

1 Title: Optimal control in the face of evolving resistance by hiding portions of the popu-
2 lation from selection.

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Abstract

Evolved resistance to xenobiotics (i.e. antibiotics, herbicides, pesticides, fungicides) is a global threat to public health and food security. In agricultural systems non-chemical control methods can be combined with xenobiotics (Integrated Pest Management; IPM) to prolong the useful life of compounds and manage pest populations after resistance has evolved. We find IPM strategies with the highest economic returns for an arable cropping system, and perform a global sensitivity analysis to find the factors that shape those strategies. The key uncertainties we find are economic in nature, and farmers have an economic incentive to be responsive to changes in weed the shape of the yield loss function. Doing so will require estimating, at a minimum, what yields would be in the absence of the pest, and how yields change with increasing pest density, with enough detail to say how much control (if any) is justified.

Significance

Integrated pest management (IPM) applies chemical and non-chemical control methods to pest populations to manage evolved resistance. However, we have a poor understanding of when different IPM strategies are incentivised. We find IPM strategies with the highest economic returns in an arable cropping system where high levels of herbicide resistance has evolved repeatedly. The best IPM strategies were dependent crop yields, yield loss caused by the weed, land tenure and levels of herbicide resistance. With the exception of herbicide resistance, all these factors are economic in nature. Knowing which IPM strategy to apply where will require, at a minimum, knowing the yield loss function for the major weeds of a farm, an economic problem rather than biological one.

Introduction

Controlling populations in the face evolving resistance to xenobiotics (i.e. antibiotics, herbicides, pesticides, fungicides) is one of the biggest challenges facing public health (Laxminarayan *et al.*, 2016; Willyard *et al.*, 2017), and food security (Denholm & Rowland, 1992; Palumbi, 2001; Hicks *et al.*, 2018). Evolved resistance also costs billions of dollars globally (Livingston *et al.*, 2016; Chesson *et al.*, 2018; Hicks *et al.*, 2018). While there have been some successes in combating resistance in public health (REX Consortium, 2013), resistance is still a major problem in health care (Willyard *et al.*, 2017) and there has been little success in other contexts, such as food production.

Current strategies to manage resistance focus on delaying the initial evolution of resistance by reducing the population (reducing the potential for *de nova* resistance mutations), and killing any resistant mutants by using a second compound (Denholm & Rowland, 1992; REX Consortium, 2013). Multiple compounds are either stacked (used at the same time) or cycled in sequence. While these strategies can be effective in delaying the initial evolution of resistance they may be counter-productive if xenobiotic resistance is already present, which is true of important pests in food production systems (Denholm & Rowland, 1992; Hicks *et al.*, 2018) and threats to human health (Willyard *et al.*, 2017). Strategies like stacking and cycling involve the continuous (and even increased) use of xenobiotics, which can help drive existing resistance through an entire population (Denholm & Rowland, 1992; Hicks *et al.*, 2018).

In agricultural systems chemical control can be used in combination with non-chemical control such as crop rotation, cultivation and spot control (e.g. hand-weeding), known as integrated pest management (IPM). IPM can be used both pro-actively to delay the evolution of resistance, and reactively to control pest populations as chemical control becomes less effective. While the concept of IPM is well established (Bottrell *et al.*, 1979), finding good IPM strategies is challenging (Dana *et al.*, 2014; Chalak & Pannell, 2015). Management tools need to be used in the correct combination and sequence to be most

effective. This results in a very large number of potential IPM strategies (i.e. different combinations and sequences), even when considering only a handful of management tools and short time horizons (Chalak & Pannell, 2015). As a result there have been few attempts to rigorously search for good IPM strategies (see Chalak & Pannell, 2015 for an exception). More commonly optimal strategies have looked for the best allocation between a few management options (Epanchin-Niell & Hastings, 2010; Meisner *et al.*, 2016; Okumura *et al.*, 2016; Büyüктаhtakın & Haight, 2017), and none have been developed where resistance could evolve to one of the primary management tools. These are important omissions for food production systems where resistance to xenobiotics has evolved numerous times (Denholm & Rowland, 1992; Palumbi, 2001) and multiple non-chemical control options are available that can be used in combination to deliver cost effective control (Chalak & Pannell, 2015).

