

Trophic transfer of pesticides: the fine line between predator-prey regulation and pesticide-pest regulation

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Running Title Sensitive line of predator/pesticide pest-regulation

Abstract

1. Understanding pesticide impacts on populations of target/non-target species and communities is a challenge to applied ecology. When predators that otherwise regulate pest densities ingest prey contaminated with pesticides, this can suppress predator populations by secondary poisoning. It is, however, unknown how species relationships and protocols of treatments (e.g. anticoagulant rodenticide (AR)) interact to affect pest regulation.
2. To tackle this issue, we modelled a heuristic non-spatialized system including montane water voles, specialist vole predators (stoats, weasels), and a generalist predator (red fox) which consume voles, mustelids and other prey. By carrying out a broad-range sensitivity analysis on poorly known toxicological parameters, we explored the impact of 5 farmer functional responses (defined by both AR quantity and threshold vole density above which AR spreading is prohibited) on predator-prey interactions, AR transfer across the trophic chain and population effects.
3. Spreading AR to maintain low vole densities suppressed mustelid and fox populations, leading to vole population dynamics being entirely regulated by AR use. Such vole-suppression treatment regimes inhibited predation ecosystem services and promoted pesticide dependence.

34 4. Keeping vole density below acceptable bounds by spreading AR while maintaining suffi-
35 cient voles as prey resources led to less AR being applied and extended periods without AR
36 in the environment, benefiting predators while avoiding episodes with high vole density.
37 This may meet farm production interests while minimizing the impact on mustelid and
38 fox populations and associated ecosystem processes. These alternating phases of mustelids
39 and farmer regulation highlight the consequence of intraguild relationship where mustelids
40 may rescue foxes from poisoning. Both global and wide-range sensitivity analysis illustrate
41 the tightrope between predator-prey regulation and pesticide-pest regulation.

42 5. *Synthesis and applications* Different pesticide protocols lead to a rich variety of predator-
43 prey dynamics in agro-ecosystems. Our model reveals the need to maintain refuges with
44 sufficient non-poisoned voles for sustaining specialist mustelids, to conserve the predator
45 community given the potential of secondary poisoning with rodenticides. We suggest that
46 long periods without pesticide treatment are essential to maintain predator populations,
47 and that practices of pesticides use that attempt to permanently suppress a pest over a
48 large scale are counterproductive.

49 **Keywords** biodiversity conservation; secondary poisoning; cyclic fluctuations; pesticides; cascade
50 effects; ecosystem service; sensitivity analysis ; ecological control

51 1 Introduction

52 Since the “green revolution” following the 1950s, pesticides use has increased to control pests
53 damaging properties, public health or crops (Tilman et al., 2002). Pesticides usage is varyingly
54 triggered by the perception/estimation of pest densities. Natural enemies (e.g., predators, para-
55 sites, competitors) also reduce pest densities and hence may preclude the need for using pesticides
56 (Michalko and Pekár, 2017). Natural enemies are, however, also affected by pesticides, either
57 by direct exposure, through ingestion of contaminated prey (Berny, 2007) or indirectly through
58 cascading consequences of effects of resource depletion (Halstead et al., 2014). Thus, under some
59 regimes of pesticides use, pest populations only become regulated by pesticides once predators have
60 collapsed. To preserve ecosystem health and the services that predators provide through regulation
61 of pest densities, we need to assess the feasibility and benefit of pesticide treatment regimes in their
62 ability to control pest species with minimal damage to predators (Halstead et al., 2014). It is how-
63 ever empirically challenging to assess the overall impact of pesticide treatments on the dynamics of
64 species linked by trophic interactions. In this context, process-based models describing simplified
65 scenarios are powerful tools to reveal hidden patterns by disentangling processes emerging from
66 pesticide impacts on predator-prey systems (e.g., Baudrot et al. (2018)).

67 Voles and other grassland rodent species undergo multi-annual population cycles (e.g., Krebs
68 (2013)). At their peaks, they may attain extremely high densities, causing substantial damage
69 to grass/cultivated crops and forestry and conflicts with humans (Delattre and Giraudoux, 2009).
70 Farmers worldwide expand financial resources to purchase and spread anticoagulant rodenticides

71 (hereafter AR), hoping to reduce vole populations and damages, and increase profits despite the
72 investment required (Stenseth et al., 2003). They do so according to protocols, equivalent to farmer
73 functional responses (hereafter FFR), that involve varying amounts of AR spread in response to
74 different thresholds in vole densities.

75 Voles and many small rodents are perceived as pests, but they are also keystone species, crucial
76 to the functioning of grassland and forest ecosystems, as well as being the prey of numerous preda-
77 tors, including species of conservation concern (Delibes-Mateos et al., 2011; Coeurdassier et al.,
78 2014). Their population cycles create pulses of resources crucial to the viability of a wide range of
79 resident predators and the aggregation of mobile avian vole predators (Korpimaki and Norrdahl,
80 1991). The smallest mustelids (e.g. weasels *Mustela nivalis*) are specialist vole predators. Their
81 numerical response has been shown theoretically to be necessary for generating predator-prey cy-
82 cles (Hanski et al., 1991). They are said to be responsible for driving 3-to-5-year vole cycles in
83 Fennoscandia (the specialist predation hypothesis) (Hanski et al., 1991). Generalist resident preda-
84 tors like foxes (*Vulpes vulpes*) are expected to have regulatory and limiting effects on voles, owing
85 to dietary plasticity that slows down vole population increase at low density (Hanski et al., 1991).
86 Foxes do not show numerical responses to vole abundance (Weber et al., 2002) but they influence
87 the food chain through occasional killing and consumption of bite-sized mustelids. Mustelids form
88 a small proportion of fox diet (0-10%) but their offtake could represent a significant portion of the
89 population (reviewed in Lambin (2018)).

90 Anticoagulant rodenticides are non-selective toxicants with deleterious effects on non-target
91 fauna (e.g. Coeurdassier et al. (2014)). Despite AR being exclusively licensed for rodent control,
92 a large number of predator species are secondarily exposed to AR (Sánchez-Barbudo et al., 2012).
93 Consumption of dead and sub-lethally intoxicated voles reduced fox abundance in farmland in east-
94 ern France (Jacquot et al., 2013) and ARs caused short-term declines in stoats in New Zealand
95 (Alterio, 1996). Rodent-eating mustelid populations are affected by ARs given the pervasive levels
96 of contamination reported (McDonald et al., 1998). Thus, there is little doubt ARs use inadvertently
97 depresses predator populations. As predators likely limit vole populations, it is essential to under-
98 stand when AR use becomes counterproductive by altering the pest population dynamics, producing
99 more frequent outbreaks and high residual vole abundance.

100 With the aim of understanding the potentially complex interactions between prey that are per-
101 ceived as pest, predators and farmers spreading rodenticide in response to vole abundance and their
102 functional responses, we studied a simplified system inspired by cyclically fluctuating montane wa-
103 ter voles (*Arvicola scherman*), small mustelids (stoats, weasels) that mostly eat voles (specialists),
104 and foxes (generalists), with voles and mustelids as food items. We used a process-based model
105 using differential equations to explore 5 FFR types of AR spread, combining population dynamics,
106 predator-prey interactions and rodenticide transfer across the trophic chain. Model parameters and
107 FFR were inspired by farming systems in the Jura Mountains, Franche-Comté (France), the region
108 of Comté cheese production. In Franche-Comté, farmers shifted from polyculture to almost exclu-

109 sively grass production for milk used to produce cheese from the early 1970s (Giraudoux et al.,
 110 1997). Due to recurrent vole outbreaks and damages to grasslands, massive rodenticide treatments
 111 were implemented from the early 80s with consequences on non-target wildlife. Practices were de-
 112 veloped technically under pressure from public opinion, farmer unions and farmer technical orga-
 113 nizations collaborating with researchers to find treatment regimes with less harmful consequences
 114 for biodiversity (Delattre and Giraudoux, 2009).

