



Is product threshold model better than additive threshold model for studying A.I. result? in sheep

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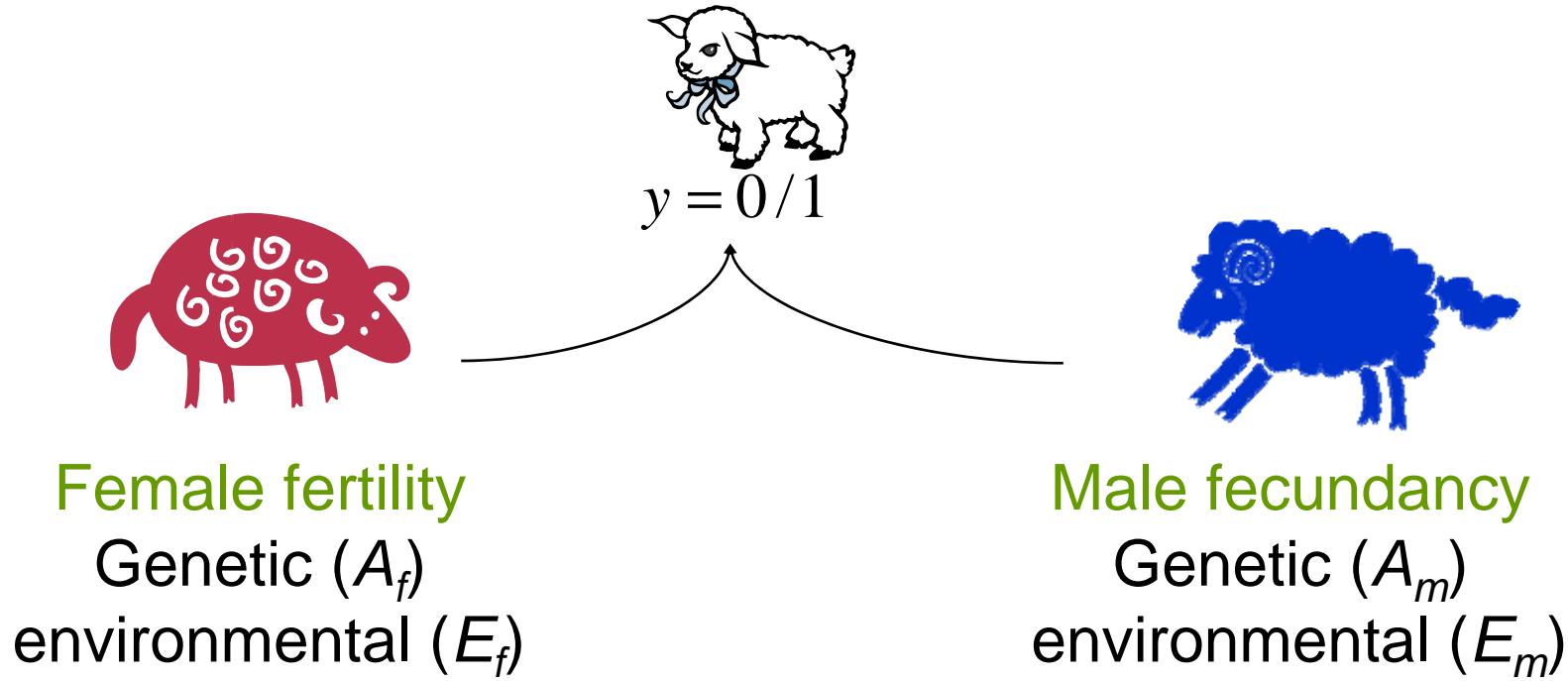
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Problem



**How to model a binary phenotype
dependent on 2 individuals ?**

Possible solution 1

Additive threshold model

$$P(\text{AI success}) = \Phi(\underbrace{A_f + E_f}_{\text{female}} + \underbrace{A_m + E_m}_{\text{male}})$$
$$\qquad\qquad\qquad \overbrace{\qquad\qquad\qquad}^{\Phi(A_f + A_m + E)}$$

Is it the right association ?

