

Technology characterisation of Gene Drives and potential applications - an introduction -

Bernd Giese¹, Johannes Frieß², Arnim von Gleich²

¹Institut für Sicherheits- und Risikowissenschaften (ISR), Universität für Bodenkultur, Wien

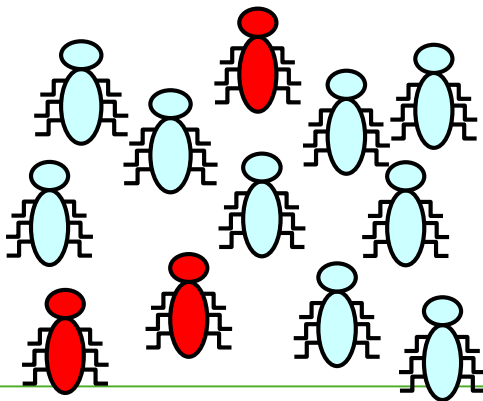
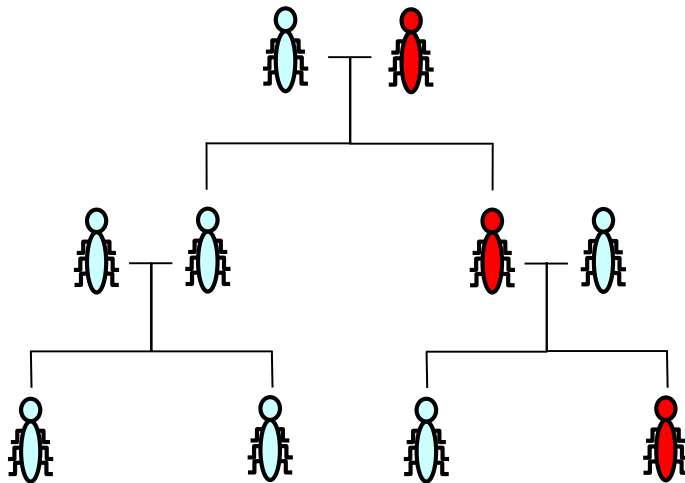
²Fachgebiet Technikgestaltung und Technologieentwicklung, Universität Bremen

Remote control of populations via a „genomic interface“?

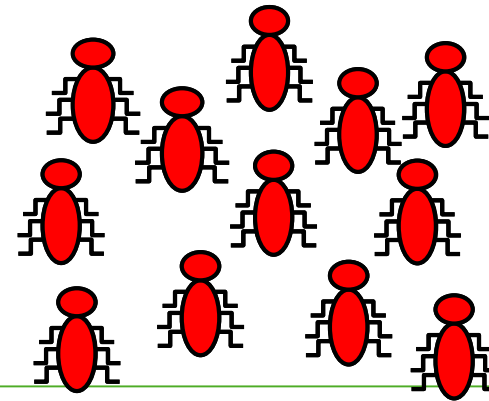
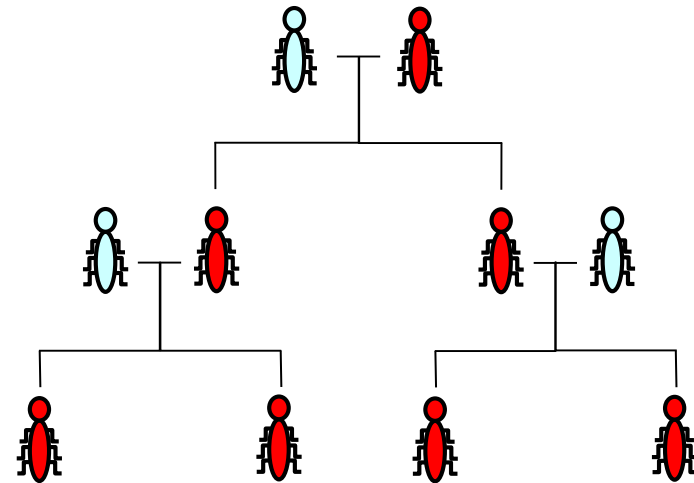


