

CRISPR/Cas9-based Gene Drive to Suppress Agricultural Pests



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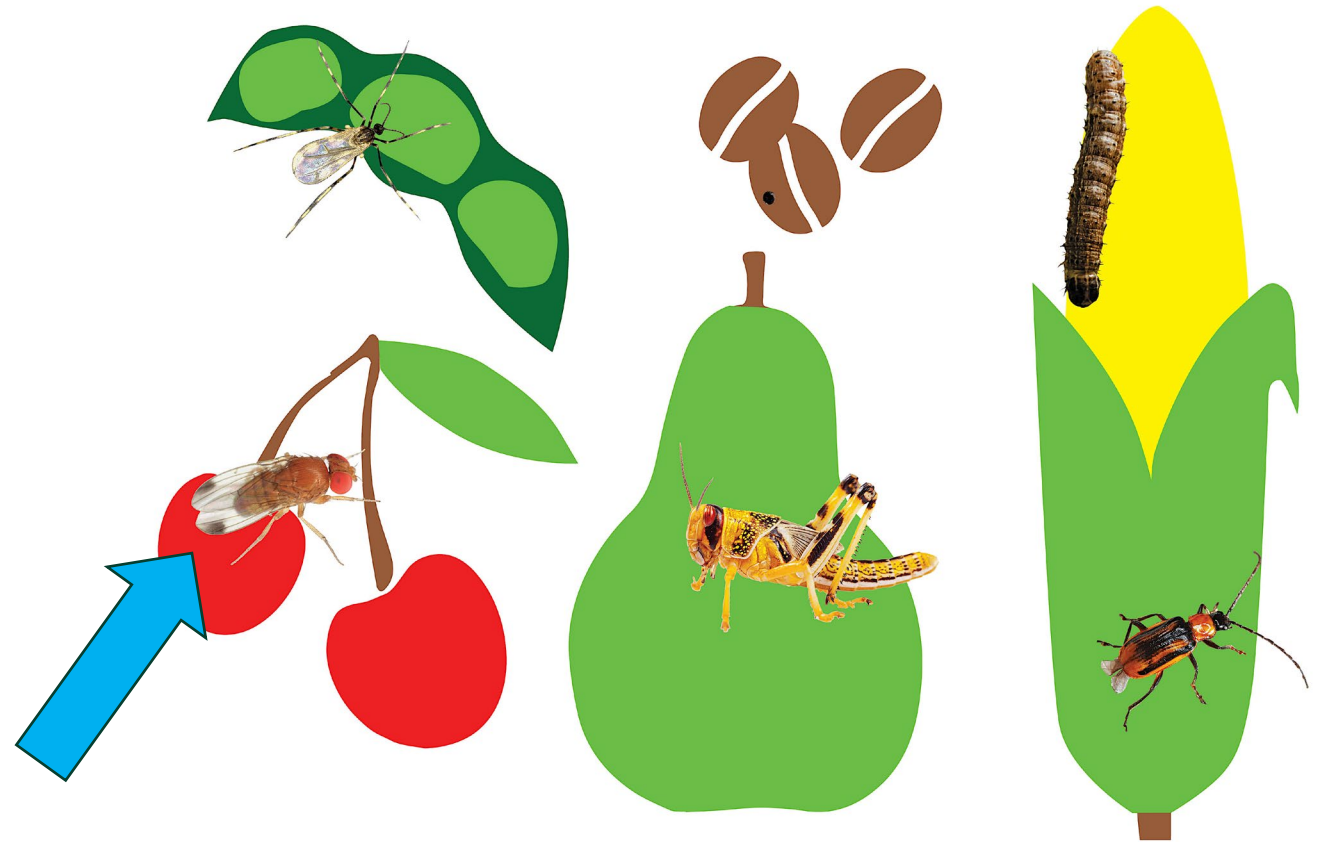
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Talk Overview

- Agricultural pests **genetic biocontrol**
- CRISPR/Cas9-based **gene drives development (in SWD)**
- **Challenges** in gene drives development and its application

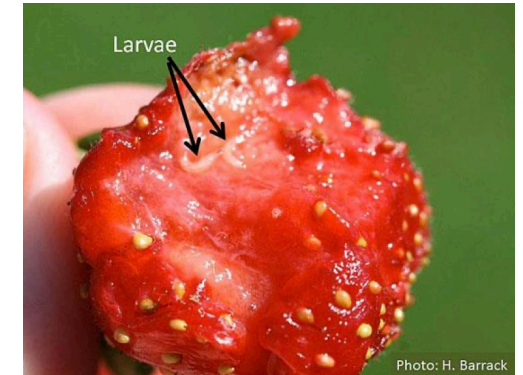
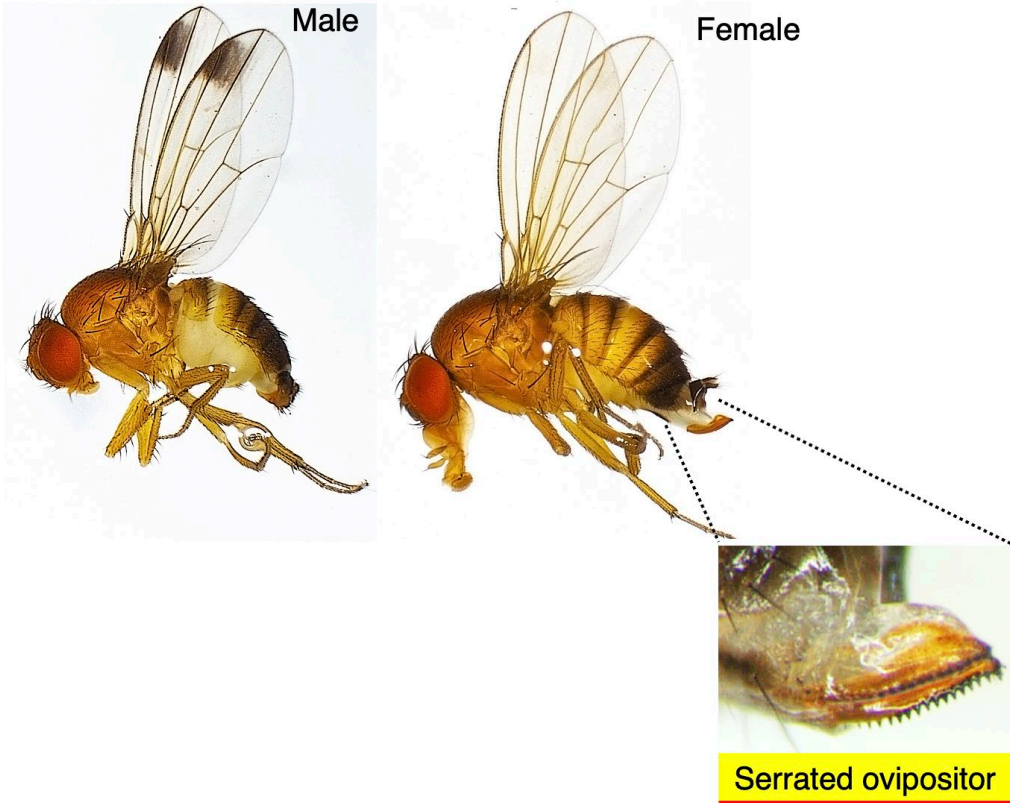
Agricultural Pests impact

- Threat to agricultural productivity and food-security
- Each year, **20% to 40%** of the global crop production lost to pests
- Annual economic loss ~\$540 billion



Spotted wing drosophila (SWD)

- Native to Southeast Asia
- Single Female can lay up to 300 eggs
- 3-16 overlapping generations in a year

