



70 years of rabbit biocontrol in Australia

An ongoing co-evolutionary
arms race

Tanja Strive | Montpellier, July 2022

Australia's National Science Agency



Outline

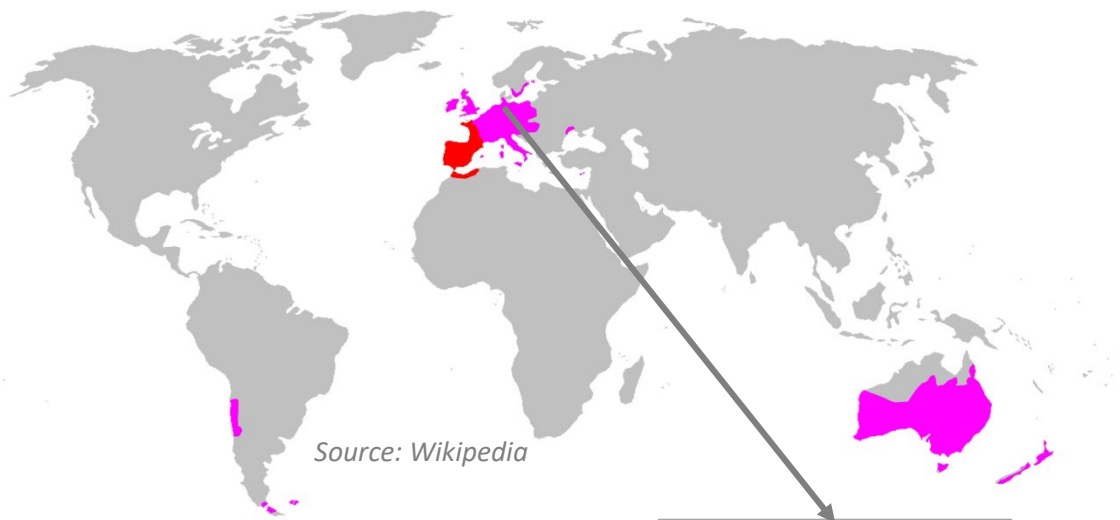
70 years of rabbit biocontrol

- Rabbits and rabbit biocontrol in Australia
 - Myxoma virus & RHDV
 - Host-pathogen co-evolution
- Impacts of rabbit biocontrol
- Current & future biocontrol initiatives in Australia





European rabbits (*Oryctolagus cuniculus*)



Source: Wikipedia



■ native ■ introduced



Iberian lynx
(*Lynx pardinus*)



Spanish
Imperial eagle
(*Aquila adalberti*)

Sacramento-Moreno et al., 2007
Delibes-Mateos et al., 2009



The European rabbit (*Oryctolagus cuniculus*): A perfect invader

- 26 rabbits (released in 1859 for hunting)
- Colonised continent in 70 years
- Ideal environment, few diseases
- Gestation period 28 days, mate same day of giving birth
- Large litters (5-7), early sexual maturity
- Peak estimate: 0.5 – 10 billion rabbits
- Massive economic and ecological impacts

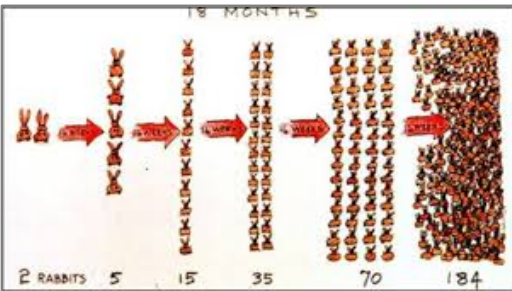
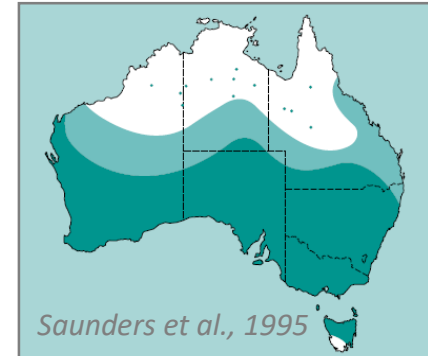


Image: Matt Probert Photography



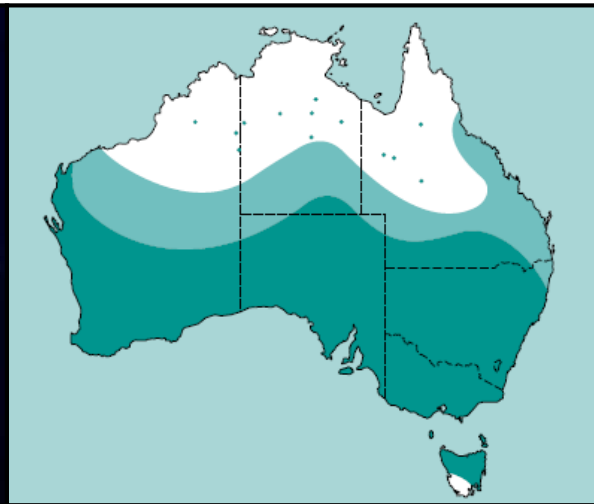
Australia in 1930:



6.7 million km²



6.5 million people
(in 1930)



Estimated
0.5 – 10 billion rabbits

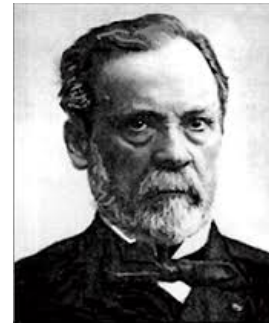
- ⇒ Conventional rabbit control (shorting, trapping, etc) ineffective
- ⇒ Self-disseminating (=biological) controls were needed for continental scale control



Biological control of the European rabbit (*Oryctolagus cuniculus*) - early days

1887: NSW-Government promised £24,000 rewards for eradication of rabbits

Pasteurella multocida (Chicken cholera)
Not species specific, did not transmit



Louis Pasteur
1822-1895



Adrien Loir
1862-1941



Myxoma virus (MYXV): an emerging disease of rabbits

Family: *Poxviridae*, Genus: *Leporipoxvirus*, ds-DNA, 160 kb

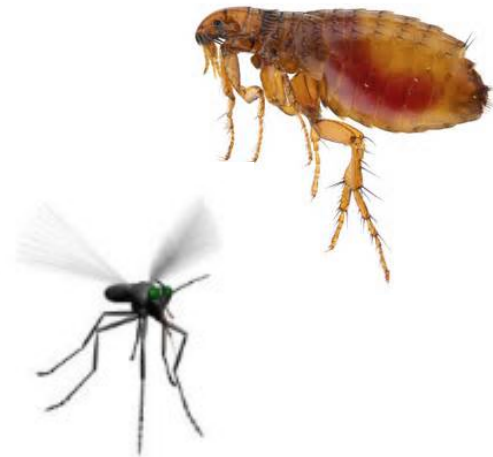
Natural host: Tapeti
(*Sylvilagus brasiliensis*)
Mild infection (cutaneous fibromas)



Jumped species to European rabbit 1896
(*Oryctolagus cuniculus*)
Highly virulent generalised disease



Transmission (mechanical) by biting insects
Fleas & mosquitoes





MYXV – release and impacts

- 1950 release: >90% mortality rates in some areas
- Within 2-3 years attenuated MYXV strains appeared (and dominated)
- Rabbits developed genetic resistance
- Rabbit numbers increased again!

- Introduction of rabbit fleas to boost transmission
 - *Spilopsyllus cuniculi* (1966)
 - *Xenopsylla cunicularis* (1995)



Dame Jean
Macnamara
1914-2010



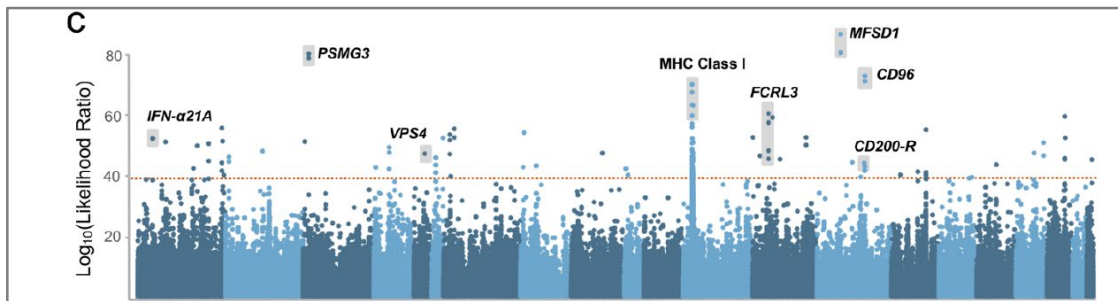


MYXV – text book example for ‘Trade-off Theory’ of Virulence Evolution of an emerging disease



Frank Fenner
1914-2010

- Selection of moderately virulent phenotype for maximum transmission by biting insects:
 - Increased survival rates and survival times
 - Longer disease period
 - Increased opportunity for mosquitoes/fleas to pick up virus from skin lesions
 - Much more severe disease & high welfare impact
- Rapid (2 year) selection for host resistance acting on standing genetic variation: Immune modulating genes



Science

RESEARCH ARTICLES

Cite as: J. M. Alves *et al.*, *Science* 10.1126/science.aau7285 (2019).

Parallel adaptation of rabbit populations to myxoma virus

Joel M. Alves^{1,2,3*}, Miguel Carneiro^{2,4*}, Jade Y. Cheng^{5,6}, Ana Lemos de Matos⁷, Masmudur M. Rahman⁷, Liisa Loog^{2,8}, Paula F. Campos^{6,9}, Nathan Wales^{6,10,11}, Anders Eriksson¹², Andrea Manica¹², Tanja Strive^{12,14}, Stephen C. Graham⁵, Sandra Afonso^{6,9}, Diana J. Bell¹⁵, Laura Belmont¹⁶, Jonathan P. Day¹, Susan J. Fuller¹⁷, Stéphane Marchandeaun¹⁸, William J. Palmer¹⁹, Guillaume Queney²⁰, Alison K. Surridge²¹, Filipe G. Vieira², Grant McFadden², Rasmus Nielsen^{2,2}, M. Thomas P. Gilbert^{2,21}, Pedro J. Esteves^{2,22}, Nuno Ferrand^{2,23}, Francis M. Jiggins^{1*}



The race continues.....

- 1990s MYXV field strains are hypervirulent in unselected domestic rabbits, causing a completely new disease phenotype

PNAS

Next step in the ongoing arms race between myxoma virus and wild rabbits in Australia is a novel disease phenotype

Peter J. Kerr^{a,b}, Isabella M. Cattadori^{c,d}, June Liu^b, Derek G. Sim^{c,d}, Jeff W. Dodds^e, Jason W. Brooks^e, Mary J. Kennett^e, Edward C. Holmes^{a,f}, and Andrew F. Read^{c,d,g,1}

^aMarie Bashir Institute for Infectious Diseases and Biosecurity, Charles Perkins Centre, School of Life and Environmental Sciences, University of Sydney, Sydney, NSW 2006, Australia; ^bCSIRO Health and Biosecurity, Black Mountain Laboratories, Acton, ACT 2601, Australia; ^cCenter for Infectious Disease Dynamics, The Pennsylvania State University, University Park, PA 16802; ^dDepartment of Biology, The Pennsylvania State University, University Park, PA 16802; ^eDepartment of Veterinary and Biomedical Sciences, The Pennsylvania State University, University Park, PA 16802; ^fSydney Medical School, University of Sydney, Sydney, NSW 2006, Australia; and ^gDepartment of Entomology, The Pennsylvania State University, University Park, PA 16802

Edited by Simon A. Levin, Princeton University, Princeton, NJ, and approved July 14, 2017 (received for review June 23, 2017)

Peter Kerr



“Now, *here*, you see, it takes all the running you can do, to keep in the same place”
(Lewis Carroll, Through the looking glass)





New hope for Australia's feral rabbit problem

Rabbit Haemorrhagic
Disease Virus (RHDV)





Rabbit Haemorrhagic Disease Virus (RHDV)

