

Gene technology and its potential for pest control

Prof Paul Thomas

Director

Genome Editing Program

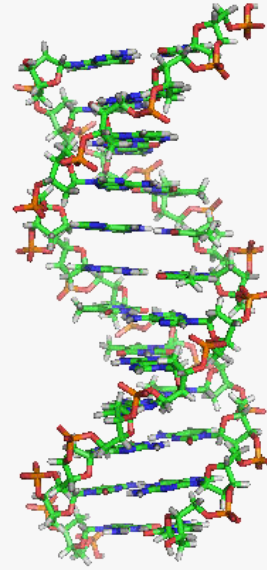
SA Genome Editing Facility



THE UNIVERSITY
of ADELAIDE



Overview



1. Genetic biocontrol (gene drives)

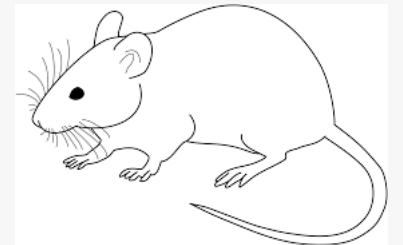
-what are they?

-how do they work (CRISPR)

2. In what species have gene drives been developed?

-invertebrates

-mammals



3. Could gene drives be developed in cats?

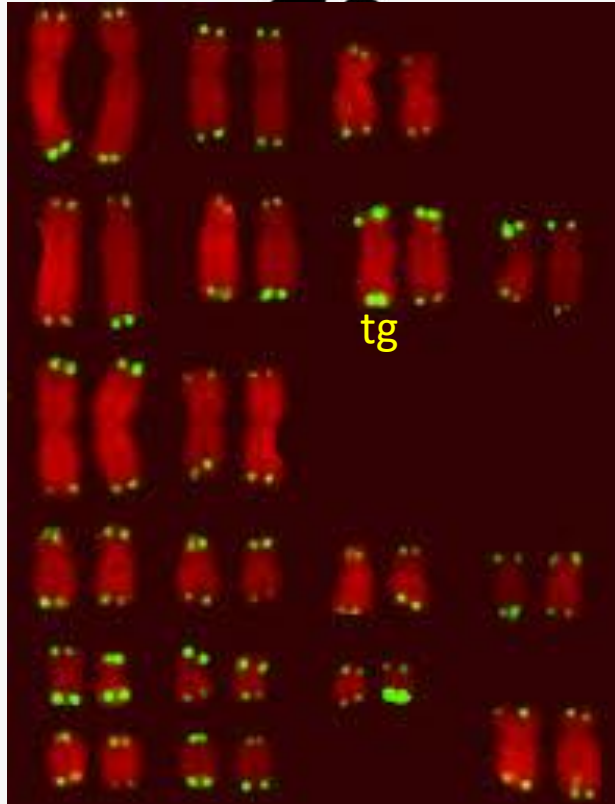
-potential for suppression (modelling)

-challenge/barriers



Genetics and transgenic animals 101

Cat genome on 19 pairs of chromosomes



“Transgenic” cat



Bbc.com

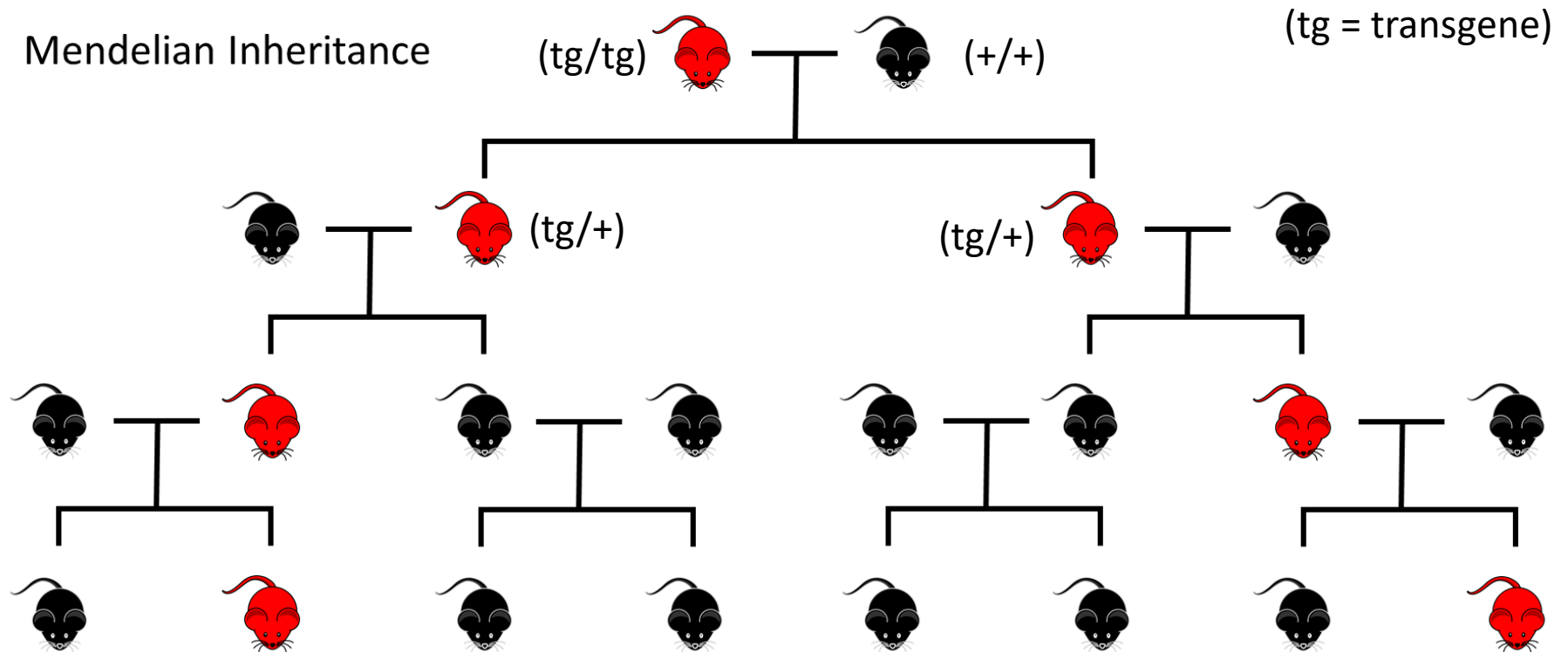
Every cell contains the blueprint for life.... (2.4) billions of DNA building blocks → 20,000 genes

Transgenesis is adding a “foreign” gene into the genome → new “phenotype”

(synthetic) gene drives are a type of transgenic animal which have biased inheritance

