV<sub>Nakayama</sub> and JEV<sub>FU</sub> (Fig. 1J), consistent with significantly lower mortality compared with JEV<sub>Nakayama</sub> (Fig. 1I), with no C57BL/6J mice infected with JEV<sub>NSW/22</sub> succumbing to infection (Fig. 1F, I). Overall, this suggests that JEV<sub>NSW/22</sub> was significantly less virulent in C57BL/6J mice.

#### C57BL/6J mice infected with JEV or MVEV led to viral neuroinvasion, apoptosis, and reactive astrogliosis

The brains of C57BL/6J mice were analyzed by immunohistochemistry (IHC) using the pan-flavivirus anti-NS1 antibody (4G4). This antibody detects the viral non-structural protein 1, which is a highly conserved multifunctional protein important for flavivirus replication<sup>48</sup>. Staining was consistently seen in the cortex for JEV and MVEV-infected mice, with staining also seen in the thalamus (Fig. 2A, Supplementary Fig. 3A–C). Prevailing infection of the cerebral cortex and thalamus parallels IHC data from post-mortem human brains<sup>6,49</sup>. In some mice, virus was detected in other brain regions, including the hippocampus, anterior olfactory nucleus (Fig. 2A, JEV<sub>Nakayama</sub>), and caudate putamen (Supplementary Fig. 3B, JEV<sub>FU</sub>). A JEV<sub>FU</sub>-infected mouse that lost ~15% body weight and then recovered (Fig. 1E) had residual virus staining in the cortex at day 29 post infection (Supplementary Fig. 3D), while a mouse that survived infection with minimal weight loss had no detectable viral antigen staining (Supplementary Fig. 3E). As expected, 4G4 staining was associated primarily with cells showing neuronal morphology (Fig. 2A, Supplementary Fig. 3D).

ApopTag staining was evident in areas that were positive for virus staining, particularly in the cortex, and was again associated with cells showing neuronal morphology (Fig. 2B and Supplementary Fig. 4A–C). Apoptotic cells were also evident in the mouse that showed ~15% body weight loss and recovered (Supplementary Fig. 4D). Mice with signs of disease from JEV or MVEV infection (Supplementary Fig. 2) thus showed high levels of viral antigen and apoptosis in the brain. In addition, like in humans, animals can recover from brain infection, despite apoptotic damage, although neurological sequelae may ensue.



**Fig. 1 | JEV and MVEV infection in C57BL/6J mice.** **A–D** Female  $\approx$ 6 week old C57BL/6J mice were infected s.c. with  $5 \times 10^5$  (black) or  $5 \times 10^3$  (red) CCID<sub>50</sub> of the indicated viruses. Violin plots for  $n = 6$  per group over 5 days is shown. The horizontal line within the violin plot represents the median. All mice recorded a detectable viremia for at least one timepoint. Limit of detection is  $2 \log_{10}$ CCID<sub>50</sub>/ml of serum. Statistics represent *t* test or Kolmogorov–Smirnov exact test at the indicated timepoint or for area under the curve (AUC) values (see methods section). **E** Percent body weight change of individual mice after infection with the indicated virus at  $5 \times 10^5$  CCID<sub>50</sub> compared to each mouse’s weight on day zero. Four mice lost  $\geq 20\%$  body weight and were euthanized (†). **F** Kaplan–Meier plot showing percent survival ( $n = 6$  for each virus/virus isolate inoculated at  $5 \times 10^3$  CCID<sub>50</sub>). **G** Viral tissue titers in brains and spleens of seven euthanized mice at the time when the

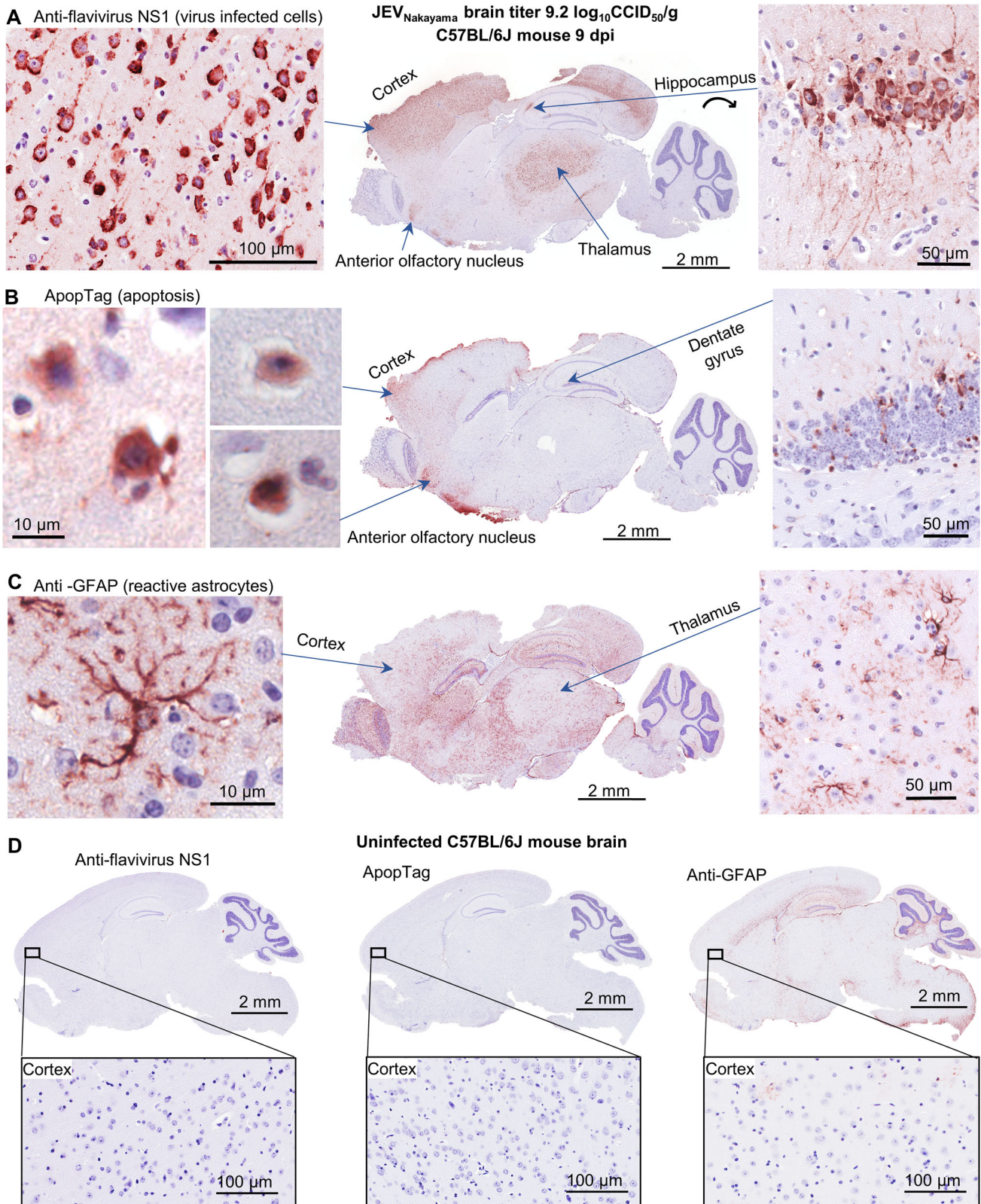
criteria for humane euthanasia was met (see Supplementary Fig. 2) ( $n = 4$  JEV<sub>Nakayama</sub>—black circles,  $n = 2$  JEV<sub>FU</sub>—blue square,  $n = 1$  MVEV<sub>TC123130</sub> green downward triangle). Tissue titers determined by CCID<sub>50</sub> assay (limit of detection  $\sim 2 \log_{10}$ CCID<sub>50</sub>/g). **H** Percent body weight change of individual mice after infection with the indicated virus at  $5 \times 10^3$  CCID<sub>50</sub> compared to each mouse’s weight on day zero. Six mice lost  $\geq 20\%$  body weight and were euthanized (†). **I** Kaplan–Meier plot showing percent survival ( $n = 6$  for each virus/virus isolate inoculated at  $5 \times 10^3$  CCID<sub>50</sub>). **J** Viremia comparing JEV<sub>Nakayama</sub> (black circles), JEV<sub>FU</sub> (blue squares), JEV<sub>NSW/22</sub> (red triangles) and MVEV<sub>TC123130</sub> (green downward triangles) at  $5 \times 10^5$  inoculation dose; data is a reanalysis of data presented in Fig. 1A–D. Data are mean of  $n = 6$  per group and error bars represent standard error. Statistics are *t* test for JEV<sub>Nakayama</sub> versus JEV<sub>NSW/22</sub> on day 2 and JEV<sub>FU</sub> versus JEV<sub>NSW/22</sub> on day 3.

Astrocytes are a type of glial cell that provides physical and chemical support to neurons. Astrocytes become activated (reactive astrogliosis) in response to brain pathology, including from JEV infection<sup>50–52</sup>. Reactive astrocytes are characterized by the upregulated expression of glial fibrillary acidic protein (GFAP) and the growth/extension of cellular processes<sup>53</sup>. Brains from mice with signs of disease from JEV or MVEV infection (Supplementary Fig. 2) had significant upregulation of GFAP-positive astrocytes throughout the brain, including the heavily infected areas of the cortex, thalamus and anterior olfactory nucleus (Fig. 2C, Supplementary Fig. 5). GFAP is constitutively expressed in astrocytes of the hippocampus,

corpus callosum, and cerebral peduncle<sup>54</sup>, consistent with staining seen in uninfected brains (Fig. 2D, right). Widespread reactive astrogliosis is thus a feature of lethal JEV and MVEV encephalitis in C57BL/6J mice.

**JEV<sub>NSW/22</sub> is less virulent than JEV<sub>Nakayama</sub> and JEV<sub>FU</sub> in *Irf7*<sup>-/-</sup> mice**

C57BL/6J mice had a relatively low viremia and low penetrance of lethal brain infection for JEV<sub>NSW/22</sub> (Fig. 1). A model with a higher penetrance of brain infection is clearly desirable, given encephalitis is the key clinical concern for JEV infections. Interferon regulatory factor 7 knockout (*Irf7*<sup>-/-</sup>



mice) were thus chosen as these mice enhanced neuroinvasion for WNV<sup>55</sup>, as well as another arbovirus<sup>30</sup>. We first sought to optimize the inoculation dose for JEV<sub>NSW/22</sub> in *Irf7*<sup>-/-</sup> mice by comparing  $5 \times 10^5$ ,  $5 \times 10^4$ ,  $5 \times 10^3$ , and  $5 \times 10^2$  CCID<sub>50</sub>. The inoculation dose of  $5 \times 10^3$  CCID<sub>50</sub> provided the highest viremia area under the curve, which was significantly higher than  $5 \times 10^4$  and  $5 \times 10^5$  CCID<sub>50</sub> inoculation doses (Fig. 3A). The highest

infection dose ( $5 \times 10^5$ ) led to the highest peak viremia on day 1 post infection, but was cleared significantly more quickly than lower infection doses (Fig. 3A,  $p = 0.004$ ). Our data agrees with the concept of an ‘optimal dose’ that is not too high as to excessively stimulate type I IFN, but is high enough to establish a robust viremia<sup>47</sup>. One mouse infected with  $5 \times 10^3$  CCID<sub>50</sub> JEV<sub>NSW/22</sub> succumbed to infection (Fig. 3C, D), while all other mice

**Fig. 2 | IHC for viral antigen, apoptosis, and reactive astrocytes in JEV-infected C57BL/6J mouse brain.** IHC of JEV<sub>Nakayama</sub> infected brain (required euthanasia on 9 dpi, brain virus titer  $9.2 \log_{10} \text{CCID}_{50}/\text{g}$ ), which is representative of all other C57BL/6J brains from infected mice requiring euthanasia ( $n = 10$ ). **A** Staining for flavivirus NS1 using the 4G4 monoclonal antibody. High magnification images show cells with neuronal morphology in the cortex (left) and hippocampus (right). The latter also shows staining of dendrites and axons (fibrillar patterns above and densely staining cells). 4G4 staining of the brains of the other mice requiring euthanasia (marked by † in Fig. 1E) is shown in Supplementary Fig. 3. **B** ApopTag staining of the same brain

as in 'A'. High magnification images show cells with neuronal morphology in the cortex (left) and apoptotic cells in the dentate gyrus (right). ApopTag staining of the brains of the other mice requiring euthanasia is shown in Supplementary Fig. 4. **C** Staining for GFAP, a marker of reactive astrocytes, for the same brain as in 'A'. High magnification images show a typical reactive astrocyte in the cortex (left) and reactive astrocytes in the thalamus (right). GFAP IHC for the brains of the other mice requiring euthanasia is shown in Supplementary Fig. 5. **D** IHC negative controls; uninfected mouse brain stained with 4G4 (left), ApopTag (middle), and anti-GFAP (right).

survived JEV<sub>NSW/22</sub> infection. An inoculation dose of  $5 \times 10^3 \text{CCID}_{50}$  was thus chosen to compare JEV and MVEV strains in *Irf7*<sup>-/-</sup> mice.

