



Not All Gene Drives Are Created Equal

Fred Gould Genetic Engineering & Society Center NC State University

Drive Category	Threshold	Can spread high fitness cost	Temporally limited	Spatially limited	Resistance to drive
Standard Homing Drive	No (except special cases)	Yes	No	No	Yes In most cases
Cleave/Replace TARE/Medea	Low- if no fitness cost	No	No	Depends on fitness cost	Possible
Wolbachia	Low- if no fitness cost	No	No	Depends on fitness cost	Yes
Killer-Rescue Split-Homing Daisy Chain	No	No	Yes	No	Possible
Underdominance	At least 27%	No	No	Yes	Possible
Tethered Homing (with Underdominance)	At least 27%	Yes	No	Yes	Yes In most cases

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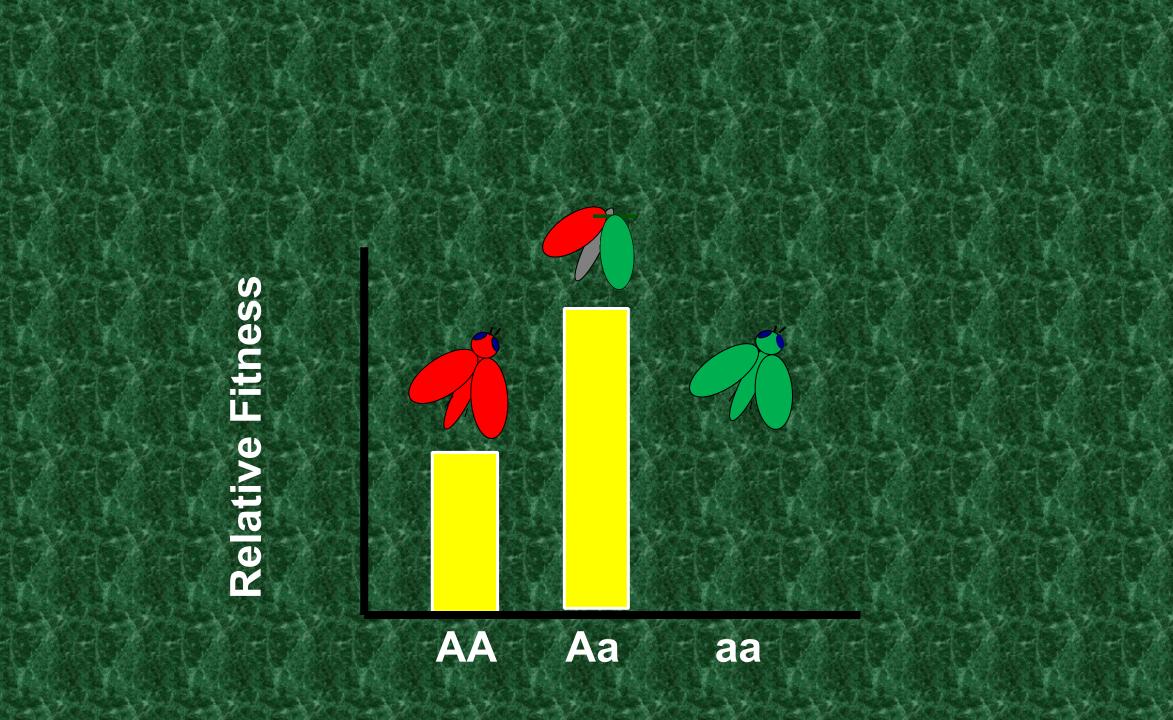
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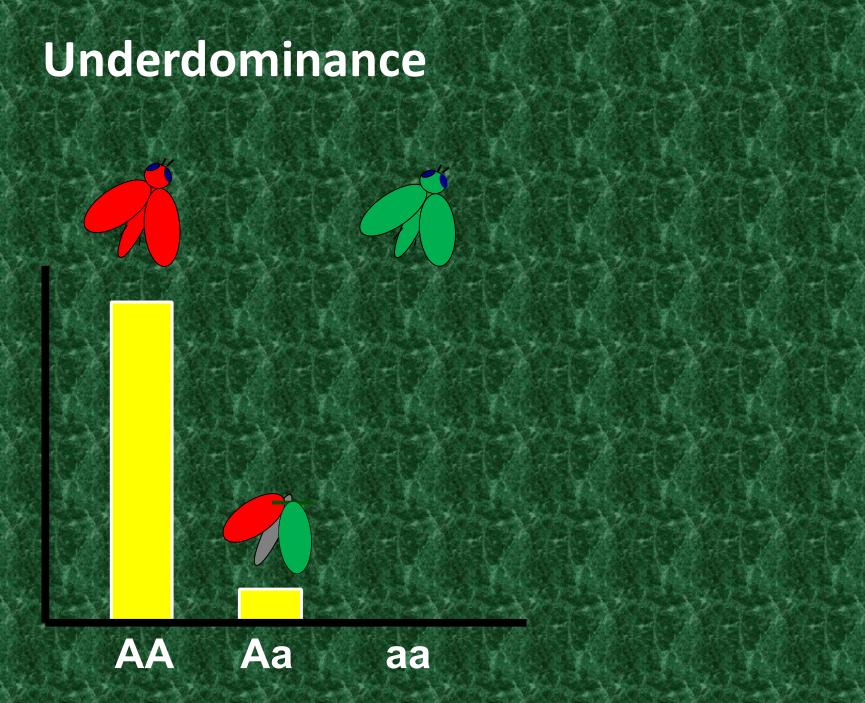
Underdominance for Gene Drive

