# THE LANCET

# Supplementary appendix

This appendix formed part of the original submission and has been peer reviewed. We post it as supplied by the authors.

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# **Appendix 1**

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## 1. WHO regional groupings and country list

#### WHO African Region: 47

Algeria, Angola, Benin, Botswana, Burkina Faso, Burundi, Cabo Verde, Cameroon, Central African Republic, Chad, Comoros, Congo, Côte d'Ivoire, Democratic Republic of the Congo, Equatorial Guinea, Eritrea, Eswatini, Ethiopia, Gabon, Gambia, Ghana, Guinea, Guinea-Bissau, Kenya, Lesotho, Liberia, Madagascar, Malawi, Mali, Mauritania, Mauritius, Mozambique, Namibia, Niger, Nigeria, Rwanda, Sao Tome and Principe, Senegal, Seychelles\*, Sierra Leone, South Africa, South Sudan, Togo, Uganda, United Republic of Tanzania, Zambia, Zimbabwe.

#### WHO Region of the Americas: 35

Antigua and Barbuda\*, Argentina, Bahamas\*, Barbados\*, Belize, Bolivia (Plurinational State of), Brazil, Canada\*, Chile\*, Colombia, Costa Rica, Cuba, {Dominica}, Dominican Republic, Ecuador, El Salvador, Grenada, Guatemala, Guyana, Haiti, Honduras, Jamaica, Mexico, Nicaragua, Panama, Paraguay, Peru, {Saint Kitts and Nevis\*}, Saint Lucia, Saint Vincent and the Grenadines, Suriname, Trinidad and Tobago\*, United States of America\*, Uruguay\*, Venezuela (Bolivarian Republic of).

#### WHO South-East Asia Region: 11

Bangladesh, Bhutan, Democratic People's Republic of Korea, India, Indonesia, Maldives, Myanmar, Nepal, Sri Lanka, Thailand, Timor-Leste.

#### WHO European Region: 53

Albania, {Andorra\*}, Armenia, Austria\*, Azerbaijan, Belarus, Belgium\*, Bosnia and Herzegovina, Bulgaria, Croatia\*, Cyprus\*, Czechia\*, Denmark\*, Estonia\*, Finland\*, France\*, Georgia, Germany\*, Greece\*, Hungary\*, Iceland\*, Ireland\*, Israel\*, Italy\*, Kazakhstan, Kyrgyzstan, Latvia\*, Lithuania\*, Luxembourg\*, Malta\*, {Monaco\*}, Montenegro, Netherlands\*, North Macedonia, Norway\*, Poland\*, Portugal\*, Republic of Moldova, Romania, Russian Federation, {San Marino\*}, Serbia, Slovakia\*, Slovenia\*, Spain\*, Sweden\*, Switzerland\*, Tajikistan, Türkiye, Turkmenistan, Ukraine, United Kingdom of Great Britain and Northern Ireland\*, Uzbekistan.

#### WHO Eastern Mediterranean Region: 21

Afghanistan, Bahrain\*, Djibouti, Egypt, Iran (Islamic Republic of), Iraq, Jordan, Kuwait\*, Lebanon, Libya, Morocco, Oman\*, Pakistan, Qatar\*, Saudi Arabia\*, Somalia, Sudan, Syrian Arab Republic, Tunisia, United Arab Emirates\*, Yemen.

#### WHO Western Pacific Region: 27

Australia\*, Brunei Darussalam\*, Cambodia, China, {Cook Islands}, Fiji, Japan\*, Kiribati, Lao People's Democratic Republic, Malaysia, {Marshall Islands}, Micronesia (Federated States of), Mongolia, {Nauru\*}, New Zealand\*, {Niue}, {Palau\*}, Papua New Guinea, Philippines, Republic of Korea\*, Samoa, Singapore\*, Solomon Islands, Tonga, {Tuvalu}, Vanuatu, Viet Nam

\* high-income  $(57)^1$ , countries in curly brackets indicate countries for which no total disease burden figures are available through the WHO Global Health Observatory (GHO)<sup>2</sup>

#### 2. Multilevel modeling of population exposures to different levels of WASH services

	Number of countries	Number of data points
	(urban/rural)#	(urban/rural)
Drinking water		
Improved water source	170/167	1839/1803
Water source within 30 minutes	140/139	609/608
Water source accessible on	169/166	787/774
premises		
Water available when needed	107/91	357/247
Water free from contamination	82/65	266/166
POU chlorination	103/103	190/190*
POU solar treatment	102/102	190/190*
POU filtration/boiling	103/103	190/190*
Sanitation		
Improved sanitation facility	169/166	1863/1854
Shared sanitation	155/153	759/752
Sanitation connected to sewer	155/152	1290/1277
Hygiene		
Basic handwashing facility with	98/99	193/199
water and soap at home		

Table S.1: Number of countries covered with  $\geq 1$  national data point and number of data points by WASH exposure category

POU: point of use; # out of 195 countries (WHO member states and territories), \*includes POU treatment on different water exposures categories (e.g. POU chlorination on surface water, unimproved and limited drinking water, POU chlorination on improved, not on premises, POU chlorination, on premises)

#### **Random slope model**

This model is used for different drinking water and sanitation services.

$$logit(Y_{ij}) = \beta_0 + \beta_1 X_{1ij} + \sum_{k=2}^{18} \beta_k X_{kj} + \mu_j + u_{ij} X_{1ij} + \varepsilon_{ij}$$

where 'logit(Y<sub>ij</sub>)' is the proportion of use of a specific drinking water or sanitation service on the logit scale for year *i* and country *j*, ' $\beta_0$ ' is the overall mean intercept of Y across countries, ' $\beta_1$ ' is the regression coefficient for the predictor year  $X_{1ij}$  (defined at level 1: individual observations-level), ' $\beta_k$ ' are the regression coefficients for the regional covariates (' $X_{kj}$ ' – 17 regions), ' $\mu_j$ ' are the residuals on the country level (group-level residuals), ' $u_{ij}X_{1ij}$ ' is the interaction between country and year and ' $\varepsilon_{ij}$ ' are the residuals at the level of the observations (individual-level residuals).

