779 STAR METHODS

780 CONTACT FOR REAGENT AND RESOURCE SHARING.

781 Requests should be addressed and will be fulfilled by lead author Michael S. Diamond;
782 diamond@wusm.wustl.edu.

783

784 EXPERIMENTAL MODEL AND SUBJECT DETAILS

785 Ethics statement. This study was carried out in strict accordance with the 786 recommendations in the Guide for the Care and Use of Laboratory Animals of the National 787 Institutes of Health. The protocols were approved by the Institutional Animal Care and Use 788 Committee (IACUC) at the Washington University School of Medicine (Assurance Number: 789 A3381-01), the IACUC at the La Jolla Institute for Allergy & Immunology under protocol # 790 AP028-SS1-0615, and the IACUC at Utah State University under protocol 2598. Dissections 791 and footpad injections were performed under anesthesia that was induced and maintained with 792 ketamine hydrochloride and xylazine, and all efforts were made to minimize suffering.

793 Mouse experiments. BALB/c and C57BL/6 mice were purchased from The Jackson 794 Laboratory, and AG129 mice were bred in the animal facilities at Utah State University, 795 Washington University, or the La Jolla Institute for Allergy and Immunology. All mice were 796 housed in pathogen-free mouse facilities. For immunizations, mice were inoculated via an intramuscular route with 50 µl of the indicated vaccine LNP constructs. For challenge studies, 797 mice were inoculated subcutaneously with 10⁴ PFU of ZIKV P6-740 or 10⁶ FFU of mouse-798 adapted ZIKV Dakar 41519 in 50 µl of HBSS + 0.1% FBS. For ZIKV challenge studies in 799 800 BALB/c and C57BL/6 mice, 2 mg of anti-IFNAR1 blocking antibody (MAR1-5A3 (Sheehan et al., 801 2006)) was administered via intraperitoneal injection 24 hours prior to viral infection. For DENV 802 challenge studies in AG129 mice, animals were passively transferred pooled vaccine immune sera or PBS one day prior to infection with ~10⁵ FFU of DENV-2 S221. Animals were monitored 803

for mortality and clinical score (1 = heathy; 2 = slightly ruffled fur; 3 = very ruffled fur; 4 = mild
lethargy, decreased scurrying activity; 5 = very sick, slow to no movement; 6 = very sick,
euthanize (in distress); 7 = deceased) as described previously (Tang et al., 2016).

807

828

808 METHOD DETAILS

809 Viruses and cells. ZIKV strain Dakar 41519 (Senegal, 1984) and P6-740 (Malaysia, 810 1966) were provided by the World Reference Center for Emerging Viruses and Arboviruses (R. 811 Tesh and S. Weaver, University of Texas Medical Branch). To create a mouse-adapted more pathogenic variant of ZIKV Dakar 41519, it was passaged twice in Rag1^{-/-} mice (Sapparapu et 812 813 al., 2016; Zhao et al., 2016). ZIKV strain Paraiba 2015 (Brazil) was provided by S. Whitehead 814 (NIH, Bethesda, MD) (Tsetsarkin et al., 2016). DENV-2 strain S221 is a mouse-adapted strain 815 that has been described previously (Yauch et al., 2009). Virus stocks were propagated in 816 mycoplasma-free Vero cells and titrated by focus-forming assay (FFA), as described previously 817 (Lazear et al., 2016). Experiments with ZIKV and DENV were conducted under biosafety level 2 818 (BSL2) containment at Washington University School of Medicine or under BSL3 containment at 819 Utah State University with Institutional Biosafety Committee approval.

820 Generation of modified mRNA and LNP The mRNA was synthesized in vitro using T7 821 polymerase-mediated DNA-dependent RNA transcription where the UTP was substituted with 1-822 methylpseudoUTP, using a linearized DNA template, which incorporates 5' and 3' untranslated 823 (UTRs) includes tail regions and а poly-A (5'-UTR: 824 TCAAGCTTTTGGACCCTCGTACAGAAGCTAATACGACTCACTATAGGGAAATAAGAGAGAA 825 AAGAAGAGTAAGAAGAAATATAAGAGCCACC; 3'-UTR: and 826 TGATAATAGGCTGGAGCCTCGGTGGCCATGCTTCTTGCCCCTTGGGCCTCCCCCAGCCC 827 CTCCTCCCCTTCCTGCACCCGTACCCCCGTGGTCTTTGAATAAAGTCTGAGTGGGCGGC).