JEV<sub>NSW/22</sub>, JEV<sub>Nakayama</sub>, and JEV<sub>FU</sub> infection of *Irf7*<sup>-/-</sup> mice (dose  $5 \times 10^3 \text{CCID}_{50}$ ) resulted in robust viremias, while MVEV<sub>TC123130</sub> viremia was significantly lower than JEV (Fig. 3B,  $p < 0.0001$ ). JEV<sub>NSW/22</sub> had a significantly lower viremia (area under the curve) compared to JEV<sub>FU</sub> (Fig. 3B,  $p = 0.004$ ). Infection of *Irf7*<sup>-/-</sup> mice with the different JEV isolates and MVEV ( $n = 14$  mice per group,  $n = 56$  total infected mice) illustrated a mean weight loss of 5–10% for surviving mice across all four groups (Fig. 3C), with the exception of 20 mice that reached ethically defined endpoints for weight loss and/or disease manifestations (Supplementary Fig. 6). Kaplan-Meier curves illustrated significantly higher survival for *Irf7*<sup>-/-</sup> mice infected with JEV<sub>NSW/22</sub> or MVEV<sub>TC123130</sub> when compared with JEV<sub>Nakayama</sub> and JEV<sub>FU</sub>, with only 1/14 JEV<sub>NSW/22</sub> or MVEV<sub>TC123130</sub> infected *Irf7*<sup>-/-</sup> mice requiring euthanasia (Fig. 3D). These data are consistent with the significantly lower viremias of JEV<sub>NSW/22</sub> and MVEV<sub>TC123130</sub> infected *Irf7*<sup>-/-</sup> mice (Fig. 3B). On average, mice that succumbed to infection (either JEV<sub>Nakayama</sub>, JEV<sub>FU</sub>, JEV<sub>NSW/22</sub> or MVEV<sub>TC123130</sub>) had a more robust viremia (Fig. 3E), further implicating a higher viremia in a higher chance of lethal neuropenetrance in *Irf7*<sup>-/-</sup> mice.

Brain titers in euthanized *Irf7*<sup>-/-</sup> mice were  $\sim 4\text{--}9 \log_{10} \text{CCID}_{50}/\text{gram}$ , while spleen titers were  $< 2\text{--}5 \log_{10} \text{CCID}_{50}/\text{g}$  (Fig. 3F). IHC staining for viral antigen confirmed JEV infection in the brains of *Irf7*<sup>-/-</sup> mice, with reduced levels of staining reflecting lower brain titers (Fig. 3G, H, Supplementary Fig. 7). Viral antigen staining was consistently present in the cortex, with staining also seen in the medulla, thalamus, and hippocampus (Fig. 3G, H, Supplementary Fig. 7). Brain regions showing ApopTag staining broadly overlapped with staining for viral antigen (Fig. 3G, H). Reactive astrogliosis was also evident (Supplementary Fig. 5). Overall, this suggests that JEV infection of *Irf7*<sup>-/-</sup> mice produced a consistent viremia, with mice showing brain infection and apoptotic damage.

### JEV<sub>NSW/22</sub> and JEV<sub>FU</sub> are more sensitive to type I IFN compared to JEV<sub>Nakayama</sub> and MVEV<sub>TC123130</sub>

Viremia and survival were compared between C57BL/6J and *Irf7*<sup>-/-</sup> mouse strains infected with  $5 \times 10^3 \text{CCID}_{50}$  virus to determine whether differences in viremia may be associated with the differences in viral neuropenetrance and survival. Area under the curve analyses showed a significantly increased/prolonged viremia in *Irf7*<sup>-/-</sup> mice compared to C57BL/6J mice for all JEV strains (Fig. 4A–C). However, MVEV<sub>TC123130</sub> viremia was not significantly improved in *Irf7*<sup>-/-</sup> mice (Fig. 4D). Kaplan–Meier survival curves indicated that JEV<sub>FU</sub> had a significantly increased mortality in *Irf7*<sup>-/-</sup> mice compared to C57BL/6J mice (Fig. 4F), while there was no significant difference between mouse strains for the other viruses (Fig. 4E, G–H).

Differences in sensitivity to type I IFN between virus strains may explain the different outcomes in C57BL/6J versus *Irf7*<sup>-/-</sup> mice. To examine type I IFN sensitivity in the absence of *in vivo* adaptive immune responses, mouse embryonic fibroblasts (MEFs) with (wild-type) and without (*Irf3/7*<sup>-/-</sup>) functional type I IFN production were infected with JEV<sub>Nakayama</sub>, JEV<sub>FU</sub>, JEV<sub>NSW/22</sub> and MVEV<sub>TC123130</sub>. In wild-type MEFs, JEV<sub>FU</sub> and JEV<sub>NSW/22</sub> had significantly lower virus replication kinetics compared to JEV<sub>Nakayama</sub> and MVEV<sub>TC123130</sub> (Fig. 4I). In *Irf3/7*<sup>-/-</sup> MEFs, the difference in replication between virus strains was diminished, with JEV<sub>FU</sub> not significantly different to JEV<sub>Nakayama</sub> (Fig. 4J). This suggests

that JEV<sub>FU</sub> and JEV<sub>NSW/22</sub> were more sensitive to type I IFN compared to JEV<sub>Nakayama</sub> and MVEV<sub>TC123130</sub>, consistent with an increase in mortality in *Irf7*<sup>-/-</sup> mice (partially defective type I IFN responses) compared to C57BL/6J mice for JEV<sub>FU</sub> but not for JEV<sub>Nakayama</sub> or MVEV<sub>TC123130</sub>. Although not statistically significant, this trend was also observed for JEV<sub>NSW/22</sub>, which caused mortality in one *Irf7*<sup>-/-</sup> mouse, compared to zero C57BL/6J mice (Fig. 4G).

### *Ifnar*<sup>-/-</sup> mice are a model of lethal JEV viremia without neuroinvasion

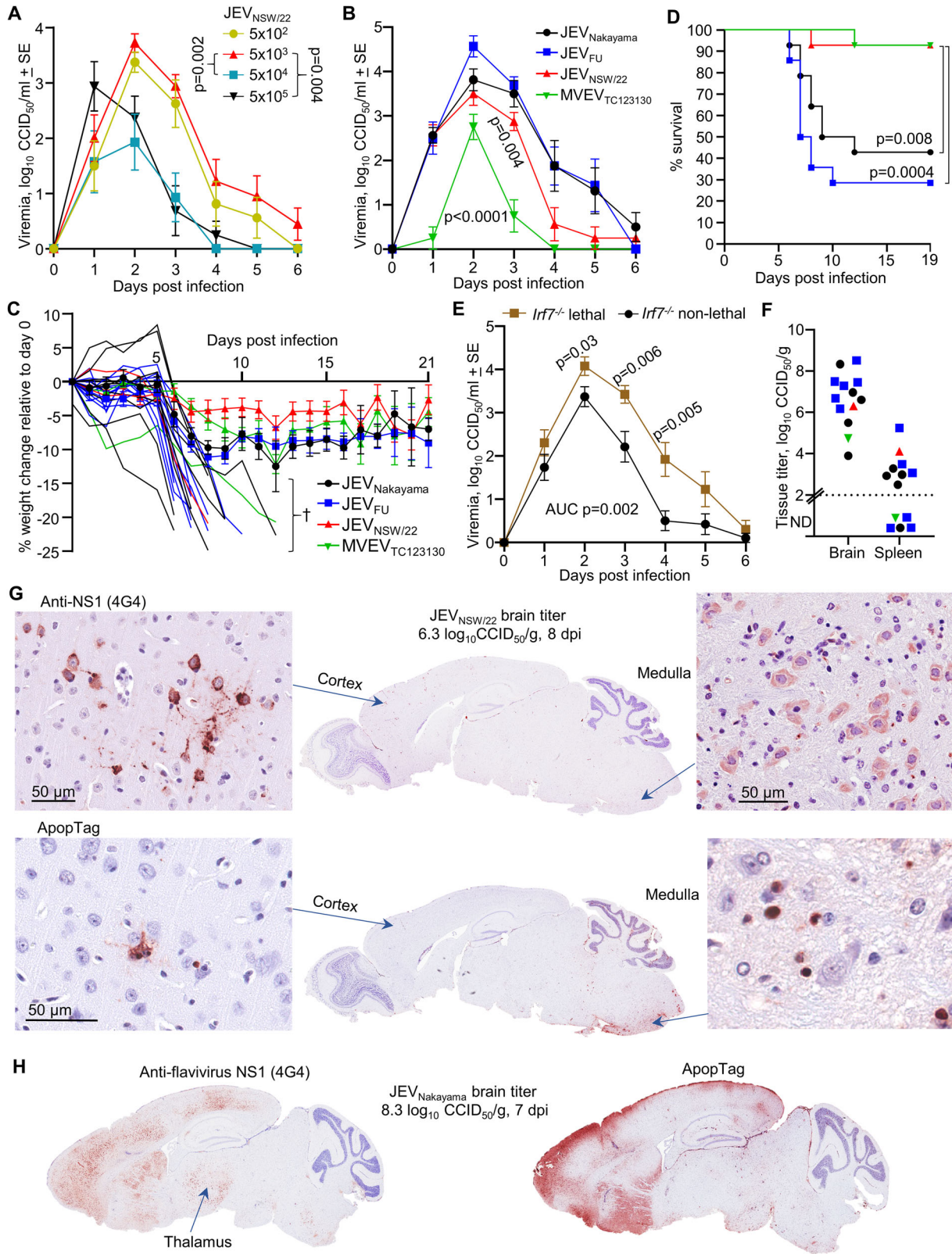
As JEV<sub>NSW/22</sub> did not lead to fatal neuroinvasion in C57BL/6J mice, and fatal neuroinvasion in *Irf7*<sup>-/-</sup> mice was rare, type I interferon receptor knockout (*Ifnar*<sup>-/-</sup>) mice were evaluated. *Ifnar*<sup>-/-</sup> mice have been used extensively in studies of pathogenic flaviviruses such as Zika virus (ZIKV), West Nile virus (WNV), and Yellow fever virus (YFV), and generally provide a robust viremia and a lethal outcome<sup>23–25,35</sup>. Adult *Ifnar*<sup>-/-</sup> mice were infected with  $5 \times 10^5 \text{CCID}_{50}$  of JEV<sub>Nakayama</sub>, JEV<sub>FU</sub>, JEV<sub>NSW/22</sub>, or MVEV<sub>TC123130</sub> via s.c. injection. A robust viremia developed, reaching 6–8  $\log_{10} \text{CCID}_{50}/\text{ml}$  of serum by 2 dpi (Fig. 5A). JEV<sub>NSW/22</sub> infected mice had  $\approx 2 \log$  lower viremia compared to JEV<sub>Nakayama</sub> and JEV<sub>FU</sub> on 3 dpi (Fig. 5A). Mice displayed a rapid loss of body weight, which was slightly delayed for JEV<sub>NSW/22</sub> and MVEV<sub>TC123130</sub> at 2 dpi (Fig. 5B), requiring euthanasia 2/3 dpi (Fig. 5C). Mice infected with any of the JEV strains also displayed varying levels of abnormal posture (hunching), reduced activity, and fur ruffling (Supplementary Fig. 8A).

Consistent with the robust viremia (Fig. 5A), spleen tissue titers reached  $\sim 6\text{--}9 \log_{10} \text{CCID}_{50}/\text{g}$  at 2–3 dpi (Fig. 5D). Similar levels were seen in brains (Fig. 5E); however, 4G4 staining illustrated that brain cells were not infected (Fig. 5F). Viremia and brain titers correlated strongly (Supplementary Fig. 8B), arguing that the brain titers (Fig. 5E) likely arose from virus in the blood vessels of the brain. Interestingly, anti-flavivirus NS1 antibody staining was clearly present in blood vessels (Fig. 5G, H), with extracellular JEV NS1 also found in serum of JEV patients<sup>36</sup>. Secreted NS1 from JEV-infected cells is reported to increase vascular leakage and may contribute to mortality<sup>57</sup>, although overt vascular leakage of NS1 was not evident in IHC, with lack of type I IFN signaling potentially involved<sup>58,59</sup>. Neither apoptosis (Supplementary Fig. 8C), nor significantly increased staining for reactive astrocytes (Supplementary Fig. 5, *Ifnar*<sup>-/-</sup>) was apparent for the brains of JEV-infected *Ifnar*<sup>-/-</sup> mice. Therefore, *Ifnar*<sup>-/-</sup> mice infected with JEV or MVEV represent a model of lethal viremia without viral neuroinvasion.