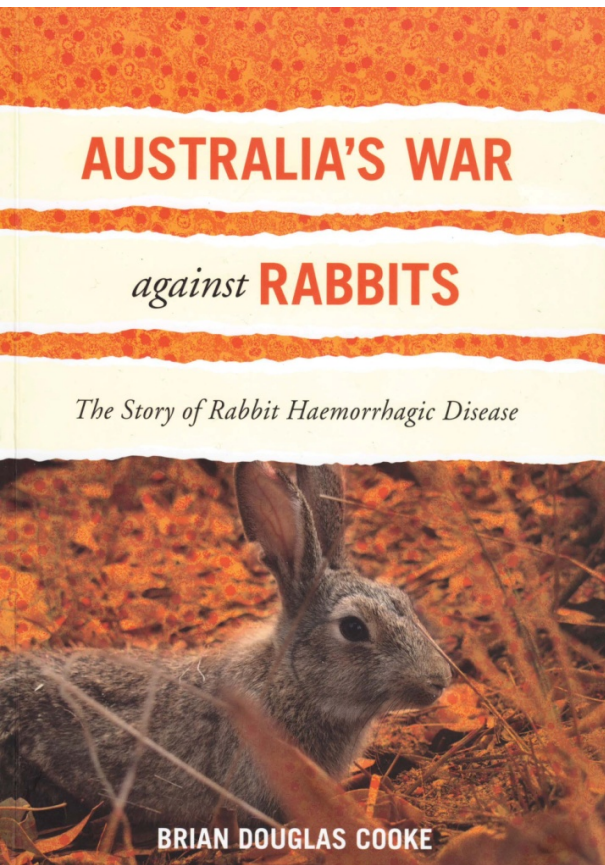
Family: *Caliciviridae*, Genus: *Lagovirus*, +ss-RNA, 7.5 kb



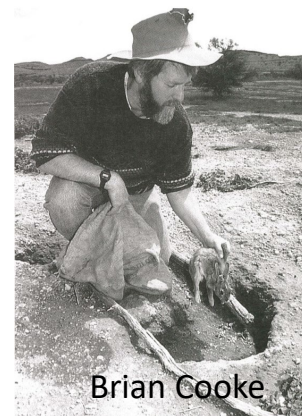
- Emerging disease of rabbits (China 1984)
- Infectious hepatitis with >95% case fatality rates
- Death 36-72 h.p.i.
- **Short period of clinical signs (~12h)**
- Young rabbits innately resistant to lethal disease
- Does not grow in conventional cell culture
- Species specific (infects only European rabbits)



Bringing RHDV to Australia



- Quarantine tests (CSIRO AAHL) commenced in 1991
 - Later: field transmission trials on Wardang Island
 - Escaped in 1995 (Fly transmission!)
 - Highly successful in reducing rabbit populations (> 95% in parts of Australia)
 - In 1997 illegally introduced into New Zealand
-
- some evidence of genetic resistance
 - No significant reduction in virulence observed



Brian Cooke

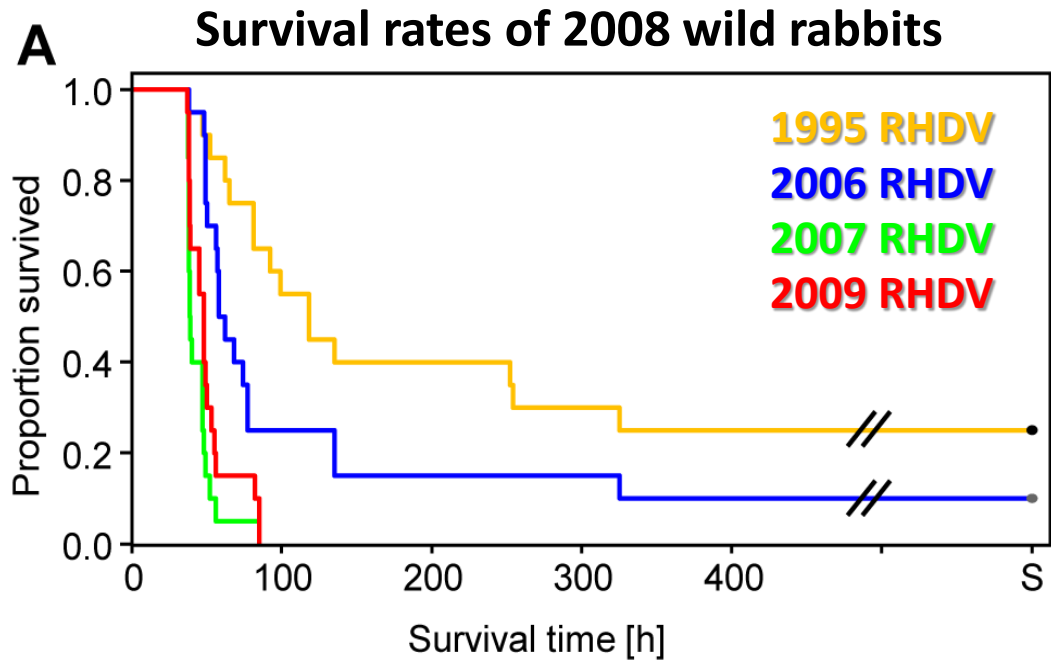


Virulence evolution of RHDV is the opposite of MYXV

=> More recent field strains kill a higher proportion of rabbits faster

Virology 464-465 (2014) 415-423
Contents lists available at ScienceDirect
Virology
journal homepage: www.elsevier.com/locate/yviro

Increased virulence of rabbit haemorrhagic disease virus associated with genetic resistance in wild Australian rabbits (*Oryctolagus cuniculus*)
Peter Elsworth^{a,b}, Brian D. Cooke^{b,c}, John Kovaliski^{b,d}, Ronald Sinclair^{b,d}, Edward C. Holmes^e, Tanja Strive^{b,f,g,*}



WHY?



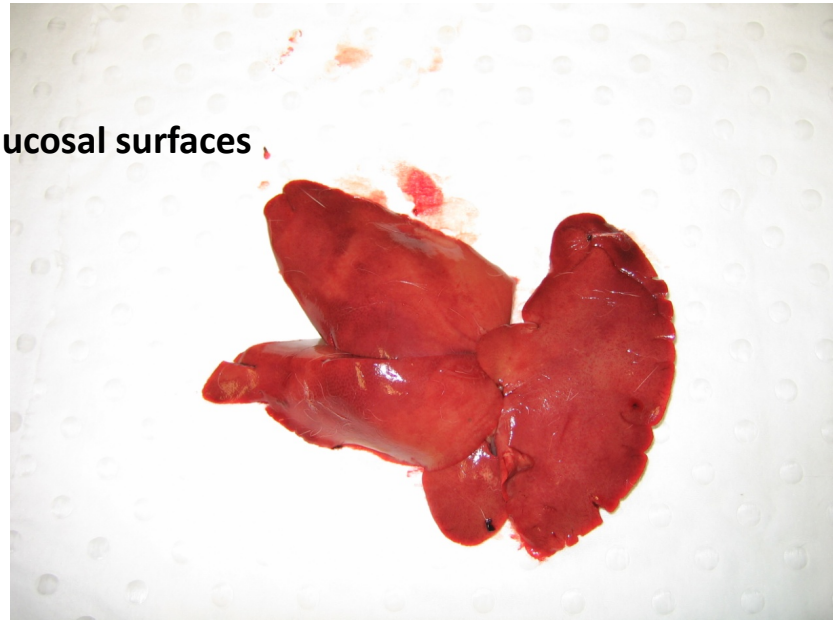
Transmission of RHDV:

Direct contact, fomites AND flies

- Rabbit dies
- VERY high concentration of virus in dead body (10e9 copies/mg, ~10e15/carcass)
- Body decays or is opened up by scavengers
- Fly feeds on carcass
- Flies to next rabbit warren
- Lands on grass and/or rabbit's mucosal surfaces

Plays a key role in distance transmission between non-connected rabbit populations

Asgari et al., 1998
McColl et al., 2002
Elsworth et al., 2014
Schwensow et al., 2014
Hall et al., 2018





Transmission of RHDV:

Direct contact, fomites AND flies

Asgari et al., 1998
McColl et al., 2002
Elsworth et al., 2014
Schwensow et al., 2014
Hall et al., 2018

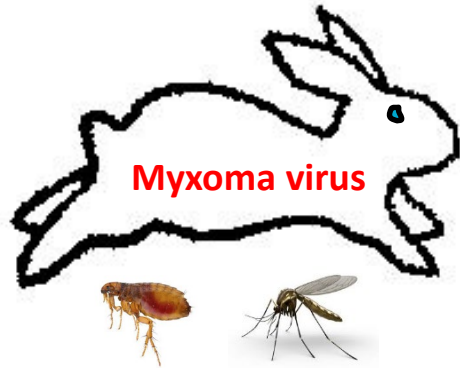
- Rabbit dies
- VERY high concentration of virus in dead body (10e9 copies/mg, ~10e15/carcass)
- Body decays or is opened up by scavengers
- Fly feeds on carcass
- Flies to next rabbit warren
- Lands on grass and/or rabbit's mucosal surfaces

Plays a key role in distance transmission between non-connected rabbit populations

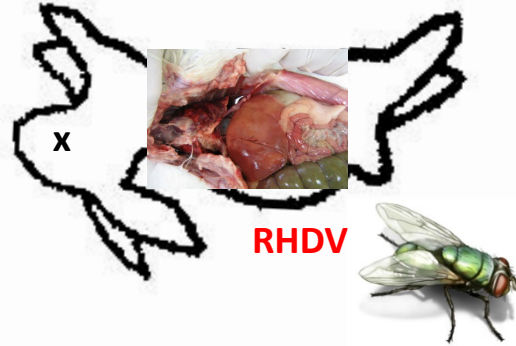




⇒ Both evolve towards maintain virulence levels optimal for mechanical insect transmission



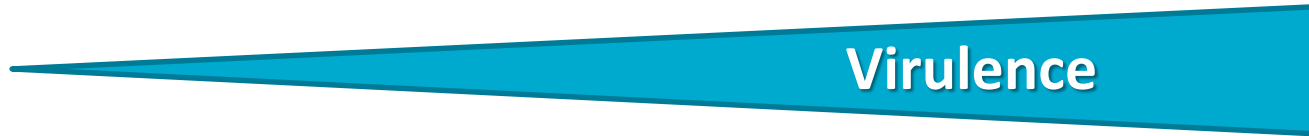
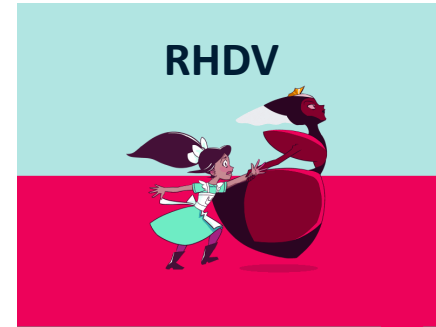
Biting insects
Moderate virulence



Carrion feeding flies
High virulence

For RHDV the dead animal is (not the sick animal) is an important source of distance transmission via insects

⇒ Both evolve towards maintain virulence levels optimal for mechanical insect transmission



Moderate

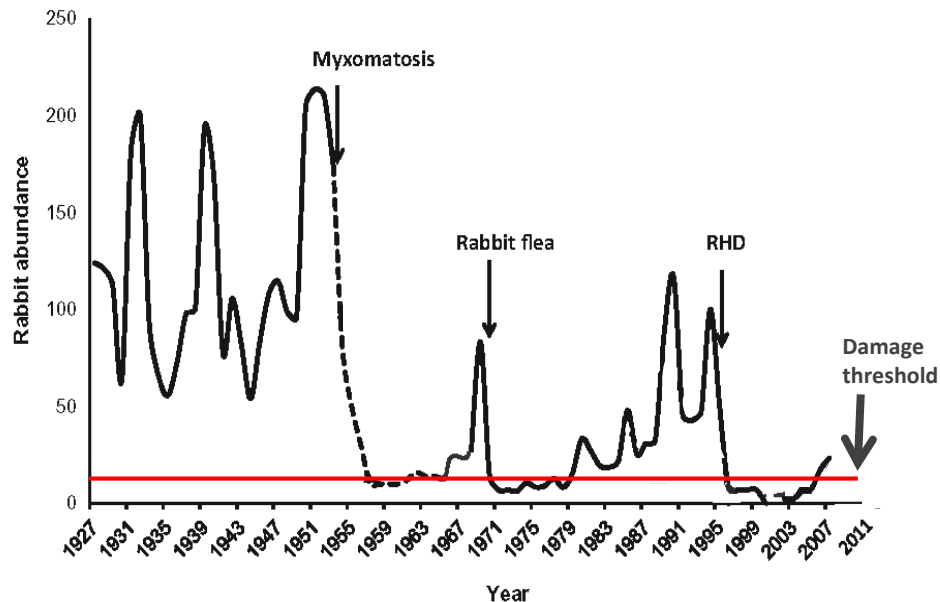


High



Has biocontrol worked?