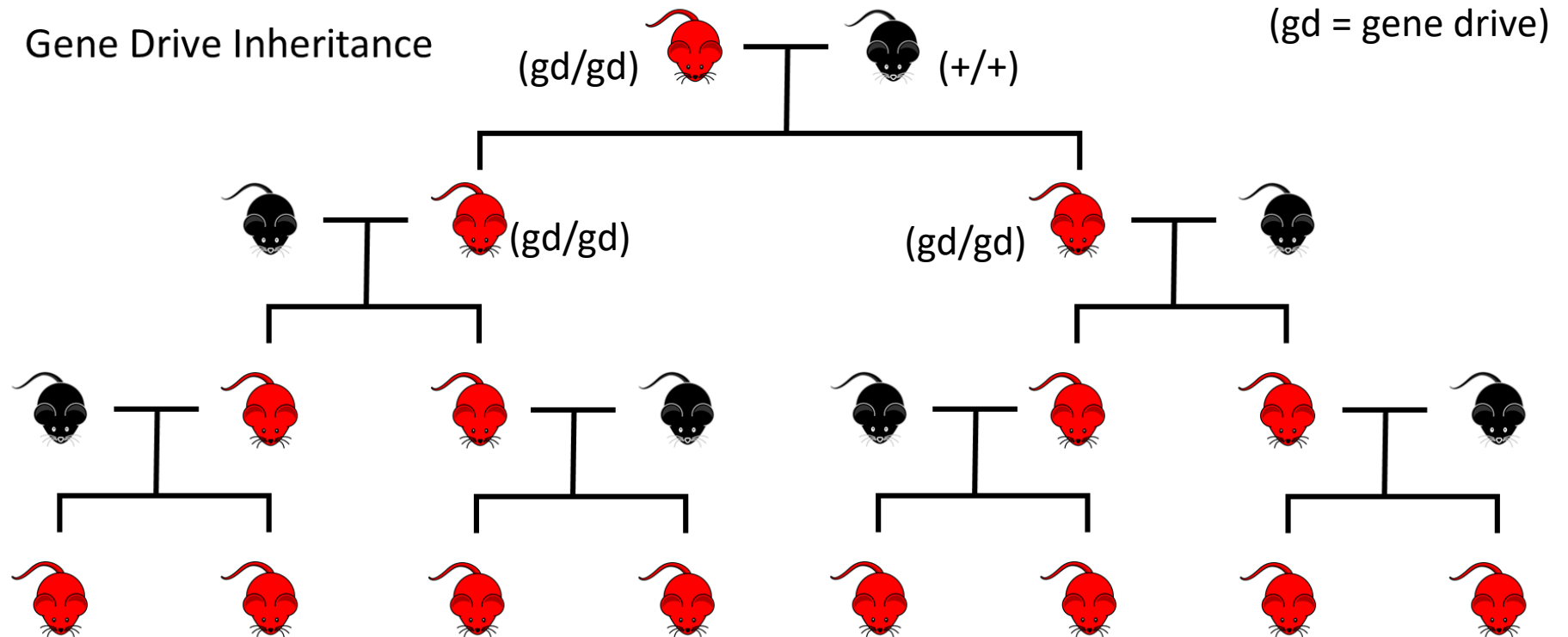
What is a Gene Drive?

- Genetic construct (transgene) that promotes its own inheritance at a rate greater than Mendelian inheritance
- Potentially spreads through entire population and allows population-level genetic engineering (modification or **suppression (fertility or sex bias)**)



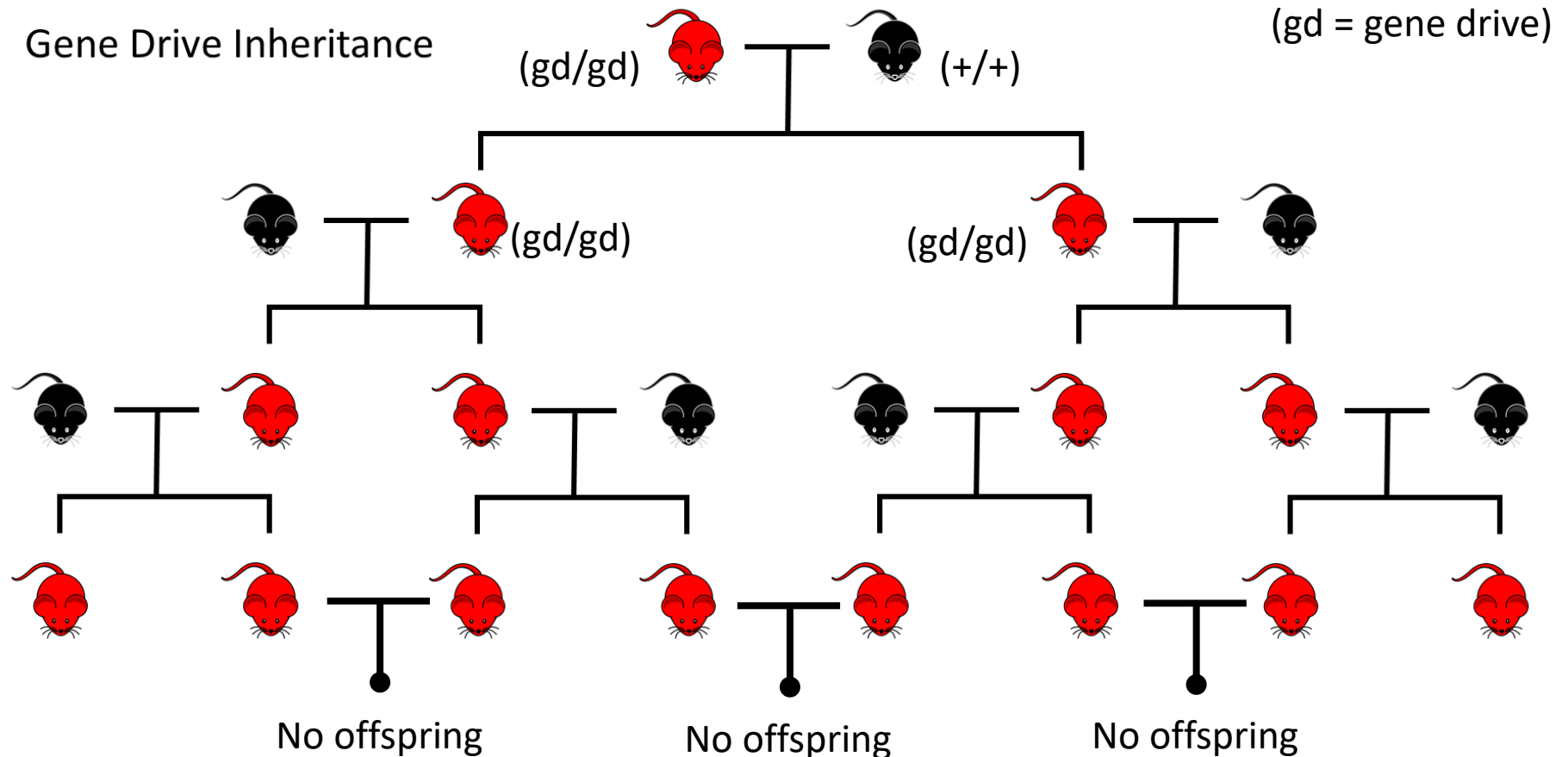
What is a Gene Drive?

