

1 Rainfall as a driver of epidemic cholera: comparative  
2 model assessments of the effect of intra-seasonal  
3 precipitation events

4 Joseph Lemaitre<sup>a</sup>, Damiano Pasetto<sup>a</sup>, Javier Perez-Saez<sup>a</sup>, Carla Sciarra<sup>b</sup>,  
5 Joseph Francis Wamala<sup>c</sup>, Andrea Rinaldo<sup>a,d,\*</sup>

6 <sup>a</sup>*Laboratory of Ecohydrology, École Polytechnique Fédérale de Lausanne (EPFL), 1015*  
7 *Lausanne (CH)*

8 <sup>b</sup>*Dipartimento di Ingegneria dell'Ambiente, del Territorio e delle Infrastrutture,*  
9 *Politecnico di Torino, Corso Duca degli Abruzzi, 24, 10129 Torino, (IT)*

10 <sup>c</sup>*WHO South Sudan*

11 <sup>d</sup>*Dipartimento ICEA, Università di Padova, 35100 Padova (IT)*

---

12 **Abstract**

13 The correlation between cholera epidemics and climatic drivers, in particular  
14 seasonal tropical rainfall, has been studied in a variety of contexts owing to its  
15 documented relevance. Several mechanistic models of cholera transmission  
16 have included rainfall as a driver by focusing on two possible transmission  
17 pathways: either by increasing exposure to contaminated water (e.g. due  
18 to worsening sanitary conditions during water excess), or water contamina-  
19 tion by freshly excreted bacteria (e.g. due to washout of open-air defeca-  
20 tion sites or overflows). Our study assesses the explanatory power of these  
21 different modeling structures by formal model comparison using determinis-  
22 tic and stochastic models of the type susceptible-infected-recovered-bacteria  
23 (SIRB). The incorporation of rainfall effects is generalized using a nonlinear  
24 function that can increase or decrease the relative importance of the large  
25 precipitation events. Our modelling framework is tested against the daily

---

\*to whom correspondence should be sent

*Preprint submitted to Joseph Lemaitre@epfl.ch (Joseph Lemaitre), November 3, 2018*  
*damiano.pasetto@epfl.ch (Damiano Pasetto), javier.perezsaez@epfl.ch (Javier*  
*Perez-Saez), carla.sciarra@polito.it (Carla Sciarra), walamaj@who.int (Joseph*  
*Francis Wamala), andrea.rinaldo@epfl.ch (Andrea Rinaldo)*

*URL: <https://echo.epfl.ch/> (Andrea Rinaldo)*

26 epidemiological data collected during the 2015 cholera outbreak within the  
27 urban context of Juba, South Sudan. This epidemic is characterized by a  
28 particular intra-seasonal double peak on the incidence in apparent relation  
29 with particularly strong rainfall events. Our results show that rainfall-based  
30 models in both their deterministic and stochastic formulations outperform  
31 models that do not account for rainfall. In fact, classical SIRB models are  
32 not able to reproduce the second epidemiological peak, thus suggesting that  
33 it was rainfall-driven. Moreover we found stronger support across model  
34 types for rainfall acting on increased exposure rather than on exacerbated  
35 water contamination. Although these results are context-specific, they stress  
36 the importance of a systematic and comprehensive appraisal of transmission  
37 pathways and their environmental forcings when embarking in the modelling  
38 of epidemic cholera.

39 *Keywords:* epidemiological drivers, waterborne disease epidemics, Juba,  
40 South Sudan, environmental exposure

---

## 41 **1. Introduction**

42 Two main exposure pathways fuel cholera transmission across endemic  
43 and epidemic settings. First, as famously discovered by John Snow during  
44 the 1854 London cholera outbreak, an *indirect* exposure occurs from con-  
45 sumption of unsafe water contaminated by raw sewage [1]. Here, rainfall and  
46 the ensuing hydrologic transport processes play a major role in water con-  
47 tamination, for instance through the washout of open-air defecation sites and  
48 raw sewage circulation in the environment which is thought to have caused  
49 the revamping of the Haitian 2010 outbreak [2]. *Direct*, or human-to-human

50 exposure occurs when the bacteria is transmitted from an infected directly  
51 to a healthy person, for example via contaminated food. In this case, en-  
52 vironmental factors do not play a major role, except for possibly enhanced  
53 transmission due to (over-)crowding [3, 4]. It is known that the combination  
54 of environmentally-mediated and direct exposures shapes the spatio-temporal  
55 distribution of cases during cholera epidemics [5, 6, 7, 8].

56 The importance of climatic and environmental factors in the transmis-  
57 sion of cholera, namely temperature and rainfall, has been highlighted across  
58 settings. Indeed, the relationship between cholera and climate has long been  
59 studied, moving from seminal works linking cholera outbreaks to anomalies in  
60 the El Niño Southern Oscillation [9, 10] that have paved the way for a new  
61 field in epidemiological research. For large-scale infection patterns, many  
62 studies highlighted the role of climatic drivers on cholera dynamics, mostly  
63 focusing on climate change effects on disease spread [11, 12, 13, 14, 15, 16, 8]  
64 or on the impacts of spatial and temporal heterogeneities [17, 18, 19, 20, 21,  
65 22, 23]. While the effect of temperature on cholera transmission has been  
66 well unravelled, mainly regarding bacterial biology and ecology in natural  
67 environments, that of rainfall remains to be fully elucidated, possibly due  
68 to the multiple ways in which it can influence transmission at the local and  
69 regional scales [2, 24, 25]. Indeed, intense rainfall events have been shown  
70 to alter infection risk through a variety of potential mechanisms, including:  
71 flooding, leading to raw sewage contamination of water sources [26, 11]; in-  
72 creased hydrologic transport-driven iron availability in environmental waters  
73 that enhances pathogen survival and the expression of toxins [27, 28, 29]; dry  
74 spells inducing persistent low water levels leading to increased use of unsafe

75 water sources [30]; and crowding during strong flood events [17].

76 Most countries where associations between rainfall and cholera risk have  
77 been studied experience endemic cholera transmission [8]. Empirical studies  
78 have shown a range of correlations, both positive and negative, endowed  
79 with time lags ranging from weeks to months [26, 31, 12]. In general, rainfall  
80 has been found to enhance cholera transmission, but there is evidence that  
81 propagation buffer effects in wet regions may be due to pathogen dilution [26].  
82 Such variability reflects the variety of potential mechanisms whereby rainfall  
83 may alter infection risk. Similarly, a clear empirical correlation between  
84 intense rainfall and enhanced transmission is found in several regions hit by  
85 cholera epidemics [32, 30, 33, 34, 35]. The Haitian case, which has been in  
86 the midst of a major outbreak since October 2010 [36, 35, 37, 38, 39, 40, 41],  
87 has been studied under that angle, but its patterns have been argued to  
88 require a specific understanding [42]. Rainfall therein is empirically known  
89 to be directly associated with sudden resurgence of cholera infections via  
90 the analysis of reported cases [35], but a direct, causal relationship has only  
91 begun to be quantitatively examined [2, 24, 43]. Indeed, cholera case counts  
92 tend to rise sharply at the onset of seasonal heavy rains [44, 45, 46]. Notably,  
93 for the Haitian outbreak, such nexus has been addressed theoretically [2, 24].  
94 Results therein showed that at all spatial scales and locations examined, the  
95 tropical storms were significantly correlated with increased cholera incidence  
96 with lags of the order of a few days. As a consequence, accounting for  
97 the related forcing of dynamic models resulted in improved fits of reported  
98 incidence.

99 Properly incorporating the effects of rainfall in mathematical models of

100 cholera transmission is thus paramount to discriminate among the above-  
101 mentioned alternative transmission pathways, thus unlocking a predictive  
102 framework to evaluate the potentially rainfall-sensitive efficacy of available in-  
103 tervention strategies in endemic and epidemic settings including vaccination,  
104 antibiotics, and improved access to water sanitation and hygiene (WaSH),  
105 leveraging the numerous solutions that exist for rainfall forecasting [2, 43].  
106 This becomes critically important when evaluating the number of averted  
107 infections by deploying vaccines, as was done in the aftermath of the passage  
108 of Hurricane Matthew [40], or considering optimal deployment in space and  
109 time.

110 Rainfall has been accounted for in two main fashions in recent mathe-  
111 matical models of cholera. On one side, a contamination-centered approach  
112 suggesting that bursts of infections could be linked to increased contamina-  
113 tion of the water compartment [2]. This process conceptualizes the washout  
114 of open-air defecation sites by hydrologic transport. The same ‘transport’  
115 effect may be realized by sewer collectors’ overflows. In fact, both mecha-  
116 nisms have the net effect of charging progressively the bacterial concentration  
117 in the water reservoir [47]. Pathogens’ loads are washed out from a hydro-  
118 logic catchment enclosing human settlements and their infective individuals  
119 shedding bacteria. Therein, pathogen survival and thus the toxicity of their  
120 loads depend on hydrologic residence time distributions [2, 8]. Such loads  
121 increase as a function of rainfall, which acts as proxy of runoff volumes. The  
122 second approach is exposure-centered and employs a rainfall-dependent ex-  
123 posure rate subsuming both pathogen availability and the probability of the  
124 ingestion of contaminated water during wet spells [24]. Although both ap-

125 proaches are physically plausible, they have not been compared directly on  
126 the same datasets within a formal statistical framework, which would allow  
127 to highlight their respective merits and further recommendations for their  
128 use in different settings.