Regional grouping for modelling of drinking water and sanitation services follow m49 classification and are divided as: Australia and New Zealand; Central Asia; Eastern Asia; Eastern Europe; Latin America and the Caribbean; Melanesia; Micronesia; Northern Africa; Northern America; Northern Europe; Polynesia; Southern Asia; Southern Asia; Southern Europe; Sub-Saharan Africa; Western Asia; Western Europe.

#### **Random intercept model**

Random intercept models are simpler compared to random slope models and are used for household water treatment and access to basic handwashing facilities.

Modeling of household water treatment

$$Y_{ij} = \beta_0 + \beta_1 X_{1ij} + \sum_{k=2}^{6} \beta_k X_{kj} + \mu_j + \varepsilon_{ij}$$

where  $Y_{ij}$  is the proportion of household water treatment (for each water treatment category) for year *i* and country *j*,  $\beta_0$  is the overall mean intercept of *Y* across countries,  $\beta_1$  is the regression coefficient for the predictor year  $X_{1ij}$  (defined at level 1: individual observations-level),  $\beta_k$  are the regression coefficients for the regional covariates  $X_{kj}$  (defined at level 2: country-level),  $\mu_j$  are the residuals on the country-level (group-level residuals) and  $\varepsilon_{ij}$  are the residuals at the level of the observations (individual observations-level residuals).

Modeling of basic handwashing facility presence

$$logit(Y_{ij}) = \beta_0 + \beta_1 X_{1ij} + \sum_{k=2}^{6} \beta_k X_{kj} + \sum_{l=7}^{9} \beta_l X_{lj} + \mu_j + \varepsilon_{ij}$$

where logit( $Y_{ij}$ ) is the proportion of presence of a designated handwashing facility on the logit scale for year *i* and country *j*,  $\beta_0$  is the overall mean intercept of *Y* across countries,  $\beta_1$  is the regression coefficient for the predictor year  $X_{1ij}$  (defined at level 1: individual observations-level),  $\beta_k$  are the regression coefficients for the regional covariates  $X_{kj}$  (defined at level 2: country-level),  $\beta_l$  are the regression coefficients for the income level covariates  $X_{lj}$  (defined at level 2),  $\mu_j$  are the residuals on the country-level (group-level residuals) and  $\varepsilon_{ij}$  are the residuals at the level of the observations (individual observations-level residuals).

Regional grouping for modelling of household water treatment follow m49 classification and divided as: Africa, Americas, Asia, Europe and Oceania.

Income level covariates include: low income, lower-middle income, higher-middle income.

#### 3. Estimation of safely managed drinking water adjusted for household level and point of use quality

**Background:** Exposure to safely managed drinking water at the household level is the counterfactual minimum risk exposure level in the analysis of drinking water-attributable diarrhoea. Estimates for population exposure to safely managed drinking water by country for the year 2019 are derived using multilevel modeling of JMP data, which calculate safely managed drinking water services at the level of the urban and rural domains. These estimates of safely managed drinking water were adjusted because: i) the JMP uses data on microbiological quality of drinking water at the point of collection with no consideration of changes in water quality at point of use (e.g., through unhygienic household storage), and ii) the JMP safely managed drinking water estimates are based on the minimum of quality, availability and accessibility at urban and rural levels for each country, since data on these factors typically comes from different sources and cannot be integrated at the household level.

Methods: We adjusted the modeled exposure estimates of safely managed drinking water by integrating nationally-representative information on deterioration of water quality between point-of-collection and point-ofuse and on safely managed drinking water access at the household-level from 38 nationally-representative household surveys. We used data from these household surveys (MICS and other national surveys, available through the JMP) that collected data on all three elements of safely managed drinking water: (i) accessibility on premises, (ii) availability when needed, and (iii) freedom from faecal contamination at (a) both the urban/rural domain levels and household levels and (b) point-of-collection water quality as well as point of use water quality<sup>3</sup>. The 38 household surveys cover 17 countries from the African region, 6 countries from the region of the Americas, 5 countries from the Eastern Mediterranean Region, 2 countries from the European region, 2 countries from the South-East Asian region and 6 countries from the Western Pacific region and comprise 9 lowincome, 18 lower middle-income and 11 upper middle-income countries. Data from these surveys can be used to explore the relationship between safely managed drinking water categorized at the urban and rural domain level with no faecal contamination at the point of collection (smdw domain poc) and safely managed drinking water categorized at the household level with no faecal contamination at the point of use (smdw hh pou). smdw hh pou can then be modelled for all low- and middle-income countries for which a modeled estimate for smdw domain poc is available for the year 2019.

We analysed smdw\_hh\_pou as a function of smdw\_domain\_poc as main predictor in a linear regression model. As both the dependent variable and the main predictor are proportions, they were logit-transformed to account for their distribution between 0 and 1.

