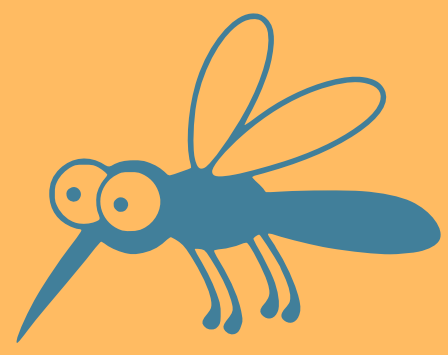
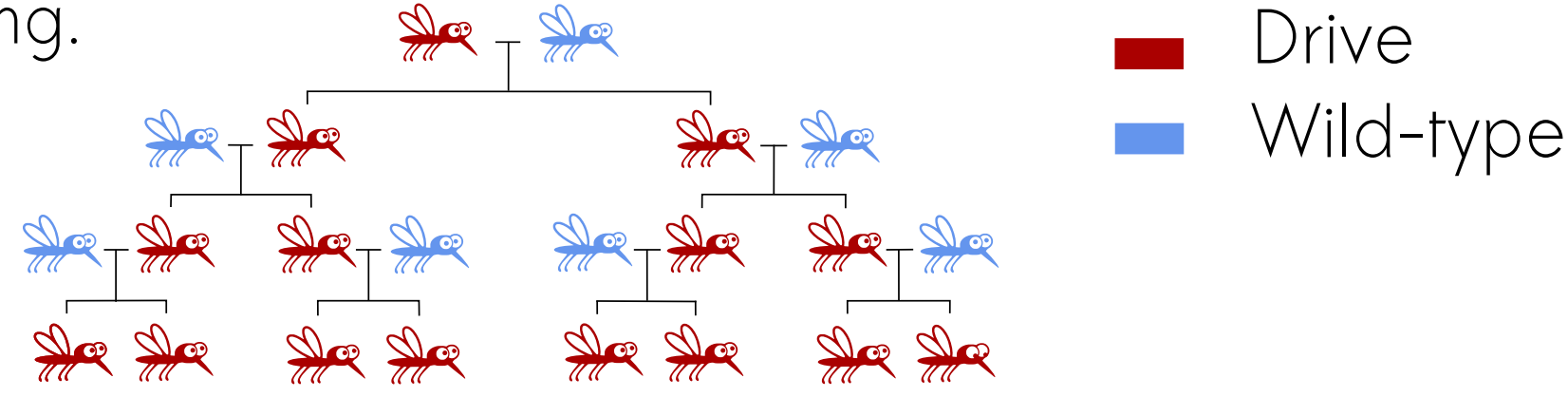


# The importance of demography in gene drive mathematical models.



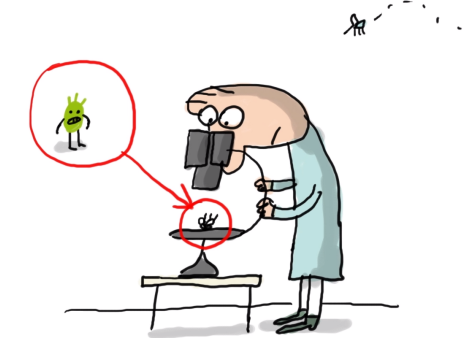
## FIRST: WHAT IS GENE DRIVE?

Gene drive is a genetic technology that biases the transmission of a targeted mutation by ensuring quasi-systematic inheritance to the offspring.



Example of a drive rapide propagation

Applications are diverse and numerous: eradicate insect-borne diseases, control invasive or pest species...