Little is known about how robust good IPM strategies are to changes in factors such as crop yield and pest population dynamics (Epanchin-Niell & Hastings, 2010). However, previous work on the optimal control of invasive populations has found general factors that shape optimal decisions. Biologically, a population's ability to escape density dependence shifts optimal control to younger age classes (Pichancourt *et al.*, 2012). The degree to which eradicated regions can be re-invaded also influences the optimal control strategies (Januchowski-Hartley *et al.*, 2011; Epanchin-Niell & Wilen, 2012), but the exact strategy depends on the way suitable habitats are connected (Chadès *et al.*, 2011). Economic factors tend to be at least as important as biological ones in shaping the optimal control strategy (Epanchin-Niell & Hastings, 2010). In particular the relationship between the density of an invasive species and the damage it does has been found to be crucial (Yokomizo *et al.*, 2009). The way that future returns are valued also strongly influences the optimal control strategy; when more value is placed on future versus present returns more intensive control is favoured (Epanchin-Niell & Hastings, 2010).

We apply a genetic algorithm to a population model of an important weed of wheat in Europe (*Alopecurus myosuroides*), where resistance to two herbicides can evolve. [ALEXA:

sentence or two here + REF on just how damaging BG is]. To allow IPM strategies we also include crop rotation, cultivation and spot control as management options. We use global sensitivity analysis to determine the minimum information required to create a field scale IPM strategy that maximizes economic return and the impact of those strategies on herbicide resistance and economic gross margin.

Results and Discussion

Parameters defining the yield loss function (see Table 1) had the most influence on shaping incentivized IPM strategies. We used a linear yield loss function to relate density of *A. myosuroides* to winter wheat yields, fit to data from 10 fields (Appendix 1). The intercept (Y_0) and slope of the yield function (Y_D) were two of the most important parameters (Fig. 1). Another set of parameters that control how large the seed bank can become (f_m , f_d and ϕ_b) were also important. The potential size of the seed bank scales the x-axis of the yield function. Although yield functions have been estimated for major weeds (Cousens, 1985; Doyle *et al.*, 1986; Swinton *et al.*, 1994), there is evidence that yield functions vary substantially between fields (Swinton *et al.*, 1994; Hicks *et al.*, 2018), and little attention has been paid this variation and understanding its causes.

While the shape of the yield function is important, it may not be necessary to know it in great detail to find IPM strategies with high gross margin. When the yield of winter wheat with no *A. myosuroides* (Y_0) was low, management intensity was lower and relied on crop rotation and tactical use of herbicide (Fig. 2, 'Y₀ low'). The strategy changed little when slope of the yield function (Y_D) increased. Although, more herbicide was used when the value of Y_D increased from a very low value to a higher value (1% to 12% losses at high densities of *A. myosuroides*; Fig. 2g,e). When Y_0 was high, Y_D showed two thresholds where IPM strategy changed. When Y_D was very low the best strategy was to do noting and live with high populations of *A. myosuroides* (Fig. 2h), since yield losses were never high enough to justify expenditure on control. Increasing Y_D slightly

parameter	rel. inf.
Y_D	16.5
f_d	13
Y_0	10.2
f_m	9.2
ϕ_D	8.1
G_{\min}	5.4
γ	4.7
η_s^0	4.4
β	4.4
\varnothing	2.9
\varnothing	2.9
G_{\max}	2.8
N_{int}	2.6
θ	2.4
η_h	2.1
\downarrow	2.1
η_b	1.8
η_s	1.6
α	1.6
ϕ_e	1.5

Figure 1: Relative influence (relative reduction in squared error; Miller *et al.*, 2016) of each parameter, averaged across dimensions of the solution space (Appendix 2). Values are scaled 0 to 100, with higher values indicating parameters with more influence on the structure of incentivized IPM strategies. See Table 1 for an explanation of parameters.

meant some herbicide use and cultivation to rotate the seed bank became advantageous (Fig. 2f). Once Y_D increased enough to justify intensive control, further increases did not change the IPM strategy (Fig. 2b,d). Thus, knowing how much management is justified by the yield loss function may only require having an estimate of the intercept (Y_0), and one or two thresholds values of Y_D , where a new IPM strategy becomes advantageous.