„Gene Drives“

(Mendelian) inheritance of a **Gene**

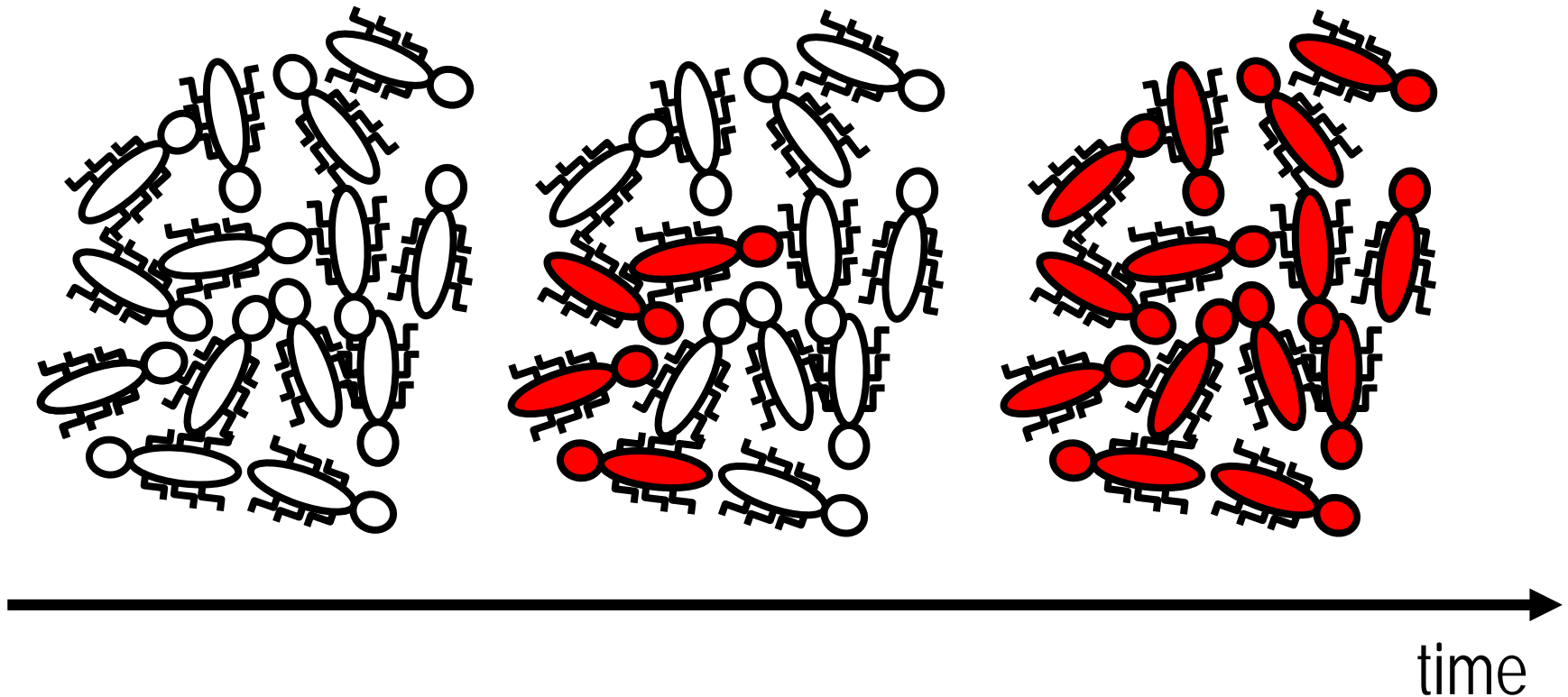


Gene Drive biased inheritance



