

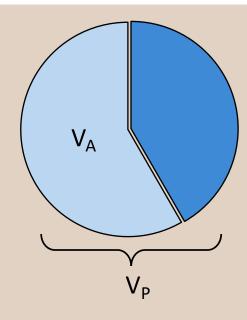
Why estimate heritability?

- Variance components are a key component of breeding
- Heritability h² will determine the best selection strategy
- Unveil if there is genetic variation for a trait
- Insights useful also for rearing and management

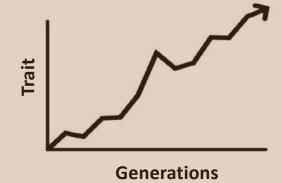
AIM

Estimate the heritability of larval size in BSF

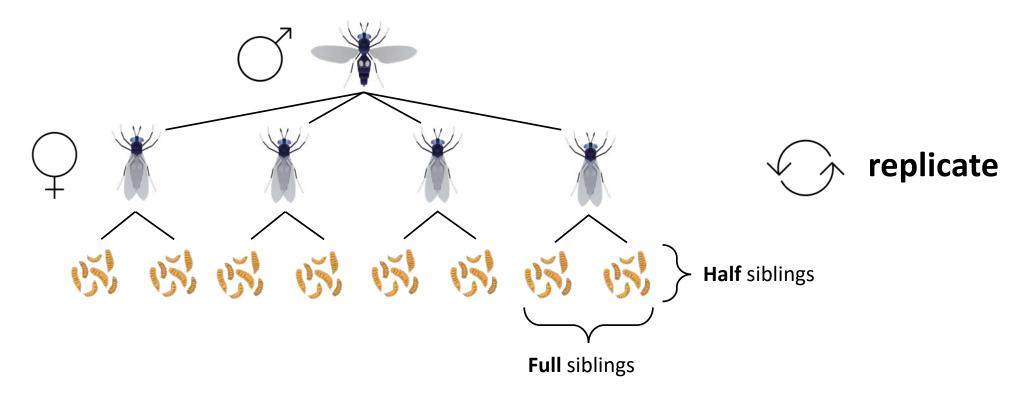




$$h^2 = \frac{V_A}{V_P}$$



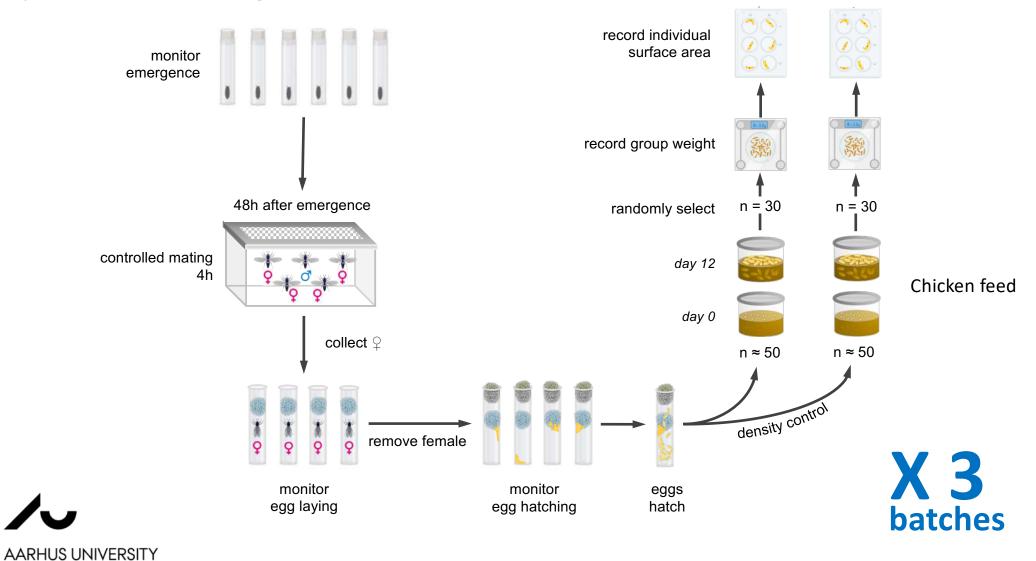
Mating and generation of family groups





BSF polygynous mating: Jensen et al. (in review – preprint: https://doi.org/10.1101/2024.09.09.611978)