Supporting previous work (Epanchin-Niell & Hastings, 2010) we found higher values on future returns (larger values for γ) lead to more intensive IPM strategies. Intensive management to reduce the seed bank was only used when discount rates were high (Figure 3). Recall that although we only show the first 10 years of IPM strategy, the discounted returns over 25 years are considered by the genetic algorithm. In agricultural systems land tenure have a crucial effect on how investments in weed control are valued. Those who own fields can benefit from long-term investments like weed control campaigns and soil conservation, whereas those who rent fields do not (Wiese *et al.*, 1996; Fraser, 2004). 35% of all agricultural land in England and Wales is under tenancy agreement (CAAV, 2017). This has important implications for the level of *A. myosuroides* control managers are incentivised to provide, and thus its spread and the evolution of resistance.

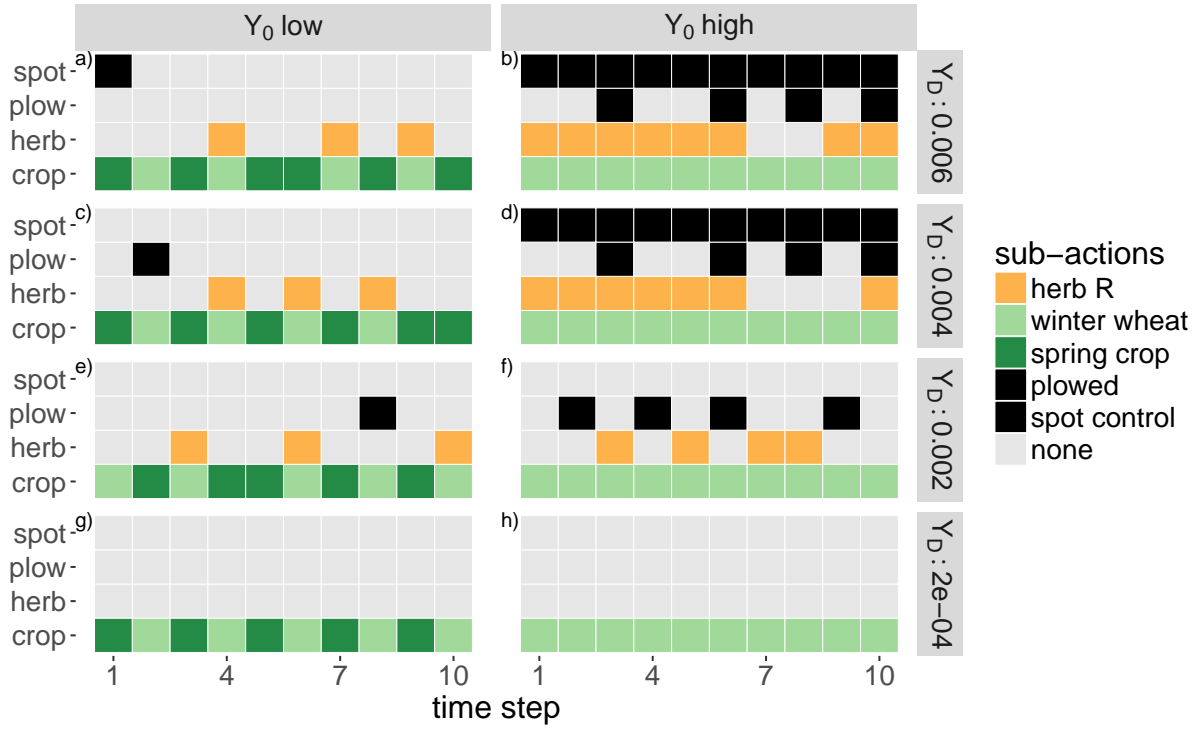


Figure 2: IPM strategies under high ($\text{£}1668 \cdot \text{ha}^{-1}$) and low ($\text{£}986 \cdot \text{ha}^{-1}$) values of Y_0 (yield of winter wheat with no *A. myosuroides*), under increasing values (rows) of Y_D (in $\text{£} \cdot \text{plant}^{-1} \cdot \text{ha}^{-1}$). At the lower limit of Y_D very high *A. myosuroides* densities result in a 1% yield loss under the high Y_0 scenario, and the upper limit implies a yield loss of 35% . There is initially one effective herbicide ($R_{\text{int}} = 0.0001$, $Q_{\text{int}} = 0.9$).

