

1 A game-theoretic model of Monkeypox to 2 assess vaccination strategies

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16 ABSTRACT

17 Monkeypox (MPX) is a zoonotic disease similar to smallpox. Its fatality rate is about 11% and it is endemic
18 to the Central and West African countries. In this paper, we analyze a compartmental model of MPX
19 dynamics. Our goal is to see whether MPX can be controlled and eradicated by voluntary vaccinations.
20 We show that there are three equilibria - disease free, fully endemic and previously neglected semi-
21 endemic (with disease existing only among humans). The existence of semi-endemic equilibrium has
22 severe implications should the MPX virus mutate to increased viral fitness in humans. We find that MPX
23 is controllable and can be eradicated in a semi-endemic equilibrium by vaccination. However, in a fully
24 endemic equilibrium, MPX cannot be eradicated by vaccination alone.

25 1 INTRODUCTION

26 Monkeypox (MPX) is a zoonotic disease that has the potential to develop into one of the most threatening
27 human *Orthopoxvirus* infections since the eradication of smallpox (Durski et al., 2018). The causative
28 agent of MPX is monkeypox virus (MPXV), found in the same genus as the variola virus (smallpox),
29 vaccinia virus, and cowpox virus (Shchelkunov et al., 2006; Sklenovská and Van Ranst, 2018). Common
30 symptoms of MPX, though relatively milder than smallpox, include fever, severe headaches, skin lesions,
31 and myalgia (CDC, 2003). Prevention of the disease has remained a challenge for poverty-stricken rural
32 areas with poor infrastructure that lack necessary sanitary supervision (Sklenovská and Van Ranst, 2018).

33 MPX is endemic to Central Africa and West Africa (Weinstein et al., 2005; Yinka-Ogunleye et al.,
34 2018). West African and Central African strains of MPXV exist, the latter of which is more virulent and
35 symptomatically severe (Likos et al., 2005; Mwamba et al., 2014). In the Democratic Republic of the
36 Congo (DRC) the mortality rate of the Central African strain is 11% (Ježek et al., 1987). Since the first
37 case of human infection in 1970, there have been numerous outbreaks in the DRC (Eteng et al., 2018).
38 Annually, the DRC reports over 2,000 cases of suspected infections (Mwamba et al., 2014). This estimate
39 may be modest, as MPX is often misdiagnosed as chickenpox or other diseases that cause rashes (Ježek
40 et al., 1988). Additionally, modern and robust surveillance of MPX is neglected as a consequence of
41 limited funds and resources (Rimoin et al., 2010), and countries other than the DRC are not required
42 to report all cases of MPX (Durski et al., 2018). Thus, the disease may be more severe than previously
43 estimated.

44 In 2003, 47 cases of MPX were reported across 5 states in the U.S., originating from a shipment of
45 animals from Ghana imported to Texas (CDC, 2003). In 2018, 3 cases of MPX were reported in the

46 United Kingdom, making it the first time since the 2003 United States outbreak that the disease had
47 reached a country outside of Africa (Eteng et al., 2018).

48 The clinical presentation of MPX can be found in Di Giulio and Eckburg (2004). The incubation
49 period for the virus ranges from 5 to 21 days. MPX infection is split into 2 distinct phases: the invasion
50 period and the skin eruption period. The invasion period starts between 0-5 days and is characterized by
51 fever, lymphadenopathy, intense asthenia, severe headaches, and myalgia. The skin eruption period occurs
52 1-3 days after the appearance of a fever or lymphadenopathy, and it is characterized by rash formation,
53 which often begins on the face and spreads to the rest of the body. The rash first appears as maculopapules
54 (lesions with flat bases) and progresses to fluid filled blisters called vesicles. The vesicles then burst,
55 forming pustules, and a crust forms over the affected area within 10 days. The number of lesions formed
56 can vary from a few to thousands across the body, with children reportedly experiencing more severe
57 symptoms than adults.

58 The predominant mode of MPX transmission is through human-animal interaction. Direct contact
59 with an infected animal's blood, bodily fluids, or lesions can lead to infection. Documented cases of MPX
60 in Central and West Africa show that transmission can occur via the handling of wild animals (Reynolds
61 et al., 2017). Cultural influences, such as consumption of "bush meat," can be a potential source of
62 transmission. Additionally, direct contact with an infected person's bodily fluids and skin lesions can lead
63 to the transmission of the disease (McCollum and Damon, 2013).

64 Despite MPX's high case fatality rate (Ježek et al., 1987) there are no known cures (Eteng et al.,
65 2018). Until recently, there were no disease-specific preventative measures such as vaccines, though
66 existing smallpox vaccines have historically been around 85% successful (Eteng et al., 2018). However,
67 administration of the smallpox vaccine has ceased since the disease's eradication in 1980, resulting in
68 lowered immunity against *Orthopoxviruses* in general. This has led to a supposed increase in population
69 susceptibility to MPXV (Sklenovská and Van Ranst, 2018). In 2019, the vaccine termed Jynneos® was
70 approved by the US FDA for protection against VARV and MPXV (Meyer et al., 2020).

71 It is possible that this lack of preventative measures is partially explained by a matching lack of
72 literature on the potential dangers of inter-human transmission of MPX (Rimoin et al., 2010; Mwamba
73 et al., 2014; Doshi et al., 2018; Sklenovská and Van Ranst, 2018). The urgency of better understanding of
74 MPX is exacerbated considering vaccination cessation and immunocompromised populations in Central
75 Africa, so a need for comprehensive preventative strategies is apparent (WHO, 2017). The factors such as
76 (1) a lack of an effective vaccination strategy from fixed bases, (2) a shortfall in the vaccine supply and (3)
77 logistical and security problems associated with the distance from the health centers, all contribute to the
78 challenges of vaccinating the whole population in Central and Western Africa (Herp et al., 2003).

79 The identity of MPXV reservoir host(s) remains unknown (Di Giulio and Eckburg, 2004; Falendysz
80 et al., 2017). The seroprevalence of MPXV was found highest in a population of moribund rope squirrel
81 (*Funisciurus anerythrus*) in Zaire (now DRC), and it was also found in sun squirrels (*Heliosciurus*
82 *rufobrachium*) and non-human primates in DRC (Khodakevich et al., 1987, 1988) as well as in Gambian
83 pouched rats (*Cricetomys gambianus*) (Doty et al., 2017; Doshi et al., 2019). In West Africa, African
84 dormice (*Graphiurus sp.*), and ground squirrels (*Xerus sp.*) were identified as additional hosts (Reynolds
85 et al., 2010). The majority of reported human cases originate from an interaction with an infected
86 animal (Arita et al., 1985). The transmission of MPX among animals can be affected by environmental
87 conditions (Brown and Leggat, 2016). Deforestation and flooding could potentially increase or decrease
88 the MPX reservoirs, depending on how the animal population is affected by these conditions (Brown and
89 Leggat, 2016). Long-distance transportation of potential MPX carriers may result in the expansion of the
90 geographical range of the MPX reservoir, as exemplified by the 2003 U.S. outbreak.

91 Currently, MPXV likely needs the animal reservoir as the human-to-human transmission chains of
92 MPX are relatively short; the maximum number of generations reported in literature is seven (Learned
93 et al., 2005). Nevertheless, as demonstrated by the case of H1N1 influenza (swine flu), some virus
94 mutations can increase viral fitness in humans (Elderfield et al., 2014). We note that poxviruses have
95 linear, double-stranded DNA genomes that vary from 130 to 230 kbp (Moss, 2013) and as such are
96 evolving much slower than H1N1. Nevertheless, they can still adapt rapidly (Elde et al., 2012) and genetic
97 engineering and modern molecular biology already turned a mousepox virus into unusually lethal strain
98 (Jackson et al., 2001; Di Giulio and Eckburg, 2004).

99 Epidemiologic compartmental models have been used to better understand the potential implications
100 of disease transmission and infection (Blackwood and Childs, 2018; Bidari and Goldwyn, 2019). For

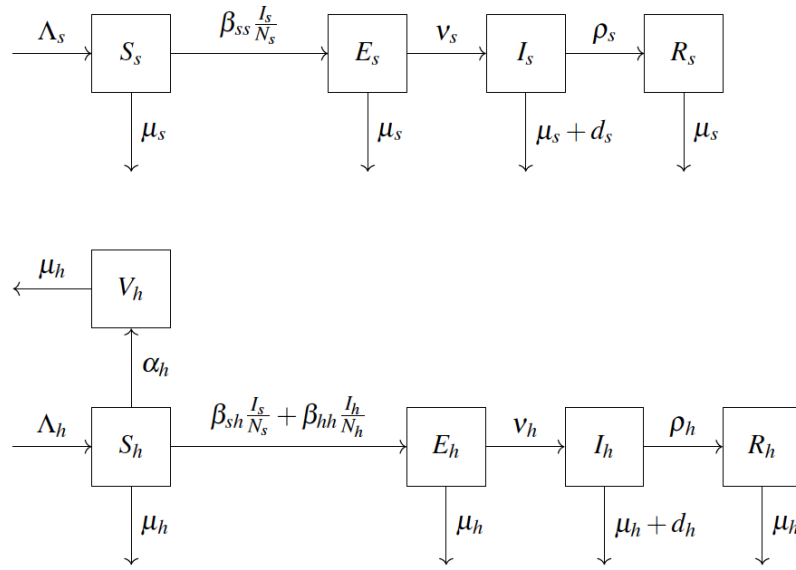


Figure 1. Scheme of mathematical model for humans and squirrels, adapted from Usman and Adamu (2017).