### JEV causes severe histological lesions in C57BL/6J and *Irf7*<sup>-/-</sup> mouse brains

H&E staining of JEV-infected mouse brains revealed a series of histopathological lesions, predominantly neuronal degeneration with pyknotic nuclei (often associated with apoptosis), neuronal vacuolation, perivascular cuffing, leukocyte infiltrates, hemorrhage, meningitis and microgliosis (Fig. 6A). IHC for the microglial marker, Iba1, confirmed the presence of microgliosis (Fig. 6B).

The presence of H&E lesions was scored for the different JEV isolates and mouse strains (Supplementary Fig. 9). C57BL/6J and *Irf7*<sup>-/-</sup> mice that were euthanized at the time when viral neuroinvasion led to the criteria for humane euthanasia being met (full score cards provided in Supplementary Figs. 2 and 7) had a broadly similar incidence of the aforementioned lesions,



whereas only hemorrhage was seen in *Ifnar*<sup>-/-</sup> mice (Supplementary Fig. 9, Acute). Leukocyte infiltration was quantified by calculating the ratio of nuclear/cytoplasmic staining, as leukocytes have a higher ratio of nuclear to cytoplasmic staining<sup>33</sup>. This confirmed that C57BL/6J and *Irf7*<sup>-/-</sup> mice that succumbed to infection had significantly increased leukocyte infiltrates (Supplementary Fig. 9B). For C57BL/6J and *Irf7*<sup>-/-</sup> mice that survived and

were euthanized later (nominally chronic phase), only hemorrhage was consistently observed (Supplementary Fig. 9, Chronic).

Lesions in C57BL/6J and *Irf7*<sup>-/-</sup> mouse brains were associated with areas of high virus infection, most prominently in the cortex (Supplementary Fig. 10); however, lesions were also found in other areas of the brain including where there was minimal viral antigen staining (Supplementary



**Fig. 3 | JEV neuroinvasive infection in *Irf7*<sup>-/-</sup> mice.** **A** *Irf7*<sup>-/-</sup> mice (15–48 weeks old) were infected s.c. with  $5 \times 10^5$ ,  $5 \times 10^4$ ,  $5 \times 10^3$ , or  $5 \times 10^2$  CCID<sub>50</sub> of JEV<sub>NSW/22</sub>. Data is the mean of  $n = 6$  per group and error bars represent standard error. Statistics are a *t* test of area under the curve for the indicated comparisons. **B** *Irf7*<sup>-/-</sup> mice (15–48 weeks old) were infected s.c. with  $5 \times 10^3$  of the indicated virus ( $n = 8$  for each virus). Statistics are a *t* test of area under the curve for JEV<sub>FU</sub> versus JEV<sub>NSW/22</sub> or MVEV<sub>TC123130</sub>. **C** Percent body weight change compared to 0 dpi. 20 mice lost >20% body weight or reached a disease score that required euthanasia (marked by †) and are plotted individually. The weight change for the remaining mice are shown as means  $\pm$  SE. Data is from three independent experiments, total  $n = 14$  mice per group. **D** Kaplan–Meier plot showing percent survival ( $n = 14$  per group, data from three independent experiments). Statistics by log-rank (Mantel–Cox) tests. Symbols as for C. **E** Viremias of *Irf7*<sup>-/-</sup> mice infected with either JEV<sub>Nakayama</sub>, JEV<sub>FU</sub>, JEV<sub>NSW/22</sub>, or MVEV<sub>TC123130</sub> were averaged for mice with non-lethal outcomes (black circles,

$n = 19$ ), versus those with lethal outcomes (brown squares,  $n = 13$ ). Statistics are comparing average viremia of mice with lethal outcomes versus mice with non-lethal outcomes at each timepoint by *t* test or Kolmogorov–Smirnov exact test. *T* test of area under the curve values for this comparison is also shown. **F** Brain and spleen tissue titers for 13 euthanized mice at the time when the criteria for humane euthanasia was met (see Supplementary Fig. 6) ( $n = 5$  JEV<sub>Nakayama</sub>,  $n = 6$  JEV<sub>FU</sub>,  $n = 1$  JEV<sub>NSW/22</sub>,  $n = 1$  MVEV<sub>TC123130</sub>). Tissue titers determined by CCID<sub>50</sub> assay (limit of detection  $\sim 2 \log_{10}$ CCID<sub>50</sub>/g). **G** IHC of JEV<sub>NSW/22</sub> infected brain (euthanized on day 8, brain virus titer  $6.3 \log_{10}$ CCID<sub>50</sub>/g) using 4G4 monoclonal antibody (top) or ApopTag (bottom). High magnification images of cortex (left) and medulla (right). **H** IHC using 4G4 (left) and ApopTag IHC (right) for the JEV<sub>Nakayama</sub> infected brain with the highest virus titer (euthanized on day 7, brain virus titer  $8.3 \log_{10}$ CCID<sub>50</sub>/g). IHC images are representative of  $n = 20$  brains of mice that succumbed to infection (some other examples are shown in Supplementary Fig. 8).

Fig. 11A). Some mice that lost >8% body weight and then recovered had persistent lesions detectable at the latest time point sampled (up to day 32 for C57BL/6J and day 21 for *Irf7*<sup>-/-</sup>) (Supplementary Fig. 11B), consistent with persistent neurological sequelae in humans that survive infection. Mice that did not lose any body weight did not show any overt brain lesions at the later time points, indicating that brain infection causing weight loss is associated with persistent lesions.

Overall, the lesions in C57BL/6J and *Irf7*<sup>-/-</sup> mouse brains are consistent with H&E detectable lesions in post-mortem human JEV-infected brains<sup>6,50,60–62</sup>. To provide additional potential insights into the nature of the cellular infiltrates (e.g., perivascular cuffing and leukocyte infiltrates, Fig. 6), immune cell type abundance estimates were obtained from reanalyzed<sup>63</sup> publicly available JEV-infected mouse brain RNA-Seq expression data<sup>64</sup> using SpatialDecon<sup>65</sup>. The dominant cell types identified were CD4 T cells, NKT cells, monocytes/macrophages, neutrophils, innate lymphoid cells, T regs, and CD8 T cells, with an increase in microglia cells also identified (Supplementary Fig. 12). GFAP was significantly upregulated 5.4-fold by infection (Supplementary Table 3, log<sub>2</sub>FC 2.4), consistent with the GFAP IHC (Fig. 2C) and reactive astrogliosis seen herein. Iba1 (Aif1) was also significantly upregulated 5.5-fold (Supplementary Table 3, log<sub>2</sub>FC 2.5), consistent with the H&E and IHC (Fig. 6) and microgliosis observed in our study.

### Productive replication of JEV in human cortical brain organoids

Human cortical brain organoids (hBOs) have been used to investigate the pathogenesis of flaviviruses, including JEV and ZIKV<sup>73,66,67</sup>. To determine JEV replication capacity in neuronal cells and the ensuing cytopathogenicity, hBOs were used as an alternative to intracranial injection of mice. We thus generated  $\sim 30$ -day-old hBOs from adult human dermal fibroblast (HDFa) cells, growing the hBOs in a CelVivo Clinostar incubator as described<sup>37</sup> (Fig. 7A). hBOs were infected with  $10^5$  CCID<sub>50</sub> of the indicated JEV and MVEV isolates, as well as (i) the IMOJEV chimeric virus vaccine (previously called ChimeriVax-JE) comprising the prME genes of the attenuated JEV<sub>SA14-14-2</sub> strain on the YFV 17D backbone<sup>38,68</sup> and (ii) the Yellow Fever live attenuated vaccine strain (YFV 17D)<sup>25</sup>, with wild-type YFV infection<sup>69</sup>, and occasionally YFV 17D vaccination<sup>70</sup>, able to cause neuropathology. hBOs were fixed in formalin 4 dpi, and IHC was undertaken using the anti-NS1 monoclonal antibody 4G4<sup>25</sup>. JEV<sub>Nakayama</sub>-infected hBOs showed the most pronounced viral antigen staining, with JEV<sub>NSW/22</sub>, JEV<sub>FU</sub>, and MVEV<sub>TC123130</sub> also showing clear staining (Fig. 7B). Viral antigen was primarily localized to the outer surface of the organoids (Fig. 7B), where cells are in direct contact with the culture medium. Viral antigen staining for IMOJEV and YFV 17D infected hBOs was also seen but was sparse (Fig. 7B and Supplementary Fig. 13A).

All viruses were able to productively infect hBOs, with JEV<sub>Nakayama</sub> infection generating higher viral titers than IMOJEV at all time points (Fig. 7C,  $p < 0.001$ ). By 96 h JEV<sub>Nakayama</sub> titers were also significantly higher than those seen for JEV<sub>NSW/22</sub> (Fig. 7C).

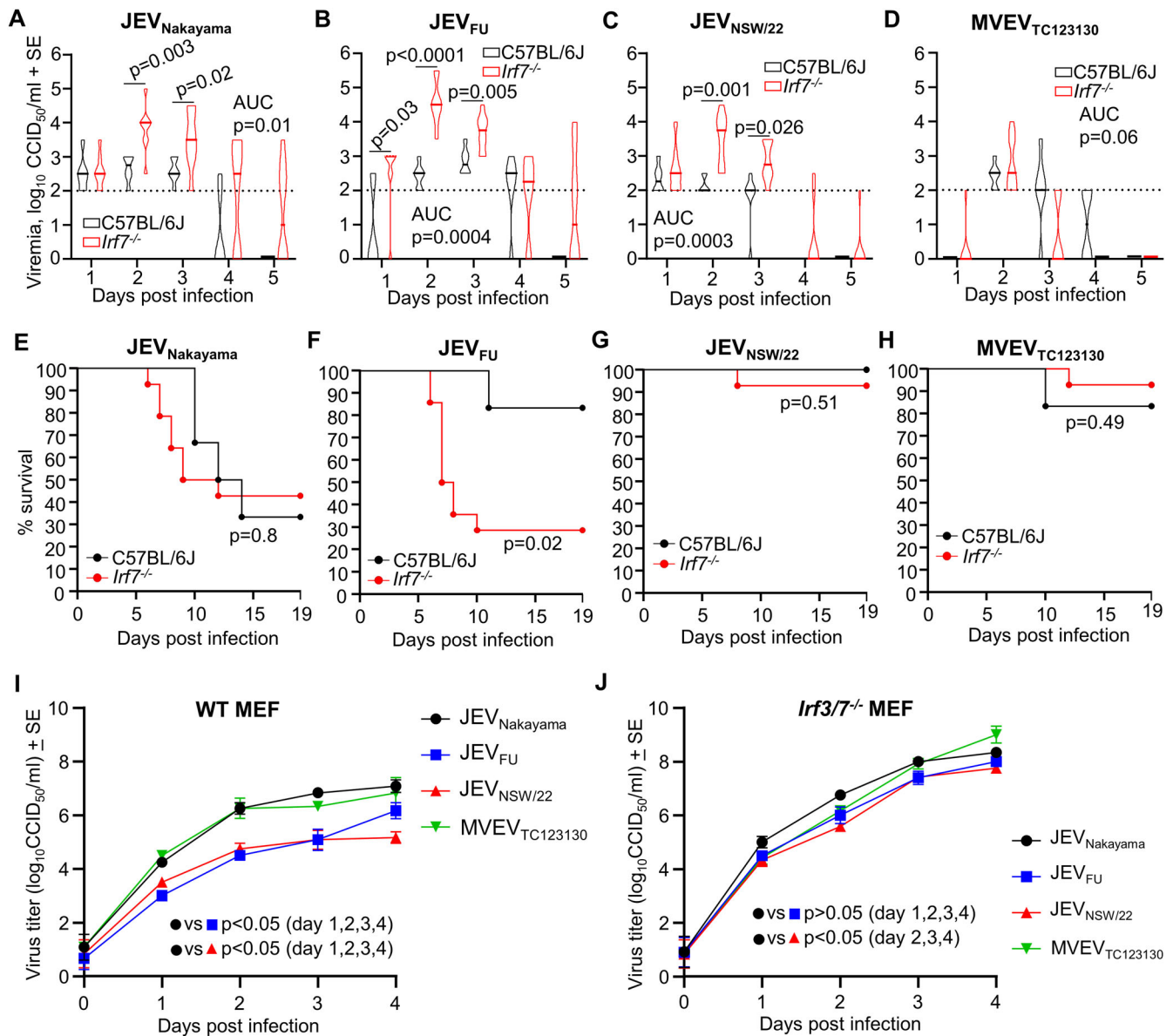
Over 11 days, uninfected hBOs grew in circumference by  $\sim 20\%$  (Fig. 7D, Uninfected). hBOs infected with JEV isolates shrank in circumference by  $\sim 5$  to  $15\%$  by 11 dpi, although when compared with uninfected organoids, this only approached significance for JEV<sub>NSW/22</sub> (Fig. 7D; Supplementary Fig. 13B, C). hBOs infected with MVEV<sub>TC123130</sub>, IMOJEV and YFV 17D infected hBO did not shrink significantly when compared with uninfected controls, although the MVEV<sub>TC123130</sub> data suggested a marginal reduction in circumference (Fig. 7C; Supplementary Fig. 13C). These data (Fig. 7D) reflect differences in viral replication (Fig. 7C) and/or viral protein immunohistochemistry (Fig. 7B) and likely reflect virus-induced CPE.