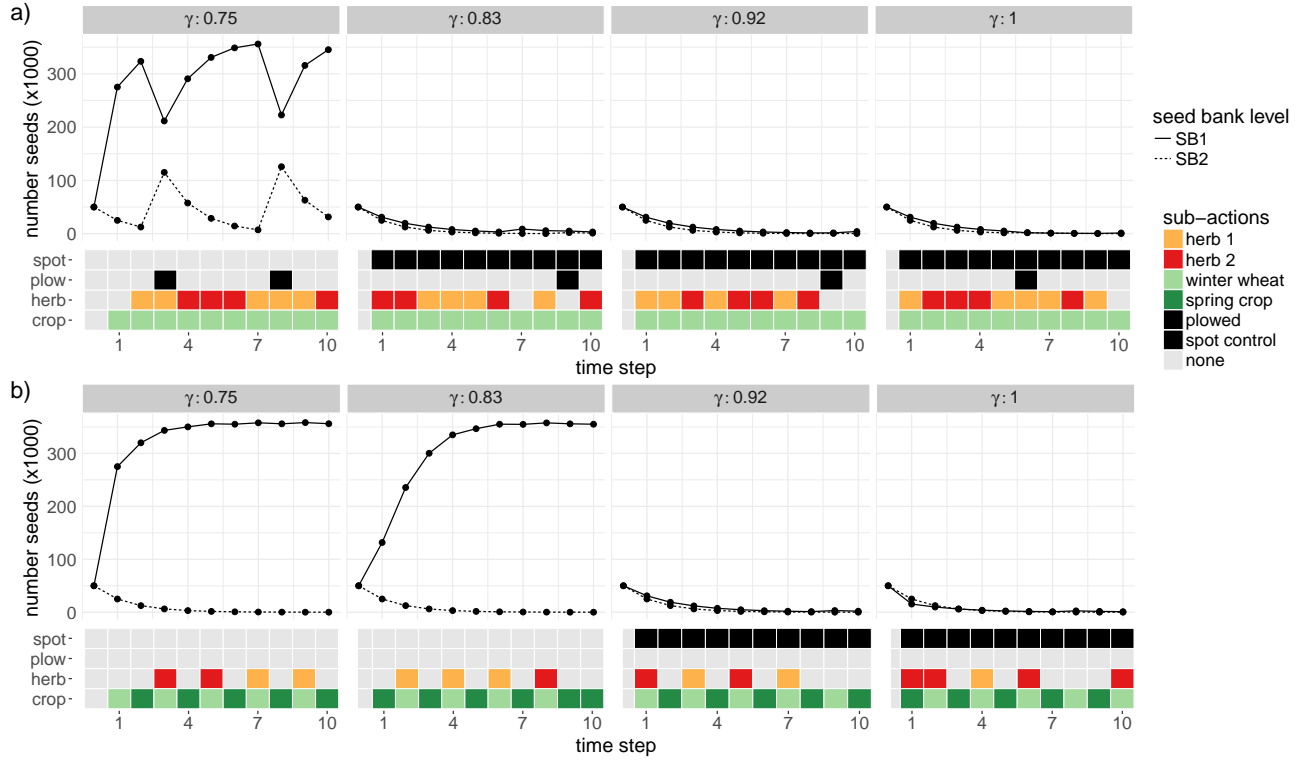


Figure 3: The effect of discount rate (γ) on the seed bank and IPM strategy (tile plots) when yields from winter wheat are high (a; $\text{£}1668 \cdot \text{ha}^{-1}$) and low (b; $\text{£}986 \cdot \text{ha}^{-1}$). In both cases the slope of the yield function (Y_D) is high ($0.006 \text{£} \cdot \text{plant}^{-1} \cdot \text{ha}^{-1}$). Initial resistance was low for both herbicides.