30

A donor methyl group S-adenosylmethionine (SAM) was added to the methylated capped RNA

(cap 0), resulting in a cap 1 structure to increase mRNA translation efficiency. The modified
mRNAs encoded the signal sequences from human IgE (MDWTWILFLVAAATRVHS) or JEV
prM (MWLVSLAIVTACAGA) and the prM and E genes from an Asian ZIKV strain (Micronesia
2007, GenBank accession number EU545988 (Lanciotti et al., 2008)), which is >99% identical
to circulating American strains.

834 LNP formulations were prepared using a modified procedure of a method described for 835 siRNA (Chen et al., 2016). Briefly, lipids were dissolved in ethanol at molar ratios of 836 50:10:38.5:1.5 (ionizable lipid: DSPC: cholesterol: PEG-lipid). The lipid mixture was combined 837 with a 50 mm citrate buffer (pH 4.0) containing mRNA at a ratio of 3:1 (aqueous:ethanol) using a 838 microfluidic mixer (Precision Nanosystems, Vancouver, BC). Formulations were dialyzed 839 against PBS (pH 7.4) in dialysis cassettes for at least 18 h. Formulations were concentrated 840 using Amicon Ultra Centrifugal Filters (EMD Millipore, Billerica, MA), passed through a 0.22-µm filter and stored at 4°C until use. All formulations were tested for particle size, RNA 841 842 encapsulation, and endotoxin and were found to be between 80 to 100 nm in size, with greater 843 than 90% encapsulation and <1 EU/ml of endotoxin.

844 Viral protein analysis. SVPs were floated on a 10-50% sucrose gradient (50 mM MES pH 5.5, 120 mM NaCl) and ultracentrifuged at 32,000 rpm at 4℃ for 4 h in a Beckman SW55 845 846 rotor. Fractions (1 ml) were collected and pelleted by a second ultracentrifugation step over a 847 20% sucrose cushion at 32,000 rpm at 4℃ for 2 h. Protein concentration of SVPs was 848 measured using a spectrophotometer (280 nm wavelength). Western blotting was performed 849 using NuPAGE MES Western blot system (Thermo Fisher) reagents as per the manufacturer's instructions. Samples were boiled at 100°C for 10 min in the absence of reducing agent. 850 851 Samples (100 or 500 ng total protein) were loaded on 4-12% NuPAGE gradient gel and 852 electrophoresed. Proteins were transferred onto nitrocellulose membranes using iBlot 2 gel 853 transfer system. Membranes were washed three times with deionized water and blocked in PBS 854 with 5% Blotto (Thermo Fisher) overnight at 4°C. Primary mAbs (WNV E60 (Oliphant et al.,

855 2006) or mouse anti-ZIKV E (Biofront, BF-1176-86)) were added at 1 µg/ml in PBS 856 supplemented with 5% Blotto and 0.2% (v/v) Tween 20 (Sigma) and incubated for 2 h at room 857 temperature with agitation. Membranes were washed three times with PBS supplemented with 858 0.2% Tween 20. Secondary antibody (horseradish peroxidase conjugated goat anti-mouse IgG 859 (Ab97023, Abcam)) was added at 0.25 µg/ml in PBS supplemented with 5% Blotto and 0.2% 860 Tween 20 for 2 h at room temperature with agitation. After final washing, blots were developed 861 using Amersham ECL prime solution (GE Healthcare Life Sciences) for 2 min and imaged. 862 SeeBlue plus 2 pre-stained protein ladders were included for molecular weight references.

To compare expression of SVPs from different mRNA vaccine constructs, HeLa cells
were transfected with 1.25 μg of mRNA constructs using Lipofectamine 2000 (Thermo Fisher),
and cells were harvested 24 h later. Cell lysates were prepared in RIPA lysis buffer (Thermo
Fisher) with phosphatase and protease inhibitors (Millipore) added.

