nature portfolio

Corresponding author(s):	Simon Hay
Last updated by author(s):	Oct 18, 2022

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.

_				
C	<u>-</u> ۱	t٠	ct	ics
_	_		\sim 1	11 \

FOI	an statistical analyses, commit that the following items are present in the figure legend, table legend, main text, or Methods section.		
n/a	Confirmed		
	\square 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		
\boxtimes	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.		
\boxtimes	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>		
	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 <u>availability of computer code</u>			

Data collection

No primary data collection was carried out for this analysis.

Data analysis

This analysis was carried out using R version 3.6.1 and using R-INLA v.20.01.29.9000. Maps were produced using ArcGIS Desktop 10.6. All code used for these analyses is publicly available online at http://ghdx.healthdata.org

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

The findings of this study were produced using data available in public online repositories: [https://malariaatlas.org/malaria-burden/], [http://ghdx.healthdata.org/record/ihme-data/africa-under-5-lri-incidence-prevalence-mortality-geospatial-estimates-2000-2017], [http://ghdx.healthdata.org/record/ihme-data/lmic-under-5-diarrhea-incidence-prevalence-and-mortality-geospatial-estimates-2000-2017], [http://ghdx.healthdata.org/record/ihme-data/gbd-2017-incidence-prevalence-and-ylds-1990-2017], data available upon request from the data provider. This study complies with

the Guidelines for Accurate and Transparent Health Estimates Reporting (GATHER) recommendations. All maps presented in this study are generated by the authors; no permissions are required for publication.

Field-specific reporting				
Please select the one below that is the best fit for your research. If you are not sure, read the appropriate sections before making your selection.				
Life sciences	ces Behavioural & social sciences Ecological, evolutionary & environmental sciences			
or a reference copy of t	he document with all sections, see nature.com/documents/nr-reporting-summary-flat.pdf			
_ife scier	nces study design			
All studies must dis	close on these points even when the disclosure is negative.			
Sample size	The individual cause-level spatio-temporally varying estimates of incidence, prevalence, and mortality for LRIs, diarrhoea, and malaria have been previously published. Within each paper, the sources, model specifications and model validations are described in detail. Our paper combines these existing estimates.			
Data exclusions	The individual cause-level spatio-temporally varying estimates of incidence, prevalence, and mortality for LRIs, diarrhoea, and malaria have been previously published. Within each paper, the sources, model specifications and model validations are described in detail. Our paper combines these existing estimates.			
Replication	All code and data are publicly available to reproduce this work.			
Randomization	This analysis is an observational mapping study and there were no experimental groups.			

Reporting for specific materials, systems and methods

Blinding was not relevant to this study, as it was an observational study using survey and surveillance data.

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 & experimental systems		Methods	
n/a	Involved in the study	n/a	Involved in the study
\boxtimes	Antibodies	\boxtimes	ChIP-seq
\boxtimes	Eukaryotic cell lines	\boxtimes	Flow cytometry
\boxtimes	Palaeontology and archaeology	\boxtimes	MRI-based neuroimaging
\boxtimes	Animals and other organisms		
\boxtimes	Human research participants		
\boxtimes	Clinical data		
\boxtimes	Dual use research of concern		

Blinding