## Relative impact of *Wolbachia* genotype and host genetic background on cytoplasmic incompatibility in *Drosophila suzukii*

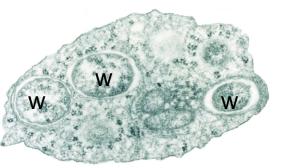


By Mountassira MATIN

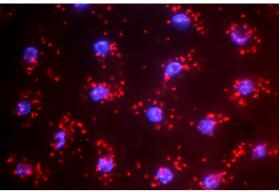


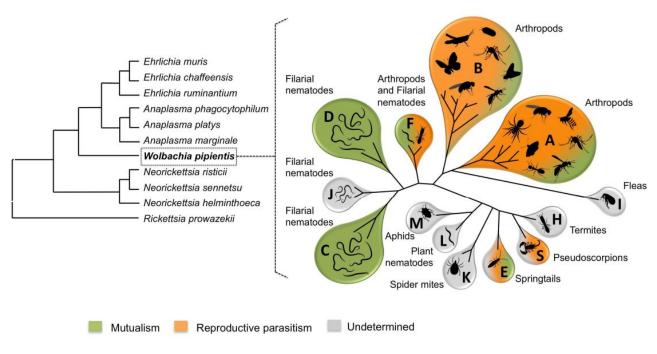
Mentors: Svitlana SERGA, Nicolas RODE Tutor: Mathieu SICARD

### Introduction: Presentation of Wolbachia pipientis



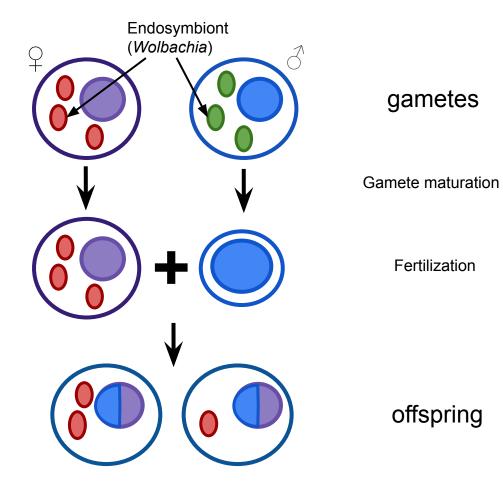
*Wolbachia pipientis* (w) under an electron transmission microscope inside of an insect cell **Example** 



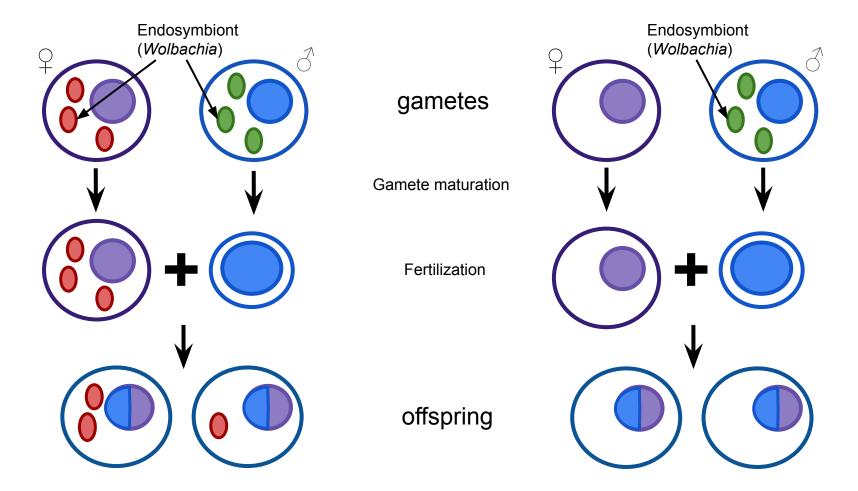


DAPI-stained Drosophila embryo tissue with the nuclear DNA appearing blue and the *Wolbachia* appearing red (Description of Careford (Description