When both herbicides were effective the preference was to cycle between them, however even this did not prolong their continued use by much. Even when both R and Q started at frequencies of 1 in 20,000 alleles (Figure 4a) continued herbicide use raised those frequencies to 1 in 50 within 10 time steps. This frequency provided enough variation for selection to rapidly act on (Figure 4c).

We present the best case that can be hoped for in reactive management, as we assume that herbicide resistance was conferred by target site mutations. This is why cycling was often favoured over stacking, as cycling prolonged the useful life of both herbicides since the application of each was spread out. However, there is growing evidence that non-target site resistance, which confers cross resistance, is widespread (Hicks *et al.*, 2018). If generalized, non-target site resistance mechanisms are present, the total amount of herbicide exposure predicts resistance level (Hicks *et al.*, 2018), and cycling will not help.

Even when initial frequencies of resistance to the remaining effective herbicide was low (1 in 200; Figure 4c) initial continual use quickly drove the evolution of resistance to levels where herbicide use was greatly reduced after just five generations. As the initial frequency of resistance to the remaining effective herbicide ('herb Q' in this case) increased, even moderate herbicide use drove the rapid evolution of herbicide resistance. With an initial frequency of resistance alleles of 1 in 20 even three herbicide applications were enough to increase resistance to the point where applying both herbicides would result in 40% survival (Figure 4d). Once this situation was reached, gross margin was reduced by a quarter compared to returns with low resistance.

This is in contrast to current management practice in this cropping system, where multiple herbicide applications a year are the norm, despite high levels of resistance (Hicks *et al.*, 2018). This disparity could arise from a number of contributing factors. Some managers may believe that even a little control (mortality of a few susceptible individuals) is better than no control and inaction is seen as the worst approach to weed management (Wilson *et al.*, 2008). In addition, IPM strategies are often seen as complex in comparison to routine application of chemicals, and having a steep learning curve (Llewellyn *et al.*,

2006). Resistance tends to be partial and build up slowly (Moss & Hull, 2009; Hull *et al.*, 2014), so farmers may be victims of a shifting baseline, lowering their expectations of efficacy of weed control. There may be a belief that new herbicides will become available, despite no new modes of action being marketed for over 20 years (Duke, 2012). Thus, current strategies are viewed as a bridging strategy until a new product is found (Hurley & Frisvold, 2016). Finally, Our population model was deterministic, so IPM strategies could not be risk averse to variability in *A. myosuroides* populations and economics factors like crop prices. Uncertainty in when herbicide resistance will emerge and the efficacy of non-chemical control can be a major impediment to adopting IPM (Hurley & Frisvold, 2016).

