# Co-evolution of house mouse and an intracellular parasite, Eimeria spp.

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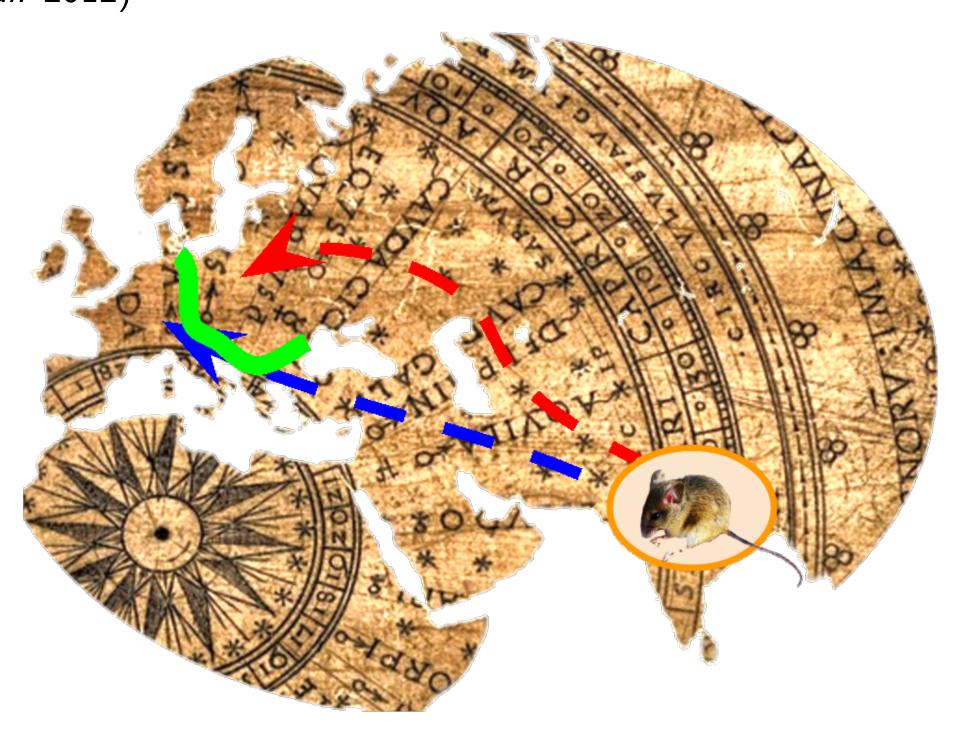
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#### Context

• House Mouse Hybrid Zone, 20km wide, formed by hybrids of Mus musculus domesticus and Mus musculus musculus. After 500,000 years in isolation, secondary contact 5000 years ago (Macholán et al. 2012)



• Eimeria spp., obligate intracellular apicomplexan parasite. Two major clades (A & B) of Eimeria spp. identified (3 markers) in the mice of the hybrid zone (Jost 2016)

## Aims of the study

- Investigating the vigor/resistance of hybrids of house mouse to their parasite Eimeria spp. using prevalence and intensity data for parasite strains throughout the House Mouse Hybrid Zone.
- 2. Looking for evidence of **local adaptation** between the murine host and its Eimerian parasite

## Material & Methods: Field study

- Annual sampling every September 104 mice in 2015 from 48 localities 165 mice in 2016 from 45 localities upcoming sampling in 2017 Brandenburg area (Germany)
- Oocyst counted in mice feces All parasite strains genotyped using 3 markers, then assigned to an haplotype
- Adaptation of the method of Stuart J.E. Baird (Baird et al. 2012): Maximum likelihood analysis explicitly linking parasite abundance to a gradient along the hybrid index (as a proxy of host heterozygosity), generalized linear model with negative binomial distribution