Electron microscopy of SVPs. The SVPs were imaged by electron microscopy and negative staining using a fee-for-service facility (University of California, Los Angeles). Briefly, purified SVPs (2.5 μl) were applied to a Glow-discharge carbon-coated grid (Ted Pella Inc.). Staining (2% uranyl acetate) was added in a drop-wise manner for 60 sec. After blotting of excess liquid and drying, the images were collected on a FEI Tecnai TF20 transmission electron microscope at an accelerating voltage of 200 kV using TVIPS EM-Menu program. The nominal magnifications used were 50,000, 29,000 and 14,500 with 2 binning.

Measurement of viral burden. At specified time points after ZIKV challenge, blood was collected and organs were recovered. Organs were weighed and homogenized using a beadbeater apparatus (MagNA Lyser, Roche), and serum was prepared after coagulation and centrifugation. Tissue samples and serum from ZIKV-infected mice were extracted with the RNeasy Mini Kit (Qiagen). ZIKV RNA levels were determined by TaqMan one-step quantitative reverse transcriptase PCR (qRT-PCR) on an ABI 7500 Fast Instrument using standard cycling

conditions. Viral burden is expressed on a log₁₀ scale as viral RNA equivalents per gram or per
milliliter after comparison with a standard curve produced using serial 5-fold dilutions of ZIKV
RNA from known quantities of infectious virus. For ZIKV, the following primer sets were used:
1183F: 5'-CCACCAATGTTCTCTTGCAGACATATTG-3'; 1268R: 5'TTCGGACAGCCGTTGTCCAACACACAAG-3'; and probes (1213F): 5'-56-FAM/AGCCTACCT
TGACAAGCAGTC/3IABkFQ-3'.

886 Neutralization assays. (a) PRNT or FRNT assays. Serial dilutions of heat-inactivated 887 sera obtained from AG129 or C57BL/6 mice were incubated with 50 to 100 FFU of ZIKV 888 (Paraiba, Brazil 2015) for 1 h at 37 °C. The serum Ab-virus complexes were added to Vero cell 889 monolayers in 96-well plates for 60 min at 37°C. Pl aque assays were performed as described 890 previously (Brien et al., 2013; Lazear et al., 2016) For FRNT assays, cells were overlaid with 1% 891 (w/v) methylcellulose in MEM supplemented with 4% heat-inactivated FBS. Plates were fixed 892 40 h later with 1% PFA in PBS for 1 h at room temperature. The plates were incubated 893 sequentially with 500 ng/ml of humanized anti-WNV E60 (Oliphant et al) and horseradish-894 peroxidase-conjugated goat anti-human IgG in PBS supplemented with 0.1% (w/v) saponin 895 (Sigma) and 0.1% BSA. ZIKV-infected cell foci were visualized using TrueBlue peroxidase 896 substrate (KPL) and quantitated on an ImmunoSpot 5.0.37 macroanalyzer (Cellular 897 Technologies). (b) RVP assays. RVPs incorporating the structural proteins of ZIKV or DENV 898 were produced by complementation of a previously described sub-genomic GFP-expressing 899 replicon derived from a lineage II strain of WNV (Dowd et al., 2016a; Dowd et al., 2015). Serial 900 dilutions of heat-inactivated sera obtained from BALB/c mice were mixed with ZIKV (strain 901 H/PF/2013; French Polynesia, 2013) or DENV-2 (strain 16681) reporter viral particles (RVPs) 902 and incubated for 1 h at 37°C. Immune complexes were added in duplicate technical replicates 903 to pre-plated Vero cells in a 96-well plate and incubated for two days. Cells were trypsinized, 904 resuspended in 4% PFA in PBS, and RVP infection scored as a function of GFP expression by 905 flow cytometry. All neutralization data were analyzed by non-linear regression to determine the

dilution of sera required to inhibit 50% (EC50) and 90% (EC90) of infection. RVP studies were
performed starting at an initial serum dilution of 1:100 (based on the final volume of cells, virus,
and sera per well), which was designated as the limit of detection.

