Derivation of a Cropping System Transfer Function for Weed Management: Part 1 – Herbicide Weed Management

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Abstract: System behaviour is described by the transfer functions, which relate the system's output to one or more input variables. No-till cropping systems depend on herbicide inputs for weed management and crop yield optimisation. This paper derives the transfer function for crop yield potential as a function of herbicide input, in the presence of herbicide resistance in the weed population, using several mathematical components for crop and weed ecology from published literature. The resulting transfer function reveals the herbicide application rate for optimal crop yield potential and highlights the growing herbicide resistance problem in no-till cropping systems.

Keywords: System analysis, weeds, herbicide, herbicide resistance, crop ecology.

1. INTRODUCTION

System analysis techniques can be applied to most agricultural system in order to better understand their operation and optimise performance. System analysis usually includes the development of transfer functions [1]. Transfer functions are mathematical equations, involving various input variables or matrices, which relate the system's output to one or more of the system's inputs. In the case of an agricultural cropping system the key output from the system is potential crop yield. Some crop ecology studies have demonstrated that competition from weeds can reduce the potential yield of some crops by 35 % to 55 % [2,3].

Modern no-till cropping depends on herbicides for weed management; therefore herbicide applications are an important system input. Unfortunately, herbicide resistance in many weed species is becoming wide spread [4] and multiple herbicide resistances in several economically important weed species has also been widely reported [5]. In time, herbicide resistant weeds may ultimately result in significant yield reductions and grain contamination; therefore this paper derives a system transfer function relating herbicide input to potential crop yield in the presence of herbicide resistance, based on various ecological models published in literature.

2. DERIVATION OF CROP SYSTEM TRANSFER FUNCTION FOR HERBICIDE WEED MANAGEMENT

The effect of weed damage on crop yields can be described by the following [6]:

$$Y = Y_o \left[1 - D(R) \right] \tag{1}$$

In equation (1), D(R) is the damage function caused by a weed density of R, which represents the number of weeds that are recruited from the seed bank (plants m⁻¹ of row). The Damage function can be described by the following equation [7]:

$$D(R) = \frac{I \cdot R}{100\left(e^{ct} + \frac{I \cdot R}{A_w}\right)}$$
(2)

Substituting equation (2) into equation (1) yields:

$$Y = Y_o \left[1 - \frac{I \cdot R}{100 \left(e^{ct} + \frac{I \cdot R}{A_w} \right)} \right]$$
(3)

2.1. Herbicide Weed Management

Weed infestations will be made up of some resistant weeds (R_R) and some weeds that can be easily controlled by herbicides (R_s). After some kind of herbicide treatment, the density of susceptible weeds will be [6]:

$$R_{s} = R_{so} \left[1 - K \left(H \right) \right] \tag{4}$$

 R_{So} is the initially susceptible weed density; K(H) is the kill function for the herbicide in this portion of the weed population for a given herbicide treatment of H. The resistant weed population will be [6]:

$$R_{R} = R_{Ro} \tag{5}$$

 R_{Ro} is the initial resistant weed density. A typical kill function for a herbicide treatment is [8]:

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$$K(H) = 1 - e^{-\lambda H} \tag{6}$$

Substituting all these components into equation (3) yields:

$$Y = Y_o \left[1 - \frac{I\left(R_{so}e^{-\lambda H} + R_{Ro}\right)}{100\left(e^{ct} + \frac{I\left(R_{so}e^{-\lambda H} + R_{Ro}\right)}{A_w}\right)} \right]$$
(7)

If p represents the portion of the population that is herbicide-resistant and R_o is the initial weed population density, then:

$$Y = Y_o \left[1 - \frac{I \cdot R_o \left[\left(1 - p \right) e^{-\lambda H} + p \right]}{100 \left(e^{ct} + \frac{I \cdot R_o \left[\left(1 - p \right) e^{-\lambda H} + p \right]}{A_w} \right)} \right]$$
(8)

The recruitment of seedlings from the seed bank can be described by the following equation [9]:

$$R_o = \frac{W}{1 + e^{-\left(\frac{t - t_o}{d}\right)}}$$
(9)

Substituting this into equation (8) yields:

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$$Y = Y_o \left[1 - \frac{I \cdot W \left[(1-p)e^{-\lambda H} + p \right]}{100 \left[1 + e^{-\left(\frac{t-t_o}{d}\right)} \right]} \left[e^{ct} + \frac{I \cdot W \left[(1-p)e^{-\lambda H} + p \right]}{A_w \left[1 + e^{-\left(\frac{t-t_o}{d}\right)} \right]} \right] \right]$$
(10)

This can be simplified to become:

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$$Y = Y_o \left[1 - \frac{I \cdot W\left[(1-p)e^{-\lambda H} + p \right]}{100 \left(e^{ct} \left[1 + e^{-\left(\frac{t-t_o}{d}\right)} \right] + \frac{I \cdot W\left[(1-p)e^{-\lambda H} + p \right]}{A_w} \right)} \right]$$
(11)

The portion of the population that is resistant to herbicide treatment will change from generation to generation depending on the selection pressure being applied by the herbicide treatments. Based on work by Gubbins and Gilligan [10], if there is a relatively constant selection pressure (s) towards herbicide resistance from generation to generation, then the following relationship will hold:

$$\frac{\partial p}{\partial g} = sp(1-p) \tag{12}$$

This partial differential equation can be solved by integration to give:

$$p = \frac{p_o e^{sg}}{1 + p_o \left(e^{sg} - 1\right)} \text{ and } 1 - p = \frac{1 - p_o e^{sg}}{1 + p_o \left(e^{sg} - 1\right)}$$
(13)

Substituting these equations into equation (11) and simplifying yields:



There is also evidence that herbicides have a toxic effect on the crop as well. Using the study by Yin *et al.* [11] as a guide, and assuming that the toxicity of the herbicide on a crop can be expressed as a polynomial of the form $Loss = aH^2 - BH$, equation (14) can be modified to become:

$$Y = Y_{o} \begin{bmatrix} 1 - \frac{I \cdot W \left[\left(1 - p_{o} e^{sg} \right) e^{-\lambda H} + p_{o} e^{sg} \right]}{100 \left(e^{ct} \left[1 + e^{-\left(\frac{t - t_{o}}{d} \right)} \right] \left[1 + p_{o} \left(e^{sg} - 1 \right) \right] + \frac{I \cdot W \left[\left(1 - p_{o} e^{sg} \right) e^{-\lambda H} + p_{o} e^{sg} \right]}{A_{w}} \right]}{A_{w}} \end{bmatrix}$$

$$(15)$$

The seed bank will be dynamic depending on factors such as natural seed mortality, immigration of seeds into the area from other locations *via* various vectors, emigration of seeds out of the area to other locations *via* various vectors, the onset of dormancy that prevents germination in the current season, and

the breaking of dormancy from previous seasons in the seed bank.



(16)

2.2. Sensitivity Analysis

The development of transfer functions is not always for the purpose of providing accurate prediction but to provide insight into system behaviours as input parameters change. The sensitivity of the output to these changes can be assessed by differentiating the transfer function equations with respect to the input parameter of interest and assessing the magnitude of the resulting differential equation.

Differentiating equation (16) with respect to any of the key parameters allows sensitivity analyses to be performed. For example, differentiating equation (16) with respect to H determines the sensitivity of crop yield to herbicide weed treatments:

$$\frac{\partial I}{\partial H} = Y_{o} \left[\frac{\lambda I \left[W(1-N-D_{o}) - E_{m} + I_{m} \right] \left[1 - p_{o} e^{sg} \right] e^{-\lambda H}}{100 \left[e^{cf} \left[1 + e^{\frac{\left(1-t_{o}\right)}{d}} \right] \left[1 + p_{o} \left(e^{sg} - 1 \right) \right] + \frac{I \left[W(1-N-D_{o}) - E_{m} + I_{m} \right] \left[\left[1 - p_{o} e^{sg} \right] e^{-\lambda H} + p_{o} e^{sg} \right] \right]}{A_{w}}} \right]}{\frac{\partial Y}{\partial H} = Y_{o} \left[\frac{\lambda I^{2} \left[W(1-N-D_{o}) - E_{m} + I_{m} \right]^{2} \left[\left(1 - p_{o} e^{sg} \right) e^{-\lambda H} + p_{o} e^{sg} \right] \left[1 - p_{o} \right]}{100 A_{w} \left[e^{cf} \left[1 + e^{\frac{\left(1-t_{o}\right)}{d}} \right] \right] \left[1 + p_{o} \left(e^{sg} - 1 \right) \right] + \frac{I \left[W(1-N-D_{o}) - E_{m} + I_{m} \right] \left[\left(1 - p_{o} e^{sg} \right) e^{-\lambda H} + p_{o} e^{sg} \right] \right]}{A_{w}} \right]^{2}} \right] + 2aH - b \right]$$
(17)

