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Reporting Summary

Nature Portfolio wishes to improve the reproducibility of the work that we publish. This form provides structure for consistency and transparency in reporting. For further information on Nature Portfolio policies, see our Editorial Policies and the Editorial Policy Checklist.

For all statistical analyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section

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n/a	Confirmed
	$oxed{\boxtimes}$ The exact sample size (n) for each experimental group/condition, given as a discrete number and unit of measurement
	🔀 A statement on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly
	The statistical test(s) used AND whether they are one- or two-sided Only common tests 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)
	For null hypothesis testing, the test statistic (e.g. <i>F</i> , <i>t</i> , <i>r</i>) with confidence intervals, effect sizes, degrees of freedom and <i>P</i> value noted <i>Give P values as exact values whenever suitable.</i>
\boxtimes	For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings
\boxtimes	For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes
	\boxtimes 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 code

Policy information about availability of computer code

Data collection

Data were collected from dog vaccination, rabies surveillance and community education field activities using the WVS App (Android versions 1.0 - 6.6) in accordance with the MOU between Mission Rabies and the Government of Goa to implement the project. Records were verified with paper-based project records.

Data on human rabies deaths and human dog bites were gathered from the Government of Goa Directorate of Health Services data reports.

Data analysis

Data were analyzed using R statistical software version 3.6.2 and 4.0.2.

Cluster analyses was performed using SaTScan™ v9.6 software through R package rsatscan.

Cost-effectiveness analysis was conducted using the MS Excel based model, RabiesEcon published by Kunkel et al (2022), with modification described in the methods.

 $Phylogenetic analysis \ was \ performed \ using \ Mega7 \ and \ BEAST \ v1.10.4 \ and \ phylogeny \ trees \ were \ generated \ using \ FigTree \ v1.4.4.$

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 and reviewers. We strongly encourage code deposition in a community repository (e.g. GitHub). See the Nature Portfolio guidelines for submitting code & software for further information.

Data

Policy information about availability of data

All manuscripts must include a <u>data availability statement</u>. This statement should provide the following information, where applicable:

- Accession codes, unique identifiers, or web links for publicly available datasets
- A description of any restrictions on data availability
- For clinical datasets or third party data, please ensure that the statement adheres to our policy

Data supporting the findings of this work are available within the paper and its Supplementary Information files.

All sequences generated as part of this study are deposited in GenBank (accession codes: MW054945-MW055041).

GenBank accession codes for reference sequences used in the phylogenetic analysis were as follows:

JQ685901, JQ685945, JQ944707, KX148105, JQ944708, JQ944706, JQ944705, KX148225, HE801587, JX987739, KF150744, MK760761, KX148226, HE802676, KC171645, KC171644, KC171643, KM272192, GU937029, KX148228.

Field-specific reporting

Ple	ease select the one below	that is the	best fit for your research	. If yo	ou are not sur	e, read the	appropriate sections	before making y	our selection.
X	Life sciences	Behav	vioural & social sciences		Ecological,	evolutionary	/ & environmental sc	iences	

For a reference copy of the document with all sections, see nature.com/documents/nr-reporting-summary-flat.pdf

Life sciences study design

All studies must disclose on these points even when the disclosure is negative.

Sample size

The vaccination approach aimed to access as high a proportion of the population as possible. A total of 426,119 dog rabies vaccination events were conducted during the study period. Postmortem sampling and rabies diagnostic testing was performed on 552 dogs with suspect rabies as part of the Goa State rabies surveillance activities coordinated by the Department for Animal Husbandry and Veterinary Services and in accordance with The Prevention of Cruelty to Animals Act, 1960. Attempts were made to sample all dogs reported to have died in Goa following signs consistent with rabies (aggression, hypersalivation, neurological signs, sudden death).

Data exclusions

No data were excluded from the study.

Replication

The study demonstrates strategies for effective human and canine rabies elimination at the sub-district administrative region (taluka), which is a standard administrative unit in India. Our logistical regression analysis assessed the relationship between canine rabies and dog vaccination coverage across twelve taluka regions. The odds of dog rabies decreased as rolling mean 12-month vaccination coverage increased, a pattern that was seen across 10 of the 12 talukas, with identified causes of the two anomalies included in the discussion. As such the outcome of human and canine rabies elimination through the described methods would be repeatable under similar epidemiological settings in India.

Randomization

As this was a state-wide initiative all regions of the state were included in the study. It was not considered appropriate or feasible to perform a randomized cohort study on the implementation aspect of the study from a financial, political or ethical standpoint. We have accounted for covariates where appropriate, e.g. with the use of multivariable logistic regression.

Samples for phylogenetic analysis were randomly selected from the bank of positive rabid dog brains spanning 2016 - 2018 using a random number generator.

Blinding

As all areas of the state were included in the rabies control program, it was not possible to blind investigators during data collection or analysis.

Individuals performing laboratory diagnosis were blinded to the initial Lateral Flow Test outcome.

Individuals performing post-vaccination surveys were blinded from the areas that had been vaccinated.

Reporting for specific materials, systems and methods

We require information from authors about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material, system or method listed is relevant to your study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response.

Materials & experime	ental systems	Methods	
n/a Involved in the study		n/a Involved in the study	
Antibodies		ChIP-seq	
Eukaryotic cell lines		Flow cytometry	
Palaeontology and a	archaeology	MRI-based neuroimaging	
Animals and other organisms			
Human research participants			
Clinical data			
Dual use research o	f concern		
1			
Animals and othe	r organisms		
,		ARRIVE guidelines recommended for reporting animal research	
Laboratory animals	Laboratory animals were no	ot involved in any part of this study.	
and sex were vaccinated in a activities were part of the stabutterfly nets by trained stafintramuscular injection, mark		e vaccination of owned and unowned domestic dogs were recorded as part of the study. Dogs of any age accordance with government and WHO guidance on mass dog vaccination for rabies control. The study state public health intervention to control the spread of rabies. Dogs were handled humanely or using aff in accordance with state and national guidelines. Dogs were vaccinated by subcutaneous or arked with a non-toxic temporary marker paint and immediately released at the same location. For owned the before conducting vaccination.	
Field-collected samples	Postmortem brain samples were collected from suspect rabid animals in Goa during the study period. Samples were taken as soon after death as possible at the Disease Investigation Unit in Tonca, Panjim. Samples were stored at 4 degrees Celsius until processing for rabies diagnosis using direct fluorescent antibody testing within days. A subset were stored a -80C for subsequent quality-control and viral gene sequencing. Samples continue to be stored by the Government of Goa as an ongoing archive of rabies virus for subsequent analysis. When samples are no longer required, they are incinerated as hazardous biological waste.		
Ethics oversight	rabies surveillance. The Go Veterinary ethical review w	epartment for Animal Husbandry oversaw the project protocols and methods for mass dog vaccination and ba Veterinary Association also had oversight of veterinary protocols on the project. Vas also provided by the University of Edinburgh Veterinary Ethical Review Committee, with ethical Dec 2014 (107 14: Investigation of efficacy of rabies vaccination in India) and 28th July 2015 (64/15: cination approaches).	

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