



Allelic dropouts, null alleles or rare sex detection in clonal organisms: simulations and application to real data sets

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32ND MEETING OF INTERNATIONAL SCIENTIFIC COUNCIL FOR TRYPANOSOMIASIS RESEARCH AND CONTROL

Session 16

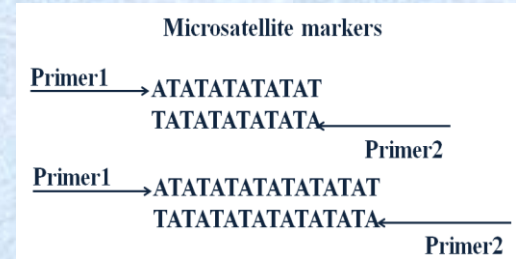
Difficulty of direct observation for Pathogens and their vectors



improvement of DNA amplification techniques during the last few decades



spatio-temporal variability of molecular markers

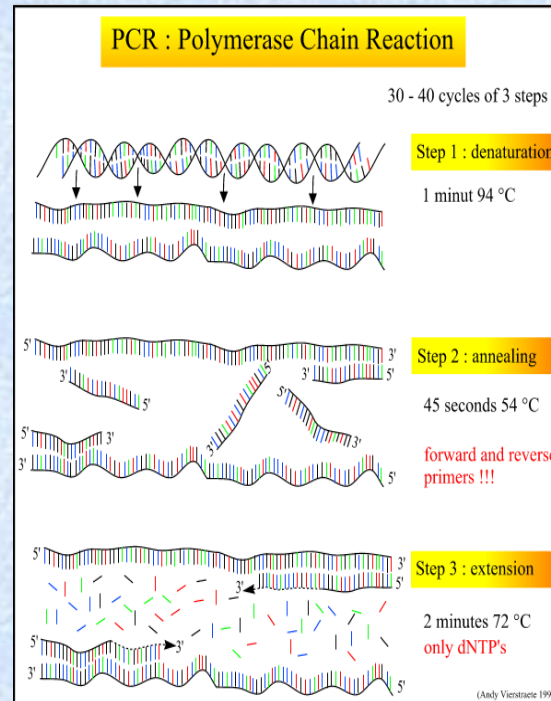


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population genetics tools

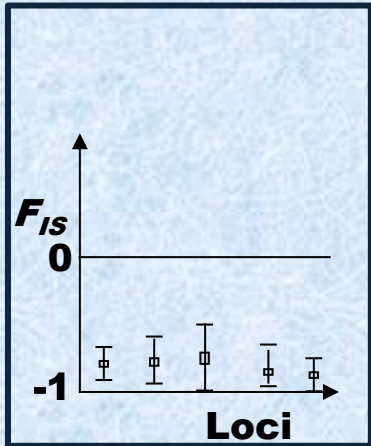


Inferring of basic ecological Parameters such as: reproduction unit size, dispersal, spatial organization and mode of reproduction of the populations

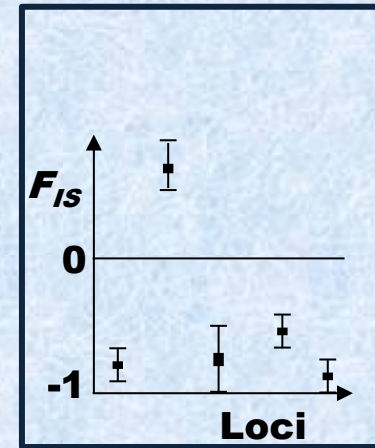


F_{IS} : a measure of the deficit of heterozygote/panmixia, resulting from the reproductive system

For a clonal rate (c) of 100%
→ strongly negative F_{IS} values are expected (Balloux et al., 2003)