The following additional covariates were explored in the linear model:

- a squared term of the main predictor smdw domain poc.
- different regional covariates: the six WHO regions (see section 3, indicator variables) and region m49, level 1 (5 indicator variables, used for SDG reporting)
- country income status (low income, lower-middle income, upper-middle income, indicator variables)
- per capita GDP (gross domestic product) for 2019 (continuous variable, natural log transformed)

The final model was chosen based on statistical tests for significance (for covariates), the value of the adjusted  $R^2$  and the Akaike's and Bayesian Information Criterion (AIC and BIC).

For countries with an actual value for smdw\_hh\_pou the modeled value of smdw\_domain\_poc based on JMP estimates was multiplied with the ratio of smdw\_hh\_pou / smdw\_domain\_poc. For 98 LMIC with no value for smdw\_hh\_pou predictions of this variable were generated through the linear regression model with CIs derived using standard errors of the forecast.

Model diagnostics were performed by plotting residuals against the linear predictions and Q-Q-plots.

**Results**: smdw\_hh\_pou shows strong positive linear correlations with smdw\_domain\_poc (Pearson correlations coefficients 0.9 for both urban and rural datasets.

The final model for the urban dataset consisted of a linear model including smdw\_domain\_poc (logit transformed) and log per capita GDP as continuous variables and smdw\_domain\_poc (logit transformed) plus its squared term as predictor for the rural dataset (Table S1.2).

Table S.2: Model summary for urban/rural model

Urban model	Rural model
Dependent variable: smdw_hh_pou (logit	Dependent variable: smdw_hh_pou (logit
transformed)	transformed)
Smdw_domain_poc (logit): p: <0.001	Smdw_domain_poc (logit): p: 003
GDP capita (2019, ln): p=0.07	Smdw_domain_poc squared (of logit
	smdw_domain): 0.02
R <sup>2</sup> : 0.76	R <sup>2</sup> : 0.81

The mean difference between the modeled and actual values for smdw\_hh\_pou for those countries (n=38) where this data was available was 0.9 percentage points for the urban and 0.5 percentage points for the rural dataset.

After the described adjustment, we estimate that 37.9% (29.1%, 49.9%) of the population in LMICs used safely managed drinking water at the household level. This compares to 56.3% (50.8%, 61.8%) without the adjustment.

#### 4. Uncertainty estimates at country, regional and global levels

We estimated uncertainty intervals for the PAFs and attributable disease burden (deaths and DALYs) at the country level with Monte Carlo simulations, taking account of uncertainty of the exposure estimates, as well as the uncertainties of the matching exposure-response relationships, and the total disease burden. For the various exposure levels and exposure-response relationships, we drew random samples of size 10,000 for every country assuming normal distributions after logit or log transformation with the standard error of the logit exposure estimate or the log relative risk as standard deviation. Each randomly generated PAF by country was multiplied with a random draw of the overall disease burden for the condition of interest (diarrhoea, ARI, protein-energy malnutrition, soil-transmitted helminthiasis) assuming a log-normal distribution with the point estimate as the mean, to correct for skewness. The latter allows for the mean of the Monte Carlo result distribution being equal to the result calculated with the point estimates directly. Confidence intervals for the PAFs and attributable disease burdens were set as the difference between the 2.5<sup>th</sup> and 97.5<sup>th</sup> percentile.

#### 5. WASH-attributable burden of malnutrition



Figure S.1: Conceptual framework for calculating the WASH-attributable burden of undernutrition mediated via the WASH-attributable burden of diarrhoea; light grey: external data inputs, dark grey: estimated in this work

WHO total disease burden figures on malnutrition include ICD 10 codes E40-E46 which are not further disaggregated into wasting, stunting or underweight (Table S.3).

Table S.3: Composition of protein-energy malnutrition as reported by the WHO Global Health Observatory (GHO)

ICD 10- codes (version 2019)			
E40	Kwashiorkor		
E41	Nutritional marasmus		
E42	Marasmic kwashiorkor		
E43	Unspecified severe protein-energy malnutrition		
E44	Protein-energy malnutrition of moderate and mild		
	degree		
E45	Retarded development following protein-energy		
	malnutrition		
E46	Unspecified protein-energy malnutrition		

Source: ICD-10, version: 2019, https://icd.who.int/browse10/2019/en#/E40

ICD-10 codes included for diarrhoea, ARI and soil-transmitted helminthiasis are listed in Tables S.6-S.8.

#### 6. Sources of exposure, exposure-response and disease data by outcome

Health outcome	Source of exposure	Source of exposure-	Source of disease
	data	response relationship	data
diarrhoea	JMP household	Wolf et al. 2022 <sup>4</sup>	WHO Global Health
	survey database		Estimates <sup>2</sup>
Acute respiratory infections	JMP household	Ross et al. 2023 <sup>5</sup>	WHO Global Health
	survey database		Estimates <sup>2</sup>
Protein energy malnutrition	JMP household	Troeger et al. 2018 <sup>6</sup>	WHO Global Health
	survey database		Estimates <sup>2</sup>
		Wolf et al. 2022 <sup>4</sup>	
Soil-transmitted helminth infections	NA	NA	WHO Global Health
			Estimates <sup>2</sup>