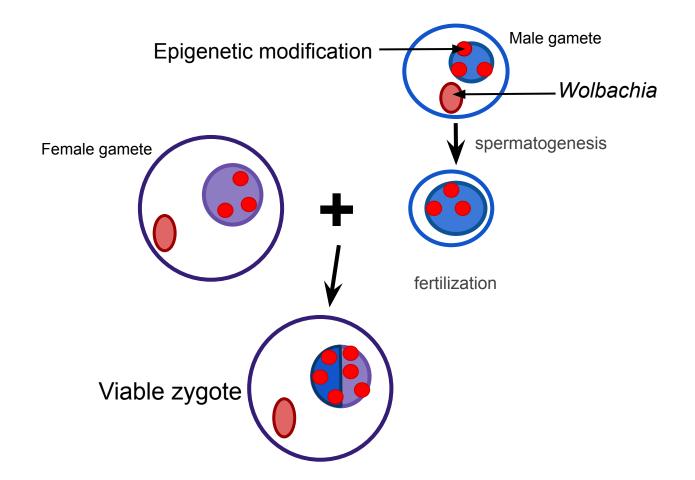
### Introduction: Maternal transmission of Wolbachia



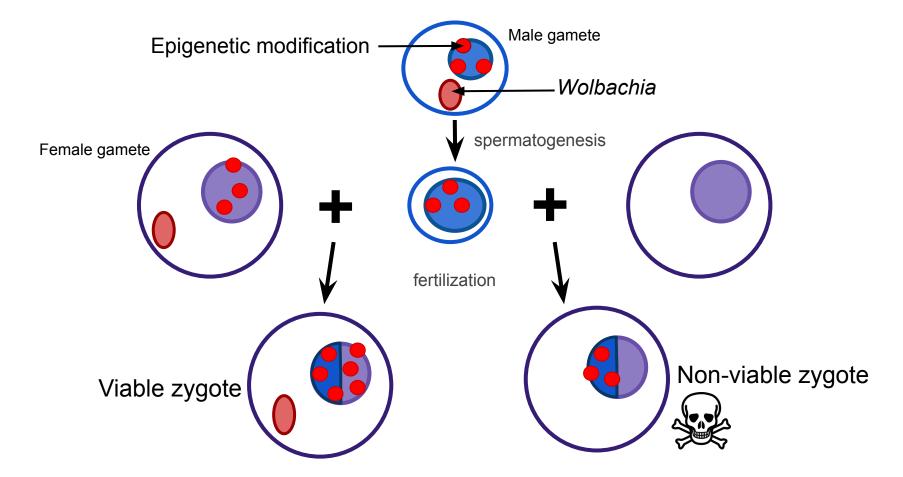
#### Introduction: Maternal transmission of Wolbachia



### Introduction: Explanation of Cytoplasmic Incompatibility (CI)

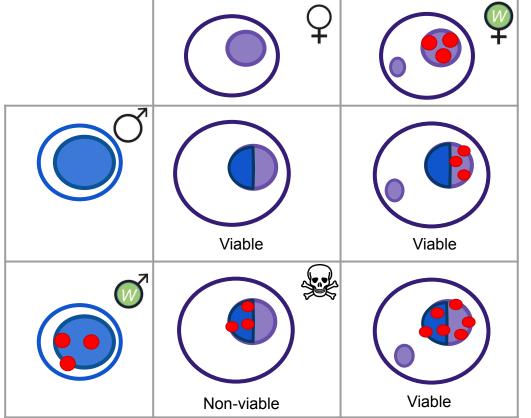


### Introduction: Explanation of Cytoplasmic Incompatibility (CI)



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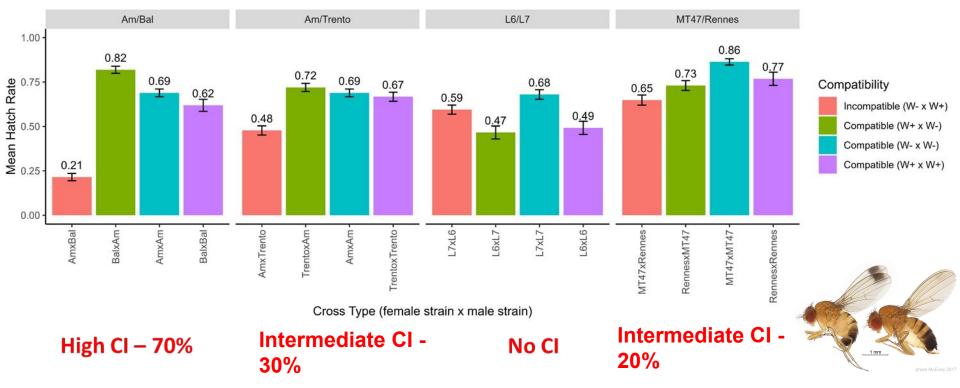
Cytoplasmic Incompatibility (CI): embryonic lethality resulting from the cross between an uninfected female and a *Wolbachia*-infected male



