

Genetic constraints underlying human reproductive timing in a pre-modern Swiss village



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Introduction

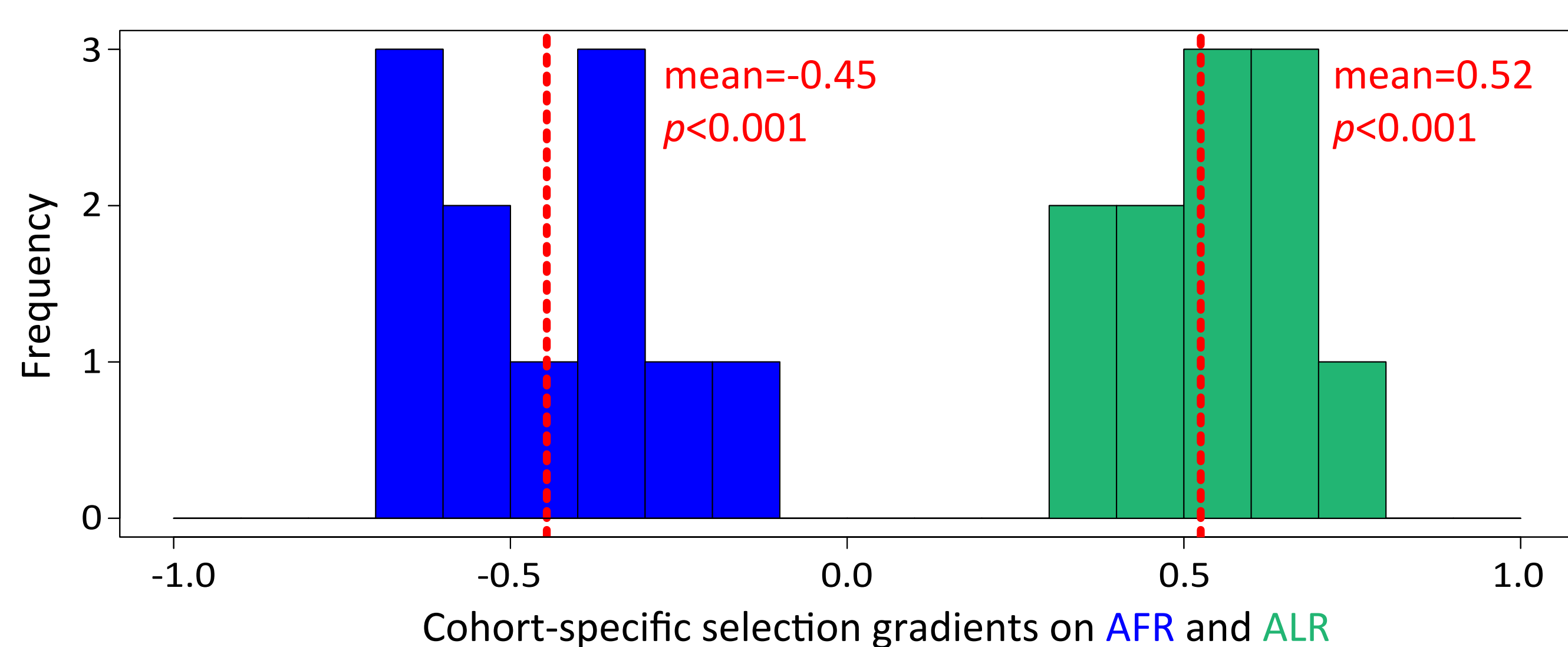
Trade-offs between reproductive output in early and late life are central to life-history theory. However, the specific trade-off between **age at first reproduction (AFR)** and **age at last reproduction (ALR)** has received little attention, especially in long-lived species with a pronounced reproductive senescence such as humans. Based on Catholic parish records we reconstructed genealogies for most inhabitants of a 19th-century, medium-sized, rural village in Switzerland (800-1300 inhabitants), and use these data to

Aims

1. quantify natural selection acting on reproductive timing
2. estimate the underlying additive genetic (co)variances
3. use these to predict evolutionary responses to selection
4. test for signs of reproductive ageing in both men and women

1.

Natural selection on **age at first reproduction (AFR)** and **age at last reproduction (ALR)**



- Strong and significant linear selection for an **earlier** first and **later** last reproduction in both men and women

2.

Additive genetic variation and covariation

		Women			Men			Narrow-sense heritabilities
		h^2	AFR	ALR	h^2	AFR	ALR	Genetic variances
AFR		0.45	1518	0.85	0.81	4977	0.91	Genetic covariance
ALR		0.22	1197	1321	0.54	4926	5855	Genetic correlation
		in bold: $p<0.05$						

- Significant genetic variation in **AFR** in both sexes
- Significant and strongly positive genetic correlation between **AFR** and **ALR** in men

3.