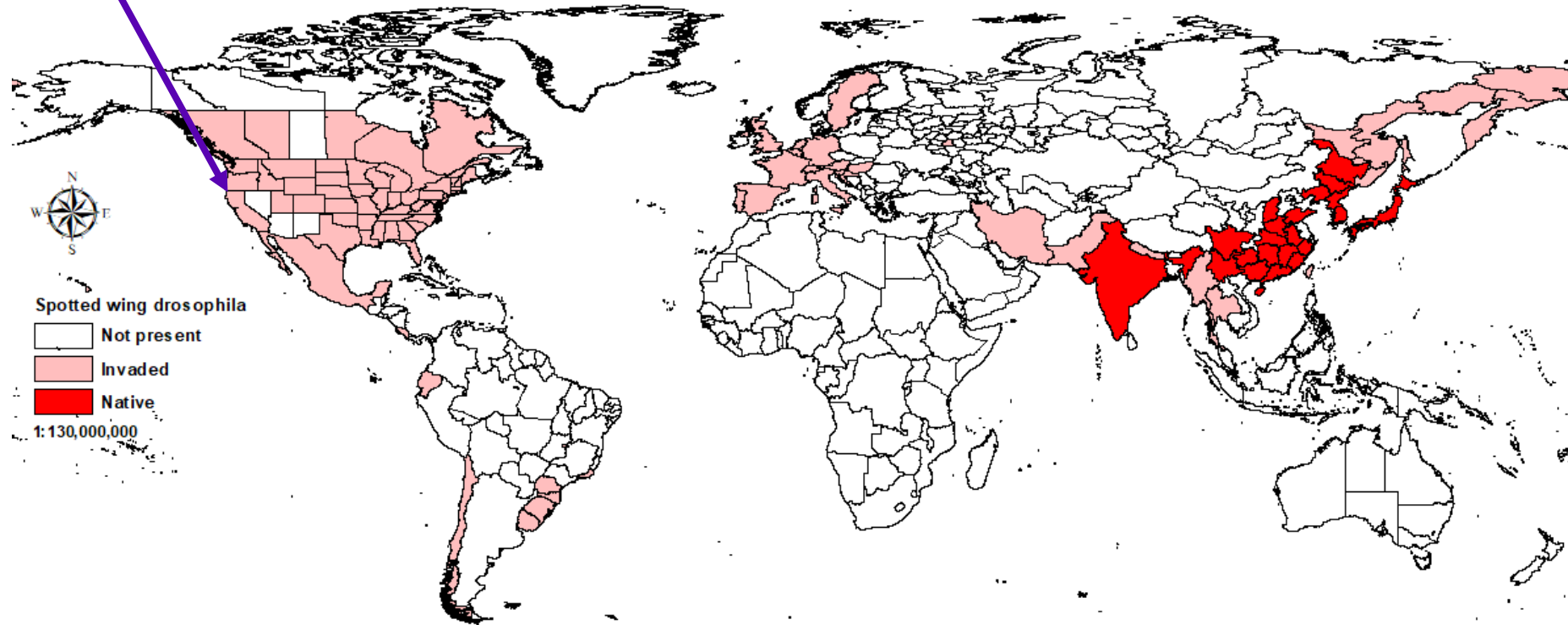


Damaged fruits, and growing larvae and Pupae inside fruits

Spotted wing drosophila: Global distribution

- Becoming a major agricultural **invasive pest** species in the USA and Europe

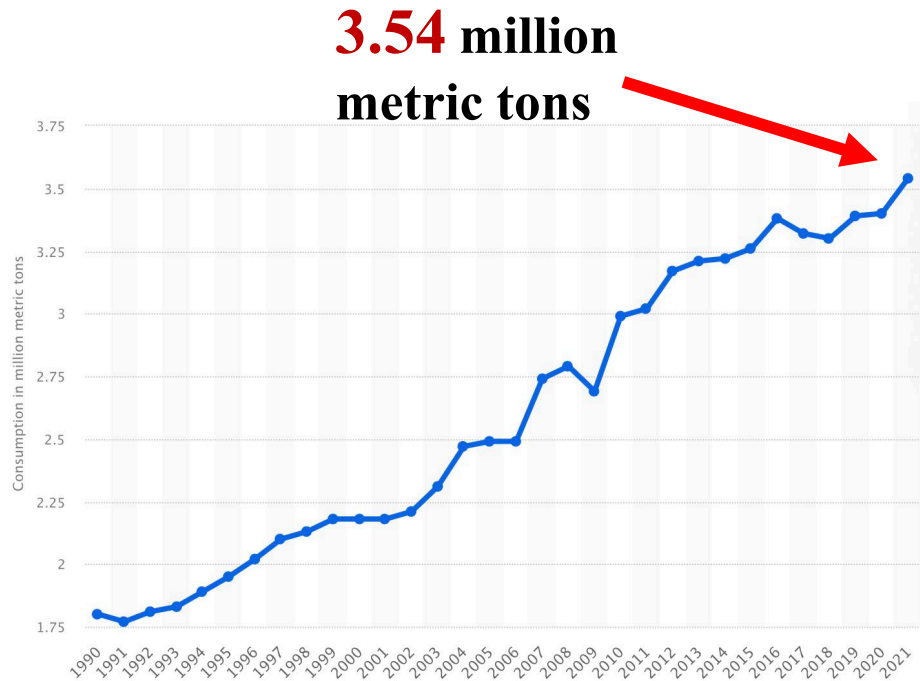
2008



Genetic Biocontrol an alternative to chemicals...

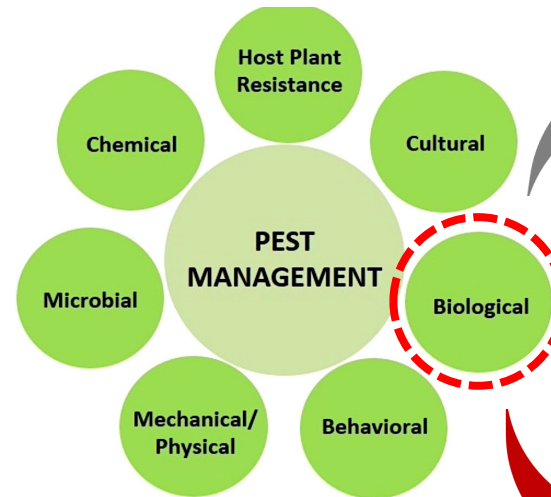
- Genetic control Reduces the reliance on pesticides

Consumption of **pesticides** worldwide from 1990 to 2021



Details: Worldwide; FAO; 1990 to 2021

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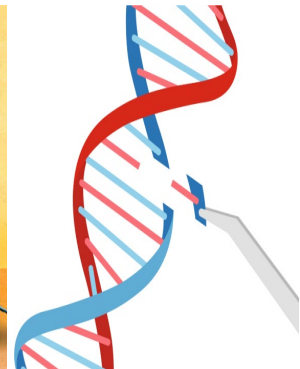
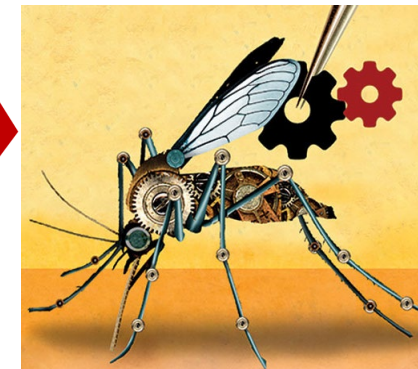


Genetic biocontrol

Natural predators

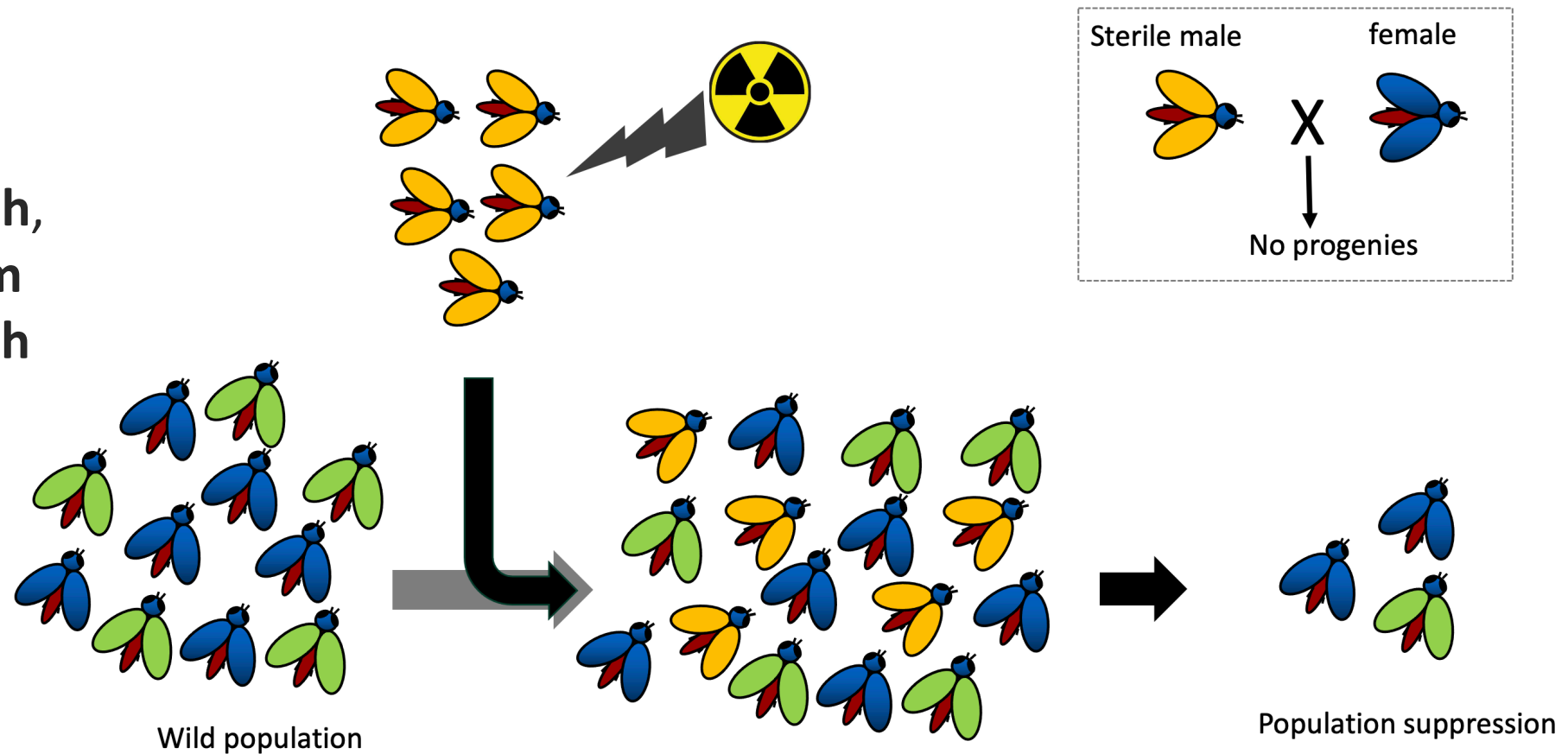


Genetically Engineered Pest Organism Or the natural incompatible variants



The Sterile Insect Technique (SIT)

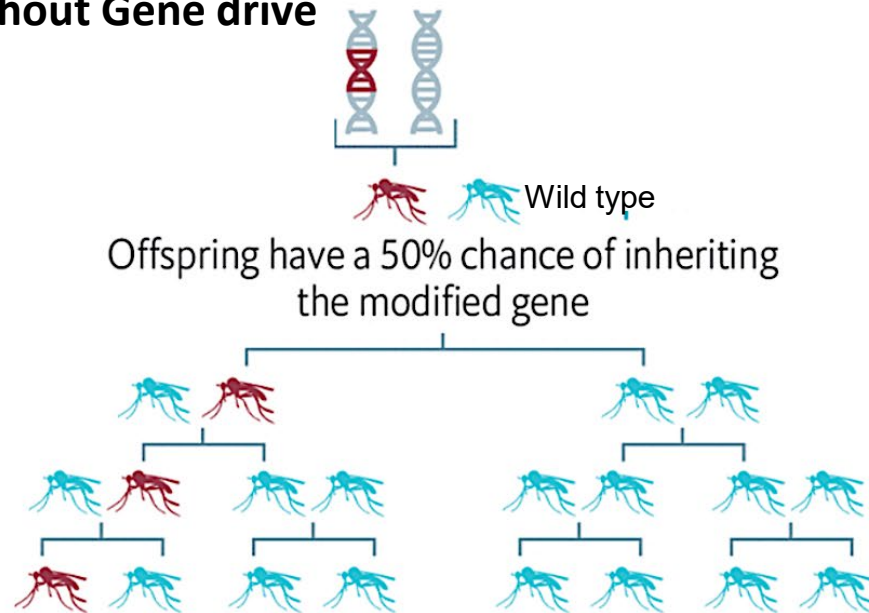
- Environmentally-friendly
- Successful eradication of **screw worms** from the southeastern United States
- Control of **codling moth**, **med fly**, **pink bollworm** and **painted apple moth** etc..
- Other methods: pgSIT, fsRIDL



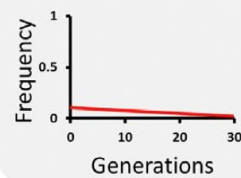
Gene drive

- A gene drive is a **phenomenon of biased inheritance** to introduce a genetic change into a population **at a rate higher** than normal inheritance.