Table S.4: Sources of exposure, exposure-response and disease data by outcome

Table S.5: Exposure response functions linking WASH and diarrhoea<sup>4</sup> and hygiene and acute respiratory infections

	diarrhoea	ARI
WATER		
Basic drinking water*	0.81 (0.70, 0.94)	
Basic, on premises, higher water	0.48 (0.26, 0.87)	
quality*		
POU chlorination*	0.66 (0.56, 0.77)	
POU solar treatment*	0.63 (0.50, 0.80)	
POU filtration*	0.50 (0.41, 0.60)	
SANITATION		
Basic sanitation, without sewer	0.79 (0.61, 1.03)	
connection#		
Basic sanitation with sewer	0.53 (0.30, 0.93)	
connection#		
HYGIENE		
Handwashing with soap§	0.7 (0.64, 0.76)	0.83 (0.76, 0.90)

POU: point of use, \* compared against surface unimproved or limited drinking water; # compared against open defecation, unimproved or limited sanitation; § comparison is handwashing with soap promotion (with or without provision of handwashing infrastructure against no handwashing with soap promotion; bold: relative risk for minimum risk exposure level in this analysis

Table S.6: Composition of diarrhoea as reported by the WHO Global Health Observatory (GHO)

ICD 10- codes (version 2019)			
A00	Cholera		
A01	Typhoid and paratyphoid fevers		
A03	Shigellosis		
A04	Other bacterial intestinal infections		
A06	Amoebiasis		
A07	Other protozoal intestinal diseases		
A08	Viral and other specified intestinal infections		
A09	Other gastroenteritis and colitis of infectious and		
	unspecified origin		

Source: ICD-10, version: 2019, https://icd.who.int/browse10/2019/en#/A00

Table S.7: Composition of acute respiratory infections as reported by the WHO Global Health Observatory (GHO)

Nonsuppurative otitis media		
Suppurative and unspecified otitis media		
Acute nasopharyngitis [common cold]		
Acute sinusitis		
Acute pharyngitis		
Acute tonsillitis		
Acute laryngitis and tracheitis		
Acute obstructive laryngitis [croup] and epiglottitis		
Acute upper respiratory infections of multiple and		
unspecified sites		
Influenza due to identified zoonotic or pandemic		
influenza virus		
Influenza due to identified seasonal influenza virus		
Influenza, virus not identified		
Viral pneumonia, not elsewhere classified		
Pneumonia due to Streptococcus pneumoniae		
Pneumonia due to Haemophilus influenzae		
Bacterial pneumonia, not elsewhere classified		
Pneumonia due to other infectious organisms, not		
elsewhere classified		
Pneumonia in diseases classified elsewhere		
Pneumonia, organism unspecified		
Acute bronchitis		
Acute bronchiolitis		
Unspecified acute lower respiratory infection		

Source: ICD-10, version: 2019, <u>https://icd.who.int/browse10/2019/en#/H65;</u> <u>https://icd.who.int/browse10/2019/en#/J00; https://icd.who.int/browse10/2019/en#/J09-J18;</u> https://icd.who.int/browse10/2019/en#/J20-J22

Table S.8: Composition of soil-transmitted helminthiasis (intestinal nematode infections) as reported by the WHO Global Health Observatory (GHO)

ICD 10- codes (version 2019)	
B76	Hookworm diseases
B77	Ascariasis
B79	Trichuriasis

Source: ICD-10, version: 2019, https://icd.who.int/browse10/2019/en#/B76

Composition of protein-energy malnutrition as reported by the GHO is listed in Table S.3.

#### 7. Exposure estimates

	Percentage of population (95% CI) using			
Region	safely managed drinking water sources at household level plus 0 <i>E-coli</i> at POU	basic drinking water sources	surface, unimproved or limited drinking water	
Sub-Saharan Africa, LMIC	8.8 (6.9, 11.2)	65.6 (63.5, 67.7)	34.4 (32.3, 36.5)	
America, LMIC	59.3 (42.3, 74.3)	96.7 (94.1, 98.2)	3.3 (1.8, 5.9)	
Eastern Mediterranean, LMIC	36.2 (28.0, 45.3)	88.7 (86.5, 90.5)	11.3 (9.5, 13.5)	
Europe, LMIC	61.9 (44.8, 76.5)	97.0 (95.1, 98.1)	3 (1.9, 4.9)	
South-East Asia, LMIC	27.2 (10.7, 53.7)	91.7 (89.0, 93.7)	8.3 (6.3, 11)	
Western Pacific, LMIC	55.6 (26.2, 81.6)	92.2 (88.0, 95.0)	7.8 (5, 12)	
Total LMIC	37.9 (27.1, 49.9)	88.0 (86.0, 89.7)	12 (10.3, 14)	

POU: point of use, LMIC: low- and middle-income countries, CI: confidence interval

Table S.10: Distribution o	f the population	to exposure lev	vels of sanitation,	by region,	for 2019
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	Percentage of population (95% CI) using/practicing					
Region	Basic sanitation connected to sewer	Basic sanitation	Open defecation, unimproved or limited sanitation			
Sub-Saharan Africa, LMIC	7.4 (6.3, 8.7)	35.0 (33.5, 36.4)	65.0 (63.6, 66.5)			
America, LMIC	59.4 (54.0, 64.5)	87.1 (85.6, 88.5)	12.9 (11.5, 14.4)			
Eastern Mediterranean, LMIC	32.2 (24.5, 41.0)	76.1 (73.2, 78.8)	23.9 (21.2, 26.8)			
Europe, LMIC	59.0 (33.9, 80.1)	96.2 (93.4, 97.9)	3.8 (2.1, 6.6)			
South-East Asia, LMIC	9.1 (5.7, 14.3)	72.7 (69.3, 75.8)	27.3 (24.2, 30.7)			
Western Pacific, LMIC	48.7 (28.2, 69.6)	84.2 (79.6, 88.0)	15.8 (12, 20.4)			
Total LMIC	29.7 (23.9, 36.1)	72.7 (70.5, 74.7)	27.3 (25.3, 29.5)			