115 Hence, our main objective was to explore the properties of FFR in relation to varying population
 116 sensitivity to AR on the global tri-trophic population dynamics made of vole outbreak frequencies
 117 and guild of interacting predators.

118 2 Materials and Methods

119 We specified and parameterized the tri-trophic system of voles-mustelids-foxes with use of AR by
 120 farmers in response to vole density. We considered several FFR depending on the AR amount and
 121 vole threshold triggering AR spreading to assess AR upward transfer through the trophic web. In
 122 all cases, parameterization units are on hectare⁻¹ and day⁻¹.

123 2.1 Model for the tri-trophic dynamic

124 We considered a tri-trophic system described by Figure 1 and equations (1-5); parameterization is
 125 provided in Table 1. Voles, denoted V , were the primary prey, Mustelids, M , intermediate preda-
 126 tors, and Foxes, F , were top predators, consuming voles and mustelids. For each species, the
 127 instantaneous variation of population size over time is:

$$\begin{cases} \frac{dV}{dt} &= Vr_V \left(1 - \frac{V}{K_V}\right) - \Phi_{V,M}(V)M - \Phi_{V,F}(V, M)F \\ \frac{dM}{dt} &= \varepsilon_M \Phi_{V,M}(V)M - m_M M - \Phi_{M,F}(V, M)F \\ \frac{dF}{dt} &= Fr_F \left(1 - \frac{F}{K_F}\right) \end{cases} \quad (1)$$

128 The vole population followed a logistic growth rate, with r_V the maximal reproduction rate,
 129 fixed at $r_V = \ln(2 \times 600)/365$ per day, since montane water vole populations can increase from 0 to
 130 600 individuals ha⁻¹ or more (Giraudoux et al., 1997), resulting in the equilibrium density being
 131 fixed at $K_V = 600$ individuals. The vole population was preyed upon by mustelid and fox popula-
 132 tions. The vole consumption rate at different vole densities was described by functional responses
 133 ($\Phi_{V,M}$ for mustelids, $\Phi_{V,F}$ for foxes), see equation (2). We assumed small mustelids behave as spe-
 134 cialist predators (King and Powell, 2006), we considered a numerical response linearly dependent
 135 on the functional response with parameter ε_M (dimensionless) as conversion efficiency of prey into
 136 newborn predator (see Supporting Information - Appendix S1). The mustelid background mortality
 137 rate (i.e. all other reasons of death: ageing, disease, etc.) was m_M parameterized has the inverse

138 of life expectancy (Table 1). We assumed foxes had a logistic growth rate function, parameterized
 139 with maximal growth rate $r_F = \ln(3)/365$ and equilibrium density $K_F = 0.03$ individuals ha^{-1}
 140 (Ruelle et al., 2003) without numerical response (Weber et al., 2002).

141 Since we treated small mustelids as vole specialist predators, we assumed a Holling Type II
 142 functional response with attack rate a_M and handling time h_M (equation (2)). We then represented
 143 foxes feeding on voles and mustelids by a multi-species functional response derived from Holling
 144 Type III, referring to generalist feeding behaviour (Baudrot et al., 2016). For that, we denoted
 145 a_{VF} and a_{MF} the fox attack rate on voles and mustelids respectively. The parameter h_F was the
 146 handling time for foxes.

$$\begin{aligned}\Phi_{V,M}(V)M &= \frac{a_M V}{1 + h_M a_M V} \\ \Phi_{V,F} &= \frac{a_{VF} V}{a_{VF} V + a_{MF} M} \times \frac{(a_{VF} V + a_{MF} M)^2}{1 + h_F (a_{VF} V + a_{MF} M)^2} \\ \Phi_{M,F}(V, M)F &= \frac{a_{MF} V}{a_{VF} V + a_{MF} M} \times \frac{(a_{VF} V + a_{MF} M)^2}{1 + h_F (a_{VF} V + a_{MF} M)^2}\end{aligned}\quad (2)$$

147 Parameterization of functional responses was estimated to fit the daily satiation level of preda-
 148 tors for handling times, and the observed 5-6 year vole cycles for attack rates (Table 1). We assumed
 149 foxes spent longer searching for voles than mustelids, based on each species diet and daily number
 150 of individuals captured. Therefore, a_{VF} was considered larger than a_{MF} and selected to produce
 151 6-year vole cycles without AR.

152 2.2 Model with rodenticide

153 Figure 1 represents the whole study system. Rodenticide is spread in grasslands during treatments,
 154 denoted $T_{Broma}(V)$, triggered by vole density V . Firstly, baits (50 mg kg^{-1} of bromadiolone, here-
 155 after AR) were spread in grasslands at quantity 7.5 to 20 $\text{kg ha}^{-1} \text{day}^{-1}$. Notation day^{-1} stands
 156 because of the daily time resolution. Such quantity, C , was available for voles, and a proportion
 157 disappeared in the environment at rate k_0 (set at $k_0 = 0.0815$) (Sage et al., 2008). The proportion
 158 consumed per vole, with rate $\kappa(C)$, was assumed to be an increasing function. The function $\kappa(C)$
 159 was characterized by a maximum ingestion M_{in} , and a half-saturation constant for ingestion D_{in} in
 160 [mg kg^{-1}]:

$$\kappa(C) = \frac{M_{in} \times C}{D_{in} + C} \quad (3)$$

161 For the toxicokinetics of AR (i.e. internal compound dynamics) leading to AR concentrations
 162 in animal body (voles, mustelids and foxes), we considered an uptake without biotransformation
 163 and time-regulated distribution, (i.e. AR concentration in the body of animals was instantly ho-
 164 mogeneous) and that the whole body was consumed or scavenged without selection/rejection of
 165 tissues-organs. We also assumed disappearance including excretion of the parent compound and
 166 metabolisation, and that metabolites were non-toxic and/or excreted in the scats. For the toxi-

167 cokinetics of AR ingested by voles, a fraction C_V was assumed to remain active and available to
 168 predators ingesting voles. The absorption rate of ARs (η) exceeds 50% in less than 24h (Jacquot
 169 et al., 2013). The excretion rate from voles, $k_{out,V}$ was 0.4 day^{-1} (Sage et al., 2008). The mortality
 170 rate through ARs was $\mu(D_V)$. Death through poisoning created a dead vole population (V_d) with AR
 171 concentration C_V . Dead voles could either be scavenged by mustelids/foxes or decompose at rate
 172 d . We assumed AR in dead voles disappeared from the system when voles decomposed (assumed
 173 in one week: $d = 1/7 \text{ day}^{-1}$). Mustelids could feed on live voles V , or non-decomposed dead voles
 174 V_d and we assumed a Type II functional response adapted for a multi-species functional response
 175 (Baudrot et al., 2016). Mustelids ingested AR with absorption rate η_M (ratio between biomasses of
 176 voles, B_V , and mustelids, B_M) and the total of ingested voles (alive V and dead V_d) was defined
 177 by function $\Theta_M(V, V_d)$ (Table 1).

178 A fraction of AR ingested was accumulated in weasels while the rest was excreted with rate
 179 $k_{out,M}$. AR contaminating weasel through vole poisoning, is denoted D_M , induces lethal effect at
 180 rate $\mu_M(D_M)$, additive to natural mortality rate m_M . Parameter definitions are further detailed
 181 in Supporting Information (Appendix S1). AR was ingested by foxes with a rate proportional to
 182 the functional response of foxes to voles, dead voles and mustelids. Foxes also accumulated AR
 183 available in their prey, resulting in upward AR transfer in the trophic chain. Foxes accumulated AR
 184 in concentration C_F . A fraction of AR was excreted by foxes at rate $k_{out,F}$ at a rate between 0.38
 185 and 0.72 day^{-1} (Sage et al., 2010), and AR caused fox mortality at rate $\mu_F(D_F)$.

