

Modelling the creation of ecological traps by environmentally persistent pathogens

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Abstract

Patterns of habitat quality are expected to play an important role in the dynamics of wildlife metapopulations, with the expectation that high quality habitat serves to stabilize and maintain occupancy. However, we might expect that the costs and benefits of different quality habitat might be modulated, or possibly even reversed, by the presence of an environmentally transmitted pathogen. We explore this possibility through the development of a stochastic patch occupancy model of host-pathogen dynamics, in which we can vary both the environmental longevity of the pathogen, and the distribution of habitat quality in the host metapopulation.

1 Introduction

Many wildlife diseases are caused by pathogens that can persist, and remain infectious, for long periods of time in the environment. Examples include chronic wasting disease ([1]), anthrax ([2]), plague ([3]), and white nose syndrome ([4]), among others. This environmental persistence creates environmental pathogen reservoirs from which hosts can become infected without direct contact with an

infectious individual. This additional transmission pathway can have important consequences for disease dynamics, with models showing that increased environmental persistence generally facilitates increased pathogen persistence and spread relative to direct transmission alone ([5], [6], [7]).

Since environmental transmission is spatially explicit (i.e. environmental reservoirs can only infect local residents), its role in disease dynamics may further depend on the spatial structure of the host population. In particular, we expect that host population structure and movement will be influenced by heterogeneity in quality among the habitat patches in a metapopulation. Indeed, the quality of a habitat patch can affect its extinction and colonization rates, as well as its contribution to the colonization of other empty patches ([8]). These processes in turn may then influence where environmental pathogen reservoirs get established and how they affect disease dynamics and host occupancy throughout the metapopulation.

Specifically, we expect that high quality habitat patches, which support greater host density and traffic than lower quality habitat, might be more likely to form pathogen reservoirs. As these reservoirs are undetectable to the host, we predict that high quality patches will continue to attract – and infect – susceptible immigrants, effectively creating an ecological trap ([5]). In addition, the greater traffic through high quality patches may further facilitate pathogen spread by positioning high quality patches as metapopulation-scale superspreaders ([9]). Similarly, we expect that low quality patches, which see relatively less host traffic, will be less likely to develop pathogen reservoirs and thus may serve as refuges on which susceptible hosts can escape infection.

Many, if not all, of the pathogens listed above affect spatially structured host populations (e.g. plague in prairie dog colonies, [10]), and thus understanding how environmental transmission interacts with patterns of habitat quality is critical to managing disease in these systems. In this study then, we seek to explore how a pathogen’s environmental longevity (how long it can persist and remain infectious in the environment), and the distribution of habitat quality in a metapopulation interact to influence pathogen spread and host occupancy, with a specific interest in the roles played by high and low quality habitat.

2 Methods

2.1 Model Structure

To address the above questions, we developed a stochastic patch occupancy model (SPOM) in which each patch can be in one of three possible states: occupied by susceptible hosts (S), occupied by infectious hosts (I), and unoccupied by the host (\emptyset). State transitions are governed by host colonization, extinction, and infection rates, where a susceptible population can become infectious either through direct contact with infectious immigrants, or through a local pathogen reservoir (Table 1).

Building on the framework developed by [11], the rate at which patch i ,

with quality A_i , is colonized by individuals in state $X \in (S, I)$ is given by its connectivity:

$$C_{Xi} = A_i^{\xi_{im}} \sum_{j \neq i} \phi_j A_j^{\xi_{em}} \exp(-D d_{ij}), \quad (1)$$

where ϕ_j is an indicator function that is 1 if patch j is in state X and is 0 otherwise; ξ_{im} and ξ_{em} control how rates of immigration and emigration, respectively, scale with patch quality; D is the inverse of mean dispersal distance; and d_{ij} is the distance between patches i and j . Essentially, C_{Xi} sums the colonization effort from all patches in state X to patch i .

The extinction rate of patch i in state X is given by:

$$E_{Xi} = \frac{\mu_X}{A_i^\alpha}, \quad (2)$$

where μ_X is the extinction rate of a patch of unit quality in state X (here we assume that $\mu_I \geq \mu_S$ to account for disease-induced mortality), and α controls how extinction rate scales with patch quality. When an infectious population goes extinct, we assume that the hosts leave behind an infectious pathogen reservoir on the patch.