101 MPX, the framework for a mathematical model has been tentatively set, but existing iterations have had
 102 shortcomings, failing to address some of the aforementioned aspects of the disease in their entirety. Bhunu
 103 and Mushayabasa (2011) introduced a basic SIR vector-borne compartmental model between humans and
 104 primates, yet deem an endemic state solely in humans as trivial. Usman and Adamu (2017) build upon
 105 this framework by introducing an SVEIR compartmental model to account for the disease’s incubation
 106 period and potential vaccine.

107 Game theoretical models attempt to study complex scenarios in which self-interested individuals will
 108 take an action based on the decisions of the rest of the population (Bauch and Earn, 2004). The model is a
 109 predictive tool in populations for extracting an optimum decision-making strategy (Chang et al., 2020).
 110 Game theory has been applied to protection strategies to control diseases such as smallpox (Bauch et al.,
 111 2003), toxoplasmosis (Sykes and Rychtář, 2015), cholera (Kobe et al., 2018), measles (Shim et al., 2012),
 112 rubella (Shim et al., 2009), influenza (Galvani et al., 2007), African sleeping sickness (Crawford et al.,
 113 2015), malaria (Orwa et al., 2018; Broom et al., 2016), zika (Padmanabhan et al., 2017; Bañuelos et al.,
 114 2019), polio (Cheng et al., 2020), Ebola (Brettin et al., 2018), chikungunya (Klein et al., 2019), meningitis
 115 (Martinez et al., 2019), typhoid (Acosta-Alonzo et al., 2020), Hepatitis C (Scheckelhoff et al., 2019) or
 116 Hepatitis B (Chouhan et al., 2019) among others. In this paper, we apply a similar approach to MPX
 117 to investigate a scenario in which individuals have the option of vaccinating to reduce the chance of
 118 contracting the virus. We further evaluate vaccination strategies on an individual and population-wide
 119 level by discussing the vaccination rates required to achieve herd immunity and Nash equilibrium.

120 In the present paper, we build on the work of Usman and Adamu (2017), see also Lauko et al. (2018)
 121 for a simplified SIR version of the model. The mathematical model of the MPX dynamics is shown in
 122 Section 2. In Section 3.1, we provide closed-form formulas for equilibrium states of MPX dynamics; the
 123 formulas provided in Usman and Adamu (2017) do not allow for direct calculations of the equilibria. We
 124 also show the existence of a “semi-endemic” equilibrium. This was not previously discussed in Usman
 125 and Adamu (2017), although it appears in Lauko et al. (2018). In Section 3.2, we apply a game-theoretic
 126 approach to evaluate individual and population-wide vaccination strategies on the basis of cost and
 127 probabilistic disease acquisition. In Section 3.3 we perform sensitivity analysis. We conclude the paper
 128 by a discussion in Section 4.

129 2 MATHEMATICAL MODEL

130 We adopt the compartmental epidemiological model introduced in Usman and Adamu (2017) and shown
 131 in Figure 1. We consider squirrels to be the primary reservoir hosts. The population is divided into

Table 1. Model parameters. The human MPX related death rate was taken as a solution to $d_h/(d_h + \rho_h) = 0.1$ where 10% is the MPX fatality (Ježek et al., 1987). Similarly, squirrel MPX related death rate was taken as a solution to $d_s/(d_s + \rho_s) = 0.6$ where 60% is an estimate for the MPX fatality found between 50-75% (Falendysz et al., 2017). We estimated the effective squirrel-to-squirrel transmission rate as $\beta_{ss} = 40$; this yields about 24% of seropositive squirrels in the population, a number that agrees with estimates from Khodakevich et al. (1988). The effective transmission rates between humans was estimated as $\beta_{hh} = 32.85$ as follows. Arita et al. (1985) provide transmission risk $p = 0.15$ amongst household contacts and $p = 0.03$ amongst other contacts. We assumed human-to-human contact rate $\gamma = 365$ (i.e. once a day) and obtained $\beta_{hh} = 365 \cdot \frac{0.15+0.03}{2} = 32.85$. The effective squirrel-to-human transmission rate was estimated to be $\beta_{sh} = 0.05$ as this yields about 1% of seropositive humans (Khodakevich et al., 1988). The actual cost of vaccine is \$4.85 (Lambert de Rouvroit and Heegaard, 2016). While the vaccine is provided for free, there are many other direct and indirect costs associated with vaccination (need to travel to the health center, associated security risk, loss of income etc., see for example Herp et al. (2003)) and we estimated the cost of vaccination to be \$4. We note that the previously approved smallpox vaccines such as ACAM2000 could cause severe side effects (Wollenberg and Engler, 2004; Nalca and Zumbrun, 2010). It is not clear if the new vaccine, JYNNEOS, is more effective to protect against MPXV infections in humans than ACAM2000 and what the potential side effects are.

Symbol	Meaning	Value	Source
Λ_h	Human birth rate	0.0328	CIA (2019)
Λ_s	Squirrel birth rate	2	Hayssen (2008)
μ_h	Human natural death rate	1/60	World Bank (2019)
μ_s	Squirrel natural death rate	0.5	Khodakevich et al. (1988)
d_h	Human MPX related death rate	3.12	Ježek et al. (1987)
d_s	Squirrel MPX related death rate	17.5	Falendysz et al. (2017)
ρ_h	Human recovery rate	28.08	Di Giulio and Eckburg (2004)
ρ_s	Squirrel recovery rate	12	Falendysz et al. (2017)
v_h	Human infection rate	30.42	Di Giulio and Eckburg (2004)
v_s	Squirrel infection rate	120	Falendysz et al. (2017)
α_h	Vaccination rate	variable	
β_{ss}	Squirrel-to-squirrel transmission rate	40	Assumed based on Khodakevich et al. (1988)
β_{sh}	Squirrel-to-human transmission rate	0.05	Assumed based on Khodakevich et al. (1988)
β_{hh}	Human-to-human transmission rate	32.85	Arita et al. (1985)
C_V	Cost of vaccination	\$4	Herp et al. (2003)
C_{MPX}	Cost of MPX infection	\$100	Adam et al. (2003)

132 squirrels and humans, denoted by s and h subscripts, respectively. Individuals are born as Susceptible
 133 (S) at rate Λ . Susceptible humans vaccinate (move to V_h) at rate α_h . Vaccinated humans are assumed to
 134 never contract the disease in the remainder of their lifetime. Susceptible squirrels become Exposed (E_s)
 135 by coming into contact with infected squirrels with effective transmission rate β_{ss} . Susceptible humans
 136 become exposed by coming into contact with either infected squirrels (with effective transmission rate β_{sh})
 137 or infected humans (with effective transmission rate β_{hh}). After an incubation period v^{-1} , the exposed
 138 individuals become Infected (I). Infected individuals are infectious. They Recover (R) at rate ρ . Any
 139 individual may die due to natural causes at rate μ . Infected individuals can also die from the disease at
 140 rate d . The notation is summarized in Table 1. The model yields the following differential equations, see
 141 for example Blackwood and Childs (2018).

$$\frac{dS_s}{dt} = \Lambda_s - \left(\mu_s + \beta_{ss} \frac{I_s}{N_s} \right) S_s \quad (1)$$

$$\frac{dE_s}{dt} = \beta_{ss} \frac{I_s}{N_s} S_s - (\mu_s + v_s) E_s \quad (2)$$

$$\frac{dI_s}{dt} = v_s E_s - (\mu_s + d_s + \rho_s) I_s \quad (3)$$

$$\frac{dR_s}{dt} = \rho_s I_s - \mu_s R_s \quad (4)$$

$$\frac{dS_h}{dt} = \Lambda_h - \left(\mu_h + \left(\beta_{sh} \frac{I_s}{N_s} + \beta_{hh} \frac{I_h}{N_h} \right) + \alpha_h \right) S_h \quad (5)$$

$$\frac{dV_h}{dt} = \alpha_h S_h - \mu_h V_h \quad (6)$$

$$\frac{dE_h}{dt} = \left(\beta_{sh} \frac{I_s}{N_s} + \beta_{hh} \frac{I_h}{N_h} \right) S_h - (\mu_h + v_h) E_h \quad (7)$$

$$\frac{dI_h}{dt} = v_h E_h - (\mu_h + d_h + \rho_h) I_h \quad (8)$$

$$\frac{dR_h}{dt} = \rho_h I_h - \mu_h R_h \quad (9)$$

142 3 RESULTS

143 3.1 Equilibrium states of the MPX dynamics

144 The basic reproduction numbers were derived by Usman and Adamu (2017) and are given by

$$R_{0ss} = \beta_{ss} \cdot \frac{1}{\mu_s + d_s + \rho_s} \cdot \frac{v_s}{\mu_s + v_s} \quad (10)$$

$$R_{0hh} = \beta_{hh} \cdot \frac{\mu_h}{\alpha_h + \mu_h} \cdot \frac{1}{\mu_h + d_h + \rho_h} \cdot \frac{v_h}{\mu_h + v_h}. \quad (11)$$

145 As shown in Appendix A1, R_{0ss} corresponds to a number of secondary squirrel infections caused by a
 146 single infected squirrel in an otherwise healthy population. The meaning of R_{0hh} is similar.

