Insecticide Resistance Management : prototype game scenarios v3 current version based on emergence

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This document demonstrates how a few simple equations can simulate vector populations and insecticide resistance over time depending on the use of insecticides. Parameters within the simple equations can be modified to generate different patterns.

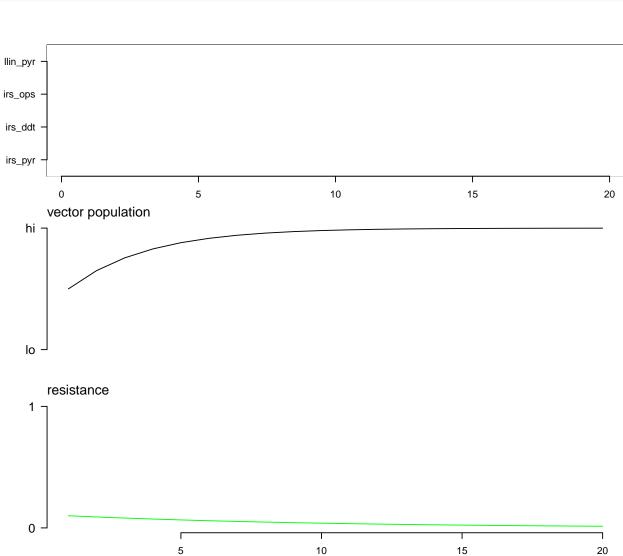
We intend to tweak the input parameters to generate reasonable scenarios. We are looking for assessment of the scenarios themselves rather than the input parameter values. The input parameters are simply a means by which we can generate reasonable scenarios.

In the following plots time is represented on the x axis, the top panel shows insecticide use, the middle panel shows vector population and the lower panel shows resistance (phenotypic). The code included is there merely to show us as developers how the scenarios were generated.

For an interactive version of the equations used to generate these plots see https://andysouth.shinyapps.io/shinyGame4.

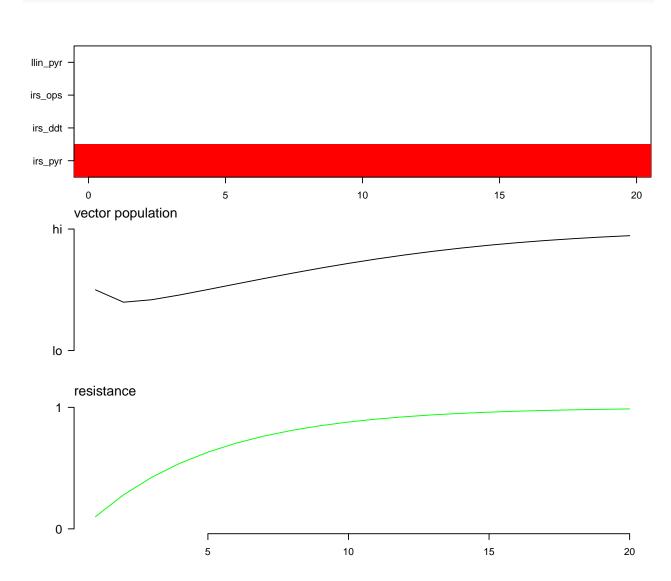
scenario 1 : no insecticide use

plot_sim(run_sim())