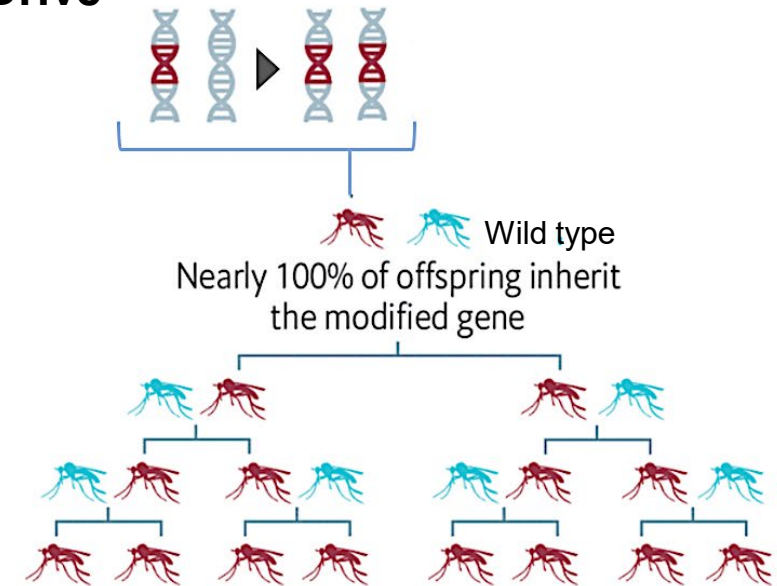
Without Gene drive



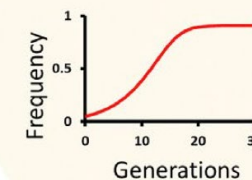
No spread



Gene Drive

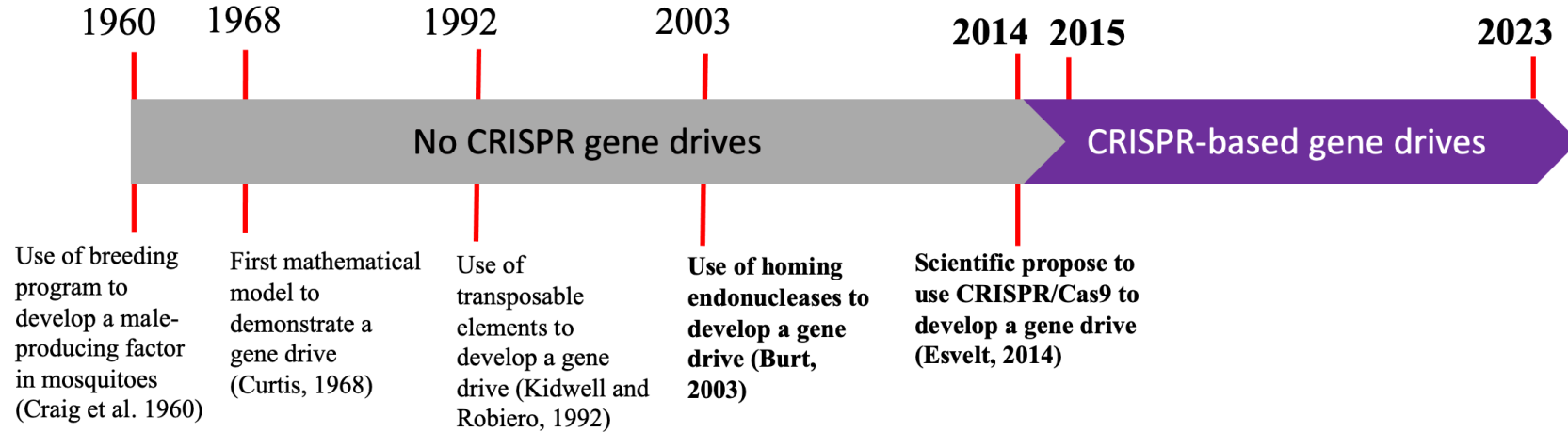


Spread

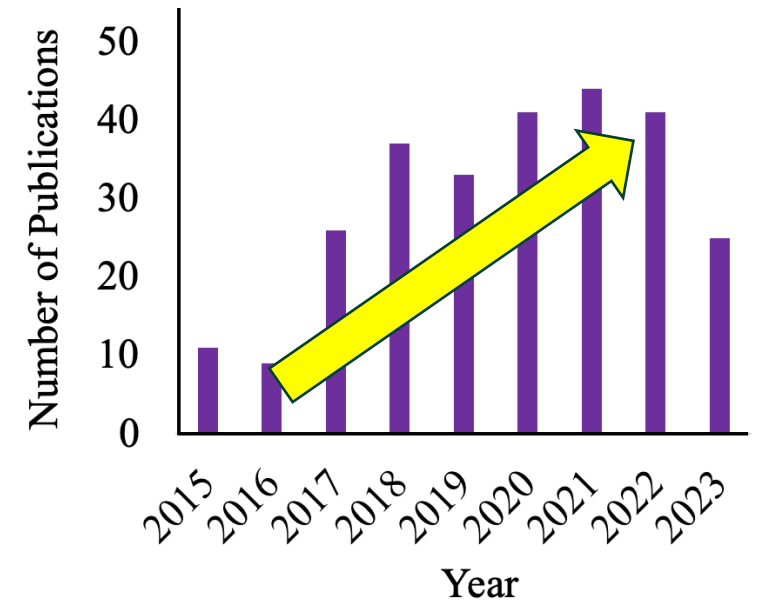


The timeline of Gene-drives

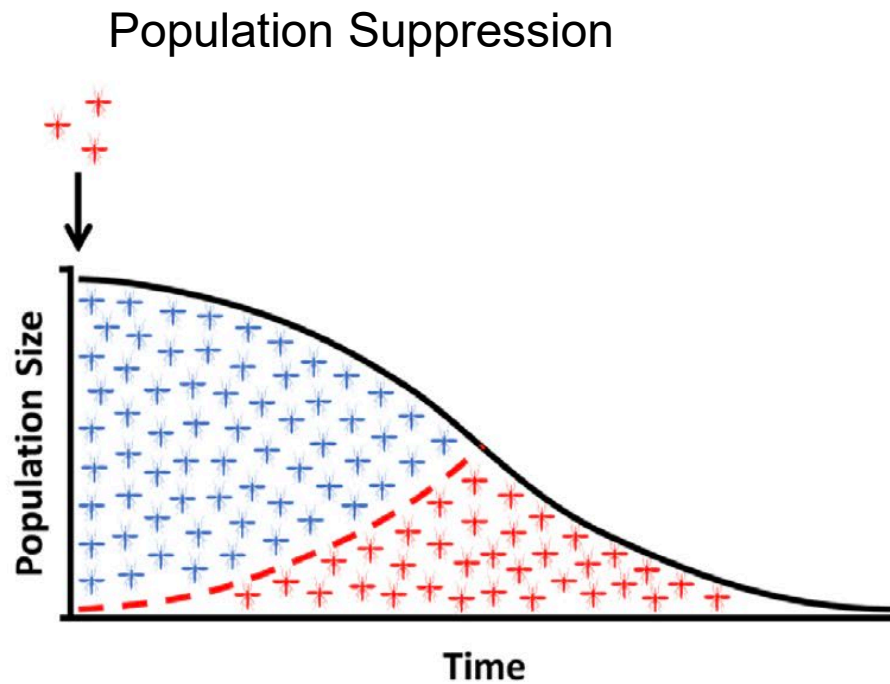
- Natural gene drives
- Synthetic gene-drives



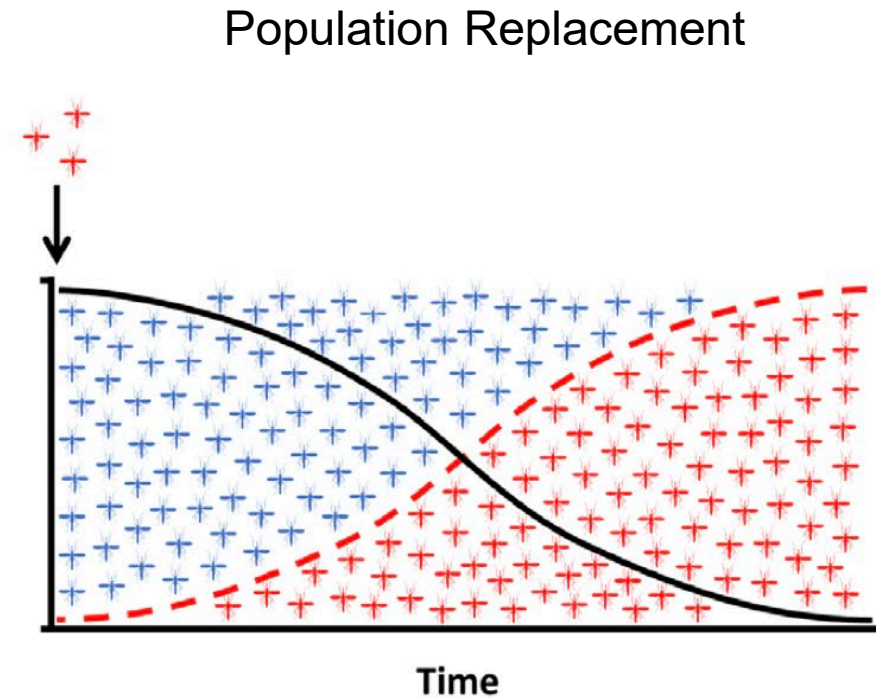
Research/Review articles on CRISPR-based gene drives since 2015