Susceptible populations are infected via direct contact with infectious colonists at rate $\delta \lambda_{Ii}$, where δ is the direct transmission probability. In addition, environmental transmission can take place when a susceptible population occupies a previously infected patch, and occurs at rate:

$$\gamma(\tau_i) = \gamma_0 \exp(-r\tau_i), \quad (3)$$

where γ_0 is the initial infection rate of the pathogen reservoir, τ_i is the time since last infectious occupancy ($\tau_i = \infty$, and thus $\gamma(\tau_i) = 0$, if the patch has never been occupied by infectious hosts), and r is the pathogen's decay rate in the environment.

2.2 Model Parameterization

The model outlined above was parameterized in five different ways to explore the consequences of different assumptions (Table 2). The default model was parameterized according to the following assumptions. (1) High quality patches maintain higher population sizes and therefore produce more colonists than low quality patches and have lower extinction rates than low quality patches ($\xi_{im} = 0.5$, $\alpha = 1$, [11]). (2) High quality patches attract more colonists than low quality patches ($\xi_{em} = 0.5$, [11]). (3) Connectivity among patches is determined entirely by quality ($d_{ij} = 1$ for all $i \neq j$). In addition to these assumptions, parameters were chosen so that without infection, approximately 0.75 of the metapopulation was occupied ($\mu_S = 0.1$, $D = 5$), and a range of epidemiological behaviors – disease-free, endemic disease, and pandemic disease (i.e. all host populations infectious) – was feasible, depending on the pathogen's environmental longevity ($\mu_I = 0.5$, $\delta = 0.5$, $\gamma_0 = 0.5$).

110 The baseline model above assumes that all patches are equally accessible
 111 from all other patches (i.e. are separated by distance 1), so to examine the effects
 112 of a more rigid spatial structure, we also implemented a model with patches
 113 arranged in a square lattice such that each patch is only accessible from its four
 114 neighboring patches (Local movement, Table 2). In addition, we explored a
 115 model with $\alpha = 0$ so that the extinction rate was constant with patch quality,
 116 simulating the effects of high environmental stochasticity, i.e. environmental
 117 stochasticity is strong enough that the population size of a patch does not
 118 affect its extinction rate (High env. stoc., Table 2). To explore the disease
 119 dynamics when transmission is driven largely by the environmental reservoir
 120 we parameterized a model with $\delta = 0.1$ to reduce the relative importance of
 121 direct transmission (Dominant reservoir trans., Table 2). Lastly, to simulate a
 122 more virulent pathogen, we parameterized a model with $\mu_I = 1$ to substantially
 123 increase disease-induced mortality (High disease mortality, Table 2).

124 **2.3 Simulation studies**

125 For each of the above parameterizations, we performed two different simulation
 126 experiments to explore the combined effects of pathogen longevity and habitat
 127 quality on disease dynamics and metapopulation occupancy. In the first of these,
 128 we investigated the effect of increasing habitat heterogeneity on the dynamics
 129 of pathogens with a range of environmental longevities. For these simulations,
 130 we explored 10 values of pathogen longevity ranging from 0.1 to 3 times the
 131 expected residence time of a healthy population on a unit quality patch (i.e. 10
 132 units of time), with values representing the half-life of the pathogen's infectivity
 133 and 10 values for the variance of the patch quality distribution, ranging from
 134 0.02 to 0.2. For each simulation (100 replicates for each of the 100 longevity-
 135 variance combinations), qualities for 100 patches were drawn from a uniform
 136 distribution with fixed mean 1 and given variance. In each case, an entirely sus-
 137 ceptible population was simulated until it reached an approximate steady state,
 138 at which point a randomly chosen occupied patch was infected. The state of
 139 the metapopulation was then tracked for 5000 time steps. We recorded the pro-
 140 portion of patches in each state at the end of each simulation and classified the
 141 simulation as host extinction, disease-free (no infectious populations after 5000
 142 time steps), endemic (both susceptible and infectious populations present), or
 143 pandemic (only infectious populations). In addition, to identify the roles played
 144 by individual patches of varying quality, for every simulation, we recorded the
 145 quality of each patch, the proportion of time spent in each state (i.e. occupied
 146 by susceptible hosts, occupied by infectious hosts, and unoccupied), and the
 147 number of transitions between each state.