186 We used log-logistic equations for describing dose-dependent mortality of animals exposed to
 187 AR. Vole and predator mortality rates due to AR $\mu_X(D_X)$ (X referring to the considered species)
 188 were expressed by equation (5):

$$\mu_X(\zeta_X) = 1 - \frac{1}{1 + (LD_{50}/\zeta_X)^H} \quad (4)$$

189 The parameter LD_{50} is the daily median lethal dose (50% of population dying), parameter H
 190 is the Hill's coefficient modulating the curve steepness (see Table 1). Parameterization of LD_{50} is
 191 important since it defines the inflexion point of the ecotoxicological effect. While it is a classical
 192 parameter targeted in experiments, empirical estimates of its value are highly uncertain and indeed
 193 span different order of magnitude for the same species and under the same experimental design
 194 (Grolleau et al., 1989; Erickson and Urban, 2004; Sage et al., 2010; Karmaus et al., 2018). We
 195 therefore carefully explored this parameterization through a wide-range sensitivity analysis detailed
 196 here-after (see also Figure 2).

197 2.3 The farmer functional responses (FFRs) explored through simulation

198 We considered a range of realistic FFRs spanning treatments during vole outbreaks only and a
 199 precautionary approach in which treatments only takes place at intermediate or low vole density
 200 threshold. These scenarios are inspired by historic and contemporary protocols of bromadiolone

201 use to control montane water voles in Franche-Comté, but also representative of practice globally
202 (Delattre and Giraudoux, 2009): (A) a scenario without AR treatment, (B) a scenario with high
203 vole density threshold triggering treatment (500 ind ha^{-1}) and high AR amount per treatment (20
204 kg ha^{-1}), (C) a third scenario with intermediate triggering threshold (250 ind ha^{-1}) and high AR
205 per treatment (20 kg ha^{-1}), (D) scenario with intermediate threshold (250 ind ha^{-1}) and low AR
206 amount (7.5 kg ha^{-1}) and (E) scenario with low threshold (50 ind ha^{-1}) and low treatment (7.5
207 kg ha^{-1}). To check the influence of predators such as foxes and intraguild predation on the system
208 dynamics, we also simulated scenarios with and without foxes. Our simulations tracked the linked
209 vole-mustelid-fox dynamics for 40 years, after a “burn-in” period of 10 years to reduce dependency
210 of results upon initial conditions, to observe several vole cycles and to characterise AR effects on
211 these species population dynamics. This burn-in period also had AR treatment triggered at specific
212 vole densities and with a given rodenticide quantity for each FFR. The burn-in phase was selected
213 according to a set of simulations with different initial conditions. Those simulations showed that in
214 a given FFR (i.e., same threshold of vole density and amount of AR spread), the dynamics of the
215 population were converging toward a similar pattern.

216 2.4 Numerical Simulation and Sensitivity Analysis

217 All numerical analysis have been done using the open language R and particularly the ODE solver
218 package *deSolve* (Soetaert et al., 2010). Model implementation and code to run analysis are avail-
219 able on a Github repository (Baudrot et al., 2020).

220 We applied a wide-range sensitivity analysis on lethal dose parameters (i.e. LD_{50}) for voles,
221 mustelids and foxes, to explore changes in dynamical patterns. While wide-range variations of each
222 parameter is likely to impact the global pattern, the lethal dose parameter is a direct characteristic
223 of the toxicological part of the model. The variability of LD_{50} from *in vivo* acute systemic toxicity
224 studies under same experimental design spans 2 to 3 order of magnitude (see rat oral acute toxicity
225 in Karmaus et al. (2018)), which makes it challenging to assess risk with exposure models. Since
226 the LD_{50} parameter of fox appears to have low impact on dynamical pattern in our model, we
227 evaluated only 4 different values (i.e. 0.5, 1.5, 2 and 7 ppm) which are consistent with a large
228 range covering such a parameter (Erickson and Urban, 2004; Sage et al., 2010). For the prey and
229 the specialist predator, this parameter was much more sensitive, so we explored 10 values for both,
230 respectively in the range $[0.7 - 7]$ ppm and $[1 - 25]$ ppm following consistent ranges (Grolleau
231 et al., 1989; Erickson and Urban, 2004). We classified the subsequent 1600 simulations according
232 to the dynamical pattern of mustelids population: either "periodic regulation by mustelid" or "AR
233 permanent regulation" where mustelids collapse.

234 Then, we performed a local sensitivity analysis to study changes in model outputs apportioned
235 to small variation around model parameters input (Saltelli et al., 2019). We applied a first-order
236 Sobol’s sensitivity index S_i defined as $S_i = \mathbb{V}(\mathbb{E}_{-i}(y|x_i))/\mathbb{V}(y)$, where $\mathbb{V}(y)$ is the variance of y when
237 all factors are allowed to vary, $\mathbb{E}_{x \sim i}(y|x_i)$ the mean of y when one factor is fixed (Sobol and Saltelli,

238 1993). We defined the domain of variation with a beta distribution within an hyperspace of plus
239 or minus 10% from the original value for all parameters. For output variable of local sensitivity
240 analysis, we estimated the following cost functions: (i) Number of treatment events per FFR; (ii)
241 Cumulative amount of AR (kg); (iii) Proportion of time when the AR-induced mortality of mustelids
242 higher than 50% (i.e. lethal exposure profile killing 50% of mustelid population); (iv) Proportion of
243 time when the mortality of mustelids was higher than 50% due to natural mortality (see Supporting
244 Information - DATA available); (v) Proportion of time when the vole density was below 50 voles
245 ha^{-1} , as a proxy for time when forage grass grows with low herbivore influence; (vi) Mean vole,
246 mustelid and fox densities.

247 **3 Results**

248 Allowing for mortality by predators ingesting AR-poisoned voles changed the outcome of predator-
249 prey dynamics involving vole, mustelid and fox populations. Secondary poisoning led to a rich
250 spectrum of emergent dynamics according to the FFR to vole abundance. Without AR (scenario
251 A), vole dynamics were regulated by mustelid predation that gave rise to a 6-year cycles (Figure 3)
252 typical of prey-specialist dynamics, while the generalist fox population remained at its carrying
253 capacity (i.e., 0.03 ind ha^{-1} , see Table 1). Based on this null model, we explored 4 scenarios of
254 treatment described earlier and denoted respectively B, C, D and E, defined by the vole density
255 threshold triggering treatment (respectively: 500, 250, 250 and 50 ind ha^{-1}) and the AR amount
256 per treatment (respectively: 20, 20, 7.5 and 7.5 kg ha^{-1}).

257 **3.1 Sensitivity analysis of model parameters**

258 The exploration of ecotoxicological uncertainties on population dynamics was performed for the
259 lethal dose parameters LD_{50} for voles, mustelids and foxes (see Figure 2). From the 1600 pat-
260 terns explored two typical patterns emerged. On the one hand (i) vole dynamics were sequentially
261 regulated by either AR treatments, which we refer to as farmer-regulated phase, or by mustelids,
262 mustelids-regulated phase (see in Figure 3 panel c. B-2). This succession of farmer-regulated and
263 mustelids-regulated phases are denoted "Mustelid periodic regulation" in Figure 2 (black pixels).
264 On the other hand, (ii) vole dynamics were permanently regulated by AR treatments, because of
265 crashes of the mustelid population we refer to as "Permanent AR regulation" in Figure 2 (grey pix-
266 els). We see that LD_{50} for foxes does not change the general pattern of dynamics. Both scenario B
267 and D exhibited the highest fractions of "Mustelid periodic regulation" compared to scenarios C and
268 E. This illustrates the tight line that exists between resource and AR impact to benefit from predator
269 regulation periods.