Gene drives potential Applications



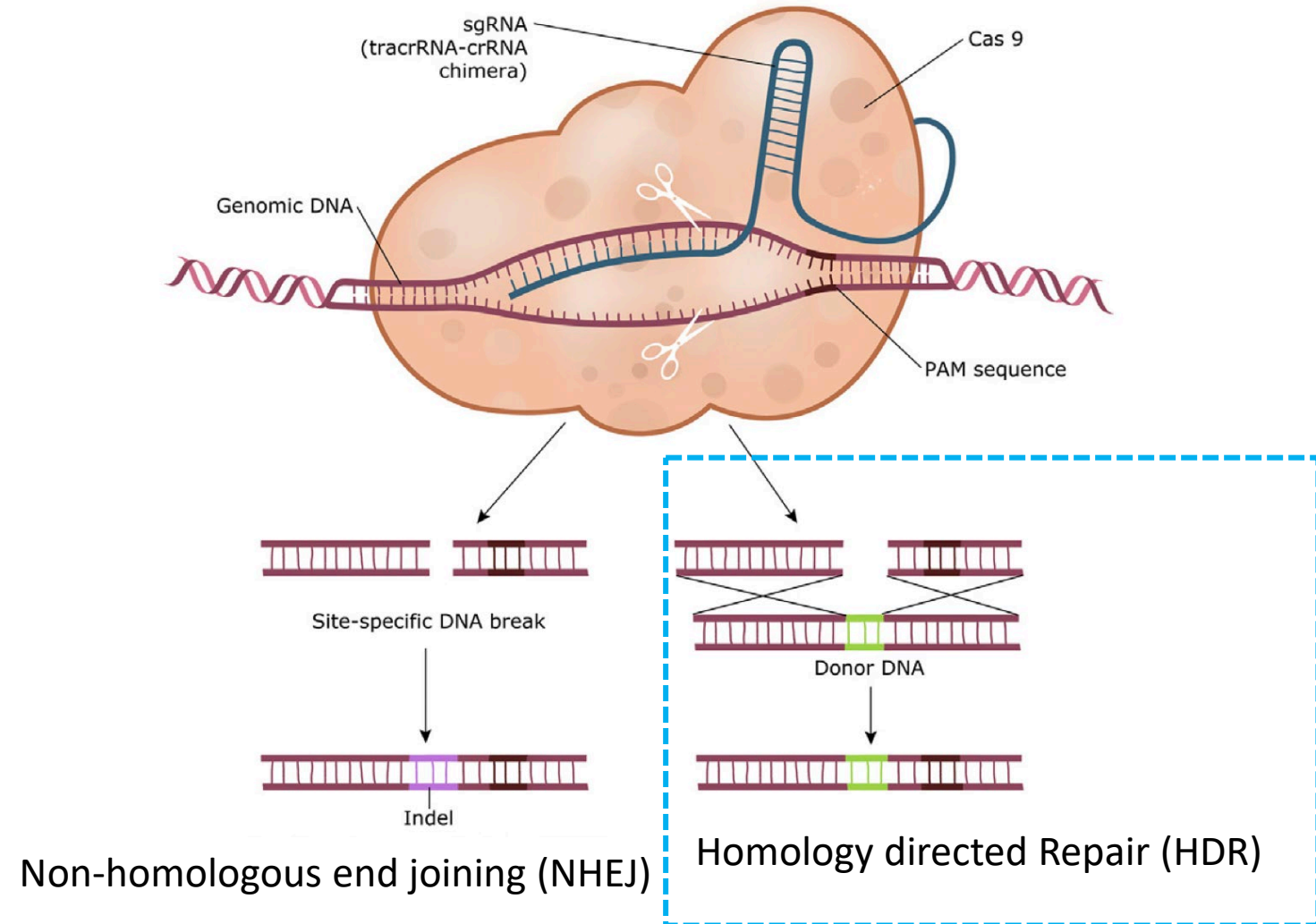
- Targets gene essential for development/female fertility
- Bias sex ratio



- Alter genes or add new gene to stop the pathogen transmission in a disease vector

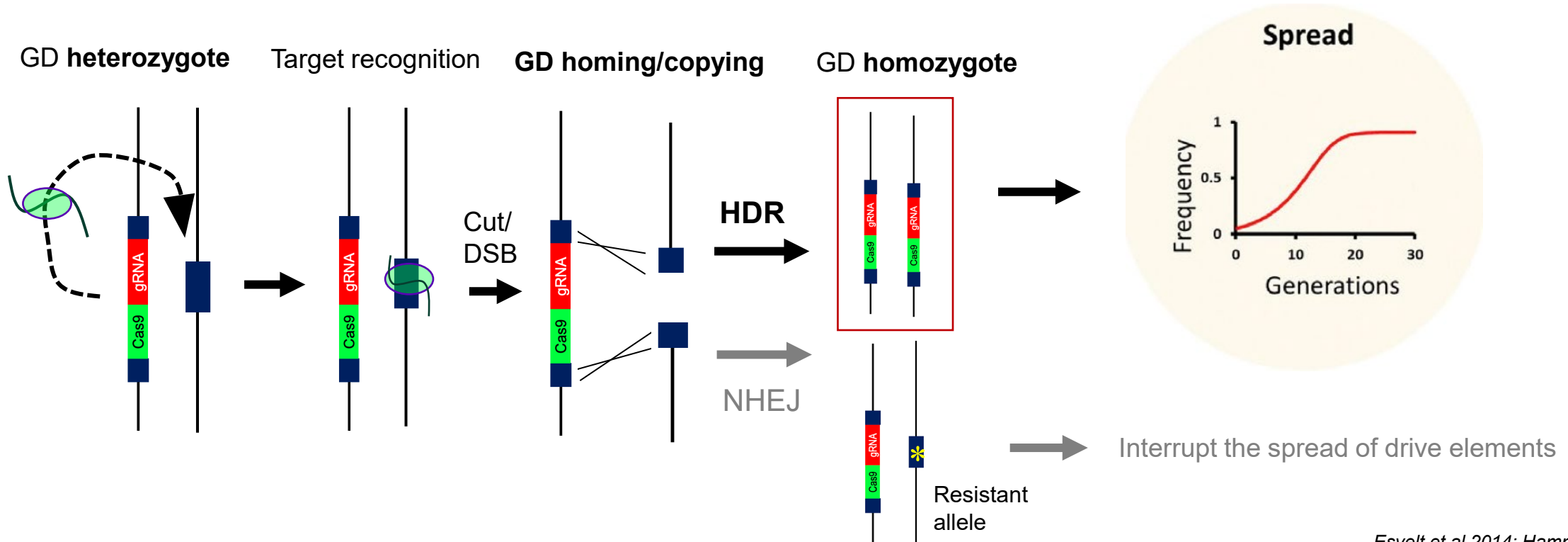
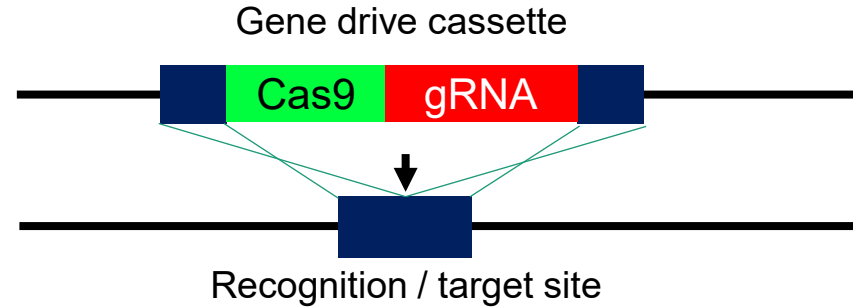
'CRISPR/Cas9' system

- Derived from the immune system of bacteria and archaea
- Uses RNA guided DNA endonuclease



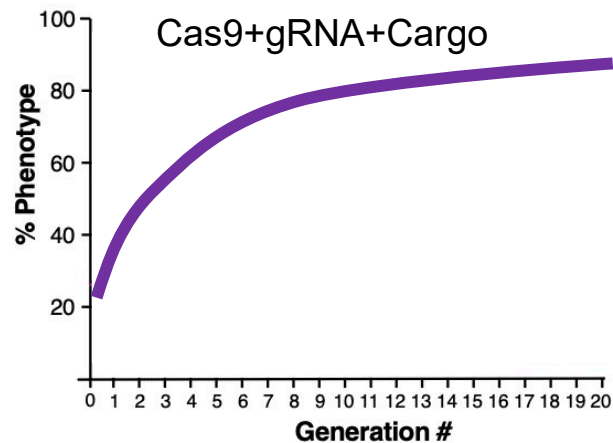
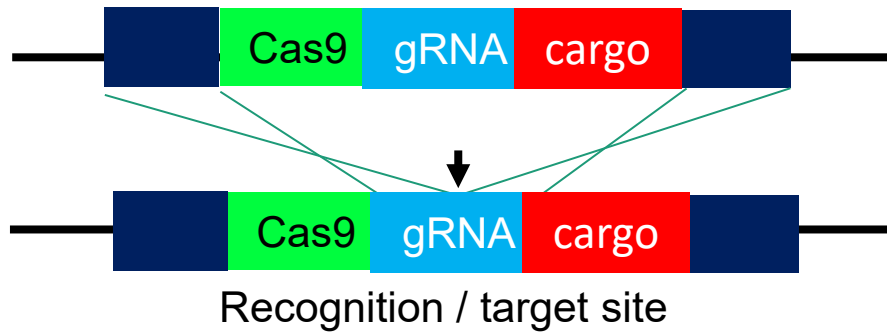
CRISPR/Cas9-based Gene drives design