148 The above simulations held the mean of the patch quality distribution con-
 149 stant and explored the effect of increasing habitat heterogeneity around that
 150 mean (i.e. by expanding the quality distribution symmetrically about 1). To
 151 further parse out the relative influence of high and low quality patches sep-
 152 arately, we performed a second series of simulations where we expanded the
 153 range of the uniform quality distribution to favor either high or low quality

154 patches (i.e. by expanding the quality distribution asymmetrically about 1).
 155 For these simulations, at each of the 10 longevities used above, we simulated
 156 100 metapopulations for each of four patch quality distributions: low variance,
 157 (patch qualities ranging 0.75 to 1.25); intermediate variance, high quality (range
 158 0.75 to 1.75); intermediate variance, low quality (range 0.25 to 1.25); and high
 159 variance (range 0.25 to 1.75).

160 Continuous time stochastic simulations of the above model were imple-
 161 mented in the R language ([12]) using the Gillespie algorithm ([13]). Code
 162 is available at <https://github.com/clint-leach/Metapop-Disease>.

163 3 Results

164 In the baseline parameterization, infectious occupancy increased substantially
 165 with longevity, while susceptible and total occupancy decreased (Supplemental).
 166 However, increasing the variance of the patch quality distribution around a mean
 167 of 1 had relatively little effect on average occupancy (Supplemental). The four
 168 alternative parameterizations (models with local movement, high environmental
 169 stochasticity, dominant reservoir transmission, and high disease-induced mor-
 170 tality) produced qualitatively similar results, though pathogen spread in these
 171 models was generally more limited (Supplemental).

172 We found that distributions with more low quality habitat (distribution
 173 ranging from 0.25 to 1.25) increased mean susceptible occupancy, while dis-
 174 tributions with more high quality habitat (range from 0.75 to 1.75) decreased
 175 mean susceptible occupancy, with the largest differences observed at interme-
 176 diate longevities (Fig 1(a)). Conversely, low quality habitat distributions de-
 177 creased infectious occupancy, while high quality distributions increased infec-
 178 tious occupancy (Fig 1(b)). These general trends were consistent across the four
 179 other parameterizations (Supplemental).

180 However, despite facilitating pathogen spread, at high pathogen longevities,
 181 high quality patch distributions had a net positive effect on total occupancy re-
 182 lative to low quality distributions (Fig 1(c)). Due to their lower extinction rates
 183 and high connectivity, high quality patches were able to maintain stable occu-
 184 pancy despite the widespread infection facilitated by high pathogen longevity.
 185 On the other hand, though low quality patch distributions inhibited pathogen
 186 spread at intermediate longevities, when high pathogen longevities led to pan-
 187 demic dynamics, low quality distributions were unable to support infectious
 188 occupancy, resulting in an increase in host extinction events and a decrease in
 189 mean total occupancy.

190 The other model parameterizations produced a similar qualitative pattern,
 191 with the differences between total occupancy on high and low quality patches
 192 generally producing a "u" shape as a function of pathogen longevity (Fig 2).
 193 However, the five parameterizations differed in the magnitude and sign of this
 194 general relationship. High environmental stochasticity (i.e. $\alpha = 0$) produced
 195 very similar results to the default model, though with a larger positive effect
 196 of low quality habitat at intermediate longevity (Fig 2(c)). In simulations with

197 reservoir transmission the dominant infection pathway ($\delta = 0.1$), the high qual-
 198 ity patch distributions always produced occupancy higher than (or equal to) the
 199 low quality patch distributions (Fig 2(d)), while in the models with local move-
 200 ment or high disease-induced mortality, low quality patch distributions actually
 201 produced higher occupancies than high quality distributions at high longevities
 202 (Fig 2(b, e)).