Develop mosquitoes resistance to Malaria

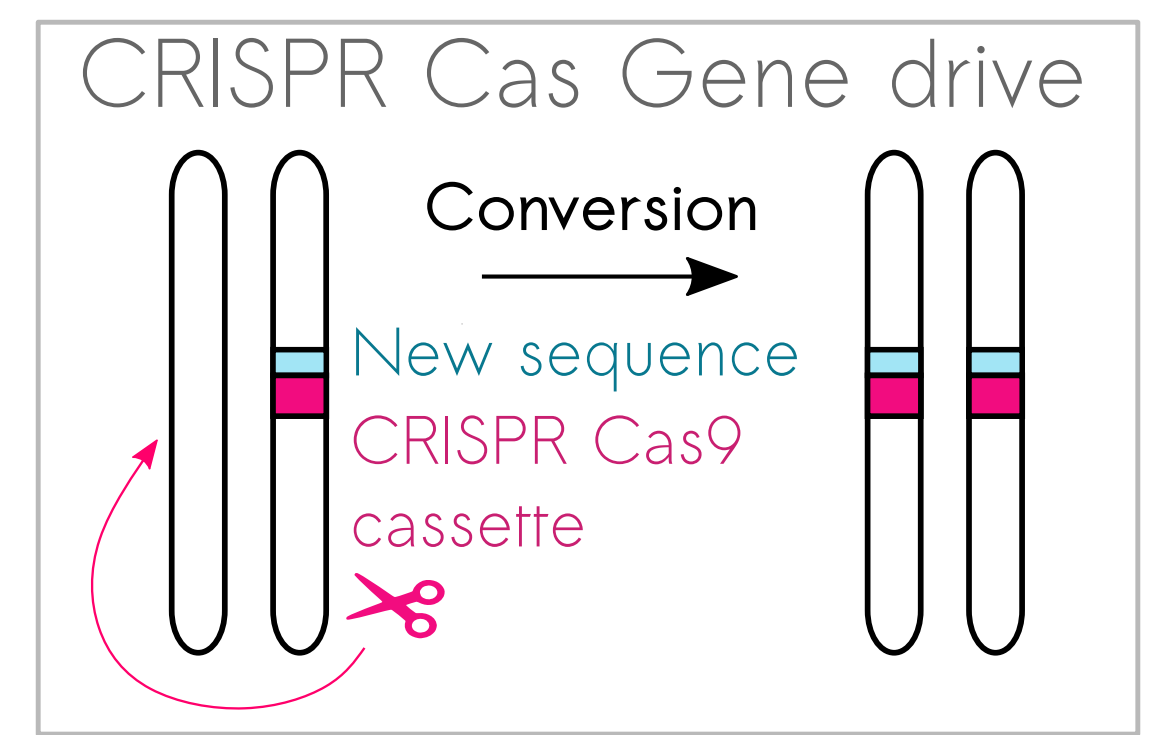


Preserve islands biodiversity



Eliminate wild resistance to herbicides

## HOW DOES IT WORK?



## MODEL WITH EARLY PERFECT CONVERSION

We study the propagation of gene drive with spatio-temporal models. The first one (below) considers perfect conversion in the zygote, resulting in the absence of heterozygous individuals.  $n_D(t,x)$ : wild-type density,  $n_W(t,x)$ : drive density,  $r$ : intrinsic growth rate,  $s$ : fitness disadvantage for drive.

$$\begin{aligned}
 & \text{Wild-type: } \partial_t n_W - \partial_{xx} n_W = 1 \left( r (1 - n_D - n_W) + 1 \right) \frac{n_W^2}{n_W + n_D} - n_W \\
 & \text{Drive: } \partial_t n_D - \partial_{xx} n_D = (1-s) \left( r (1 - n_D - n_W) + 1 \right) \frac{n_D^2 + 2n_D n_W}{n_W + n_D} - n_D
 \end{aligned}$$

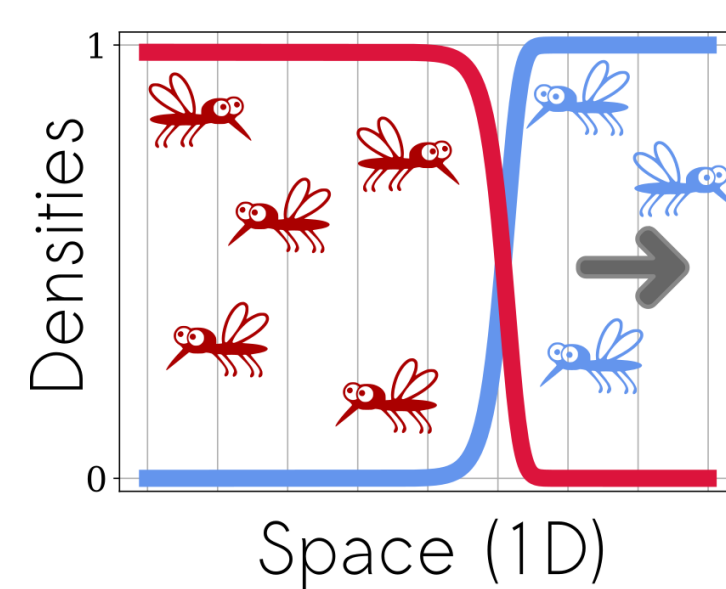
Variations in time:  $\partial_t$ , Diffusion:  $\partial_{xx}$ , Fitness Disadvantage:  $(1-s)$ , Growth term:  $r(1 - n_D - n_W) + 1$ , Mating Advantage (conversion):  $\frac{n_D^2 + 2n_D n_W}{n_W + n_D}$ , Mortality:  $-n_D$

## How important is it to consider demography?

We can rewrite the system above with  $n = n_W + n_D$  and  $p = n_D / (n_W + n_D)$ :

$$\begin{cases}
 \partial_t p - \partial_{xx} p - 2 \partial_x (\log n) \cdot \partial_x p = \left( r (1 - n) + 1 \right) s p (1 - p) \left( p - \frac{2s-1}{s} \right) \\
 \partial_t n - \partial_{xx} n = \left( r (1 - n) + 1 \right) \left( 1 - s + s(1 - p)^2 \right) n - n
 \end{cases}$$

Most gene drive models in the literature focus on the proportion (only one equation on  $p$ ) and assume a constant population size  $n$ . We wonder if this simplification is justified. To study this question we focus on travelling waves solutions (drawing on the right) and compare them over three criteria (see right column) for two values of the demographic parameter  $r$ . We observe significant differences, attesting the importance of demography.