129 Here, we compare the explanatory power of these different types of rainfall-  
130 driven mechanistic models applied to a cholera outbreak in South Sudan.  
131 We quantitatively examine the link between rainfall and cholera during the  
132 outbreak recorded in Juba in 2015, when an intra-seasonal peak of cholera  
133 cases was recorded possibly in correspondence to intense precipitation events.  
134 The analysis of the lagged relationship between rainfall rates and revamped  
135 cholera incidence is addressed via dynamical compartmental models con-  
136 sidered both in deterministic and stochastic versions incorporating direct  
137 (human-to-human) and indirect (water-to-human) disease transmission, and  
138 rainfall effects on contamination and exposure.

139 This paper is organized as follows. The Materials and Methods section  
140 introduces to the data sets employed in our exercises and the general mod-  
141 eling rationale and framework. Results and a discussion follow, highlighting  
142 the role of rainfall in this specific case study with a view to the systematic  
143 comparative analysis of the importance of intra-seasonal precipitation events  
144 in epidemic and endemic cholera in other settings. A set of conclusions build-  
145 ing on the modelling results and suggesting the way forward closes then the  
146 paper.

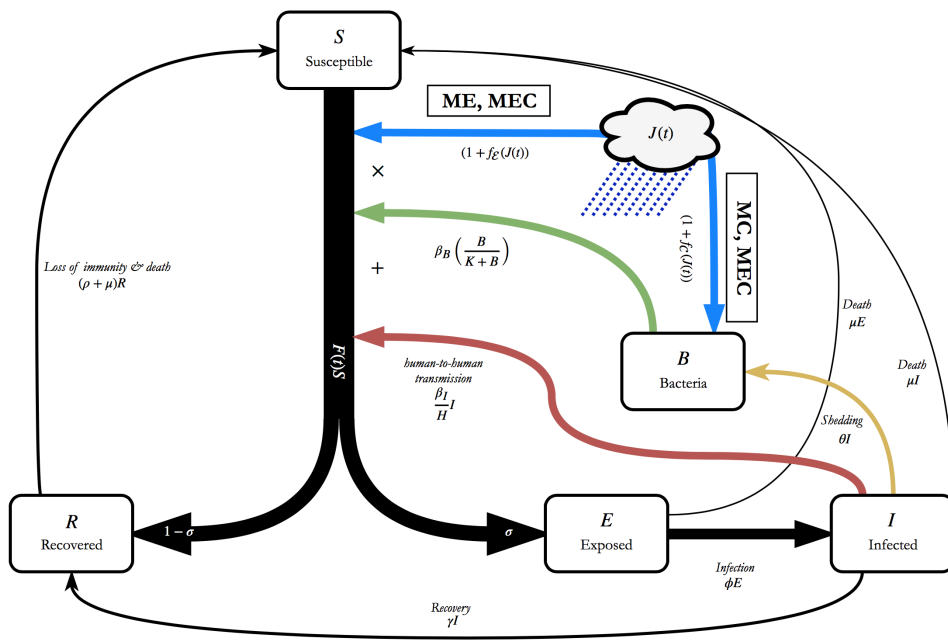


Figure 1: Flow diagram for our cholera models, with the different variations **ME**, **MR**, and **MEC** indicated.

## 147 2. Materials and methods

148 We base our analysis of the cholera outbreak in Juba on a general epi-  
149 demiological model that encompasses previous approaches to account for  
150 the effect of rainfall on cholera transmission. The proposed model builds  
151 on the classic susceptible  $S$ , infected  $I$ , and recovered  $R$  compartments for  
152 individuals, with an additional variable  $B$  describing the concentration of  
153 the bacteria in the environment (thus, the model is named SIRB). Previ-  
154 ous modelling exercises had considered rainfall intensity  $J(t)$  either to i)  
155 multiplicatively increase water contamination with bacteria shed by infected  
156 individuals [43, 40], or ii) assumed that the rainfall multiplicatively increases  
157 the exposure to contaminated water [24] (details on these two modelling  
158 frameworks are the Supplementary Information, SI, Section S.1). Aiming at  
159 a systematic comparison of the effect of rainfall through these two different  
160 transmission pathways, we here consider a generalized formulation of these  
161 cholera-forced models, wherein both formulations are nested.

162 Given the daily temporal resolution at which incidence data was available  
163 for the 2015 Juba’s epidemic (see Section 3), we here introduce in addition  
164 to the S-I-R-B variables, a compartment of exposed individuals  $E$ , to de-  
165 scribe the incubation period of the disease (from 12 h to 5 days [48]). This  
166 compartment is necessary to account the lag between the time of infection  
167 and the onset of the symptoms which result in reported cases. Moreover, in  
168 order to account for the vaccination campaigns that were deployed in Juba  
169 during August 2015, four compartments ( $V^S$ ,  $V^E$ ,  $V^I$ , and  $V^R$ ) are added to  
170 describe the dynamics of vaccinated individuals and their removal from the  
171 pool of susceptibles.



172 The proposed generalized cholera model is described in Figure 1, and  
 173 formulated as:

$$\frac{dE}{dt} = \sigma F(t)S - (\phi + \mu + \nu)E \quad (1)$$

$$\frac{dI}{dt} = \phi E - (\gamma + \mu + \alpha)I \quad (2)$$

$$\frac{dR}{dt} = (1 - \sigma)F(t)S + \gamma I - (\rho + \mu + \nu)R \quad (3)$$

$$\frac{dB}{dt} = -\mu_B B + \theta [1 + f_C(J(t))] (I + V^I) \quad (4)$$

$$\frac{dV^S}{dt} = \nu S - \mu V^S + \rho_v V^R - (1 - \eta)F(t)V^S \quad (5)$$

$$\frac{dV^E}{dt} = \nu E + \sigma(1 - \eta)F(t)V^S - (\phi + \mu)V^E \quad (6)$$

$$\frac{dV^I}{dt} = \phi V^E - (\gamma + \alpha + \mu)V^I \quad (7)$$

$$\frac{dV^R}{dt} = \nu R - (\mu + \rho_v)V^R + \gamma V^I + (1 - \sigma)(1 - \eta)F(t)V^S, \quad (8)$$

174 where  $F(t)$  takes into account both human-to-human transmission and non-  
 175 linear water-to-human transmission:

$$F(t) = \beta_B \left[ \frac{B}{K + B} \right] \left( 1 + f_\varepsilon(J(t)) \right) + \frac{\beta_I}{H} (I + V^I). \quad (9)$$

176 As few reliable data on changes in Juba's population are available for the  
 177 years of interest (2014 and 2015), the total population  $H$ , is assumed to be  
 178 constant, which implies that the number of susceptible individuals at time  $t$   
 179 is  $S(t) = H - I(t) - E(t) - R(t) - V^S(t) - V^E(t) - V^I(t) - V^R(t)$ . Individuals  
 180 are removed from the susceptible compartment  $S$  at rate  $F(t)$ , becoming ei-  
 181 ther symptomatically or asymptotically infected with probabilities  $\sigma$  and  
 182  $(1 - \sigma)$ , respectively. Symptomatically infected,  $I$ , shed *V. cholerae* into the  
 183 local watershed at rate  $\theta$ . The infectious period is governed by parameter  $\gamma$ ,

184 which determines the portion of infected individuals that enter the recovered  
 185 compartment  $R$ , joining the asymptomatic infected. Recovered individuals  
 186 return to the susceptible compartment at a rate  $\rho$ , describing the average  
 187 rate of loss of immunity for individuals that previously had been asymp-  
 188 tomatic or symptomatic infected. Parameters  $\alpha$  and  $\mu$  concern cholera re-  
 189 lated and unrelated death rates, respectively. The compartment  $B$  quantifies  
 190 the concentration of *V. cholerae* in the local (conceptualized) water reservoir,  
 191 which is used to estimate the probability of exposure to the contaminated  
 192 water in (9) through the term  $\beta_B \frac{B}{K+B}$ , where  $\beta_B$  is the maximum exposure  
 193 rate and  $K$  is the half-saturation constant of the dose-response function of  
 194 *V. cholerae* [47]. Parameter  $K$  is usually set to one by considering the change  
 195 of variable  $\tilde{B} = B/K$ . The parameter  $\mu_B$  expresses the rate of decay of bac-  
 196 teria in the environment. Exposed individuals become symptomatic infected  
 197 at a rate  $\phi$ , which corresponds to an average incubation period of  $1/\phi \approx$   
 198 1.5 days [48]. Functions  $f_C(J(t))$  and  $f_E(J(t))$  account for the rainfall effect  
 199 respectively by increasing the bacteria contamination in the water reservoir  
 200 (as the term  $\lambda J(t)$  in eq. S.3) or directly through amplifying the exposure  
 201 in the force of infection (as the term  $\lambda J(t)$  in eq. S.8).