All viruses replicated productively in a human neural progenitor cell line, RENcell VM, with all viruses except IMOJEV causing fulminant CPE (Supplementary Fig. 14A, B). Although Vero E6 and BHK-21 cells are widely used for flavivirus research, CPE induced by JEV<sub>NSW/22</sub> was considerably less pronounced in these cells (Supplementary Fig. 14C, D).

### Human post-IMOJEV vaccination sera neutralizes JEV and MVEV with titers related to envelope protein amino acid conservation

IMOJEV is one of two JEV vaccines available in Australia. The IMOJEV prME genes are derived from the genotype 3 JEV<sub>SA14-14-2</sub><sup>38,68</sup> strain, which was attenuated via extensive in vitro and in vivo passaging (Supplementary Fig. 1). Most flavivirus-neutralizing antibodies recognize epitopes on the envelope protein, particularly in the putative receptor binding domain III<sup>71</sup>. JEV<sub>Nakayama</sub> and the JEV<sub>SA14-14-2</sub> component of the IMOJEV vaccine both belong to genotype 3, but JEV<sub>Nakayama</sub> has 96.8% envelope protein amino acid identity to IMOJEV (Fig. 8A). JEV<sub>FU</sub> has 96.4% envelope protein identity, while JEV<sub>NSW/22</sub> has drifted further from the genotype 2 and 3 strains with 93.4% envelope protein identity (Fig. 8A). MVEV<sub>TC123130</sub>, which is the closest phylogenetically related flavivirus to JEV (Supplementary Fig. 1B), has 80.4% envelope amino acid identity (Fig. 1A). Alignment of the envelope amino acid differences for these strains compared to IMOJEV reveal that a disproportionate number of the non-conservative changes were in domain III (Fig. 8A).

Serum-neutralizing antibodies post-vaccination are currently viewed as the best measurable correlate of vaccine protection for JEV<sup>72</sup>. To determine if the Australian outbreak genotype 4 JEV<sub>NSW/22</sub> is neutralized by antibodies produced in response to IMOJEV vaccination, serum was taken from two human cohorts. Cohort 1 was serum collected from individuals at approximately 28 days post-vaccination, while cohort 2 serum was collected at variable times post-vaccination (2 months to 1 year). Ethical approvals for cohort 1 serum allowed for neutralization assays against the IMOJEV vaccine itself, JEV<sub>Nakayama</sub>, JEV<sub>FU</sub>, and JEV<sub>NSW/22</sub> (Fig. 8B), while ethical approval for cohort 2 additionally allowed for neutralization assays against MVEV<sub>TC123130</sub> (Fig. 8C). All serum samples from both cohorts had a measurable 50% neutralization titer (PRNT<sub>50</sub>) titer against JEV<sub>Nakayama</sub>, JEV<sub>FU</sub>, and JEV<sub>NSW/22</sub> (Fig. 8B, C), indicating that seroconversion (defined as detectable neutralizing antibodies  $>10^1$  PRNT<sub>50</sub> titer) provided neutralizing antibodies that cross-react between JEV genotypes 2, 3, and 4. In cohort 1, PRNT<sub>50</sub> titers were significantly lower against JEV<sub>Nakayama</sub> and



**Fig. 4 | Comparisons of viremia and survival between C57BL/6J versus *Irf7*<sup>-/-</sup> mice, and comparison of virus replication in wild-type versus *Irf3*<sup>-/-</sup> mouse embryonic fibroblasts (MEFs).** C57BL/6J or *Irf7*<sup>-/-</sup> mice were infected with 5 × 10<sup>3</sup> CCID<sub>50</sub> of JEV<sub>Nakayama</sub>, JEV<sub>FU</sub>, JEV<sub>NSW/22</sub>, or MVEV<sub>TC123130</sub>. Data shown in A–H are a reanalysis of data presented in Fig. 1A–D, Fig. 1I, Fig. 3B, D. A–D Female ≈6 week old C57BL/6J mice (black) or *Irf7*<sup>-/-</sup> mice (red) were infected s.c. with 5 × 10<sup>3</sup> CCID<sub>50</sub> of the indicated viruses. Violin plots for n = 6 (C57BL/6J) or n = 8 (*Irf7*<sup>-/-</sup>) per group over 5 days is shown. The horizontal line within the violin plot represents the median. Limit of detection is 2 log<sub>10</sub>CCID<sub>50</sub>/ml of serum. Statistics represent *t* test or Kolmogorov–Smirnov exact test at the indicated timepoint or for

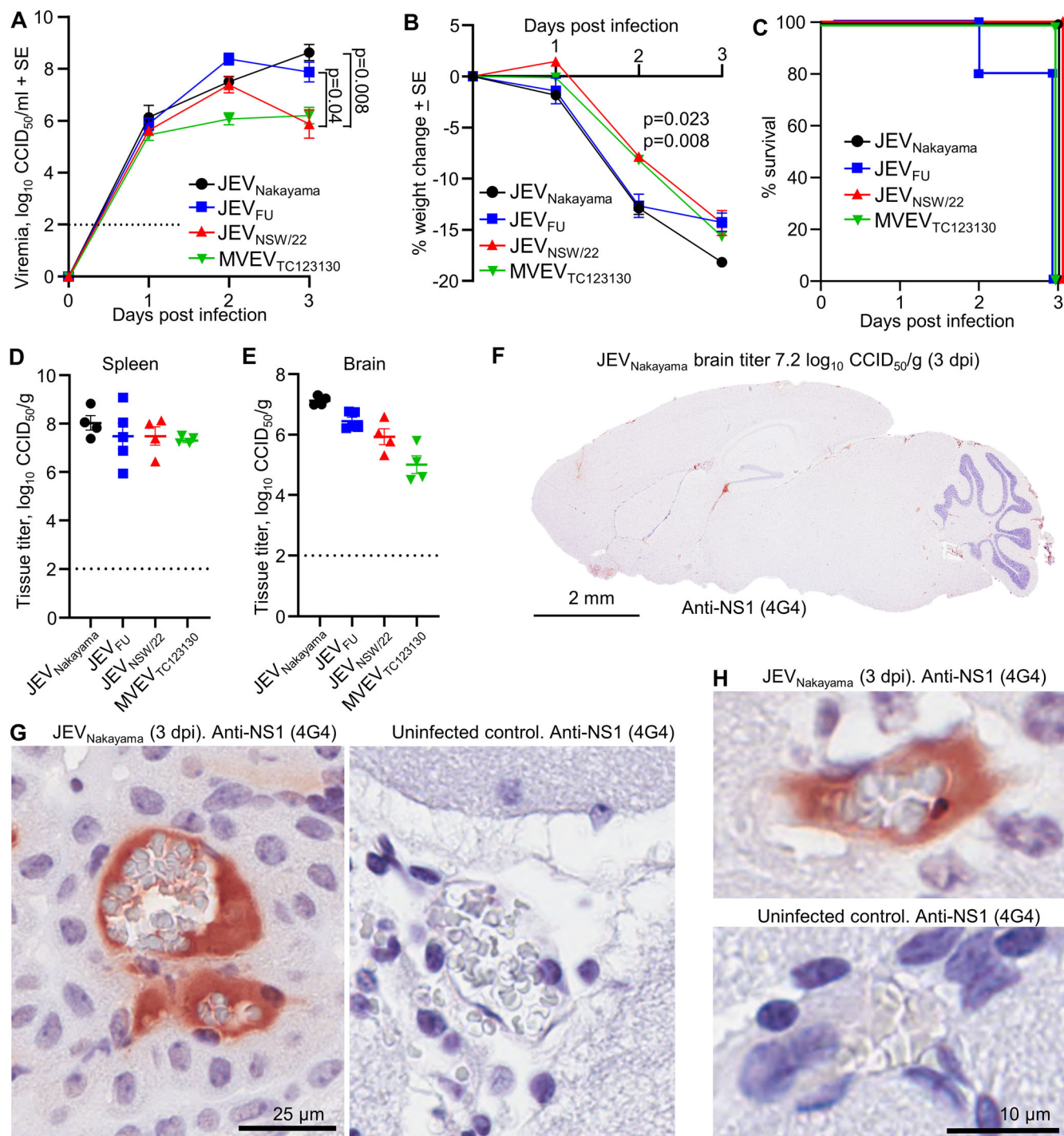
area under the curve (AUC) values (see statistics methods section). E–H Kaplan–Meier plot showing percent survival for C57BL/6J mice (black line, n = 6) and *Irf7*<sup>-/-</sup> mice (red line, n = 14) infected with 5 × 10<sup>3</sup> CCID<sub>50</sub> of the indicated virus. Statistics by log-rank (Mantel–Cox) tests. I Wild-type or J *Irf3*<sup>-/-</sup> MEFs were infected with JEV<sub>Nakayama</sub> (black circles), JEV<sub>FU</sub> (blue squares), JEV<sub>NSW/22</sub> (red triangles), or MVEV<sub>TC123130</sub> (green downward triangles) at MOI 0.1. Virus titer in the culture supernatant was monitored over 4 days. Data is the mean of two independent experiments with a total of n = 6 replicates per group. Error bars represent standard error. Statistics are by *t* test or Kolmogorov–Smirnov exact test for the indicated comparisons.

JEV<sub>NSW/22</sub> compared to against IMOJEV (Fig. 8A), suggesting key antigenic differences from the vaccine. In cohort 2, PRNT<sub>50</sub> titers against JEV<sub>Nakayama</sub> were not significantly different compared to IMOJEV, however PRNT<sub>50</sub> titers were significantly lower for JEV<sub>NSW/22</sub> (Fig. 8C). Similar conclusions were drawn when the raw percentage of plaque neutralization at a high serum dilution (1:160) was compared between virus strains (Supplementary Fig. 15). With PRNT<sub>50</sub> data from both cohorts combined, the percentage amino acid identity of the envelope protein compared to the IMOJEV vaccine significantly correlated with PRNT<sub>50</sub> titers (Fig. 8D, black line). The significance (p-value) and the correlation coefficient (rho), were similar when the same analysis was conducted, excluding the PRNT<sub>50</sub> data for

MVEV<sub>TC123130</sub> (Fig. 8D, brown line). Overall, our data indicates that IMOJEV vaccination provided neutralizing antibodies against JEV<sub>NSW/22</sub> in all individuals, but the level of cross-neutralization was related to the conservation in envelope protein amino acid sequences.