909 ADE assays. Serial dilutions of heat-inactivated sera obtained from BALB/c mice were 910 mixed with DENV-1 RVPs (Western Pacific-74 strain) and incubated for 1 h at 37°C. Immune 911 complexes were added in duplicate technical replicates to K562 cells that express the $Fc-\gamma$ 912 receptor CD32A and incubated for two days. Due to limited volumes of sera, a small number of 913 samples (four) could not be performed in duplicate. Cells were fixed with 2% PFA, and RVP 914 infection was scored as a function of GFP expression by flow cytometry. To normalize the 915 magnitude of enhancement across independent experiments, results are displayed relative to 916 the maximum infectivity observed with a control cross-reactive WNV mAb E60 (Oliphant et al., 917 2006) run in parallel. ADE studies were performed starting at an initial serum dilution of 1:60 918 (based on the final volume of cells, virus, and sera per well), which was designated as the limit 919 of detection. For calculations of peak enhancing titer, samples for which no enhancement 920 (infectivity) was observed are reported as a titer of 30 (one half the limit of detection).

921

922 **QUANTIFICATION AND STATISTICAL ANALYSIS.** All data were analyzed with GraphPad 923 Prism software. Kaplan-Meier survival curves were analyzed by the log rank test, and weight 924 losses were compared using two-way ANOVA. For neutralization antibody titers and viral 925 burden analysis, the log titers and levels of viral RNA were analyzed by a Kruskal-Wallis 2-way 926 ANOVA with a multiple comparisons correction. A *P* value of < 0.05 indicated statistically 927 significant differences.

928

929 DATA AND SOFTWARE AVAILABILITY. All data is available upon request to the lead contact
930 author. No proprietary software was used in the data analysis.

931

ADDITIONAL RESOURCES. mRNA LNP vaccines are available from Valera/Moderna upon 932 933 request and completion of appropriate Material Transfer Agreements.

- 934
- 935

SUPPLEMENTAL TITLES AND LEGENDS 936

- 937 938

939 SUPPLEMENTAL FIGURE LEGENDS

Figure S1. Production of LNPs for vaccination, Related to Figure 1. A. 940 941 Schematic representation of the process to encapsulate mRNA into LNPs. B. A 942 representative cryo-electron microscopy image of an LNP solution, following mRNA 943 encapsulation.

944 Figure S2. Serum neutralization curves from IgE_{sig}-prM-E vaccinated AG129 945 mice, Related to Figure 1. Ten AG129 mice in each group (combined from two 946 independent experiments) were immunized with 10 (Groups 1 and 2, panels A-B) or 2 947 (Groups 3 and 4, panels C-D) µgs of IgE_{sig} prM-E mRNA LNPs. Some of the mice 948 (Groups 1 and 3) were boosted with an equivalent dose 21 days later. Serum was 949 collected at 6 weeks (day 42) post initial vaccination and analyzed for ZIKV 950 neutralization activity by PRNT. Each line represents the neutralization curve from an 951 individual mouse.

Figure S3. Serum neutralization curves from IgE_{sig} prM-E vaccinated 952 C57BL/6 mice, Related to Figure 2. Ten C57BL/6 mice in each group (combined from 953 two independent experiments) were immunized with 10 µg of IgE_{sia}-prM-E mRNA LNPs 954 955 and boosted with an equivalent dose four weeks later. Serum was collected at 4 (B), 8 956 (C), and 18 (D) weeks post initial vaccination and analyzed for ZIKV neutralization

activity by FRNT. Serum from mice immunized with placebo mRNA LNPs also were
analyzed (A) Each line represents the neutralization curve from an individual mouse.
Error bars indicate the standard deviation (SD) of triplicate technical replicates.