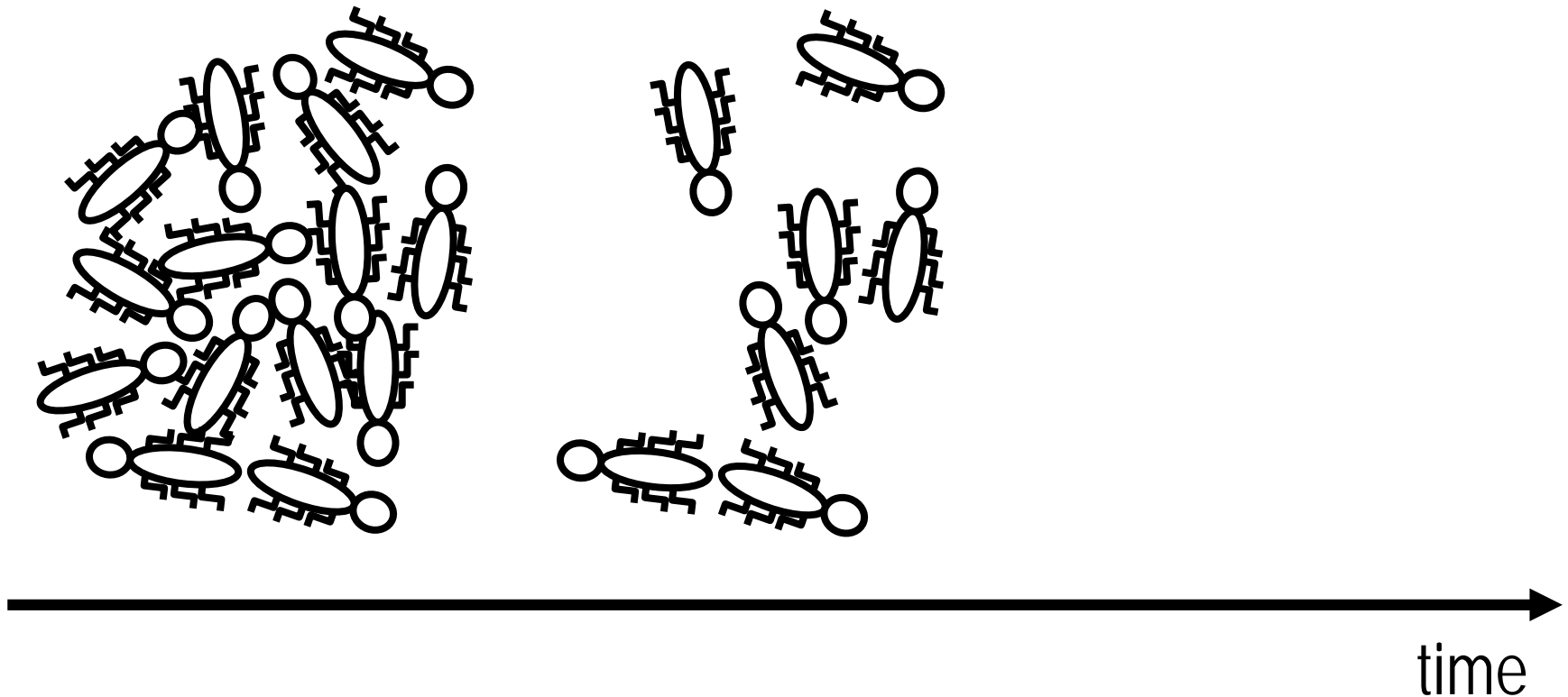
Gene Drives as Conversion Drives

→ Control by spread (of new traits)