LMIC: low- and middle-income countries, CI: confidence interval

	Percentage of population (95% CI) having access to/practicing					
Region	Handwashing with soap and water after faecal contact	Basic handwashing facility	No or limited handwashing facility			
Sub-Saharan Africa, LMIC	9.2 (5.5, 15.1)	30.0 (27.0, 33.1)	70.0 (66.9, 73.0)			
America, LMIC	35.9 (32.6, 39.3)	82.0 (69.2, 90.2)	18.0 (9.8, 30.8)			
Eastern Mediterranean, LMIC	21.8 (7.9, 47.6)	71.5 (64.3, 77.7)	28.5 (22.3, 35.7)			
Europe, LMIC	24.9 (7.6, 57.2)	95.8 (91.2, 98.0)	4.2 (2, 8.8)			
South-East Asia, LMIC	28.4 (24.6, 32.4)	74.4 (65.3, 81.7)	25.6 (18.3, 34.7)			
Western Pacific, LMIC	17.3 (15.9, 18.8)	92.3 (74.8, 98.0)	7.7 (2, 25.2)			
Total LMIC	22.0 (19.0, 25.4)	73.5 (64.9, 80.6)	26.5 (19.4, 35.1)			
Total HIC	50.6 (47.9, 53.3)*	-	-			

Table S.11: Distribution of the population to exposure levels of hygiene, by region, for 2019

LMIC: low- and middle-income countries, CI: confidence interval

	Percentage of population treating their drinking water by					
Region	filtration or boiling	chlorination	solar radiation			
Sub-Saharan Africa, LMIC	8.5	6.8	0.0			
America, LMIC	20.8	12.9	0.1			
Eastern Mediterranean, LMIC	12.4	2.5	0.1			
Europe, LMIC	38.8	1.5	0.1			
South-East Asia, LMIC	26.4	2.1	0.1			
Western Pacific, LMIC	26.6	1.5	0.1			
Total LMIC	26.1	3.7	0.1			

Table S.12: Distribution of the population to exposure levels of household water treatment, by region, for 2019

LMIC: low- and middle-income countries

#### 8. Details on calculating attributable disease burden estimates following Figure 1 of the main manuscript

For the calculation of the deaths and DALYs attributable to a risk factor, the three following data inputs are needed:

- the distribution of exposure to (different levels of) the risk factor of interest in the study population (p<sub>j</sub> in Figure 1)
- 2) the exposure-response relationship between (different levels of) the risk factor and the health outcome(s) of interest (RR<sub>j</sub> in Figure 1)
- 3) overall disease statistics, i.e. WHO Global Health Estimates in deaths and DALYs by iso3, outcome, age group and sex (total burden in Figure 1)

Exposure response functions of Table S.5 are combined with the proportion of the population in the respective exposure levels to calculate the PAFs which are then multiplied with respective WHO Global Health Estimates.

#### 9. Regional estimates for WASH-attributable diarrhoea and ARI

Table S.13: Diarrhoea burden attributable to inadequate drinking water by region, 2019

Region	PAF	(95% CI)	Deaths	(95% CI)	DALYs	(95% CI)
Sub-Saharan	0.42	(0.38, 0.46)	209.033	(188 883	14 389 811	(12 463 521
Africa, LMIC	0.12	(0.50, 0.10)	209,055	232,835)	11,505,011	17,073,440)
America,	0.20	(0.17, 0.27)	4,467	(3,724,	341,627	(285,193,
LMIC				5,871)		449,556)
Eastern	0.35	(0.29, 0.43)	53,940	(41,154,	3,453,176	(2,727,407,
Mediterranean,				70,171)		4,346,215)
LMIC						
Europe, LMIC	0.13	(0.10, 0.20)	486	(389, 693)	125,942	(93,335,
. /		· · · /		/		189,431)

South-East	0.29	(0.19, 0.42)	230,821	(145,806,	8,337,219	(5,429,207,
Asia, LMIC				342786)		12,147,332)
Western	0.20	(0.15, 0.31)	6,299	(5,043,	603,057	(440,106,
Pacific, LMIC				8,320)		921,424)
Total LMIC	0.35	(0.31, 0.40)	505,046	(418,147,	27,250,832	(23,665,814,
				622,247)		31,915,532)

PAF: population attributable fraction, CI: confidence interval, DALYs: disability-adjusted life years, LMIC: low- and middle-income countries; 132 low- and middle-income countries, regional PAF aggregates relate to diarrhoea DALYs

Table S.14: Diarrhoea burden attributable to inadequate sanitation by region, 2019

Region	PAF	(95% CI)	Deaths	(95% CI)	DALYs	(95% CI)
Sub-Saharan	0.42	(0.39, 0.47)	211,020	(190,972,	14,479,134	(12,593,570,
Africa, LMIC				235,531)		17,076,920)
America,	0.24	(0.21, 0.26)	5,273	(4,665,	398,091	(353,571,
LMIC				5,962)		450519)
Eastern	0.33	(0.28, 0.40)	49,749	(39,360,	3,261,357	(2,672,534,
Mediterranean,				63,054)		3,998,589)
LMIC						
Europe, LMIC	0.19	(0.14, 0.22)	893	(738, 997)	179,133	(128, 689,
						209,681)
South-East	0.36	(0.30, 0.44)	288,508	(221,719,	10,392,260	(8,216,933,
Asia, LMIC				372,792)		12,968,318)
Western	0.28	(0.23, 0.33)	8,865	(7,633,	838,429	(684,272,
Pacific, LMIC				10,169)		994,937)
Total LMIC	0.38	(0.35, 0.41)	564,308	(494,793,	29,548,404	(26,550,940,
				653,164)		33,222,352)