202 With the objective of assessing the importance of rainfall on cholera trans-  
 203 mission, we here propose a generalization of the linear relation in eqs. (S.3)  
 204 and (S.8) by using a nonlinear function form for  $f_{C,E}(J(t))$ , reading:

$$f_{C,E}(J(t)) = \lambda_{C,E} \left( \frac{J(t)}{\max_t J(t)} \right)^{\alpha_{C,E}} \quad (10)$$

205 where the subscripts  $C, E$  respectively denote the effect of rainfall on expo-  
 206 sure and contamination,  $\max_t J(t)$  is the maximum recorded rainfall intensity  
 207 during the epidemic, and the  $\alpha_{C,E} \geq 0$  controls for the relative importance of

208 different rainfall intensities in their effect on the force of infection. Indeed,  
 209 since the ratio  $\frac{J(t)}{\max_t J(t)} \in [0, 1]$ , for  $\alpha_{\mathcal{C}, \mathcal{E}} \gg 1$  the ratio will tend to 0 for  
 210 all small precipitation events, leaving only the effect of the strongest events,  
 211 whereas for  $\alpha_{\mathcal{C}, \mathcal{E}} < 1$  all precipitation events will be assigned a similar weight  
 212 in the FOI. We also note that by setting  $\alpha_{\mathcal{C}, \mathcal{E}} = 1$  we recover the formu-  
 213 lations in eqs. (S.3) and (S.8). The flexibility allowed by (10) is therefore  
 214 able to discriminate between rainfall effects along a continuum from acting  
 215 on disease transmission regardless of intensity to a threshold-like effect for  
 216 the largest events which could be associated to severe flooding causing dam-  
 217 ages to the city's water and sanitary system, for instance leading to sewer  
 218 overflow. Note that in (9) precipitation enters in the term  $(1 + f_E(J(t)))$   
 219 which entails water-to-human transmission also when  $J(t) = 0$  (differently  
 220 from what happens in eq. S.8).

221 During the vaccination campaign, OCV doses are assumed to be dis-  
 222 tributed with equal rate  $\nu$  to susceptible, exposed and recovered individuals,  
 223 which enter the compartments  $V^S$ ,  $V^E$  and  $V^R$ . As the OCV provides only a  
 224 partial immunity having efficacy  $\eta$ ,  $0 \leq \eta \leq 1$ , vaccinated susceptibles ( $V^S$ )  
 225 can become exposed ( $V^E$ ) through a decreased force of infection of a fac-  
 226 tor  $(1 - \eta)$  with respect to non-vaccinated individuals. Vaccinated infected  
 227 individuals behave exactly like infected ones, but are placed in a different  
 228 compartment to exclude them from future vaccination campaigns. After re-  
 229 covering at rate  $\gamma$ , they lose their vaccine protection at rate  $\rho_v$ .

### 230 *2.1. Model comparison*

231 Here we assess the relevance of the two rainfall-driven transmission path-  
 232 ways by comparing the impact the results of the following models:

233 **MN** SIRB model without rainfall:  $\lambda_C = \lambda_E = 0, \beta_I = 0$ , as the null hypoth-  
234 esis for the importance of rainfall.

235 **MC** SIRB model using the approach described in eqs. (S.1-S.4) accounting  
236 for rainfall enhances the *contamination* of the water reservoir [2, 43]:  
237  $\lambda_E = 0, \beta_I = 0$ .

238 **ME** SIRB model using the formulation rainfall as in eqs. (S.5-S.8) where  
239 rainfall increases the *exposure* to bacteria [24]:  $\lambda_C = 0, \beta_I = 0$ .

240 **MEC** SIRB model combining both approaches **MC** and **ME** ( $\beta_I = 0$ ). Both  
241 ways of accounting rainfall play a role simultaneously.

242 For each model we explore the possibility of adding explicitly human-to-  
243 human transmission ( $\beta_I > 0$ ), which is indicated with an **H** at the end of the  
244 model name: **MNH**, **MCH**, **MEH**, and **MECH**. Table 1 summarizes the  
245 different parameters associated with the considered models.

246 The results of the 8 models that arise from this setting are compared  
247 on the basis of their ability to match the time series of daily reported cases  
248 during the cholera epidemic in Juba of 2015 (see Section 3). Table S.1 sum-  
249 marizes which parameters are calibrated for each model and their prior dis-  
250 tribution. The degrees of freedom of the models,  $n_p$ , vary from  $n_p = 7$  for  
251 **MN** to  $n_p = 12$  for **MECH**. Given the low number of daily reported cases  
252 and their ensuing variability, we also implement a stochastic equivalent of  
253 the deterministic ODE system (1-8) formulated as a continuous time par-  
254 tially observed Markov process model, accounting for both demographic and  
255 disease transmission stochasticities [49] (details Section S.2).

Model	$\lambda_c$	$\alpha_c$	$\lambda_\varepsilon$	$\alpha_\varepsilon$	$\beta_I$
<b>MN</b>	-	-	-	-	-
<b>MNH</b>	-	-	-	-	X
<b>MC</b>	X	X	-	-	-
<b>ME</b>	-	-	X	X	-
<b>MCH</b>	X	X	-	-	X
<b>MEH</b>	-	-	X	X	X
<b>MEC</b>	X	X	X	X	-
<b>MECH</b>	X	X	X	X	X

Table 1: Parameters considered in the eight compared models.  $\lambda$  and  $\alpha$  characterize the functional forms considering the precipitation (eq. 10).  $\beta_I$  is the exposure for human-to-human transmission.

256 Calibration of the deterministic model is performed using a Markov Chain  
257 Monte Carlo (MCMC)-based algorithm, which draws samples from the pos-  
258 terior distribution of the parameters. Inference on the stochastic model is  
259 performed using a frequentist multiple iterated filtering algorithm. Both  
260 model were fit against the daily reported cases accounting for over- or under-  
261 reporting, and assuming a Poisson distribution. Models were then compared  
262 using the Bayesian Information criterion (BIC), Bayes factors, and the like-  
263 lihood ratio test for the nested models (details given in Sections S.3 and  
264 S.4).

### 265 **3. Case study: the 2015 Juba’s outbreak**

266 In the past years, South Sudan had been struck by several cholera out-  
267 breaks (details in S.5). Here we focus on the analysis of the outbreak in  
268 Juba during 2015, when a particular double peak of cholera cases occurred,  
269 probably associated to a strong intra-seasonal precipitation event (Figure 2).  
270 Epidemiological records for the 2014 and 2015 cholera epidemics included  
271 cholera cases and hospitalization time series at the second-lowest administra-  
272 tive level (named Payams in South Sudan) as reported by SSMoH [50, 51, 52].  
273 The recorded cases in the 7 Payams that constitute the administrative area of  
274 Juba have been aggregated to obtain the reported time series for the county  
275 level. We assigned the population values for Juba county as in the official  
276 projections provided by the South Sudan National Bureau of Statistics [53].  
277 These values were validated against the growth rate value stated by CIA [54].  
278 Daily rainfall estimates [ $\text{mm d}^{-1}$ ] were downloaded from the Climate Data  
279 Library (National Oceanic and Atmospheric Administration, NOAA) [55] for  
280 the years 2014 and 2015, with spatial resolution of  $0.1^\circ$  (approximately 10  
281 km at the equator). The precipitation considered in the model have been  
282 spatially averaged over the study area (see Figure 2). For what concerns the  
283 implementation of vaccines in 2015, we considered the 167’377 OCV doses  
284 that were distributed in the county of Juba [52] during 6 days of a mass  
285 vaccination campaign started in July 31, 2015. Additional details about the  
286 model setup can be found in [56].

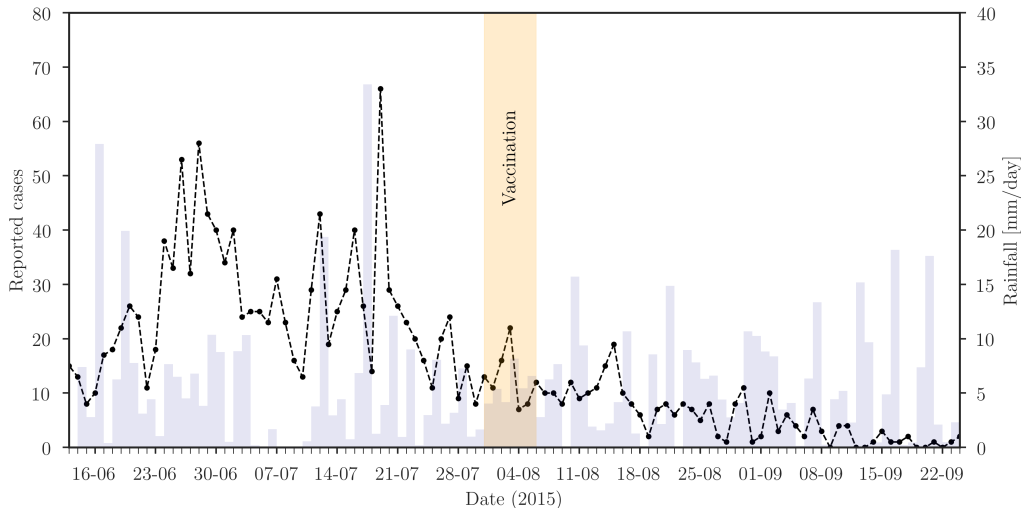


Figure 2: Reported cholera cases (dots) versus precipitation (bars) during the 2015 epidemic in Juba. The timing of the vaccination campaign is highlighted in yellow.