270 For other parameters, we performed a local sensitivity analysis where we computed a first order
271 sensitivity index (Sobol and Saltelli, 1993; Saltelli et al., 2019) given the contribution of each
272 parameter to the variance (in Figure 5 for scenario D - intermediate amount of 7.5 kg AR at threshold

273 density of 250 voles). Parameter values were moved within +/- 10% interval around the fixed
274 value as defined in scenario D (see Table 1). Crossing all cost functions to test the sensitivity of
275 model parameters (Figure 5), the influence of each parameter was quite homogeneous. There was
276 no redundant or conversely any parameters with a dominant influence on model behaviour. The
277 homogeneity of the sensitivity to parameters was particularly true for densities of predators to which
278 we focused our attention.

279 3.2 Population dynamics

280 While both "Mustelid periodic regulation" and "Permanent AR regulation" pattern occurred under
281 the 4 scenarios with AR (B, C, D and E) (see Figure 2), we illustrate in Figure 3 the most contrasting
282 population dynamics for each of scenario in order to better describe their underlying mechanism.
283 The boundary between contrasting dynamics lies for vole LD_{50} around 1.5 to 2 ppm and in the
284 range 2 to 2.25 ppm for mustelid LD_{50} .

285 From Figure 3, we see that scenario B (high vole density threshold triggering treatment and
286 high AR amount per treatment) and scenario D (intermediate threshold and low AR amount) ac-
287 counts for the most of dynamics with sequential farmer-regulated phase and mustelids-regulated
288 phase. Farmer-regulated periods started when densities of living voles triggered treatments. This
289 produced sudden declines of live voles followed by increases of dead voles. However, the vole pop-
290 ulation recovered quickly which triggered repeated further treatments and pulses of availability of
291 contaminated (both live and dead) voles. Also, in scenario D, vole declines were not as deep as
292 when pulses of AR amount were high like in scenario B (see Supporting Information), owing to the
293 reduced AR amount per treatment. Mustelids and sometimes foxes also experienced AR-induced
294 declines during this period (Figure 3-c,f,g,h and Figure 4). For the predators, mustelid-regulated
295 periods started when mustelid numbers grew slowly to a peak, which depressed vole density, pre-
296 cluding AR treatments and releasing the fox population from secondary poisoning, such that its
297 abundance rebounded. Vole depletion by mustelids and subsequent mustelid declines allowed the
298 vole population to grow again up to threshold densities that initiated a new period of regulation by
299 farmers.

300 For scenario C (intermediate threshold and high AR), vole dynamics were mostly regulated by
301 AR treatment only (e.g. Figure 3-e). Populations of live and dead voles experienced high frequency
302 fluctuations driven by AR. As AR treatments were frequent, being triggered by vole peaks, contami-
303 nated dead voles were always abundant (see Supporting Information: peaks at 90 ind ha^{-1}). With
304 scenario E (low threshold, low AR) vole populations were maintained by farmers at around 50 voles
305 ha^{-1} (See Figure 3 and Supporting Information). The population of live voles was regulated by
306 treatments. For most of simulations (except when vole LD_{50} with high, Figure 2), whenever voles
307 reached densities triggering treatment, predator populations experienced strong declines. However,
308 fox densities were higher compared to scenario C (intermediate threshold, high AR) (see Support-
309 ing Information), reflecting the reduced amount of AR used (7.5 kg) and transferred to foxes as

310 there were lower vole densities. The maximum numbers of dead voles under this scenario E was
311 relatively low (highest around 15 ind ha⁻¹) but, due to frequent treatments, there was a steady
312 replenishment of contaminated dead voles. This, in turn, induced mustelid and fox mortality and
313 population declines (Supporting Information). Additionally, low availability of live voles triggered
314 small mustelids mortality through starvation, down to abundances similar to those resulting from
315 AR use (Supporting Information).

316 **3.3 Influence of intra-guild predation**

317 Figure 4 shows the system dynamics under a scenario D, where successive farmer-regulated and
318 mustelids-regulated phases occurred. This simulation shows that the removal of foxes did not elim-
319 inate the successions of mustelids-regulated and farmer-regulated phases. However, the mustelid
320 regulated period allowed short-term peaks of voles, suggesting the emergence of a classical one-
321 predator - one-prey cycles interrupted by a farmer-regulated period. Without foxes, population
322 dynamics of mustelids presented a more chaotic behaviour, while it presented regular cyclic pattern
323 with fox occurrence. Therefore, this simple model suggests a stabilizing role of a generalist predator
324 (foxes in this system) during the mustelid-regulated period. At the end of the mustelid-regulated
325 period, foxes strongly contributed to vole mortality and, to a lesser degree, mustelid mortality. Con-
326 sequently, the removal of foxes implied less predation on voles during the mustelid regulated period,
327 and short-term vole releases from mustelid predation. The 2-year rolling mean of vole density (blue
328 lines in Figure 4) illustrates the change of regime from farmer-regulated to mustelids-regulated
329 alternating phases. Indeed, the amplitude of averaged vole densities (i.e., the amplitude of vole cy-
330 cles for the 2-year rolling mean) was relatively stable at the beginning of farmer regulation periods
331 and then suddenly decreased to become minimal before sharply increasing, announcing a regime
332 change. These changes in density amplitude may be used as an early-warning signal of the regime
333 transition.

334 **4 Discussion**

335 Considering that rodenticide kills not only voles but also their predators through secondary poison-
336 ing, our models show that AR profoundly changes the modeled outcome of predator-prey dynamics
337 involving vole, mustelid and fox populations beyond what mere intuition could elucidate. Our study
338 reveals how the dual influences of the amount of pesticide spread and the vole density threshold
339 triggering AR spread drive (i) pesticide spreading frequency, (ii) predation ecosystem service, and
340 subsequently (iii) the control of pest outbreaks. Two types of a rich spectrum of emergent dynam-
341 ics, including farmer or mustelid regulation changing from classical predator prey dynamics arose
342 because poisoned voles acted as global stressor on the food chain.

343 **4.1 Modelling farmer regulation into a classical predator-prey system**

344 The threshold functional response of farmers deciding when to apply varying amounts of rodenti-
345 cides according to prevailing vole density was crucial in selecting the emergent ecosystem dynamics,
346 resulting in much variability in ecosystem and conservation and farming production interests. In
347 the idealised ecosystem our models depict, farmers spreading rodenticide not only depleted vole
348 prey exploited by specialist and generalist predators but also created pulses of lethally or sub-
349 lethally poisoned voles that subsequently poison their predators. Arguably this set of ecological
350 interactions has similarities with circumstances where a pathogen affecting prey species also in-
351 fects predators, as in the case with the flea vectored plague (*Yersinia pestis*) infecting prairie dogs
352 (*Cynomys* spp.) and black footed ferrets (*Mustela nigripes*) in central US (Matchett et al., 2010).
353 However, to our knowledge, the behaviour of such tri-trophic model with multiple reciprocal in-
354 teractions has not been explored. This is despite the obvious relevance to the management of the
355 globally widespread circumstances where keystone small mammals are poisoned and may second-
356 darily poison their predators (Delibes-Mateos et al., 2011).