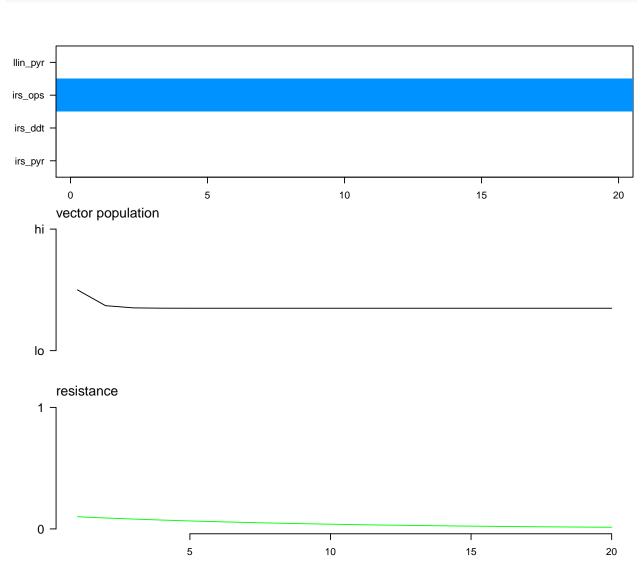
scenario 2: continuous pyr use in presence of resistance

```
l_config <- read_config()
l_config2 <- config_plan(l_config, t_strt=1, t_stop=20, control_id='irs_pyr')
plot_sim( run_sim(l_config=l_config2) )</pre>
```

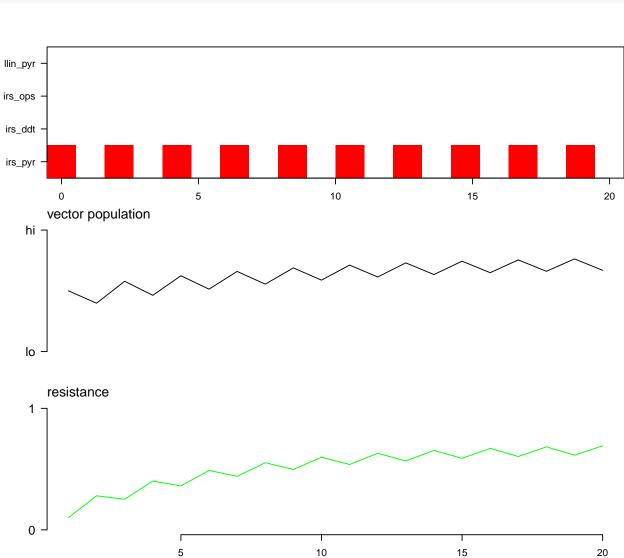


scenario 3: continuous use of ops with no resistance

```
1_config2 <- config_plan(1_config, t_strt=1, t_stop=20, control_id='irs_ops')
plot_sim( run_sim(1_config=1_config2) )</pre>
```