Experimental design

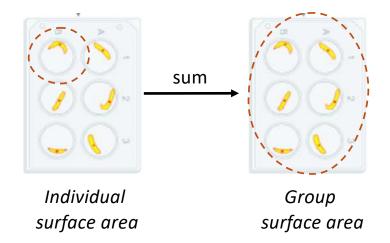


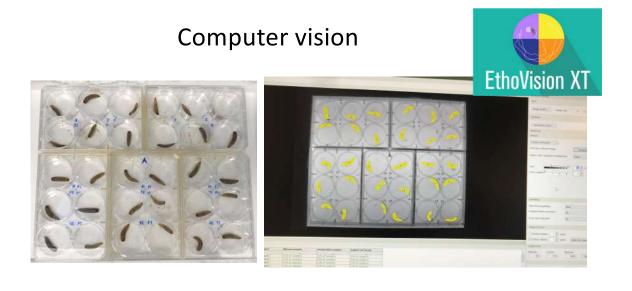
Traits and phenotyping



Manual weighing

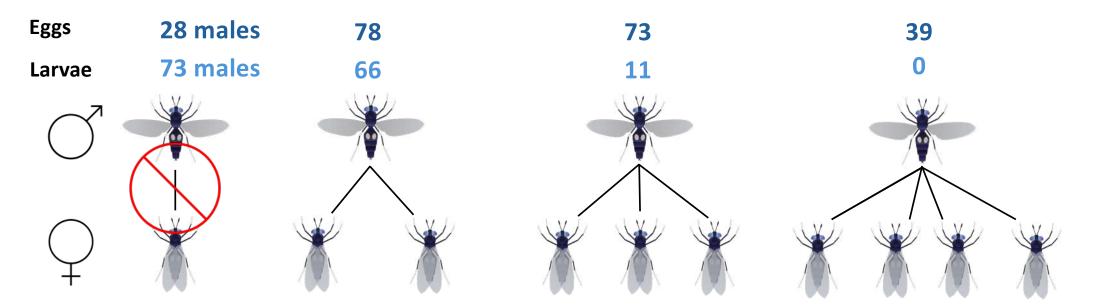
Group weight





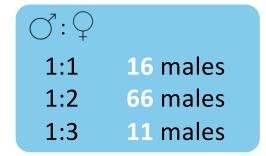
Laursen et al. 2021

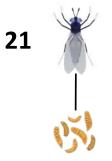
Mating success

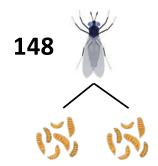


559 females with eggs \rightarrow 238 females with larvae \rightarrow 181 measured \rightarrow 169 used for analyses

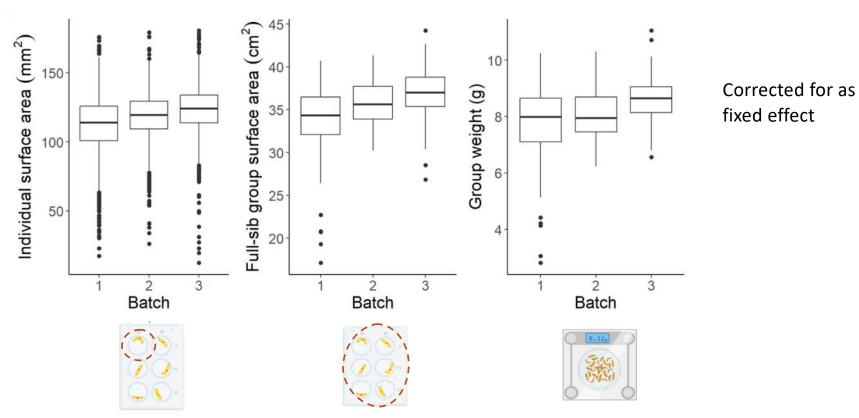








Effect of batch on the three traits





Genetic models and calculations



Individual surface area = μ + batch + sire + dam + cup + e

$$h^2 = \frac{4\sigma_S^2}{\sigma_p^2}$$





Group traits = μ + batch + *sire* + *dam* + *e*

$$h^2 = \frac{4\sigma_s^2}{\sigma_s^2 + \sigma_d^2 + \sqrt{30} * \sigma_e^2}$$



Heritability and common environment

Trait	Unit	h^2	c^2	σ_s^2	σ_d^2	σ_c^2	σ_e^2
Individual surface area	mm²	0.40 (0.11)	0.21 (0.02)	29.1 (9.9)	11.0 (8.6)	60.4 (7.7)	191.2 (2.8)
Group surface area	cm ²	0.39 (0.12)	-	3.5 (1.1)	1.6 (0.8)	-	5.6 (0.6)
Group weight	g	0.30 (0.10)	-	0.29 (0.10)	0.15 (0.09)	-	0.61 (0.07)



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Conclusion

- Strong potential for selection for larger BSF larvae
- h² estimates higher than previous findings in BSF (Bouwman et al. 2022 and 2024)
- c² align with previous findings in BSF, house fly and mealworm

Next steps

- Trait correlations
- Trait selection for breeding





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