287 *3.1. Initial conditions*

288 The past history of cholera epidemics in South Sudan, and particularly in  
 289 Juba (see Section S.5), plays an important role in the determination of the  
 290 size of susceptible and recovered compartments at the beginning of the 2015  
 291 epidemic. These two compartments are also largely impacted by the rate of  
 292 immunity loss,  $\rho$ , of the recovered individuals, which determines the duration  
 293 of the stay in the  $R$  compartment (from few months to several years), and  
 294 the probability of asymptomatic infected,  $1 - \sigma$ , which determines how many  
 295 asymptomatics enter the  $R$  compartment (values in literature range between  
 296  $\sigma = 0.5$ , meaning one asymptomatic per each symptomatic infected, to less  
 297 than  $\sigma = 0.01$ , corresponding to more than 99 asymptomatic infected per  
 298 each symptomatic infected [57]. The initial conditions must therefore be  
 299 estimated for each parameter set considered during calibration.

300 To take into account the uncertainty associated with the past epidemics  
301 and vaccination campaigns, the initial number of temporary immune indi-  
302 viduals,  $R_0$ , in April 2014 is calibrated for each model.

303 The detailed daily data of suspected cases during 2014 is used to estimate  
304 the associated number of recovered individuals. These undergo an exponen-  
305 tial decay with average time of immunity loss  $1/\rho$  (similarly to what done  
306 in [40]), thus obtaining a consistent estimate of the recovered in June 2016.  
307 Simulations are then initialized on the 5<sup>th</sup> of June 2015 considering two  
308 exposed individuals, two infected, and the associated steady-state bacteria  
309 concentration.

## 310 4. Results

### 311 4.1. Selection of rainfall effects on transmission pathways

312 The summary statistics of the deterministic and stochastic models con-  
313 sidered in the study are given in Table 2. Overall, the stochastic models  
314 outperform their deterministic counterparts for all model structures by  $\approx 40$   
315 log-likelihood units. Both model types agree in the significance of rainfall in  
316 explaining the time series of daily reported cases, in particular through the in-  
317 creased exposure pathway, although the specific ordering of the models differs  
318 between model types. Indeed, the BFs for the deterministic models suggest a  
319 strong support for model **MEC**, followed by model **ME** ( $BF_{ME,MEH}^{-1} = 0.16$ ),  
320 with very little support for all other models ( $BF_{\cdot,MEH}^{-1} < 10^{-2}$  for all other  
321 models than ME). For the stochastic model, the BFs estimated with the BIC  
322 suggest the strongest support for model **ME**, with the basic SIRB model  
323 coming in second with 5 times less evidence ( $BF_{MN,ME}^{-1} \approx 0.15$ ). When



324 considering only the BIC, model **ME** ranks first for both the determinis-  
325 tic and the stochastic formulations. Interestingly, all models that include  
326 human-to-human transmission present smaller or equal log-likelihoods than  
327 their counterparts with only the bacteria compartment, which suggests that  
328 the data does not support both environmental and human-to-human trans-  
329 mission within the set of the models we considered here.

330 The results of the nested LR-tests confirm the statistical significance of  
331 including rainfall in the cholera transmission models, with the effect on ex-  
332 posure better supported by the data in both model types than the effect on  
333 contamination. In the deterministic case, the extension of the basic SIRB  
334 (model **MN**) with rainfall effects were significant for all direct comparisons  
335 (Fig. 3 A). The addition of human-to-human transmission was not signifi-  
336 cant mostly due to the above-mentioned lower estimate of the log-likelihood  
337 in these models. When considering only a single effect of rainfall (either  
338 increasing exposure or contamination), **ME** outperforms **MC** in terms of  
339 likelihood for the same number of parameters. Interestingly, the inclusion  
340 of rainfall-induced contamination in model **ME** is rejected due to the very  
341 limited increase of the estimated log-likelihood of **MEC**, in contrast with  
342 the BFs favouring the latter. Model **ME** is thus the one retained by the  
343 LR-tests in the deterministic set of models. In the case of the stochastic  
344 models, the LR-tests also highlight the importance of the effect of rainfall on  
345 exposure rather than on contamination (Fig. 3 B). In fact, the much stronger  
346 performance of **MN** in comparison with its deterministic counterpart rela-  
347 tive to all other model structures imposes a stronger condition for retaining  
348 additional transmission processes. Indeed, both models **MC** and **MCH** were

Model	Deterministic				Stochastic			
	$n$	$\hat{\ell}$	BIC	$BF^{-1}$	$n$	$\hat{\ell}$ (s.e.)	BIC	$BF^{-1}$
<b>MN</b>	7	-368.62	770.27	3.1E-05	8	-326.45 (0.105)	690.65	1.5E-01
<b>MNH</b>	8	-368.95	775.64	1.1E-09	9	-327.52 (0.052)	697.51	4.7E-03
<b>MC</b>	9	-358.32	759.11	5.5E-03	10	-323.50 (0.037)	696.01	2.5E-02
<b>MCH</b>	10	-359.06	765.30	1.7E-04	11	-324.89 (0.041)	701.68	5.9E-04
<b>ME</b>	9	-356.96	<b>756.40</b>	1.6E-01	10	<b>-319.81</b> (0.035)	<b>687.38</b>	<b>1</b>
<b>MEH</b>	10	-358.06	763.30	6.3E-04	11	-320.64 (0.030)	693.18	4.1E-02
<b>MEC</b>	11	<b>-356.87</b>	765.64	<b>1</b>	12	-320.17 (0.031)	696.96	6.2E-03
<b>MECH</b>	12	-357.55	771.73	2.4E-06	13	-320.38 (0.024)	702.09	4.8E-04

Table 2: Model comparison statistics. We report for each model its number of parameters  $n$ , the associated estimated log-likelihood  $\hat{\ell}$  (and its Monte Carlo standard error for the stochastic model), and the inverse of the Bayes Factor ( $BF^{-1}$ ) with respect to the model with the largest evidence. The BFs for the deterministic models were computed directly from the parameters posteriors, whereas for the stochastic models they were estimated with the Bayesian Information Criterion (BIC) as  $BF_i \approx e^{\frac{1}{2}(BIC_i - BIC_{min})}$ . The BIC for the deterministic models was computed using the maximum log-likelihood value visited with the MCMC algorithm across chains. Best values in each column are indicated in bold.

349 rejected when compared to **MN**, thus only models with rainfall-driven ex-  
350 posure were retained. As in the deterministic case, model **ME** is the one  
351 finally retained due to the lack of significance of the inclusion of additional  
352 transmission processes. We here note that the conclusion based on the LR-  
353 tests for the deterministic models should be taken with caution because the  
354 MCMC algorithm used for calibration does not directly aim at maximizing  
355 the likelihood, but rather at sampling from the posterior distribution of the  
356 parameters given the data and the model. Moreover, the best likelihood  
357 visited by the chains when sampling the posteriors that we here use in the  
358 LR-tests is not a formal estimate of the models' likelihood. However, the  
359 fact that the LR-tests applied to both model types agree with the selection  
360 of **ME** supports their use in both cases.

361 Both statistical methods for model comparison therefore agree about the  
362 importance of the effect of intra-seasonal rainfall on the exposure to trans-  
363 mission during the 2015 cholera epidemic in Juba. For the deterministic  
364 type of models the BFs suggest a stronger support for model **MEC**, and  
365 the LR-tests for **ME**, whereas for the stochastic models both the BIC-based  
366 estimates of BFs and the LR-tests favor **ME**.

#### 367 *4.2. Intra-seasonal rainfall events and the 2015 Juba epidemic*

368 The comparison between the estimated output cases computed by the  
369 basic SIRB model (**MN**) and the most significant rainfall-based processes  
370 (**MEC** and **ME** for the deterministic and stochastic types, respectively)  
371 highlight the importance of rainfall in retrieving the second epidemiological  
372 peak (Figure 4). Both deterministic and stochastic SIRB fit well the gen-  
373 eral trend of the data, but they clearly underestimate the large number of