## MODEL WITH EARLY OR LATE PARTIAL CONVERSION

In a second part, we consider a more realistic model: the conversion is not systematic anymore and takes place in the zygote or in the germline. The conversion rate  $c$  is between 0 and 1. These changes result in two models more complex than the previous one, containing 3 equations as we now need to follow heterozygous individuals. The fitness of heterozygotes is  $1-sh$ .

Early partial conversion occurring (in the zygote)

$$\begin{cases}
 \partial_t n_D - \partial_{xx} n_D = (1-s)(r(1-n)+1) \frac{c n_W n_H + 2 c n_W n_D + (0.5 c + 0.25) n_H^2 + (c+1) n_H n_D + n_D^2}{n} - n_D \\
 \partial_t n_H - \partial_{xx} n_H = (1-sh)(r(1-n)+1) \frac{(1-c) n_W n_H + 2 n_W n_D + 0.5 n_H^2 + n_H n_D}{n} - n_H \\
 \partial_t n_W - \partial_{xx} n_W = (r(1-n)+1) \frac{n_W^2 + n_W n_H + 0.25 n_H^2}{n} - n_W
 \end{cases}$$

First case  $(1-c)(1-h) < 1/2$ : drive invades, bistability or wild-type invades depending on  $s$ .  
 Second case  $(1-c)(1-h) > 1/2$ : drive invades, coexistence or wild-type invades depending on  $s$ .

Late partial conversion occurring (in the germline)

$$\begin{cases}
 \partial_t n_D - \partial_{xx} n_D = (1-s)(r(1-n)+1) \frac{0.25(1+c)^2 n_H^2 + (1+c) n_H n_D + n_D^2}{n} - n_D \\
 \partial_t n_H - \partial_{xx} n_H = (1-sh)(r(1-n)+1) \frac{(1+c) n_W n_H + 2 n_W n_D + 0.5(1-c^2) n_H^2 + (1-c) n_H n_D}{n} - n_H \\
 \partial_t n_W - \partial_{xx} n_W = (r(1-n)+1) \frac{n_W^2 + (1-c) n_W n_H + 0.25(1-c)^2 n_H^2}{n} - n_W
 \end{cases}$$

... with  $n = n_D + n_H + n_W$ .

First case  $h > 1/2$ : drive invades, bistability or wild-type invades depending on  $s$ .  
 Second case  $h < 1/2$ : drive invades, coexistence or wild-type invades, depending on  $s$ .

As the models are complex to study, we are searching for conditions implying a pulled monostable wave (of which we know the speed) again for  $r \rightarrow 0$  and  $r \rightarrow +\infty$ . Surprisingly, in the second case, the speed of the drive wave always seems independent of  $r$ , even if the final equilibrium states differ.

## CONCLUSION

- Taking into account the variations in population size can drastically change the conclusion (which type invades, at which speed, etc...). It also indicates us if there will be an extinction event.
- However under certain conditions (see the second results box), the growth rate  $r$  does not seem to impact the wave: similar waves are leading to totally different densities in the end.

- Final equilibrium**  
What is left in the environment?
- Monostable or bistable wave**  
One or two stable steady state(s)?
- Pulled or pushed wave, and speed?**  
Which individuals drive the movement?

## RESULTS

$r \rightarrow +\infty$  Model without demography<sup>(a)</sup>

$$\partial_t p - \partial_{xx} p = \frac{s p (1-p) (p - \frac{2s-1}{s})}{1-s+s(1-p)^2}$$

	$0 < s < 0.36$	$0.36 < s < 1/2$	$1/2 < s < 0.70$	$0.70 < s < 1$
1)	Drive invasion	Drive invasion	Drive invasion	WT invasion
2)	Monostable	Monostable	Bistable	Bistable
3)	Pulled wave	Pulled wave	Pushed wave	Pushed wave

$r \rightarrow 0$  Nearly an SI epidemiological model<sup>(b)</sup>

$$\begin{cases}
 \partial_t p - \partial_{xx} p - 2 \partial_x (\log n) \cdot \partial_x p = s p (1-p) \left( p - \frac{2s-1}{s} \right) \\
 \partial_t n - \partial_{xx} n = s p (p-2) n
 \end{cases}$$