VERY effective, but NOT a silver bullet !



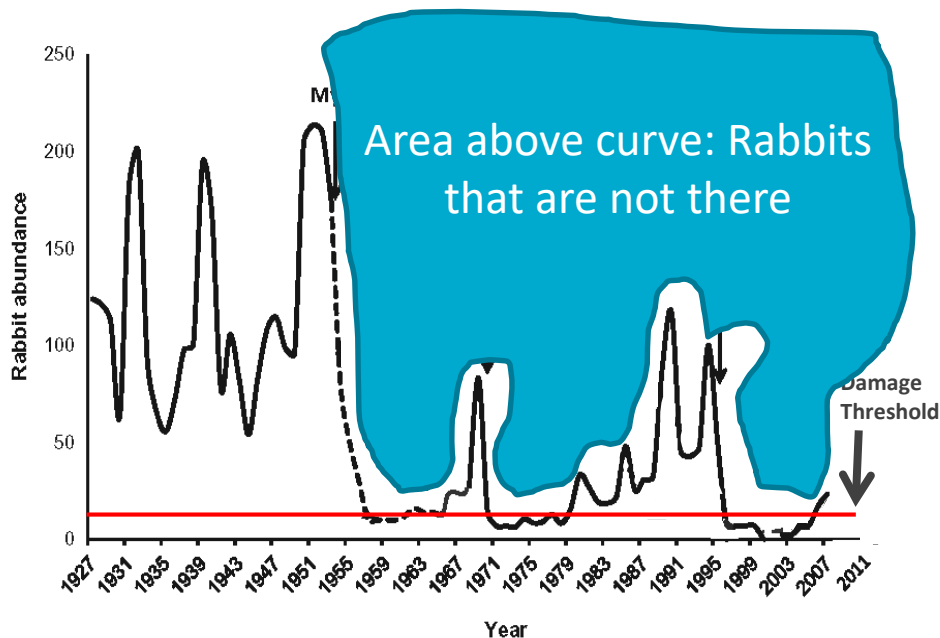
- Must be followed up with conventional control tools
- Virus will not eradicate the host
- ongoing host-pathogen co-evolution
- Repeated adjustments required to maintain gains

(Cooke et al., 2013)



Has biocontrol worked?

VERY effective, but NOT a silver bullet !



(Cooke et al., 2013)

- Must be followed up with conventional control tools
- Virus will not eradicate the host
- ongoing host-pathogen co-evolution
- Repeated adjustments required to maintain gains
- **NOT** about killing lots of viruses
- **BUT** about preventing rabbit impacts

Economic benefits of rabbit biocontrol



**Myxoma virus
1950**



**Rabbit fleas
1968 and 1993**



**Calicivirus
(RHDV)
1996**

1950-2010 (60 years): Combined cumulative benefits estimated at **\$70 billion** to the Australian agricultural industries
(.....and counting....)

(Cooke et al., 2013)

Ecological benefits

=> Top down and bottom up trophic effects



Sustains high population numbers of feral cats and foxes



Key drivers for mammalian extinction in Australia



Competition and land degradation



Plains mouse
(*Pseudomys australis*)



© Peter D Canty

Dusky hopping-mouse
(*Notomys fuscus*)



Crest-tailed mulgara (*Dasycercus cristicauda*)

Contributed Paper

Rabbit biocontrol and landscape-scale recovery of threatened desert mammals

Reece D. Pedler,* ¶ Robert Brandle,* John L. Read,†‡ Richard Southgate,§ Peter Bird,** and Katherine E. Moseby†‡

- Recovery of 3 mammalian species through 20 years of sustained rabbit suppression from RHDV
- Removal from the IUCN red list

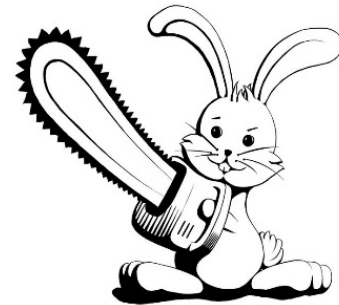


Less than 1 rabbit/ha can completely abolish the recruitment of palatable plant species

Photo: Tanja Strive
Hattah Kulkyn National Park 2009

- Long term suppression of rabbits has led to the recovery of many native plants, e.g. Sheoak & Cypress pine (*Allocasuarina sp.*, *callitris sp.*)

Sandell 2002



Today in Australia >300 threatened species and ecological communities are impacted by competition and land degradation caused by rabbits.

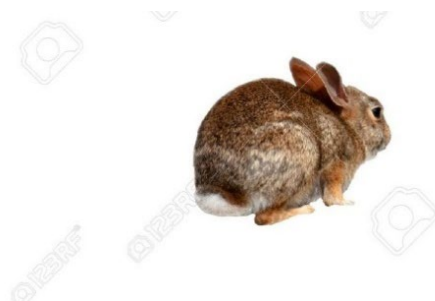
Rabbit exclosure on Macquarie Island

Image: Brian Cooke





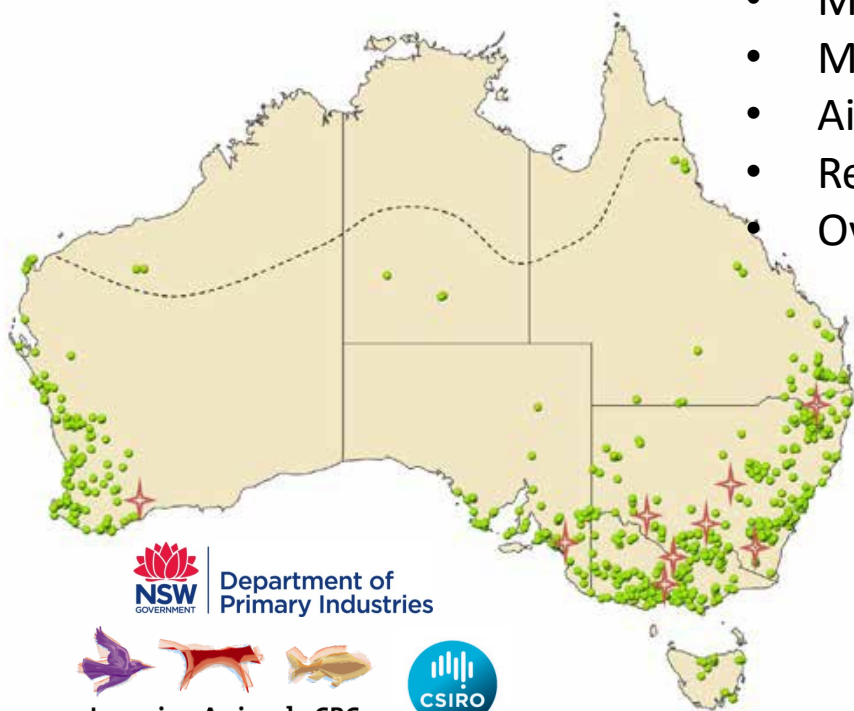
Recent rabbit biocontrol developments in Australia



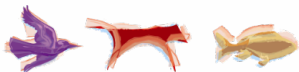


March 2017: Nationwide release of a new RHDVa variant from Korea (K5, GI.1a):

- More effective in genetically resistant rabbits
- More effective against endemic benign caliciviruses
- Aimed at slowing down rate of rabbit increase
- Released at >300 sites across Australia
- Overshadowed by the arrival of another RHDV.....



Department of Primary Industries



Invasive Animals CRC



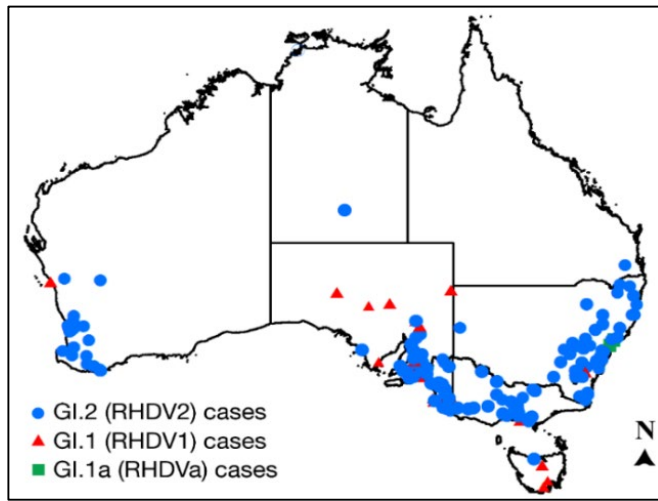


RHDV2 (GI.2) incursion (NOT released)

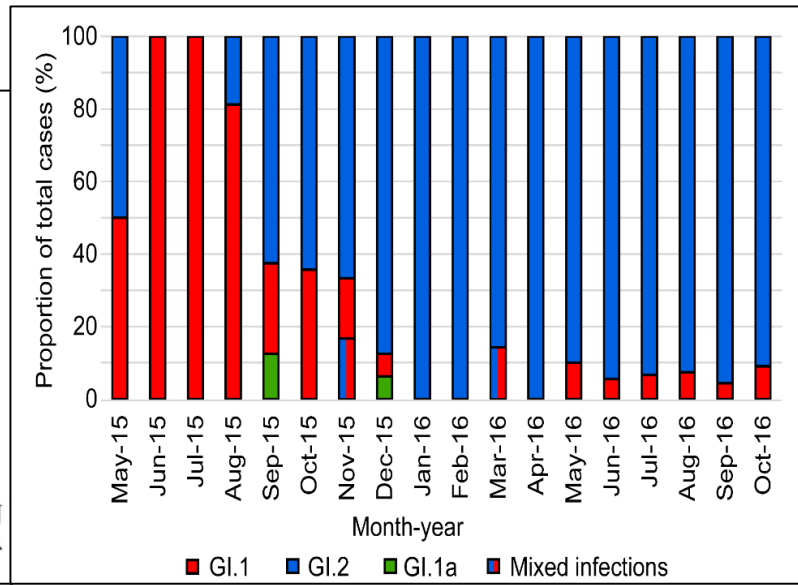
RHDV2: A new rabbit calicivirus

- Emerged in Europe in 2010
- Can overcome immunity to other strains (to some degree)
- Can kill young rabbits and other lagomorphs (incl. European hares)
- First reported in Australia in May 2015
- Has become dominant strain in Australia

Global spread
(all continents)
'lagomorph covid'