- CRISPR/Cas9-based gene drive relies on **Homology Directed Repair** in Germ cells

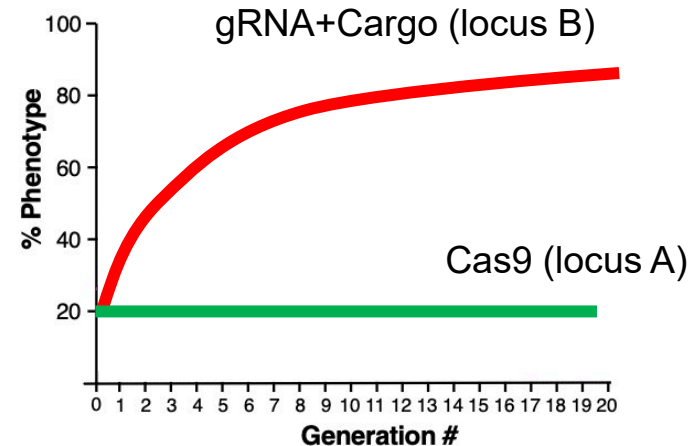
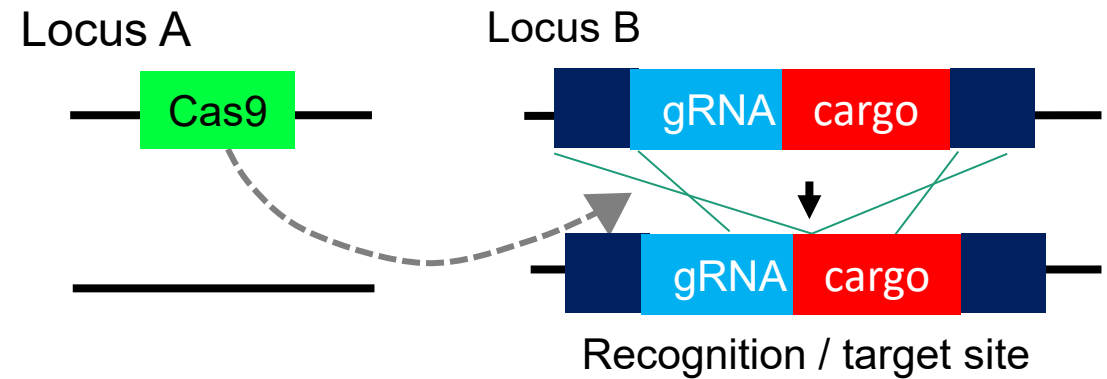


homing Gene drive: types

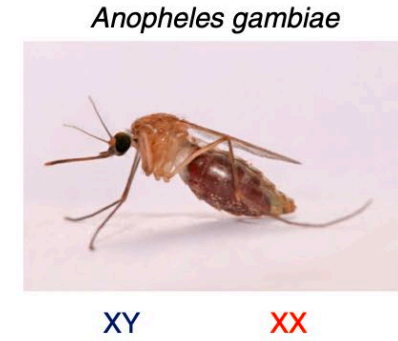
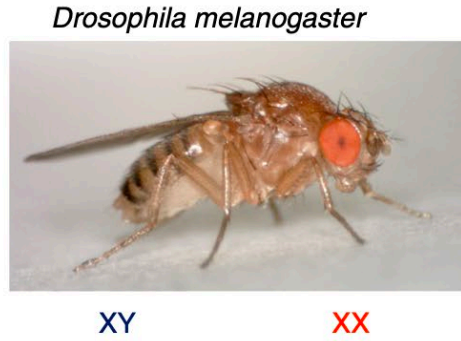
A. Gene drive (All in one)



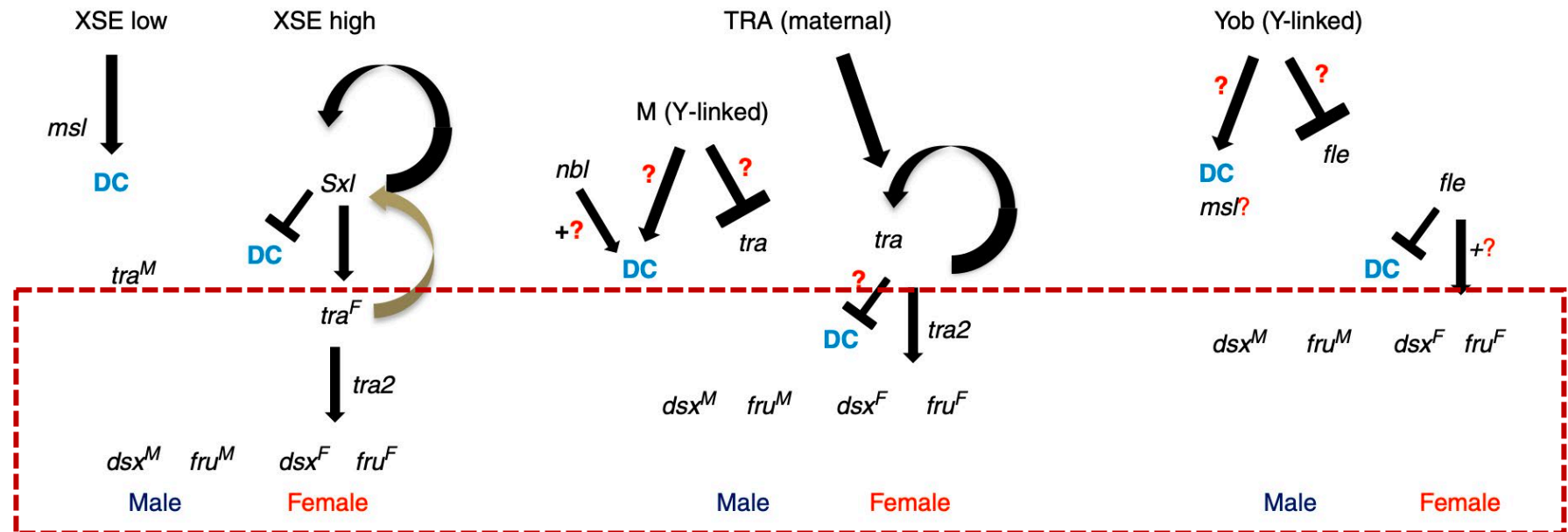
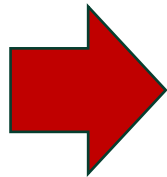
B. Gene drive (Split drive)



Sex determination genes as target for the suppression drives



The bottom of the regulatory pathway is conserved across all species.



Homing Gene drives demonstrated

D. melanogaster



<u>Target</u>	<u>Drive</u>
yellow	97%
yellow-g	86.4 [♀] -90.4% [♂]
RpL35A	91%
rab5	76.1 [♂] -97.1% [♀]
prosalpha2	99.3 [♀] -99.9% [♂]
tra	78%
dsx	70% [♂] -85% [♂]

....many more!