 $Parasite\ load \sim mouse\ heterozygosity\ level*parasite\ strain$ 

• R package under development: https://github.com/alicebalard/Parasite\_Load

## Material & Methods: Cross infection

#### Pilote experiment

- Parasite strains :
- Eimeria haplotype A laboratory strain Eimeria falciformis (Heitlinger et al. 2014)
- 2. Eimeria haplotype B strain isolated in the wild
- Host strains :
- . WSB Wild-derived inbred strain. Isolated from wild Mus musculus domesticus Region of capture: Eastern Shore, Maryland
- 2. PWD Wild-derived inbred strain. Isolated from wild Mus musculus musculus Region of capture : near Prague, Czech Republic
- 3. **WP** Hybrids between the 2 previous commercial strains

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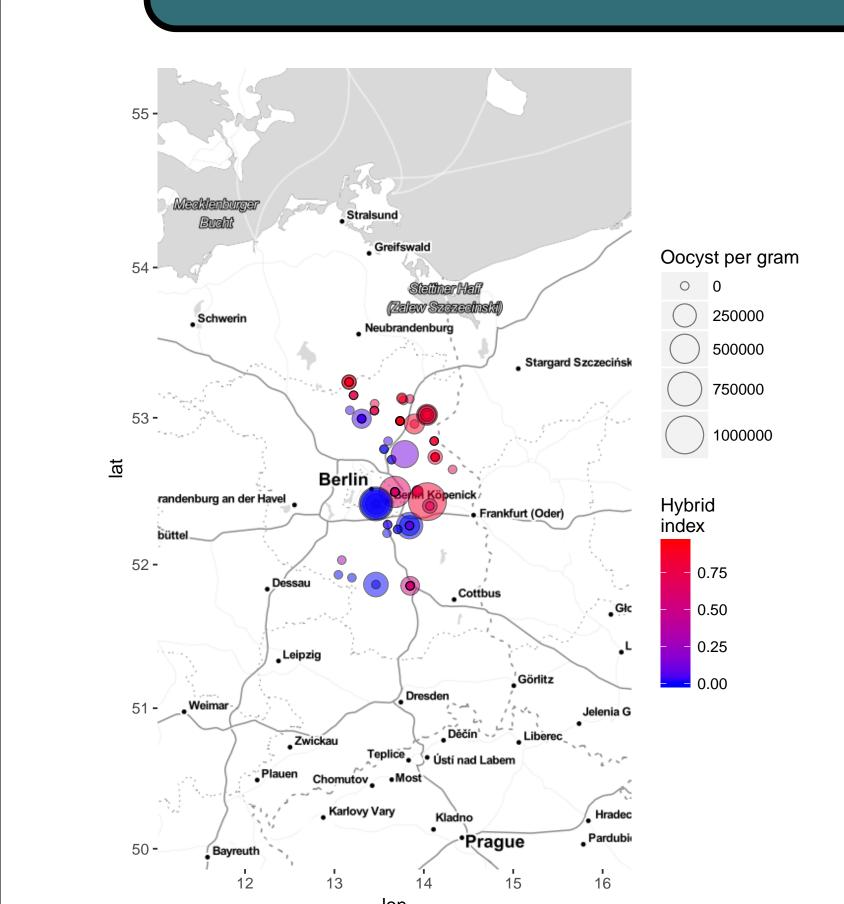




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# Preliminary results: exploring hybrid vigor/resistance in the wild