- Genetic construct that promotes its own inheritance at a rate greater than Mendelian inheritance
- Potentially spreads through entire population and allows population-level genetic engineering (modification or **suppression (fertility or sex bias)**)



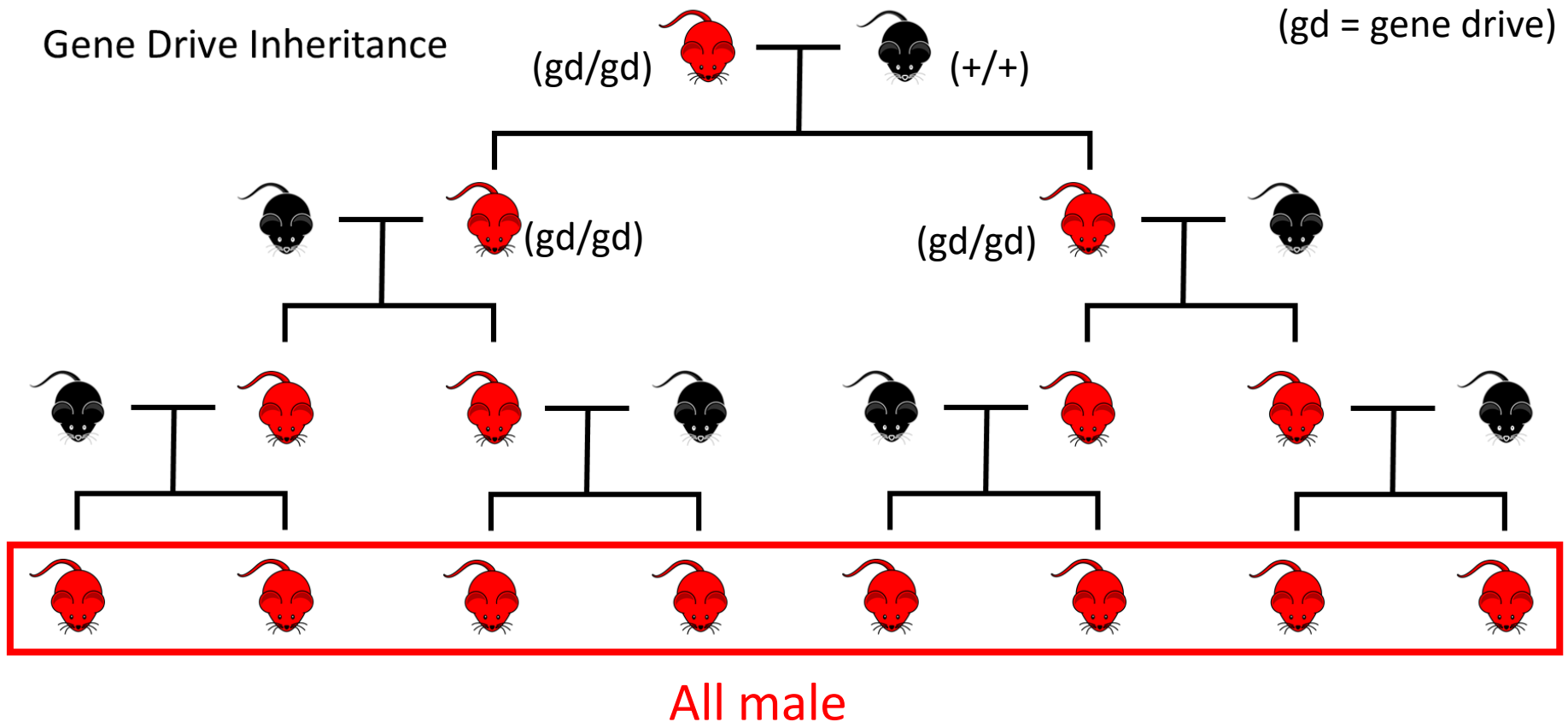
What is a Gene Drive?

- Genetic construct that promotes its own inheritance at a rate greater than Mendelian inheritance
- Potentially spreads through entire population and allows population-level genetic engineering (modification or **suppression (fertility or sex bias)**)



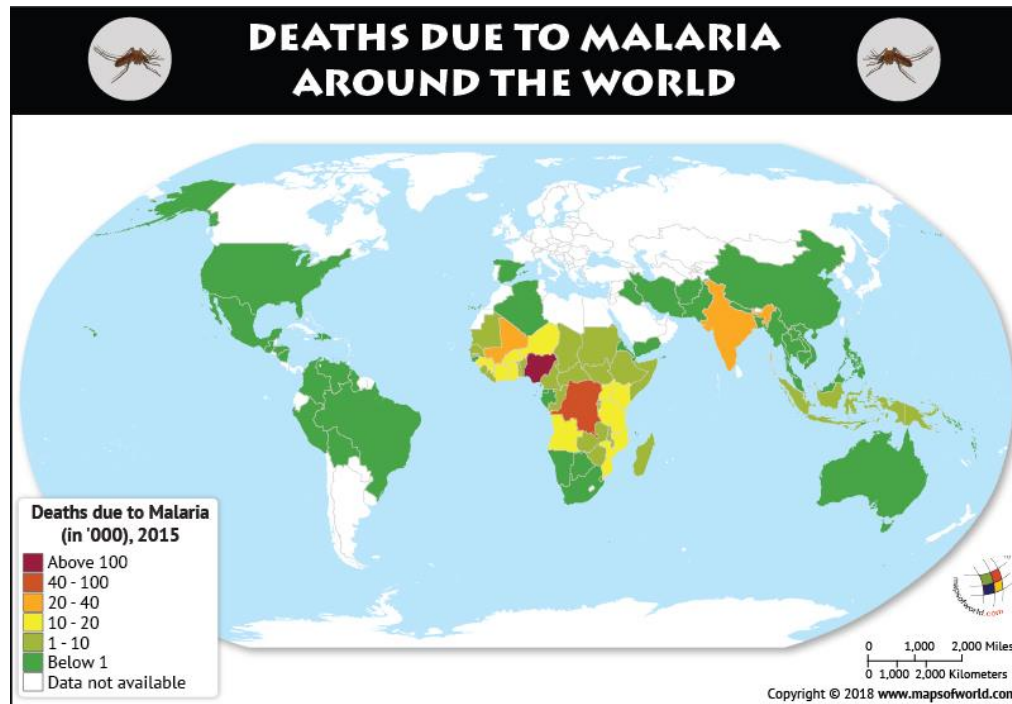
What is a Gene Drive?

- Genetic construct that promotes its own inheritance at a rate greater than Mendelian inheritance
- Potentially spreads through entire population and allows population-level genetic engineering (modification or **suppression (fertility or sex bias)**)



Why develop gene drives?