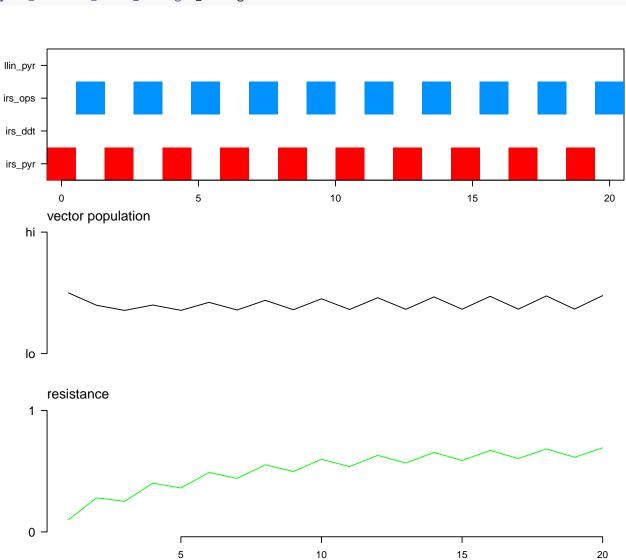


scenario 4 : pyr used in alternate steps

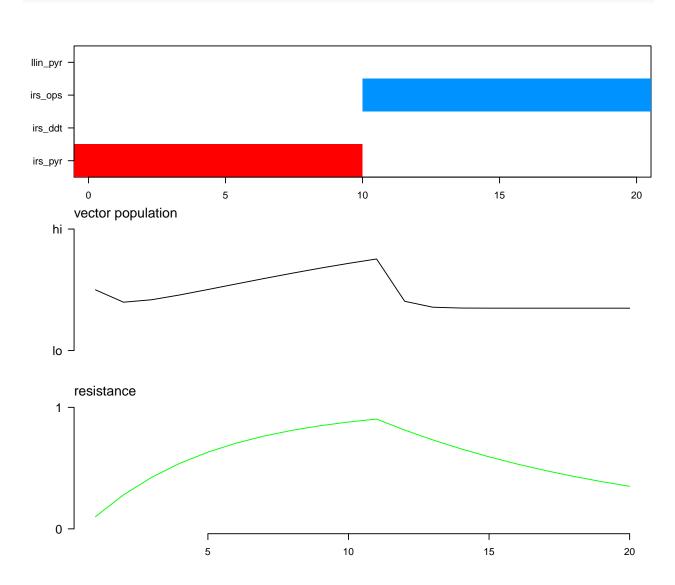


scenario 5: alternate use of ops and pyr

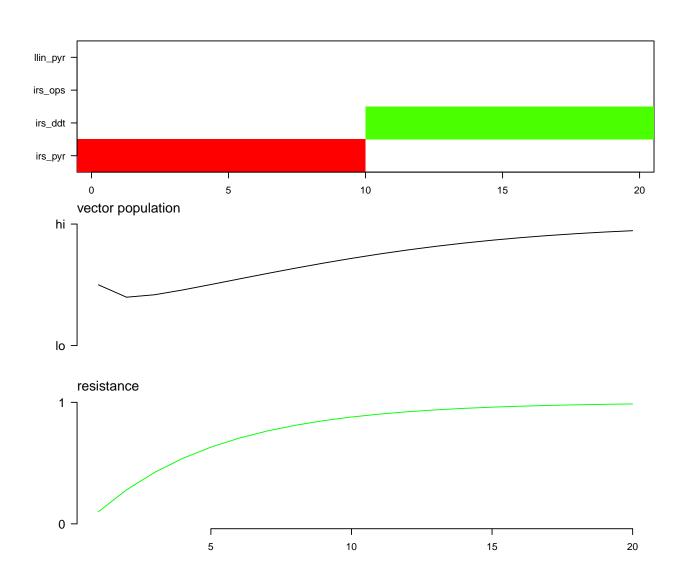
```
l_config2 <- config_plan(l_config, t_strt=seq(1,20), t_stop=seq(1,20), control_id=c('irs_pyr','irs_ops'
plot_sim( run_sim(l_config=l_config2) )</pre>
```