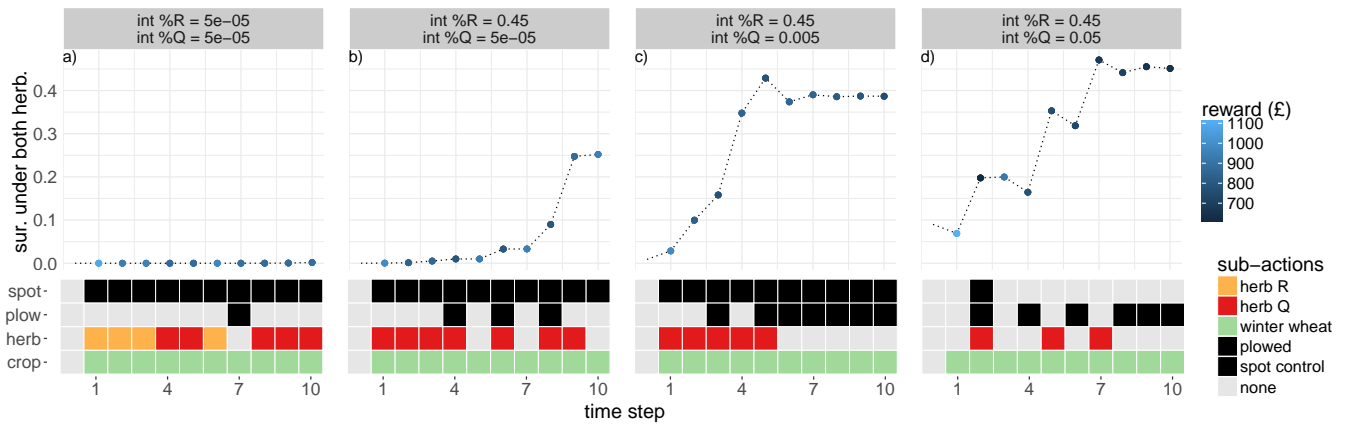


Figure 4: The effect of initial resistance on the selected IPM strategy (tile plots) and the evolution of herbicide resistance (% survival to under both herbicides). Lighter coloured points indicate higher reward (gross margin) obtained in that time step. In this case $Y_0 = 1668$ (high winter wheat yield) and $Y_D = 0.0062$ (high yield penalty).

We assume that herbicide is the only action that drives the evolution of resistance. Any effective management tool will impose selection pressure, and so drive resistance to that tool. In reality, the spring cropping and spot control sub-actions make heavy use of glyphosate to control *A. myosuroides*. Glyphosate resistance has evolved on many separate occasions in response to prolonged, heavy use (Sammons & Gaines, 2014). As glyphosate becomes a more important part of weed management (Hicks *et al.*, 2018) resistance is likely.

Conclusion

Combating xenobiotic resistance is ultimately a problem of behaviour change, and thus how individuals are incentivized to act (Hurley & Frisvold, 2016). Our results show that farmers have an economic incentive to be responsive to changes in weed the shape of the yield loss function. Doing so will require estimating, at a minimum, what yields would be in the absence of the pest, and how yields change with increasing pest density, with enough detail to say how much control (if any) is justified.

[ALEXA: would be nice to conclude with a nice summary statement on what a failure to be responsive will cost farmers, the economy as whole and the environment, are at least as much as we can say at this point. Also be a good place to flag your up coming work for any things that are still unresolved. If it is all unresolved that is an important message as well I think.]

Data Archival

PNAS must be able to archive the data essential to a published article. Where such archiving is not possible, deposition of data in public databases, such as GenBank, Array-Express, Protein Data Bank, Unidata, and others outlined in the Information for Authors, is acceptable.

Supporting Information (SI)

Appendices

Appendix 1: Reward Function

Appendix 2: Finding Which Initial Conditions and Model Parameters Lead to Which IPM Strategies

Appendix 3: Population Model

Table 1: Parameter descriptions

Parameter	Description
<i>Population Model see Appendix 3</i>	
ϕ_e	germination probability
ϕ_b	Probability a seed survives one year
s_0	Survival probability without herbicide
θ	Proportion of <i>A. myosuroides</i> population exposed to herbicide under sub-action a_h = herb R, herb Q or both. plants may be missed spatially or temporally, or spraying may be affected by rain.
s_h	Survival of susceptible <i>A. myosuroides</i> exposed to herbicide (a_h). Herbicide assumed to be effective.
α	Survival probability under the alternative crop (a_k = alt), spring barley.
β	Survival under spot control (sub-action a_s), for example because plants are missed.
f_m	Seeds \cdot plant $^{-1}$ when density is 0 (f_m) and the effect of
f_d	density on seed production (f_d) interact to determine maximum population size. Values chosen to keep the max population close to the maximum population seen in Queenborough <i>et al.</i> (2011) so yield is not extrapolated outside the observed range.
I	The proportion of seed moved between seed bank levels by ploughing (sub-action a_b)
<i>Reward function see Appendix 1</i>	
γ	discount rate on future returns.
Y_0	Yield in $\text{£}\cdot\text{ha}^{-1}$ from winter wheat when <i>A. myosuroides</i> is absent.
Y_D	reduction in yield caused by each <i>A. myosuroides</i> (in $\text{£}\cdot\text{plant}\cdot\text{ha}^{-1}$).
ϑ	Yield of spring barley in $\text{£}\cdot\text{ha}^{-1}$, an alternative crop commonly used to control <i>A. myosuroides</i> .
ϖ	Proportion of yield achieved if crop a_k is repeated
η_h	Cost of a single herbicide application in $\text{£}\cdot\text{ha}^{-1}$.
η_b	Cost of ploughing in $\text{£}\cdot\text{ha}^{-1}$.
η_s^0	Cost of spot control even when <i>A. myosuroides</i> density is 0, in $\text{£}\cdot\text{ha}^{-1}$.
η_s	Increase in spot control cost for each <i>A. myosuroides</i> in $\text{£}\cdot\text{plant}^{-1}\cdot\text{ha}^{-1}$
η_k	Costs of crop a_k , in $\text{£}\cdot\text{ha}^{-1}$, not associated with the other sub-actions targeted at <i>A. myosuroides</i> control.
<i>Initial Conditions</i>	
$G_{\min/\max}$	Initial allele frequency of mutants resistant to the herbicide with the lowest (min) and highest (max) population level resistance.
N_{int}	Initial number of seeds in each level of the seed bank