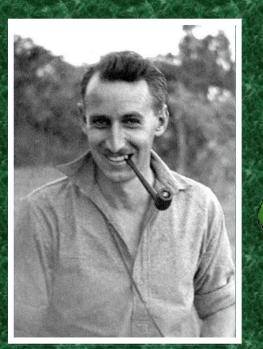
Spatially Limited Spread

Relative Fitness AA Aa aa









F.L. Vanderplank 1940's Hybrids of Tsetse fly species

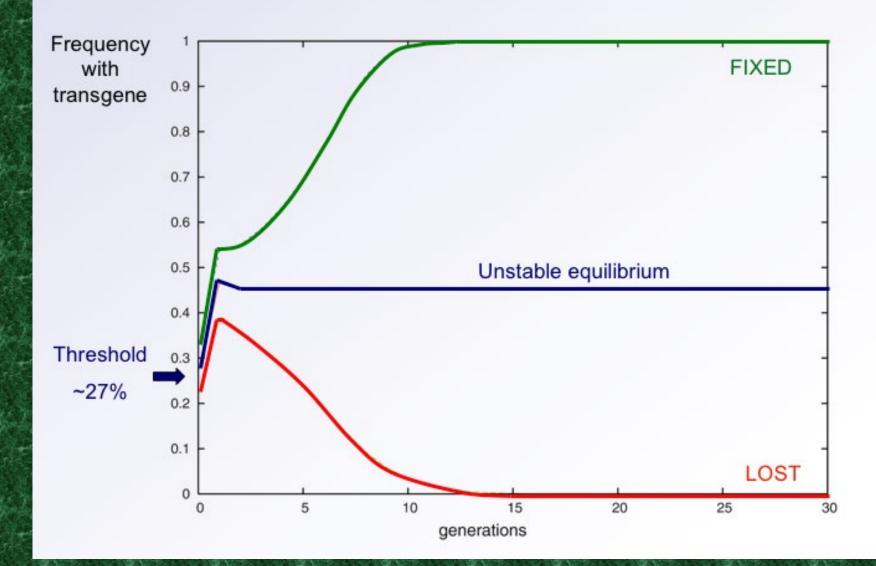




Chris Curtis UK 1960's

A. S. Serebrovski Soviet Union - Lysenko era 1940's Translocations