357 Under the "reference" scenario without AR spreading (A), we assumed a predator-prey cycle
358 which is a plausible pattern thoroughly explored theoretically (Hanski et al., 2001), though with
359 debated empirical support (Lambin, 2018). There is no controversy on the role of small mustelids
360 tracking vole dynamics, though it is not yet well understood whether there is sufficient lag between
361 predators and prey for predation to drive steep declines (King and Powell, 2006). Parameters of
362 the reference scenario for our predator-prey model were biologically realistic and tuned to generate
363 population fluctuations similar to those observed in the studied cyclic system (Delattre and Girau-
364 doux, 2009). The addition of pulses of rodenticide and their toxicokinetics in vole and predators
365 are based on previous experiments with bromadiolone, a widely used AR, ensuring biologically
366 realistic functional forms and their parameterization. Irrespective of the FFR considered, the fre-
367 quency of vole cycles dramatically increased compared to the reference scenario, except during
368 mustelid-regulated phases emerging under some FFR scenarios.

369 **4.2 How specialist predators may protect generalists from poisoning**

370 An interesting model behaviour with farmer- and mustelid-regulated phases alternating with low
371 frequency was seen with scenarios B (high vole threshold, high AR) and D (intermediate vole thresh-
372 old, intermediate AR), and to a lesser extent scenario C (intermediate vole threshold, high AR) and
373 rarely with scenario E (lowest vole density threshold). Such flipping between alternative states in
374 population dynamics has been previously described in predator-prey model where weasels rely on
375 a primary prey and entrain the dynamics of secondary prey (Hanski and Henttonen, 1996) but not
376 for the kind of indirect interaction we explore here. It further demonstrates that adding biologically
377 realistic complexity to simple models may drastically change the emergent properties of trophic
378 interactions. From these scenarios, we understand that the emergence of successive farmer- and
379 mustelid-regulated phases is neither driven by vole density threshold alone nor by AR amount, but

380 instead by a subtle combination of both. Also, under scenarios with low LD_{50} for fox, the mod-
381 elling description of these patterns uncovered the dual key roles of mustelids on fox dynamics, as
382 intraguild competitors and as a vector for poisoning. This led to a surprising form of facilitation
383 for foxes: mustelids protect foxes from collapses. The establishment of such a response can be
384 described in 3 steps. Firstly, low mustelid densities inhibit their regulation of voles and contribute
385 to farmer AR use. In line with empirical evidence, the latter directly impacts foxes by poisoning
386 (Jacquot et al., 2013). Secondly, fox predation on mustelids is reduced, and with an intermediate
387 AR amount, this allows mustelids to slowly recover. Vole outbreaks and subsequently farmer AR
388 treatments are then gradually delayed, benefitting mustelids recovery. This is the point of transi-
389 tion from farmer to mustelids regulation regimes, starting the third step: mustelids increase faster,
390 suppressing vole densities and precluding the need for AR treatments, and eventually indirectly
391 allowing the fox population growth.

392 Our findings that complexities in trophic interaction, induced by the poisoning of predator by
393 poisoned prey, may cause the system to flip between alternative states is novel and robust. However,
394 given we only explored deterministic and spatially homogeneous versions of our model, any infer-
395 ence on the frequency of flipping between states should be cautious given the inherent stochastic
396 nature of natural and farmland environments. If such dynamics occur within real farming sys-
397 tems, flipping between states is unlikely to emerge with regularity where many other factors impact
398 population dynamics. While generalists are known to have stabilising effect (Hanski et al., 1991),
399 the benefit of specialist predators imparted to generalist predator and resulting increase in the
400 prevalence of intraguild predation would be difficult to detect in empirical studies. Nevertheless,
401 other generalist predators such as the endangered red kite (*Milvus milvus*) which feed on voles op-
402 portunistically, occupy areas with bromadiolone treatments and are also affected by rodenticides
403 (Coourdassier et al., 2014) and may therefore also benefit from the presence of mustelids in the
404 ecosystem.

405 **4.3 How region-wide vole suppression may inhibit ecosystem services**

406 In many situations, notably scenario E and C and when vole and mustelids are highly sensitive
407 to AR, the whole system was solely driven by farmer regulation, whereby the chronic use of AR
408 completely suppressed the pest-regulation ecosystem service of predators. It has previously been
409 shown empirically that repeated rodenticide treatments are highly detrimental to the populations of
410 predators and reduce their densities (Jacquot et al., 2013). Secondary poisoning of predators is an
411 established reality (Berny, 2007). Through modelling, we formalised the insight that some poison
412 deployment protocols, including those presently used in the empirical system which motivated our
413 study, are counterproductive if employed on a large scale, suppressing natural predator regulation
414 of pest rodents. It has been long known that poisoning rodents with AR permeates the food chain
415 at peak abundance, achieves little in terms of protecting crops and may have strong deleterious
416 impact (Olea et al., 2009). In Franche-Comté, a change in treatment protocols, from controlling

417 voles at high densities to low-intermediate densities, has reduced the mortality of non-target species
418 (including foxes) (Jacquot et al., 2013). Nevertheless, deployment regimes of pesticides that can
419 contaminate the food chain should also include periods of time which permit predator populations
420 to rebound and avoid extirpation from the ecosystem. We have shown that, over time, farmers who
421 strictly maintain voles at low density thresholds would likely suppress predation services provided
422 by vole predators and, in so doing, instigate pesticide dependence. In addition, small mammals
423 like voles certainly have ecosystem functions. Our results also suggest that the presence of small
424 mustelids in ecosystems is beneficial for biodiversity conservation (see above) and agriculture inter-
425 ests. Given the importance of vole cycles and their trophic interactions, it is desirable to maintain
426 vole population fluctuations of sufficient amplitude to maintain ecosystem processes.

427 **4.4 Managing rodents and ecosystems**

428 Presently, in Franche-Comté, farmers relying on bromadiolone alone can only treat pre-emptively
429 when voles are at low densities (scenario E) whereas those also using alternative methods (i.e.,
430 mechanical, not pesticide-based) are allowed to spread AR in low quantity up to intermediate vole
431 densities (scenario D). Spreading AR in low quantity seems superficially desirable, but our heuris-
432 tic model, assuming an idealised homogenous landscape, shows this is associated with frequent
433 treatments. Consequently, it would induce a near permanent availability of a small number of in-
434 toxicated voles which, combined with low availability of non-contaminated voles, would reduce
435 predator populations. Therefore, the extreme situation of using a low vole density threshold (sce-
436 nario E) at a large scale is undesirable because it depletes the prey resources of foxes and mustelids
437 and their populations. Triggering treatment at intermediate vole density with a low amount of AR
438 (scenario D) allows for temporal refuges, i.e. longer periods free of rodenticide necessary for preda-
439 tor densities to rebound while simultaneously avoiding episodes with high vole density, as required
440 by farm production interests. Under a landscape management approach, such temporal refuges
441 could arguably be substituted for by spatial refuges, with parts of the landscape free of pesticides
442 and fuel the recolonisation of the farming landscape by predator populations.

443 Our key result and the basis for management prescriptions is that allowing for refuges where
444 voles are not poisoned and allowed to persist at medium-high densities such that they can be
445 exploited by mustelids is crucial for predator population recovery and preserving the ecosystem
446 services mustelids deliver. Treatment regimes allowing temporal and/or spatial refuges seems com-
447 patible with both conservation and farming interests. A critical insight is to avoid potential side
448 effects of chronic low-dose AR prescription (e.g., depletion of community services, stimulation of
449 resistances), as is well known with antibiotics, by demanding regularly long-term period without
450 treatment. However, combining chronic treatments and long periods free of AR may be difficult
451 to achieve in real systems. Our model only considers temporal refuges, and the conceptualization
452 of untreated areas as equivalent to triggering treatment at intermediate vole density cannot pro-
453 vide guidance on the size of these spatial refuges. Nevertheless, while management of voles is

454 implemented at the scale of fields, mustelids and foxes roams over much larger areas (King and
455 Powell, 2006), such that large refuges with medium-high vole densities voles would be required
456 while maintaining low vole density at local scale.