Appendix 4: Action Space

Appendix 5: Genetic Algorithm

Methods

We frame IPM as a combinatorial optimisation problem where the goal is to find a good combination of management tools, used in sequence. We use a genetic algorithm to solve this combinatorial problem (Taylor & Hastings, 2004; Carrasco *et al.*, 2010). Genetic algorithms cannot be checked to have found the globally optimal solution, as this would require already knowing the solution. However, genetic algorithms are efficient at weeding out comparatively poor solutions, so that over successive iterations the regions of the solution space being explored gets progressively better, resulting in a set of good (often near optimal) solutions.

Our goal is to find good IPM strategies in the face of rapidly evolving resistance, and how those strategies change in response to biological and management parameters. This problem has four parts: i) A reward function that measures how good a given IPM strategy is based on how much that strategy cost and its effectiveness, we use net present economic value. ii) A population model that translates a given IPM strategy into a population, and thus a reward. iii) An algorithm that finds IPM strategies with higher rewards, the genetic algorithm. iv) Finally we need to relate changes in the best IPM strategy found to changes in initial conditions and model parameters. We use a meta-modelling global sensitivity analysis (Coutts & Yokomizo, 2014) based on multi-variate boosted regression trees (Miller *et al.*, 2016).

Population model

The population model links management actions to the response of the *A. myosuroides* population, and thus wheat yields. The action a_j is how the manager effects the population model, and thus the reward they get. Each action is a tuple of four sub-actions $a_j = \langle a_h, a_b, a_k, a_s \rangle$, see Appendix 4 for a description of the sub-actions and all eligible combinations of these sub-actions (i.e. the full actions space, \mathbf{A}).

The processes included in the population model limit the scope of the IPM strategies found. We use a deterministic model, and so our IPM strategies can only deal with average expected population responses, ignoring demographic uncertainty, and environmental and market variability. Also, we only model herbicide resistance that is already present in the population because *de nova* mutation is a fundamentally stochastic process.

A commonly recommended (REX Consortium, 2013) and applied (Hicks *et al.*, 2018) strategy to combat resistance is to apply xenobiotics that impair different cellular pathways (i.e. modes of action), either sequentially (cycling) or concurrently (stacking). To allow this behaviour we use a discrete time, spatially implicit model, where two independent alleles (R and Q), each confer target site resistance to a separate herbicide. The model must also be flexible enough to accommodate non-chemical control. We include a two level seed bank (to allow plowing to take seeds out of the germinating population) and model survival as a function of resistance, herbicide choice, crop choice and spot control (where the cost increases with *A. myosuroides* density). The model tracks the number of seeds in each level of the seed bank in each of nine genotypes G , starting at the beginning of the growing season before any seeds have emerged. See Appendix 3 for a full description of the model and how each sub-action affects the population.