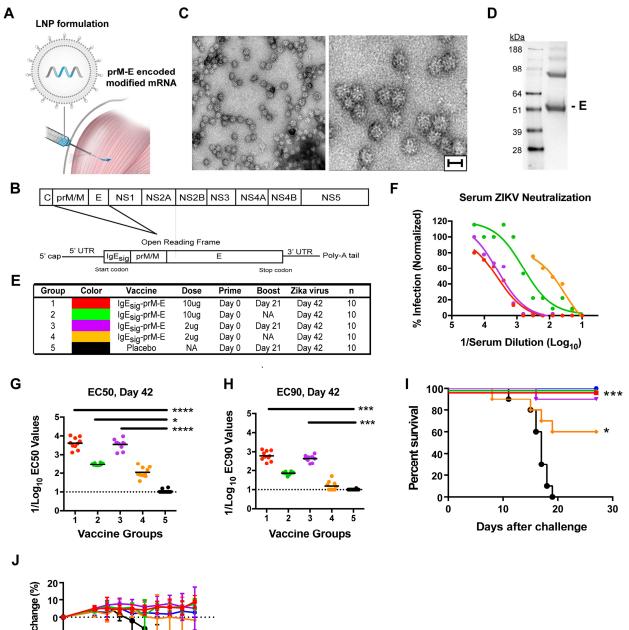
Figure S4. Serum neutralization curves from LNP vaccines containing WT 960 or mutant FL sequences, Related to Figure 3. Ten BALB/c mice in each group 961 962 (combined from two independent experiments) were immunized with 2 or 10 µg of prM-963 E mRNA LNP vaccines containing IgE or JEV signal sequences and WT or mutant 964 fusion loop (FL) sequences or placebo mRNA LNPs (Groups 1-9, A-I). Animals were 965 boosted with the equivalent dose of the same vaccine 4 weeks later. At week 8, serum 966 was harvested and analyzed for ZIKV neutralization capacity by incubating serial 967 dilutions of serum with GFP-expressing ZIKV RVPs, followed by infection of Vero cells. Infected cells were quantified 2 days later by flow cytometry. Each curve represents the 968 969 data from an individual mouse analyzed by non-linear regression analysis. Error bars 970 indicate the range of duplicate technical replicates.

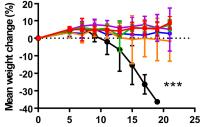
Figure S5. ADE curves from LNP vaccines containing WT or FL mutant 971 sequences, Related to Figure 4. Ten BALB/c mice in each group (combined from two 972 973 independent experiments) were immunized with 2 or 10 µg of prM-E mRNA LNP 974 vaccines containing IgE or JEV signal sequences and WT or mutant FL sequences or 975 placebo mRNA LNPs (Groups 1-9, A-I). Animals were boosted with the equivalent dose 976 of the same vaccine 4 weeks later. At week 8, serum was harvested from vaccinated 977 mice. Serial dilutions of sera were mixed with GFP-expressing DENV-1 (strain West-978 Pac 74) RVPs and incubated with FcyR-expressing K562 cells. Infected cells were 979 quantified by flow cytometry. In each experiment, the cross-reactive FL-specific mAb

980 E60 was included as a control (J). To normalize the magnitude of enhancement across 981 independent experiments, results are displayed relative to the maximum infectivity 982 observed with mAb E60 run in parallel (designated by the dotted line at 100). Each line 983 represents the enhancement curve from an individual mouse. Error bars indicate the 984 range of duplicate technical replicates.

985 Figure S6. ZIKV mRNA LNP vaccines containing mutant FL sequences 986 showed reduced ADE against DENV in AG129 mice, Related to Figure 4. A. 987 Neutralization of DENV-2 RVPs by sera pooled from placebo or JEVsig-prM-E (2 µg dose of WT or FL mutant) vaccinated mice. Error bars indicate the range of duplicate 988 989 technical replicates. B. Enhancing effects of ZIKV immune serum on DENV-2 infection in AG129 mice. Recipient AG129 mice were passively transferred PBS or 10 µl of 990 pooled serum from BALB/c mice vaccinated with WT or FL mutant JEV_{sig}-prM E LNPs. 991 992 One day later, animals were challenged with 10⁴ FFU of DENV-2 (strain S220) and followed for mortality. Results are pooled from two independent challenge experiments 993 994 (numbers of animals indicated beneath graph). Survival curves between serum 995 transfers from JEV_{sig}-prM-E (WT and FL mutant LNPs) vaccinated mice were 996 statistically different (***, P < 0.001, log-rank test).

997





Days after challenge