457 **4.5 Conclusion and perspectives**

458 Our process-based model revealed pesticides that permeate the food chain upward can lead to di-
459 verse population dynamics with alternative states regulated by predators and farmers. It also shows
460 that the practice currently promoted to use low-dose AR treatments at low vole density could have
461 the undesirable side-effects of leading to chronic application of AR on a large scale, in the absence
462 of refuges, and the depletion of the vole predator community. This emerging question would benefit
463 from a landscape modelling approach to characterize spatial refuges. An other prerequisite for this
464 work becoming applied and guiding management practice would be to further explore toxicologi-
465 cal lethal and sublethal (e.g. growth, reproduction, behaviour) effects of pesticides on population
466 dynamics. We have also uncovered a counterintuitive mechanism whereby, owing to intraguild
467 predation, mustelids could rescue foxes from poisoning. This suggest that contemporary Environ-
468 mental Risk Assessment of pesticides that mostly consider one-species - one-compound experiments
469 fail to capture the impact of pesticides on trophic links. Assessing risk at the ecosystem level is
470 empirically challenging such that process-based modelling can play a critical role.

471 **Authors' contributions**

472 XL conceived the initial idea; all authors developed the concept; VB and JF developed the models
473 and led manuscript writing; VB implemented the model and ran simulations. GC contributed treat-
474 ment protocols; MC, JF and VB explored model parameters; XL, PG and MC contributed critically
475 to drafts; all authors gave final approval for publication.

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482 **DATA availability statement**

483 All code and data used for this manuscript are available on Github [https://zenodo.org/badge/](https://zenodo.org/badge/latestdoi/233555669)
484 [latestdoi/233555669](https://zenodo.org/badge/latestdoi/233555669) (Baudrot et al., 2020).

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Table 1 – Populations dynamics and toxicological parameters used in the model simulations. Notation ppm stands for parts-per-million [mg kg^{-1}] and n.d. denotes a dimensionless parameter. See description in Methods.

Parameters	Definitions	Units	Value
Population dynamics parameters			
r_V	Maximal growth rate of voles	day^{-1}	$\ln(2 \times 600)/365$
K_V	Carrying capacity of voles	ind ha^{-1}	600
a_M	Attack rate of mustelids on voles	day^{-1}	1/30
h_M	Handling time of mustelids on voles	day	1/3.5
a_{VF}	Attack rate of foxes on voles	day^{-1}	1/10
a_{MF}	Attack rate of foxes on mustelids	day^{-1}	$1/10 \times 80/300$
h_F	Handling time of foxes on voles	day	1/6
ε_M	Conversion efficiency of ingested food	n.d.	0.0025
m_M	Natural mortality rate of mustelids	day^{-1}	$1/(0.8 \times 365)$
r_F	Maximal growth rate of foxes	day^{-1}	$\ln(3)/365$
K_F	Carrying capacity of foxes	ind ha^{-1}	0.03
B_V	Mean biomass of a vole individual	g	80
B_M	Mean biomass of a mustelid individual	g	300
B_F	Mean biomass of a fox individual	g	5800
d	Degradation rate of dead vole	ind.day^{-1}	1/7
Toxicological parameters			
$T_{broma}(V)$	Farmer input AR function of vole density	$\text{mg ha}^{-1} \text{day}^{-1}$	Scenarios
k_0	Disappearance of AR in the field	day^{-1}	$(0.106 + 0.057)/2$
M_{in}	Maximal intake rate of vole	ppm day^{-1}	$6 \times 1/5$
D_{in}	Half saturation intake rate of vole	ppm	100
H_V, H_M, H_F	Hill's coefficient dose-response curve	n.d.	4
$LD_{50,X}$	Lethal Dose for 50% of individuals of X	ppm	see Text
η_M	AR uptake rate in mustelids	n.d.	0.5
η_F	AR uptake rate in foxes	n.d.	0.5
$k_{out,V}$	Excretion rate of AR by voles	day^{-1}	0.4
$k_{out,M}$	Excretion rate of AR by mutelids	day^{-1}	0.6
$k_{out,F}$	Excretion rate of AR by foxes	day^{-1}	0.6

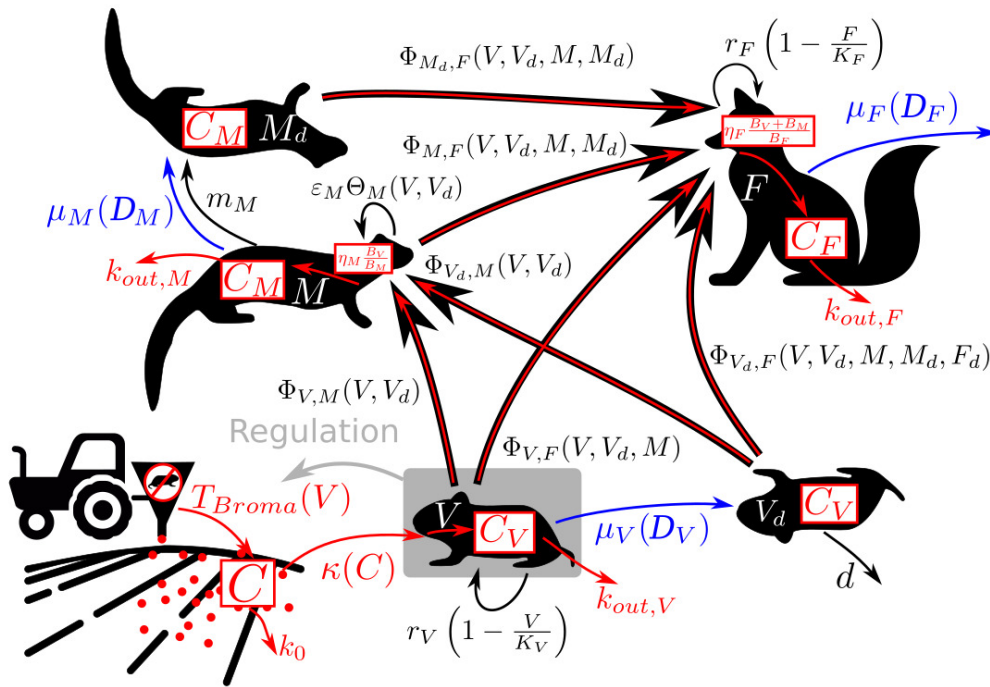


Figure 1 – Flow diagram of the fate and impacts of anticoagulant rodenticide (AR, bromadiolone) in a tri-trophic food web. Black arrows and equations correspond to natural dynamics with trophic interactions, red arrows and equations represent the transfer of AR through the system and its accumulation into the different compartments. The arrows and equations in blue correspond to its impact (i.e., death of individuals) on the three species populations.