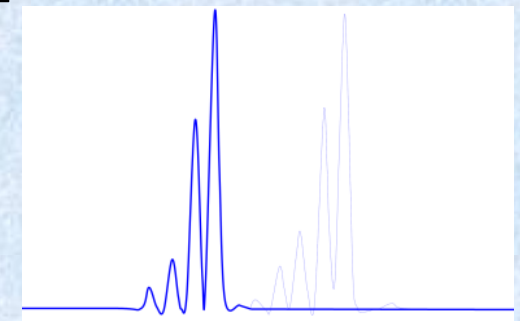
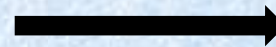
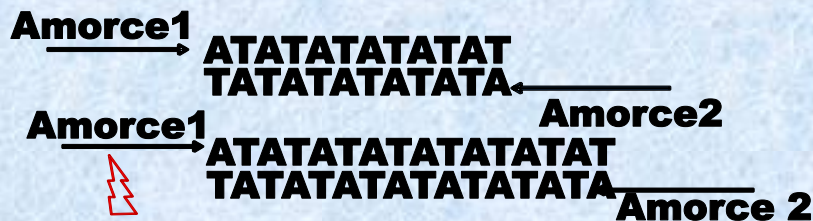
For $99.99 \leq c \leq 95$
→ Large variance F_{IS} values are observed between loci



A useful criterion for detecting very low rates of recombination in clonal organisms

Allelic dropouts and null alleles

Genetic markers (microsatellite)