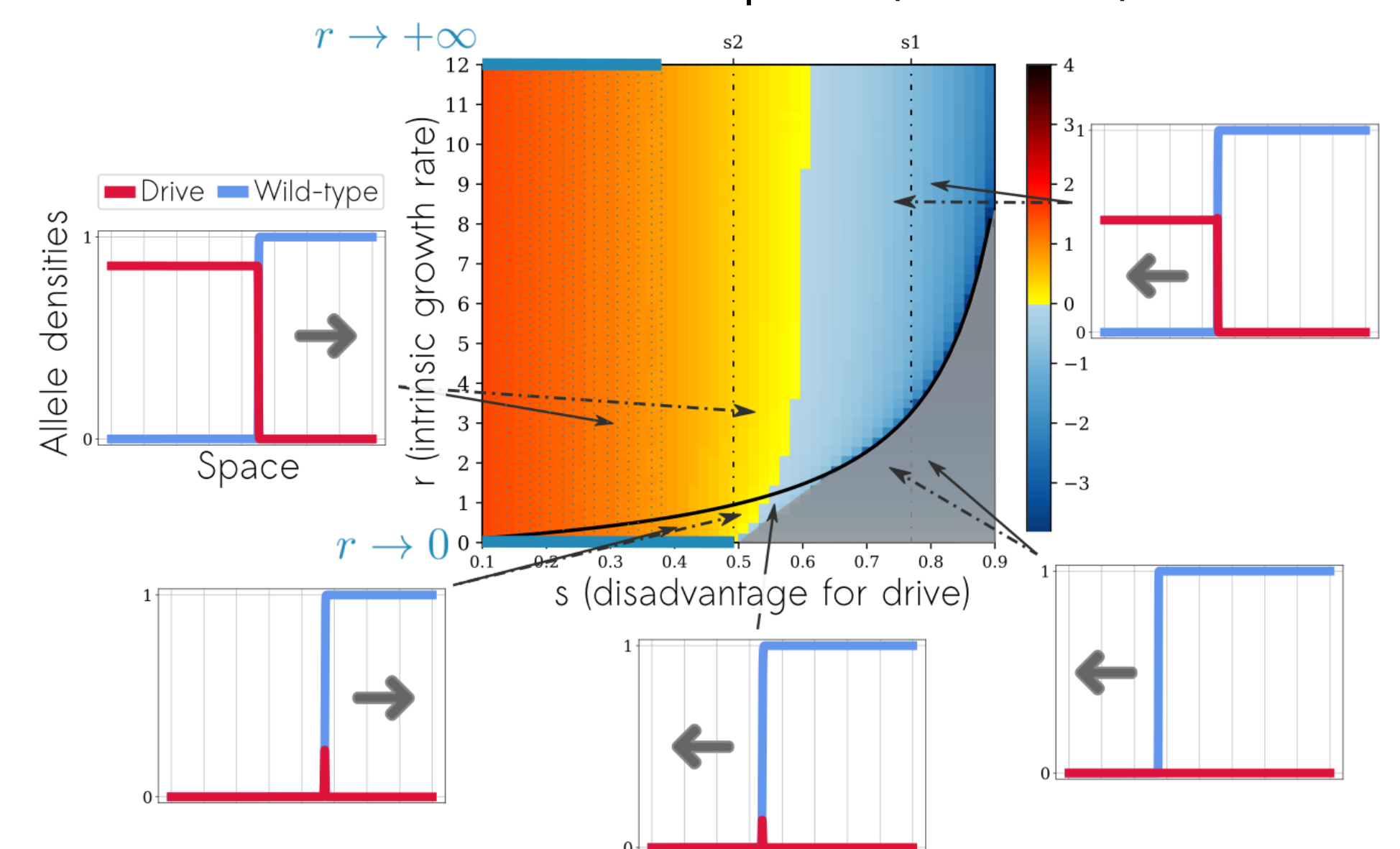
	$0 < s < 0.36$	$0.36 < s < 1/2$	$1/2 < s < 0.70$	$0.70 < s < 1$
1)	Drive invasion	Drive invasion	No invasion	No invasion
2)	Monostable	Monostable	Continuum	Continuum
3)	Pulled wave	Pulled wave	No wave	No wave

(a) M. Strugarek and N. Vauchele "Reduction to a Single Closed Equation for 2-by-2 Reaction-Diffusion Systems of Lotka-Volterra Type". In: *SIAM Journal on Applied Mathematics* 76.5 (Jan. 2016), pp.2060-2080.  
 (b) J. Zhou et al. "Critical Traveling Waves in a Diffusive Disease Model". In: *Journal of Mathematical Analysis and Applications* 476.2 (Aug. 2019), pp.522-538.

## RESULTS

The conditions implying a pulled monostable wave for  $r \rightarrow 0$  and  $r \rightarrow +\infty$  are colored in blue here below.

Drive wave speed (first case)



Drive wave speed (second case)