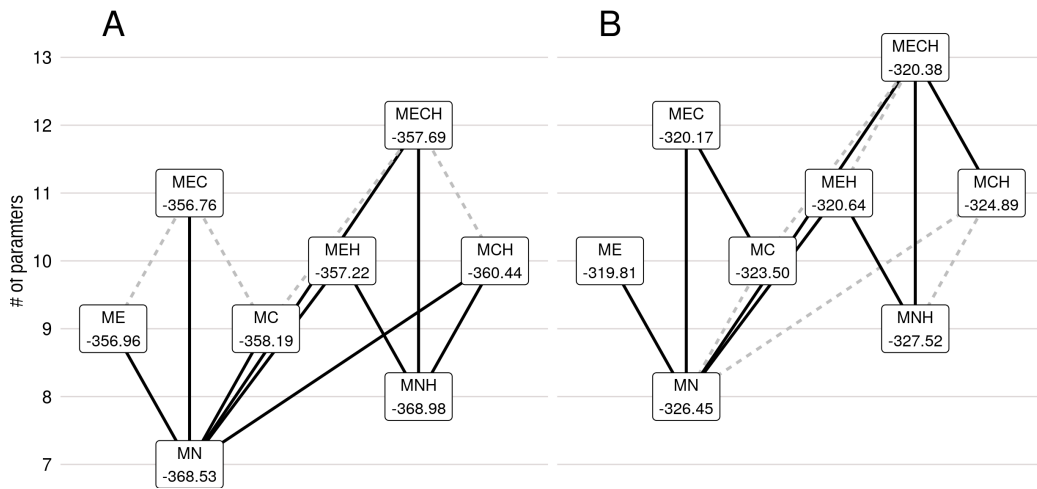


Figure 3: Likelihood ratio tests of model nesting. The LL-tests were computed for each nested pair of models  $\{\mathcal{M}_0, \mathcal{M}_1\}$ , with parameter vectors  $\theta^0, \theta^1$ , for which at least one of the parameters that is not null in  $\theta^1$  is equal to 0 in  $\theta^0$ . Each model is labeled with its associated estimated maximum log-likelihood value,  $\hat{\ell}$ , for the deterministic (A) and the stochastic (B) models, and linked based on whether the likelihood ratio is significantly (full black lines) or not (dashed gray lines) at the 5% level. The absence of lines indicates a lower  $\hat{\ell}$  for the more complex model.

374 reported cases on the 19th of July (65 cases). Instead, the more complex  
375 models **ME** and **MEC** follow the SIRB dynamics and then are forced by the  
376 precipitation occurred in the 18th of July (33 mm/d) toward the epidemio-  
377 logical peak.

378 Model calibration results suggest that precipitations with smaller inten-  
379 sities did not have a strong impact on cholera transmission during the 2015  
380 epidemic in Juba. Indeed, the exponents  $\alpha_C$  and  $\alpha_E$  were found to be sys-  
381 tematically larger than 1 (as shown by posteriors of the deterministic models  
382 in Figure S.2 and the Monte Carlo confidence intervals for the stochastic  
383 **ME** in Figure S.4 of the SI). Thus, in the considered epidemic, the nonlinear  
384 function used to account for rainfall in the model (eq. 10) helps isolating the  
385 contribution of the largest rainfall.

386 The best measures of fit computed for the stochastic **ME** (see Table 2)  
387 are thus explained by a larger sensitivity to precipitation, which causes the  
388 match between the mean of the simulated cases and the data during the  
389 second peak.

390 Comparing the two model types, stochastic results have a larger 95%  
391 confidence interval, which better encompass most of the data. In particular,  
392 both epidemiological peaks are well captured by the stochastic models, while  
393 the deterministic results systematically underestimate them. Two factors  
394 contribute to this result: the intrinsic stochastic nature of the model, that  
395 requires the simulation of various model runs for the same set of parameters,  
396 and the noise that necessarily perturbs the force of the infection yielding an  
397 overdispersion in infections. The standard deviation of such (assumed white)  
398 noise is estimated in each stochastic model, and it is interesting to note that

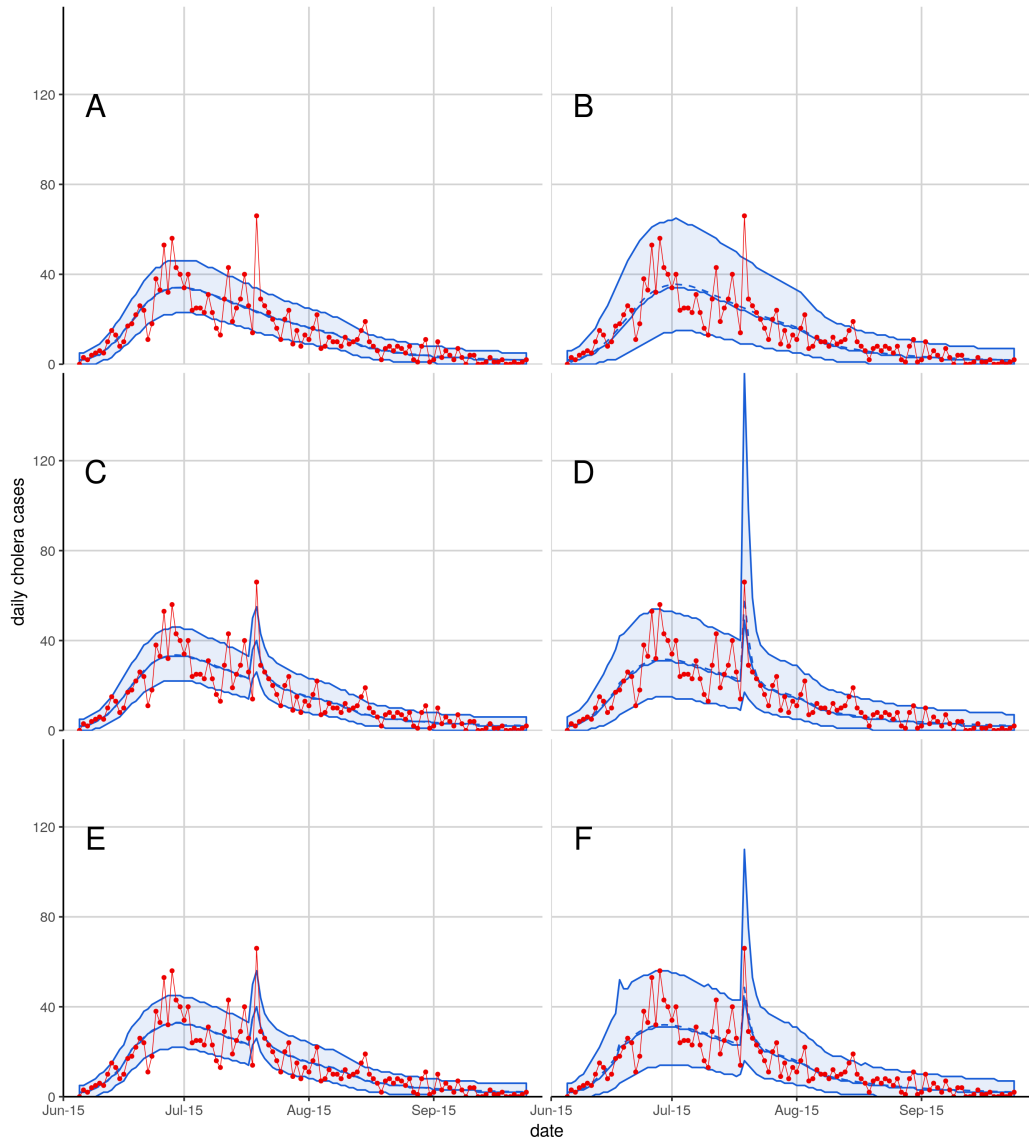


Figure 4: Simulations of the **MN** (A-B), **ME** (C-D) and **MEC** (E-F) models. Simulations for the deterministic versions (A,C,E) are given by the mean (blue dashed line), median (blue full line) and 95% simulation envelopes (blue ribbon) of 100 simulations of the measurement model for each trajectory from 100 samples from the posteriors of model parameters against reported daily cholera cases (red line and dots). Simulations from the stochastic models (B, D, F) are given for 10'000 simulations of the stochastic process and measurement models.

399 the MLE obtained for **ME** is slightly smaller than in **MN** (0.028 versus  
400 0.022), highlighting again that the data are retrieved with a lower uncertainty  
401 when rainfall is included in the model. This is evident in Figure 4, where  
402 the width of the 95% confidence interval of models **ME** and **MEC** is smaller  
403 with respect to **MN**.

404 Finally, despite having different BFs, the deterministic models **ME** and  
405 **MEC** are qualitatively similar in terms of output response, indicating that  
406 the recorded changes in the log-likelihood function do not correspond to  
407 qualitative changes in the output.

## 408 **5. Discussion and conclusions**

409 In this study we developed a general mechanistic SIRB-based epidemi-  
410 ological model to evaluate the relevance of rainfall in the amplification of  
411 cholera transmission, focusing on the 2015 Juba outbreak. Two rainfall-based  
412 transmission processes were compared: the direct increase of the exposure to  
413 the contaminated water (model **ME**) following [24], and the increase of water  
414 contamination by flooded open defecation sites (model **MC**) following [2]. In  
415 addition, we also considered human-to-human transmission (models' name  
416 with **H**).

417 Regarding the epidemiological model, this study introduced two innova-  
418 tions with respect to previous modeling attempts of cholera epidemics (see,  
419 e.g., [43, 40]. First, the focus on daily incidence data, as opposed to weekly  
420 epidemiological reports commonly used in modelling works, motivated the  
421 introduction of a compartment of exposed individuals (eq. 1) to account for  
422 the incubation period of the disease and, thus, the lag between the possibly

423 rainfall-driven infection process and the manifestation of the symptoms re-  
424 sulting in the timeseries of daily reported cases at our disposal. Second, a  
425 non-linear version of rainfall driver, in the form of a power-law controlled by  
426 a single parameter, was introduced to generalize the previous linear depen-  
427 dence. Such formulation has the flexibility to either emphasize the impact  
428 of the largest rainfall events, or give equal weight to all non-zero rainfall  
429 intensities.