Diamondback Moth



<u>Target</u>	<u>Drive</u>
yellow	~50%
kmo	~50%

Medfly



<u>Target</u>	<u>Drive</u>
white	*85.6%
tra	83.1%

Spotted wing drosophila



<u>Target</u>	<u>Drive</u>
dsx	95-99%

Anopheles gambiae



<u>Target</u>	<u>Drive</u>
dsx	95.9-99.4%

Anopheles stephensi



<u>Target</u>	<u>Drive</u>
kh	99.7%

Culex quinquefasciatus



<u>Target</u>	<u>Drive</u>
white	56%
kmo	57.5%

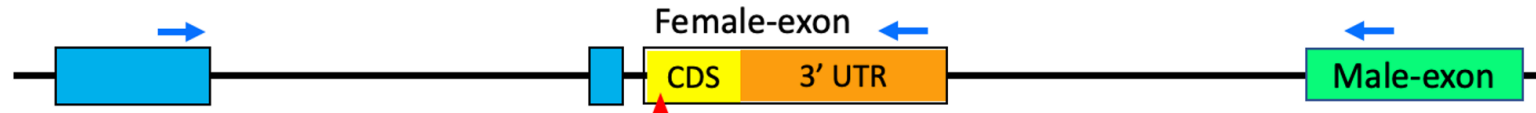
Aedes aegypti



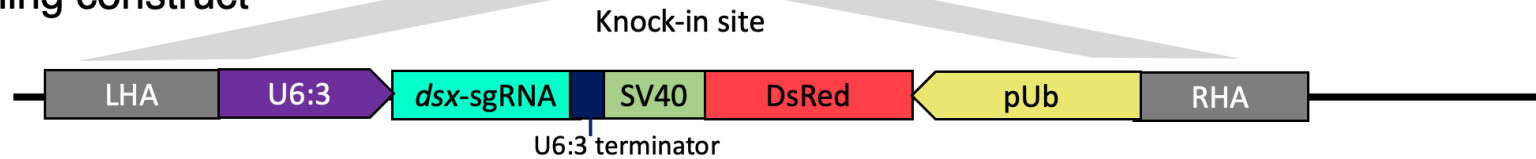
<u>Target</u>	<u>Drive</u>
white	*94%

SWD homing strains development

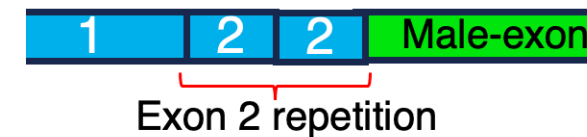
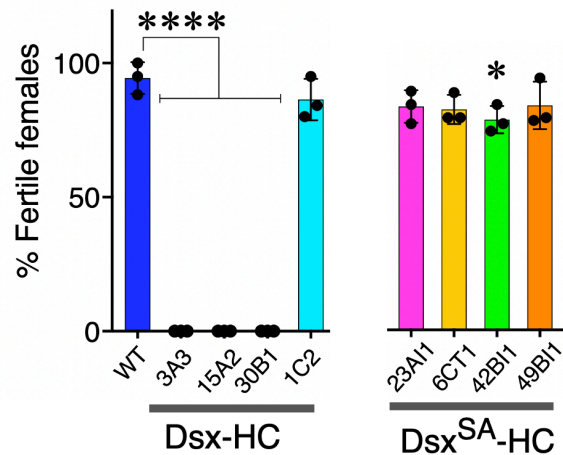
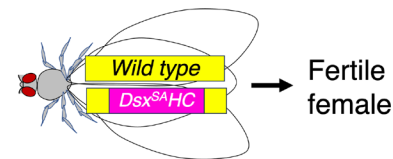
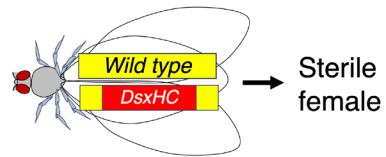
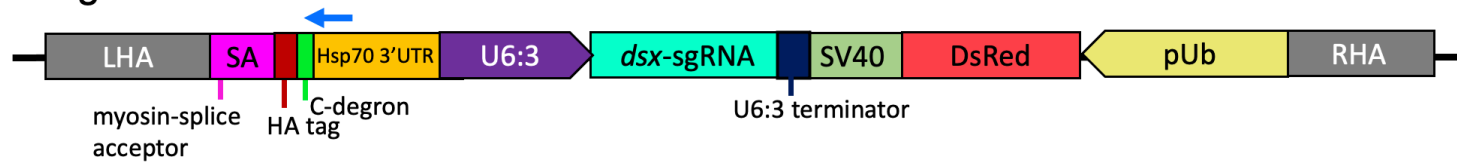
A *dsx* target site



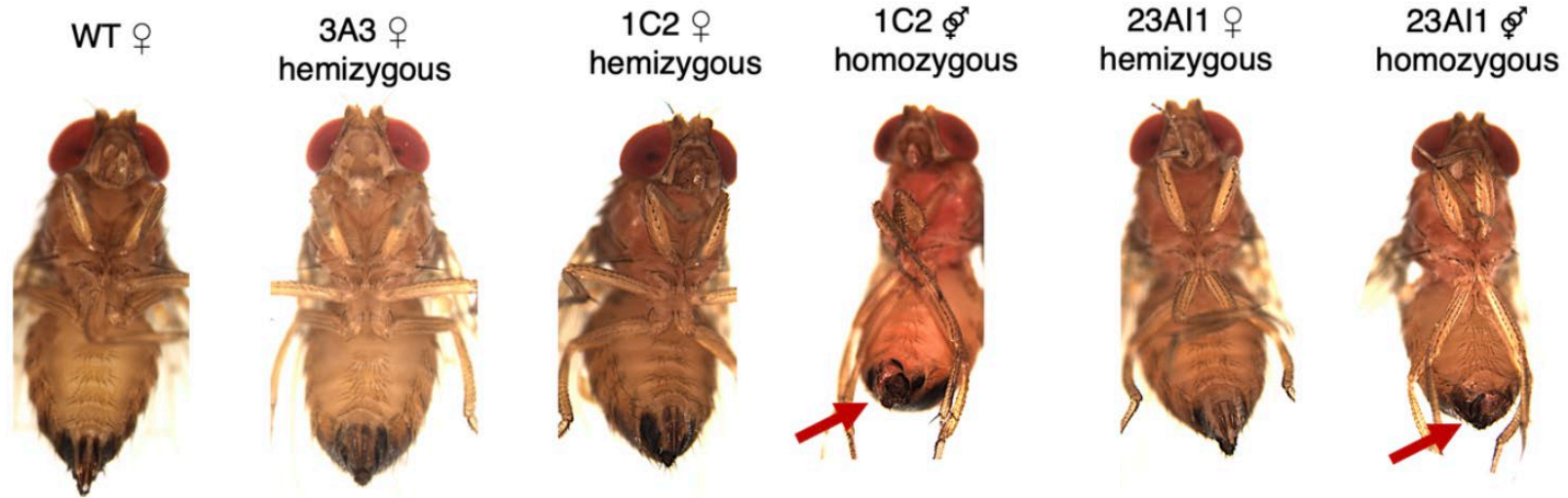
B *Dsx*-homing construct



C *Dsx*^{SA}-homing construct



SWD Homing strains morphology

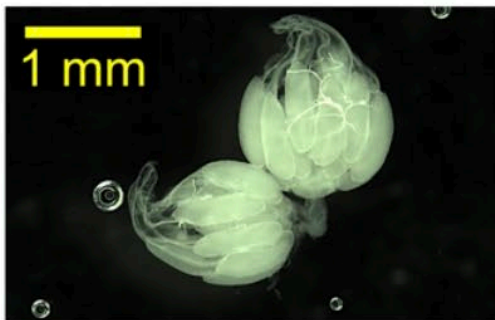


Wild type

3A3

15A2

30B1



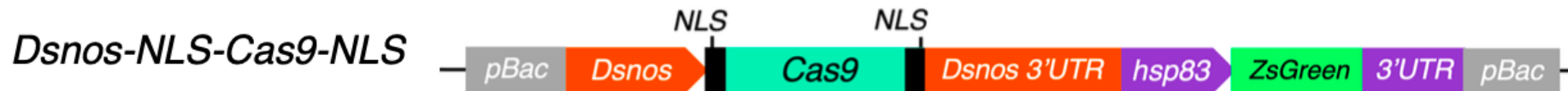
Ovaries

Dominant sterile hemizygous females

SWD Cas9 strains established

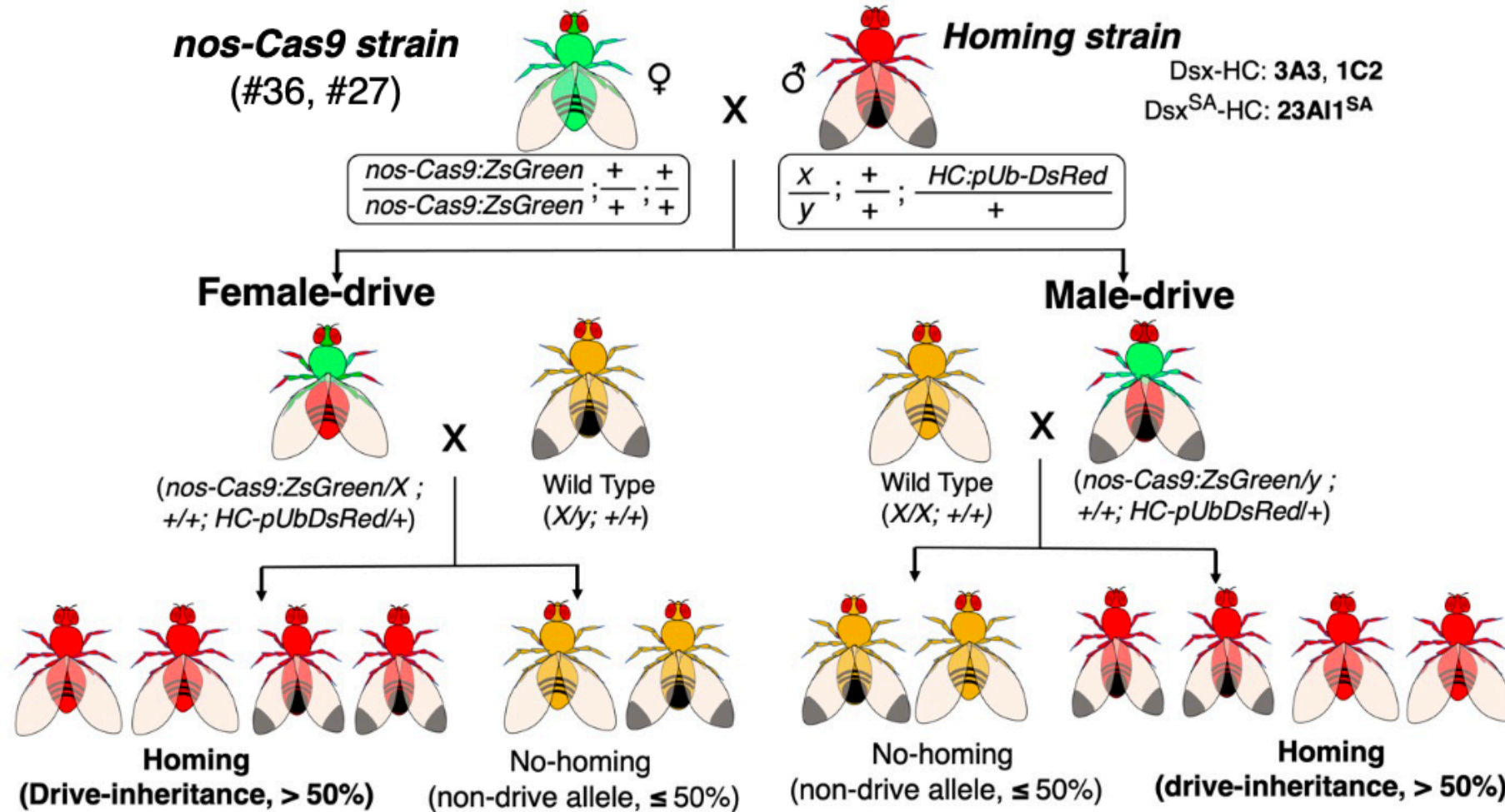


- X chromosome linked
- Cas9 expression under *D. melanogaster nanos* promoter and 3'UTR
- Cas9 has single NLS
- Low editing efficiency relatively