147 There are three qualitatively distinct equilibria of the dynamics (1) - (9). First, ε^0 is the disease free
 148 equilibrium. It occurs when $R_{0ss} < 1$ and $R_{0hh} < 1$. Second, ε^* is the fully endemic equilibrium with
 149 disease occurring amongst humans as well as squirrels. The equilibrium is stable when $R_{0ss} > 1$. Finally,
 150 ε^\dagger is a semi-endemic equilibrium with disease prevalent only amongst human population. It is stable
 151 when $R_{0ss} < 1$ and $R_{0hh} > 1$.

152 The closed form formulas are given in the Table 2. Step-by-step derivation can be found in Ap-
 153 pendix A1.

154 3.2 Herd immunity and Nash equilibrium vaccination rates

155 The average cost of not vaccinating when the population vaccination rate is α_h is denoted $C_{notV}(\alpha_h)$ and
 156 it is given as a product of the cost of the MPX infection (C_{MPX}) and the probability of moving from the S_h

Table 2. Different equilibria of the MPX dynamics. The formulas for N_h^* and N_h^\dagger are too long for the table and are given in Appendix A1.

	Disease-free (ε^0)	Fully Endemic (ε^*)	Semi-endemic (ε^\dagger)
N_s	$\frac{\Lambda_s}{\mu_s}$	$\frac{\Lambda_s \cdot \left(\frac{\mu_s + d_s + \rho_s}{v_s} + 1 + \frac{\rho_s}{\mu_s} \right)}{\mu_s \cdot \left(\frac{\mu_s + d_s + \rho_s}{v_s} \right) + d_s \cdot \left(1 - \frac{1}{R_{0ss}} \right) + \mu_s + \rho_s}$	$\frac{\Lambda_s}{\mu_s}$
S_s	N_s^0	$N_s^* \cdot \frac{1}{R_{0ss}}$	N_s^\dagger
E_s	0	$\left(\frac{\mu_s + d_s + \rho_s}{v_s} \right) \cdot I_s^*$	0
I_s	0	$\frac{\Lambda_s - \mu_s N_s^*}{d_s}$	0
R_s	0	$\frac{\rho_s}{\mu_s} \cdot I_s^*$	0
N_h	$\frac{\Lambda_h}{\mu_h}$	(45)	(59)
S_h	$\frac{\mu_h}{\mu_h + \alpha_h} \cdot N_h^0$	$\frac{(\mu_h + v_h) \left(\frac{\mu_h + d_h + \rho_h}{v_h} \right) \cdot I_h^*}{\beta_{sh} \left(\frac{I_s^*}{N_s^*} \right) + \beta_{hh} \left(\frac{I_h^*}{N_h^*} \right)}$	$\frac{(\mu_h + d_h + \rho_h)(\mu_h + v_h)}{v_h \beta_{hh}} \cdot N_h^\dagger$
V_h	$\frac{\alpha_h}{\mu_h + \alpha_h} \cdot N_h^0$	$\frac{\alpha_h}{\mu_h} \cdot S_h^*$	$\left(\frac{\alpha_h}{\mu_h} \right) \cdot S_h^\dagger$
E_h	0	$\left(\frac{\mu_h + d_h + \rho_h}{v_h} \right) \cdot I_h^*$	$\left(\frac{\mu_h + d_h + \rho_h}{v_h} \right) \cdot I_h^\dagger$
I_h	0	$\frac{\Lambda_h - \mu_h N_h^*}{d_h}$	$\frac{\Lambda_h - \mu_h N_h^\dagger}{d_h}$
R_h	0	$\frac{\rho_h}{\mu_h} \cdot I_h^*$	$\frac{\rho_h}{\mu_h} \cdot I_h^\dagger$

157 compartment to the I_h compartment, i.e.

$$C_{notV}(\alpha_h) = C_{MPX} \cdot \left(\frac{\left(\beta_{sh} \frac{I_s}{N_s} + \beta_{hh} \frac{I_h}{N_h} \right)}{\left(\beta_{sh} \frac{I_s}{N_s} + \beta_{hh} \frac{I_h}{N_h} \right) + \mu_h} \right) \cdot \left(\frac{v_h}{v_h + \mu_h} \right). \quad (12)$$

158 As the vaccination rate α_h increases, R_{0hh} decreases by (11). Furthermore, the fraction $\frac{I_h}{N_h}$ at the
159 appropriate equilibrium also decreases. Consequently, the cost of not vaccinating decreases. In the
160 semi-endemic equilibrium, the cost eventually becomes 0 when the vaccination rate reaches

$$\alpha_{HI} = \max \left\{ 0, \frac{v_h \beta_{hh} \mu_h}{(\mu_h + d_h + \rho_h)(\mu_h + v_h)} - \mu_h \right\}. \quad (13)$$

161 At that point the herd immunity is achieved and the disease is eradicated from the population. In the fully
162 endemic equilibrium, there is always a reservoir of MPX in the squirrel population. This reservoir causes
163 an influx of MPX infections amongst humans. Therefore the disease can never be fully eradicated and the
164 cost of not vaccinating will never reach 0, see Figure 2.

165 When the vaccination rate is such that $C_{notV}(\alpha_h) = C_V$, the vaccination rate is at Nash equilibrium,
166 α_{NE} . When $\alpha_h < \alpha_{NE}$, it is beneficial for the individual to vaccinate; when $\alpha_h > \alpha_{NE}$, it is beneficial for
167 the individual not to vaccinate.

168 Figure 3 shows a scenario where $\beta_{hh} = 60$. While this value is unrealistically high, we investigated
169 this hypothetical scenario to see what would happen if MPXV mutates as was the case of H1N1 influenza
170 (swine flu) or is genetically engineered as was the case of mousepox (Jackson et al., 2001; Di Giulio and
171 Eckburg, 2004). When β_{hh} is large enough, specifically when

$$\beta_{hh} > \frac{\alpha_h + \mu_h}{\alpha_h} \cdot (\mu_h + d_h + \rho_h) \cdot \frac{\mu_h + v_h}{v_h} \quad (14)$$

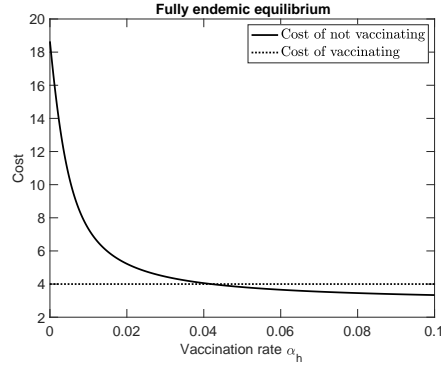


Figure 2. Cost versus vaccination rate. The vaccination rate (α_h) is varied while all other parameter values are as specified in Table 1.

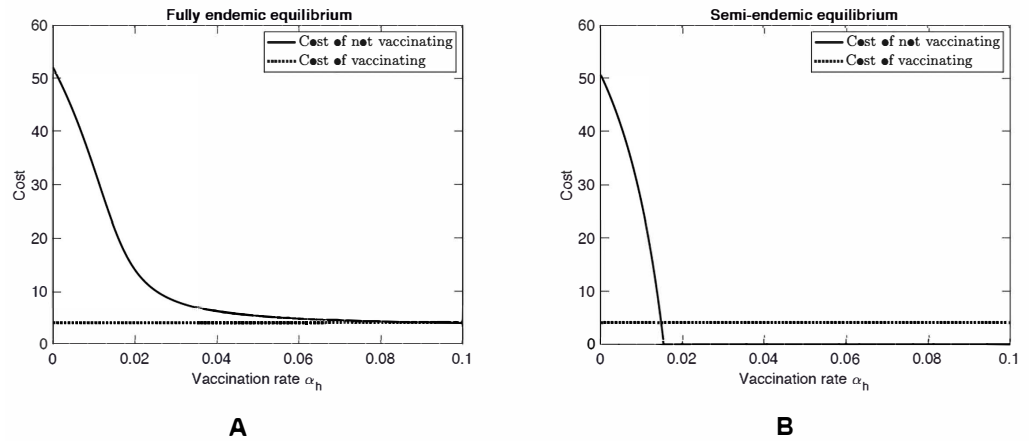


Figure 3. Costs vs. Vaccination rate when the effective human-to-human transmission rate is high, $\beta_{hh} = 60$. **A** the fully endemic state. **B** the semi-endemic state ($\beta_{ss} = 30 < (\mu_s + d_s + \rho_s) \cdot \frac{\mu_s + v_s}{v_s}$). The same scenario occurs when $\beta_{sh} = 0$ and β_{ss} is arbitrary. In both figures, the vaccination rate (α_h) is varied while all other parameter values are as specified in Table 1.