430 All model assumptions were compared for both deterministic and stochas-  
431 tic model's types, in order to draw more general conclusions. The statistics  
432 and tests used to compare the model results (Table 2 and Figure 3) sup-  
433 ported the significance of rainfall effects during the 2015 epidemic in Juba.  
434 In fact, results showed that for both model types there exists a significant  
435 positive effect of including rainfall drivers, in particular because standard  
436 SIRB models were not able to reproduce the second epidemiological peak  
437 of reported cases occurred in July during the recession period. All models  
438 considering rainfall, instead, showed an increase of the number of cases in  
439 correspondence of the second epidemiological peak, which was due to the  
440 large rainfall rates occurred during the previous days (Fig. 4). This differ-  
441 ence in the simulated responses of models considering (or not) rainfall lead  
442 to stronger support for rainfall-based models. Due to the small variations  
443 among the likelihoods of rainfall-based models, however, (Table 2), it is not  
444 straightforward to draw conclusions on the best way to include the rainfall  
445 effect. Models with the minimum BIC were those considering the increase in  
446 exposure (model **ME**) for both the stochastic and the deterministic model  
447 types. For the deterministic models, the computation of the Bayes Factors



448 (BFs), which should provide a direct estimation of the model probability,  
449 suggested the selection of the model combining exposure and contamina-  
450 tion processes (model **MEC**). However, this information criterion might be  
451 unstable due to numerical issues and oscillations in the MCMC used for cal-  
452 ibration [58]. By considering the fact that the models' outputs were similar  
453 for **MEC** and **ME** (4), we advise to select the approach endowed with less  
454 parameters, in this case **ME**, as indicated by the BIC. Note that the inclu-  
455 sion of human-to-human transmission was not statistically relevant in this  
456 modeling exercise.

457 The comparison between the likelihoods of the two models' types (de-  
458 terministic and stochastic) showed that considering the stochasticity of the  
459 processes improves the model results (Table 2). This suggests that also de-  
460 terministic models should include a stochastic term in the computation of  
461 the force of infection (eq. 9), which might increase the flexibility of the  
462 outputs.

463 Several limitations should be considered when analyzing the present re-  
464 sults. The calibration exercise attempted in this study considered daily rain-  
465 fall and cholera reported cases, which are characterized by significant random  
466 fluctuations that might partly cloud the description of the underlying infec-  
467 tion processes. Small random delays in reporting could change the infection  
468 curve and thus the effect of rainfall. This issue was partially addressed by  
469 considering the exposed compartment (1) for simulation of the incubation  
470 period, and unknown reporting rate  $\epsilon$  for the observed cases.

471 Here, we strove to reproduce the epidemic by modelling epidemiological  
472 transmission processes. While we took into account non-linear rainfall effects

473 and possible over-reporting, we did not consider human mobility effect [59,  
474 60, 61, 23], which could help modeling the arrival of infected individual.  
475 Moreover, in our model asymptomatic infected individuals did not contribute  
476 to the bacterial concentration in the environment, while they might impact  
477 the infection cycle due to the presence of bacteria in their feces. From a  
478 modeling viewpoint, these unaccounted processes were compensated by the  
479 calibration procedure, at the loss of predictive power.

480 The prior bounds to be assigned to parameters are typically wide [62]  
481 because the rates governing transmission processes are highly dependent on  
482 the specific epidemiological context, so that somewhat contradictory values  
483 had been estimated in literature. These considerations, together with the  
484 intrinsic noise affecting recorded cases, underlie the possibility that some of  
485 the model parameter might be unidentifiable [63], in the sense that different  
486 parameter combinations would yield the same model output (also called equi-  
487 finality). The exploration of the posterior parameter distribution using an  
488 MCMC approach allowed us to evince the possible correlation among paramete-  
489 rs that were well identified by the data, with the main risk of the algorithm  
490 getting trapped in a local minimum of the fitness landscape (the distribution  
491 of parameters). The posterior probability distributions of the model paramete-  
492 rs (see SI, Section S4) are associated with the model uncertainty, and were  
493 here explored by the chains of the MCMC calibration.

494 The lack of available data prevented us to include the effects of the overall  
495 efforts towards WaSH improvement in this study. This assumption is reason-  
496 able in the case at hand, however, given the short time-frame of the study.  
497 Despite these limitations, our model comparison using both a deterministic

498 and a stochastic model gave coherent results. The agreement of the two  
499 modeling types strengthened our results regarding the importance of rainfall  
500 patterns to significantly affect the development of cholera cases in time.

501 Overall, the findings of the study are consistent with the lessons learned  
502 in South Sudan with most of the transmission starting with the onset of the  
503 rainy season. In 2016 and 2017, cases in the dry season were observed and  
504 associated to the overexploitation of scarce water resources by nomadic herds-  
505 men (cattle camps). This suggests that, as already observed, a general as-  
506 sessment of the relationship between precipitation and general waterborne or  
507 water-based disease infections is far from obvious and surely case-dependent.  
508 It has been argued, for example, that in the domain of water-based parasitic  
509 infections (see e.g. [64, 8]) rainfall could not only boost disease transmis-  
510 sion (especially in dry climates where it is a key driver of habitat formation  
511 for possible intermediate hosts) but also reduce it substantially, e.g. by in-  
512 creasing water flow (which in turn decreases habitat suitability for both the  
513 intermediate and the human hosts). Rainfall patterns may also drastically  
514 affect human activities related to water contacts, thus potentially altering  
515 exposure and transmission risk [65]. To that end, a hydrology-driven assess-  
516 ment cannot ignore certain characteristics, in particular the ephemeral or  
517 permanent nature of the waterways fostering contacts among pathogens and  
518 hosts [66]. Also, temporal fluctuations of rainfall patterns may be particularly  
519 important in determining the seasonality of transmission [67, 68, 69, 66].

520 Subsuming the results obtained by this major computational exploration  
521 focused on the analysis of Juba's 2015 epidemics, we conclude that rainfall  
522 patterns are fundamental drivers for epidemic cholera models, whether de-

523 terministic or stochastic, not only to capture seasonal trends, but also to  
524 describe short-term fluctuations in the number of reported cases.

## 525 **Acknowledgements**

526 JL, JPS, DP, and AR acknowledge funds provided by the Swiss National  
527 Science Foundation ([www.snf.ch](http://www.snf.ch)), via the projects: "Dynamics and controls  
528 of large-scale cholera outbreaks" (DYCHO CR23I2 138104) and "Optimal  
529 control of intervention strategies for waterborne disease epidemics" (200021-  
530 172578). The authors wish to thank Andrew Azman, Johns Hopkins Univer-  
531 sity, for insightful discussions on data collection and model inputs.

## 532 **References**

- 533 [1] J. Snow, On the Mode of Communication of Cholera, John Churchill,  
534 1855.
- 535 [2] A. Rinaldo, E. Bertuzzo, L. Mari, L. Righetto, M. Blokesch, M. Gatto,  
536 R. Casagrandi, M. Murray, S. Vesenbeckh, I. Rodriguez-Iturbe, Re-  
537 assessment of the 2010-2011 Haiti cholera outbreak and rainfall-driven  
538 multiseason projections, Proceedings of the National Academy of Sci-  
539 ences USA 109 (17) (2012) 6602–6607.
- 540 [3] E. Boelee, Y. Mekonnen, J. N. Poda, M. McCartney, P. Checchi, S. Ki-  
541 bret, F. Hagos, H. Laamrami, Options for water storage and rainwater  
542 harvesting to improve health and resilience against climate change in  
543 Africa, Regional Environmental Change 13 (2013) 509–519.

- 544 [4] F. Finger, T. Genolet, L. Mari, G. C. de Magny, N. M. Manga, A. Ri-  
545 naldo, E. Bertuzzo, Mobile phone data highlights the role of mass gath-  
546 erings in the spreading of cholera outbreaks, Proceedings of the National  
547 Academy of Science USA 113 (23) (2016) 6421–6426.
- 548 [5] J. D. Sugimoto, A. A. Koepke, E. E. Kenah, M. Halloran, F. Chowdhury,  
549 et al., Household Transmission of *Vibrio cholerae* in Bangladesh, PLoS  
550 Neglected Tropical Diseases 8 (2014) e3314.
- 551 [6] Q. Bi, A. S. Azman, S. M. Satter, A. I. Khan, D. Ahmed, et al., Micro-  
552 scale spatial clustering of cholera risk factors in urban Bangladesh, PLoS  
553 Neglected Tropical Diseases 10 (2016) e0004400.
- 554 [7] J. Lessler, H. Salje, M. Grabowski, D. Cummings, Measuring spatial  
555 dependence for infectious disease, Epidemiology 11(5) (2016) e0155249.
- 556 [8] A. Rinaldo, E. Bertuzzo, L. Mari, M. Blokesch, M. Gatto, Mod-  
557 eling key drivers of cholera transmission dynamics provides new  
558 perspectives for parasitology, Trends in Parasitology (2017) 1–  
559 13doi:10.1016/j.pt.2017.04.002.
- 560 [9] R. Colwell, Global climate and infectious disease: the cholera paradigm,  
561 Science 274 (1996) 2025–2031.
- 562 [10] M. Pascual, X. Rodo, S. Ellner, R. Colwell, M. Bouma, Cholera dynam-  
563 ics and El Nino-Southern oscillation, Science 289 (2000) 1766–1769.
- 564 [11] M. Hashizume, B. Armstrong, S. Hajat, Y. Wagatsuma, A. S. Faruque,  
565 T. Hayashi, D. A. Sack, The effect of rainfall on the incidence of cholera  
566 in Bangladesh, Epidemiology 19 (1) (2008) 103–110.