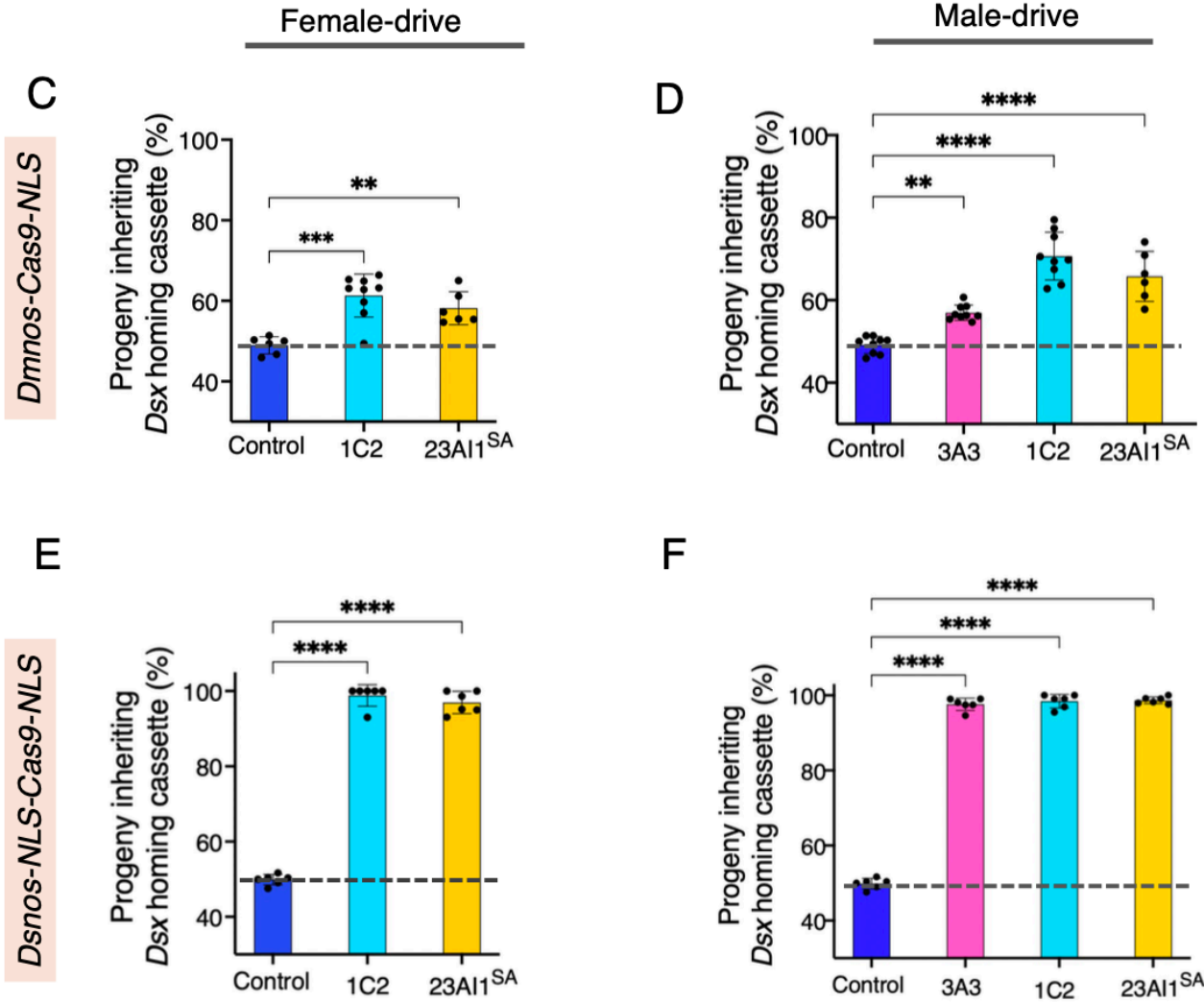
- X chromosome linked
- Cas9 expression under *D. suzukii nanos* promoter and 3' UTR
- Cas9 has two NLS
- High editing efficiency relatively

Genetic crosses to score the homing



dsx homing gene drive inheritance

- Improved Drive: 94-99% inherit *dsx* with DsRed and sgRNA gene!

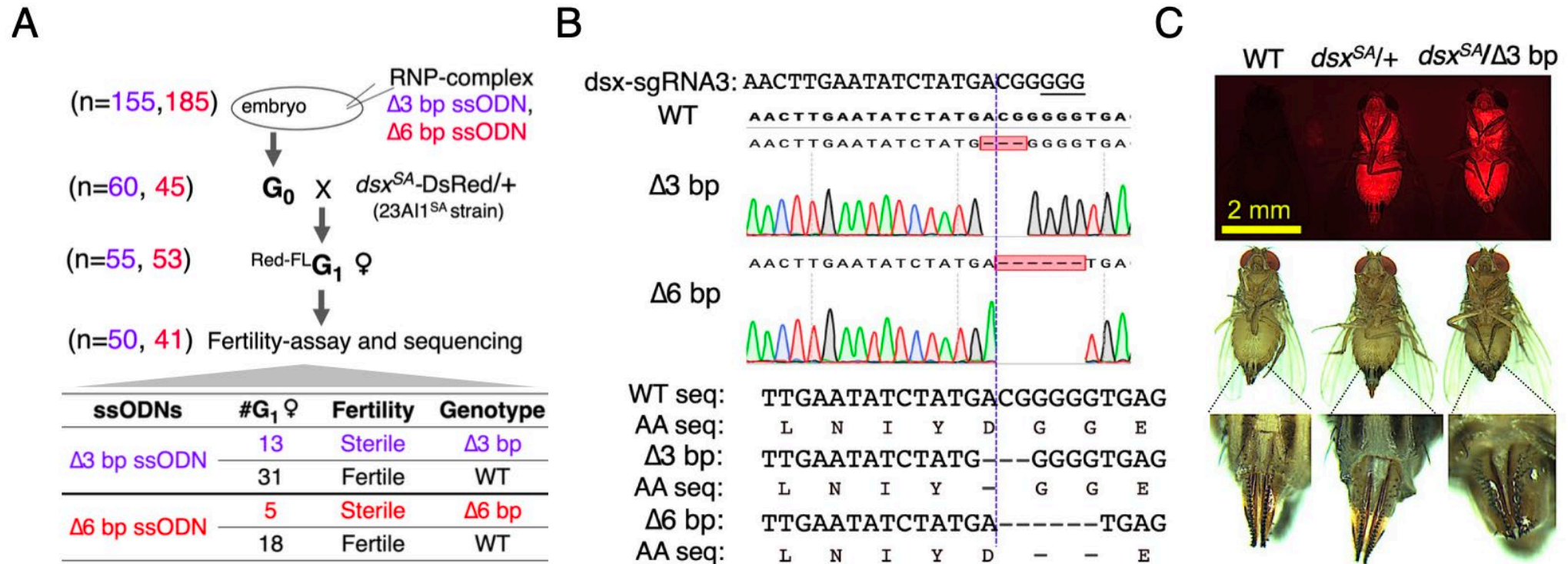


Potential Resistance Alleles Development

- NHEJ repair pathways can generate a resistance allele
- Possible, in-frame deletions or insertion would resistant to Cas9/sgRNA3 cleavage but produce functional protein!