scenario 6 : 10 steps pyr, 10 steps ops

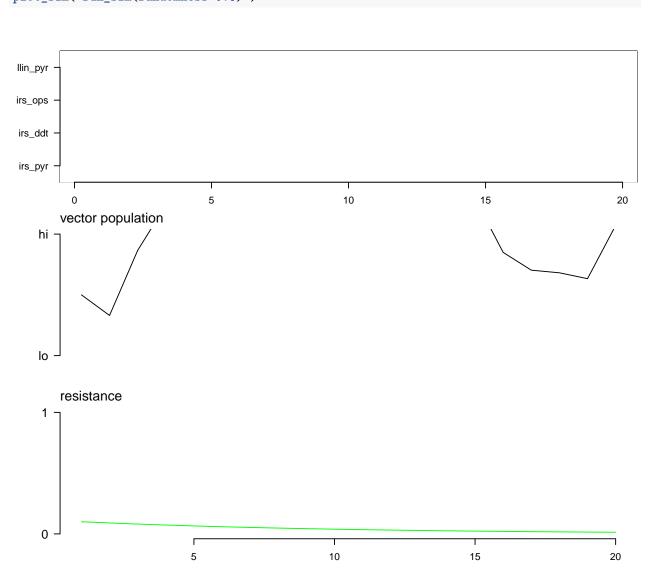


scenario 7:10 steps pyr, 10 steps ddt



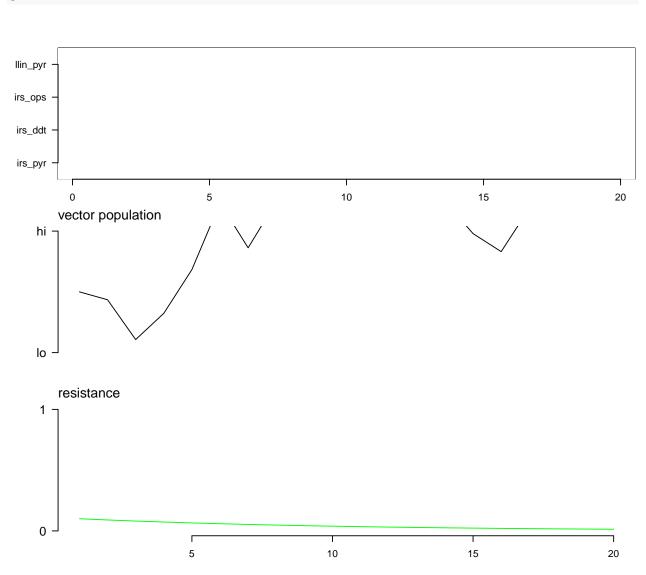
scenario 8 : no insecticide use, 50% randomness added

plot_sim(run_sim(randomness=0.5))



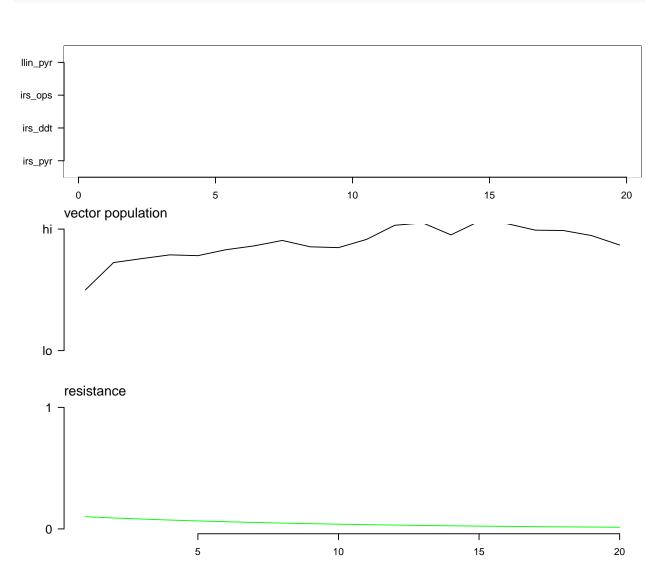
scenario 9 : same as previous but just a different randomisation

plot_sim(run_sim(randomness=0.5))



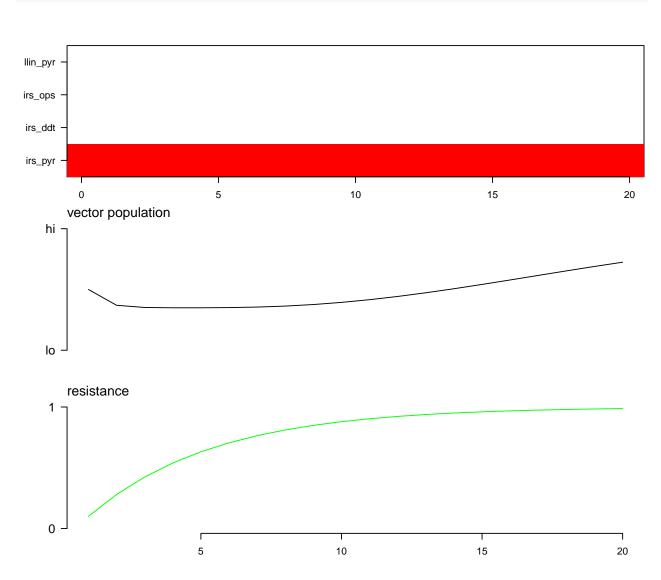
scenario 10 : no insecticide use, 10% randomness added





