

## Appendix 2 (extension) – Details of Resistance Gene Models

This is an extended version of the "Appendix 2" that appears in the published paper.

We consider (in separate models) two alternative resistance genes; a maternal-effect gene ( $R_m$ ) and a filial-effect gene ( $R_f$ ). The latter is discussed in more detail in the paper and referred to there as  $R$ . Both resistance genes act in the diploid host to reduce male-killing, and are inherited in an autosomal fashion. The resistance genes differ in their sex-dependent effects (Figure 1). The recursion equations for the two models (with two male-killer strains) are given below.

The maternal-effect resistance gene acts in a dominant fashion in the female host to reduce the transmission of male-killers to her eggs, with the proportion of eggs infected reduced from  $\alpha$  to  $d\alpha$  (where  $0 < d < 1$ ). We assume that  $R_m$  imposes a cost  $c$  on females, and that this cost acts multiplicatively so that the fitness of females homozygous for  $R_m$  is  $(1 - c)^2$ .

The filial-effect resistance gene ( $R_f$ ) acts in a dominant fashion in the male host to completely eliminate male-killers from a proportion  $1 - g$  (where  $0 < g < 1$ ) of infected male embryos, so that the proportion of males killed is  $g\alpha$  rather than  $\alpha$ . We assume that  $R_f$  imposes a cost  $c$  on males, and that this cost acts multiplicatively so that the fitness of males homozygous for  $R_f$  is  $(1 - c)^2$ .

### Mathematical Details of the Models

The genotype frequencies are represented as follows:

males:  $y$ , females:  $x$

	No MK	MK <sub>1</sub> (females only)	MK <sub>2</sub> (females only)
No $R$ gene	$y_1, x_1$	$x_4$	$x_7$
Heterozygous for $R$ gene	$y_2, x_2$	$x_5$	$x_8$
Homozygous for $R$ gene	$y_3, x_3$	$x_6$	$x_9$

### Invasion of the Maternal-Effect Resistance Gene

Recursion equations for the dynamics of a two male-killer system with the maternal-effect resistance gene were obtained by following the scheme in Figure 1. It is necessary to carry out recursions on individuals because fitness compensation affects broods as a whole.

Females in the next generation then suffer the viability costs of carrying the resistance gene and the male-killer, dependent on their own genotype.



	$y_1$	$y_2$	$y_3$
$x_1$	$x_1 + y_1$	$\frac{1}{2}(x_1 + y_1 + y_2 + x_2)$	$y_2 + x_2$
$x_2$	$\frac{1}{2}(x_1 + y_1 + y_2 + x_2)$	$\frac{1}{4}(x_1 + y_1 + 2y_2 + 2x_2 + y_3 + x_3)$	$\frac{1}{2}(y_2 + x_2 + y_3 + x_3)$
$x_3$	$y_2 + x_2$	$\frac{1}{2}(y_2 + x_2 + y_3 + x_3)$	$y_3 + x_3$
$x_4$	$x_1(1-\alpha_1) + y_1(1-\alpha_1) + x_4\alpha_1$	$\frac{1}{2}((1-\alpha_1)(x_1 + y_1 + y_2 + x_2) + \alpha_1(x_4 + x_5))$	$y_2(1-\alpha_1) + x_2(1-\alpha_1) + x_5\alpha_1$
$x_5$	$\frac{1}{2}((1-d\alpha_1)(x_1 + y_1 + y_2 + x_2) + d\alpha_1(x_4 + x_5))$	$\frac{1}{4}((1-d\alpha_1)(x_1 + y_1 + 2y_2 + y_3 + 2x_2 + x_3) + d\alpha_1(x_4 + 2x_5 + x_6))$	$\frac{1}{2}((1-d\alpha_1)(x_2 + y_2 + y_3 + x_3) + d\alpha_1(x_5 + x_6))$
$x_6$	$y_2(1-d\alpha_1) + x_2(1-d\alpha_1) + x_5d\alpha_1$	$\frac{1}{2}((1-d\alpha_1)(x_2 + y_2 + y_3 + x_3) + d\alpha_1(x_5 + x_6))$	$y_3(1-d\alpha_1) + x_3(1-d\alpha_1) + x_6d\alpha_1$
$x_7$	$x_1(1-\alpha_2) + y_1(1-\alpha_2) + x_7\alpha_2$	$\frac{1}{2}((1-\alpha_2)(x_1 + y_1 + y_2 + x_2) + \alpha_2(x_7 + x_8))$	$y_2(1-\alpha_2) + x_2(1-\alpha_2) + x_8\alpha_2$
$x_8$	$\frac{1}{2}((1-\alpha_2)(x_1 + y_1 + y_2 + x_2) + \alpha_2(x_7 + x_8))$	$\frac{1}{4}((1-\alpha_2)(x_1 + y_1 + 2y_2 + 2x_2 + y_3 + x_3) + \alpha_2(x_7 + 2x_8 + x_9))$	$\frac{1}{2}((1-\alpha_2)(x_2 + y_2 + y_3 + x_3) + \alpha_2(x_8 + x_9))$
$x_9$	$y_2(1-\alpha_2) + x_2(1-\alpha_2) + x_8\alpha_2$	$\frac{1}{2}((1-\alpha_2)(x_2 + y_2 + y_3 + x_3) + \alpha_2(x_8 + x_9))$	$y_3(1-\alpha_2) + x_3(1-\alpha_2) + x_9\alpha_2$

