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Reporting Summary

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Statistics		
	es, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.	
n/a Confirmed		
☐ ☐ The exact sam	uple size (n) for each experimental group/condition, given as a discrete number and unit of measurement	
A statement of	on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly	
	test(s) used AND whether they are one- or two-sided ests should be described solely by name; describe more complex techniques in the Methods section.	
A description of all covariates tested		
A description of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons		
A full description of the statistical parameters including central tendency (e.g. means) or other basic estimates (e.g. regression coefficient) AND variation (e.g. standard deviation) or associated estimates of uncertainty (e.g. confidence intervals)		
	thesis testing, the test statistic (e.g. F , t , r) with confidence intervals, effect sizes, degrees of freedom and P value noted exact values whenever suitable.	
For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings		
For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes		
\square Estimates of effect sizes (e.g. Cohen's d , Pearson's r), indicating how they were calculated		
'	Our web collection on <u>statistics for biologists</u> contains articles on many of the points above.	
Software and c	code	
Policy information abou	ut <u>availability of computer code</u>	
Data collection	n/a	
Data analysis	Data was graphed and analyzed using GraphPad Prism software.	
For manuscripts utilizing custom algorithms or software that are central to the research but not yet described in published literature, software must be made available to editors/reviewers. We strongly encourage code deposition in a community repository (e.g. GitHub). See the Nature Research guidelines for submitting code & software for further information.		
Data		
- Accession codes, un - A list of figures that	ut <u>availability of data</u> include a <u>data availability statement</u> . This statement should provide the following information, where applicable: ique identifiers, or web links for publicly available datasets have associated raw data restrictions on data availability	
All data generated or analyzed during this study are included in this published article.		
Field-speci	fic reporting	
Please select the one b	elow that is the best fit for your research. If you are not sure, read the appropriate sections before making your selection.	
Life sciences Behavioural & social sciences Ecological, evolutionary & environmental sciences		
For a reference copy of the do	ocument with all sections, see <u>nature.com/documents/nr-reporting-summary-flat.pdf</u>	

Life sciences study design

	a on those points even when the disclosure is possible
	e on these points even when the disclosure is negative.
	sample size (n=5 per group) in the mouse immunogenicity studies has been chosen based on previously published studies where we lyzed DENV subunit immunogenicity.
Data exclusions No o	data has been excluded
	samples have been analyzed in duplicate. Wildtype mouse immunizations has been replicated once in order to generate sufficient sera for sive transfer studies.
Randomization mice	e were randomly attributed to experimental groups.
Blinding	hors were not blinded to group allocation in this study
We require information from	n/a Involved in the study ChIP-seq Plow cytometry MRI-based neuroimaging MRI-based neuroimaging
Antibodies used	4G2, 1M7, C8, C10, A9E, G9E, ZKA-230, 2D22, a-mouse IgG-AP, a-human IgG-AP
Validation	4G2> Henchal et al., 1985; Am J Trop Med Hyg 1M7> Smith et al., 2014; J Vir C8, C10> Rouvinsky et al., 2015; Nature A9E, G9E> Collins et al., 2019; JCI insight ZKA-230> Stettler et al., 2016; Science 2D22> Firbriansah et al., 2015; Science a-Mouse IgG-AP: Sigma (A9316) a-human IgG-AP: Sigma (A1543)
Eukaryotic cell	lines
Policy information abou	t <u>cell lines</u>
Cell line source(s)	Vero-81 cell lines. Purchased from ATCC
Authentication	Cell lines used were not authenticated after purchase
Mycoplasma contami	nation Cell lines used are mycoplasma free
Commonly misidentifi (See <u>ICLAC</u> register)	ed lines n/a

Animals and other organisms

Policy information about studies involving animals; ARRIVE guidelines recommended for reporting animal research

Laboratory animals

Female C57BL/6 mice were purchased from Jackson Laboratory .

supported by NIH)

N/a

Field-collected samples

n/a

All mouse experiments were performed under protocols approved by the University of North Carolina Institutional Animal Care and Use Committee, in compliance with federal regulations (the Public Health Service Policy on Humane Care and Use of Laboratory Animals).

Note that full information on the approval of the study protocol must also be provided in the manuscript.