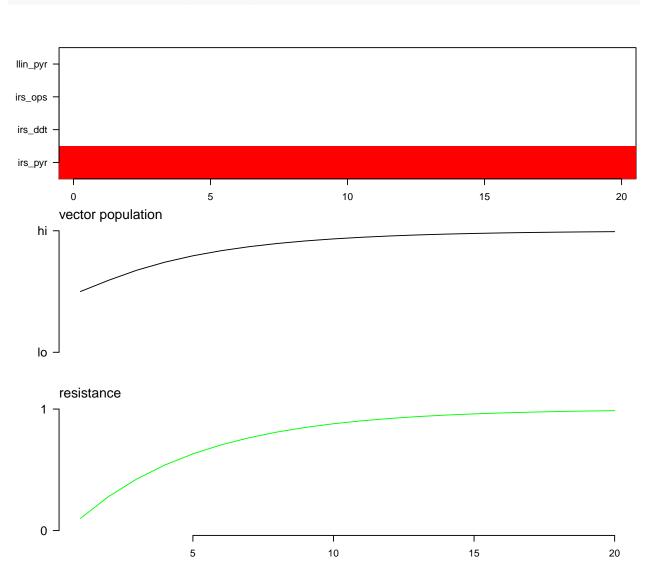
scenario 11 : continuous pyr use, resistance_modifier <1 decreases effect of resistance

```
1_config2 <- config_plan(1_config, t_strt=1, t_stop=20, control_id='irs_pyr')
plot_sim( run_sim(1_config=1_config2, resistance_modifier=0.1) )</pre>
```

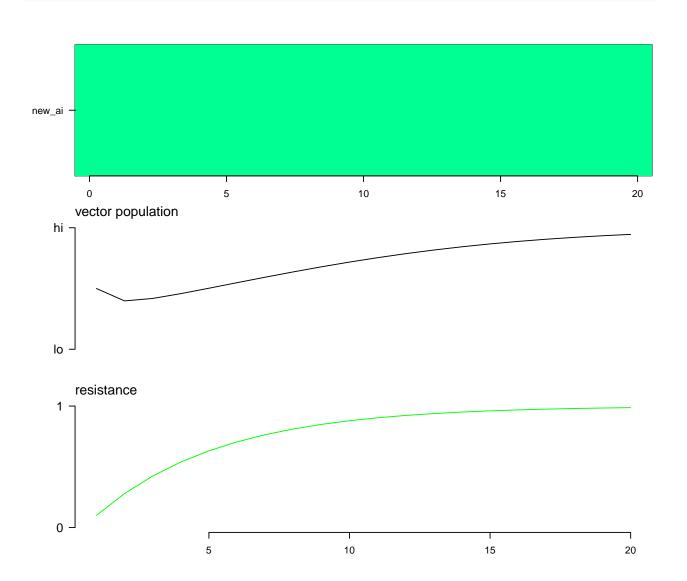


scenario 12 : continuous pyr use, resistance_modifier > 1 increases effect of resistance

plot_sim(run_sim(l_config=l_config2, resistance_modifier=10))