modification

### Introduction: Preliminary work on CI and Drosophila suzukii

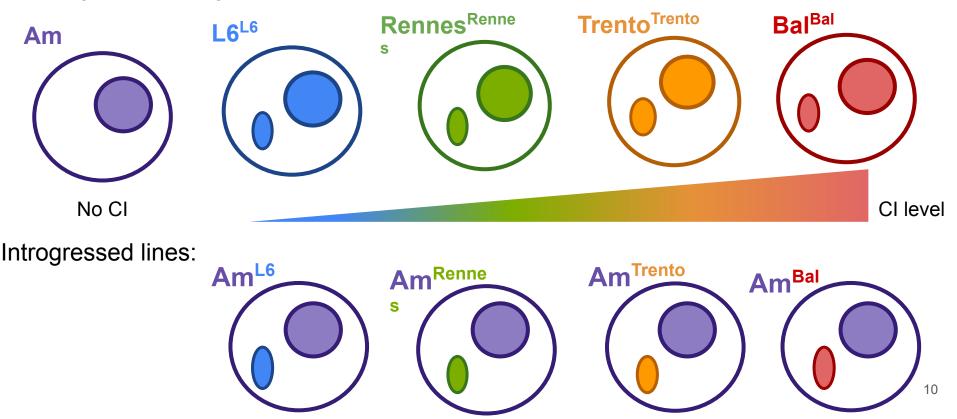
hatch rates for crossings between *D. suzukii* strains Am(uninfected), Baladran(infected), Trento (infected), L6 and L7 (infected) (unpublished data)

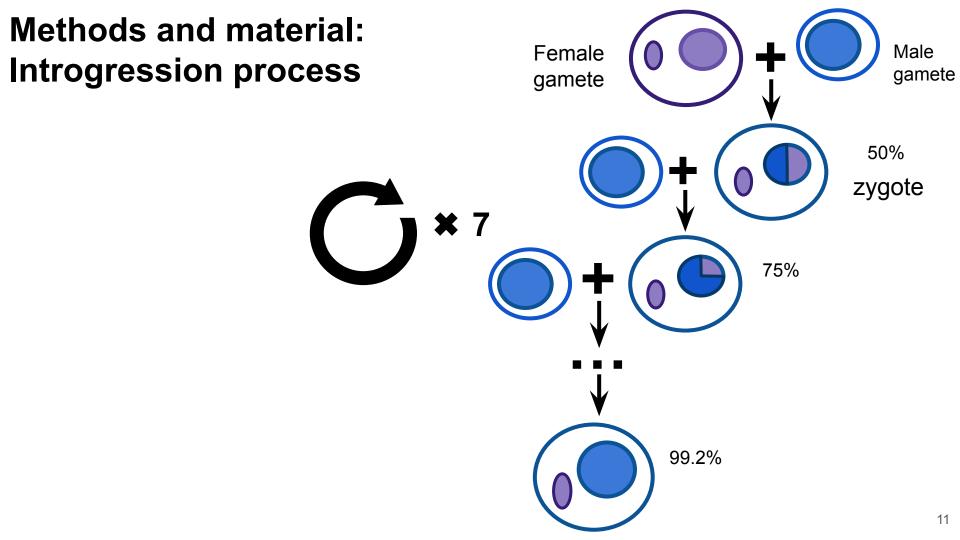


# What causes these variations in CI levels?

### **Introduction: Presentation of the experiment**

**Objective**: test the relative effect of *Wolbachia*'s genetic background versus the male host's genetic background on CI