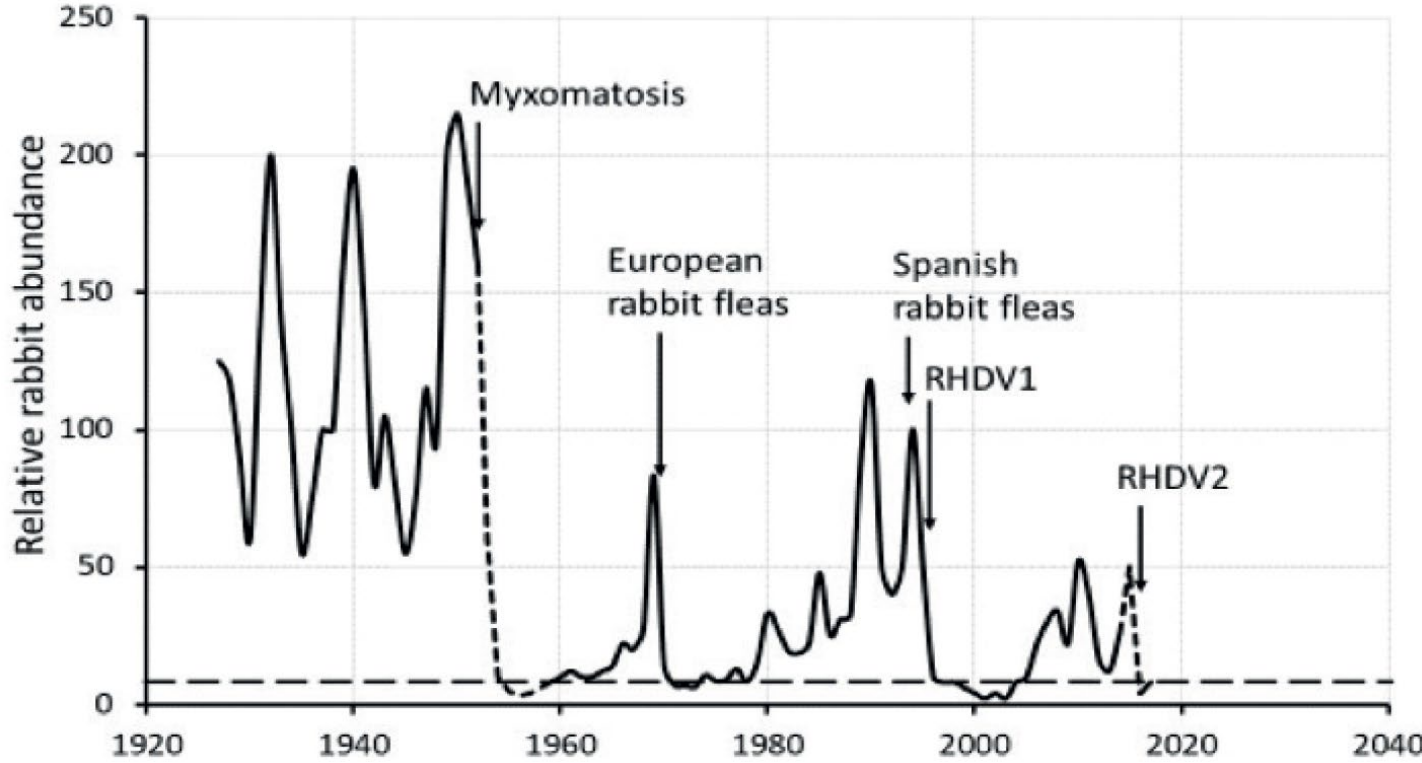


Mahar, Hall et al 2018





RHDV2 reduced Australian rabbit populations by ~60% (average)



Ramsey et al., 2020
Cooke et al., 2018

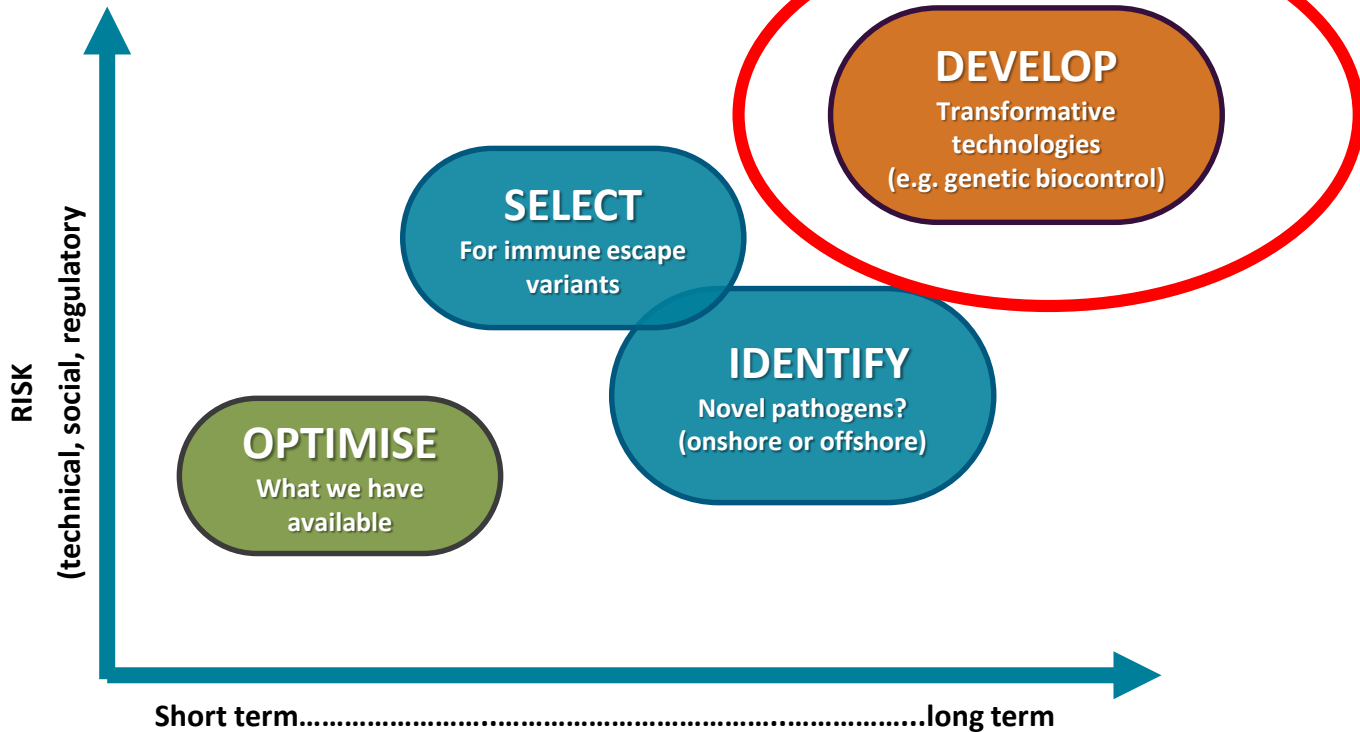


What's next in the Biocontrol pipeline?





AUSTRALIA'S RABBIT BIOCONTROL PIPELINE STRATEGY FOR 2022-27





Genetic biocontrol of invasive animals:

1. Precision genetic engineering of the
pest species
- +
2. 'driving' the engineered trait (genes)
into populations to control them



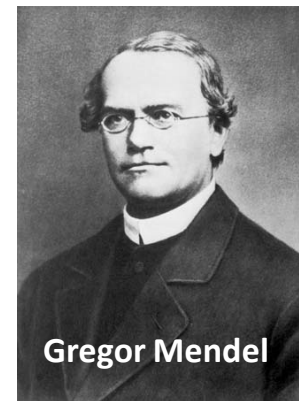
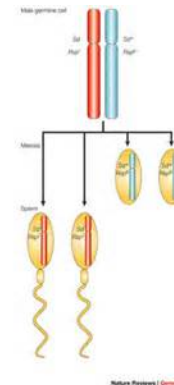
Example “Gene Drive”

During sexual reproduction of diploid organisms, the alleles segregate and the embryo gets one from each parent (50:50)

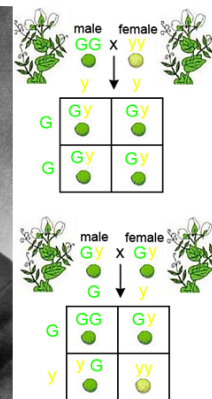
Gene drives break the laws of Mendelian inheritance (>50% inheritance of certain alleles)

Natural gene drive:
meiotic drives, ‘selfish genes’, >50% inheritance

- *sd* in drosophila
- *t complex* in mice
- Homing endonucleases (HEGs)



Gregor Mendel





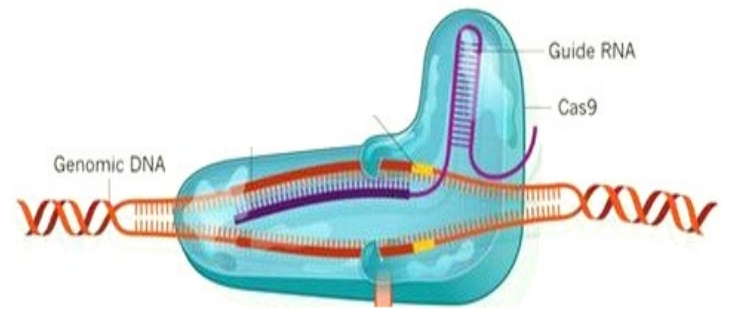
The 'Precision engineering' component: CRISPR-Cas9

Genome engineering tool derived from bacterial "immunity" system

CRISPR – clustered regularly interspaced short palindromic repeat

Cas9 - CRISPR associated protein 9 nuclease - cuts DNA

Guide RNA – tells Cas9 where to cut



How often will this cut a piece of DNA (at random)
Frequency of a **20 nt sequence** =
 $4^{20} = 10^{12} =$ **1 in 1000 billion**
(human genome has approx. 6 billion nt)

Nobel Prize:

Jennifer Doudner, Emmanuelle
Charpentier 2012:



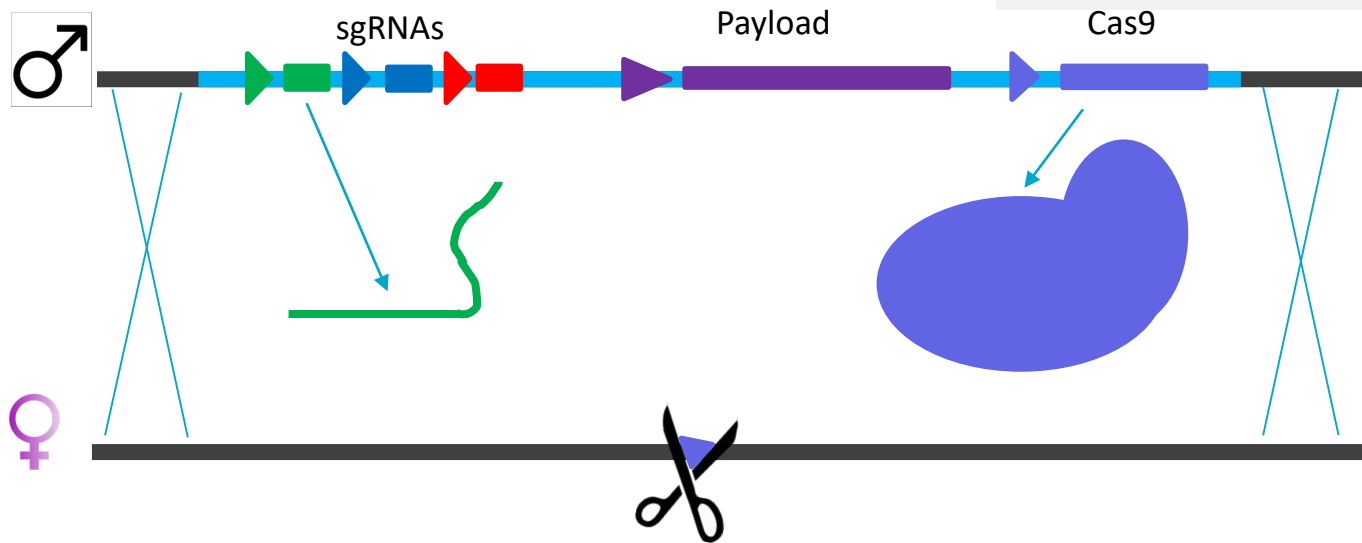
The components of a gene drive system:

Genes encoding the Guide RNA(s)