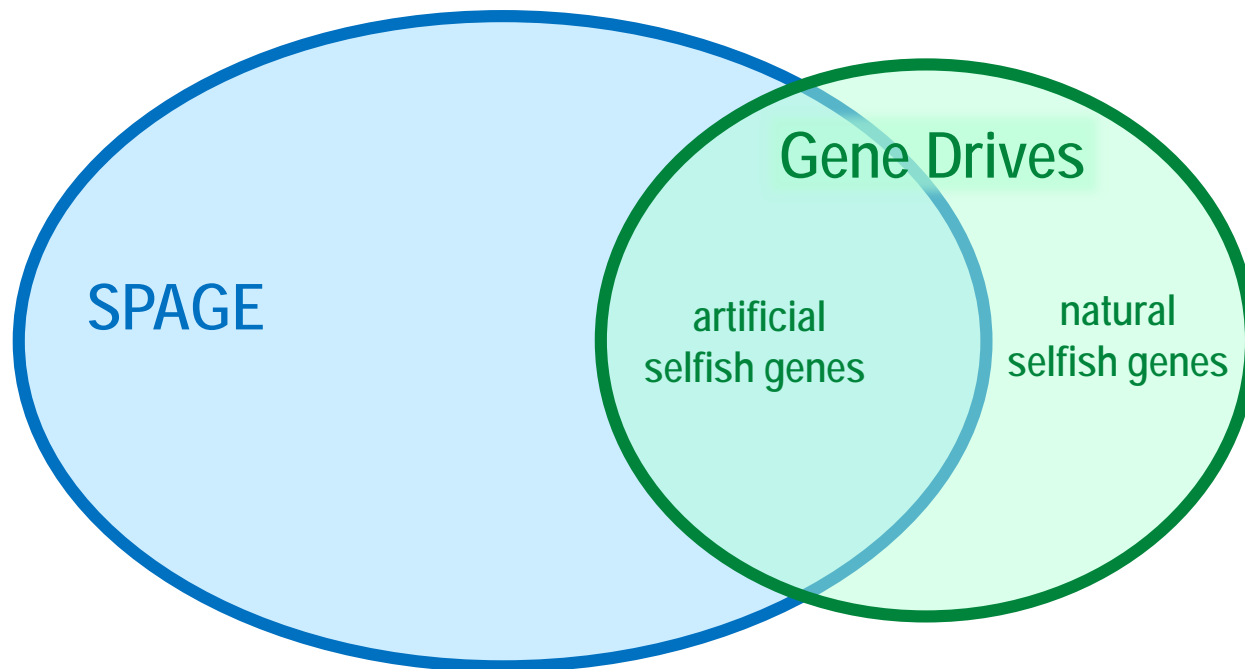
Gene Drives as Suppression Drives

→ Suppression or even eradication of populations



SPAGE-Technologies and Gene Drives

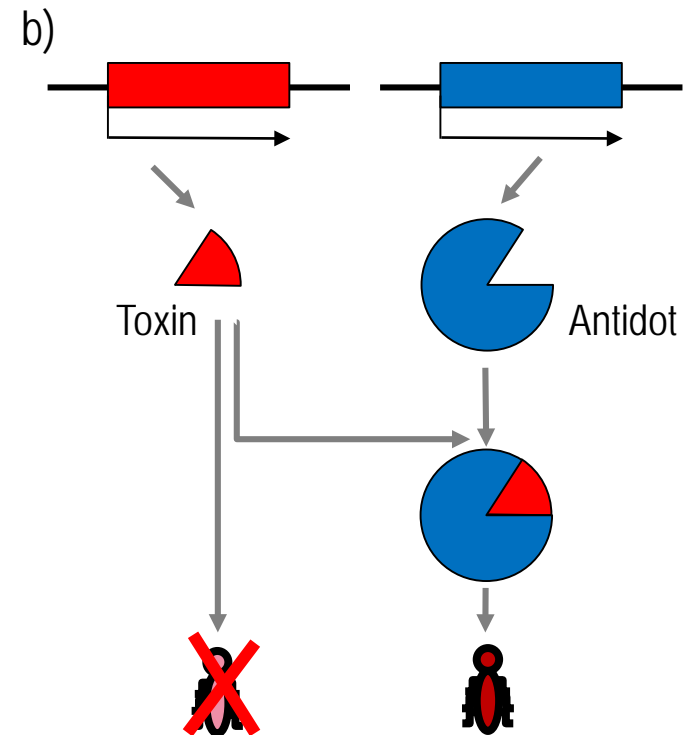
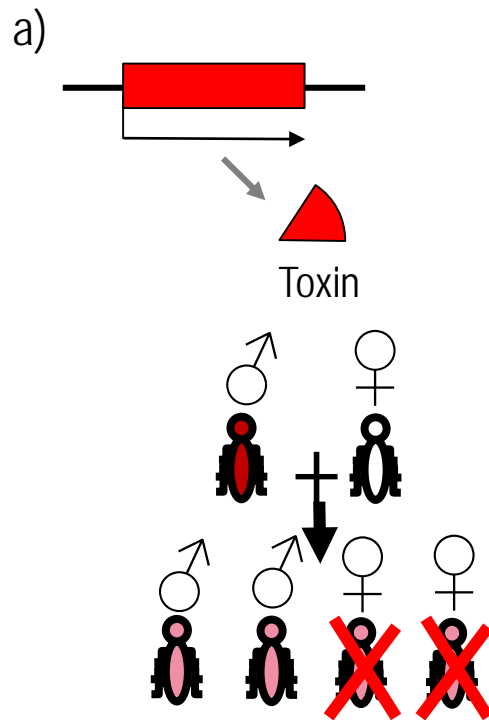
SPAGE= self-propagating artificial genetic element



Types of SPAGE/Gene Drives

Passive:

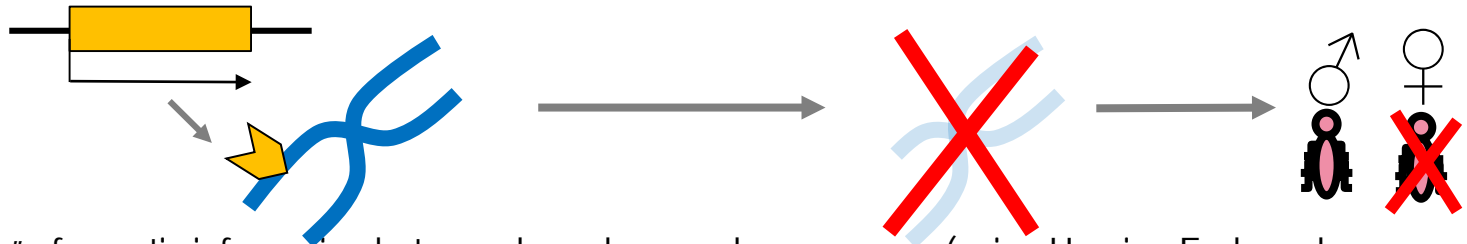
- Only offspring not targeted by a toxin or carrying a particular combination of genetic elements (toxin + antidot) survives



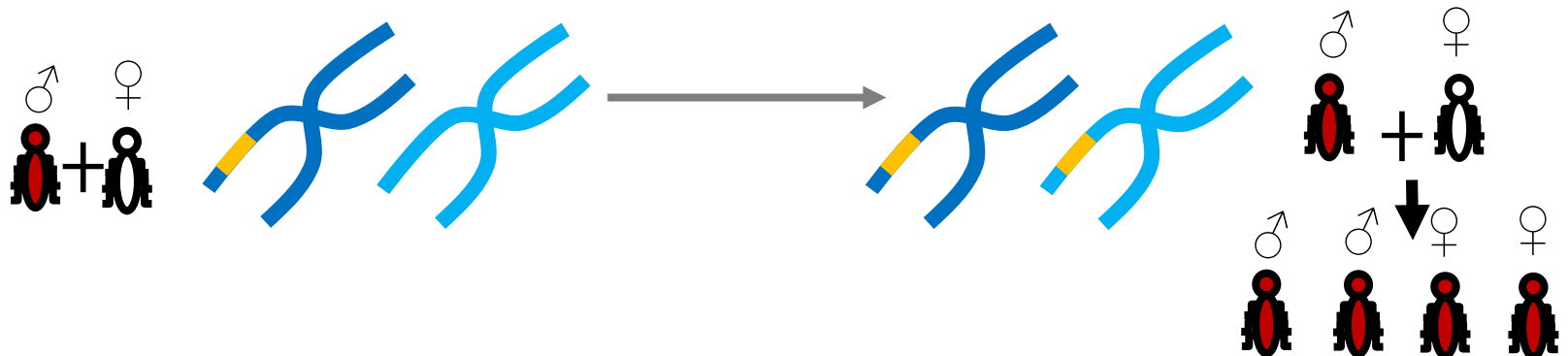
Types of SPAGE/Gene Drives

Active (by genome modifications):