Male death occurs in broods from mothers  $x_4 - x_9$ . In each case, the surviving brood members receive fitness compensation depending on the mother's genotype. Hence, the number of progeny in each row of the table above must be multiplied by the appropriate fitness compensation term:

Maternal Genotype	Appropriate fitness compensation term	Abbreviation in invasion term
$x_4$	$1 + \frac{\phi\alpha_1}{2 - \alpha_1}$	$f^4$
$x_5, x_6$	$1 + \frac{\phi d\alpha_1}{2 - d\alpha_1}$	$f^5$
$x_7, x_8, x_9$	$1 + \frac{\phi\alpha_2}{2 - \alpha_2}$	$NA$

Females in the next generation then suffer the viability costs of carrying the resistance gene and the male-killer, dependent on their own genotype.



Female Genotype	Viability costs
$x_1$	-
$x_2$	$(1-c)$
$x_3$	$(1-c)^2$
$x_4$	$(1-U_1)$
$x_5$	$(1-U_1)(1-c)$
$x_6$	$(1-U_1)(1-c)^2$
$x_7$	$(1-U_2)$
$x_8$	$(1-U_2)(1-c)$
$x_9$	$(1-U_2)(1-c)^2$

The invasion conditions for  $R_m$  in the presence of  $MK_1$  only were found by modifier analysis. Linearised recursions were obtained for the three genotypes in which  $R_f$  is heterozygous and  $MK_2$  is absent ( $y_2, x_2, x_5$ ).  $x_1, y_1$  and  $x_4$  were taken to be at their equilibrium frequencies in the absence of  $R_f$ . In matrix form, the linearised recursions become:

$$\begin{bmatrix} y_2' \\ x_2' \\ x_5' \end{bmatrix} = \begin{bmatrix} a & b & c \\ d & e & f \\ g & h & i \end{bmatrix} \begin{bmatrix} y_2 \\ x_2 \\ x_5 \end{bmatrix} = \lambda \begin{bmatrix} y_2 \\ x_2 \\ x_5 \end{bmatrix}$$

Hence for invasion, the leading eigenvalue,  $\lambda$  of the resultant 3x3 matrix must be greater than one. It therefore follows that:

$$1 < a + bd + e - ae + cg - ceg + bfg + cdh + fh - afh + i - ai - bdi - ei + aei$$

This revealed that for invasion:

$$1 < (-4 + \alpha_1 c^2 d f_5 (U_1 - 1) (p^* - 1 - p^* f_4 + \alpha_1 p^* f_4) + 2 p^{*3} (1 - f_4 + \alpha_1 f_4) (1 - f_4 + \alpha_1 f_4 U_1)^2 + p^{*2} (1 - f_4 + \alpha_1 f_4 U_1) (8 f_4 - 8 - 6 \alpha_1 f_4 - \alpha_1 d f_5 + \alpha_1 f_4 f_5 + \alpha_1 d f_4 f_5 - 2 \alpha_1^2 d f_4 f_5 - 2 \alpha_1 f_4 U_1 + \alpha_1 d f_5 U_1 - \alpha_1 f_4 f_5 U_1 - \alpha_1 d f_4 f_5 U_1 + 2 \alpha_1^2 d f_4 f_5 U_1) + p^* (10 - 10 f_4 + 4 \alpha_1 f_4 + \alpha_1 d f_5 - \alpha_1 f_4 f_5 - \alpha_1 d f_4 f_5 + \alpha_1^2 d f_4 f_5 + 6 \alpha_1 f_4 U_1 - \alpha_1 d f_5 U_1 + \alpha_1 f_4 f_5 U_1 + \alpha_1 d f_4 f_5 U_1 - \alpha_1^2 d f_4 f_5 U_1^2) + c (2 - \alpha_1 d f_5 + \alpha_1 d f_5 U_1 - p^{*2} (1 - f_4 + \alpha_1 f_4 U_1) (2 f_4 - 2 - 2 \alpha_1 f_4 - \alpha_1 d f_5 + \alpha_1 f_4 f_5 + \alpha_1 d f_4 f_5 - 2 \alpha_1^2 d f_4 f_5 + \alpha_1 d f_5 U_1 - \alpha_1 f_4 f_5 U_1 - \alpha_1 d f_4 f_5 U_1 + 2 \alpha_1^2 d f_4 f_5 U_1) + p^* (4 f_4 - 4 - 2 \alpha_1 f_4 + \alpha_1 f_4 f_5 - 2 \alpha_1 f_4 U_1 - \alpha_1 f_4 f_5 U_1 - \alpha_1^2 d f_4 f_5 U_1 + \alpha_1^2 d f_4 f_5 U_1^2))) / (4(p^* - 1 - p^* f_4 + \alpha_1 p^* f_4) (p^* - 1 - p^* f_4 + \alpha_1 p^* f_4 U_1)^2)$$

The conditions for invasion of  $R_m$  when  $\phi = 0.5$  and  $U_1 = 0.01$  are plotted in Figure 2.



## Invasion of The Filial-Effect Resistance Gene

Recursion equations for the dynamics of a two male-killer system with the filial-effect resistance gene:

	$y_1$	$y_2$	$y_3$
$x_1$	$y_1 + x_1$	$\frac{1}{2} (y_1 + y_2 + x_1 + x_2)$	$y_2 + x_2$
$x_2$	$\frac{1}{2} (y_1 + y_2 + x_1 + x_2)$	$\frac{1}{4} (1 + y_2 + x_1 + 2x_2 + x_3)$	$\frac{1}{2} (y_2 + y_3 + x_2 + x_3)$
$x_3$	$y_2 + x_2$	$\frac{1}{2} (y_2 + y_3 + x_2 + x_3)$	$y_3 + x_3$
$x_4$	$y_1(1-\alpha_1) + x_4\alpha_1 + x_1(1-\alpha_1)$	$\frac{1}{2} ((1-\alpha_1)(y_1 + x_1 + x_2) + \alpha_1(x_4 + x_5) + y_2(1-g\alpha_1))$	$y_2(1-g\alpha_1) + x_5\alpha_1 + x_2(1-\alpha_1)$
$x_5$	$\frac{1}{2} ((1-\alpha_1)(y_1 + x_1 + x_2) + \alpha_1(x_4 + x_5) + y_2(1-g\alpha_1))$	$\frac{1}{4} ((1-\alpha_1)(y_1 + x_1 + 2x_2 + x_3) + (1-g\alpha_1)(2y_2 + y_3) + \alpha_1(x_4 + 2x_5 + x_6))$	$\frac{1}{2} ((1-\alpha_1)(x_2 + x_3) + \alpha_1(x_5 + x_6) + (1-g\alpha_1)(y_2 + y_3))$
$x_6$	$y_2(1-g\alpha_1) + x_5\alpha_1 + x_2(1-\alpha_1)$	$\frac{1}{2} ((1-g\alpha_1)(y_2 + y_3) + (1-\alpha_1)(x_2 + x_3) + \alpha_1(x_5 + x_6))$	$y_3(1-g\alpha_1) + x_6\alpha_1 + x_3(1-\alpha_1)$
$x_7$	$y_1(1-\alpha_2) + x_7\alpha_2 + x_1(1-\alpha_2)$	$\frac{1}{2} ((1-\alpha_2)(y_1 + y_2) + \alpha_2(x_7 + x_8) + (1-\alpha_2)(x_1 + x_2))$	$y_2(1-\alpha_2) + x_8\alpha_2 + x_2(1-\alpha_2)$
$x_8$	$\frac{1}{2} ((1-\alpha_2)(y_1 + y_2 + x_1 + x_2) + \alpha_2(x_7 + x_8))$	$\frac{1}{4} ((1-\alpha_2)(1 + y_2 + x_1 + 2x_2 + x_3) + \alpha_2(x_7 + 2x_8 + x_9))$	$\frac{1}{2} ((1-\alpha_2)(y_2 + y_3 + x_2 + x_3) + \alpha_2(x_8 + x_9))$
$x_9$	$y_2(1-\alpha_2) + x_8\alpha_2 + x_2(1-\alpha_2)$	$\frac{1}{2} ((1-\alpha_2)(y_2 + y_3 + x_2 + x_3) + \alpha_2(x_8 + x_9))$	$y_3(1-\alpha_2) + x_9\alpha_2 + x_3(1-\alpha_2)$