172 MPX no longer needs squirrels to persist in the human population. In particular, it can become endemic
 173 even in countries without natural squirrel population (i.e. even when $\beta_{sh} = 0$). At the same time, in the
 174 semi-endemic equilibrium, the disease can be controlled through vaccination. Note that there is almost no
 175 difference between the Nash equilibrium rate α_{NE} (a solution to $C_{notV}(\alpha_h) = C_V$) and the rate α_{HI} needed
 176 for the herd immunity (a solution to $C_{notV}(\alpha_h) = 0$).

177 3.3 Sensitivity analysis

178 As shown in Figure 2, as α_h increases, $C_{notV}(\alpha_h)$ approaches an asymptote. Consequently, the value of
 179 α_{NE} , a solution to $C_{notV}(\alpha_h) = C_V$ can be very sensitive to C_V when $C_V \approx 3$. Any small decrease of C_V
 180 can cause a significant increase of α_{NE} . The same sensitivity is demonstrated in Figure 3.

181 Figure 4 shows the sensitivity analysis and how α_{NE} depends on variation of different parameters.
 182 We can see the high sensitivity of α_{NE} on the squirrel-to-human transmission rate, β_{sh} , and on the cost
 183 of vaccination, C_V , for low values of C_V . Moreover, the figure demonstrates that α_{NE} can be quite
 184 sensitive on the effective transmission rate amongst squirrels, β_{ss} and the squirrels recovery rate, ρ_s . For
 185 $\beta_{ss} < (\mu_s + d_s + \rho_s) \cdot \frac{\mu_s + v_s}{v_s}$, there is a semi-endemic equilibrium and $\alpha_{NE} = 0$. However, as β_{ss} increases
 186 above that threshold value, α_{NE} rapidly increases and, when $\beta_{ss} > 45$, there is no Nash equilibrium
 187 vaccination rate. Similarly, when ρ_s is large enough to have a semi-endemic equilibrium, the optimal
 188 vaccination rate is 0. However, for small ρ_s , there is no Nash equilibrium and the change is relatively
 189 abrupt as in the case of β_{ss} .

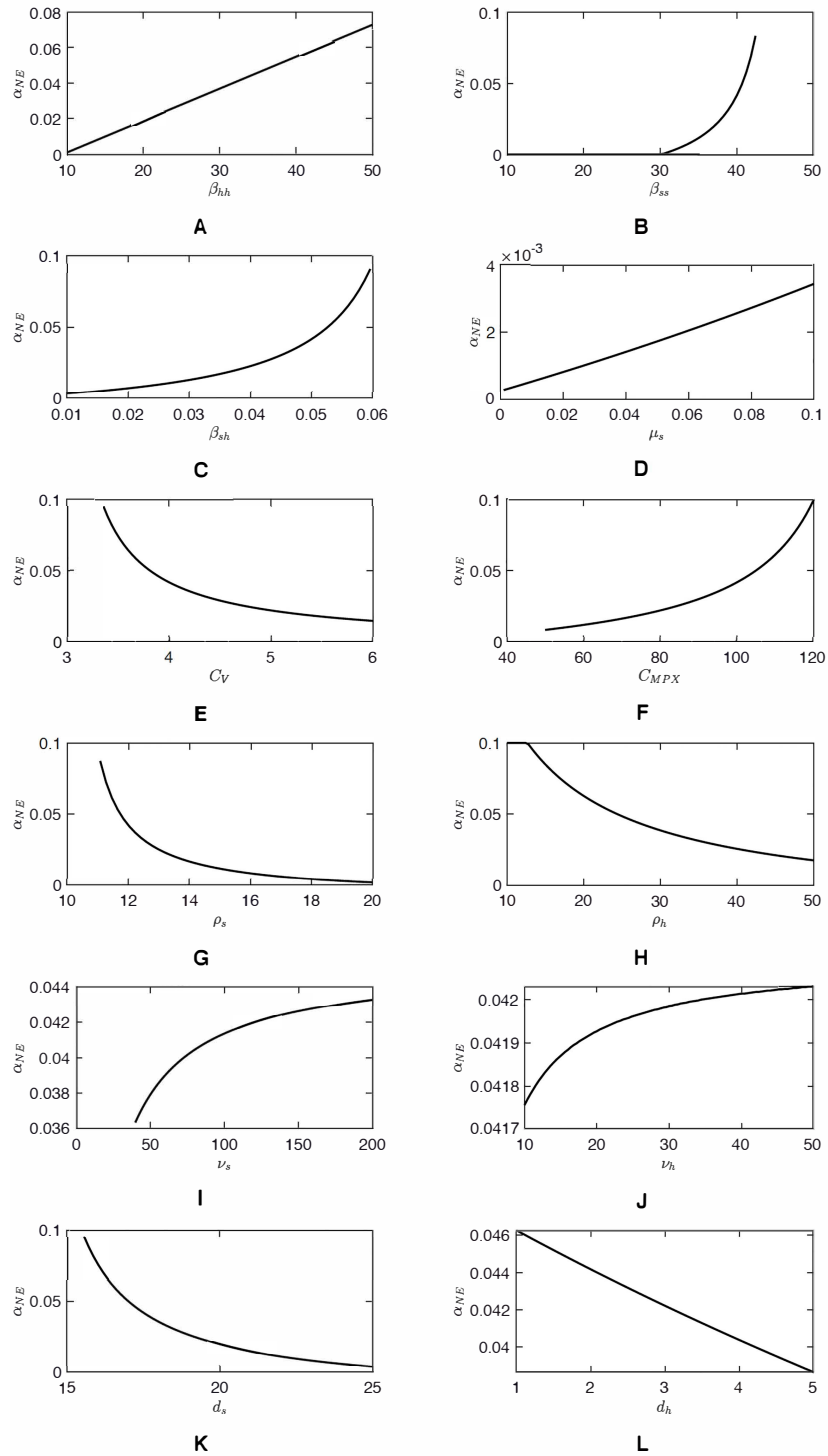


Figure 4. Dependence of α_{NE} on different parameter values. Unless varied, the parameter values are as specified in Table 1. For those parameters, $\alpha_{NE} = 0.0419$ and the sensitivity index, $SI_x = \left(\frac{x}{\alpha_{NE}}\right) \left(\frac{\partial \alpha_{NE}}{\partial x}\right)$ of α_{NE} with respect to parameters are as follows: $SI_{\beta_{hh}} = 1.415$, $SI_{\beta_{sh}} = 3.472$, $SI_{\beta_{ss}} = 10.911$, $SI_{\mu_s} = 0.932$, $SI_{\rho_s} = -7.120$, $SI_{\rho_h} = -1.258$, $SI_{\nu_s} = 0.077$, $SI_{\nu_h} = 0.004$, $SI_{d_s} = -5.917$, $SI_{d_h} = -0.141$, $SI_{C_V} = -3.518$, $SI_{C_{MPX}} = 3.603$.

3.4 Model validation

For the parameters in Table 1, the proportion of seropositive squirrels, given as $\frac{I_s+R_s}{N_s}$, is 24.44% which generally agrees with Khodakevich et al. (1988). Also, the proportion of seropositive people, $\frac{I_h+R_h}{N_h}$, is 1.06%, again agreeing with Khodakevich et al. (1988).

4 DISCUSSION

The phylogenetic relatedness between MPXV and variola virus grants the smallpox vaccine an 85% effectiveness in preventing MPX (Reynolds and Damon, 2012). Poxviruses from the *Orthopoxvirus* genus have cross-reactive antibodies, meaning that vaccinated individuals would have a much lower risk of infection and mortality compared to unvaccinated individuals (Louten, 2016). The imperfect prevention rate of the vaccine was omitted in the design of the mathematical model for the sake of simplicity. As noted in Wu et al. (2011), imperfect protection aggravates the dilemma of voluntary protective actions as lower vaccine effectiveness can lead to better vaccine coverage and smaller free-riding effects; however, the impact of the epidemic can be harder to mitigate.

It is of interest to identify and evaluate possible preventative measures in addition to vaccination that would have a measurable effect on the transmission of MPX. For instance, decreasing the animal-to-human contact and launching an education campaign about dangers of eating raw meat which seems to be the main culprit behind squirrel-to-human transmission Khodakevich et al. (1988) would significantly decrease the animal-to-human transmission rate. It could still come at a considerable individual cost (such as decrease of meat supply) but it would not require a complex or well-developed healthcare infrastructure needed for the vaccination, thus providing the general population with an easily accessible preventative measure. The mathematical model for such a measure would become more complex. The main idea would follow the spirit of Kobe et al. (2018) that investigated a situation for cholera prevention where individuals could either vaccinate or avoid drinking potentially contaminated water.