- Health, conservation & agriculture



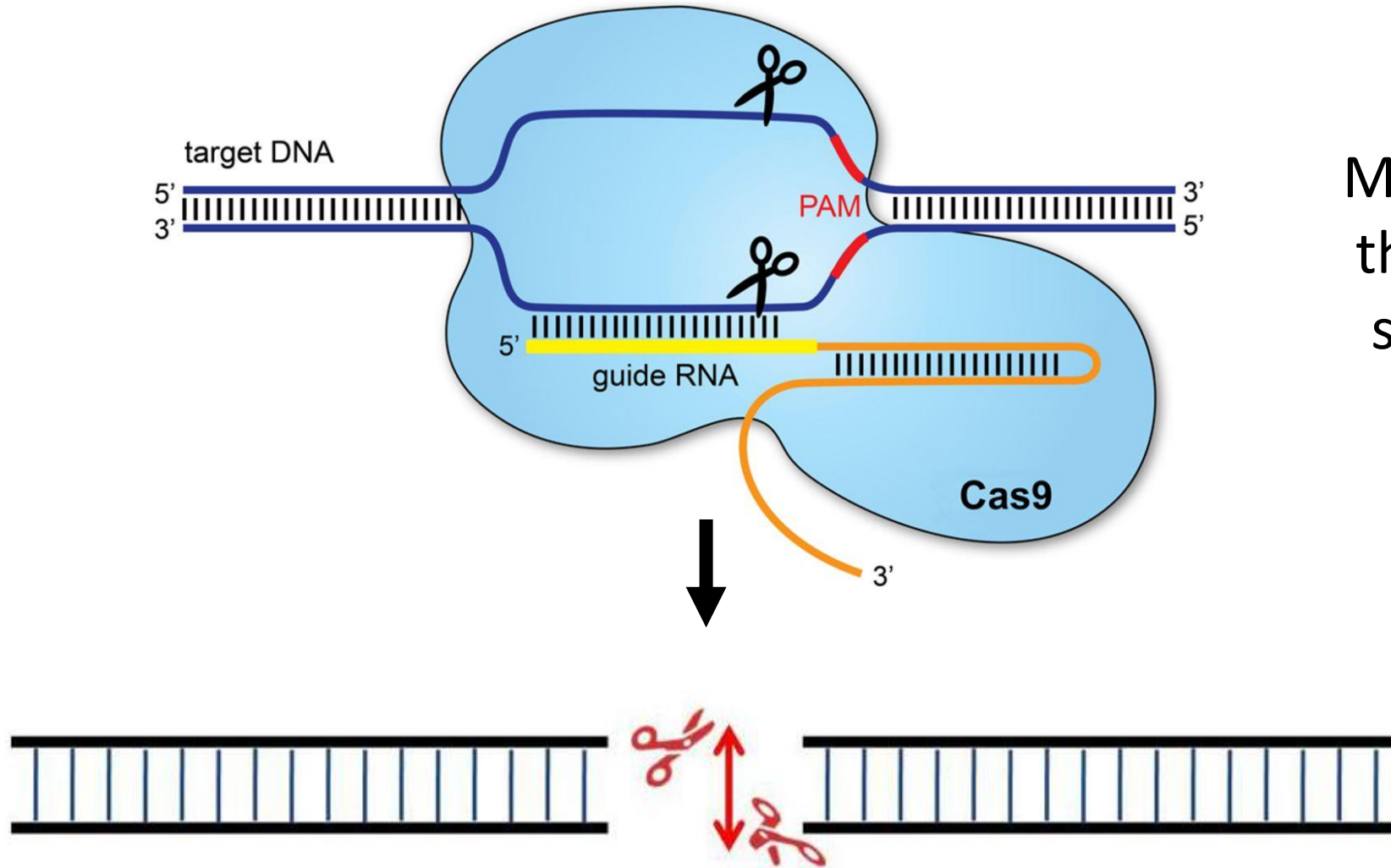
Malaria is responsible for >400,000 deaths per year



Hundreds of mice that have been trapped during the plague on Qld's Darling Downs. (Supplied: Vicki Green)

Environmental damage/loss of biodiversity
Agricultural loss of productivity/societal impact

CRISPR/CAS9 Genome Editing



Molecular scissors
that cut DNA at a
specific location

CRISPR enables generation of gene drive (transgenic) animals and gene drive activity

In 2021...

ARTICLE

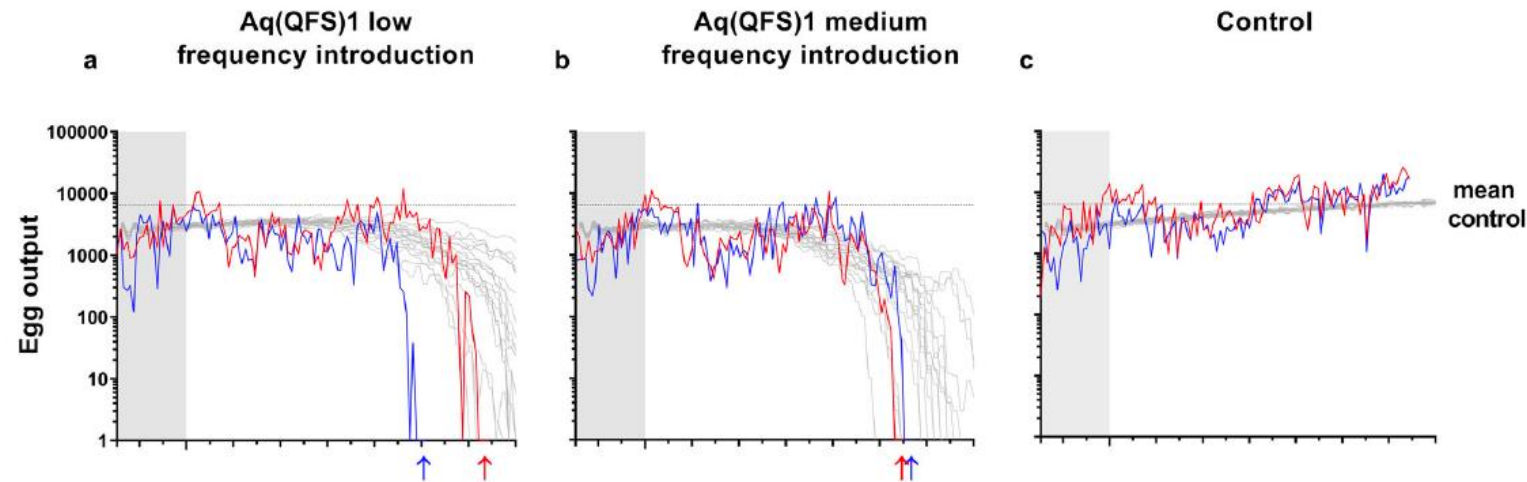


<https://doi.org/10.1038/s41467-021-24790-6>

OPEN

Gene-drive suppression of mosquito populations in large cages as a bridge between lab and field

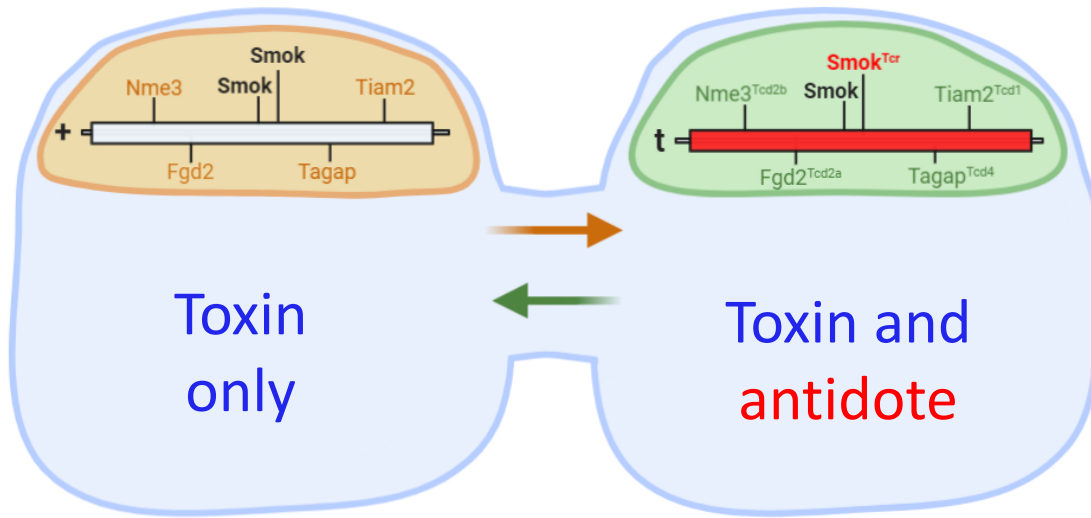
Andrew Hammond^{1,2,9}, Paola Pollegioni^{3,4,9}, Tania Persampieri^{3,9}, Ace North⁵, Roxana Minuz³, Alessandro Trusso³, Alessandro Bucci³, Kyros Kyrou¹, Ioanna Morianou¹, Alekos Simoni^{1,3}, Tony Nolan^{1,6,10}, Ruth Müller^{3,7,8,10} & Andrea Crisanti^{1,10}