Again, male death occurs in broods from mothers  $x_4 - x_9$ , but in this case, the number of males that die is not solely dependent on the mother's genotype. Infected males have a chance of surviving infection if they carry  $R_f$ , so the proportion of males dying is also affected by the father's genotype. The fitness compensation received by the brood therefore depends on the combination of the father and mother's genotype.

The conditions for invasion of  $R_f$  when  $\beta = 0.5$  and  $U_f = 0.01$  are plotted in Figure 3.



Parental Genotypes	Appropriate fitness compensation term	Abbreviation in invasion term
$y_1 x_4$	$1 + \frac{\phi\alpha_1}{2 - \alpha_1}$	$f1$
$y_1 x_6, y_2 x_6, y_3 x_4, y_3 x_5, y_3 x_6$	$1 + \frac{\phi g\alpha_1}{2 - g\alpha_1}$	NA
$y_2 x_5$	$1 + \frac{(0.75g\alpha_1 + 0.25\alpha_1)\phi}{2 - (0.75g\alpha_1 + 0.25\alpha_1)}$	NA
$y_1 x_5, y_2 x_4$	$1 + \frac{(0.5g\alpha_1 + 0.5\alpha_1)\phi}{2 - (0.5g\alpha_1 + 0.5\alpha_1)}$	$f5$
All matings involving $x_7, x_8$ and $x_9$	$1 + \frac{\phi\alpha_2}{2 - \alpha_2}$	NA

Females in the next generation then suffer viability costs dependent on their genotype. In females it is the cost imposed by the male-killer, while males suffer the viability cost of  $R_f$ .

Genotype	Viability costs
$x_1 x_2 x_3 y_1$	-
$y_2$	$(1-c)$
$y_3$	$(1-c)^2$
$x_4 x_5 x_6$	$(1-U_1)$
$x_7 x_8 x_9$	$(1-U_2)$

Invasion conditions for  $R_f$  were obtained by modifier analysis in the same way as for  $R_m$ .

This revealed that for invasion:

$$1 < (-4 + 2p^*^3(1 - f5 + \alpha_1 g f5)(1 - f1 + \alpha_1 f1 U_1)^2 + p^*^2(1 - f1 + \alpha_1 f1 U_1)(4f1 - 8 - 2\alpha_1 f1 + 4f5 - 2\alpha_1 f5 + 2\alpha_1 f5 f1 - 2\alpha_1^2 f5 f1 - 3\alpha_1 f5 g - 2\alpha_1 f1 U_1 + \alpha_1 f5 U_1 - 2\alpha_1 f1 f5 U_1 + 2\alpha_1^2 f1 f5 U_1) + p^*(10 - 8f1 + 2\alpha_1 f1 - 2f5 + 2\alpha_1 f5 - 2\alpha_1 f1 f5 + \alpha_1^2 f1 f5 + \alpha_1 f5 g + 6\alpha_1 f1 U_1 - \alpha_1 f5 U_1 + 2\alpha_1 f1 f5 U_1 - \alpha_1^2 f1 f5 U_1^2) - c(p^* - 1 - p^* f1 + \alpha_1 p^* f1 U_1)(2 - \alpha_1 f5 + \alpha_1 f5 U_1 + 2p^*^2(1 - f5 + \alpha_1 f5 g)(1 - f1 + \alpha_1 f1 U_1) - p^*(4 - 2f1 - 2f5 + \alpha_1 f5 g + 2\alpha_1 f1 U_1 + \alpha_1 f5 U_1))) / (4(p^* - 1 - p^* f1 + \alpha_1 p^* f1)(p^* - 1 - p^* f1 + \alpha_1 p^* f1 U_1)^2)$$

The conditions for invasion of  $R_f$  when  $\phi = 0.5$  and  $U_1 = 0.01$  are plotted in Figure 3.