The reservoir host for monkeypox remains unclear (Di Giulio and Eckburg, 2004; Falendysz et al., 2017). We focused on a moribund rope squirrel, *Funisciurus anerythrus*, but we note that the disease has also been confirmed in other animals (Arita et al., 1985; Khodakevich et al., 1988; Reynolds et al., 2010). As noted in Falendysz et al. (2017), in a recent outbreak of MPXV in DRC, no association was found between contact with rope squirrels and human infection (Nolen et al., 2015). Additionally, a recent survey of 34 villages in the Tshuapa region of DRC did not detect contact with a rope squirrel carcass in the previous 30 days, although they reported contacts with red-legged sun squirrel, *Heliosciurus rufobrachium*, (Monroe et al., 2015) which was identified in Khodakevich et al. (1988) as another frequent host of monkeypox.

5 CONCLUSIONS

We modeled MPX dynamics using the compartmental model of Usman and Adamu (2017). As one of our major contributions, we provided closed form formulas of the equilibrium states of the dynamics. Moreover, we also showed a potential existence of the semi-endemic equilibrium, in which there is no infection in the squirrel population and the disease still persists in the human population. Currently, MPX does not seem to have the viral fitness to become endemic solely through human transmission. Yet, simple mutations in viral proteins could still occur and increase successive inter-human cases as seen in the H1N1 virus outbreak (Le et al., 2009). Should this mutation occur, a careful understanding of the semi-endemic equilibrium will be needed.

In addition, we applied a game-theoretical approach to assess vaccination decision-making developed by Bauch and Earn (2004). Individuals in any population susceptible to MPX have the choice to vaccinate against the disease or risk the possibility of contracting the disease. Naturally, it is in the individual's best interest to choose the option with the smaller expected cost. The model quantifies the costs and benefits of getting smallpox vaccine. We found that the optimal vaccination rate is about 0.04, i.e. individuals should vaccinate about once every 25 years.

We must note that the parameter values we used are only estimates based on available literature. In reality, the parameters may be quite different, in large part because the reservoir hosts are different as discussed above. The performed sensitivity analysis allows us to gain insight into how our results depend on the specific parameter values. We observed that the optimal vaccination rate, α_{NE} , is about 10 times more sensitive to parameters related to animal hosts than to a corresponding parameter related to humans.

242 It is therefore important to establish more accurate parameters. Consequently, greater efforts are needed
243 to track the true prevalence and recurrent cases of MPX in all populations rather than relying on suspected
244 cases.

245 Though not perfect in practice, mathematical modeling of diseases remains a powerful tool that grants
246 a more profound understanding how MPX operates under certain conditions. The scope of epidemiological
247 modeling and game-theoretic cost analysis is wide. As cases of MPX become increasingly reported
248 among humans (Antwerpen et al., 2019), we hope that the models may serve as a predictive tool to better
249 study the spread of MPX.

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253 REFERENCES

- 254 Acosta-Alonzo, C., Erovenko, I., Lancaster, A., Oh, H., Rychtář, J., and Taylor, D. (2020). High endemic
255 levels of typhoid fever in rural areas of Ghana may stem from optimal voluntary vaccination behavior.
256 Preprint.
- 257 Adam, T., Evans, D. B., and Murray, C. J. (2003). Econometric estimation of country-specific hospital
258 costs. *Cost Effectiveness and Resource Allocation*, 1(1):3.
- 259 Antwerpen, M. H., Georgi, E., Nikolic, A., Zoeller, G., Wohlsein, P., Baumgärtner, W., Peyrefitte, C.,
260 Charrel, R., and Meyer, H. (2019). Use of next generation sequencing to study two cowpox virus
261 outbreaks. *PeerJ*, 7:e6561.
- 262 Arita, I., Ježek, Z., Khodakevich, L., and Ruti, K. (1985). Human monkeypox: a newly emerged
263 orthopoxvirus zoonosis in the tropical rain forests of Africa. *The American Journal of Tropical*
264 *Medicine and Hygiene*, 34(4):781–789.
- 265 Bañuelos, S., Martinez, M., Mitchell, C., and Prieto-Langarica, A. (2019). Using mathematical modelling
266 to investigate the effect of the sexual behaviour of asymptomatic individuals and vector control measures
267 on zika. *Letters in Biomathematics*, 6(1):1–19.
- 268 Bauch, C. T. and Earn, D. J. (2004). Vaccination and the theory of games. *Proceedings of the National*
269 *Academy of Sciences*, 101(36):13391–13394.
- 270 Bauch, C. T., Galvani, A. P., and Earn, D. J. (2003). Group interest versus self-interest in smallpox
271 vaccination policy. *Proceedings of the National Academy of Sciences*, 100(18):10564–10567.
- 272 Bhunu, C. and Mushayabasa, S. (2011). Modelling the transmission dynamics of pox-like infections.
273 *IAENG International Journal of Applied Mathematics*, 41(2):141–149.
- 274 Bidari, S. and Goldwyn, E. E. (2019). Stochastic models of influenza outbreaks on a college campus.
275 *Letters in Biomathematics*, pages 1–14.
- 276 Blackwood, J. C. and Childs, L. M. (2018). An introduction to compartmental modeling for the budding
277 infectious disease modeler. *Letters in Biomathematics*, 5(1):195–221.
- 278 Brettin, A., Rossi-Goldthorpe, R., Weishaar, K., and Erovenko, I. V. (2018). Ebola could be eradicated
279 through voluntary vaccination. *Royal Society Open Science*, 5(1):171591.
- 280 Broom, M., Rychtář, J., and Spears-Gill, T. (2016). The game-theoretical model of using insecticide-
281 treated bed-nets to fight malaria. *Applied Mathematics*, 7(09):852–860.
- 282 Brown, K. and Leggat, P. A. (2016). Human monkeypox: current state of knowledge and implications for
283 the future. *Tropical Medicine and Infectious Disease*, 1(1):8.
- 284 CDC (2003). Multistate outbreak of monkeypox—Illinois, Indiana, and Wisconsin, 2003.
- 285 Chang, S. L., Piraveenan, M., Pattison, P., and Prokopenko, M. (2020). Game theoretic modelling of
286 infectious disease dynamics and intervention methods: a review. *Journal of Biological Dynamics*,
287 14(1):57–89.
- 288 Cheng, E., Gambhirrao, N., Patel, R., Zhouandai, A., Rychtář, J., and Taylor, D. (2020). A game-
289 theoretical analysis of Poliomyelitis vaccination. *Journal of Theoretical Biology*.
- 290 Chouhan, A., Maiwand, S., Ngo, M., Putalapattu, V., Rychtář, J., and Taylor, D. (2019). Game-theoretical
291 model of retroactive hepatitis B vaccination in China. preprint.
- 292 CIA (2019). The world factbook - birth rate. <https://www.cia.gov/library/>

293 [publications/the-world-factbook/fields/345.html](https://www.cia.gov/library/publications/the-world-factbook/fields/345.html). Accessed December 13,
294 2019.

295 Crawford, K., Lancaster, A., Oh, H., and Rychtář, J. (2015). A voluntary use of insecticide-treated cattle
296 can eliminate african sleeping sickness. *Letters in Biomathematics*, 2(1):91–101.

297 Di Giulio, D. B. and Eckburg, P. B. (2004). Human monkeypox: an emerging zoonosis. *The Lancet*
298 *infectious diseases*, 4(1):15–25.

299 Doshi, R. H., Guagliardo, S. A. J., Doty, J. B., Babeaux, A. D., Matheny, A., Burgado, J., Townsend,
300 M. B., Morgan, C. N., Satheshkumar, P. S., and Ndakala, N. (2019). Epidemiologic and ecologic
301 investigations of monkeypox, Likouala Department, Republic of the Congo, 2017. *Emerging Infectious*
302 *Diseases*, 25(2):273.

303 Doshi, R. H., Guagliardo, S. A. J., Dzabatou-Babeaux, A., Likouayoulou, C., Ndakala, N., Moses,
304 C., Olson, V., McCollum, A. M., and Petersen, B. W. (2018). Strengthening of surveillance during
305 monkeypox outbreak, Republic of the Congo, 2017. *Emerging Infectious Diseases*, 24(6):1158.

306 Doty, J. B., Malekani, J. M., Kalemba, L. N., Stanley, W. T., Monroe, B. P., Nakazawa, Y. U., Mauldin,
307 M. R., Bakambana, T. L., Liyandja Dja Liyandja, T., and Braden, Z. H. (2017). Assessing monkeypox
308 virus prevalence in small mammals at the human–animal interface in the Democratic Republic of the
309 Congo. *Viruses*, 9(10):283.