□ Genetic tools : H_s and F_{IS}

H_s : Genetic diversity

F_{IS} – obs/ analysed data : Local heterozygote deficit

For purely clonal population

$$F_{IS} = \frac{H_s - 1}{H_s} = F_{IS \text{ - exp/ clonal.reproduction}}$$

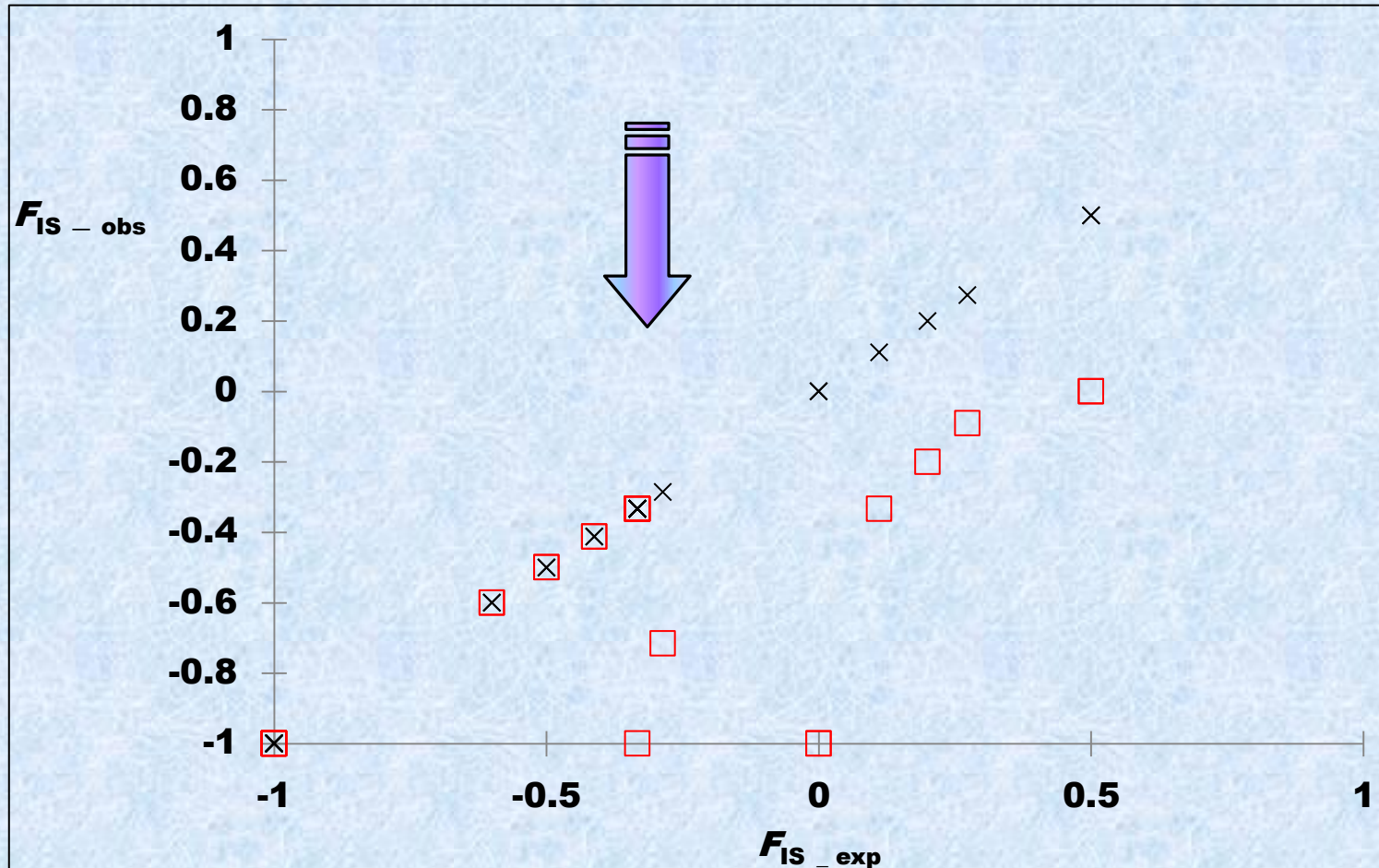


$$\Delta F_{IS} = F_{IS \text{ - exp}} - F_{IS \text{ - obs}}$$



$$|\Delta F_{IS}| \leq 0.05 \times |F_{IS \text{ - exp}}| = \text{Superimposition des } F_{IS}$$