Possible solution 2

Additive threshold model

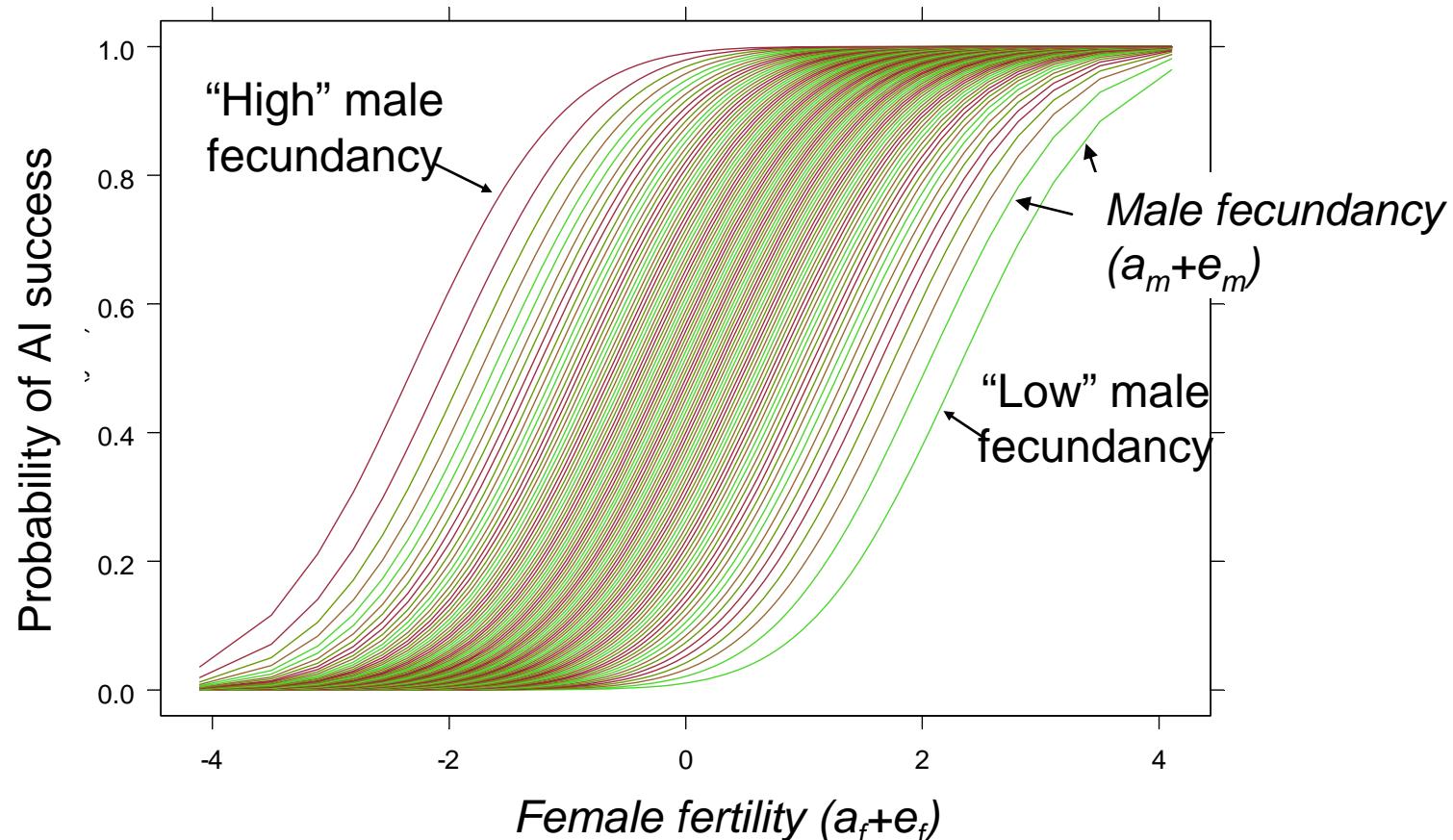
$$P(\text{AI success}) = \Phi(\underbrace{A_f + E_f}_{\text{female}} + \underbrace{A_m + E_m}_{\text{male}})$$