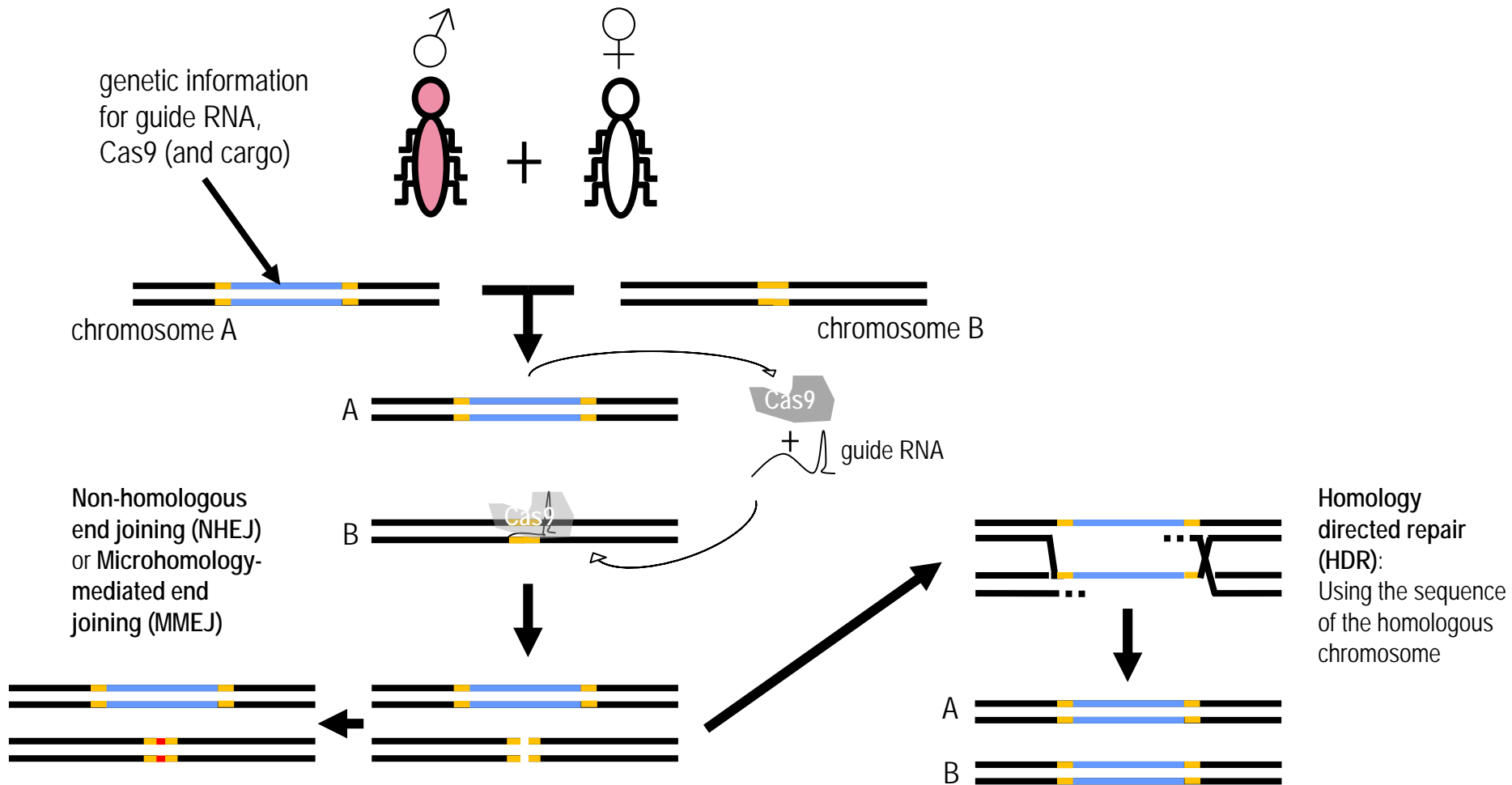
- Bias of sex ratio by (enzymatic) mechanisms influencing meiotic segregation (inhibition of sperm development) / „shredding“ of the X-chromosome



- “Copying” of genetic information between homologous chromosomes (using Homing Endonuclease Genes (HEG))



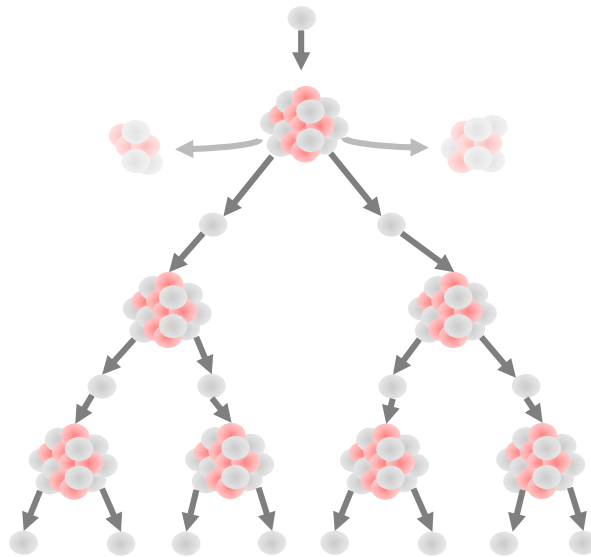
Mechanism of CRISPR/Cas9 – based Gene Drives



High power by Homing Endonuclease based drives

- for CRISPR/Cas9-based gene drives copying efficiencies of 91–99% can be achieved

(Gantz and Bier 2015; Gantz et al. 2015; Hammond et al. 2015 cited in Lin and Potter 2016)



A “mutagenic chain reaction”?