203 To understand the mechanisms behind these effects, we further examined
 204 within patch infection dynamics. To compare across the full range of patch
 205 qualities, and to ensure the opportunity for both susceptible and infectious oc-
 206 cupancy, we limit this analysis to high variance metapopulations (with qualities
 207 ranging from 0.25 to 1.75) that produced endemic dynamics. Because of their
 208 lower extinction rate and higher recolonization rate, high quality patches sup-
 209 ported more consistent occupancy than low quality patches (Fig 3). However,
 210 as the epidemic progressed, high quality patches increasingly supported infec-
 211 tious occupants, while low quality patches supported infectious hosts only briefly
 212 and infrequently (Fig 3). In addition, while high quality patches did support
 213 susceptible occupancy, they experienced a greater number of infection events
 214 (susceptible to infectious transitions) than low quality patches (Fig 4). A simi-
 215 lar trend was observed across the other parameterizations, though the observed
 216 relationship between patch quality and number of infection events was shallower
 217 for the lattice and weak direct infection models (Supplemental).

218 4 Discussion

219 This work demonstrates that the distribution of habitat quality in a metapopu-
 220 lation can have substantial impacts on the dynamics of environmentally trans-
 221 mitted pathogens and the resulting patterns of host occupancy. These impacts
 222 are driven largely by the contrasting effects of low and high quality habitat, and
 223 their relative balance in the metapopulation. These contrasting effects derive
 224 from differences in colonization and extinction rates, which in turn drive how
 225 low and high quality patches interact with and modulate pathogen transmission,
 226 either augmenting it in the case of high quality patches, or dampening it in the
 227 case of low quality habitat.

228 When a patch's connectivity is determined by its quality (Eqn 1), high qual-
 229 ity patches function as highly connected hubs in the metapopulation. Attract-
 230 ing colonists helps position high quality patches as infectious "traps", wherein
 231 susceptible hosts repeatedly colonize high quality patches and subsequently be-
 232 come infected (Fig 4). This increased infection pressure results from both trans-
 233 mission from a persistent environmental reservoir and from direct transmission
 234 from infectious immigrants attracted to the high quality patch, reflecting results
 235 from the contact network literature that show that highly connected nodes have
 236 higher infection risk ([14], [15]).

237 The relatively low extinction rate on high quality patches can help to cre-
 238 ate a stable platform from which the pathogen can spread through the rest of
 239 the metapopulation. Indeed, high quality patches are effectively metapopula-

tion scale superspreaders ([16], [9]). Once the pathogen infects high quality patches, the larger number of host colonists produced by these patches allow it to spread to the rest of the metapopulation relatively easily. This process helps to maintain infectious occupancy throughout the metapopulation, which in turn feeds back on high quality patches to ensure a steady stream of infectious colonists that help maintain the environmental reservoir and the trap effect. Thus, through these two interacting mechanisms – the trap and superspreader effects – the presence of high quality patches serves to significantly increase pathogen spread and infectious occupancy in the metapopulation (Fig 1).

In contrast to high quality habitat, low quality patches help to limit pathogen spread and increase susceptible occupancy (Fig 1). Even though individual low quality patches, due to their high extinction rates and low colonization rates, are unable to support consistent occupancy (Fig 3), the presence of low quality patches increases overall susceptible occupancy relative to a lower variance metapopulation (Fig 1). This phenomenon represents the other side of the hub role played by high quality patches in that low quality patches are relatively weakly connected to the metapopulation and are therefore infrequently colonized by infectious individuals. Moreover, the high extinction rate, together with the increased mortality of infectious populations, means that infectious populations do not persist long on low quality patches and the environmental reservoir left behind by these infectious populations may decay before it has the opportunity to infect new susceptible colonists. As a result, low quality patches effectively represent a dead-end for the pathogen, reducing the number of patches from which it can spread, and thereby reducing infectious occupancy (Fig 1).