“homing drives” not efficient enough in mammals

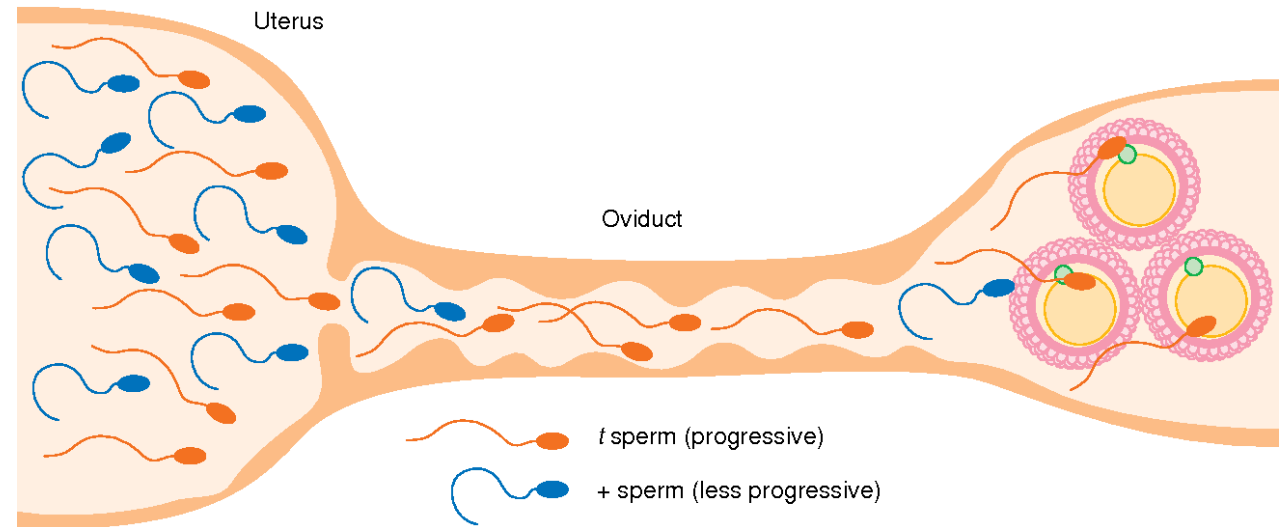
The *t* haplotype – a natural gene drive in male mice

Developing sperm



Motility ☹️

Motility 😊



Olds-Clarke et al 1997

- Male heterozygotes pass on up to 95% (females 50%)
- Male homozygotes infertile (t^{w2})

Can we modify the *t* haplotype to create a suppression gene drive?

Model framework

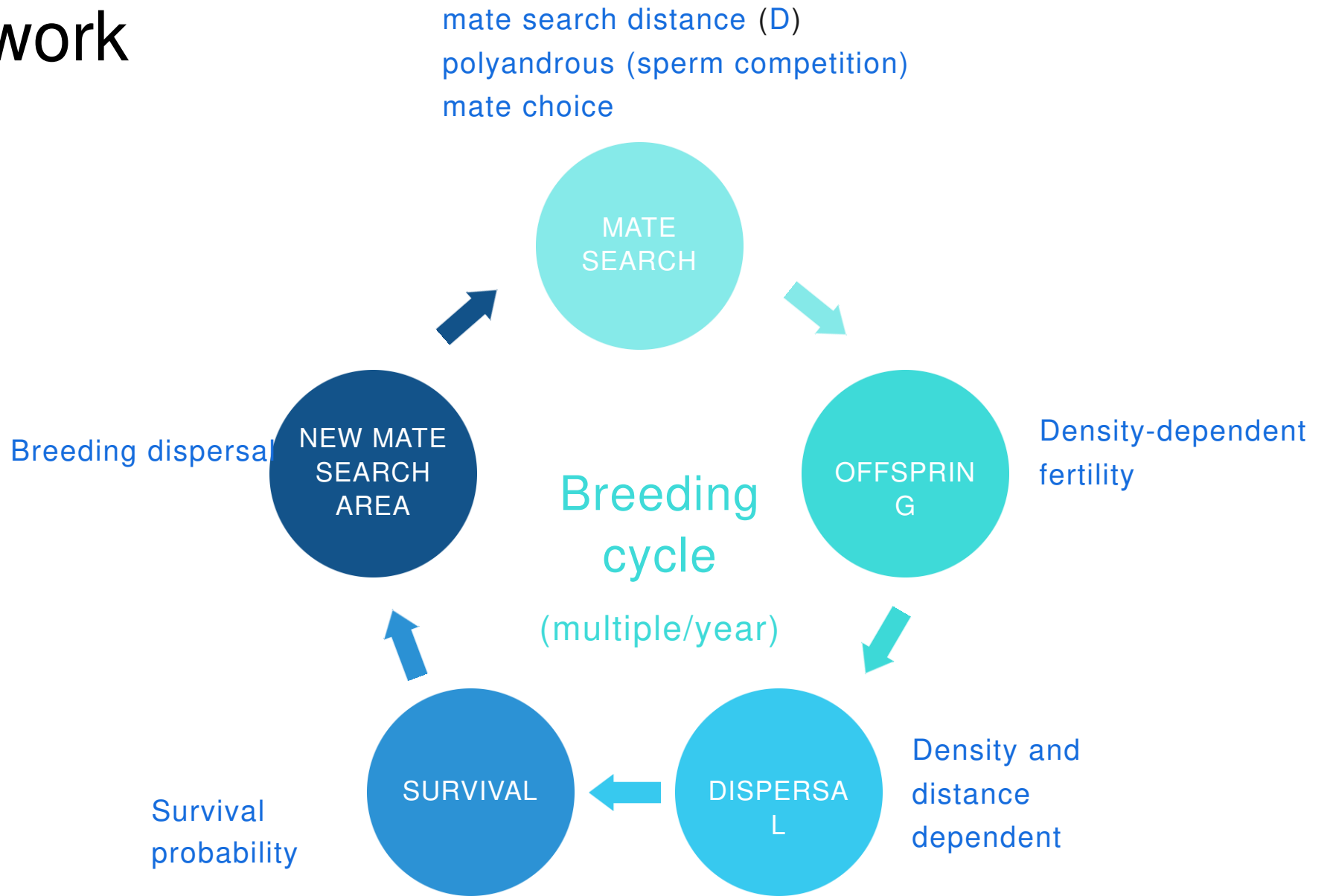
Individual based,
spatially explicit,
stochastic