- 567 [12] G. Constantin de Magny, W. Thiaw, V. Kumar, N. Manga, B. M. Diop,  
568 L. Gueye, M. Kamara, B. Roche, R. Murtgudde, R. R. Colwell, Cholera  
569 outbreak in Senegal in 2005: Was climate a factor?, PLoS ONE 7 (2012)  
570 e44577.
- 571 [13] M. Hashizume, L. F. Chaves, A. S. G. Faruque, A differential effect of  
572 Indian Ocean Dipole and El Niño on cholera dynamics in Bangladesh,  
573 PLoS ONE 8 (2013) e60001.
- 574 [14] X. Rodó, M. Pascual, F. J. Doblas-Reyes, A. Gershunov, D. A. Stone,  
575 F. Giorgi, P. J. Hudson, J. Kinter, M.-A. Rodríguez-Arias, N. C.  
576 Stenseth, D. Alonso, J. García-Serrano, A. P. Dobson, Climate change  
577 and infectious diseases: Can we meet the needs for better prediction?,  
578 Climatic Change 118 (2013) 625–40.
- 579 [15] I. J. Ramírez, S. C. Grady, El Niño, climate, and cholera associations in  
580 Piura, Peru, 1991–2001: A wavelet analysis, EcoHealth 13 (2016) 83–99.
- 581 [16] L. Vezzulli, C. Grande, P. C. Reid, P. Hélaouët, M. Edwards, M. G.  
582 Höfle, I. Brettar, R. R. Colwell, C. Pruzzo, Climate influence on *Vib-*  
583 *rio* and associated human diseases during the past half-century in the  
584 coastal North Atlantic, Proceedings of the National Academy of Sciences  
585 USA 113 (2016) 5062–5071.
- 586 [17] R. C. Reiner, A. A. King, M. Emch, M. Yunus, A. Faruque, M. Pascual,  
587 Highly localized sensitivity to climate forcing drives endemic cholera in  
588 a megacity, Proceedings of the National Academy of Sciences 109 (6)  
589 (2012) 2033–2036.

- 590 [18] C. Baker-Austin, J. A. Trinanes, N. G. H. Taylor, R. Hartnell, A. Si-  
591 itonen, J. Martinez-Urtaza, Emerging *vibrio* risk at high latitudes in  
592 response to ocean warming, *Nature Climate Change* 3 (2013) 73–77.
- 593 [19] L. Vezzulli, R. Colwell, C. Pruzzo, Ocean warming and spread of  
594 pathogenic vibrios in the aquatic environment, *Microbial Ecology* 65  
595 (2013) 817–825.
- 596 [20] B. Cash, X. Rodó, M. Emch, M. Yunus, A. Faruque, M. Pascual, Cholera  
597 and shigellosis: Different epidemiology but similar responses to climate  
598 variability, *PLoS ONE* 9 (2014) e107223.
- 599 [21] L. E. Escobar, S. J. Ryan, A. M. Stewart-Ibarra, J. L. Finkelstein, C. A.  
600 King, H. Qiao, M. E. Polhemus, A global map of suitability for coastal  
601 *Vibrio cholerae* under current and future climate conditions, *Acta Trop-*  
602 *ica* 149 (2015) 202–211.
- 603 [22] L. Vezzulli, E. Pezzati, I. Brettar, M. Höfle, C. Pruzzo, Effects of global  
604 warming on *Vibrio* ecology, *Microbiology Spectrum* 3 (2015) 0004–2014.
- 605 [23] J. Perez-Saez, A. A. King, A. Rinaldo, M. Yunus, A. S. Faruque, M. Pas-  
606 cual, Climate-driven endemic cholera is modulated by human mobil-  
607 ity in a megacity, *Advances in Water Resources* 108 (2017) 367 – 376.  
608 doi:10.1016/j.advwatres.2016.11.013.
- 609 [24] M. Eisenberg, G. Kujbida, A. Tuite, D. Fisman, J. Tien, Examining  
610 rainfall and cholera dynamics in Haiti using statistical and dynamic  
611 modeling approaches, *Epidemics* 5 (4) (2013) 197–207.

- 612 [25] T. Baracchini, A. A. King, M. J. Bouma, X. Rodó, E. Bertuzzo, M. Pas-  
613 cual, Seasonality in cholera dynamics: A rainfall-driven model explains  
614 the wide range of patterns in endemic areas, *Advances in Water Re-*  
615 *sources* 108 (2016) 357–366.
- 616 [26] D. Ruiz-Moreno, M. Pascual, M. Bouma, A. Dobson, B. Cash, Cholera  
617 seasonality in madras (1901–1940): Dual role for rainfall in endemic and  
618 epidemic regions, *EcoHealth* 4 (2007) 52–62.
- 619 [27] E. Lipp, A. Huq, R. Colwell, Effects of global climate on infectious  
620 disease: The cholera model, *Clinical Microbiology Reviews* 15 (4) (2002)  
621 757–770.
- 622 [28] S. M. Faruque, M. J. Islam, Q. S. Ahmad, A. S. G. Faruque, D. A. Sack,  
623 G. B. Nair, J. J. Mekalanos, Self-limiting nature of seasonal cholera  
624 epidemics: Role of host-mediated amplification of phage, *Proceedings*  
625 *of the National Academy of Sciences USA* 102 (2005) 6119–6124.
- 626 [29] V. R. Hill, N. Cohen, A. M. Kahler, J. L. Jones, C. A. Bopp, N. Marano,  
627 C. L. Tarr, N. M. Garrett, J. Boncy, A. Henry, G. A. Gomez, M. Well-  
628 man, M. Curtis, M. M. Freeman, M. Turnsek, R. A. Benner, Jr., G. Da-  
629 hourou, D. Espey, A. DePaola, J. W. Tappero, T. Handzel, R. V. Tauxe,  
630 Toxigenic *Vibrio cholerae* O1 in water and seafood, Haiti, *Emerging In-*  
631 *fectious Diseases* 17 (11) (2011) 2147–2150.
- 632 [30] S. Rebaudet, B. Sudre, B. Faucher, R. Piarroux, Environmental deter-  
633 minants of cholera outbreaks in inland Africa: a systematic review of



- 634 main transmission foci and propagation routes, *The Journal of Infectious*  
635 *Diseases* 208 (2013) S46–S54.
- 636 [31] M. Emch, C. Feldacker, M. S. Islam, M. Ali, Seasonality of cholera  
637 from 1974 to 2005: a review of global patterns, *International Journal of*  
638 *Health Geographics* 7 (1) (2008) 31–39. doi:10.1186/1476-072X-7-31.
- 639 [32] G. Constantin de Magny, R. Murtugudde, M. R. P. Sapiano, A. Nizam,  
640 C. W. Brown, A. J. Busalacchi, M. Yunus, G. B. Nair, A. I. Gil, C. F.  
641 Lanata, J. Calkins, B. Manna, K. Rajendran, M. K. Bhattacharya,  
642 A. Huq, R. B. Sack, R. R. Colwell, Environmental signatures associ-  
643 ated with cholera epidemics, *Proceedings of the National Academy of*  
644 *Sciences USA* 105 (2008) 17676–17681.
- 645 [33] S. Rebaudet, B. Sudre, B. Faucher, R. Piarroux, Cholera in coastal  
646 africa: a systematic review of its heterogeneous environmental determi-  
647 nants, *The Journal of Infectious Diseases* 208 (2013) S98–S106.
- 648 [34] A. Jutla, A. Akanda, A. Huq, A. Faruque, R. Colwell, S. Islam, A wa-  
649 ter marker monitored by satellites to predict seasonal endemic cholera,  
650 *Remote Sensing Letters* 4 (8) (2013) 822–831.
- 651 [35] J. Gaudart, S. Rebaudet, R. Barraï, J. Boncy, B. Faucher, M. Piarroux,  
652 R. Magloire, G. Thimothe, R. Piarroux, Spatio-temporal dynamics of  
653 cholera during the first year of the epidemic in Haiti, *PLoS Neglected*  
654 *Tropical Diseases* 7 (4) (2013) e2145.
- 655 [36] R. R. Frerichs, P. S. Keim, R. Barraï, R. Piarroux, Nepalese origin