Though there is a consistent trend in the effect of low and high quality habitat on pathogen spread, the net consequences of these effects for total occupancy vary depending on the structure and dynamics of the metapopulation. Specifically, the findings that low quality habitat inhibits disease spread while high quality habitat enhances it suggest the presence of a trade-off between between processes that facilitate metapopulation stability at the expense of disease spread, and processes that hinder disease spread at the expense of metapopulation stability. This work thus expands on previous studies of disease spread in a metapopulation that focused on movement rate as the driver of this trade-off, with increased movement both improving metapopulation stability and increasing disease spread ([17], [18], [19]). Shifting the patch quality distribution towards high quality patches in our model is in many ways analogous to increasing movement in the models of Hess, Gog, and Park, but by including habitat heterogeneity and metapopulation structure, we are able to investigate the role of individual patches and their contribution to total occupancy.

These contributions, and the relative strength of the above trade-offs, depend largely on the role of connectivity, how it is affected by patch quality, and the resulting strength and structure of the different avenues of pathogen transmission. Broadly, we see three different scenarios that roughly correspond to the three sections of the "u"-shaped relationships in Fig 2: conditions that support very weak pathogen spread, conditions that facilitate intermediate/endemic pathogen

spread, and conditions that produce easy/pandemic pathogen spread. When pathogen spread is weak (e.g. due to low longevity; the top left of the "u"), high quality patches are a net benefit to the metapopulation, as their connectivity and stability facilitate greater occupancy with little additional pathogen spread.

However, in cases where the pathogen is able to spread more easily (i.e. at an endemic level; the trough of the "u"), the addition of high quality patches functions similarly to increasing movement in the model developed by [17]: increasing pathogen spread and as a consequence reducing overall occupancy (Fig 1c). In these cases, the reduction in occupancy from the effect of high quality habitat on pathogen spread outweighs the positive consequences of increased metapopulation connectivity and stability. The shift in this trade-off can arise from conditions that change either side of that balance. High quality habitat distributions require lower pathogen longevities for pathogen spread to begin to take off, relative to low quality habitat distributions. Thus, for intermediate longevities, high quality distributions experience considerably greater pathogen spread and lower occupancy than low quality distributions.

The trade-off can also switch in cases that disrupt the positive benefits of high quality habitat for metapopulation stability, i.e. pathogens with high disease-induced mortality or when dispersal is limited by metapopulation structure. In these cases, rather than serving as sources that increase metapopulation occupancy, high quality patches become sinks. More importantly, they become ecological traps, which can be even more detrimental to the overall population ([20]). From a management perspective, it is critical to understand this shift, as previous theoretical work suggests that management should focus on maintaining patches where conditions are most favorable ([21]). Generally, this means that managers should focus on maintaining high quality habitat, but Strasser et al. (2010) show that stochastic disturbance (e.g. disease-induced mortality) can lead to cases where focusing on low quality patches is more effective in increasing population growth rate. Indeed, in these situations, our model suggests that low quality patch distributions, by hindering the spread of the pathogen and providing refuges for susceptible hosts, can produce higher occupancy than high quality distributions.

As pathogen longevity increases and environmental transmission becomes more important relative to direct transmission (i.e. the top right of the "u") the trade-off shifts back and the addition of high quality patches is again a net benefit to overall occupancy. In these situations, environmental reservoirs overwhelm the susceptible refuges on low quality patches, such that they remain contaminated even across infrequent colonization events. Under these conditions of widespread environmental transmission, high quality patches help to maintain metapopulation stability in the face of disease-induced mortality, thereby increasing total occupancy relative to more balanced or low quality patch distributions.

These dynamics reflect the findings of [18], who found that under strong background infection from an alternative host, increasing movement increased total occupancy, despite facilitating pathogen spread. [19], by adding environ-

mental transmission to the mix, found more generally that in cases where background and environmental transmission were strong relative to direct transmission, increasing movement increased total occupancy monotonically, i.e. movement was a net benefit to the metapopulation. These studies, coupled with the results presented here, suggest that when there is a persistent source of infection throughout the metapopulation (e.g. when the pathogen is long-lived in the environment and direct infection is relatively less important), factors that facilitate metapopulation stability (i.e. increased movement, high quality patches) are a greater benefit than factors that inhibit pathogen spread (i.e. decreased movement, low quality patches).