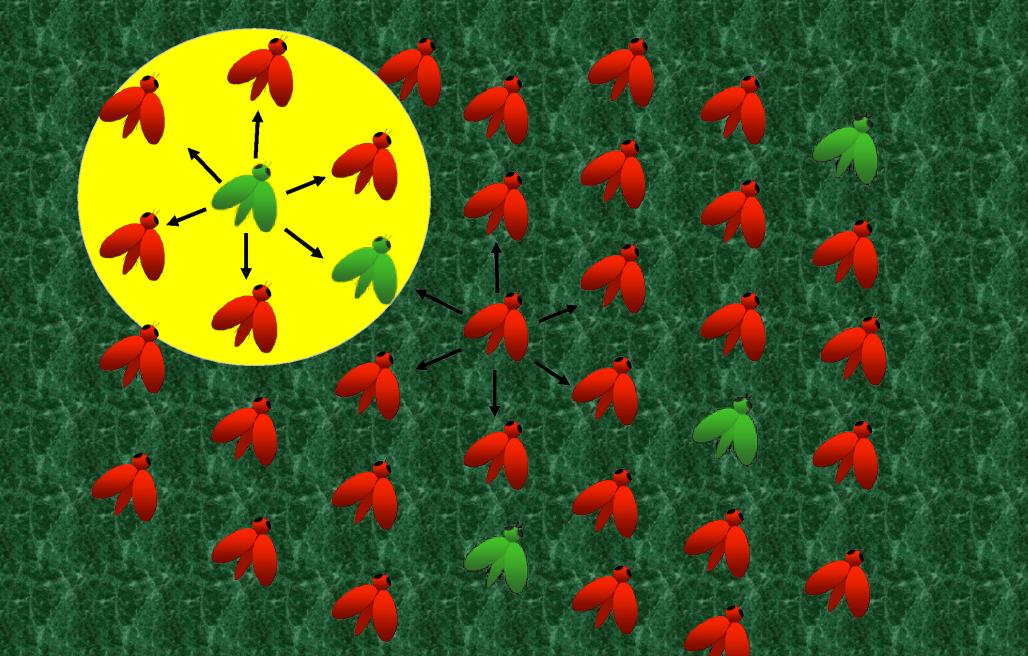
Engineered Underdominance – No cost

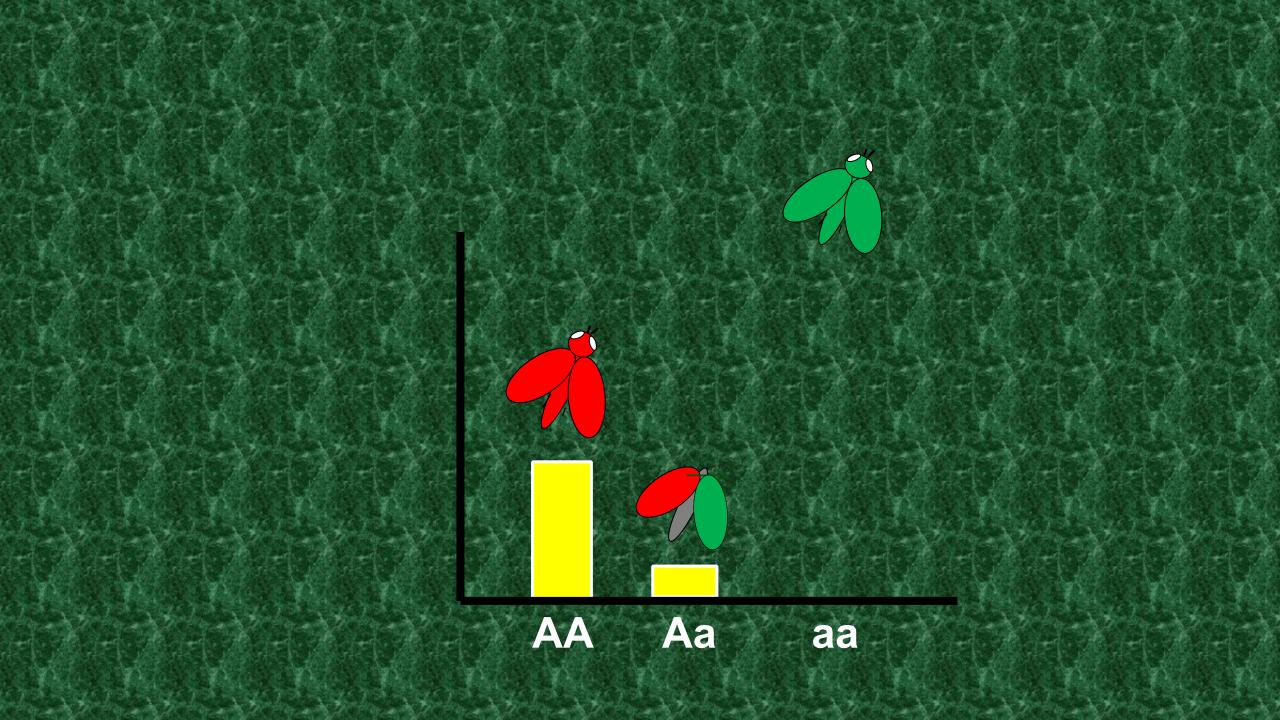


Large numbers of transgenic genotypes released

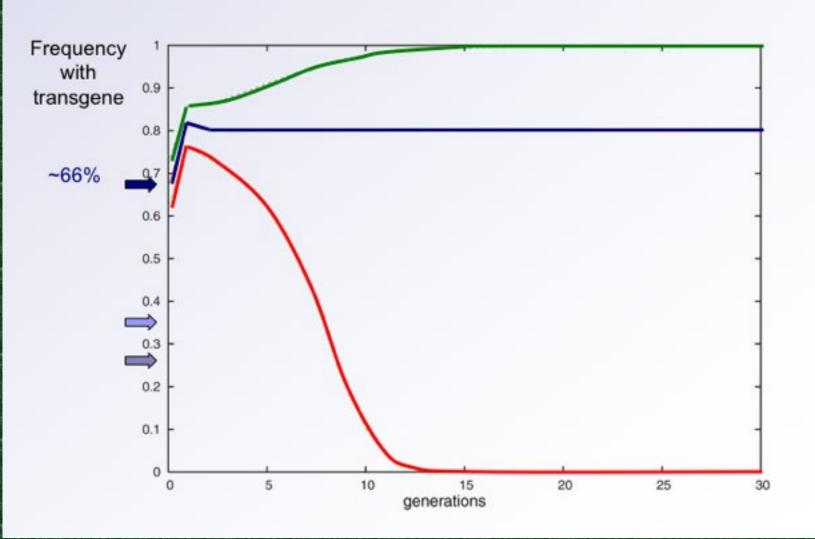
A transgenic Individual is likely to mate with another transgenic individual