## Discussion

Herein we provide a comprehensive *in vivo* and *in vitro* characterization of the genotype 4 JEV<sub>NSW/22</sub> isolate from the recent Australian outbreak and illustrate mouse models of infection and rare CNS neuropathological manifestations that recapitulate many aspects of human and primate disease<sup>6,49–52,73</sup>. The capacity of JEV<sub>NSW/22</sub> to cause lethal neuroinvasive

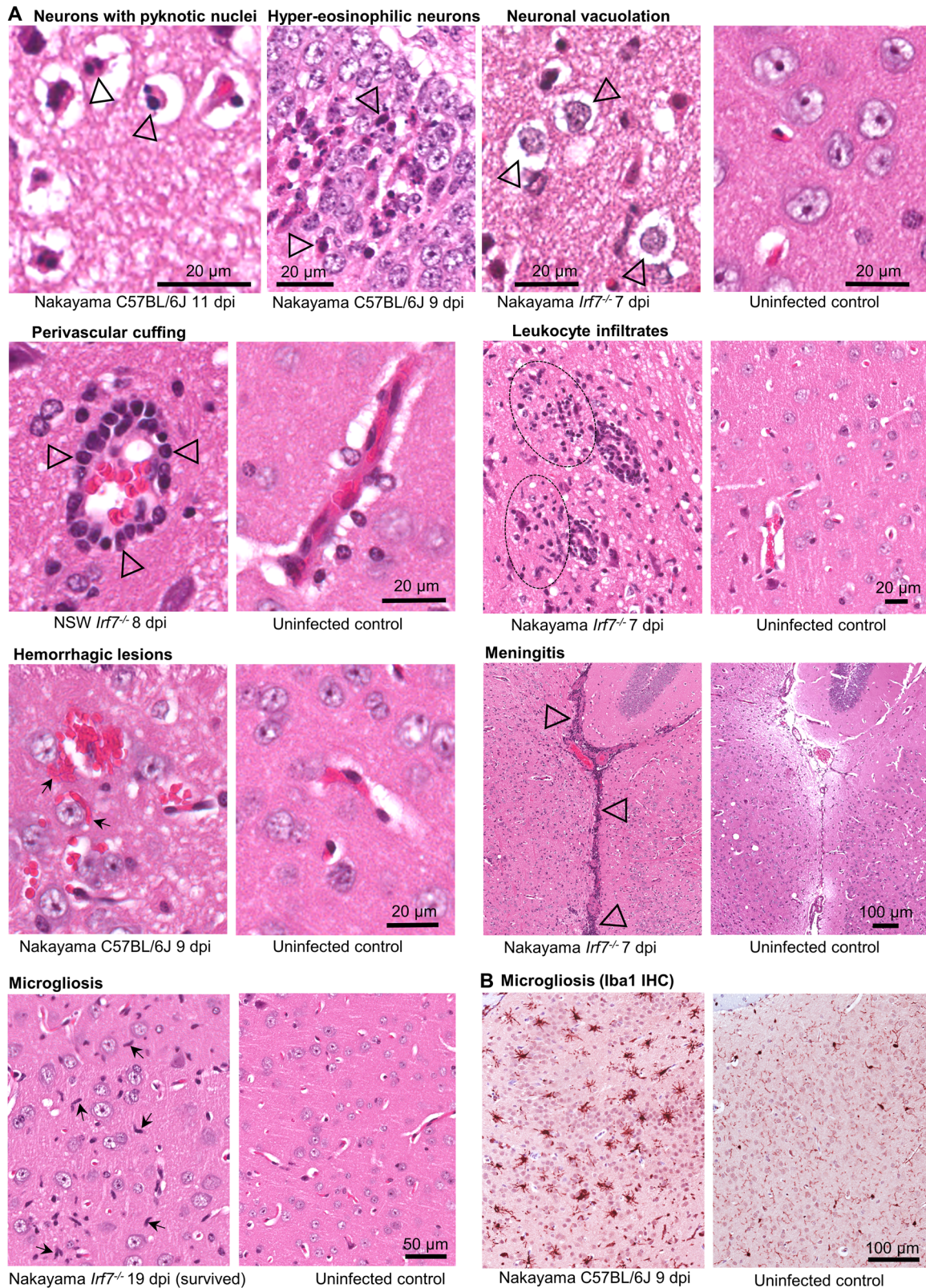


**Fig. 5 | JEV and MVEV lethal viremia in *Ifnar*<sup>-/-</sup> mice.** Female *Ifnar*<sup>-/-</sup> mice (9–24 week old) were infected s.c. with  $5 \times 10^5$  CCID<sub>50</sub> of the indicated virus ( $n = 4$  for JEV<sub>Nakayama</sub>, JEV<sub>NSW/22</sub> and MVEV<sub>TC123130</sub>, and  $n = 5$  for JEV<sub>FU</sub>). **A** Mean viremia determined by CCID<sub>50</sub> assay (limit of detection  $2 \log_{10}$  CCID<sub>50</sub>/ml). Statistics by *t* tests. **B** Mean percent body weight change compared to 0 dpi. Statistics 2 dpi JEV<sub>NSW/22</sub> versus JEV<sub>Nakayama</sub> (Kolmogorov–Smirnov test,  $p = 0.023$ ) and for JEV<sub>NSW/22</sub> versus JEV<sub>FU</sub> (*t* test,  $p = 0.008$ ). **C** Kaplan–Meier plot showing percent

survival. **D** Viral tissue titers in spleens harvested at euthanasia (3 dpi for all mice except for 1 JEV<sub>FU</sub> mouse at 2 dpi), determined by CCID<sub>50</sub> assay (limit of detection  $\sim 2 \log_{10}$  CCID<sub>50</sub>/g). **E** Viral titers in brains harvested at euthanasia. **F** IHC staining for flavivirus NS1 using the 4G4 monoclonal antibody. The brain shown was infected with JEV<sub>Nakayama</sub> (titer  $7.2 \log_{10}$  CCID<sub>50</sub>/g, 3 dpi). Staining was representative of all other JEV brains. **G, H** High magnification images from **F** showing NS1 staining in blood vessels surrounding the pale gray, biconcave shaped, red blood cells.

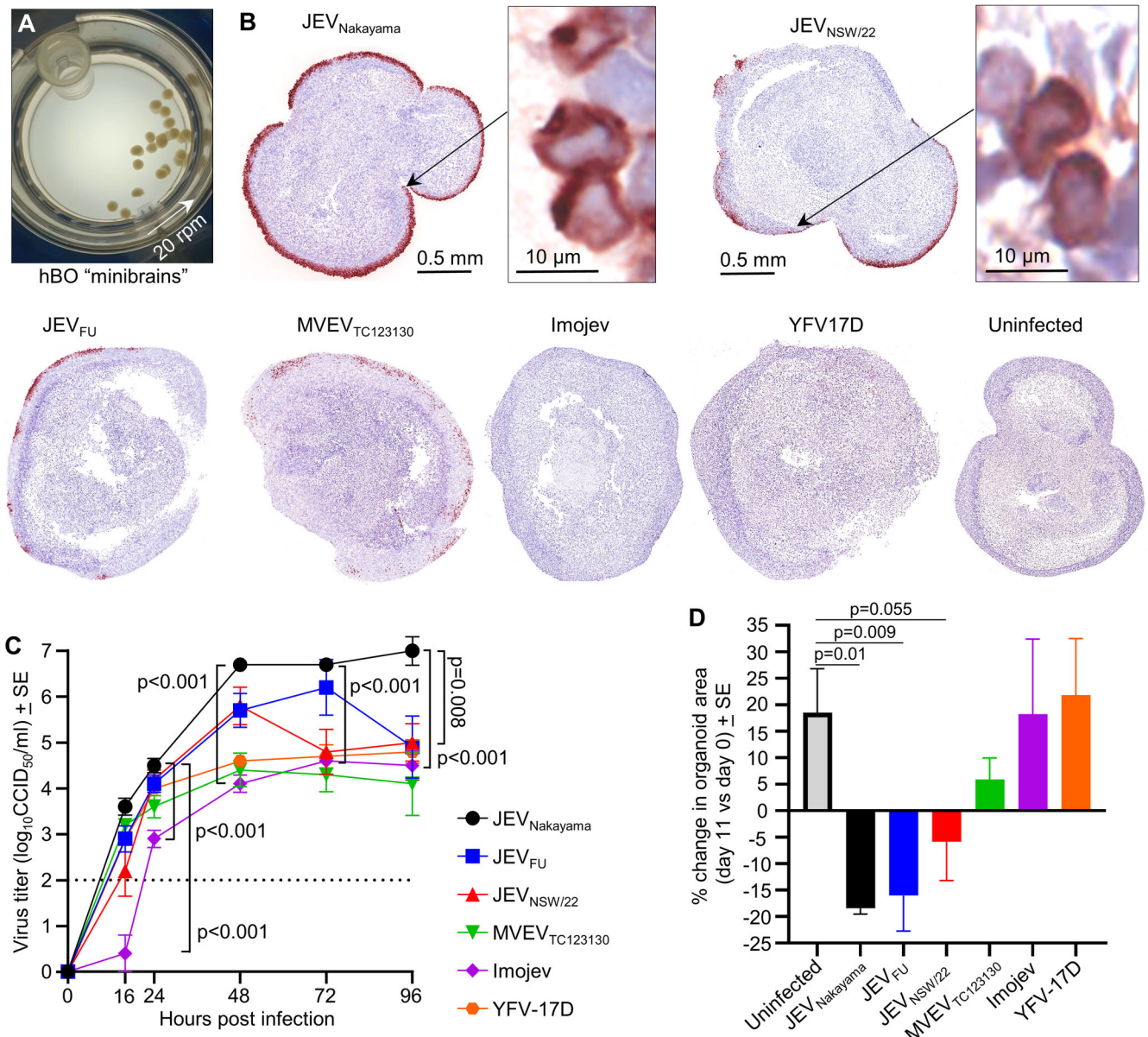
infection in mice was significantly diminished compared to JEV<sub>Nakayama</sub> and JEV<sub>FU</sub>, with only one *Irf7*<sup>-/-</sup> mouse succumbing to JEV<sub>NSW/22</sub> infection out of 63 infected C57BL/6J or *Irf7*<sup>-/-</sup> mice. Such rare lethal neuroinvasion recapitulates what is seen in humans, with  $\sim 1$  in 750 infections causing fatality<sup>2,3</sup>. Serosurvey data, albeit limited, suggests the ratio of human symptomatic to asymptomatic infections is not particularly different for

JEV<sub>NSW/22</sub>. To date, 45 clinical cases have been notified for the recent outbreak in Australia<sup>74</sup>, with serosurveys in Victoria ( $n = 820$  participants) and New South Wales (NSW) ( $n = 917$  participants) reporting 3.3% and 8.7% of participants as seropositive for JEV, respectively<sup>75,76</sup>. Although somewhat dated, population data for the primary recruitment locations for the serosurveys is available from Australian Bureau of Statistics 2016, with



**Fig. 6 | Histological lesions in brains of JEV-infected mice. A** Examples of indicated lesions stained with H&E. Degeneration of neurons indicated by (i) pyknotic nuclei (black unfilled arrowheads indicating condensation and fragmentation of nuclei, staining dark blue), and (ii) hyper-eosinophilic cytoplasm of degenerating neurons in the cornu ammonis of the hippocampus (area also staining for viral antigen Supplementary Fig. 10). Neuronal vacuolation indicated by fluid accumulation around the neurons (arrowheads). Perivascular cuffing is indicated by leukocytes aggregating in blood vessels (arrowheads). Leukocyte infiltrates

(extravascular) are indicated by dashed ovals. Hemorrhagic lesions are indicated by extravascular red blood cells (arrows). Microgliosis is indicated by accumulation of microglia, which have elongated rod-shaped nuclei (arrows). Meningitis is indicated by accumulation of leukocytes around the meninges (arrowheads). Images of uninfected controls accompany each image(s) of lesions. Histology scores for all mouse brains are shown in Supplementary Figs. 9, 10. **B** IHC using anti-Iba1, a microglial marker.