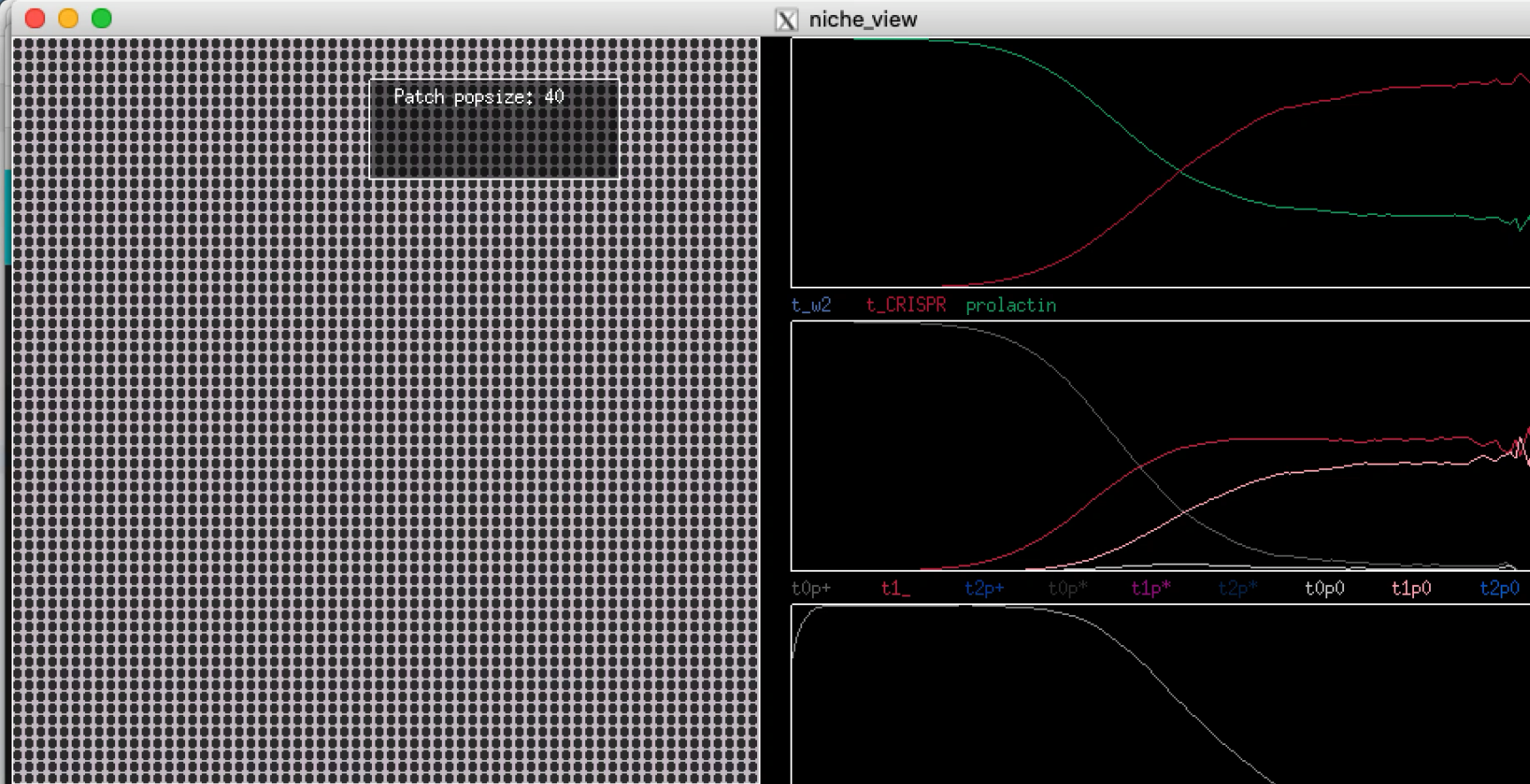
Landscape

Array of patches
Individuals use multiple patches
N ~ 200,000

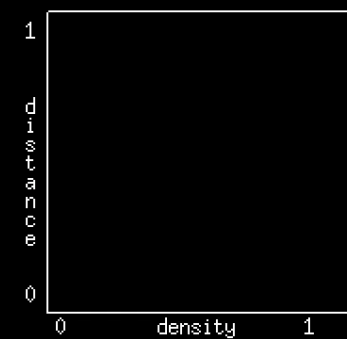
Individuals

Diploid
Discrete sexes (XX and XY)
Genetically controlled traits





Speed: 0
 Years: -2.33 (0/159)
 Mating cycles/year: 6
 T-CRISPR: 0.0000
 T-w2: 0.0000
 Prolactin: 1.0000
 N(max): 208385
 T(reduction): 22.33
 Multiple Paternity: nan



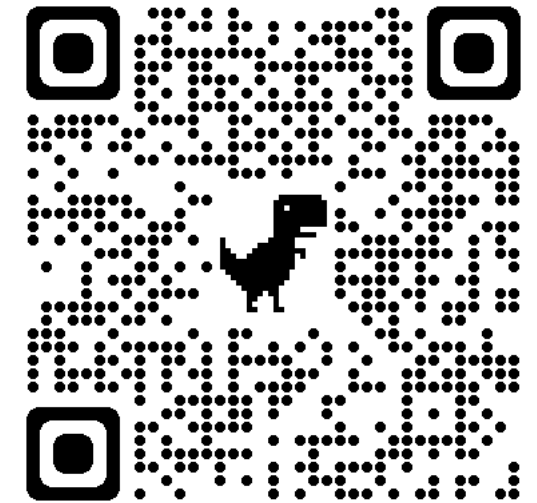
Stochastic individual-based modelling
 Island population of 200,000 mice
 Deploy 256 *t*-CRISPR mice (1/patch)
 Proof of concept in lab mice

Leveraging a natural murine meiotic drive to suppress invasive populations

Luke Gierus^{a,b,1} , Aysegul Birand^{c,1} , Mark D. Bunting^{a,b} , Gelshan I. Godahewa^{b,d}, Sandra G. Piltz^{a,b}, Kevin P. Oh^{e,f} , Antoinette J. Piaggio^g, David W. Threadgill^h , John Godwinⁱ , Owain Edwards^{e,j} , Phillip Cassey^c, Joshua V. Ross^k , Thomas A. A. Prowse^c and Paul Q. Thomas^{a,b,2}