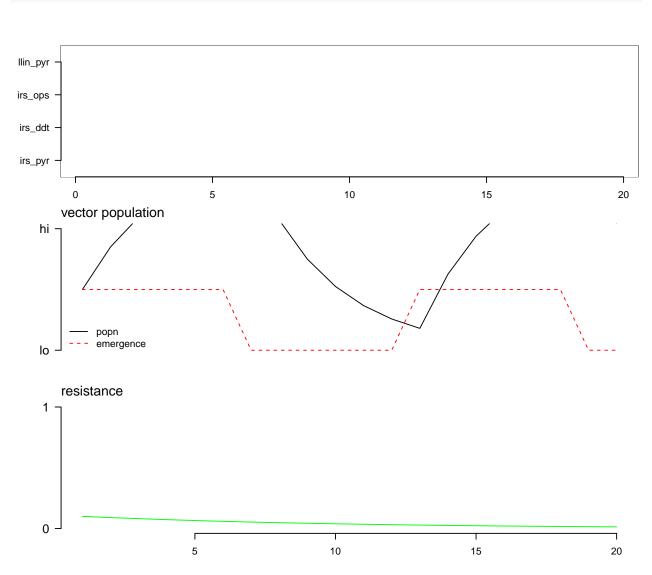


scenario 13: set a different control and resistance mechanism and control plan



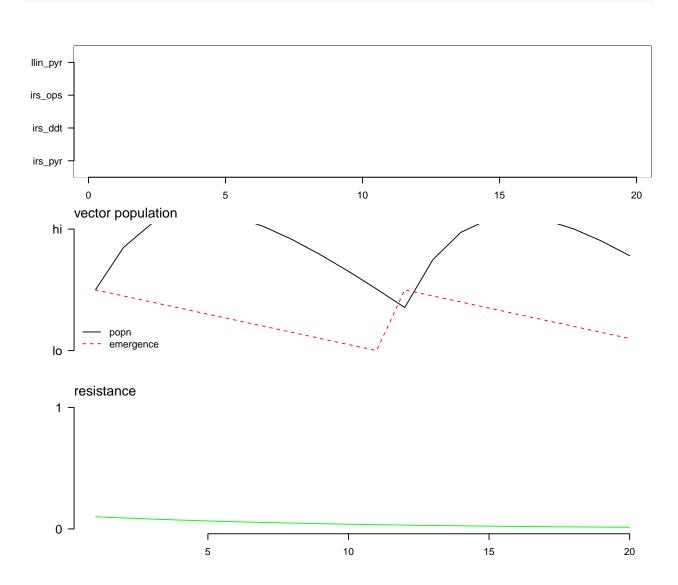
scenario 14: change carrying capacity over time (6 tstep(month) cycle). High growth rate.

```
emergence = c(rep(0.5,6),rep(0,6))
plot_sim( run_sim(emergence=emergence, survival=0.7), plot_emergence=TRUE )
```



scenario 15: carrying capacity gradual change. High growth rate.

```
emergence = seq(0.5,0,-0.05)
plot_sim( run_sim(emergence=emergence, survival=0.7), plot_emergence=TRUE )
```



How controls and resistance mechanisms can be specified.

Our generic approach allows us to specify any combination of controls and resistance mechanisms. The controls cause a specified kill rate(s) on specified vector(s). The resistance mechanisms specify which controls they apply to and how fast resistance increases and decreases in the presence and absence respectively of that control. Cross resistance can be specified simply by specifying multiple controls for one resistance mechanism.

The relationships between vectors, controls and resistance mechanisms are specified in simple configuration files. Here is a simple example of a collection of such configuration files:

places.csv

vectors.csv

```
## vector_id vector_name vector_desc vector_growth
## 1 an_gamb Anopheles gambiae NA 0.4
```

controls.csv

##		control_id	control_name	control_desc	vector_id	control_kill_rate
##	1	irs_pyr	IRS pyrethroid	NA	an_gamb	0.4
##	2	irs_ddt	IRS ddt	NA	an_gamb	0.5
##	3	irs_ops	IRS organophosphates	NA	an_gamb	0.3
##	4	llin_pyr	pyrethroid bednet	NA	an_gamb	0.2

resistances.csv

```
resistance id
                                 resistance name control id
## 1
      met_pyr_ddt metabolic pyrethroids and ddt
                                                     irs_pyr
       met_pyr_ddt metabolic pyrethroids and ddt
                                                    llin pyr
       met_pyr_ddt metabolic pyrethroids and ddt
                                                     irs_ddt
     resistance_strength resistance_incr resistance_decr
                     1.0
                                    0.20
## 1
                                                      0.1
## 2
                     1.0
                                    0.05
                                                      0.1
## 3
                     0.9
                                    0.20
                                                      0.1
```

${\bf control_plan.csv}$