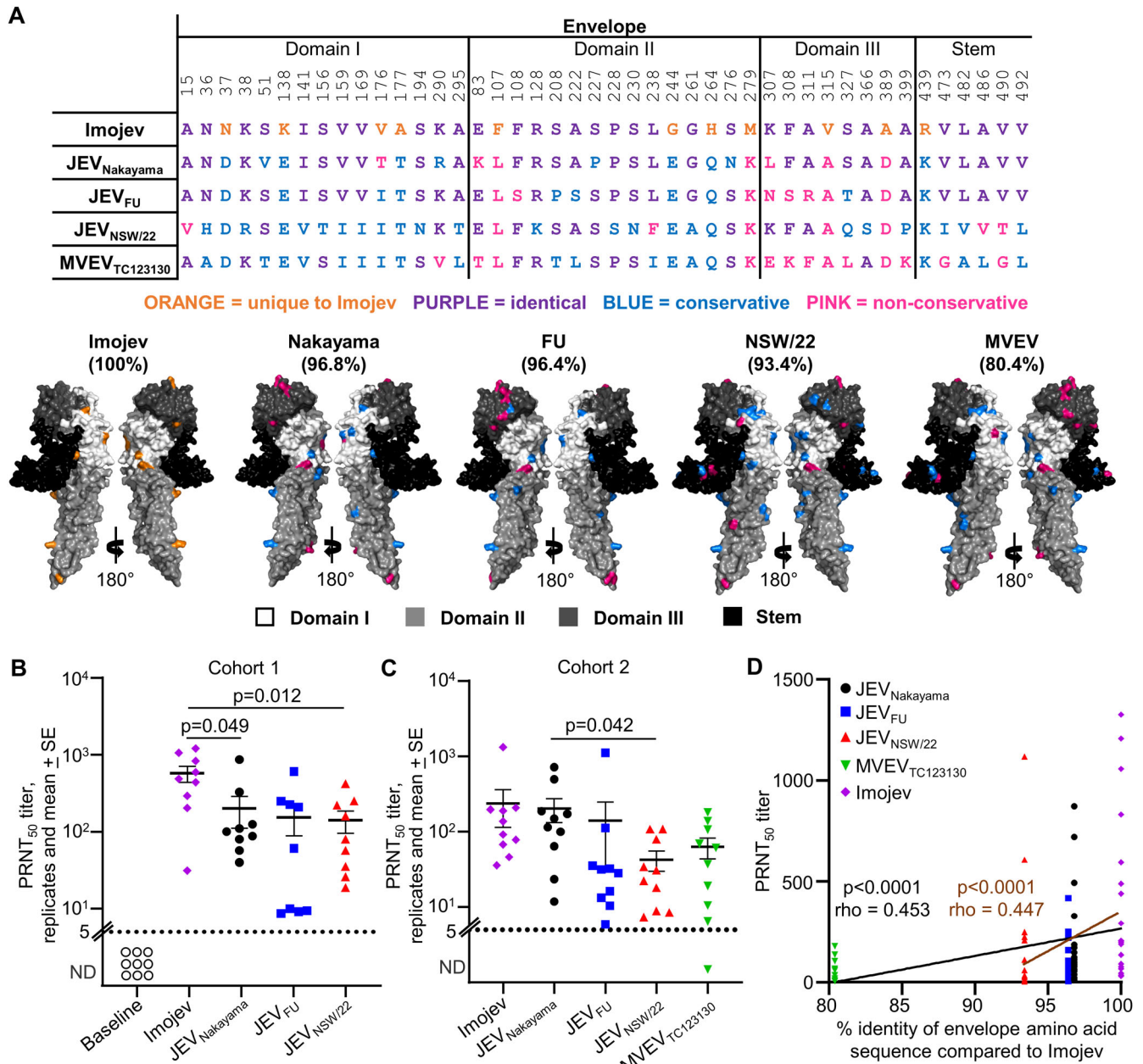
**Fig. 7 | Infection of human cortical brain organoids (hBOs).** **A** Photograph of “mini-brains” cultured in a rotating CelVivo Clinostar incubator. **B** IHC of viral antigen (4G4) for hBOs at 4 dpi. Images are representative of  $n = 4$  hBOs for each group. Magnified images of sparse IMOJEV and YFV 17D infected cells are shown in Supplementary Fig. 12A. **C** Viral growth kinetics up to 4 dpi determined by CCID<sub>50</sub> assays of culture supernatants at the indicated hours post infection; limit of detection

is  $2 \log_{10}$ CCID<sub>50</sub>/ml. At all time points JEV<sub>Nakayama</sub> vs. IMOJEV, and at 96 h JEV<sub>Nakayama</sub> vs. JEV<sub>NSW/22</sub> were significant ( $t$  tests,  $n = 5$  organoids per group). **D** Mean percentage change in organoid area at 11 dpi vs. 9 dpi for each organoid ( $n = 8$  for uninfected and MVEV<sub>TC123130</sub>, otherwise  $n = 4$ ). Statistics are by Kolmogorov–Smirnov exact test for uninfected versus JEV<sub>Nakayama</sub>, and  $t$  test for uninfected versus JEV<sub>FU</sub> or JEV<sub>NSW/22</sub>.

Victorian recruitment locations providing a population total of 160,294 (Mildura, Lockington, Shepparton, Cobram, Yarrowonga, Rutherglen, Wodonga, Wangaratta, Rochester), and NSW locations a total of 68,431 (Balranald, Corowa, Dubbo, Griffith, Temora). As  $[0.033 \times 160,294] + [0.087 \times 68,431]/250 = 45$ , the serosurvey data is consistent with the expected symptomatic to asymptomatic ratio of  $\approx 1$  in 250 for JEV and thus provides no compelling evidence for overt virulence differences for JEV<sub>NSW/22</sub> in human populations.

Our *Irf7*<sup>-/-</sup> mouse model of JEV<sub>NSW/22</sub> provides for a more robust viremia, and a slightly higher chance of lethal neuroinvasive infection. Increased lethal neuropenetrance in *Irf7*<sup>-/-</sup> mice was associated with prolonged viremia, possibly via increased inflammation-driven blood-brain barrier breakdown as a result<sup>5,77</sup>. The use of *Irf7*<sup>-/-</sup> mice to increase lethal neuroinvasive infection compared to C57BL/6J was only suitable for JEV<sub>FU</sub>

and JEV<sub>NSW/22</sub>. This was likely due to higher sensitivity to type I IFN for these isolates, demonstrated using MEF cells, compared to JEV<sub>Nakayama</sub> and MVEV<sub>TC123130</sub>. The partially defective type I IFN responses in *Irf7*<sup>-/-</sup> mice<sup>30</sup> thus provide a benefit for JEV<sub>FU</sub> and JEV<sub>NSW/22</sub> lethal neuropenetrance, but not for JEV<sub>Nakayama</sub> or MVEV<sub>TC123130</sub>. When type I IFN responses were completely defective (*Ifnar*<sup>-/-</sup> mice and *Irf3/7*<sup>-/-</sup> MEFs), differences between virus replication and/or lethality were minimal. Overall, these results suggest that JEV<sub>NSW/22</sub> may be more sensitive to, or less able to suppress, type I IFN responses. Inhibition of type I IFN responses is mediated by NS5 for inhibition of STAT2 and NS3<sup>78</sup> and subgenomic flavivirus RNA (sRNA)/NS5<sup>23</sup> for inhibition of STAT1. Although the latter likely operates for WNV in mice<sup>79</sup> and is involved in promoting apoptosis<sup>23</sup>, the efficiency of these systems during JEV infection of humans and mice remains to be determined. sRNA is derived from the 3'UTR<sup>23</sup>, where



**Fig. 8 | Human post-IMOJEV vaccination sera neutralizes JEV and MVEV with titers related to envelope protein amino acid conservation. A** Envelope protein (domains I, II, III, and STEM) amino acid sequences for IMOJEV, JEV<sub>Nakayama</sub>, JEV<sub>FU</sub>, JEV<sub>NSW/22</sub>, and MVEV<sub>TC123130</sub> (refer to Supplementary Fig. 1 for GenBank accession numbers). Sequences for isolates were aligned using MEGA-X and the ClustalW plugin with default parameters. Coloring indicates amino acid category compared to IMOJEV (orange = unique to IMOJEV, purple = identical, blue = conservative amino acid difference, pink = non-conservative amino acid difference<sup>43</sup>). Crystal structure of JEV envelope (PDB: 5WSN) with amino acid differences for JEV<sub>Nakayama</sub>, JEV<sub>FU</sub>, JEV<sub>NSW/22</sub>, and MVEV<sub>TC123130</sub> compared to IMOJEV colored as described in the table. Percentages indicate percent sequence identity relative to IMOJEV. **B** Human serum taken at day 0 and day 28 post-

IMOJEV vaccination ( $n = 9$ , cohort 1) was used in plaque reduction neutralization assays against JEV<sub>Nakayama</sub>, JEV<sub>FU</sub>, and JEV<sub>NSW/22</sub>, and the plaque reduction neutralization 50% titer (PRNT<sub>50</sub>) was calculated. Mean and standard errors are shown. Statistics are paired  $t$  test comparing IMOJEV with JEV<sub>Nakayama</sub> or JEV<sub>NSW/22</sub>. **B** Human serum taken 2–12 months post-IMOJEV vaccination ( $n = 10$ , cohort 2) was used in plaque reduction neutralization assays against JEV<sub>Nakayama</sub>, JEV<sub>FU</sub>, JEV<sub>NSW/22</sub>, and MVEV<sub>TC123130</sub> and the plaque reduction neutralization 50% titer (PRNT<sub>50</sub>) was calculated. Mean and standard errors are shown. Statistics are paired  $t$  test comparing JEV<sub>Nakayama</sub> with JEV<sub>NSW/22</sub>. **D** PRNT<sub>50</sub> titers in ‘B’ and ‘C’ plotted against percentage envelope protein amino acid identity in ‘A’. Curve fit is shown, and statistics calculated by Spearman correlation with  $p$  and  $\rho$  values shown (black line represents all data, brown line excludes MVEV<sub>TC123130</sub> from analysis).

JEV<sub>NSW/22</sub> does show a small number of nucleic acid changes, but which are unlikely to affect sRNA production (Supplementary Table 2). Mouse models of flavivirus pathogenesis frequently use *Ifnar*<sup>-/-</sup> mice<sup>25,80,81</sup> as flaviviruses often replicate poorly in wild-type mice as the ability of NS5 to suppress the antiviral type I IFN responses in humans often fails to operate in mice<sup>82</sup>. We show herein that *Ifnar*<sup>-/-</sup> mice are not a good model for JEV neuropathology, as they reach ethically defined endpoints before brain

infection can occur. However, *Ifnar*<sup>-/-</sup> mice are a good model of robust and lethal viremia, which may provide a useful and stringent model for vaccine testing.

JEV<sub>NSW/22</sub> does not contain known attenuating mutations that would explain its reduced virulence in mice. There are 9 amino acids in the IMOJEV vaccine envelope gene that have been associated with attenuated virulence; F107, K138, V176, A177, G244, H264, M279, V315, and R439<sup>83–93</sup>.

At these positions, JEV<sub>Nakayama</sub>, JEV<sub>FU</sub>, JEV<sub>NSW/22</sub>, and MVEV<sub>TC123130</sub> all have the same amino acids (L107, E138, T177, E244, Q264, K279, A315, and K439), except for position 176 where JEV<sub>Nakayama</sub> has T176, and JEV<sub>FU</sub>, JEV<sub>NSW/22</sub>, and MVEV<sub>TC123130</sub> have I176 (Supplementary Table 2). JEV<sub>NSW/22</sub> retains E at position 138, and this amino acid has been identified by several studies as a principal virulence determinant<sup>94–98</sup>, with a role in neuronal cell binding hypothesized<sup>42</sup>. YFV 17D has a valine (V) at this residue, possibly contributing to reduced virulence<sup>99</sup>. NS1 and NS2A have been implicated in JEV virulence<sup>92</sup>. Among the six changes in NS1 associated with attenuation of virulence are R147H and R339M<sup>92</sup>, of which H147 is present in both JEV<sub>Nakayama</sub> and JEV<sub>NSW/22</sub>, with K339 (a conserved substitution for R) present in JEV<sub>NSW/22</sub>. JEV non-structural protein 4B (NS4B) alone can induce apoptosis and encephalitis<sup>100</sup>, however, NS4B is completely conserved between JEV<sub>Nakayama</sub>, JEV<sub>FU</sub>, and JEV<sub>NSW/22</sub> (Supplementary Table 2). prM has been reported to influence the virulence of ZIKV<sup>80</sup> and JEV virulence in mice<sup>101</sup>. JEV<sub>NSW/22</sub> has a number of unique changes in prM (Supplementary Table 2), although their functional implications remain unclear. JEV<sub>NSW/22</sub> also has an additional N-linked glycosylation site at position 175 in NS1 that is lacking in JEV<sub>Nakayama</sub> and JEV<sub>FU</sub> (Supplementary Table 2). However, this N-linked glycosylation site is reported to increase WNV virulence in wild-type mice<sup>102,103</sup>, a trend not seen in our JEV data (Fig. 1F). Thus, JEV<sub>NSW/22</sub> shows no obvious sequence characteristics that can be readily associated with the reduced virulence in mice, and mutagenesis experiments would be required to fully understand these differences.

All human participants were vaccinated with the IMOJEV vaccine-induced neutralizing antibodies to JEV<sub>NSW/22</sub>, suggesting that this vaccine, which is available in Australia, is likely to afford some protection against Australian outbreak genotype 4 JEV. However, the divergence of envelope protein amino acid sequences from that of the IMOJEV vaccine affected the PRNT<sub>50</sub> titers, although it is unclear how this may translate to the impact on vaccine efficacy, especially given that in vitro neutralization assays do not capture the full range of protective mechanisms mobilized in vivo<sup>104</sup>. Nonetheless, this provides a strong rationale for the development of updated JEV vaccines that use antigen sequences from currently circulating JEV strains, such as genotype 4 in Australia<sup>14</sup>, genotype 5 in Republic of Korea<sup>12</sup>, and genotype 1 in most other areas of South East Asia<sup>105</sup>. IMOJEV vaccination also produced neutralizing antibodies against MVEV<sub>TC123130</sub>, which is consistent with previous studies using other JEV vaccines<sup>106,107</sup>, and is consistent with cross-reactivity in serology-based diagnostic assays<sup>13,108</sup>. There is also some evidence that JEV vaccination or infection provides partial cross-protection against MVEV and vice versa<sup>109–111</sup>.