PNAS 2022 Vol. 119 No. 46 e2213308119

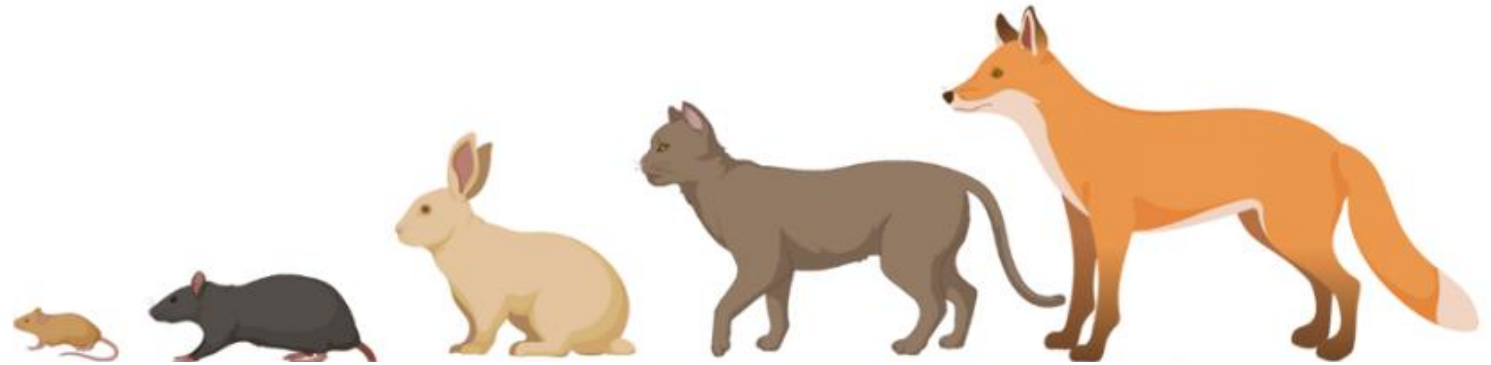
First proof of concept for a mammalian gene drive



What about cats (and other invasive pest mammals)?

What about other vertebrates?

$N \sim 200,000$

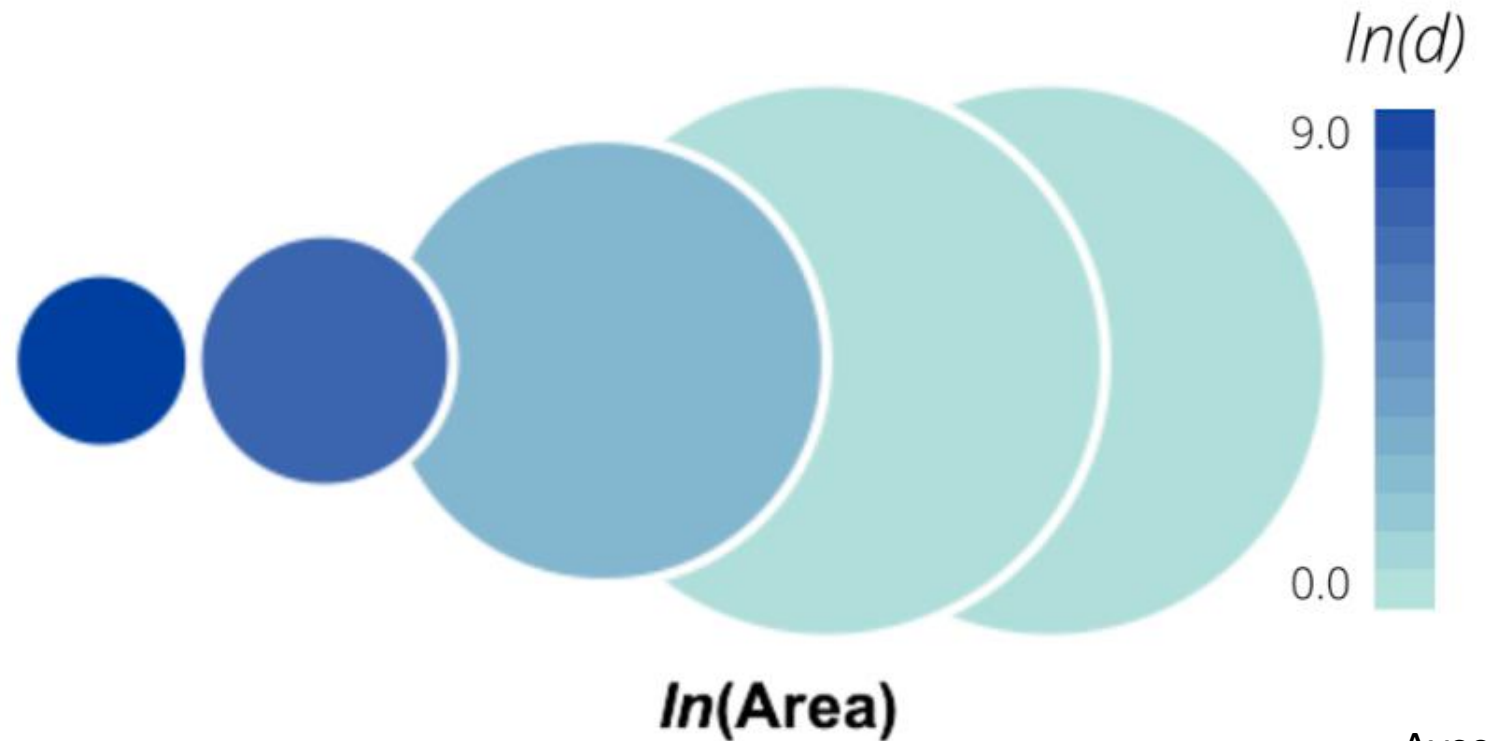


X-SHREDDER

Male biasing drive

HOMING

female infertility drive



Life-history parameters

Survival probability Probability of polyandry Dispersal

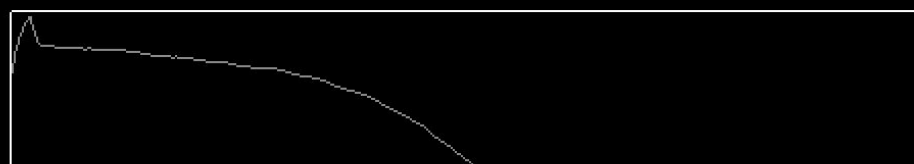
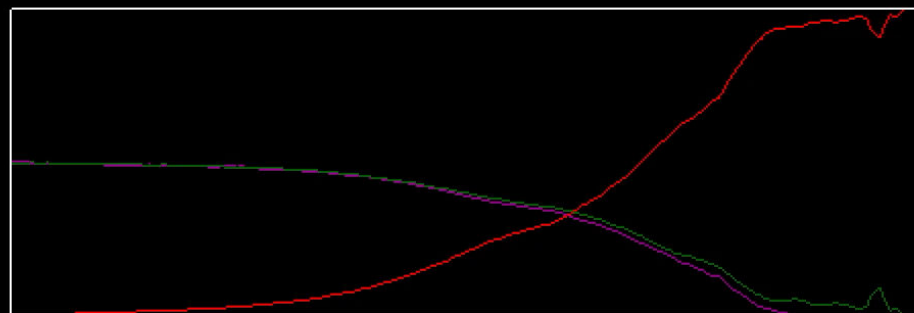
Parameters:

Species	b	n_c	age_m	ω	p_m	d	A	Δ_i	D
mouse	6	6	2	0.53	0.46	5000	40	0.4	3
black rat	4	6	2	0.62	0.68	1000	200	2	8
rabbit	4	4	3	0.82	0.20	25	8000	12.5	8
cat	4	2	5	0.85	0.25	2	100000	25	4
fox	4	2	5	0.88	0.76	2	100000	45	8