Reward function

The reward function measures how good an IPM strategy is, given a initial starting condition and parameter set that the model is run under. The reward function encodes the goals of a manager. We assume farmers are primarily driven by economic returns. The economic return consists of two parts, the income made from the crop and the costs of producing that crop. We assume that usual farm costs, such as buildings and machinery as constant from year to year, so we focus on gross margin, i.e. income - variable costs (Redman, 2016, pp. 3–4).

To explicitly link the above ground population to the reward function we define $N''(\mathbf{a}, n_0, t)$, the total above ground population after all control actions, at time t given an initial population n_0 and a sequence of actions

$$\mathbf{a} = \{a_j^0, a_j^1, \dots, a_j^T\} \quad (1)$$

where a_j^t is the action $a_j \in \mathbf{A}$ taken at time t and T is the time horizon over which management is run. We assume all returns after T are ignored. The reward function is

$$R(\mathbf{a}, n_0) = \sum_{t=0}^T \gamma^t \left(Y(N''(\mathbf{a}, n_0, t)) - C(a_j^t) \right) \quad (2)$$

where $R(\mathbf{a}, n_0)$ is the time discounted reward for action sequence \mathbf{a} given starting population n_0 , $\gamma \in [0, 1]$ is the discount rate. When $\gamma = 0$ only the reward in the first time step is considered, when $\gamma = 1$ returns in all future time steps up to T are valued equally. $Y(N''(\mathbf{a}, n_0, t))$ is the income (in £/ha) from the crop chosen at time t given start stating n_0 and following action sequence \mathbf{a} . $C(a_j)$ is the cost of taking action a_j , and is composed of the cost of controlling *A. myosuroides* and other costs that depend in the crop being grown (a_k).

See Appendix 1 for the yield and cost models for each sub-action and parameter estimation.

Finding good IPM strategies

Our goal is to find good strategists to manage black grass in the face of evolving resistance, however it is not feasible to test every combination of management options over more than a handful of years. Genetic algorithms have been used to find good solutions to this class of problem (Taylor & Hastings, 2004; Carrasco *et al.*, 2010). The genetic algorithm starts with an randomly generated set of action sequences, these action sequences are then iteratively improved to find a set of action sequences with a high gross margin. Genetic algorithms rely on the fact that even though the number of possible action sequences is large, many perform very poorly. The genetic algorithm explores better performing regions of the solution space more intensely. While genetic algorithms are not guaranteed to find the optimal action sequence they will find a set of actions sequences that perform well, often close to the optimal solution.

To find good action sequences we use a genetic algorithm with knock out tournament selection, where each action sequence in a set of 1000 actions sequences is randomly paired with another, and the action sequence with the highest $R(\mathbf{a}, n_0)$ survives to help generate new action sequences. We used pair mating between survivors and N-point cross-over to produce new action sequences. After new action sequences are created there is a process of random mutation where each a_j^t is changed to another $a_j^t \in \mathbf{A}$ with probability $m = 0.03$. The algorithm used is given in Appendix 5.

Finding which initial conditions and model parameters lead to which IPM strategies

It is unlikely a given IPM strategy will perform well in all scenarios. To find the parameters and initial conditions (n_0) that shaped the IPM strategy with the highest reward, we extend the meta-modelling approach to global sensitivity analysis outlined in Coutts & Yokomizo (2014), to multivariate time series outputs (i.e. the sequences of the four sub actions). We: i) ran the genetic algorithm under 15000 different parameter sets and

initial conditions, generated with Latin hyper-cube sampling (see Table S1 in Appendix 2 for upper and lower limits of each parameter), ii) used Longest Common Sub-Sequence (Toohey, 2015) as a measure of distance between these action sequences, iii) projected the resulting distance matrix into an 8D solution space using non-metric multi-dimensional scaling (implemented in the 'ecodist' R package; Goslee & Urban 2007), iv) predicted where each IPM solution sat in the solution space using multi-variate boosted regression tree (Miller *et al.*, 2016), where the model parameters and initial conditions were predictors. See Appendix 2 for details.

It was this multi-variate boosted regression tree we interrogated to find which parameters and initial conditions were important for changing the best IPM strategy found—using relative influence and partial dependence plots (Miller *et al.*, 2016).

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