PAF: population attributable fraction, CI: confidence interval, DALYs: disability-adjusted life years, LMIC: low- and middle-income countries; 132 low- and middle-income countries, regional PAF aggregates relate to diarrhoea DALYs

Table S.15: Diarrhoea burden attributable to inadequate hygiene by region, 2019

Region	PAF	(95% CI)	Deaths	(95% CI)	DALYs	(95% CI)
Sub-Saharan	0.28	(0.23, 0.31)	140,058	(116,147,	9,618,663	(7,722,377,
Africa, LMIC				155,565)		11,285,958)
America, LMIC	0.22	(0.20, 0.25)	4,830	(4,301,	371,792	(331,350,
				5,487)		421,931)
Eastern	0.25	(0.13, 0.32)	37,489	(15,754,	2,487,854	(1,237,616,
Mediterranean,				51,020)		3,201,902)
LMIC						
Europe, LMIC	0.25	(0.16, 0.29)	1,014	(679, 1,201)	232,565	(147,527,
						276,334)
South-East Asia,	0.24	(0.18, 0.30)	187,773	(138,350-	6,762,433	(5,156,334,
LMIC				249,183)		8,692,073)
Western Pacific,	0.26	(0.23, 0.31)	7,331	(6,462,	784,762	(671,909,
LMIC				8,403)		922,322)
High-income	0.18	(0.15, 0.21)	5,290	(4,585,	237,797	(208,576,
countries				6,224)		277,249)
Total	0.26	(0.22, 0.28)	383,786	(322,397,	20,495,866	(17,557,832,
				445,046)		22,961,088)

PAF: population attributable fraction, CI: confidence interval, DALYs: disability-adjusted life years, LMIC: low- and middle-income countries; 183 low-, middle- and high-income countries, regional PAF aggregates relate to diarrhoea DALYs

Region	PAF	(95% CI)	Deaths	(95% CI)	DALYs	(95% CI)
Sub-Saharan	0.76	(0.72, 0.79)	377,170	(350,598,	25,918,546	(22,931,680,
Africa, LMIC				403,477)		29,713,824)
America,	0.52	(0.49, 0.57)	11,569	(10,631,	884,798	(814,503,
LMIC				12,634)		966,667)
Eastern	0.68	(0.60, 0.73)	100,905	(83,842,	6,598,741	(5,622,273,
Mediterranean,				118,038)		7,510,623)
LMIC						
Europe, LMIC	0.47	(0.40, 0.52)	1,976	(1,690,	444,995	(372,649,
				2,200)		493,610)
South-East	0.66	(0.57, 0.74)	521,292	(420,992,	18,784,186	(15,789,982,
Asia, LMIC				639,897)		22,156,116)
Western	0.58	(0.52, 0.64)	16,969	(15,290,	1,721,318	(1,521,799,
Pacific, LMIC				18,859)		1,952,561)
High-income	0.18	(0.15, 0.21)	5,290	(4,585,	237,797	(208,576,
countries				6,224)		277,249)
Total	0.69	(0.65, 0.72)	1,035,170	(929,178,	54,590,384	(50,033,488,
				1,159,750)		59,561,844)

Table S.16: Diarrhoea burden attributable to the cluster of inadequate drinking water, sanitation and hygiene by region, 2019

PAF: population attributable fraction, CI: confidence interval, DALYs: disability-adjusted life years, LMIC: low- and middle-income countries; 183 low-, middle- and high-income countries, regional PAF aggregates relate to diarrhoea DALYs

Table S.17: Burden of acute respiratory infections attributable to inadequate hygiene by region, 2019

Region	PAF	(95% CI)	Deaths	(95% CI)	DALYs	(95% CI)
Sub-Saharan	0.16	(0.13, 0.20)	124,856	(100,818,	8,305,466	(6,408,695,
Africa, LMIC				150,232)		10,602,986)
America,	0.12	(0.10, 0.15)	29,224	(23,486,	772,123	(637,583,
LMIC				38,290)		991,966)
Eastern	0.14	(0.08, 0.20)	28,295	(16,500,	1,928,087	(1,077,763,
Mediterranean,				39,487)		2,833,195)
LMIC						
Europe,	0.14	(0.08, 0.19)	11,819	(6783,	442,039	(262,473,
LMICs				16,337)		605,155)
South-East	0.13	(0.09, 0.19)	71,155	(49,405,	3,041,998	(2,079,221,
Asia, LMIC				103,679)		4,469,792)
Western	0.15	(0.12, 0.20)	51,557	(39.445,	1,470,715	(1,133,788,
Pacific, LMIC				69,130)		1,975,246)
High-income	0.09	(0.08, 0.12)	38,628	(32,552,	617,391	(525,496.
countries				49,372)		783,303)
Total	0.14	(0.13, 0.17)	355,533	(319,625,	16,577,818	(14,256,955,
				404,826)		19,481,266)

PAF: population attributable fraction, CI: confidence interval, DALYs: disability-adjusted life years, LMIC: low- and middle-income countries; 183 low-, middle- and high-income countries, regional PAF aggregates relate to diarrhoea DALYs