Island population of 200,000 cats

256 gene drive cats introduced

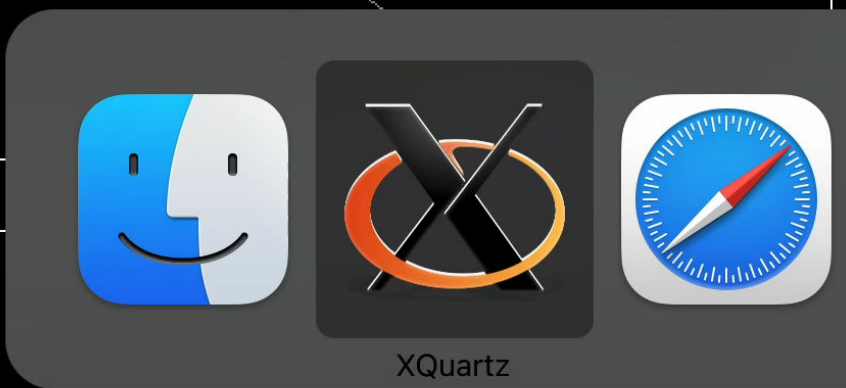
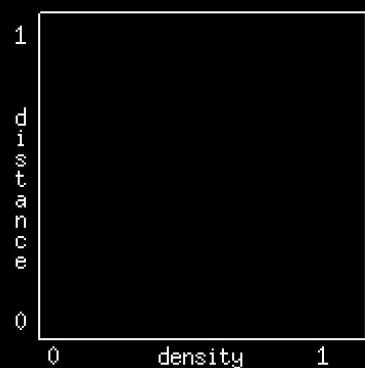
X-Shredder male biasing drive



Eradication in ~200 years!

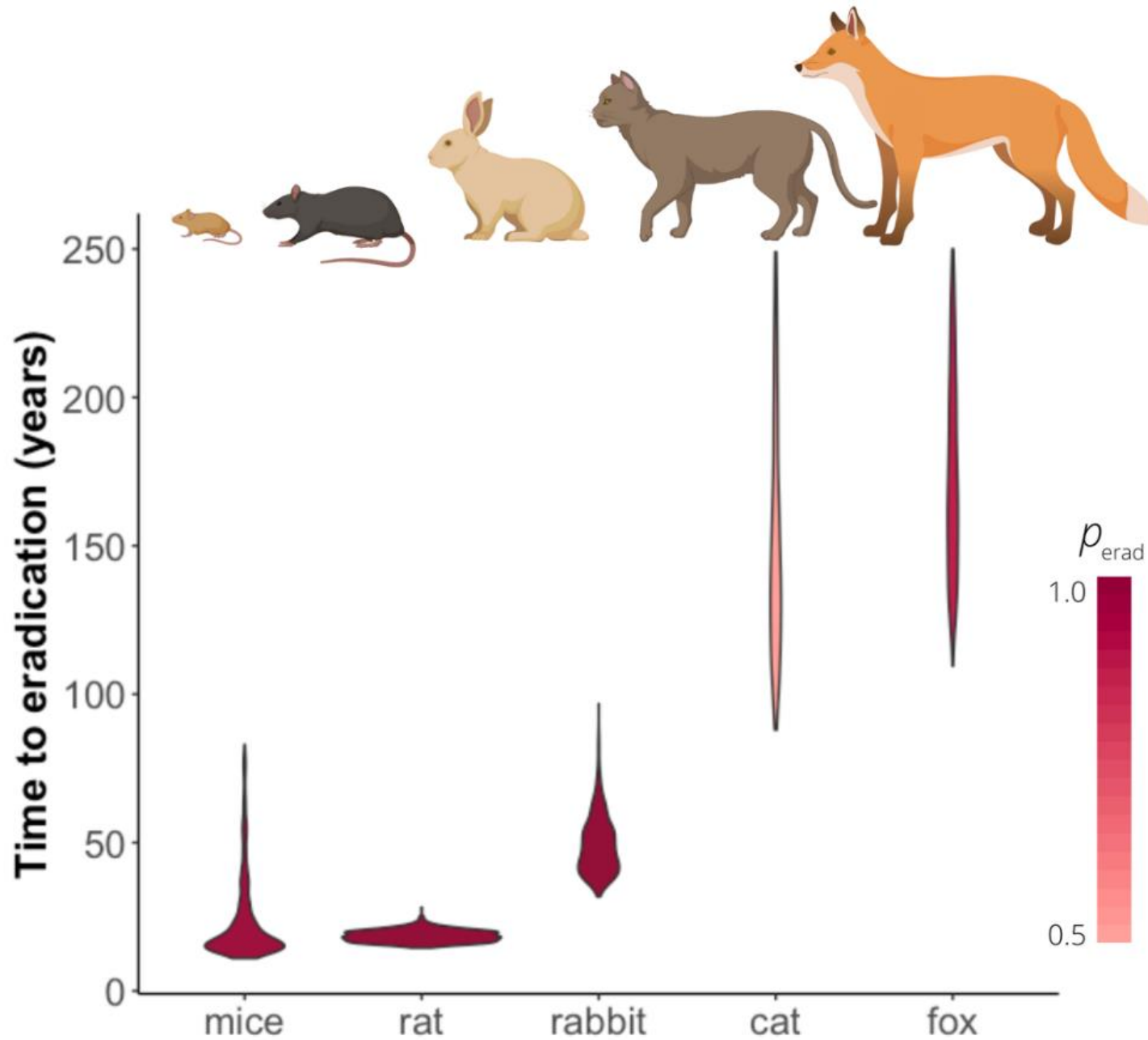


Speed: 0
Years: -7.00 (0/435)
Mating cycles/year: 2
Male: 81920
Male*: 0
Female: 81920
N/T/P(inoc): 1/1/256
N(max): 215466
T(reduction): 175.00
Multiple Paternity: nan



Results

(1000 sim.
per species)



Species	50%	90%	100%
Mouse	6.7	9.2	17.7
Black rat	9.0	11.7	18.5
Rabbit	16.8	24.1	48.0
Cat	71.0	92.0	143.2
Fox	74.0	103.5	169.0



Conclusions and Challenges

Genetic biocontrol (gene drive) technology is progressing in insects and mice – potential for disease control, conservation and agriculture

-stakeholder engagement, regulation, technical hurdles (inc. target population specificity)

Cat genetic biocontrol

-long timeframes

-technical challenges (transgenesis, facilities, genetics, reproductive technology)

-domesticated non-model animal

Stakeholder engagement (cf. CSIRO/Aditi Mankad stakeholder engagement survey (hypothetical “cat gene drive” scenario)

Modelling informed by more accurate field data (CSIRO)

Acknowledgements

CRISPR Therapeutics

Fatwa Adikusuma
Ashleigh Geiger
Jayshen Arudkumar
Joshua Chey
Caleb Lushington
Jesse Kennedy
Lachlan Staker

Genetic Biocontrol

Luke Gierus
Gelshan Godahewa
Mark Bunting

PCDH19 Epilepsy

Stefka Tasheva
Michaela Scherer

SA Genome Editing

Sandra Piltz
Melissa White

Lab Manager

Suraiya Onnesha

Modeling (Uni Adelaide)

Aysegul Briand
Thomas Prowse
Josh Ross
Phill Cassey

GBIRd consortium

Royden Saah

t mice

David Threadgill
John Godwin

X-shredder CSIRO

Owain Edwards
Mark Tizard

Gene Drive Funding

Australian Research Council Linkage
Grant with CAGT Ltd. (NZ)
Australian Research Council Discovery
Grant
CSIRO Post-doctoral Fellowship
NSW and **SA** State Governments



Thomas lab