310 Durski, K. N., McCollum, A. M., Nakazawa, Y., Petersen, B. W., Reynolds, M. G., Briand, S., Djingarey,
311 M. H., Olson, V., Damon, I. K., and Khalakdina, A. (2018). Emergence of monkeypox — West and
312 Central Africa, 1970–2017. *Morbidity and Mortality Weekly Report*, 67(10):306.

313 Elde, N. C., Child, S. J., Eickbush, M. T., Kitzman, J. O., Rogers, K. S., Shendure, J., Geballe, A. P.,
314 and Malik, H. S. (2012). Poxviruses deploy genomic accordions to adapt rapidly against host antiviral
315 defenses. *Cell*, 150(4):831–841.

316 Elderfield, R. A., Watson, S. J., Godlee, A., Adamson, W. E., Thompson, C. I., Dunning, J., Fernandez-
317 Alonso, M., Blumenkrantz, D., Hussell, T., and Zambon, M. (2014). Accumulation of human-adapting
318 mutations during circulation of A (H1N1) pdm09 influenza virus in humans in the United Kingdom.
319 *Journal of Virology*, 88(22):13269–13283.

320 Eteng, W.-E., Mandra, A., Doty, J., Yinka-Ogunleye, A., Aruna, S., Reynolds, M. G., McCollum, A. M.,
321 Davidson, W., Wilkins, K., and Saleh, M. (2018). Notes from the field: Responding to an outbreak
322 of monkeypox using the one health approach—Nigeria, 2017–2018. *Morbidity and Mortality Weekly*
323 *Report*, 67(37):1040.

324 Falendysz, E. A., Lopera, J. G., Doty, J. B., Nakazawa, Y., Crill, C., Lorenzsonn, F., Lem’s, N. K.,
325 Ronderos, M. D., Mejia, A., and Malekani, J. M. (2017). Characterization of Monkeypox virus infection
326 in African rope squirrels (*Funisciurus* sp.). *PLoS Neglected Tropical Diseases*, 11(8):e0005809.

327 Galvani, A. P., Reluga, T. C., and Chapman, G. B. (2007). Long-standing influenza vaccination policy is
328 in accord with individual self-interest but not with the utilitarian optimum. *Proceedings of the National*
329 *Academy of Sciences*, 104(13):5692–5697.

330 Hayssen, V. (2008). Reproductive effort in squirrels: ecological, phylogenetic, allometric, and latitudinal
331 patterns. *Journal of Mammalogy*, 89(3):582–606.

332 Herp, M. V., Parqué, V., Rackley, E., and Ford, N. (2003). Mortality, violence and lack of access to
333 healthcare in the Democratic Republic of Congo. *Disasters*, 27(2):141–153.

334 Jackson, R. J., Ramsay, A. J., Christensen, C. D., Beaton, S., Hall, D. F., and Ramshaw, I. A. (2001).
335 Expression of mouse interleukin-4 by a recombinant ectromelia virus suppresses cytolytic lymphocyte
336 responses and overcomes genetic resistance to mousepox. *Journal of Virology*, 75(3):1205–1210.

337 Ježek, Z., Szczeniowski, M., Paluku, K., Mutombo, M., and Grab, B. (1988). Human monkeypox:
338 confusion with chickenpox. *Acta Tropica*, 45(4):297–307.

339 Ježek, Z., Szczeniowski, M., Paluku, K., and Mutombo, M. (1987). Human monkeypox: clinical features
340 of 282 patients. *Journal of infectious diseases*, 156(2):293–298.

341 Khodakevich, L., Ježek, Z., and Messinger, D. (1988). Monkeypox virus: ecology and public health
342 significance. *Bulletin of the World Health Organization*, 66(6):747.

343 Khodakevich, L., Szczeniowski, M., Ježek, Z., Marennikova, S., Nakano, J., and Messinger, D. (1987).
344 The role of squirrels in sustaining monkeypox virus transmission. *Tropical and Geographical Medicine*,
345 39(2):115–122.

346 Klein, S. R. M., Foster, A. O., Feagins, D. A., Rowell, J. T., and Erovenko, I. V. (2019). Optimal voluntary
347 and mandatory insect repellent usage and emigration strategies to control the chikungunya outbreak on

- 348 Reunion Island. *Preprint*.
- 349 Kobe, J., Pritchard, N., Short, Z., Erovenko, I. V., Rychtář, J., and Rowell, J. T. (2018). A game-theoretic
350 model of cholera with optimal personal protection strategies. *Bulletin of Mathematical Biology*,
351 80(10):2580–2599.
- 352 Lambert de Rouvroit, A. and Heegaard, E. D. (2016). Total costs associated with replicating and
353 non-replicating smallpox vaccines. *Global Security: Health, Science and Policy*, 1(1):3–9.
- 354 Lauko, I., Pinter, G., and TeWinkel, R. E. (2018). Equilibrium analysis for an epidemic model with a
355 reservoir for infection. *Letters in Biomathematics*, 5(1):255–274.
- 356 Le, Q. M., Sakai-Tagawa, Y., Ozawa, M., Ito, M., and Kawaoka, Y. (2009). Selection of H5N1 influenza
357 virus PB2 during replication in humans. *Journal of Virology*, 83(10):5278–5281.
- 358 Learned, L. A., Reynolds, M. G., Wassa, D. W., Li, Y., Olson, V. A., Karem, K., Stempora, L. L., Braden,
359 Z. H., Kline, R., and Likos, A. (2005). Extended interhuman transmission of monkeypox in a hospital
360 community in the Republic of the Congo, 2003. *The American Journal of Tropical Medicine and*
361 *Hygiene*, 73(2):428–434.
- 362 Likos, A. M., Sammons, S. A., Olson, V. A., Frace, A. M., Li, Y., Olsen-Rasmussen, M., Davidson, W.,
363 Galloway, R., Khristova, M. L., and Reynolds, M. G. (2005). A tale of two clades: monkeypox viruses.
364 *Journal of General Virology*, 86(10):2661–2672.
- 365 Louten, J. (2016). *Essential human virology*. Academic Press.
- 366 Martinez, A., Machado, J., Sanchez, E., and Erovenko, I. (2019). Optimal vaccination strategies to reduce
367 endemic levels of meningitis in Africa. *Preprint*.
- 368 McCollum, A. M. and Damon, I. K. (2013). Human monkeypox. *Clinical infectious diseases*, 58(2):260–
369 267.
- 370 Meyer, H., Ehmann, R., and Smith, G. L. (2020). Smallpox in the post-eradication era. *Viruses*, 12(2):138.
- 371 Monroe, B. P., Doty, J. B., Moses, C., Ibata, S., Reynolds, M., and Carroll, D. (2015). Collection and
372 utilization of animal carcasses associated with zoonotic disease in tshuapa district, the democratic
373 republic of the congo, 2012. *Journal of Wildlife Diseases*, 51(3):734–738.
- 374 Moss, B. (2013). Poxvirus DNA replication. *Cold Spring Harbor Perspectives in Biology*, 5(9):a010199.
- 375 Mwamba, D., Kebela, B., Shongo, R., Pukuta, E., and Kayembe, N. (2014). Profil épidémiologique du
376 monkeypox en RDC, 2010-2014. *Ann African Med*, 8:1855–60.
- 377 Nalca, A. and Zumbrun, E. E. (2010). ACAM2000: the new smallpox vaccine for United States Strategic
378 National Stockpile. *Drug Design, Development and Therapy*, 4:71.
- 379 Nolen, L. D., Osadebe, L., Katomba, J., Likofata, J., Mukadi, D., Monroe, B., Doty, J., Malekani, J.,
380 Kabamba, J., and Bomponda, P. L. (2015). Introduction of monkeypox into a community and household:
381 risk factors and zoonotic reservoirs in the Democratic Republic of the Congo. *The American Journal of*
382 *Tropical Medicine and Hygiene*, 93(2):410–415.
- 383 Orwa, T. O., Mbogo, R. W., and Luboobi, L. S. (2018). Mathematical model for the in-host malaria
384 dynamics subject to malaria vaccines. *Letters in Biomathematics*, 5(1):222–251.
- 385 Padmanabhan, P., Seshaiyer, P., and Castillo-Chavez, C. (2017). Mathematical modeling, analysis and
386 simulation of the spread of zika with influence of sexual transmission and preventive measures. *Letters*
387 *in Biomathematics*, 4(1):148–166.
- 388 Reynolds, M. G., Carroll, D. S., Olson, V. A., Hughes, C., Galley, J., Likos, A., Montgomery, J. M.,
389 Suu-Ire, R., Kwasi, M. O., and Root, J. J. (2010). A silent enzootic of an orthopoxvirus in Ghana,
390 West Africa: evidence for multi-species involvement in the absence of widespread human disease. *The*
391 *American Journal of Tropical Medicine and Hygiene*, 82(4):746–754.
- 392 Reynolds, M. G. and Damon, I. K. (2012). Outbreaks of human monkeypox after cessation of smallpox
393 vaccination. *Trends in Microbiology*, 20(2):80–87.
- 394 Reynolds, M. G., McCollum, A. M., Nguete, B., Shongo Lushima, R., and Petersen, B. W. (2017).
395 Improving the care and treatment of monkeypox patients in low-resource settings: applying evidence
396 from contemporary biomedical and smallpox biodefense research. *Viruses*, 9(12):380.
- 397 Rimoin, A. W., Mulembakani, P. M., Johnston, S. C., Smith, J. O. L., Kisalu, N. K., Kinkela, T. L.,
398 Blumberg, S., Thomassen, H. A., Pike, B. L., and Fair, J. N. (2010). Major increase in human
399 monkeypox incidence 30 years after smallpox vaccination campaigns cease in the Democratic Republic
400 of Congo. *Proceedings of the National Academy of Sciences*, 107(37):16262–16267.
- 401 Scheckelhoff, K., Ejaz, A., and Erovenko, I. V. (2019). A game-theoretic model of optimal clean
402 equipment usage to prevent hepatitis C among injecting drug users. *Preprint*.