# 10. Comparison of WASH-attributable disease burden between WHO and the IHME Global Burden of Disease (GBD)

Year of estimation	WASH-attributable diarrhoeal deaths estimates (in 1000s) (PAF)		WASH-attributable ARI deaths estimates (in 1000s) (PAF)		
	WHO	GBD*	WHO	GBD*	
2010		337 (0.23)			
2012	842 (0.58)				
2013		1,399 (0.95)			
2015		1,766			
2016	829 (0.6)	1,481 (0.89)	370 (0.13)	179 (0.08)	
2017		1,422 (0.91)		188 (0.07)	
2019	1,035 (0.69)	1,387 (0.91)	356 (0.14)	270 (0.11#)	

#### Table S.18: WASH-attributable burden of diarrhoea and ARI estimates from the WHO and GBD, over time

WASH: drinking water, sanitation, hygiene; PAF: population attributable fraction; GBD: global burden of disease.

\* GBD PAFs calculated from attributable deaths divided by all deaths for that condition (both numbers as reported by the respective GBD study

*#* disease burden for lower respiratory infections

Table S.19: Counterfactual minimum exposure levels for WASH-attributable burden of diarrhoea and ARI estimates, over time and by institution

Year of estimation	Counterfactual minimum exposure level			
risk factor	WHO	GBD*		
2010 (GBD)/2012 (WHO)				
drinking water	household water treatment using	improved drinking water facilities		
	filters or boiling			
sanitation	improved sanitation facilities	improved sanitation facilities		
hygiene	handwashing with soap and water	[hygiene not covered as risk		
	after potential faecal contact	factor]		
2016				
drinking water	household water treatment using	high quality piped water that has		
	filters or boiling	been boiled or filtered before		
		drinking		
sanitation	basic sanitation in a	sanitation facilities with sewer		
	community>75% sanitation	connection or septic tank		
	coverage			
hygiene	handwashing with soap and water	access to handwashing station		
	after potential faecal contact	after any contact with excreta		
		including children's excreta		
2019				
drinking water	safely managed drinking water	high quality piped water that has		
	(adjusted for safely managed at	been boiled or filtered before		
	household level)	drinking		
sanitation	basic sanitation connected to	sanitation facilities with sewer		
	sewer	connection or septic tank		
hygiene	handwashing with soap and water	access to handwashing station		
	after potential faecal contact	after any contact with excreta		
		including children's excreta		

ARI: acute respiratory infections; GBD: global burden of disease, \* GBD provides WASH-attributable burden of disease estimates for other years (2013, 2015, 2017), which are not listed here, # hygiene-attributable ARI estimated in 2016 and 2019

# 11. List of diseases/adverse health outcomes that are linked to inadequate WASH or related risks

Table S.20: List of adverse health	outcomes link	ked to inadequate	e WASH,	, unsafe w	ater resource	management,
and unsafe recreational water use.						

Health outcome	Exposure	PAF	attributable deaths (in 1000s)	attributable DALYs (in 1000s)
Diarrhoeal diseases	Inadequate WASH	69%	1,035	54,590
Acute respiratory infections	Inadequate hygiene	14%	356	16,578
Undernutrition	Inadequate WASH	10%	8	825
Soil-transmitted helminthiasis	Inadequate WASH	100%	2	1,942
Trachoma	Inadequate WASH	100% <sup>7,8</sup>	-	194
Schistosomiasis	Inadequate WASH	43%-82% <sup>7,8</sup>	5	700
Malaria	Unsafe water resource management	42%-80% <sup>7,8</sup>	329	26,718
Lymphatic filariasis	Unsafe water resource management, unsafe water bodies	67%8	-	1,083
Onchocerciasis	Unsafe water resource management	10%8	-	121
Dengue	Unsafe water resource management, unsafe water bodies	95%8	29	1,855
Japanese encephalitis	Unsafe water resource management, unsafe water bodies	95%* <sup>8</sup>	no overall disease burden data	no overall disease burden data
Drowning	Unsafe recreational water use	not estimated/ insufficient data		
Arsenicosis	Unsafe drinking water	not estimated/ insufficient data		
Fluorosis	Unsafe drinking water	not estimated/ insufficient data		
Legionellosis	Contaminated aerosols from water sprays, jets or mists	not estimated/ insufficient data		
Leptospirosis	Unsafe water resource management, unsafe water bodies, unsafe drinking water	not estimated/ insufficient data		
Hepatitis A and hepatitis E	Inadequate WASH	not estimated/ insufficient data		
Intoxication from cyanobacteriae	Unsafe water resource management, inadequate drinking water	not estimated/ insufficient data		
Lead poisoning	Unsafe drinking water	not estimated/ insufficient data		
Scabies	Inadequate hygiene, inadequate water supply	not estimated/ insufficient data		

Spinal injury	Unsafe recreational	not estimated/	
	water use	insufficient	
		data	
Poliomyelitis	Inadequate WASH	not estimated/	
		insufficient	
		data	
Adverse neonatal	Inadequate WASH	not estimated/	
conditions and		insufficient	
maternal outcomes		data	
Cancer (bladder)	Unsafe drinking water	not estimated/	
	_	insufficient	
		data	
Musculoskeletal	Inadequate drinking	not estimated/	
diseases	water	insufficient	
		data	

Table adapted from <sup>9</sup>, includes results from <sup>8</sup>; \* PAF for environmental risks including inadequate WASH

### 12. Gather checklist

Table S.21: Checklist of information that should be included in new reports of global health estimates<sup>10</sup>