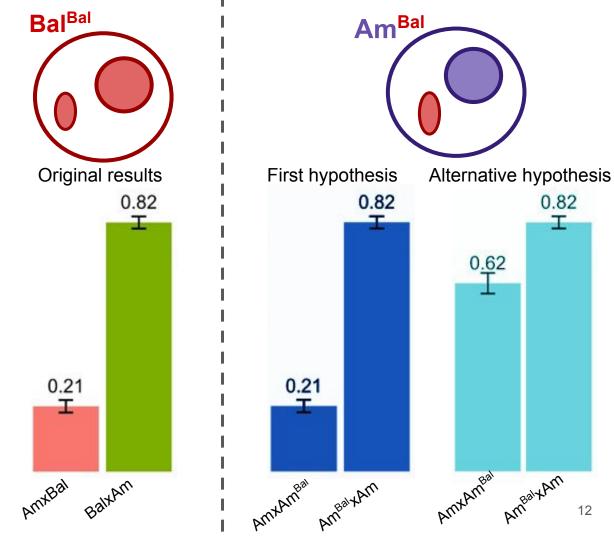
### **Hypotheses**

Incompatible crossing

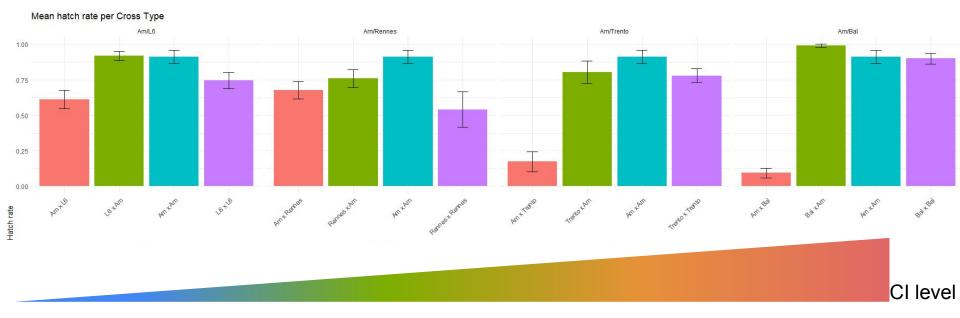
Compatible crossing

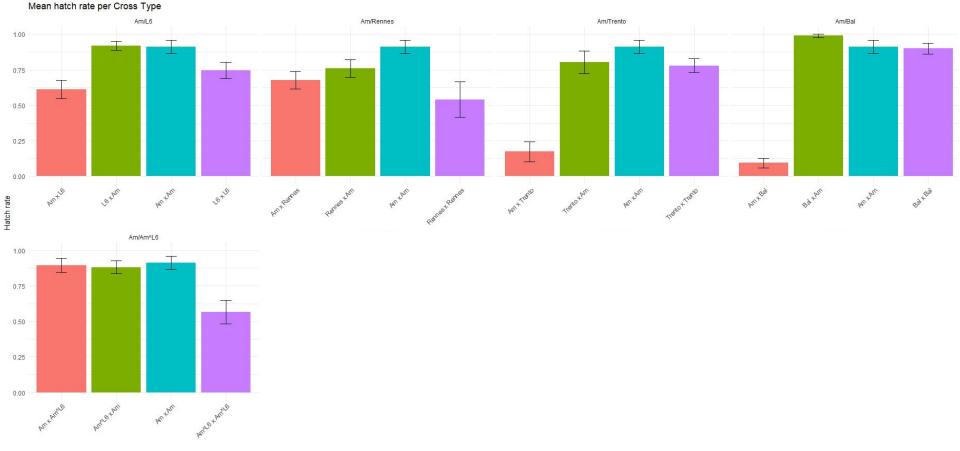
Expected hatch rates with introgressed lines if CI levels only depend on the variant of Wolbachia (same CI levels as expected results)

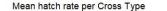
Expected hatch rates with introgressed lines if CI levels are influenced by male host genetic background (different results than expected results)

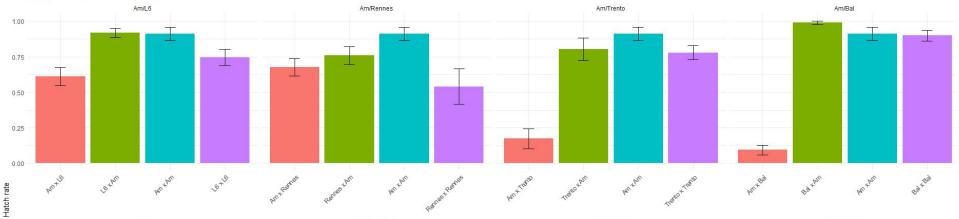


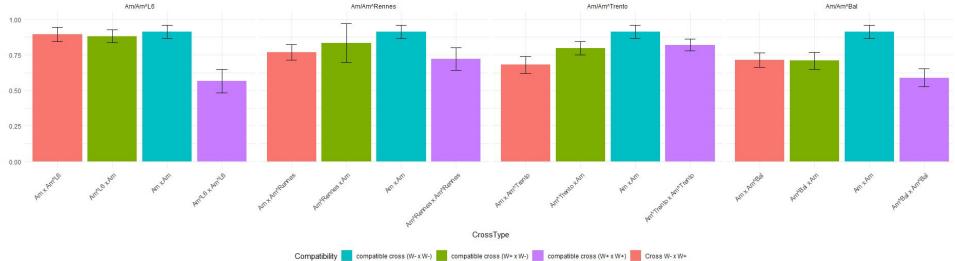
### **Results**











### Model used for statistical analyses

The hatch rate  $(Y_{ij})$  for a cross between a female strain *i* and a male strain *j* carrying a *Wolbachia* variant *k* is:

$$Y_{ijk} = \mu_{ij} + \delta_k + \varepsilon_{jk}$$

 $\mu_{ij}$  = effect of the female genotype *i*, male genotype *j*, and the genetic interaction between genotypes *i* and *j* 

 $\delta_k$  = degree of *Wolbachia*-induced incompatibility in crosses between the *i*th female strain and the *j*th male strain for a *Wolbachia* of *k* genotype

 $\varepsilon_{jk}$  = interaction between *Wolbachia* and male genotypes k and j respectively

### **Conclusion and perspectives**

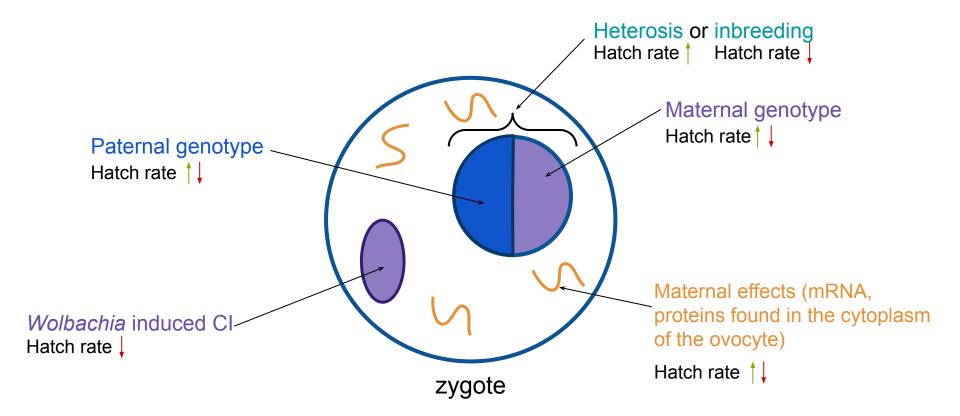
- Cytoplasmic incompatibility is heavily influenced by the male host's genetic background
  - L6 genotype contributes little variation to CI levels (weak interaction between host genotype and CI)
  - **Rennes** genotype contributes little variation to CI levels (weak interaction between host genotype and CI)
  - Trento genotype contributes a lot to variation in CI levels (strong interaction between host genotype and CI)
  - Bal genotype contributes a lot to variation in CI levels (very strong interaction between host genotype and CI)

• Next step: Find the genetic architecture of susceptibility to *Wolbachia*-induced incompatibility (either one strong-effect locus or several weak-effect loci)

Thank you for your attention !

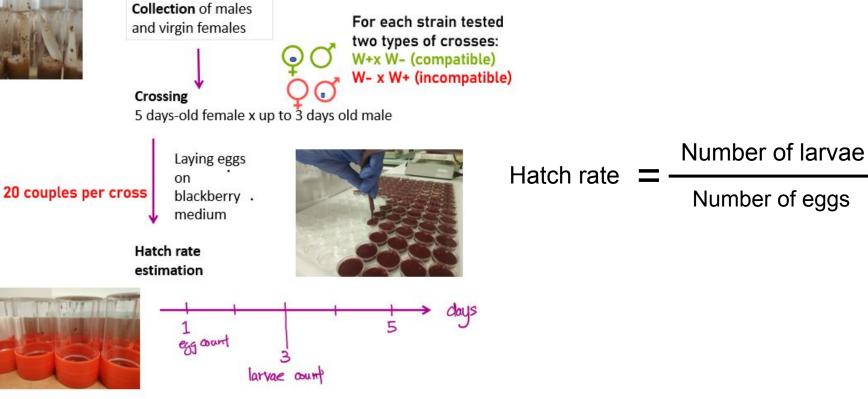


### Hatch rate can be influenced by other factors



### Methods and material: CI assay





5 Petri dishes per couple