Cp. Gantz & Bier, Science, 24 APRIL 2015 •
VOL 348 ISSUE 6233

Potential unintended effects and/or reduced power by:

- Non-homologous end joining (NHEJ) and Microhomology-mediated end joining (MMEJ) instead of Homology-directed repair (HDR) reduces the conversion rate and may cause resistance due to mutations, deletions etc.
- Resistance due to genomic variations (sequence polymorphisms), homing resistant alleles
- Maternal effects due to Cas9 deposit may cause resistance
- On-target misinsertions
- Off target-effects (unspecific binding of gRNA causes unintended insertions at different locations)
- Intragenomic interactions

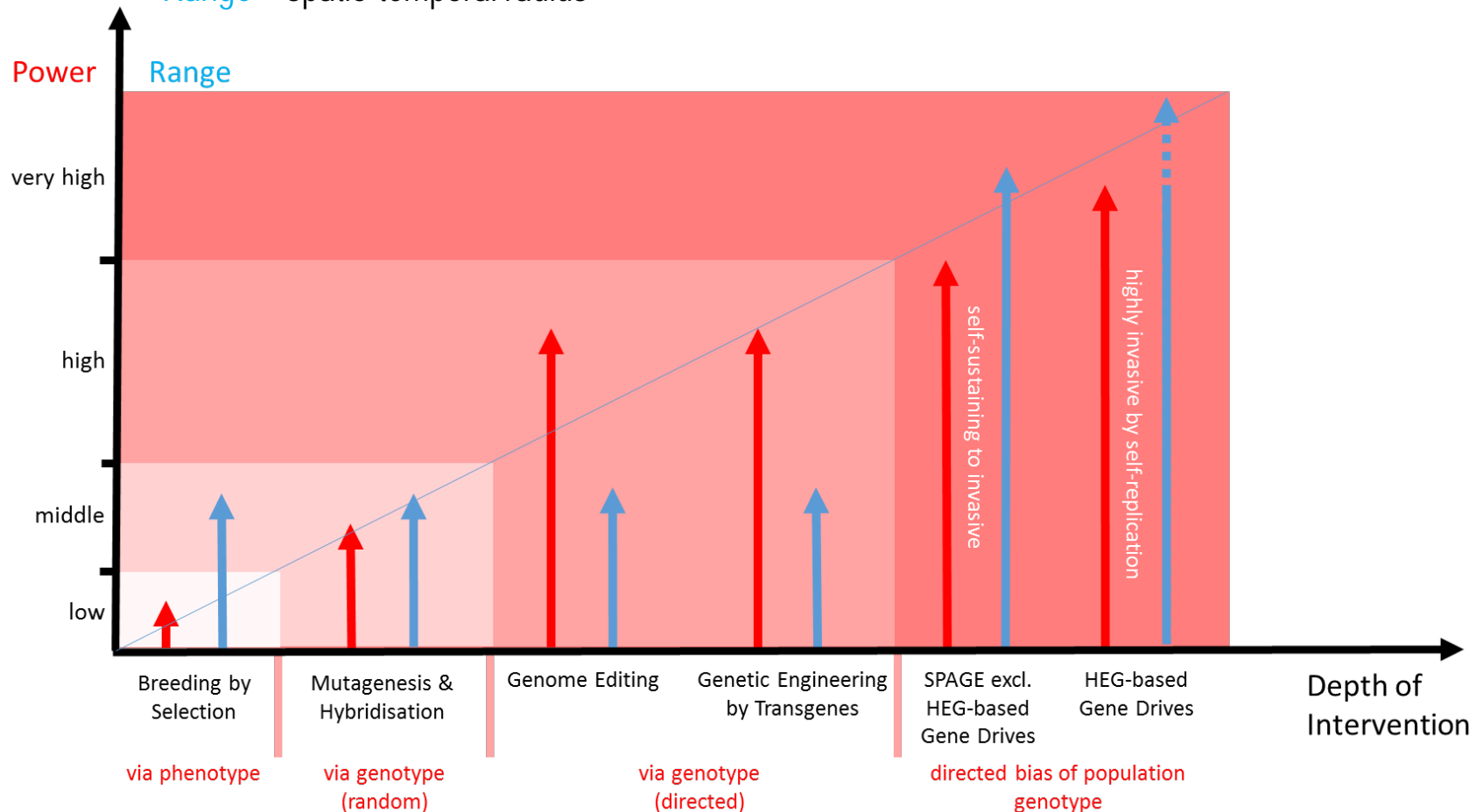
No claim to be complete ...