- 656 of cholera epidemic in Haiti, *Clinical Microbiology and Infection* 18 (6)  
657 (2012) E158–E163.
- 658 [37] A. Kirpich, T. Weppelmann, Y. Yang, A. Ali, J. Morris Jr., I. Longini,  
659 Cholera transmission in ouest department of haiti: Dynamic modeling  
660 and the future of the epidemic, *PLoS Neglected Tropical Diseases* 9 (10)  
661 (2015) e0004153. doi:10.1371/journal.pntd.0004153.
- 662 [38] ERCC ECHO, Haiti - Hurricane Matthew - Damage assessment and  
663 ECHO Response, Emergency Response Coordination Centre (ERCC),  
664 European Commission, Humanitarian Aid and Civil Protection (2016  
665 (accessed on 2017-14-04)).  
666 URL <http://erccportal.jrc.ec.europa.eu/getdailymap/docId/1774>
- 667 [39] A. Camacho, D. Pasetto, F. Finger, E. Bertuzzo, S. Cohuet,  
668 F. Grandesso, E. Lynch, F. Luquero, Prediction of cholera dynamics  
669 in Haiti following the passage of Hurricane Matthew, Tech. rep., Epi-  
670 centre, Paris, France (2016).
- 671 [40] D. Pasetto, F. Finger, A. Rinaldo, E. Bertuzzo, Real-time projections  
672 of cholera outbreaks through data assimilation and rainfall forecasting,  
673 *Advances in Water Resources* 108 (2017) 345–356.
- 674 [41] R. Khan, R. Anwar, S. Akanda, M. D. McDonald, A. Huq, A. Jutla,  
675 R. Colwell, Assessment of risk of cholera in haiti following hurricane  
676 matthew, *The American Journal of Tropical Medicine and Hygiene*  
677 97 (3) (2017) 896–903. doi:10.4269/ajtmh.17-0048.

- 678 [42] R. Piarroux, R. Barraix, B. Faucher, R. Haus, M. Piarroux, J. Gau-  
679 dart, R. Magloire, D. Raoult, Understanding the cholera epidemic, Haiti,  
680 *Emerging Infectious Diseases* 17 (2011) 1161–1168.
- 681 [43] E. Bertuzzo, F. Finger, L. Mari, M. Gatto, A. Rinaldo, On the  
682 probability of extinction of the Haiti cholera epidemic, *Stochastic*  
683 *Environmental Research and Risk Assessment* 30 (2016) 2043–2055.  
684 doi:10.1007/s00477-014-0906-3.
- 685 [44] P. Adams, Haiti prepares for cholera vaccination but concerns remain,  
686 *The Lancet* 379 (2012) 16.
- 687 [45] M. Periago, T. Frieden, J. Tappero, K. De Cock, B. Aasen, J. Andrus,  
688 Elimination of cholera transmission in haiti and the dominican republic,  
689 *The Lancet* 379 (2012) E12–E13.
- 690 [46] P. Adams, Cholera in Haiti takes a turn for the worse, *The Lancet*  
691 381 (9874) (2013) 1264.
- 692 [47] C. Codeço, Endemic and epidemic dynamics of cholera: the role of the  
693 aquatic reservoir, *BMC Infectious Diseases* 1 (1).
- 694 [48] A. S. Azman, K. E. Rudolph, D. A. Cummings, J. Lessler, The incuba-  
695 tion period of cholera: A systematic review, *Journal of Infection* 66 (5)  
696 (2013) 432 – 438. doi:10.1016/j.jinf.2012.11.013.
- 697 [49] C. Bretó, D. He, E. L. Ionides, A. A. King, et al., Time series analysis  
698 via mechanistic models, *The Annals of Applied Statistics* 3 (1) (2009)  
699 319–348.

- 700 [50] A. Abubakar, A. S. Azman, J. Rumunu, I. Ciglenecki, T. Helderman,  
701 H. West, J. Lessler, D. A. Sack, S. Martin, W. Perea, D. Legros, F. J.  
702 Luquero, The First Use of the Global Oral Cholera Vaccine Emergency  
703 Stockpile: Lessons from South Sudan, *PLoS Medicine* 12 (11) (2015)  
704 e1001901. doi:10.1371/journal.pmed.1001901.
- 705 [51] A. S. Azman, L. A. Parker, J. Rumunu, F. Tadesse, F. G. et al., Effec-  
706 tiveness of one dose of oral cholera vaccine in response to an outbreak: a  
707 case-cohort study, *The Lancet Global Health* 4 (11) (2016) e856 – e863.  
708 doi:10.1016/S2214-109X(16)30211-X.
- 709 [52] L. Parker, J. Rumunu, C. Jamet, Y. Kenyi, R. Laku Lino, J. F. Wamala,  
710 A. M. Mpairwe, F. J. Ciglenecki, I. Luquero, A. S. Azman, J.-C. Cabrol,  
711 Adapting to the global shortage of cholera vaccines: targeted single  
712 dose cholera vaccine in response to an outbreak in South Sudan, *Lancet*  
713 *Infectious Diseases* 2017 17 (4) (2017) e123e127. doi:10.1016/S1473-  
714 3099(16)30472-8.
- 715 [53] SSNBS, Population projections for South Sudan by County 2015 - 2020,  
716 Tech. rep., South Sudan National Bureau of Statistics (2015).
- 717 [54] CIA, South Sudan, Central Intelligence Agency. Available online., 2015.
- 718 [55] IRI/LDEO, <http://iridl.ldeo.columbia.edu/>, climate Data Li-  
719 brary. Available online; accessed on 7-April-2016 (2016).
- 720 [56] C. Sciarra, A. Rinaldo, F. Laio, D. Pasetto, Mathematical modeling of  
721 cholera epidemics in south sudan, arXiv preprint arXiv:1801.03125.

- 722 [57] I. C.-H. Fung, Cholera transmission dynamic models for public health  
723 practitioners, *Emerging Themes in Epidemiology* 11 (1).
- 724 [58] A. E. Raftery, M. A. Newton, J. M. Satagopan, P. N. Krivitsky, Esti-  
725 mating the integrated likelihood via posterior simulation using the har-  
726 monic mean identity, in: *Bayesian Statistics*, Oxford University Press,  
727 New York, 2007, pp. 1–45.
- 728 [59] M. Gatto, L. Mari, E. Bertuzzo, R. Casagrandi, L. Righetto,  
729 I. Rodriguez-Iturbe, A. Rinaldo, Generalized reproduction numbers and  
730 the prediction of patterns in waterborne disease, *Proceedings of the Na-  
731 tional Academy of Sciences USA* 48 (2012) 19703–19708.
- 732 [60] E. Bertuzzo, R. Casagrandi, M. Gatto, I. Rodriguez-Iturbe, A. Rinaldo,  
733 On spatially explicit models of cholera epidemics, *Journal of the Royal  
734 Society Interface* 7 (2010) 321–333.
- 735 [61] L. Mari, E. Bertuzzo, F. Finger, R. Casagrandi, M. Gatto, A. Ri-  
736 naldo, On the predictive ability of mechanistic models for the haitian  
737 cholera epidemic, *Journal of the Royal Society Interface* 12 (104) (2015)  
738 20140840. doi:10.1098/rsif.2014.0840.
- 739 [62] O. Akman, M. R. Corby, E. Schaefer, Examination of models for cholera:  
740 insights into model comparison methods, *Letters in Biomathematics*  
741 3 (1) (2016) 93–118. doi:10.1080/23737867.2016.1211495.
- 742 [63] M. C. Eisenberg, S. L. Robertson, J. H. Tien, Identifiability and es-  
743 timation of multiple transmission pathways in cholera and waterborne  
744 disease, *Journal of theoretical biology* 324 (2013) 84–102.

- 745 [64] N. McCreesh, M. Booth, Challenges in predicting the effects of climate  
746 change on *Schistosoma mansoni* and *Schistosoma haematobium* trans-  
747 mission potential, Trends Parasitol. 29 (2013) 548–555.
- 748 [65] Y. Lai, P. Biedermann, U. Ekpo et al., Spatial distribution of schistoso-  
749 miasis and treatment needs in sub-Saharan Africa: A systematic review  
750 and geostatistical analysis, Lancet Infect. Dis. 15 (2015) 927–940.
- 751 [66] J. Perez-Saez, T. Mande, N. Ceperley, E. Bertuzzo, L. Mari, M. Gatto,  
752 A. Rinaldo, Hydrology and density feedbacks control the ecology of  
753 the intermediate hosts of schistosomiasis across habitats in seasonal cli-  
754 mates, Proc. Natl. Acad. Sci. USA 113 (2016) 6427–6432.
- 755 [67] E. Bertuzzo, L. Mari, L. Righetto, M. Gatto, R. C. and I. Rodriguez-  
756 Iturbe, A. Rinaldo, Hydroclimatology of dual-peak annual cholera inci-  
757 dence: Insights from a spatially explicit model, Geophysical Research  
758 Letters 39 (2012) L05403.
- 759 [68] E. Bertuzzo, L. Mari, L. Righetto, M. Gatto, R. Casagrandi,  
760 I. Rodriguez-Iturbe, A. Rinaldo, Prediction of the spatial evolution and  
761 effects of control measures for the unfolding Haiti cholera outbreak, Geo-  
762 phys. Res. Lett. 38 (2011) L06403.
- 763 [69] N. McCreesh, G. Nikulin, M. Booth, Predicting the effects of climate  
764 change on *Schistosoma mansoni* transmission in eastern Africa, Para-  
765 site. Vect. 8 (2015) 4.