Payload gene with biological impact
e.g. **SRY (sex bias)**

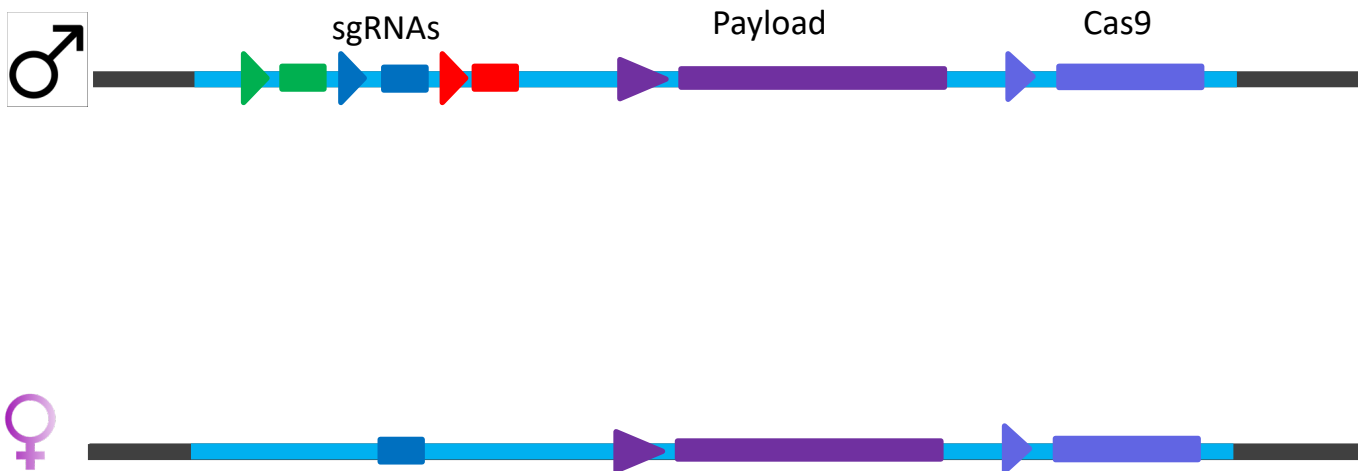
Gene for Cas9
Timing of Cas9 expression can be regulated

- Always 'on'
- Only in germline



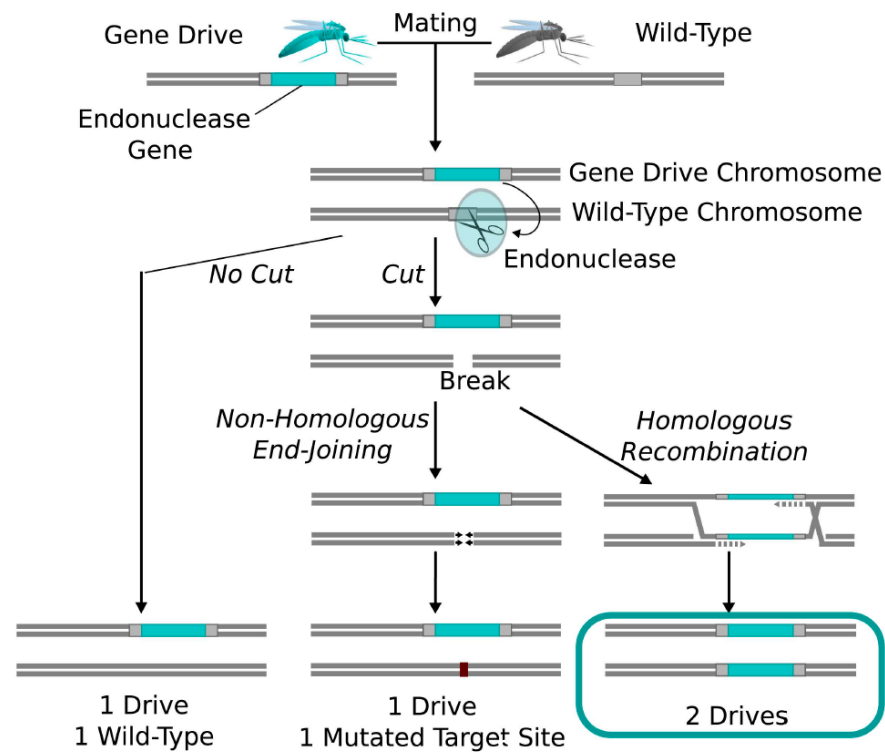


The components of a gene drive system



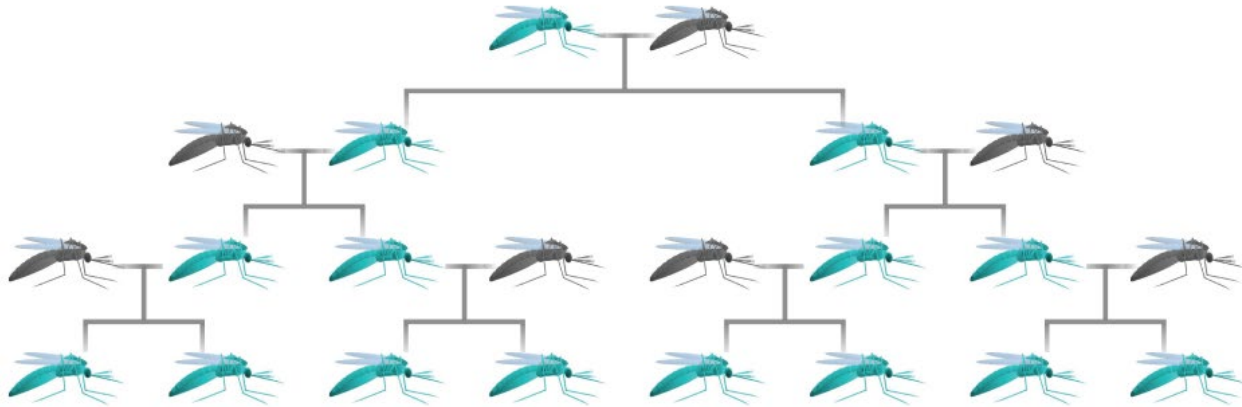


Gene drive mode of action (3 possible outcomes)



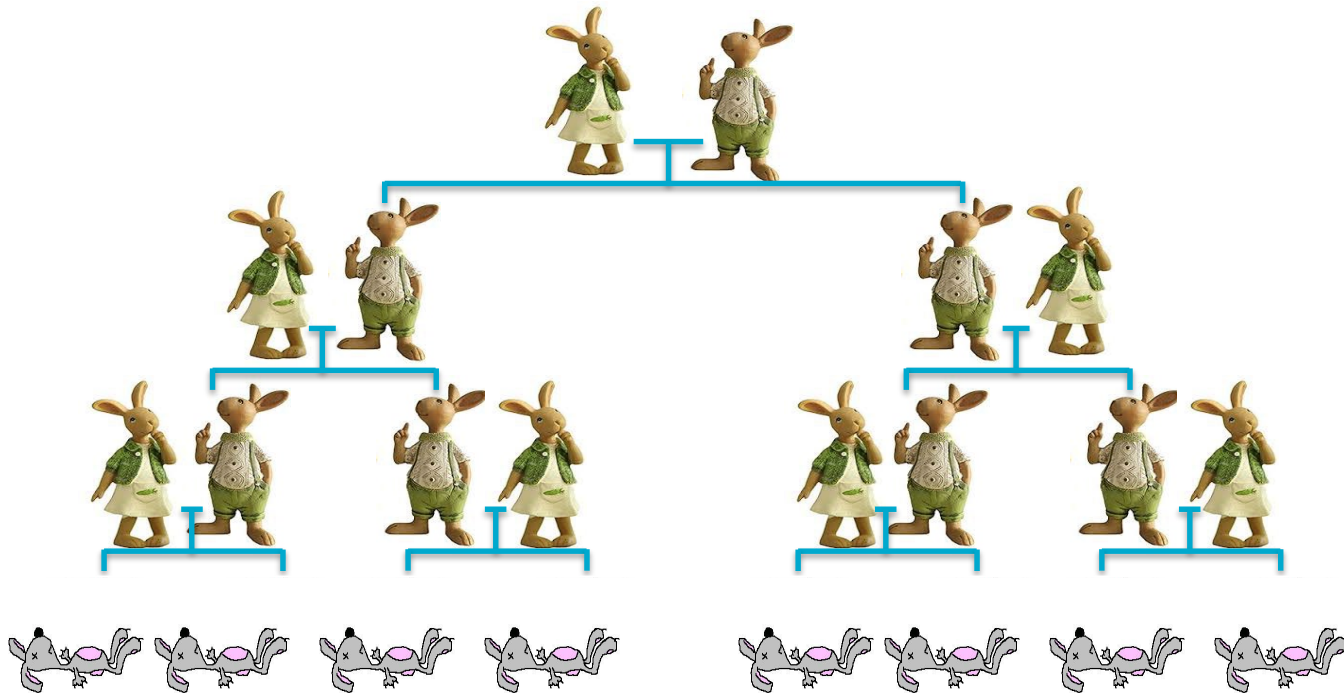
Esvelt et al., 2014 elife

Cas9-based gene drives



- 100% homing efficiency would lead to all offspring inheriting the trait (data from **insects and nematodes** show the real efficiency is between 60%-95%)
- In theory enables the drive to spread until it is present in all members of the population – even if it is mildly deleterious to the organism
- If trait was gene that turned all offspring into phenotypic males – population collapse

Cas9-based gene drives – biasing sex





Recent break through in mice !



bioRxiv

THE PREPRINT SERVER FOR BIOLOGY

bioRxiv posts many COVID19-related papers. A reminder: they have not been formally peer-reviewed and should not guide health-related behavior or be reported in the press as conclusive.

New Results

[Follow this preprint](#)

Leveraging a natural murine meiotic drive to suppress invasive populations

Luke Gierus, Aysegul Birand, Mark D. Bunting, Gelshan I. Godahewa, Sandra G. Piltz, Kevin P. Oh, Antoinette J. Piaggio, David W. Threadgill, John Godwin, Owain Edwards, Phillip Cassey, Joshua V. Ross, Thomas A.A. Prowse, Paul Q. Thomas

doi: <https://doi.org/10.1101/2022.05.31.494104>

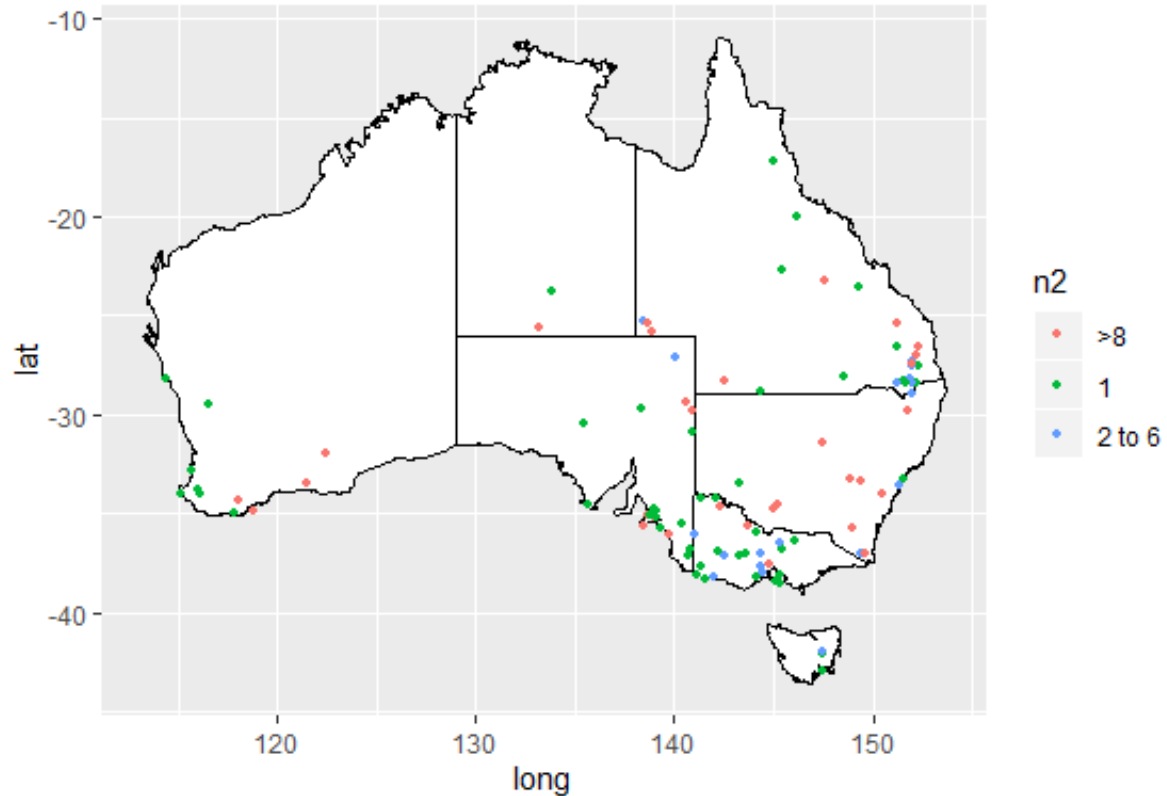
=> Transfer to rabbits?