A wildtype individual is likely to mate with a transgenic individual





Engineered Underdominance – 20% cost per construct

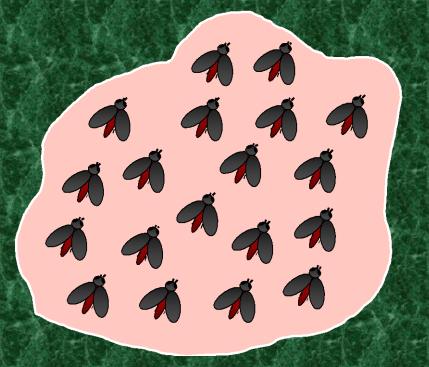


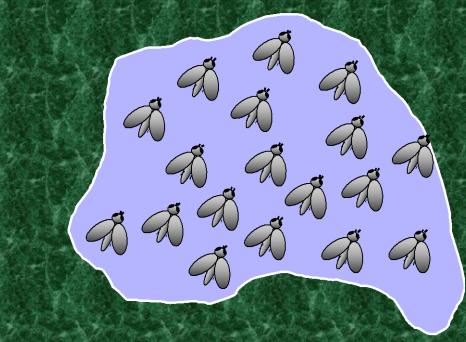
Using Underdominance as gene drive

Spread only when present above a certain proportion in the population

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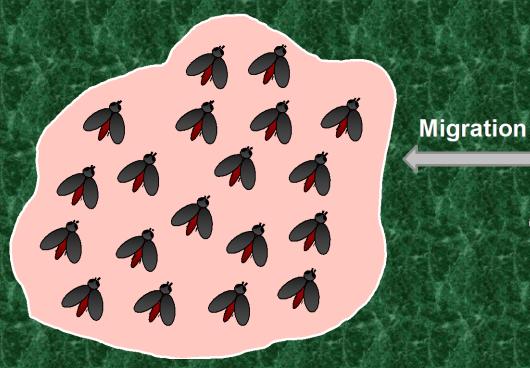
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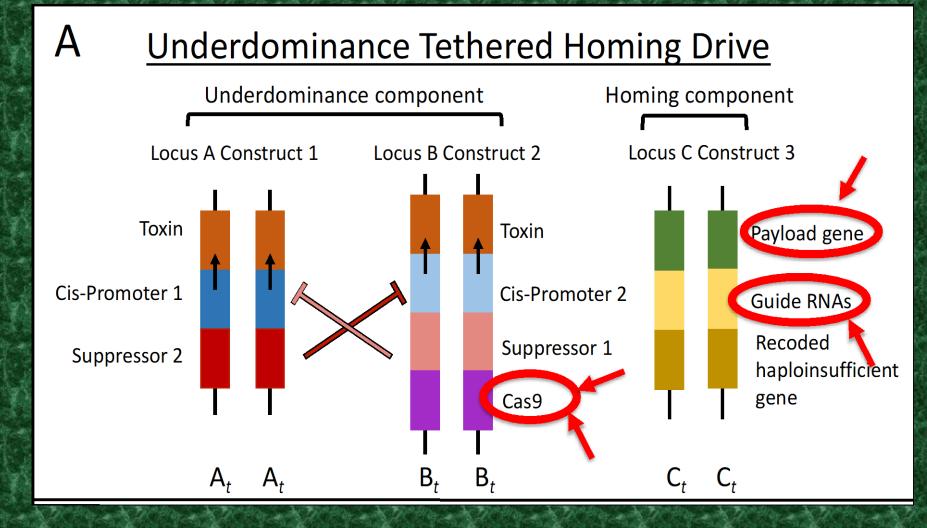




Using Underdominance as gene drive

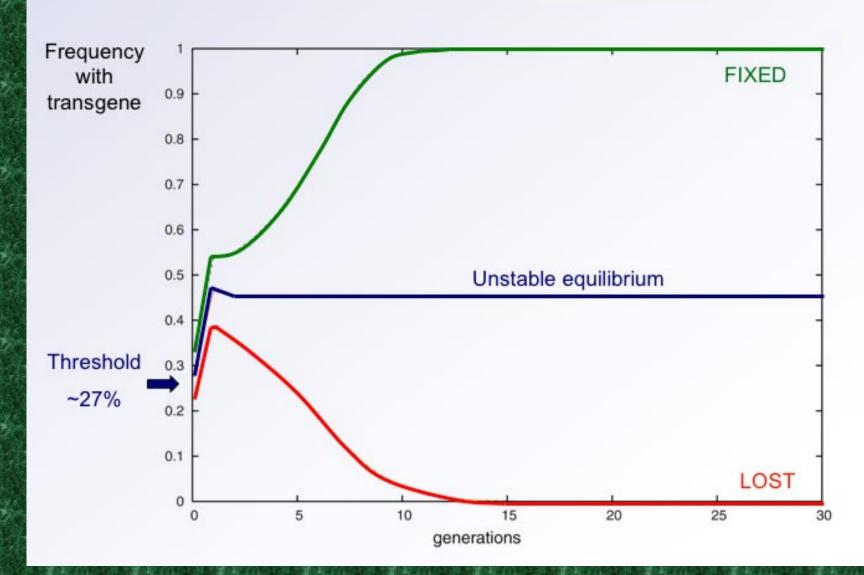
Depending on how high the threshold is, they can remain localized







Engineered Underdominance



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PHASE 1: STERILE MALE

PHASE 2: SELF-LIMITING

PHASE 3: SELF-SUSTAINING

Supression

Replacement

Resistance to Drive

Breakdown in Pathogen Control