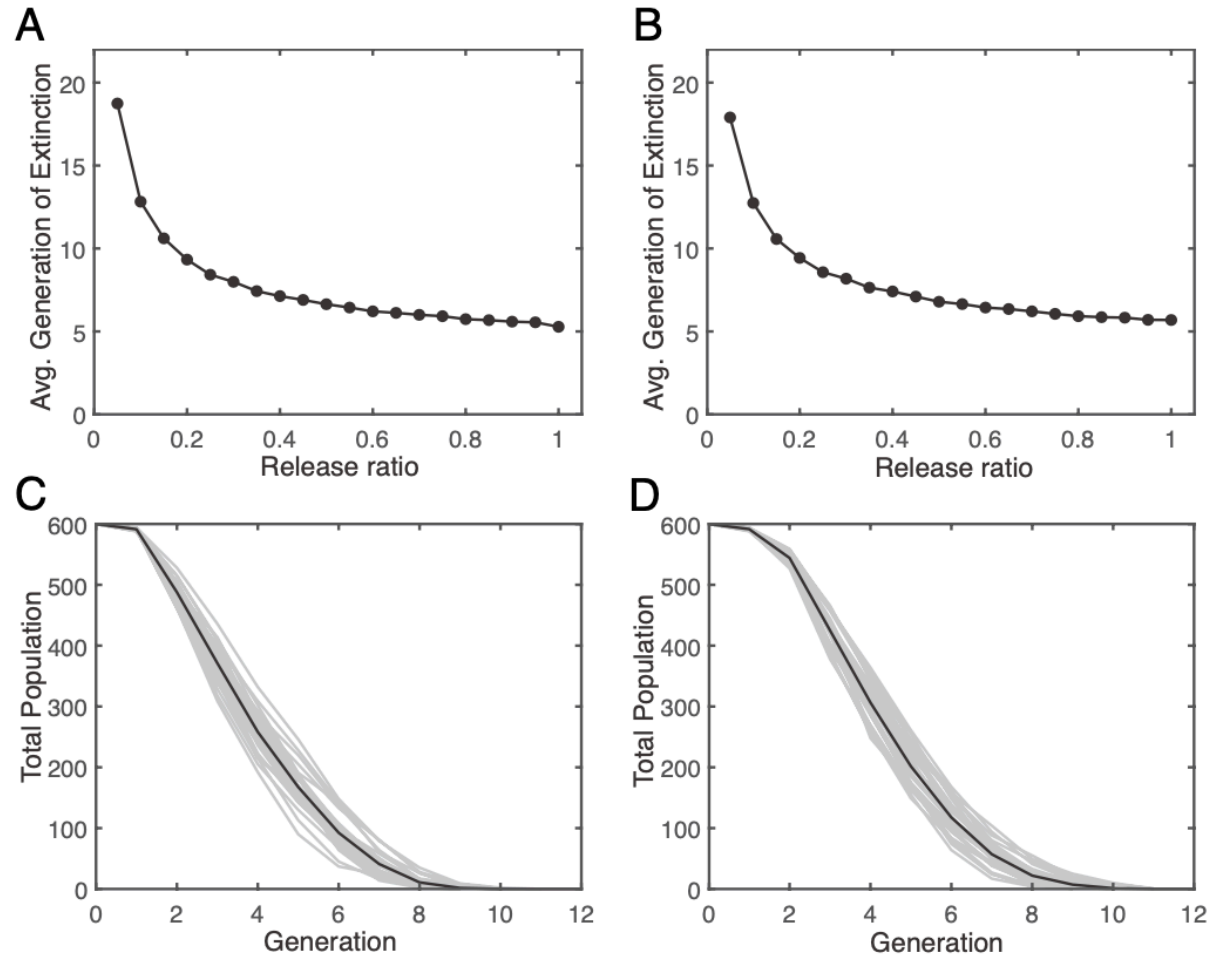
sgRNA3	AACTTGAATATCTATGACGGGGG	Indel	n/total
WT	AACTTGAATATCTATGACGGGGG		
3A3 male drive	AACTTGAATATCTATGACGGGGG	WT	7/13
	AACTTGAATATCTATG-----GGG	-4 bp	2/13
	AACTTGA-----GCTTCGCAA	-19 bp	1/13
	AACTTGAATA-----TGAGCTTCGCAA	-13 bp	1/13
	AACTTGAATATCTATGANNNCGGGGG	+3 bp	1/13
1C2 male drive	AACTTGAATATCTA---CGGGGG	-3 bp	1/13
	AACTTGAATATCTATGACGGGGG	WT	1/9
	AACTTGAATATCTATG-----GGG	-4 bp	1/9
	AACTTGA-----GCTTCGCAA	-19 bp	4/9
1C2 female drive	AACTTGAATATCTATG-CGGGGG	-1 bp	3/9
	AACTTGAATATCTATGA-GGGGG	-1 bp	2/3
23A11^{SA} male drive	AACTTGAATATCTATGACGGGGG	WT	7/8
	AACTTGAATATCTAT-----GGGGG	-3 bp	1/8
23A11^{SA} female drive	AACTTGAATATCTA---CGGGGG	-3 bp	3/12
	AACTTGAATATC-----CGGGGG	-5 bp	3/12
	AACTTGAATATCTATGA-----GGG	-3 bp	1/12
	AACTTGAATATCTATGA-----TGAGCTTCGCAA	-6 bp	1/12
	AACTTGAATATCTATGA-----GCTTCGCAA	-9 bp	1/12
	AACTTGAATATCTATGANNNNNNNCGGGGG	+7 bp	1/12
	CGTCAGCA-----GGGGG	-19 bp	1/12
AACTTGAATAT-----GGGGG	-7 bp	1/12	
GCAT-----GGTGAGCTTCG	-21 bp	1/12	

In-frame deletion mutations are resistant to cleavage but are not functional



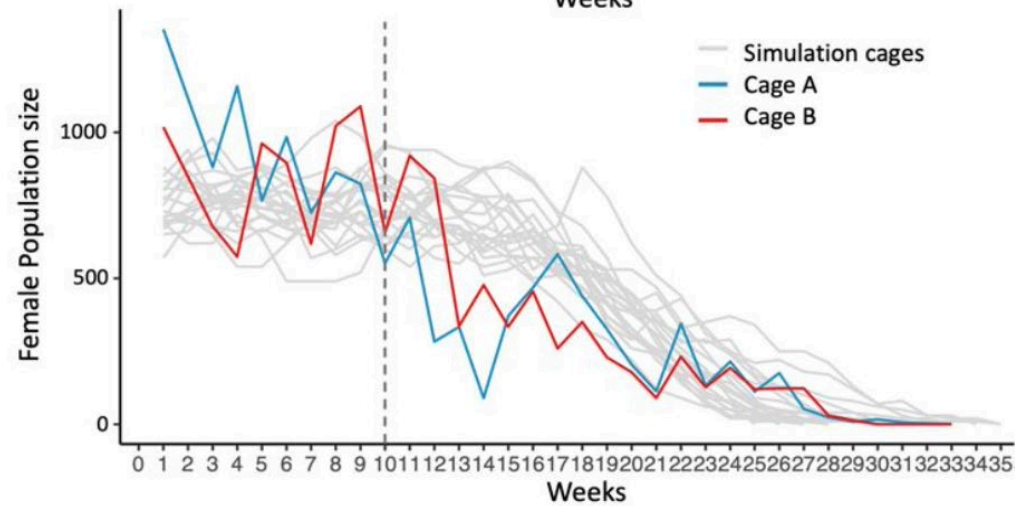
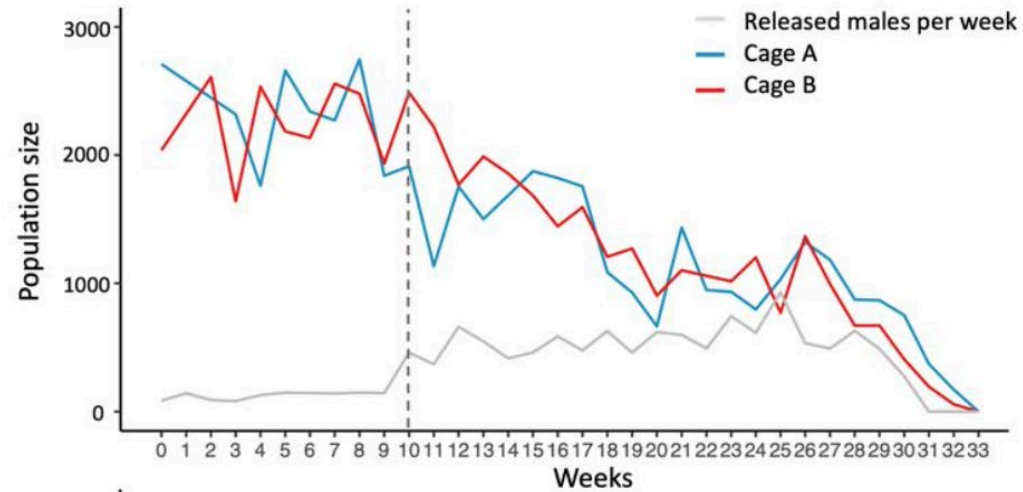
Mathematical Modelling of split drives

- Discrete generation lab cage population
- Repeated Releases, 1: 4 ratio (transgene: wild type)
- Suppression within 10 generations



Population dynamics of Dominant sterile Drive in caged populations of *D. melanogaster*

- Cage Population size: 2500
- Weekly releases of drive
- Population collapsed in 29 weeks



Challenges

- Precision at genetic and molecular level
 - Efficiency and fitness cost**
- Population dynamics of the target pest
 - Pest population size and its distribution**
- Mass rearing and field releases

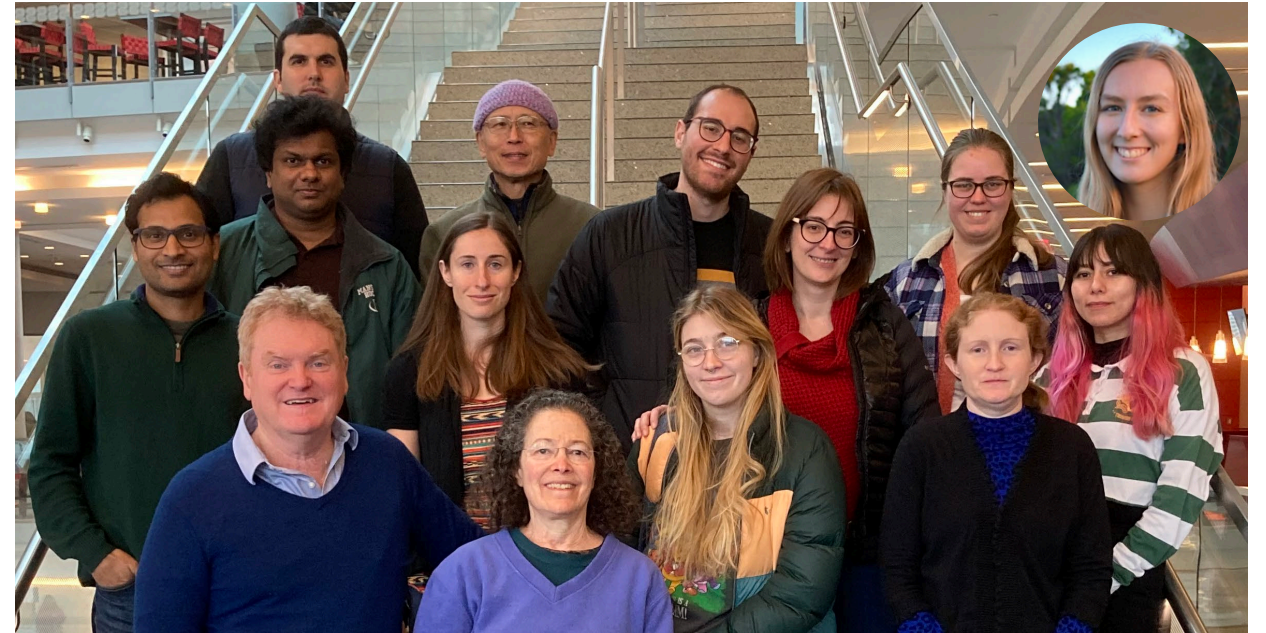
Acknowledgements

Scott lab members

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Ariel Terrand
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