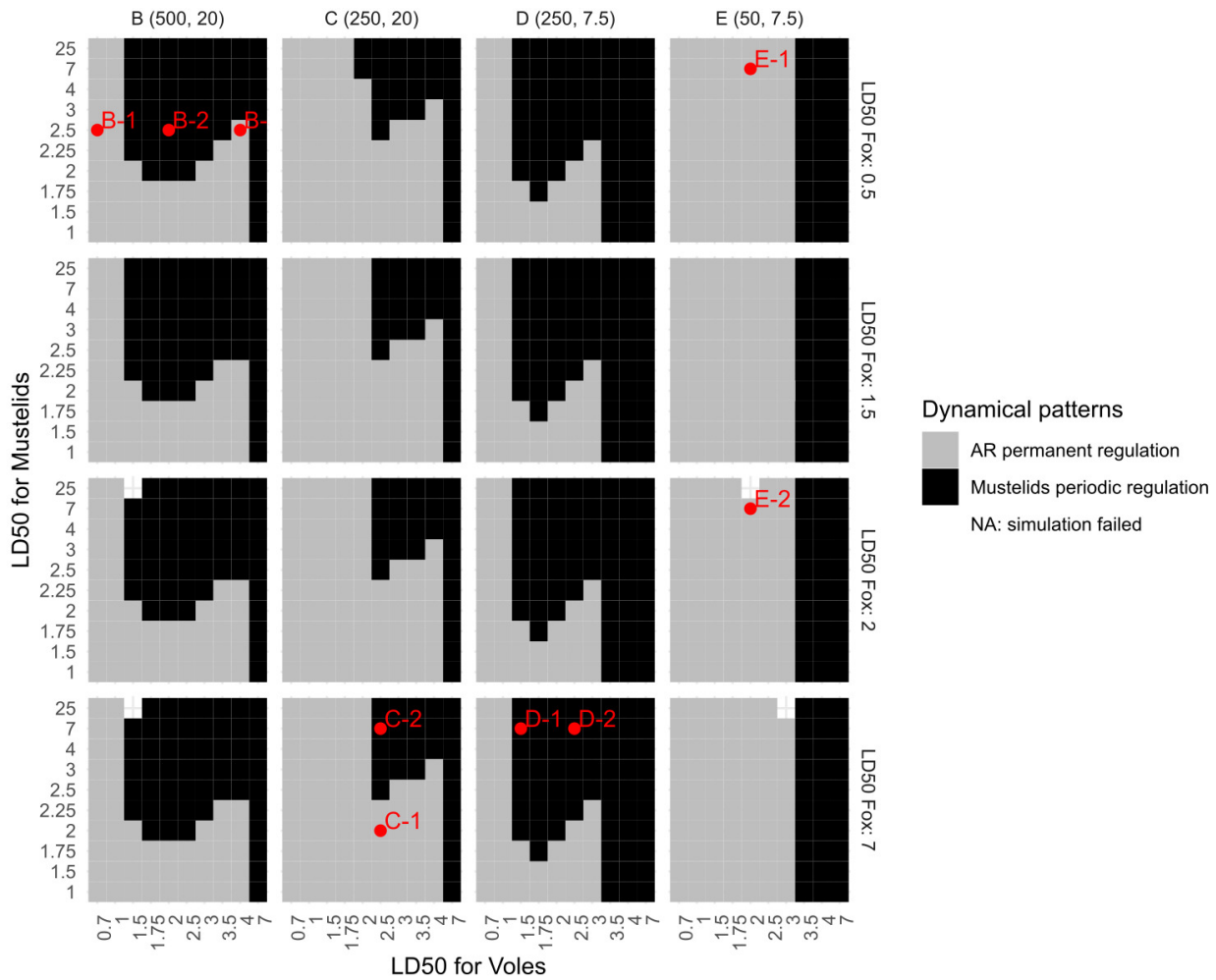


Figure 2 – Global sensitivity analysis of Lethal Dose for voles, mustelids and predators on the population dynamics of mustelids: "Mustelids periodic regulation" (see Figure 3-c,f,g,h) or "Permanent AR regulation" (see Figure 3-b,d,e,i,j). Points are simulations in Figure 3.

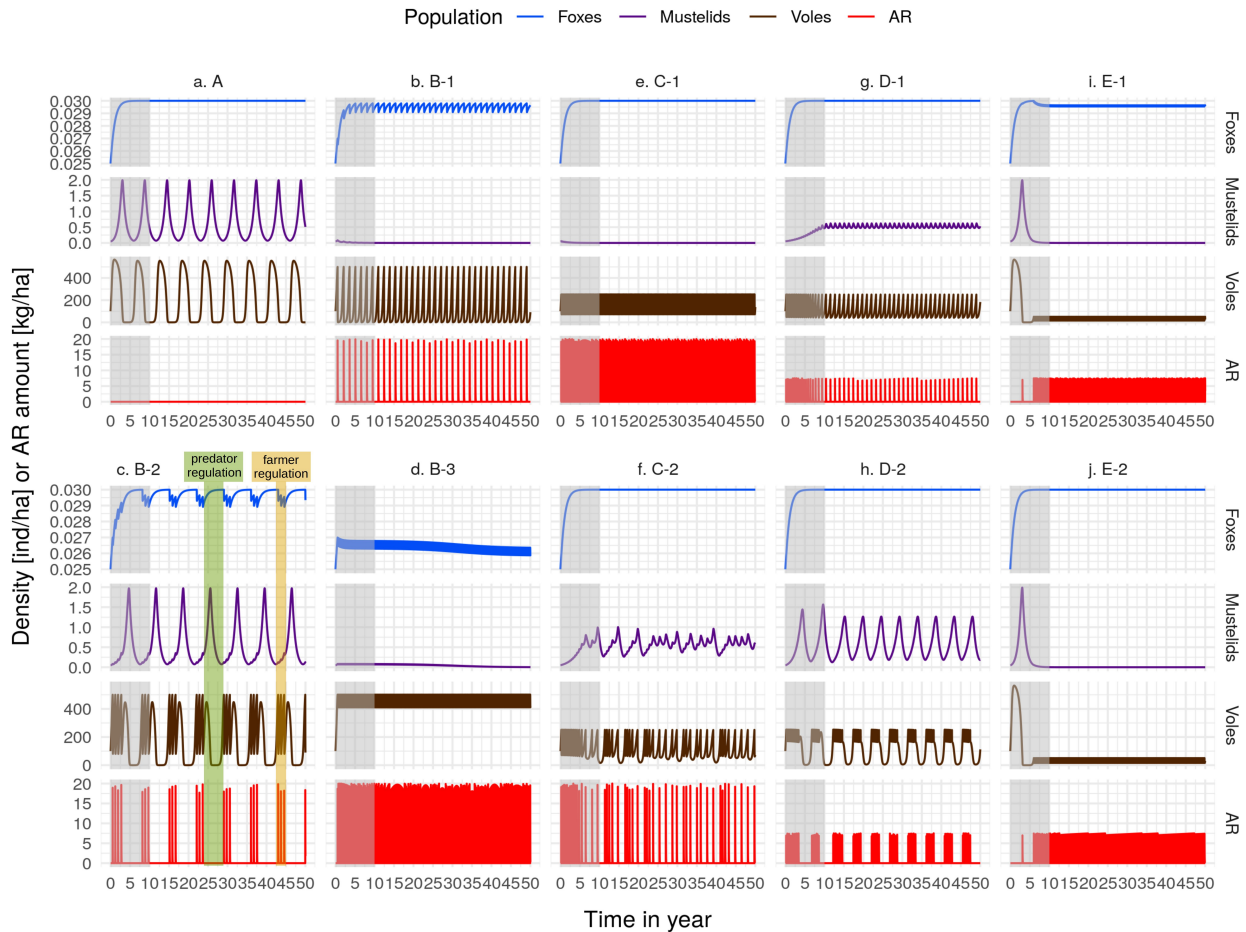


Figure 3 – Variation of population densities simulated over 50 years (10 years of burn-in period in grey area). Capital letter corresponds to farmer functional response: A the null model (no AR), B (vole threshold: 500in/ha, AR 20kg/ha), C(250,20), D(250, 7.5) and E(50,7.5). Numbers correspond to points in Figure 2. Panel c (B-2) illustrates predator-regulation and farmer-regulation phases, that can also be found in other graphics (C-2, D-1, D-2).