Although one limitation of this study may be that JEV<sub>NSW/22</sub> was isolated from a pig, there are only 4 amino acid differences between this isolate and a JEV G4 sequence from a human brain in the Tiwi Islands (Northern Territory, Australia) in 2021 (Genbank accession OM867669<sup>14</sup>). The differences are; envelope-238 F vs. L, NS2A-71 I vs. T, NS2B-59 E vs. G, NS3-436 E vs. G. In addition, JEV<sub>Nakayama</sub> was passaged in suckling mouse brains, which may contribute to the increased virulence in C57BL/6J mice, although the adaptive mutations acquired during passaging, if any, are currently unknown. Furthermore, JEV<sub>FU</sub> has not been passaged in mice, but was still more lethal than JEV<sub>Nakayama</sub> in *Irf7*<sup>-/-</sup> mice. The use of C57BL/6J mice that lack a functional nicotinamide nucleotide transhydrogenase (*Nnt*) may be another issue, as background and *Nnt* are able to affect viral immunopathogenesis<sup>31,63</sup>. However, we found that neither *Nnt* nor a C57BL/6N genetic background significantly impacted JEV replication or immunopathology (Supplementary Fig. 16).

In conclusion, we show that JEV<sub>NSW/22</sub> has reduced virulence in mice but retains the capacity for rare lethal neuroinvasion, consistent with reported human fatalities in the 2022 Australian outbreak.

## Data availability

All data are provided in the manuscript and accompanying supplementary files.

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## References

- van den Hurk, A. F., Ritchie, S. A. & Mackenzie, J. S. Ecology and geographical expansion of Japanese encephalitis virus. *Annu. Rev. Entomol.* **54**, 17–35 (2008).
- Ashraf, U. et al. Pathogenicity and virulence of Japanese encephalitis virus: neuroinflammation and neuronal cell damage. *Virulence* **12**, 968–980 (2021).
- Solomon, T. et al. Japanese encephalitis. *J. Neurol. Neurosurg. Psychiatry* **68**, 405 (2000).
- Quan, T. M., Thao, T. T. N., Duy, N. M., Nhat, T. M. & Clapham, H. Estimates of the global burden of Japanese encephalitis and the impact of vaccination from 2000–2015. *eLife* **9**, e51027 (2020).
- Hsieh, J. T., Rathore, A. P. S., Soundarajan, G. & St John, A. L. Japanese encephalitis virus neuropenetrance is driven by mast cell chymase. *Nat. Commun.* **10**, 706 (2019).
- Johnson, R. T. et al. Japanese encephalitis: immunocytochemical studies of viral antigen and inflammatory cells in fatal cases. *Ann. Neurol.* **18**, 567–573 (1985).
- Hegde, N. R. & Gore, M. M. Japanese encephalitis vaccines: Immunogenicity, protective efficacy, effectiveness, and impact on the burden of disease. *Hum. Vaccin. Immunother.* **13**, 1–18 (2017).
- Gao, X. et al. Changing geographic distribution of Japanese encephalitis virus genotypes, 1935–2017. *Vector Borne Zoonotic Dis.* **19**, 35–44 (2019).
- Kang, B. K. et al. Comparison of the antigenic relationship between Japanese encephalitis virus genotypes 1 and 3. *Clin. Exp. Vaccine Res.* **5**, 26–30, (2016).
- Han, N. et al. Comparison of genotypes I and III in Japanese encephalitis virus reveals distinct differences in their genetic and host diversity. *J. Virol.* **88**, 11469–11479 (2014).
- Mulvey, P. et al. The ecology and evolution of Japanese Encephalitis Virus. *Pathogens* **10**, <https://doi.org/10.3390/pathogens10121534> (2021).
- Lee, A. R., Song, J. M. & Seo, S. U. Emerging Japanese encephalitis virus genotype V in Republic of Korea. *J. Microbiol. Biotechnol.* **32**, 955–959 (2022).
- Pham, D. et al. Emergence of Japanese encephalitis in Australia: a diagnostic perspective. *Pathology* **54**, 669–677 (2022).
- Sikazwe, C. et al. Molecular detection and characterisation of the first Japanese encephalitis virus belonging to genotype IV acquired in Australia. *PLoS Negl. Trop. Dis.* **16**, e0010754 (2022).
- Mackenzie, J. S., Williams, D. T., van den Hurk, A. F., Smith, D. W. & Currie, B. J. Japanese Encephalitis virus: the emergence of genotype IV in Australia and its potential endemicity. *Viruses* **14** (2022).
- Yakob, L. et al. Japanese encephalitis emergence in Australia: the potential population at risk. *Clin. Infect. Dis.* **76**, 335–337 (2022).
- Gyawali, N. et al. Neglected Australian arboviruses associated with undifferentiated febrile illnesses. *Front. Microbiol.* **10**, 2818 (2019).
- Furuya-Kanamori, L. et al. The emergence of Japanese Encephalitis in Australia and the implications for a vaccination strategy. *Trop. Med. Infect. Dis.* **7**, 85 (2022).
- Bharucha, T. et al. Mouse models of Japanese encephalitis virus infection: a systematic review and meta-analysis using a meta-regression approach. *PLoS Negl. Trop. Dis.* **16**, e0010116 (2022).
- Tajima, S. et al. Growth, pathogenesis, and serological characteristics of the Japanese encephalitis virus genotype IV recent strain 19CxBa-83-Cv. *Viruses* **15**, 239 (2023).
- Beasley, D. W. C. et al. Protection against Japanese encephalitis virus strains representing four genotypes by passive transfer of sera raised against ChimeriVax™-JE experimental vaccine. *Vaccine* **22**, 3722–3726 (2004).

22. Johnson, B. J. et al. Heat shock protein 10 inhibits lipopolysaccharide-induced inflammatory mediator production. *J. Biol. Chem.* **280**, 4037–4047 (2005).
23. Slonchak, A. et al. Zika virus noncoding RNA cooperates with the viral protein NS5 to inhibit STAT1 phosphorylation and facilitate viral pathogenesis. *Sci. Adv.* **8**, eadd8095 (2022).
24. Hazlewood, J. E. et al. The chimeric Binjari-Zika vaccine provides long-term protection against ZIKA virus challenge. *Vaccines (Basel)* **10**, <https://doi.org/10.3390/vaccines10010085> (2022).
25. Yan, K. et al. A yellow fever virus 17D infection and disease mouse model used to evaluate a chimeric binjari-yellow fever virus vaccine. *Vaccines* **8**, 368 (2020).
26. Nguyen, W. et al. Arthritogenic alphavirus vaccines: serogrouping versus cross-protection in mouse models. *Vaccines (Basel)* **8**, 209 (2020).
27. Ramakrishnan, M. A. Determination of 50% endpoint titer using a simple formula. *World J. Virol.* **5**, 85–86 (2016).
28. Honda, K. et al. IRF-7 is the master regulator of type-I interferon-dependent immune responses. *Nature* **434**, 772–777 (2005).
29. Sato, M. et al. Distinct and essential roles of transcription factors IRF-3 and IRF-7 in response to viruses for IFN-alpha/beta gene induction. *Immunity* **13**, 539–548 (2000).
30. Rudd, P. A. et al. Interferon response factors 3 and 7 protect against Chikungunya virus hemorrhagic fever and shock. *J. Virol.* **86**, 9888–9898 (2012).
31. Rawle, D. J. et al. Widespread discrepancy in Nnt genotypes and genetic backgrounds complicates granzyme A and other knockout mouse studies. *eLife* **11**, e70207 (2022).
32. Dumenil, T. et al. Warmer ambient air temperatures reduce nasal turbinate and brain infection, but increase lung inflammation in the K18-hACE2 mouse model of COVID-19. *Sci. Total Environ.* **859**, 160163 (2023).
33. Prow, N. A. et al. Lower temperatures reduce type I interferon activity and promote alphaviral arthritis. *PLoS Pathog.* **13**, e1006788 (2017).
34. Prow, N. A. et al. A vaccinia-based single vector construct multi-pathogen vaccine protects against both Zika and chikungunya viruses. *Nat. Commun.* **9**, 1230 (2018).
35. Hobson-Peters, J. et al. A recombinant platform for flavivirus vaccines and diagnostics using chimeras of a new insect-specific virus. *Sci. Transl. Med.* **11**, eaax7888 (2019).
36. Oikari, L. E. et al. Altered brain endothelial cell phenotype from a familial Alzheimer mutation and its potential implications for amyloid clearance and drug delivery. *Stem Cell Rep.* **14**, 924–939 (2020).
37. Stewart, R. et al. Increased neurovirulence of omicron BA.5 over BA.1 in human brain organoids and K18-hACE2 mice. *Res. Sq.* <https://www.biorxiv.org/content/10.1101/2022.12.22.521696v2> (2023).
38. Guirakhoo, F. et al. Immunogenicity, genetic stability, and protective efficacy of a recombinant, chimeric yellow fever-Japanese encephalitis virus (ChimeriVax-JE) as a live, attenuated vaccine candidate against Japanese encephalitis. *Virology* **257**, 363–372 (1999).
39. Schneider, C. A., Rasband, W. S. & Eliceiri, K. W. NIH Image to ImageJ: 25 years of image analysis. *Nat. Methods* **9**, 671–675 (2012).
40. Nasveld, P. E. et al. Long term immunity to live attenuated Japanese encephalitis chimeric virus vaccine: randomized, double-blind, 5-year phase II study in healthy adults. *Hum. Vaccin.* **6**, 1038–1046 (2010).
41. Furuya-Kanamori, L. et al. Immunogenicity of a single fractional intradermal dose of Japanese encephalitis live attenuated chimeric vaccine. *J. Travel Med.* **30**, taac122 (2023).
42. Wang, X. et al. Near-atomic structure of Japanese encephalitis virus reveals critical determinants of virulence and stability. *Nat. Commun.* **8**, 14 (2017).
43. Miyata, T., Miyazawa, S. & Yasunaga, T. Two types of amino acid substitutions in protein evolution. *J. Mol. Evol.* **12**, 219–236 (1979).
44. Williams, D. T., Wang, L. F., Daniels, P. W. & Mackenzie, J. S. Molecular characterization of the first Australian isolate of Japanese encephalitis virus, the FU strain. *J. Gen. Virol.* **81**, 2471–2480 (2000).
45. Poidinger, M., Hall, R. A. & Mackenzie, J. S. Molecular characterization of the Japanese encephalitis serocomplex of the flavivirus genus. *Virology* **218**, 417–421 (1996).
46. Williams, D. T. et al. The molecular epidemiology and evolution of murray valley encephalitis virus: recent emergence of distinct sub-lineages of the dominant genotype 1. *PLOS Negl. Trop. Dis.* **9**, e0004240 (2015).
47. Aoki, K. et al. Type-I interferon response affects an inoculation dose-independent mortality in mice following Japanese encephalitis virus infection. *Virol. J.* **11**, 105 (2014).
48. Rastogi, M., Sharma, N. & Singh, S. K. Flavivirus NS1: a multifaceted enigmatic viral protein. *Virol. J.* **13**, 131 (2016).
49. Iwasaki, Y., Zhao, J. X., Yamamoto, T. & Konno, H. Immunohistochemical demonstration of viral antigens in Japanese encephalitis. *Acta Neuropathol.* **70**, 79–81 (1986).
50. German, A. C. et al. A preliminary neuropathological study of Japanese encephalitis in humans and a mouse model. *Trans. R Soc. Trop. Med. Hyg.* **100**, 1135–1145 (2006).
51. Siva Venkatesh, I. P., Bhaskar, M. & Basu, A. Japanese encephalitis viral infection modulates proinflammatory cyto/chemokine profile in primary astrocyte and cell line of astrocytic origin. *Metab. Brain Dis.* **37**, 1487–1502 (2022).
52. Li, Y. et al. Microarray analysis identifies the potential role of long non-coding rna in regulating neuroinflammation during japanese encephalitis virus infection. *Front. Immunol.* **8**, <https://doi.org/10.3389/fimmu.2017.01237> (2017).
53. Wilhelmsson, U. et al. Redefining the concept of reactive astrocytes as cells that remain within their unique domains upon reaction to injury. *Proc. Natl. Acad. Sci.* **103**, 17513–17518 (2006).
54. Zhang, Z. et al. The appropriate marker for astrocytes: comparing the distribution and expression of three astrocytic markers in different mouse cerebral regions. *BioMed Res. Int.* **2019**, 9605265 (2019).
55. Daffis, S. et al. Interferon regulatory factor IRF-7 induces the antiviral alpha interferon response and protects against lethal West Nile virus infection. *J. Virol.* **82**, 8465–8475 (2008).
56. Roberts, A., Kesarwani, V., Gupta, R. & Gandhi, S. Electroactive reduced graphene oxide for highly sensitive detection of secretory non-structural 1 protein: a potential diagnostic biomarker for Japanese encephalitis virus. *Biosens. Bioelectron.* **198**, 113837 (2022).
57. Puerta-Guardo, H. et al. Flavivirus NS1 triggers tissue-specific vascular endothelial dysfunction reflecting disease tropism. *Cell Rep.* **26**, 1598–1613.e1598 (2019).
58. Baccala, R. et al. Type I interferon is a therapeutic target for virus-induced lethal vascular damage. *Proc. Natl. Acad. Sci.* **111**, 8925–8930 (2014).
59. Iannacone, M. et al. Platelets prevent IFN- $\alpha$ / $\beta$ -induced lethal hemorrhage promoting CTL-dependent clearance of lymphocytic choriomeningitis virus. *Proc. Natl. Acad. Sci.* **105**, 629–634 (2008).
60. Miyake, M. The pathology of Japanese encephalitis. a review. *Bull. World Health Organ.* **30**, 153–160 (1964).
61. Waller, C. et al. Japanese Encephalitis in Australia — a sentinel case. *N. Engl. J. Med.* **387**, 661–662 (2022).
62. Maamary, J. et al. New detection of locally acquired Japanese Encephalitis virus using clinical metagenomics, New South Wales, Australia. *Emerg. Infect. Dis. J.* **29**, 627 (2023).
63. Bishop, C. R. et al. Mouse models of COVID-19 recapitulate inflammatory pathways rather than gene expression. *PLoS Pathog.* **18**, e1010867 (2022).