- 403 Shchelkunov, S. N., Marennikova, S. S., and Moyer, R. W. (2006). *Orthopoxviruses pathogenic for*
404 *humans*. Springer Science & Business Media.
- 405 Shim, E., Grefenstette, J. J., Albert, S. M., Cakouros, B. E., and Burke, D. S. (2012). A game dynamic
406 model for vaccine skeptics and vaccine believers: measles as an example. *Journal of Theoretical*
407 *Biology*, 295:194–203.
- 408 Shim, E., Kochin, B., and Galvani, A. (2009). Insights from epidemiological game theory into gender-
409 specific vaccination against rubella. *Mathematical Biosciences and Engineering: MBE*, 6(4):839–854.
- 410 Sklenovská, N. and Van Ranst, M. (2018). Emergence of monkeypox as the most important orthopoxvirus
411 infection in humans. *Frontiers in public health*, 6.
- 412 Sykes, D. and Rychtář, J. (2015). A game-theoretic approach to valuating toxoplasmosis vaccination
413 strategies. *Theoretical Population Biology*, 105:33–38.
- 414 Usman, S. and Adamu, I. I. (2017). Modeling the transmission dynamics of the monkeypox virus infection
415 with treatment and vaccination interventions. *Journal of Applied Mathematics and Physics*, 5(12):2335.
- 416 Van den Driessche, P. and Watmough, J. (2002). Reproduction numbers and sub-threshold endemic
417 equilibria for compartmental models of disease transmission. *Mathematical Biosciences*, 180(1-2):29–
418 48.
- 419 Weinstein, R. A., Nalca, A., Rimoin, A. W., Bavari, S., and Whitehouse, C. A. (2005). Reemergence of
420 monkeypox: prevalence, diagnostics, and countermeasures. *Clinical Infectious Diseases*, 41(12):1765–
421 1771.
- 422 WHO (2017). Weekly bulletin on outbreaks and other emergencies: week 18: 29 April–05 May 2017.
423 *Weekly Bulletin on Outbreak and other Emergencies*, pages 1–14.
- 424 Wollenberg, A. and Engler, R. (2004). Smallpox, vaccination and adverse reactions to smallpox vaccine.
425 *Current Opinion in Allergy and Clinical Immunology*, 4(4):271–275.
- 426 World Bank (2019). Life expectancy at birth. [https://data.worldbank.org/indicator/SP.](https://data.worldbank.org/indicator/SP.DYN.LE00.IN?cid=GPD_10)
427 [DYN.LE00.IN?cid=GPD_10](https://data.worldbank.org/indicator/SP.DYN.LE00.IN?cid=GPD_10). Accessed December 13, 2019.
- 428 Wu, B., Fu, F., and Wang, L. (2011). Imperfect vaccine aggravates the long-standing dilemma of voluntary
429 vaccination. *PloS One*, 6(6):e20577.
- 430 Yinka-Ogunleye, A., Aruna, O., Ogoina, D., Aworabhi, N., Eteng, W., Badaru, S., Mohammed, A.,
431 Agyenyi, J., Etebu, E., and Numbere, T.-W. (2018). Reemergence of human monkeypox in Nigeria,
432 2017. *Emerging Infectious Diseases*, 24(6):1149.

433 APPENDIX A1 STEP-BY-STEP SOLUTIONS TO EQUILIBRIUM STATES

434 In this section we find the equilibrium states of the system (1) - (9). We will look for solutions of the
435 following system.

$$0 = \Lambda_s - \left(\mu_s + \beta_{ss} \frac{I_s}{N_s} \right) S_s \quad (15)$$

$$0 = \beta_{ss} \frac{I_s}{N_s} S_s - (\mu_s + \nu_s) E_s \quad (16)$$

$$0 = \nu_s E_s - (\mu_s + d_s + \rho_s) I_s \quad (17)$$

$$0 = \rho_s I_s - \mu_s R_s \quad (18)$$

$$0 = \Lambda_h - \left(\mu_h + \left(\beta_{sh} \frac{I_s}{N_s} + \beta_{hh} \frac{I_h}{N_h} \right) + \alpha_h \right) S_h \quad (19)$$

$$0 = \alpha_h S_h - \mu_h V_h \quad (20)$$

$$0 = \left(\beta_{sh} \frac{I_s}{N_s} + \beta_{hh} \frac{I_h}{N_h} \right) S_h - (\mu_h + \nu_h) E_h \quad (21)$$

$$0 = \nu_h E_h - (\mu_h + d_h + \rho_h) I_h \quad (22)$$

$$0 = \rho_h I_h - \mu_h R_h \quad (23)$$

436 where (15) - (18) are equations for the squirrels and (19) - (23) are for the humans.

437 We will distinguish three equilibrium states ε^0 , ε^* and ε^\dagger depending on the existence of infection
438 among squirrels and humans. However, by (16), (18), (20), (21) and (23), no matter which equilibrium

439 state, the following formulas will always be valid.

$$E_s = \frac{\mu_s + d_s + \rho_s}{v_s} I_s \quad (24)$$

$$R_s = \frac{\rho_s}{\mu_s} I_s \quad (25)$$

$$V_h = \frac{\alpha_h}{\mu_h} S_h \quad (26)$$

$$E_h = \frac{\mu_h + d_h + \rho_h}{v_h} I_h \quad (27)$$

$$R_h = \frac{\rho_h}{\mu_h} I_h \quad (28)$$

440 **Disease-free equilibrium, ε^0**

441 Assume $I_h^0 = I_s^0 = 0$. It follows from (24) that $E_s^0 = 0$, from (25) that $R_s^0 = 0$ and from (15) that $S_s^0 = \frac{\Lambda_s}{\mu_s}$.

442 Also, by (27), $E_h^0 = 0$ and, by (28), $R_h^0 = 0$. It follows from (19) that

$$S_h^0 = \frac{\Lambda_h}{\alpha_h + \mu_h}. \quad (29)$$

443 and thus, by (26),

$$V_h^0 = \frac{\Lambda_h}{\mu_h} \cdot \frac{\alpha_h}{\alpha_h + \mu_h}. \quad (30)$$

444 The stability of ε^0 was discussed and the basic reproduction numbers were derived in Usman and
445 Adamu (2017) using the next-generation matrix method of Van den Driessche and Watmough (2002).
446 Here we present an alternative derivation of the basic reproduction numbers.

447 Assume there is an infected squirrel in an otherwise disease-free population. The squirrel stays infected
448 for a period of $(\mu_s + d_s + \rho_s)^{-1}$ during which it exposes susceptible individuals at the rate $\beta_{ss} \frac{S_s^0}{N_s^0} = \beta_{ss}$.
449 The newly exposed individuals end up in the I_s compartment with probability $\frac{v_s}{\mu_s + v_s}$. Consequently, the
450 number of secondary infections from a single infected squirrel in an otherwise disease free equilibrium is
451 given by

$$R_{0ss} = \beta_{ss} \cdot \left(\frac{1}{\mu_s + d_s + \rho_s} \right) \cdot \left(\frac{v_s}{\mu_s + v_s} \right). \quad (31)$$

452 Similarly, we can derive that the number of secondary infections caused by a single infected human in
453 otherwise disease-free population is

$$R_{0hh} = \beta_{hh} \frac{S_h^0}{N_h^0} \cdot \left(\frac{1}{\mu_h + d_h + \rho_h} \right) \cdot \left(\frac{v_h}{\mu_h + v_h} \right) = \frac{v_h \beta_{hh} \mu_h}{(\mu_h + d_h + \rho_h)(\mu_h + v_h)(\alpha_h + \mu_h)}. \quad (32)$$