- Gene Drives in general represent a **high depth of intervention** due to
 - the directed modification of the genome
 - the ability to bias the genotype of populations
 - Accordingly, it holds the potential for a
 - high **technological power** by its toxic / enzymatic mode of action and the option of cargo genes
(→ **hazard potential**)
 - high **range** due to its influence on the pattern of heredity or even the capability of self-replication
(→ **exposure potential/potential of contamination**)

Depth of Intervention, Power and Range of generations of breeding technologies

Power = directedness and character of the implemented trait

Range = spatio-temporal radius



Gene Drives increase the range of genetic engineering

- intended to spread in wild populations
- transform wild populations or even
- eliminate wild populations
 - technological tipping point
 - shift of paradigms in the handling of GMOs

GMO = genetically modified organism

Applications of Gene Drives

- Medical:
 - Immunization or elimination of disease vectors

- Agricultural:
 - Elimination of plant and animal pests
 - Control of pesticide-resistant weeds

- Ecosystem 'engineering':
 - Invasive species (e.g. in New Zealand)
 - Immunization of endangered species

Malaria and Dengue



By James D. Gathany - The Public Health Image Library



By James Gathany, CDC - This media comes from the Centers for Disease Control and Prevention's Public Health Image Library (PHIL)

Malaria

- pathogen: 5 *Plasmodium* species
- vector: mainly 3 *Anopheles* mosquito species
- endemic in 91 countries
- 216 million new cases in 2016
- 445,000 deaths
- insecticide and drug resistances evolve

Dengue

- pathogen: Dengue Virus (4 serotypes)
- vector: *Aedes* mosquito genus
- endemic in ~100 countries
- ~93 million clinical cases/year
- ~22,000 deaths
- mostly preventive vector control

Invasive Species

New Zealand:

- initiative Predator Free 2050
- especially rats, stoats and possums
- gene drive application considered
- problem of confinement (global diffusion)



Spotted-Wing Fruit Fly in California:

- deposits eggs in cherries with an ovipositor
- Medea Gene Drive is considered to eradicate the fly / alter its ovipositor



By Martin Cooper from Ipswich, UK - Spotted-wing
Drosophila (*Drosophila suzukii*) male, CC BY 2.0

„Weed“- Control

- e.g., Palmer Amaranth
- infested agricultural fields (cotton etc.) in the Southern United States
- related species are cultivated crops in Mexico and South America (in China and India as well)



NAS-Report (2016): *„What mechanisms are in place for dialogue with the Mexican national government? How will any concerns raised by the Mexican government be incorporated into US decision-making processes?“*

Potential ecological effects

- **Evolutionary effects:**
 - gene flow
 - intraspecific (may influence fitness of GE- and wild population)
 - interspecific (interspecific mating may influence fitness of other species)
 - mutations of genetic information
 - evolution of resistance (→ blockade of gene drive spread)
- **Ecological interactions:**
 - How many ecosystem processes are dependent upon the target species?
 - role of the species
 - resource/consumer (predator/prey), symbiont/parasite, disease vector, competitor,
 - Loss of ecosystem services? e.g., as pollinators
 - Cascading effects?
 - Loss of diversity?
 - Niche filling by alternative species?

Potential ecological effects depend

... on the structure and vulnerability
of the affected (eco-)system

Important aspects to consider:

- Critical elements in affected system?
- Structural instabilities?
- Potential tipping points?
- Adaptive capacity?
- Potential for self-repair?

→ structural analysis of vulnerability

... on the character of the drive

Important aspects to consider:

- Exposure potential?
- Hazard potential?

→ effect-related analysis of
vulnerability

- gene drives represent a „tipping point“ of genetic engineering, because
 - power and range potentially exceed capabilities of previous stages in genetic engineering („active genetics“*)
 - gene drives are designed to
 - actively shape the genotype of wild populations and
 - engineer ecosystems by the conversion or suppression/elimination of wild populations
 - in particular actively replicating drives based on homing endonuclease genes (e.g., CRISPR/Cas9) are probably highly invasive
 - with potentially extreme power and range many open questions arise regarding effects, control and reversibility
- urgent demand for prospective analysis of impact, side effects, countermeasures and the feasibility of low-risk approaches**