64. Tripathi, A. et al. Lack of interferon (IFN) regulatory factor 8 associated with restricted IFN-gamma response augmented Japanese encephalitis virus replication in the mouse brain. *J Virol* **95**, e0040621 (2021).
65. Danaher, P. et al. Advances in mixed cell deconvolution enable quantification of cell types in spatial transcriptomic data. *Nat. Commun.* **13**, 385 (2022).
66. Setoh, Y. X. et al. Determinants of Zika virus host tropism uncovered by deep mutational scanning. *Nat. Microbiol.* **4**, 876–887 (2019).
67. Zhang, B. et al. Differential antiviral immunity to Japanese encephalitis virus in developing cortical organoids. *Cell Death Dis.* **9**, 719 (2018).
68. Chambers, T. J., Nestorowicz, A., Mason, P. W. & Rice, C. M. Yellow fever/Japanese encephalitis chimeric viruses: construction and biological properties. *J. Virol.* **73**, 3095–3101 (1999).
69. Frassetto, F. P. & Roseberg, S. Neuropathology of yellow fever autopsy cases. *Trop. Dis. Travel Med. Vaccines* **9**, 1 (2023).
70. Cohen, M. et al. Case report: yellow fever vaccine-associated neurotropic disease and associated MRI, EEG, and CSF findings. *Front Neurol* **12**, 779014 (2021).
71. Pierson, T. C., Fremont, D. H., Kuhn, R. J. & Diamond, M. S. Structural insights into the mechanisms of antibody-mediated neutralization of flavivirus infection: implications for vaccine development. *Cell Host Microbe* **4**, 229–238 (2008).
72. Hombach, J., Solomon, T., Kurane, I., Jacobson, J. & Wood, D. Report on a WHO consultation on immunological endpoints for evaluation of new Japanese encephalitis vaccines, WHO, Geneva, 2–3 September, 2004. *Vaccine* **23**, 5205–5211 (2005).
73. Myint, K. S. et al. Neuropathogenesis of Japanese encephalitis in a primate model. *PLoS Negl. Trop. Dis.* **8**, e2980 (2014).
74. Japanese encephalitis virus (JEV). Australian Government Dept of Health and Aged Care. <https://www.health.gov.au/health-alerts/japanese-encephalitis-virus-jev/japanese-encephalitis-virus-jev> (accessed April 2023).
75. Serosurvey for JEV in northern Victoria. <https://www.health.vic.gov.au/infectious-diseases/serosurvey-for-japanese-encephalitis-in-northern-victoria> (accessed April 2023).
76. Summary of NSW JEV serosurvey results. <https://www.health.nsw.gov.au/environment/pests/vector/Documents/jev-serosurvey-report.pdf> (accessed April 2023).
77. Hsieh, J. T. & St John, A. L. Japanese encephalitis virus and its mechanisms of neuroinvasion. *PLoS Pathog.* **16**, e1008260 (2020).
78. Setoh, Y. X. et al. Helicase domain of west Nile virus NS3 protein plays a role in inhibition of type I interferon signalling. *Viruses* **9**, 326 (2017).
79. Laurent-Rolle, M. et al. The NS5 protein of the virulent West Nile virus NY99 strain is a potent antagonist of type I interferon-mediated JAK-STAT signaling. *J. Virol.* **84**, 3503–3515 (2010).
80. Nakayama, E. et al. Neuroinvasiveness of the MR766 strain of Zika virus in IFNAR<sup>-/-</sup> mice maps to prM residues conserved amongst African genotype viruses. *PLoS Pathog.* **17**, e1009788 (2021).
81. Zheng, W. et al. Liver transcriptomics reveals features of the host response in a mouse model of dengue virus infection. *Front. Immunol.* **13**, 892469 (2022).
82. Grant, A. et al. Zika virus targets human STAT2 to inhibit type I interferon signaling. *Cell Host Microbe* **19**, 882–890 (2016).
83. Yang, H. et al. A novel amino acid site closely associated with the neurovirulence of live, attenuated Japanese encephalitis vaccine (SA14-14-2 strain). *Vaccine* **38**, 2636–2642 (2020).
84. Yun, S. I. et al. A molecularly cloned, live-attenuated Japanese encephalitis vaccine SA14-14-2 virus: a conserved single amino acid in the ij Hairpin of the Viral E glycoprotein determines neurovirulence in mice. *PLoS Pathog.* **10**, e1004290 (2014).
85. Arroyo, J. et al. Molecular basis for attenuation of neurovirulence of a yellow fever Virus/Japanese encephalitis virus chimera vaccine (ChimeriVax-JE). *J. Virol.* **75**, 934–942 (2001).
86. Yang, D. et al. Characterization of live-attenuated Japanese encephalitis vaccine virus SA14-14-2. *Vaccine* **32**, 2675–2681 (2014).
87. Liu, X. et al. The structure differences of Japanese encephalitis virus SA14 and SA14-14-2 E proteins elucidate the virulence attenuation mechanism. *Protein Cell* **10**, 149–153 (2019).
88. Yang, J. et al. Envelope protein mutations L107F and E138K are important for neurovirulence attenuation for Japanese encephalitis virus SA14-14-2 strain. *Viruses* **9**, 20 (2017).
89. Ni, H., Chang, G. J., Xie, H., Trent, D. W. & Barrett, A. D. Molecular basis of attenuation of neurovirulence of wild-type Japanese encephalitis virus strain SA14. *J. Gen. Virol.* **76**, 409–413 (1995).
90. Zhou, Y. et al. Mutation of I176R in the E coding region weakens Japanese encephalitis virus neurovirulence, but not its growth rate in BHK-21 cells. *Arch. Virol.* **163**, 1351–1355 (2018).
91. Anwar, M. N. et al. Phenotypic and genotypic comparison of a live-attenuated genotype I Japanese encephalitis virus SD12-F120 strain with its virulent parental SD12 strain. *Viruses* **12**, 552 (2020).
92. Gromowski, G. D., Firestone, C. Y. & Whitehead, S. S. Genetic determinants of Japanese encephalitis virus vaccine strain SA14-14-2 that govern attenuation of virulence in mice. *J. Virol.* **89**, 6328–6337 (2015).
93. Monath, T. P. et al. Single mutation in the flavivirus envelope protein hinge region increases neurovirulence for mice and monkeys but decreases viscerotropism for monkeys: relevance to development and safety testing of live, attenuated vaccines. *J. Virol.* **76**, 1932–1943 (2002).
94. Chen, L. K. et al. Generation and characterization of organ-tropism mutants of Japanese encephalitis virus in vivo and in vitro. *Virology* **223**, 79–88 (1996).
95. Sumiyoshi, H., Tignor, G. H. & Shope, R. E. Characterization of a highly attenuated Japanese encephalitis virus generated from molecularly cloned cDNA. *J. Infect. Dis.* **171**, 1144–1151 (1995).
96. Zheng, X. et al. Acidity/alkalinity of Japanese encephalitis virus E protein residue 138 alters neurovirulence in mice. *J. Virol.* **92**, e00108–e00118 (2018).
97. Chiou, S. S. & Chen, W. J. Phenotypic changes in the Japanese encephalitis virus after one passage in Neuro-2a cells: generation of attenuated strains of the virus. *Vaccine* **26**, 15–23 (2007).
98. Zhao, Z. et al. Characterization of the E-138 (Glu/Lys) mutation in Japanese encephalitis virus by using a stable, full-length, infectious cDNA clone. *J. Gen. Virol.* **86**, 2209–2220 (2005).
99. Xia, Q. et al. Virulence and cross-protection conferred by an attenuated genotype i-based chimeric Japanese encephalitis virus strain harboring the E protein of genotype V in mice. *Microbiol. Spectr.* **10**, e01990–01922 (2022).
100. Wang, Q. et al. Japanese encephalitis virus induces apoptosis and encephalitis by activating the PERK pathway. *J. Virol.* **93**, <https://doi.org/10.1128/jvi.00887-19> (2019).
101. Tajima, S. et al. E and prM proteins of genotype V Japanese encephalitis virus are required for its increased virulence in mice. *Heliyon* **5**, e02882 (2019).
102. Whiteman, M. C. et al. Multiple amino acid changes at the first glycosylation motif in NS1 protein of West Nile virus are necessary for complete attenuation for mouse neuroinvasiveness. *Vaccine* **29**, 9702–9710 (2011).
103. Whiteman, M. C. et al. Development and characterization of non-glycosylated E and NS1 mutant viruses as a potential candidate vaccine for West Nile virus. *Vaccine* **28**, 1075–1083 (2010).
104. Burton, D. R. Antiviral neutralizing antibodies: from in vitro to in vivo activity. *Nat. Rev. Immunol.* **23**, 720–734 (2023).

105. Li, F. et al. Tracing the spatiotemporal phylodynamics of Japanese encephalitis virus genotype I throughout Asia and the western Pacific. *PLoS Negl. Trop. Dis.* **17**, e0011192 (2023).
106. Broom, A. K., Wallace, M. J., Mackenzie, J. S., Smith, D. W. & Hall, R. A. Immunisation with gamma globulin to murray valley encephalitis virus and with an inactivated Japanese encephalitis virus vaccine as prophylaxis against australian encephalitis: evaluation in a mouse model. *J. Med. Virol.* **61**, 259–265 (2000).
107. Bielefeldt-Ohmann, H. et al. Safety and immunogenicity of a delta inulin-adjuvanted inactivated Japanese encephalitis virus vaccine in pregnant mares and foals. *Vet. Res.* **45**, 130 (2014).
108. Rawle, D. J. et al. Sequencing of historical isolates, k-mer mining and high serological cross-reactivity with ross river virus argue against the presence of Getah virus in Australia. *Pathogens* **9**, <https://doi.org/10.3390/pathogens9100848> (2020).
109. Lobigs, M., Pavy, M. & Hall, R. Cross-protective and infection-enhancing immunity in mice vaccinated against flaviviruses belonging to the Japanese encephalitis virus serocomplex. *Vaccine* **21**, 1572–1579 (2003).
110. Williams, D. T. et al. Experimental infections of pigs with Japanese encephalitis virus and closely related Australian flaviviruses. *Am. J. Trop. Med. Hyg.* **65**, 379–387 (2001).
111. Lobigs, M., Larena, M., Alsharifi, M., Lee, E. & Pavy, M. Live chimeric and inactivated Japanese encephalitis virus vaccines differ in their cross-protective values against Murray Valley encephalitis virus. *J. Virol.* **83**, 2436–2445 (2009).

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## Author contributions

Conceptualization: D.J.R. Methodology: D.J.R., W.N., A.S., G.D., N.G., and N.W. Formal analysis: W.N., D.J.R., and A.S. Investigation: W.N., B.T., R.S., A.C., K.Y., N.G., T.L., and D.J.R. Resources: R.S., N.G., G.D., N.W., and A.S. Data curation: W.N., D.J.R., N.G., and A.S. Writing, original draft—D.J.R. Writing, review and editing: A.S. and W.N. Visualization: D.J.R., W.N., and A.S. Supervision: D.J.R., G.D. and A.S. Project administration: D.J.R., G.D., N.W., and A.S. Funding acquisition: A.S., D.J.R., and G.D., and N.W.

## Competing interests

The authors declare no competing interests.

## Additional information

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**Correspondence** and requests for materials should be addressed to Daniel J. Rawle.

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