There is also replenishment of the seed bank due to seed set from survivors. Therefore equation (16) is only a static response transfer function. It is apparent that any crop-weed ecological modelling exercise must be performed in an iterative way, with the previous state of the weed seed bank influencing the current weed status [12]. The weed seed bank at the start of any cropping cycle, in simplified terms, can be understood as the sum of the dormant seed bank and the seed set from survivors of the previous season's weed management strategies; therefore an iterative approach to weed studies must be adopted [12,13]. This can be approximated by the following equuation:

$$W_{i} = \frac{\left[W_{i-1}\left(1 - N_{i-1} + D_{b}\right) - E_{m} + I_{m}\right] \left[\left(1 - p_{o}e^{sg_{i-1}}\right)e^{-\lambda H} + p_{o}e^{sg_{i-1}}\right] \cdot S_{s}}{1 + e^{-\left(\frac{t-t_{o}}{d}\right)}} + D_{o}W_{i-1}$$
(10)

(18)

Herbicide resistance in many weed species is becoming more prevalent [4,14]. Thornby and Walker [15] simulated continuous summer fallows using glyphosate. Their modelling showed that barnyard grass (*Echinochloa colona*) could become resistant to glyphosate in about 15 years. Validation of their model against paddock history data for glyphosate-resistant population of barnyard grass showed that their model correctly predicted resistance development to within a few years of the real situation.

Selection pressure for genetic traits depends on the initial efficacy of the herbicide to remove susceptible individuals from the population, leaving only the resistant individuals to reproduce. This is reinforced by the adoption of a single herbicide over a long period of time to sustain the selection pressure on the population.

3. METHOD

Equations (16), (17) and (18) were coded into a simple cropping system model using the MatLab software platform. Using data published by Bosnić and Swanton [8] and Yin *et al.* [11] for Rimsulfuron herbicide and assuming: an initially small resistant population ($p_0 = 1 \times 10^{-8}$); an average seed set of 700 seeds per weed plant; a seed mortality rate of 10 % each year; and a slightly positive selection coefficient of (s = 1×10^{-4}) for herbicide resistance [16], the system transfer function was used to analyse the effect of a single herbicide application on crop yield potential. The transfer function was also used to forecast the viable weed seed bank and long term crop yield potential, assuming that only a single herbicide type was used during this time.

4. RESULTS AND DISCUSSION

Figure **1** shows the expected crop response as a function of the herbicide's application rate. There is an optimal application rate of about 0.009 kg ha⁻¹, based



Figure 1: Normalised crop yield (blue line) and rate of change (green line) of crop yield response as a function of applied herbicide energy, based equations (16) and (17).

on these parameters, while the maximum rate of crop yield response (where $\frac{\partial Y}{\partial H} = 0$) occurs at about 0.001 kg ha⁻¹. The transfer function also predicts that significant herbicide resistance will occur within 15 generations (Figure 2), as was also predicted by Thornby and Walker [15]. Herbicide rotations can forestall the development of a resistant population; however several weed species have developed multiple resistance to several herbicide groups [5].

A growing herbicide resistance problem is already evident in most Australian cropping systems [17,18]. There is evidence that glyphosate resistance has already developed in some weed populations [17] and multiple herbicide resistances has been widely reported in several weed species [5,19-21]; therefore significant crop yield (Figure **2**) losses can be expected. Alternative weed management strategies that are compatible with no-till cropping systems need to be developed.

5. CONCLUSION

This paper has developed a cropping system transfer function relating herbicide application to potential crop yield. The transfer function also allows for herbicide resistance in the weed population and can be used to investigate the potential long term implications of herbicide weed control.



Figure 2: Generational impact of herbicide resistant weeds on potential crop yield (blue line) and viable seed bank (green line) under continuous herbicide weed management, based on equations (16) and (18).

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NOMENCLATURE

- A_w Is the percentage yield loss as weed density approaches ∞ (= 38.0 [8])
- c Is the speed of light (m s⁻¹) or the rate at which I approaches zero as t approaches ∞ (= 0.017 [8])
- d Is the slope of the seed bank recruitment curve at $t_{\rm o}$
- D_o Fraction of the seed population developing dormancy (Note: this is expressed as a fraction of the initial seed bank population W_o)
- D_b Fraction of the seed population from previous seasons breaking dormancy (Note: this is expressed as a fraction of the initial seed bank population W_o)
- E_m Seed emigration from the area of interest
- g Is the generational number
- H Is the herbicide dose
- I Is the percentage yield loss as the weed density tends towards zero (= 0.38 [8])
- Im Seed immigration into the area of interest
- N Is the natural death rate for the whole population (Note: this is expressed as a fraction of the initial seed bank population W_o)
- po Is the initial frequency of herbicide resistant plants
- s Is the selection pressure for herbicide resistance
- S_s Viable seed set per plant from surviving volunteers in the weed population
- t Is the time difference between crop emergence and weed emergence
- t_{o} $% t_{\text{o}}$ Is the time for 50 % germination of the viable seed bank

- W Is the viable seed bank
- Y_o Is the theoretical yield with no weed infestations
- λ Is the efficacy of the herbicide killing action

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