□ Criterion for allelic dropouts, null alleles and rare sex detection



Proportion of superimposed points (%) => Criterion

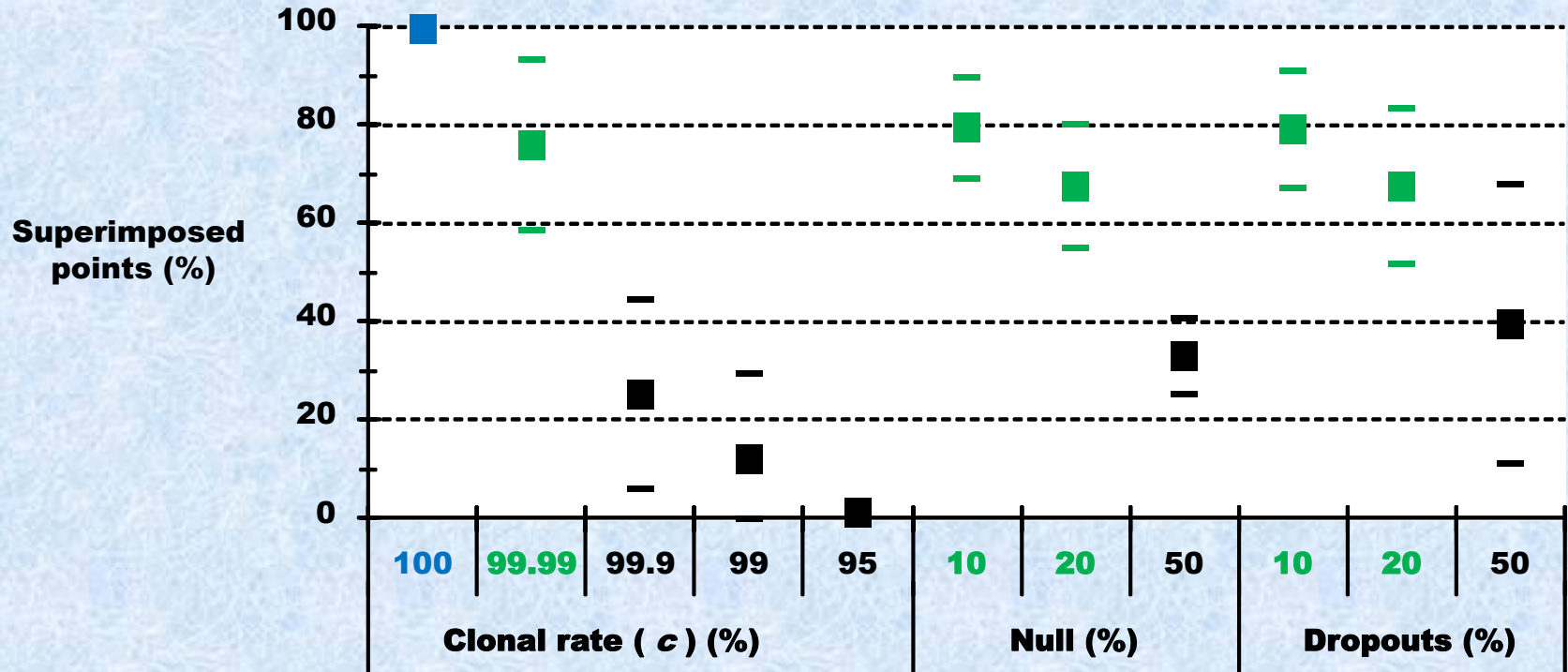
□ Simulations

- **Different models of populations**
- **Variable proportions of allelic dropouts, null alleles and rare sex**

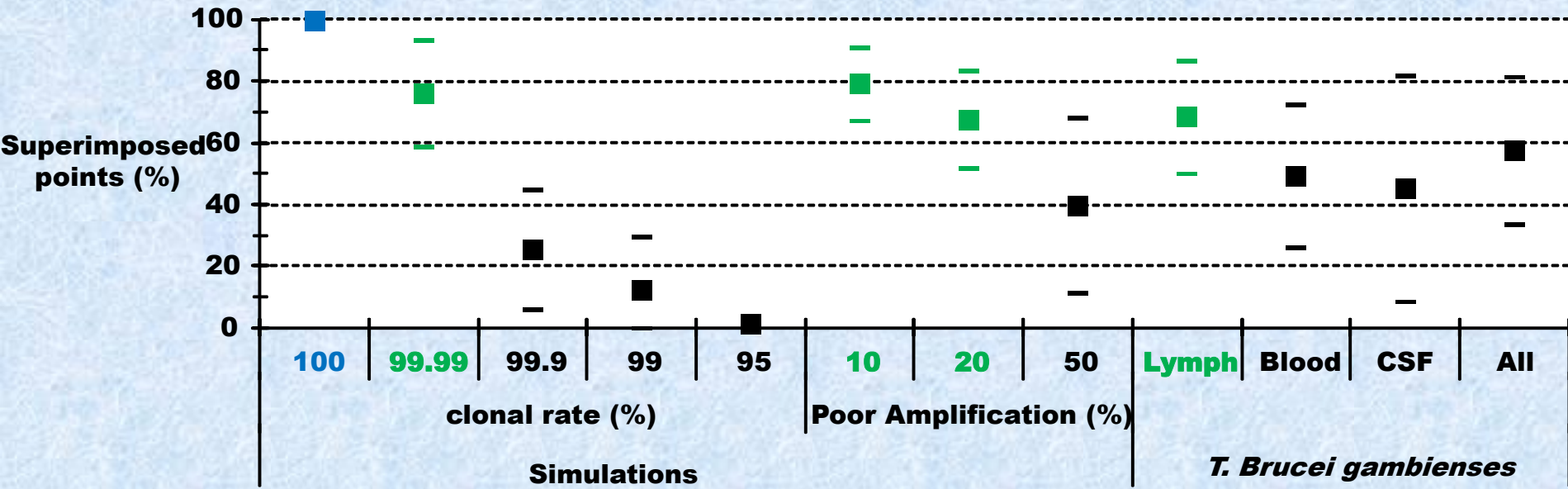
□ Reinterpretation of real datasets

- ***T. brucei gambienses* of Guinean foci (Kaboré *et al.* 2011)**
- ***T. evansi* from Sudan (Salim *et al.*, 2011)**