454 **Case when $I_s > 0$.**

By adding (15) - (18), we get $0 = \Lambda_s - \mu_s N_s^* - d_s I_s^*$ which yields

$$I_s^* = \frac{\Lambda_s - \mu_s N_s^*}{d_s}. \quad (33)$$

455 By (24), (16) becomes

$$0 = \frac{\beta_{ss} I_s^* S_s^*}{N_s^*} - (\mu_s + v_s) \cdot \frac{\mu_s + d_s + \rho_s}{v_s} \cdot I_s^* \quad (34)$$

456 and since we are assuming $I_s^* > 0$, we can divide by I_s and get

$$S_s^* = N_s^* \cdot \frac{(\mu_s + \nu_s)(\mu_s + d_s + \rho_s)}{\beta_{ss} \nu_s} = \frac{N_s^*}{R_{0ss}}. \quad (35)$$

457 Consequently,

$$\begin{aligned} N_s^* &= S_s^* + E_s^* + I_s^* + R_s^* \\ &= \frac{N_s^*}{R_{0ss}} + \left(\frac{\mu_s + d_s + \rho_s}{\nu_s} + 1 + \frac{\rho_s}{\mu_s} \right) \cdot \left(\frac{\Lambda_s - \mu_s N_s^*}{d_s} \right) \end{aligned} \quad (36)$$

458 which yields

$$N_s^* = \frac{\Lambda_s \cdot \left(\frac{\mu_s + d_s + \rho_s}{\nu_s} + 1 + \frac{\rho_s}{\mu_s} \right)}{\mu_s \cdot \left(\frac{\mu_s + d_s + \rho_s}{\nu_s} \right) + \mu_s + \rho_s + d_s \left(1 - \frac{1}{R_{0ss}} \right)}. \quad (37)$$

459 **Fully endemic equilibrium, ε^* , human population**

460 Adding (19) - (23) yields

$$0 = \Lambda_h - \mu_h N_h^* - d_h I_h^* \quad (38)$$

461 and consequently

$$I_h^* = \frac{\Lambda_h - \mu_h N_h^*}{d_h}. \quad (39)$$

462 Substituting (39) into (23), (21), (19), (20) we get

$$R_h^* = \frac{\rho_h}{\mu_h} \cdot \left(\frac{\Lambda_h - \mu_h N_h^*}{d_h} \right) \quad (40)$$

$$E_h^* = \left(\frac{\mu_h + d_h + \rho_h}{\nu_h} \right) \cdot \left(\frac{\Lambda_h - \mu_h N_h^*}{d_h} \right) \quad (41)$$

$$S_h^* = \frac{(\mu_h + \nu_h) \cdot \left(\frac{\mu_h + d_h + \rho_h}{\nu_h} \right) \cdot \left(\frac{\Lambda_h - \mu_h N_h^*}{d_h} \right)}{\beta_{sh} \left(\frac{I_s^*}{N_s^*} \right) + \beta_{hh} \frac{\Lambda_h - \mu_h N_h^*}{d_h}} \quad (42)$$

$$V_h^* = \frac{\alpha_h}{\mu_h} \cdot \left(\frac{(\mu_h + \nu_h) \cdot \left(\frac{\mu_h + d_h + \rho_h}{\nu_h} \right) \cdot \left(\frac{\Lambda_h - \mu_h N_h^*}{d_h} \right)}{\beta_{sh} \left(\frac{I_s^*}{N_s^*} \right) + \beta_{hh} \frac{\Lambda_h - \mu_h N_h^*}{d_h}} \right). \quad (43)$$

463 Since $N_h^* = S_h^* + V_h^* + E_h^* + I_h^* + R_h^*$, we get

$$\begin{aligned}
N_h^* &= \frac{\rho_h}{\mu_h} \left(\frac{\Lambda_h - \mu_h N_h^*}{d_h} \right) + \left(\frac{\mu_h + d_h + \rho_h}{v_h} \right) \cdot \left(\frac{\Lambda_h - \mu_h N_h^*}{d_h} \right) \\
&+ \frac{\Lambda_h - \mu_h N_h^*}{d_h} + \frac{(\mu_h + v_h) \cdot \left(\frac{\mu_h + d_h + \rho_h}{v_h} \right) \cdot \left(\frac{\Lambda_h - \mu_h N_h^*}{d_h} \right)}{\beta_{sh} \left(\frac{I_s^*}{N_s^*} \right) + \beta_{hh} \frac{\Lambda_h - \mu_h N_h^*}{d_h}} \\
&+ \frac{\alpha_h}{\mu_h} \cdot \left(\frac{(\mu_h + v_h) \cdot \left(\frac{\mu_h + d_h + \rho_h}{v_h} \right) \cdot \left(\frac{\Lambda_h - \mu_h N_h^*}{d_h} \right)}{\beta_{sh} \left(\frac{I_s^*}{N_s^*} \right) + \beta_{hh} \frac{\Lambda_h - \mu_h N_h^*}{d_h}} \right).
\end{aligned} \tag{44}$$

464 This yields that N_h^* is a positive root of the equation $AN_h^2 + BN_h + C = 0$, i.e.

$$N_h^* = \frac{-B + \sqrt{B^2 - 4AC}}{2A} \tag{45}$$

465 where

$$A = g \cdot d_h + k \cdot c \cdot d_h \cdot \mu_h \left(1 + \frac{\alpha_h}{\mu_h} \right) + g \cdot \mu_h (1 + z + k) \tag{46}$$

$$B = d_h \cdot \beta_{hh} \cdot \Lambda_h - (g \cdot \Lambda_h - \beta_{hh} \cdot \Lambda_h \cdot \mu_h) (1 + z + k) - k \cdot c \cdot d_h \cdot \Lambda_h \left(1 + \frac{\alpha_h}{\mu_h} \right) \tag{47}$$

$$C = -\Lambda_h^2 \cdot \beta_{hh} (1 + z + k) \tag{48}$$

466 and

$$k = \frac{\mu_h + d_h + \rho_h}{v_h} \tag{49}$$

$$g = d_h \beta_{sh} \left(\frac{I_s^*}{N_s^*} \right) - \beta_{hh} \mu_h \tag{50}$$

$$c = \mu_h + v_h \tag{51}$$

$$z = \frac{\rho_h}{\mu_h}. \tag{52}$$

467 The positive solution for N_h^* from (45) can then be recursively substituted into (39), (40), (41), (42),
468 and (43) to get closed-form formulas for equilibrium values.

469 **Semi-endemic equilibrium**

470 Here, $I_s = 0$. As above, by adding all equations (19) - (23) we get $0 = \Lambda_h - \mu_h N_h^\dagger - d_h I_h^\dagger$ which yields

$$I_h^\dagger = \frac{\Lambda_h - \mu_h N_h^\dagger}{d_h}. \tag{53}$$

471 Substituting (53) into (23), (21), (19) we get

$$R_h^\dagger = \frac{\rho_h}{\mu_h} \cdot \left(\frac{\Lambda_h - \mu_h N_h^\dagger}{d_h} \right) \quad (54)$$

$$E_h^\dagger = \left(\frac{\mu_h + d_h + \rho_h}{v_h} \right) \cdot \left(\frac{\Lambda_h - \mu_h N_h^\dagger}{d_h} \right) \quad (55)$$

$$S_h^\dagger = \frac{(\mu_h + v_h) \cdot \left(\frac{\mu_h + d_h + \rho_h}{v_h} \right) \cdot \left(\frac{\Lambda_h - \mu_h N_h^\dagger}{d_h} \right)}{\beta_{sh} \left(\frac{I_s^\dagger}{N_s^\dagger} \right) + \beta_{hh} \frac{\Lambda_h - \mu_h N_h^\dagger}{N_h^\dagger}}. \quad (56)$$

472 Since $I_s^\dagger = 0$, we get

$$S_h^\dagger = N_h^\dagger \cdot \frac{(\mu_h + d_h + \rho_h)(\mu_h + v_h)}{v_h \beta_{hh}} \quad (57)$$

473 and consequently, by (20),

$$V_h^\dagger = N_h^\dagger \cdot \left(\frac{\alpha_h}{\mu_h} \right) \cdot \frac{(\mu_h + d_h + \rho_h)(\mu_h + v_h)}{v_h \beta_{hh}}. \quad (58)$$

474 Finally, summing up (53) - (58), we obtain the closed-form formula

$$N_h^\dagger = \frac{\Lambda_h \cdot \left(\frac{\rho_h}{\mu_h} + \frac{\mu_h + d_h + \rho_h}{v_h} + 1 \right)}{d_h + \mu_h \cdot \left(\frac{\rho_h}{\mu_h} + \frac{\mu_h + d_h + \rho_h}{v_h} + 1 \right) - d_h \cdot \left(\frac{\mu_h + d_h + \rho_h}{v_h} \left(\frac{\mu_h + v_h}{\beta_{hh}} \right) + \frac{\alpha_h}{\mu_h} \left(\frac{\mu_h + d_h + \rho_h}{v_h} \right) \left(\frac{\mu_h + v_h}{\beta_{hh}} \right) \right)} \quad (59)$$

475 which can be used to calculate equilibrium values.