Predicted phenotypic responses to selection in months per generation

	Direct response		Total response	
	AFR	ALR	AFR	ALR
Women	-12.0	9.1	-3.8	-0.4
Men	-33.7	35.1	-4.1	1.8

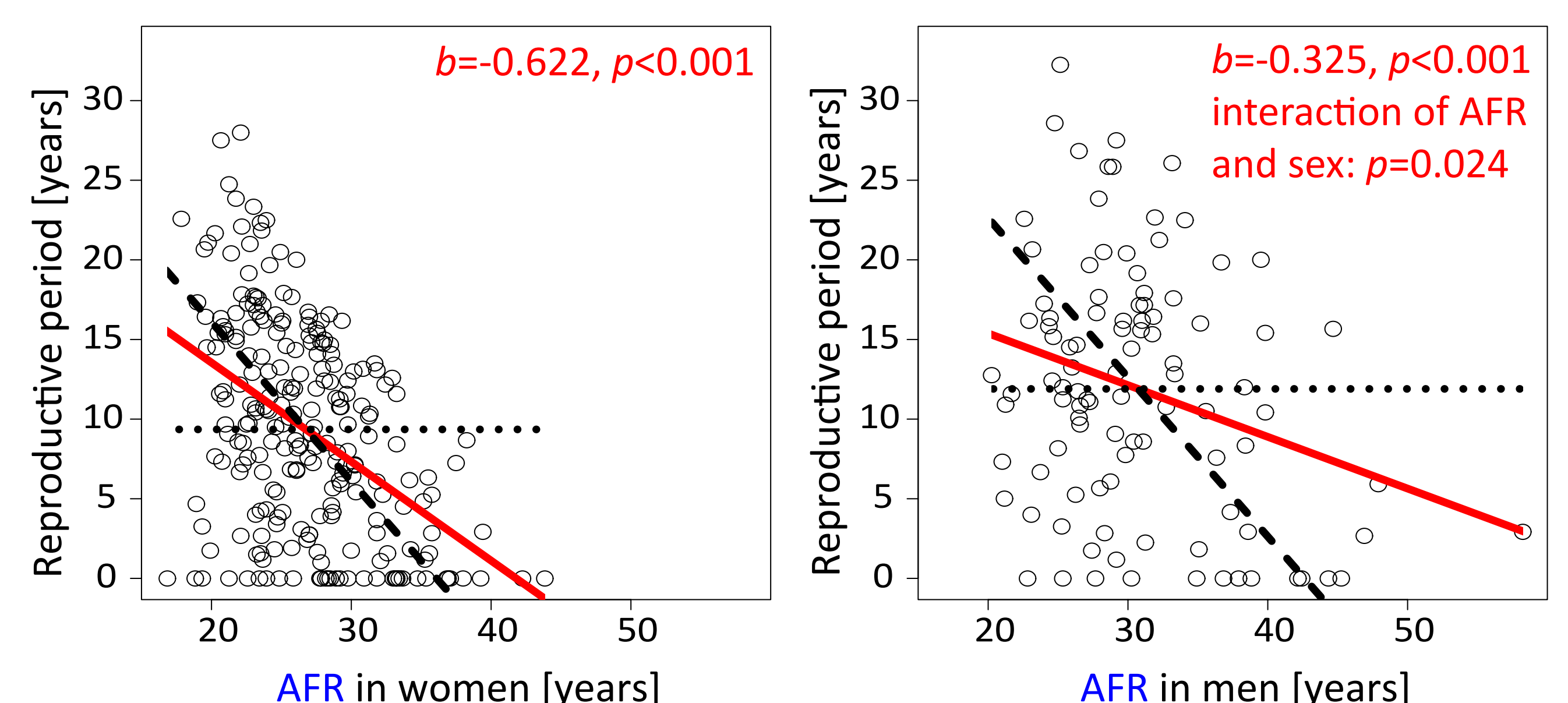
- **AFR** predicted to decrease in both sexes (adaptive)
- **ALR** predicted to decrease in women (maladaptive) and increase in men (adaptive)
- Substantial genetic constraints in both **AFR** and **ALR** because direct predicted responses exceed total responses by far

Conclusions

- Due to substantial selection on **AFR** and **ALR** (1) and a genetic correlation between the two traits (2), reproductive timing in the study population is strongly constrained (3)
- This constraint might provide an ultimate explanation for the signs of reproductive ageing we found (4)
- By increasing the costs of reproducing late, a genetic correlation between **AFR** and **ALR** will cause a further decrease in **ALR**

4.

Evidence for reproductive ageing



- Postponing **AFR** shortens the reproductive period in both sexes
- However, this reduction is twice as strong in women
- When correcting for variation in the reproductive period, selection favors both an early **AFR** and **ALR** (data not shown)
- This indicates reproductive senescence also before the onset of reproductive cessation

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Photograph of study village: www.ugs-linth.ch