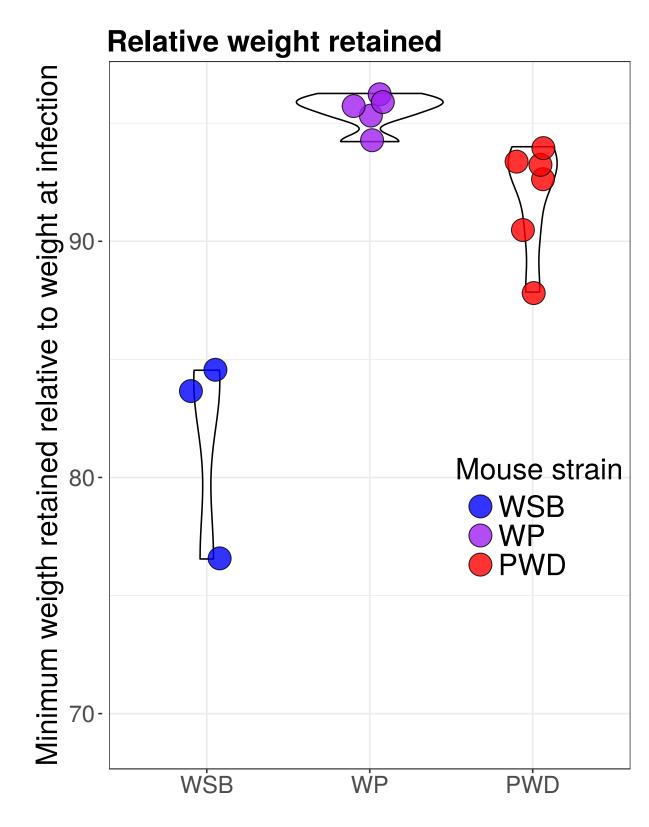
- Prevalence 19% infected farms in 2015 2016: pending analysis... 2017: upcoming sampling
- Hybrid index assigned to all mice 0 = Mus musculus domesticus 1 = Mus musculus musculus in between = hybrids

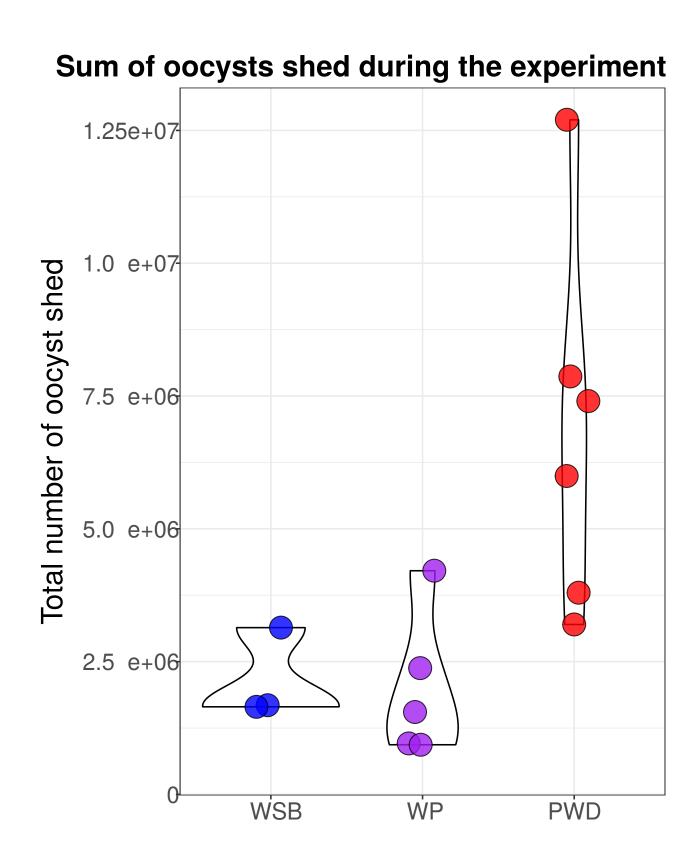
Goal using our glm.hybrid model to assess the existence of hybrid vigor/resistance, taking into account the parasite strains

## Evidence of local adaptation

Results of the infection experiment :

- Eimeria strain haplotype B has lowest parasite shedding in mice strains WSB compared to PWD, for a highest relative weight loss: indication of local adaptation
- Mice hybrids lost less weight and were less infected than the pure strains Possible hybrid vigor (limitation : unknow effect of general heterosis)





### Perspective

- Next cross infection experiment: verify our hypotheses (hybrid vigor, local adaptation), measure the effect of heterosis (within subspecies heterosis vs between subspecies)
- Assess local adaptation in other parasite strains
- Analyse of divergence scenarios for *Eimeria* spp. based on whole genomes and comparison of models of coalescence and cospeciations with their murine hosts (beyond the house mouse).
- Investigation of loci of coevolution, identifying parasite genes under divergent selection in the two house mouse subspecies. The coevolving loci corresponding on the host side will be investigated.

#### References

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