Investigating the theoretical feasibility of genetic biocontrol for Australian wild rabbits

Know your genetics!

- Whole genome sequencing of ~300 individual Australian rabbits
- + ~180 from European collaborators





Key questions the dataset can answer:

Safety:

- Understand genetic diversity of Australia's wild rabbits
- Look for unique target sites (“Private alleles”) that an Australian rabbit specific
 - shared amongst all Australian populations or just sub-populations?
 - absent (or at low frequency only) in Rabbits from the European continent?

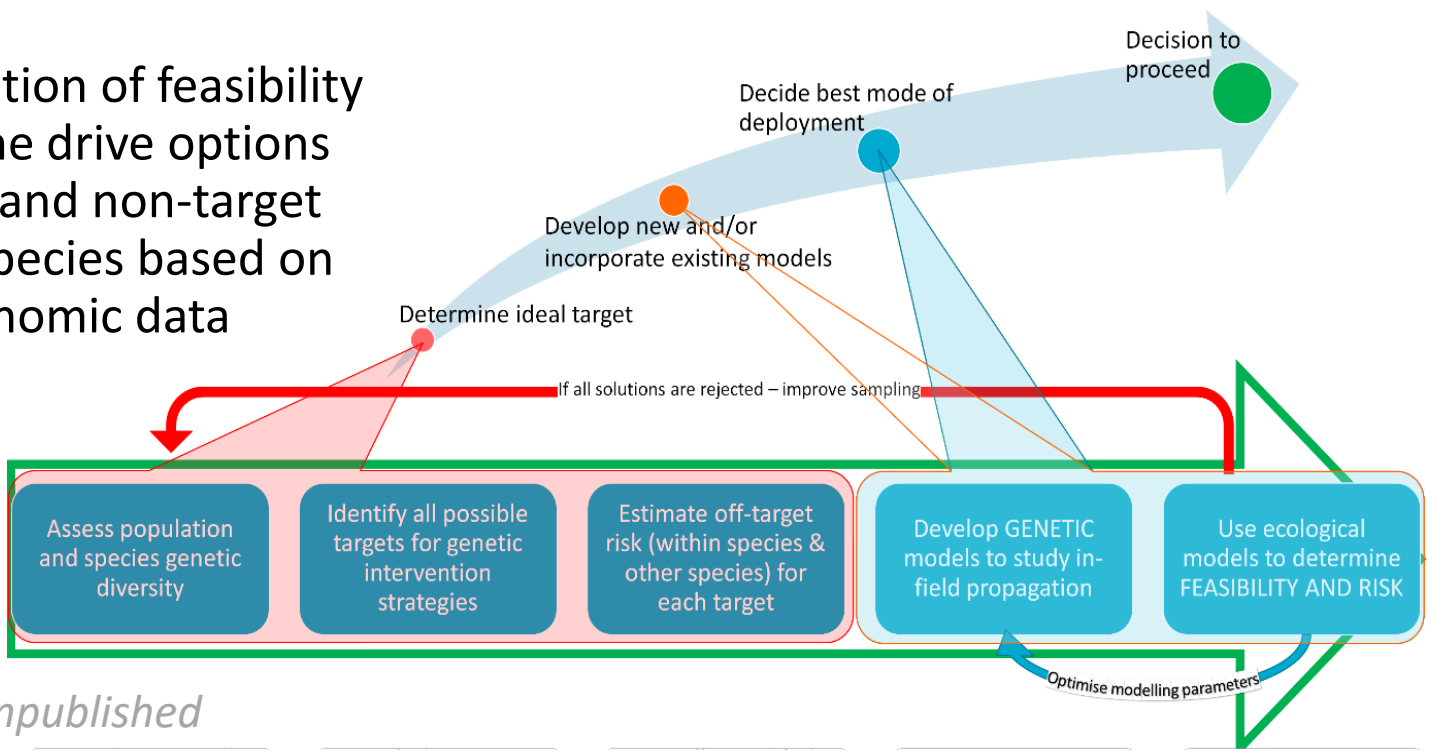
Potential efficacy:

- Based on the genetic structure of the population, the spread of a theoretical genetic biocontrol can be modelled



Gene drive Utility And Risk Determination pipeline (GUARD)

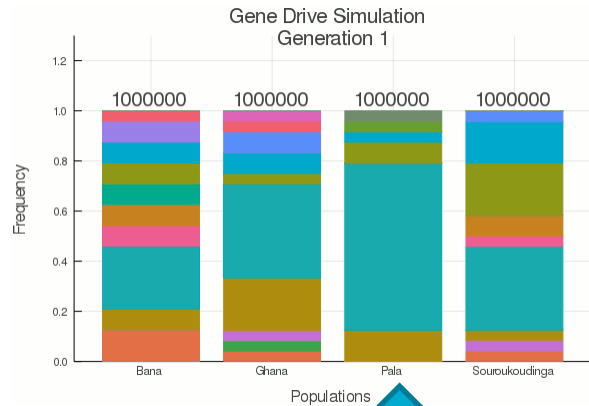
- *In silico* evaluation of feasibility and risk of gene drive options against target and non-target populations/species based on population genomic data



Yeap, Rane et al, unpublished



Example: Population sensitive gene drives for mosquitoes



Population data to analyse

Gene drive Utility And Risk Determination pipeline (GUARD)

Input data

1. PopGen Stats

2. On and off-target assessment

3. Genomic modelling

Select Data

- Anopheles coluzzi
 - Angola
 - Luanda
 - Burkina Faso
 - Bana
 - Pala
 - Souroukoudinga
 - Cote d'Ivoire
 - Ghana
 - Guinea
 - Anopheles gambiae
 - Burkina Faso
 - Bana
 - Pala
 - Souroukoudinga
 - Cameroon
 - Equatorial Guinea
 - France
 - Gabon
 - Gambia, The
 - Ghana
 - Guinea
 - Guinea-Bissau
 - Kenya
 - Uganda

Show Selected Pop

Metadata to select

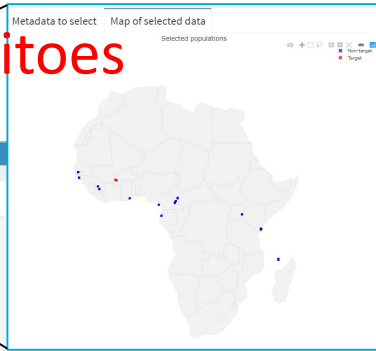
Map of selected data

Select target population OR unselect non-target species

	species	country	location
<input checked="" type="checkbox"/>	Anopheles gambiae	Burkina Faso	Bana
<input checked="" type="checkbox"/>	Anopheles gambiae	Burkina Faso	Pala
<input checked="" type="checkbox"/>	Anopheles gambiae	Burkina Faso	Souroukoudinga
<input type="checkbox"/>	Anopheles gambiae	Ghana	Madina
<input type="checkbox"/>	Anopheles gambiae	Guinea	Koundara
<input type="checkbox"/>	Anopheles gambiae	Guinea	Koraboh
<input type="checkbox"/>	Anopheles gambiae	Cameroon	Daiguene
<input type="checkbox"/>	Anopheles gambiae	Cameroon	Gado Badzere
<input type="checkbox"/>	Anopheles gambiae	Cameroon	Mayos
<input type="checkbox"/>	Anopheles gambiae	Cameroon	Zembe Borongo
<input type="checkbox"/>	Anopheles gambiae	Equatorial Guinea	Bioko

Modules in tabs

Selecting target population



Gene drive Utility And Risk Determination pipeline (GUARD)

Input data

1. PopGen Stats

2. On and off-target assessment

3. Genomic modelling

Show 10 entries

	Chromosome	start	end	sequence	on.target.score	target.gene	GC.content	OffTargetScore
35	2R	58693860	58693882	ATGCTGATATCGAAACTGGGAGG	5.38120254177425	AGAP0132218	52.381	0.0135676879552193
40	2R	57471525	57471547	ACATACCAGCAGACTGCGTGGG	5.35937531367001	AGAP004541	61.9048	0.049745944153983
7	2R	55882950	55882612	CGATAGATGATTCCTCTGTGGGG	5.61287979545557	AGAP004424	57.1429	0.0213360564783216
11	2R	55842373	55842395	TACCAGTAGAAGGAACCTGAGAG	5.52144932448955	AGAP004411	52.381	0.0309387238579802
24	2R	54122382	54122392	GATGTTTAGCTGCAACTGCGAGG	5.423648762365	AGAP004288	57.1429	0.0409402310731821
27	2R	53250864	53250886	AAGGTACTACAATCTCGGAGGG	5.41583722199569	AGAP004249	52.381	0.022623475452001
31	2R	51134310	51134330	CAGGTCCAATGGGAATCGGTGG	5.38781493941361	AGAP004169	66.6667	0.004200314939995378
17	2R	49544051	49544073	AAGTCGAATTATGGCACTGCGCGG	5.49740306821081	AGAP004090	61.9048	0.0303508668555878
51	2R	48714637	48714659	GTTTAACACAGGCTCAAGGCGGG	1.8226323	AGAP004050	52.1739	0.00817233264679089
21	2R	47592356	47592378	CAAATATTGCACGAGCTGCGCGG	5.45339385747321	AGAP013163	61.9048	0.0102652599220164

Showing 1 to 10 of 51 entries

Previous 1 2 3 4 5 6 Next





Future work:

Can we transfer the functionality of the mouse proof of concept into rabbits?

=> Internal CSIRO investment into GBC (3 year).....

watch this space !



Proceed – but with caution!

Science

Social
License,
Liaison

Risk
assessment

Gene
technology
regulators

Public perspectives towards using gene drive for invasive species management in Australia

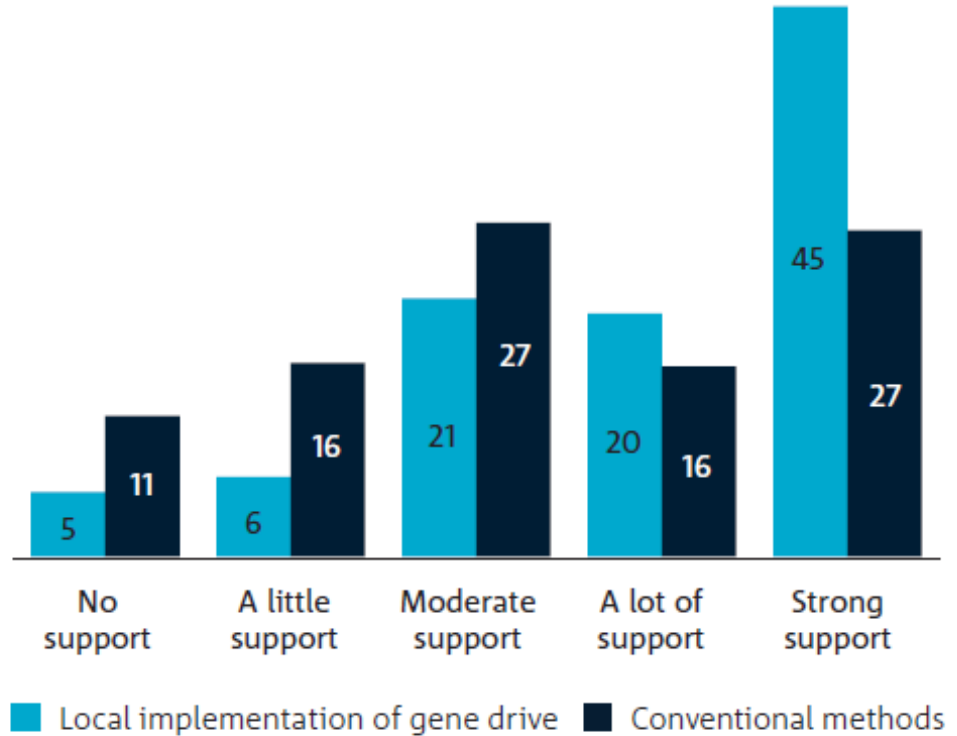


Figure 8. Levels of support for the local implementation of gene drive technology and conventional cat control methods, as a percentage (%) of total responses.