Item #	Checklist item	Reported on page #		
Objectives and funding				
1	Define the indicator(s), populations (including age, sex, and geographic entities), and time period(s) for which estimates were made.	panel 1; methods, 2 <sup>nd</sup> and 3 <sup>rd</sup> paragraph (above section on comparative risk assessment)		
2	List the funding sources for the work.	section: Role of the funding source		
Data I	nputs			
For a	Il data inputs from multiple sources that are synthesized as part of the	study:		
3	Describe how the data were identified and how the data were accessed.	method sections: "Population exposed", "Relative risks linking exposure and health outcome"; overall disease burden statistics and PAFs of malnutrition attributable to diarrhoea in Supplementary tables		
4	Specify the inclusion and exclusion criteria. Identify all ad-hoc exclusions.	Inclusion of disease outcomes: methods, 3 <sup>rd</sup> paragraph. Risk factor categorization: Methods, 2 <sup>nd</sup> paragraph.		
5	Provide information on all included data sources and their main characteristics. For each data source used, report reference information or contact name/institution, population represented, data collection method, year(s) of data collection, sex and age range, diagnostic criteria or measurement method, and sample size, as relevant.	Exposure data: Supplementary table, tab: "exposure_input", Supplementary material, section 6 Exposure-response data: Table 1, Supplementary material, section 6. Overall disease statistics: methods section "Population Attributable Fraction (PAF)		

		and attributable burden estimates", Supplementary material, section 6; supplementary tables with country disease
		burden statistics
6	Identify and describe any categories of input data that have potentially important biases (e.g., based on characteristics listed in	methods section "Population exposed", discussion section
Г	1tem 5).	on limitations
For a	lata inputs that contribute to the analysis but were not synthesized as p	oart of the study:
/	Describe and give sources for any other data inputs.	NA
For a		
8	Provide all data inputs in a file format from which data can be efficiently extracted (e.g., a spreadsheet rather than a PDF), including all relevant meta-data listed in item 5. For any data inputs that cannot be shared because of ethical or legal reasons, such as third-party ownership, provide a contact name or the name of the institution that retains the right to the data.	Supplementary tables; table
Data a	nalysis	
9	Provide a conceptual overview of the data analysis method. A diagram may be helpful.	Figure 1 on CRA, Figures 2- 4 for WASH analysis frameworks
10	Provide a detailed description of all steps of the analysis, including mathematical formulae. This description should cover, as relevant, data cleaning, data pre-processing, data adjustments and weighting of data sources, and mathematical or statistical model(s).	methods and supplementary materials
11	Describe how candidate models were evaluated and how the final model(s) were selected.	Supplementary material, section 2 and 3
12	Provide the results of an evaluation of model performance, if done, as well as the results of any relevant sensitivity analysis.	NA
13	Describe methods for calculating uncertainty of the estimates. State which sources of uncertainty were, and were not, accounted for in the uncertainty analysis.	Supplementary material, section "Uncertainty estimates at country, regional and global levels"
14	State how analytic or statistical source code used to generate estimates can be accessed.	Data sharing statement at the end of the manuscript
Result	s and Discussion	
15	Provide published estimates in a file format from which data can be efficiently extracted.	Supplementary tables with country burden of disease estimates
16	Report a quantitative measure of the uncertainty of the estimates (e.g. uncertainty intervals).	Confidence intervals are provided for all burden of disease estimates.
17	Interpret results in light of existing evidence. If updating a previous set of estimates, describe the reasons for changes in estimates.	Discussion, paragraphs 2, 3, 4; Supplementary material section 10
18	Discuss limitations of the estimates. Include a discussion of any modelling assumptions or data limitations that affect interpretation of the estimates.	Discussion, paragraphs 5 onwards.

## 13. Participants of the initial expert meeting on WASH-attributable disease burden estimates

"Consultancy on Assessment of the burden of disease attributable to water, sanitation and hygiene", London School of Hygiene and Tropical Medicine, 8-9 April 2013

• Prof. Jamie Bartram, Gillings School of Global Public Health, University of North Carolina at Chapel Hill, USA

- Prof. Sandy Cairneross, Department of Disease Control, London School of Hygiene and Tropical Medicine, UK
- Prof. Thomas Clasen, Department of Disease Control, London School of Hygiene and Tropical Medicine, UK
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- Mr Oliver Cumming, Department of Disease Control, London School of Hygiene and Tropical Medicine, UK
- Prof. Valerie Curtis, Department of Disease Control, London School of Hygiene and Tropical Medicine, UK
- Prof. Alan Dangour, Department of Population Health, London School of Hygiene and Tropical Medicine, UK
- Prof. Paul Hunter, Norwich Medical School, University of East Anglia, UK
- Dr Richard Johnston, Water Supply and Treatment, EAWAG, Switzerland
- Dr Lorna Fewtrell, Centre for Research into Environment and Health, Aberystwyth University, UK
- Mr Bruce Gordon, Water Sanitation and Hygiene, Department of Public Health and Environment, World Health Organization
- Ms Jennifer de France, Water Sanitation and Hygiene, Department of Public Health and Environment, World Health Organization
- Dr Annette Prüss-Üstün, Evidence and Policy for Environmental Health, Department of Public Health and Environment, World Health Organization
- Dr Sophie Bonjour, Evidence and Policy for Environmental Health, Department of Public Health and Environment, World Health Organization
- Dr Jennyfer Wolf, Evidence and Policy for Environmental Health, Department of Public Health and Environment, World Health Organization

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