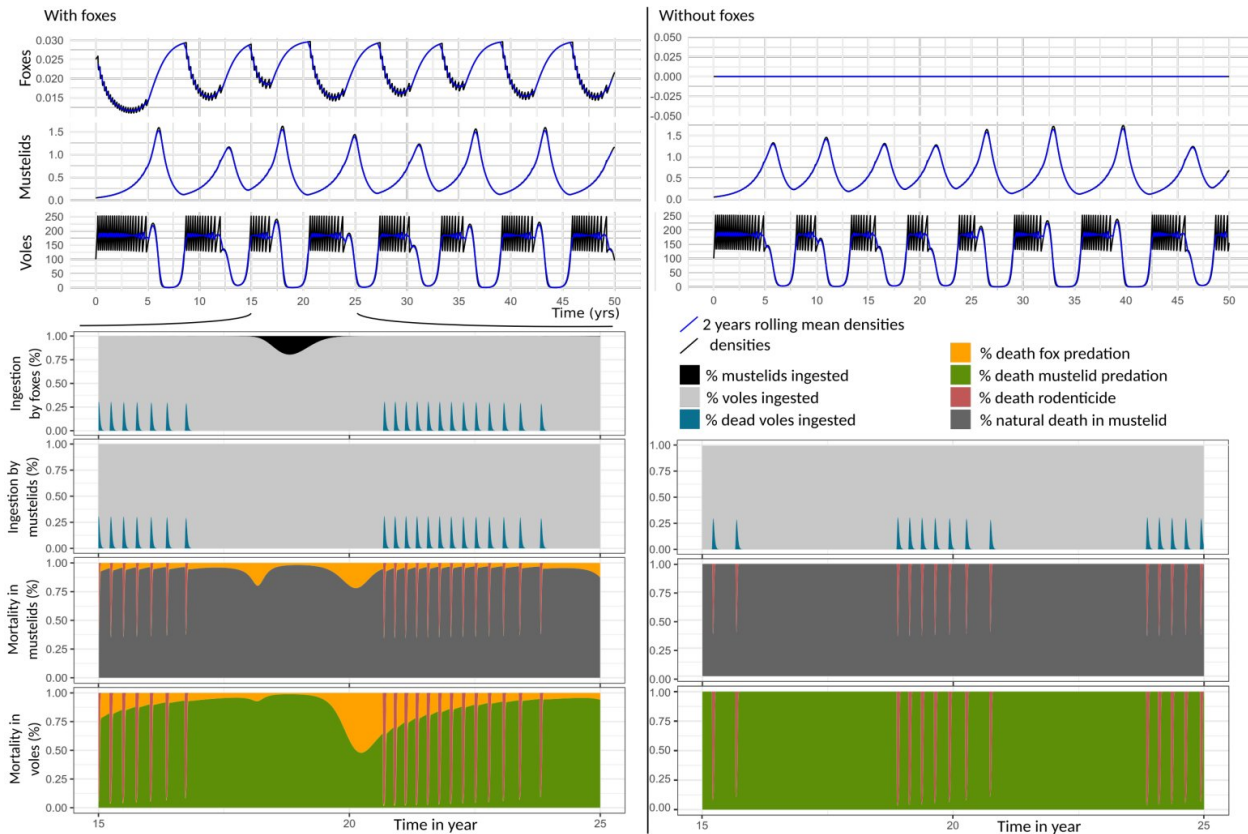


Figure 4 – Effects of intraguild predation. Top panels: daily densities of voles (in black) and 2-years rolling means of densities (in blue) for the farmer functional response d (i.e., 7.5kg of rodenticide at threshold density of 250 voles) with and without foxes (left and right respectively). Bottom graphics: stacked charts of ingestion and mortality proportion.

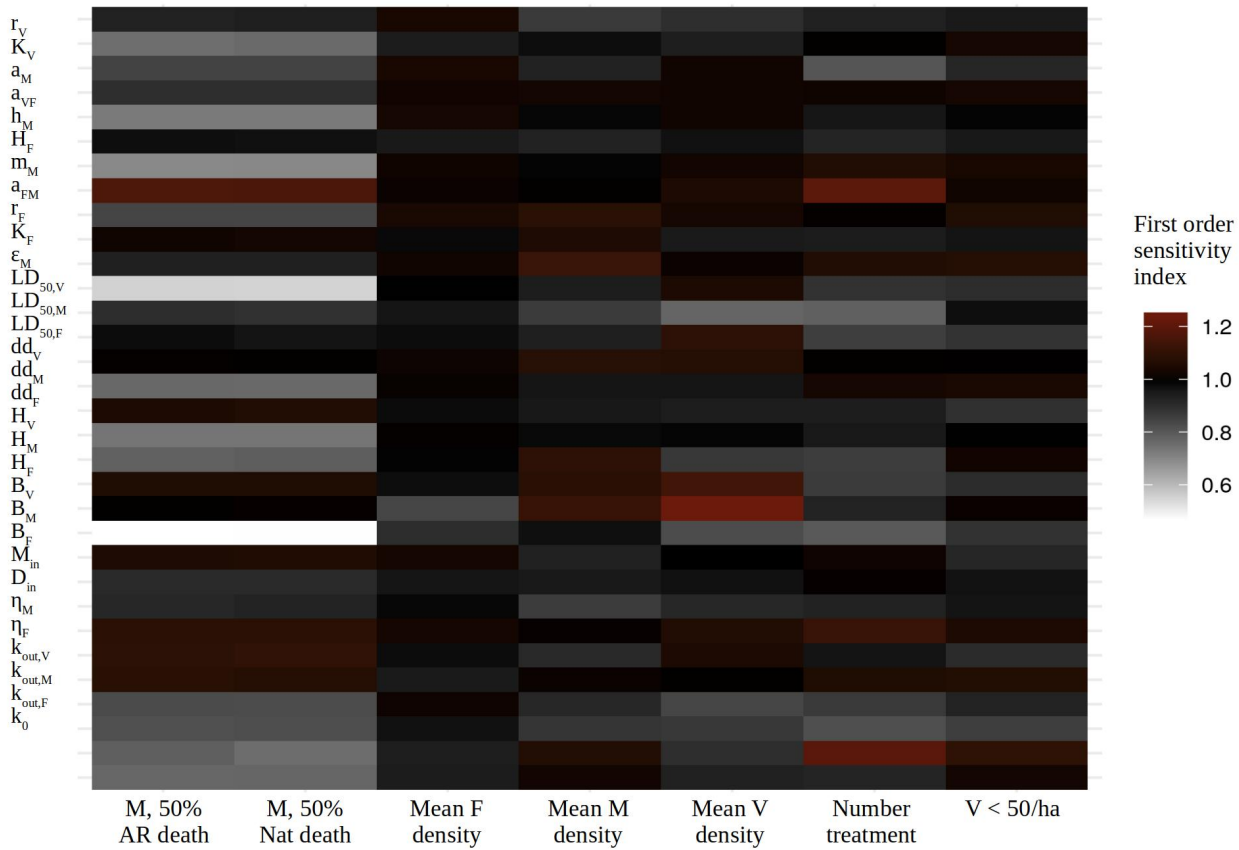


Figure 5 – Local sensitivity analysis: First-order Sobol's sensitivity index (denoted S_p for parameter p) providing the global sensitivity of cost functions to parameters of the model (Sobol and Saltelli, 1993) for the model d (i.e., 7.5kg of AR at threshold density of 250 voles). Analysis is based on 4200 simulations. Values reflect the expected fractional reduction in the variance for each cost functions that would be achieved if a specific parameter is fixed. Values of S_p may be greater than one due to the potential correlation between variables.