Genetic control technologies – unique *challenges and opportunities for Australian pest animal management*

Is it worth taking this risk? What is the risk of not trying?





Thank You

Recent RHDV and/or MYXV work



Robyn Hall
 Egi Kardia
 Maria Jenckel
 Lena Smertina
 Nina Huang
 Ina Smith
 Matt Neave
 Rahul Rane
 HengLin Yeap
 Angel Popa
 Peter Kerr
 June Liu
 Ros Mourant
 Tegan King
 Mel Piper

Hugh Mason
 Megan Pavy
 Madi Rutherford
 Maria Jenckel
 Omid Fakhri
 Tom Walsh
 Tek Tay
 Raghu
 Sathyamurthy
 Mark Tizard
 Agus Sunarto
 Caitlin Cooper



THE UNIVERSITY OF
SYDNEY

Eddie Holmes
 Jackie Mahar
 JS-Eden
 Mang Shi



UNIVERSITY OF
CANBERRA
AUSTRALIA'S CAPITAL UNIVERSITY

Michael Frese
 Brian Cooke
 Peter Elsworth



Department of
Primary Industries

Tarnya Cox
 Emma Sawyers
 Glen Saunders
 Pat Taggart



Government
of South Australia
Primary Industries
and Regions SA

Greg Mutze
 Dave Peacock
 John Kovaliski
 Kandarp Patel

BCM
Baylor College of Medicine

Mary K Estes



CENTRE FOR
INVASIVE SPECIES SOLUTIONS



Australian Government
Department of Agriculture
and Water Resources



Economic Development,
Jobs, Transport
and Resources

Dave Ramsey
 Carlo Pacioni

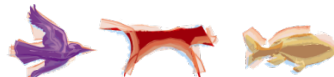


Lorenzo Capucci



UNIVERSITY OF
CAMBRIDGE

Joel Alves
 Frank Jiggins



Invasive Animals CRC



AWI Australian Wool
Innovation Limited



PennState
College of Agricultural Sciences

Andrew Read
 Isabella Cattadori



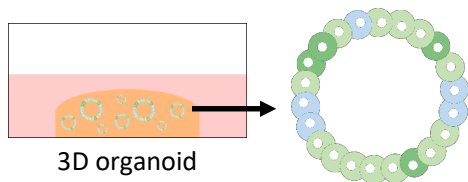
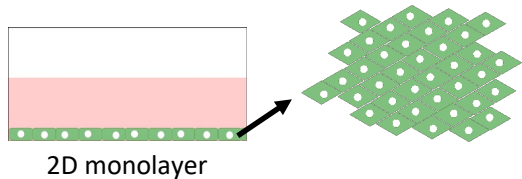
Miguel Carneiro







We need a robust cell culture system/reverse genetics



3D culture

- Multicellular
- More relevant model
- Mimic the architecture of the parental tissue



➤ **Developing rabbit intestinal and liver organoids to study enterotropic and hepatotropic lagoviruses**

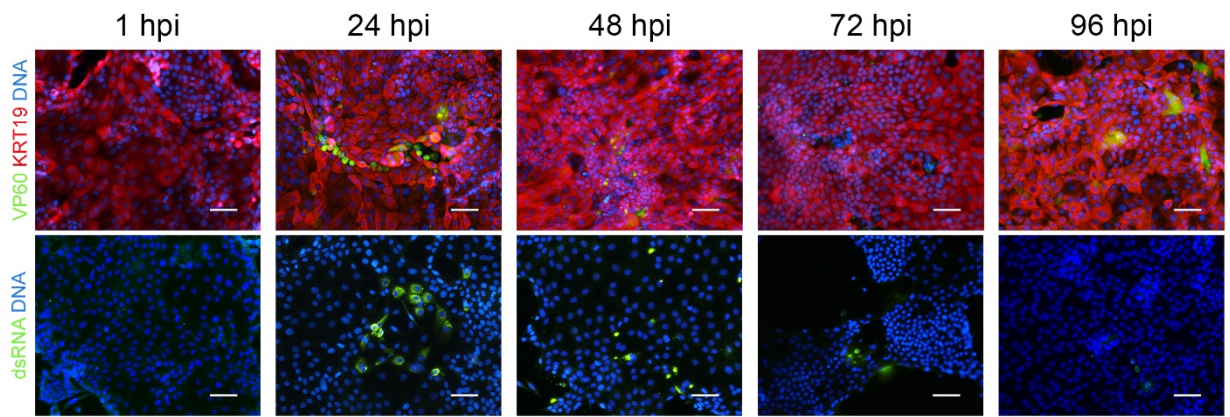
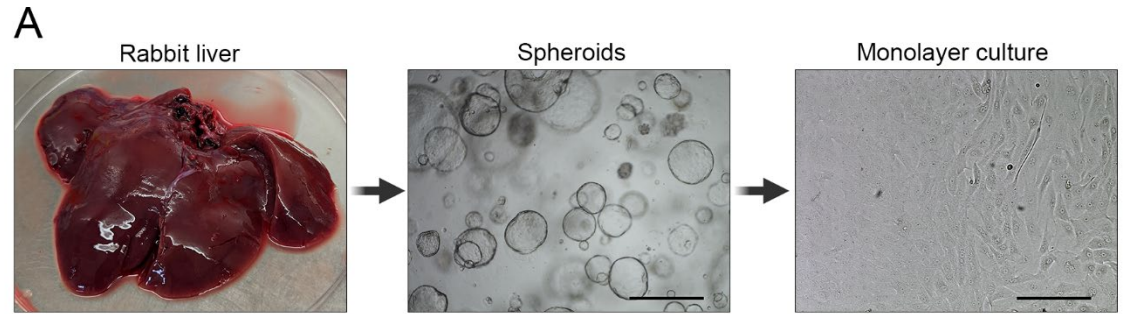
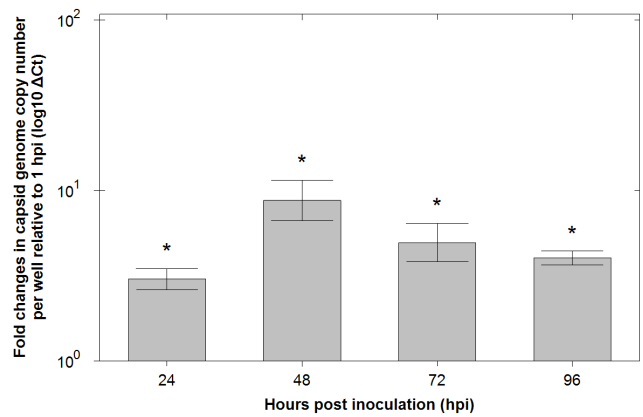
- Replication mechanisms
- Virulence mechanisms
- Tissue tropism/virulence evolution
- Accelerated natural selection of immune variants



Mary K Estes



Rabbit liver organoids: replication proof of concept



bioRxiv posts many COVID-19-related papers. A reminder: they have not been formally peer-reviewed and should not guide health-related behavior or be reported in the press as conclusive.

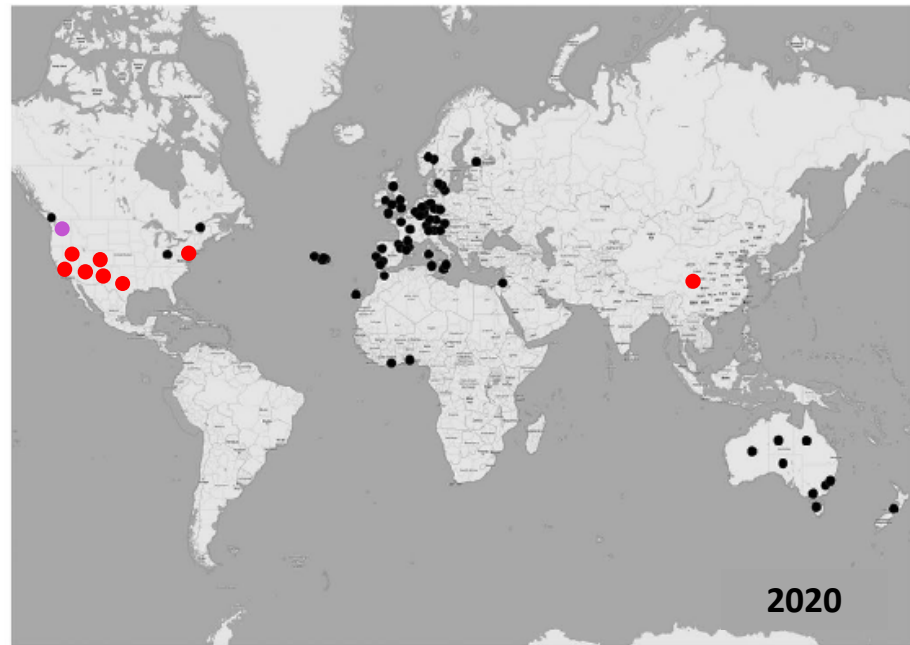
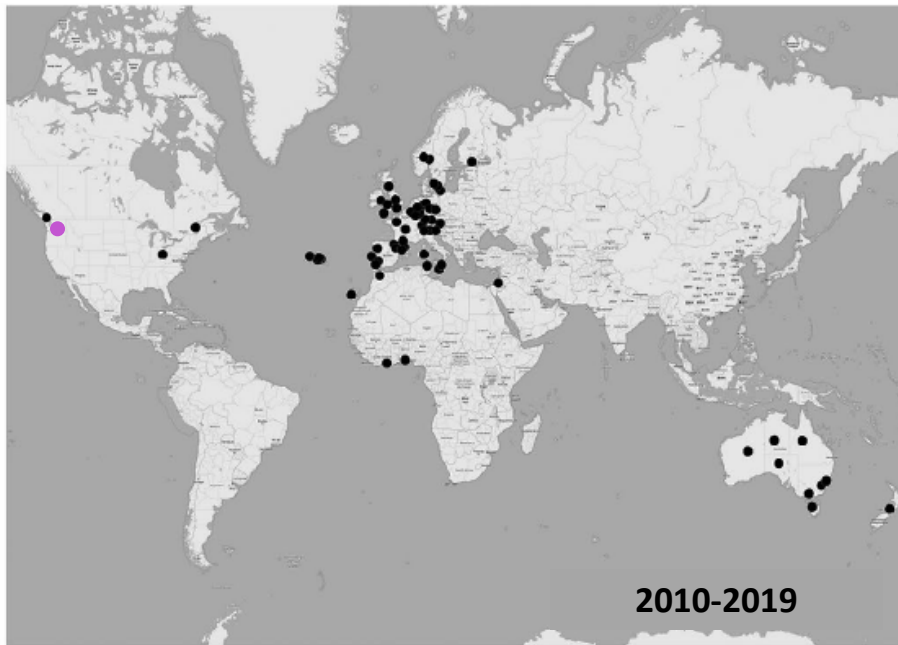
New Results [Follow this preprint](#)

Hepatobiliary organoids derived from leporids support the replication of hepatotropic lagoviruses

Egi Kardia, Omid Fakhri, Megan Pavy, Hugh Mason, Nina Huang, Elena Smertina, Mary K. Estes, Tanja Strive, Michael Frese, Robyn N. Hall
doi: <https://doi.org/10.1101/2022.04.07.487566>