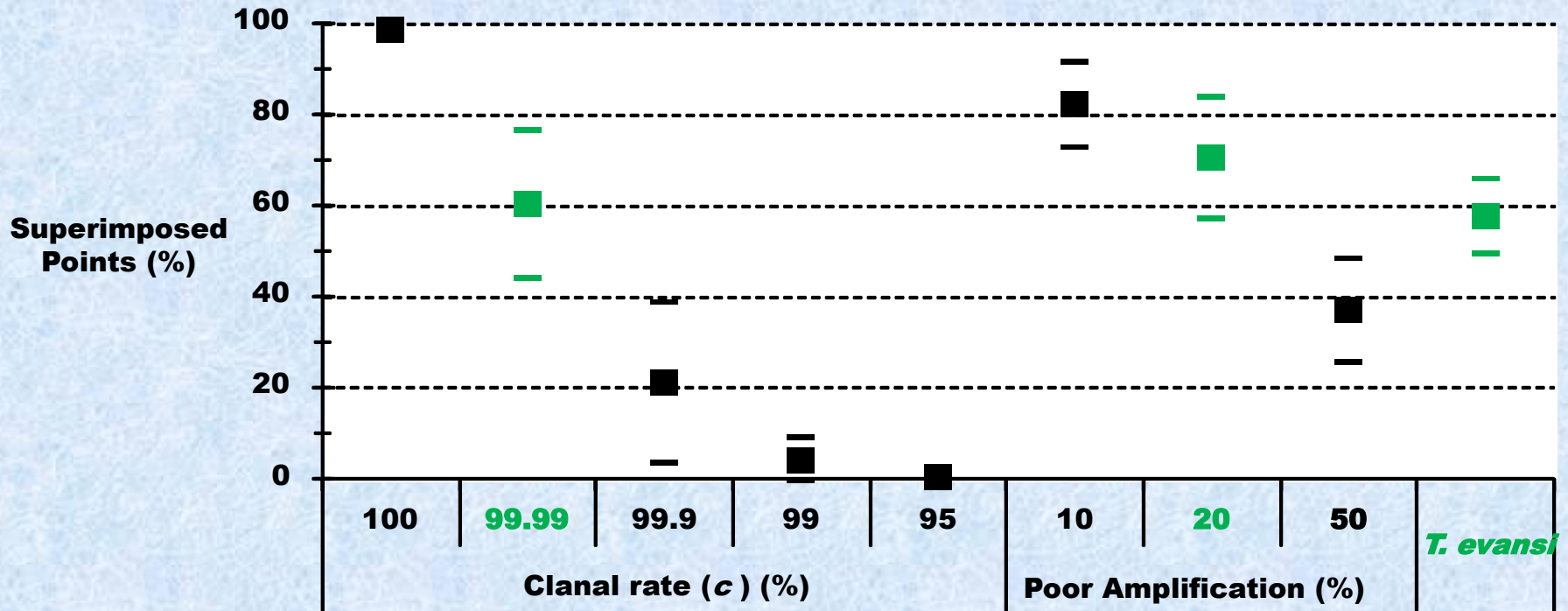
Comparison of allelic dropouts, null allele and rare sex effects



Reinterpretation: *T. brucei gambienses* of Guinean foci (Kaboré *et al.* 2011)



Reinterpretation: *T. evansi* from Sudan (Salim *et al.*, 2011)



- ✓ **The method based on the relationship between H_s and F_{IS} under the hypothesis of clonal reproduction is useful**
- ✓ **It is valid for less than 50% of poorly amplified alleles**
- ✓ **It is not a palliative but represent a useful decision criterion for regenotyping problematic data**



شكرا

Thank you

Merci

For your

Grazie

Attention

Gracias