The different scenarios and parameterizations explored here represent a broad sample of the metapopulation dynamics that we might expect to observe in empirical systems. The consistent "u"-shaped trend in the relative benefit of low and high quality habitat as a function of pathogen longevity suggest that these trade-offs between metapopulation stability and pathogen spread are likely to be fairly general. By evaluating the relative importance of direct versus environmental transmission, along with the effect of habitat quality on connectivity and extinction, we can begin to diagnose where empirical systems (e.g. plague in prairie dogs, or white nose syndrome in bats) fall on this spectrum and the corresponding consequences for management.

Spatial structure and disease spread represent two major factors influencing the dynamics of wildlife populations. As we have seen, in cases where the pathogen is able to persist and remain infectious in the environment, these two factors can interact in complex ways. Failing to consider these interactions and their effect on the roles played by high and low quality habitat risks overlooking the importance of low-quality habitat for the persistence of wildlife populations and the importance of high-quality habitat for the persistence of wildlife disease.

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Figure 1: The effect of the presence of high and low quality patches on (a) median susceptible occupancy, (b) median infectious occupancy, and (c) median total occupancy. Red lines show high quality metapopulations (patch qualities roughly between 0.75 and 1.75), blue lines show low quality metapopulations (patch qualities roughly between 0.25 and 1.25).

Figure 2: Differences between the median total occupancy of high quality patch distributions (quality between 0.75 and 1.75) and low quality patch distributions (quality between 0.25 to 1.25) across the five different model parameterizations. (a) default model, (b) local movement, (c) high environmental stochasticity, (d) dominant reservoir transmission, (e) high disease-induced mortality.

Figure 3: Results from a single representative simulation with variance 0.2 and longevity of 0.3 times the expected residence time of a susceptible population on a unit quality patch. The top panel shows total occupancy of susceptible (blue) and infectious (red) patches over time. The bottom panel shows the state – susceptible (blue), infectious (red), or unoccupied (white) – of individual patches through time. Patches are stacked vertically with the lowest quality (~ 0.25) at the bottom and the highest quality (~ 1.75) at the top.

Figure 4: The number of infection events (S to I transitions) for individual patches in endemic simulations. Light gray lines show smoothed fit for single simulations, while the thick black line shows the smoothed trend over all simulations.

Table 1: State transitions and their rates for patch i in a metapopulation simulation. S denotes occupied by susceptible hosts, I denotes occupied infectious hosts, and \emptyset denotes unoccupied by the host. All patches are characterized by an environmental infection rate, $\gamma(\tau_i)$, where τ_i is the time since the patch was last occupied by infectious individuals.

State Transition	Process	Rate
$S \rightarrow I$	Infection (from contact or reservoir)	$\delta C_{Ii} + \gamma(\tau_i)$
$S \rightarrow \emptyset$	Extinction (of susceptible)	E_{Si}
$I \rightarrow \emptyset$	Extinction (of infectious)	E_{Ii}
$\emptyset \rightarrow S$	Colonization (by susceptibles)	C_{Si}
$\emptyset \rightarrow I$	Colonization (by infectious)	C_{Ii}

Table 2: Parameters of the SPOM model, their meaning, and the values assigned under different parameterizations. Empty cells indicate the same value as the default parameterization.

Parameter	Interpretation	Default	Local move- ment	High env. stoch.	Dominant reservoir trans.	High dis- ease mor- tality
ξ_{im}	Scaling parameter for effect of target patch quality on immigration	0.5	—	—	—	—
ξ_{em}	Scaling parameter for effect of target patch quality on emigration	0.5	—	—	—	—
D	Inverse of mean dispersal distance	5	2	—	—	—
d_{ij}	Distance between patch i and j	1 $\forall i \neq j$	1 $\forall i, j$ neighbors	—	—	—
μ_S	Extinction rate of unit quality susceptible patch	0.1	—	—	—	—
μ_I	Extinction rate of unity quality infectious patch	0.5	—	—	—	1
α	Strength of environmental stochasticity	1	—	0	—	—
δ	Probability of direct infection	0.5	—	—	0.1	—
γ_0	Initial rate of infection from reservoir patch	0.5	—	—	—	—