Global Spread of RHDV2: 2010 - 2020





RHDV2 in the southwestern US

- First reported in Spring 2020
- A series of native lagomorphs affected



Sylvilagus floridanus



Lepus californicus



Sylvilagus audubonii

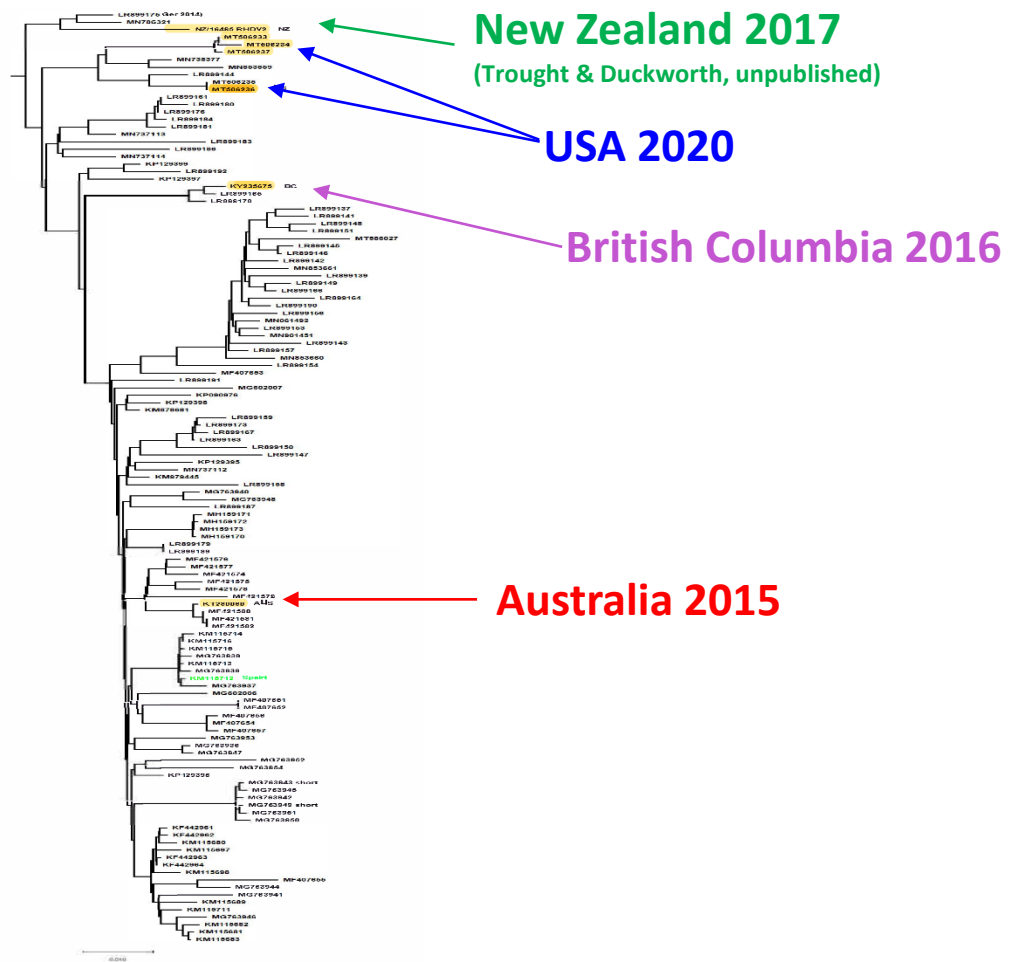


Sylvilagus nuttallii



Lepus alleni





Central Europe

Southern Europe/
Iberian Peninsula





Evolution of virulence in lagoviruses

Capucci et al 1996

Strive et al 2009

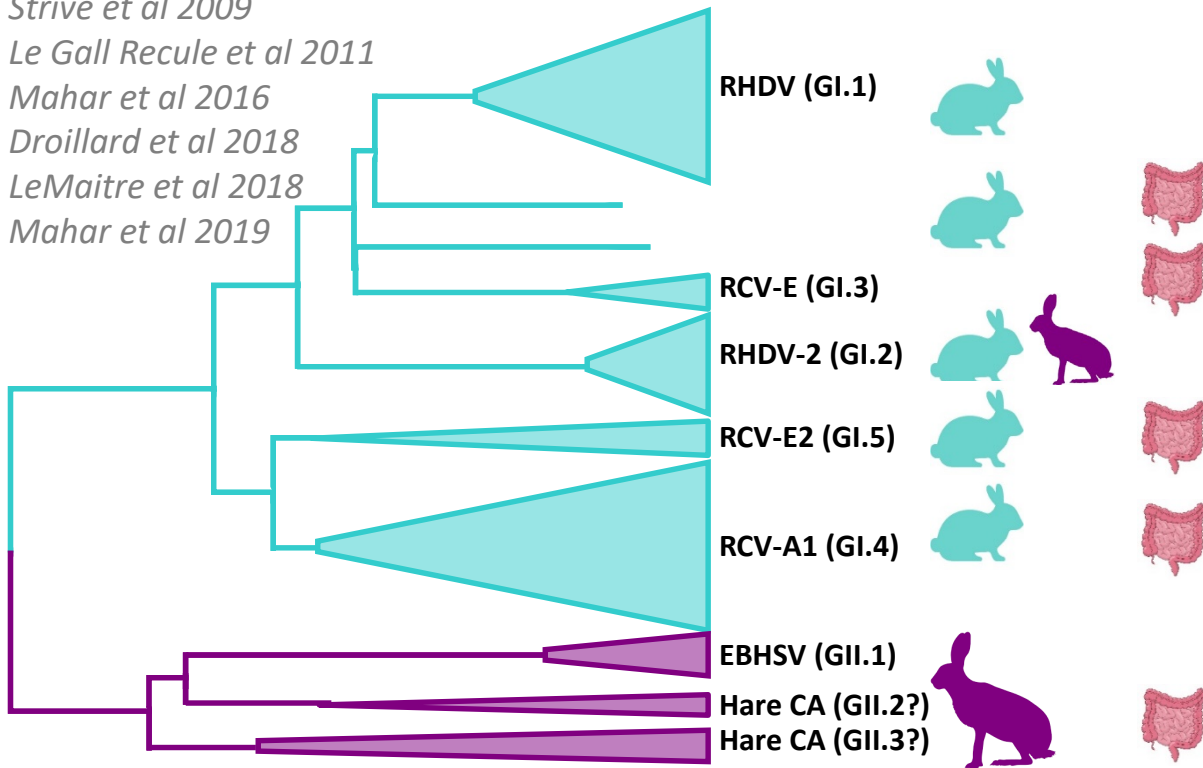
Le Gall Recule et al 2011

Mahar et al 2016

Droillard et al 2018

LeMaitre et al 2018

Mahar et al 2019



Mahar et al., 2016, Mahar et al., 2019



Evolution of virulence in lagoviruses

Capucci et al 1996

Strive et al 2009

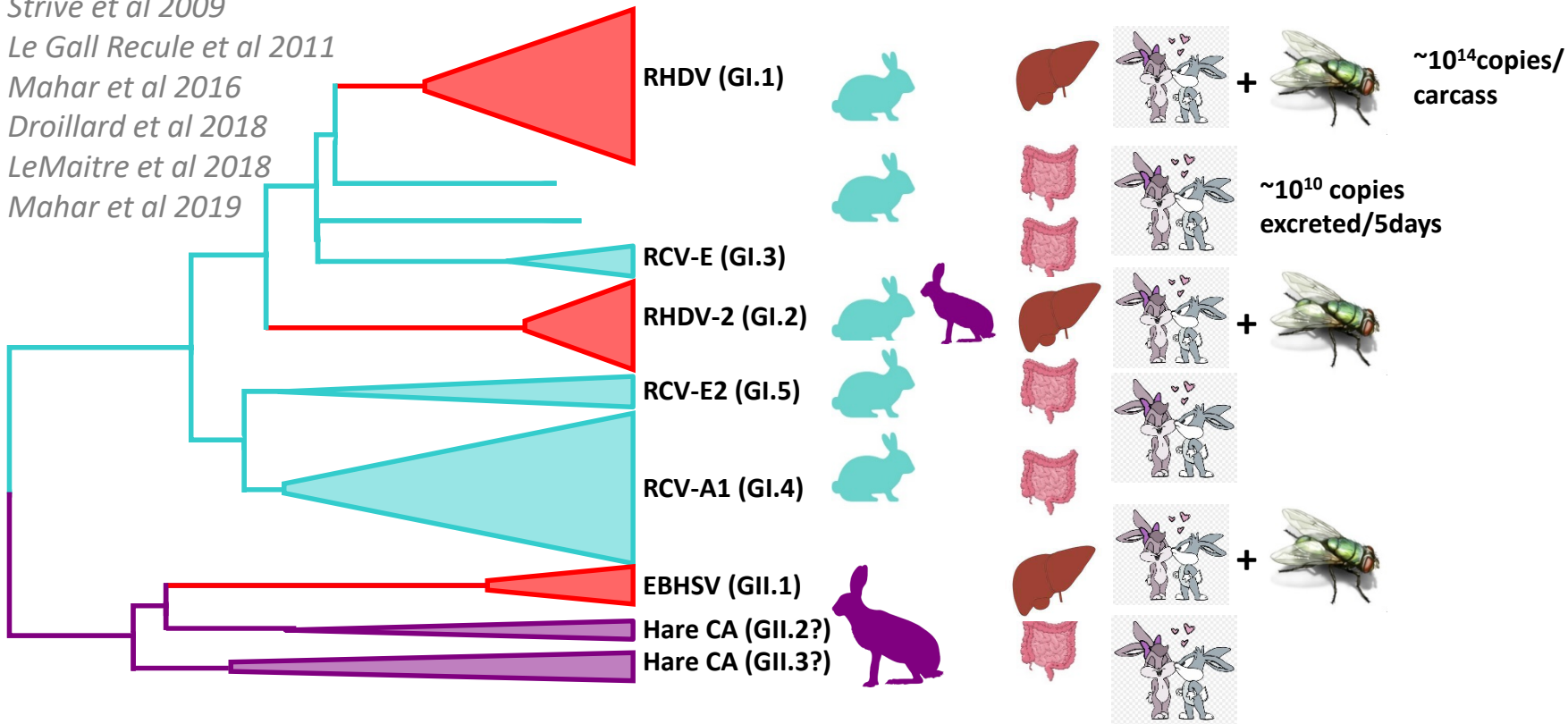
Le Gall Recule et al 2011

Mahar et al 2016

Droillard et al 2018

LeMaitre et al 2018

Mahar et al 2019





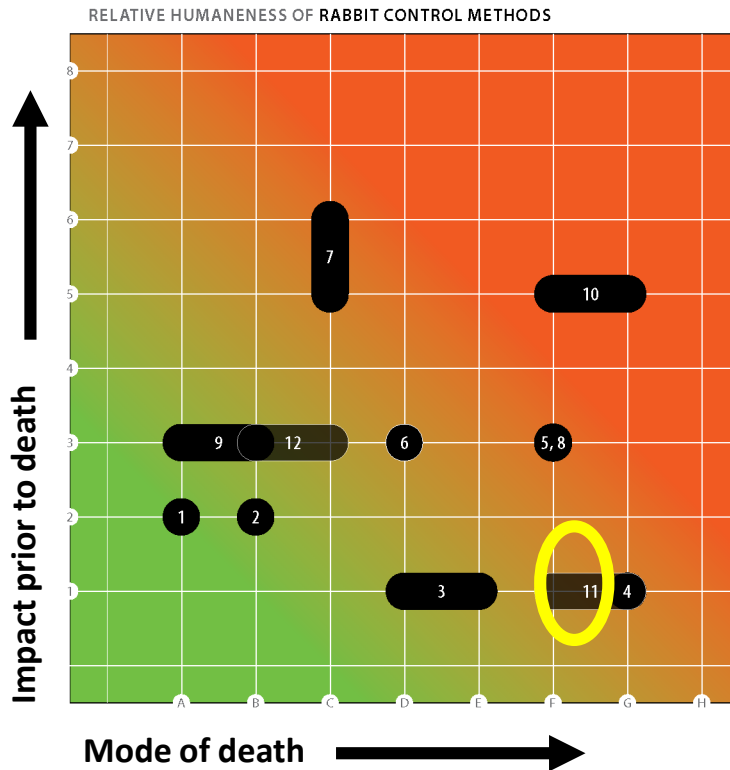
Animal Welfare

Is an integral component of pest animal control

Factor to be considered when choosing the appropriate tool for pest animal management

What is the consequence of doing nothing?

Scoring Matrix to assess the welfare impacts of rabbit control methods approved in Australia



Sharp and Saunders 2012

- Shooting (head)
- Shooting (chest)
- (Warren blasting)
- LPG Warren fumigation
- 1080 poison
- Phosphine fumigation
- Baits RHDV
- Chloropicrin fumigation
- Warren ripping
- Pindone anticoagulant
- RHDV inoculation
- Padded foot hold trap