Product threshold model

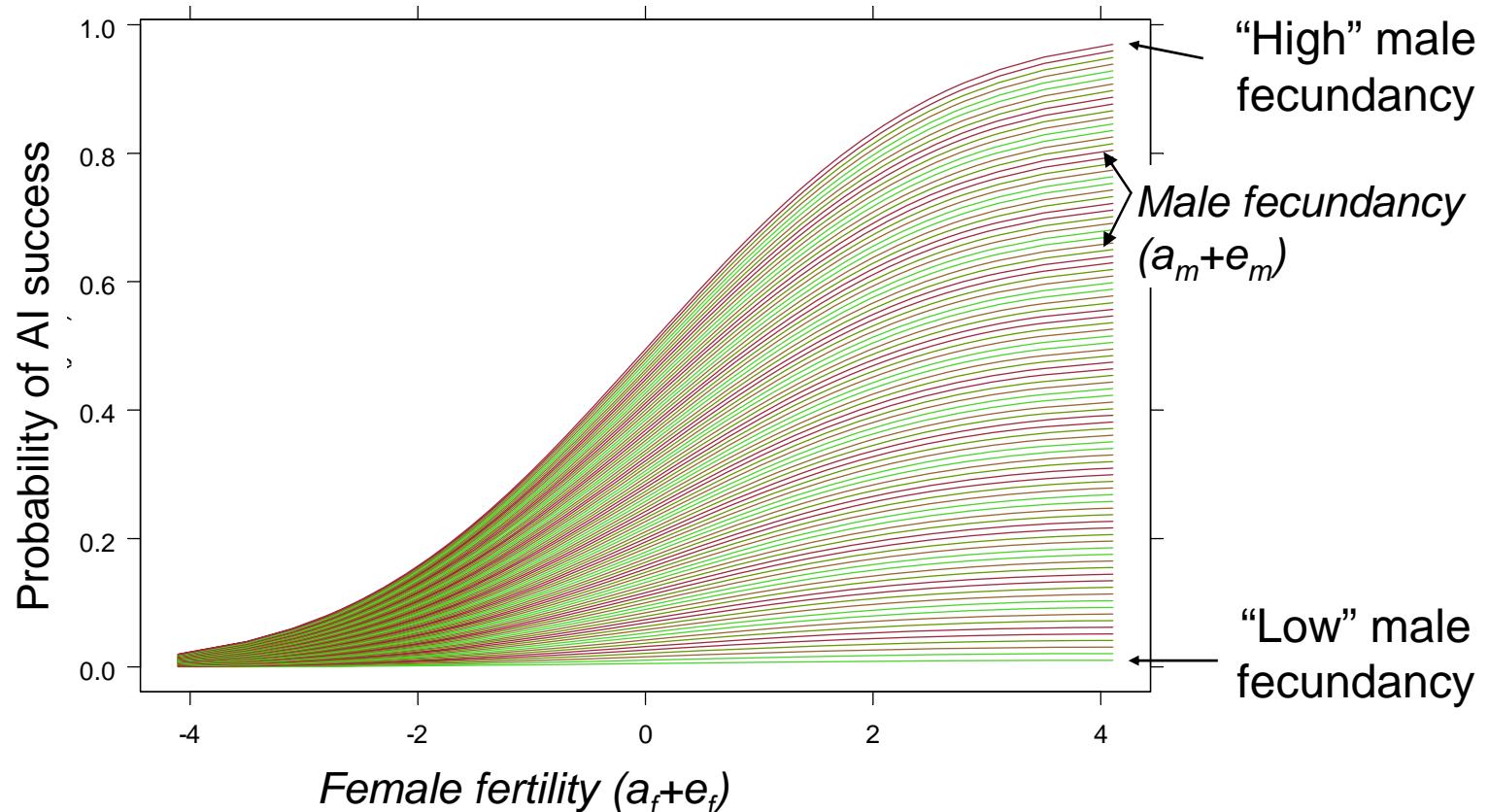
$$P(\text{AI success}) = P(\text{female fertility}) \times P(\text{male fecundancy})$$

$$\underbrace{\Phi(A_f + E_f)}_{\text{female}} \quad \times \quad \underbrace{\Phi(A_m + E_m)}_{\text{male}}$$

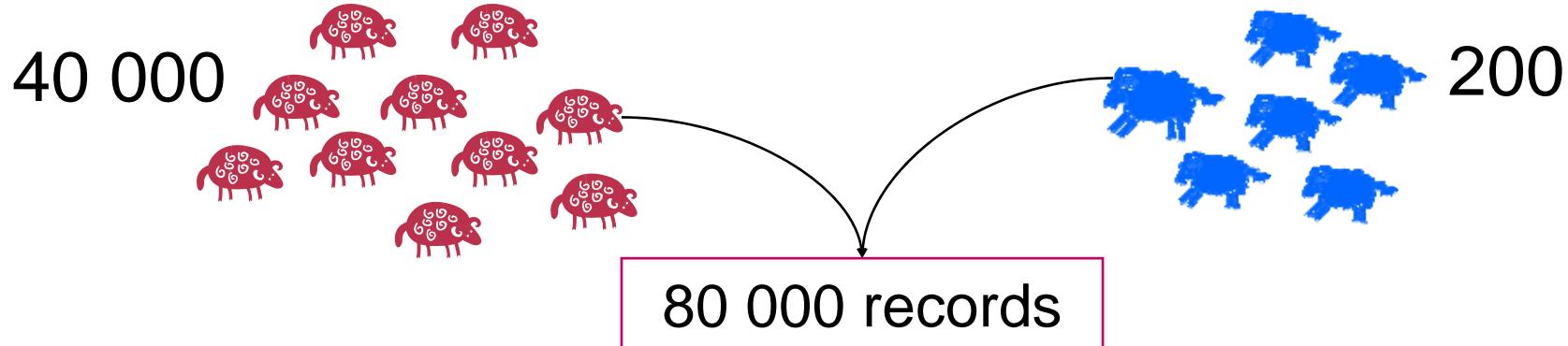
Consequences of the additive threshold model



Consequences of the product threshold model



Simulation design



$$P_f \left\{ \begin{array}{l} a_f \\ \text{year} \end{array} \right. \quad P_m \left\{ \begin{array}{l} a_m \\ \text{year} \\ \text{age of the male} \end{array} \right.$$

$$P(\text{AI success}) = 50\%$$



Parameter estimations

Bayesian approach, Gibbs sampling,
Modification *TM program* (Andres Legarra)

- Product threshold model
 - Parameters estimable?
 - Program works?

Good estimations

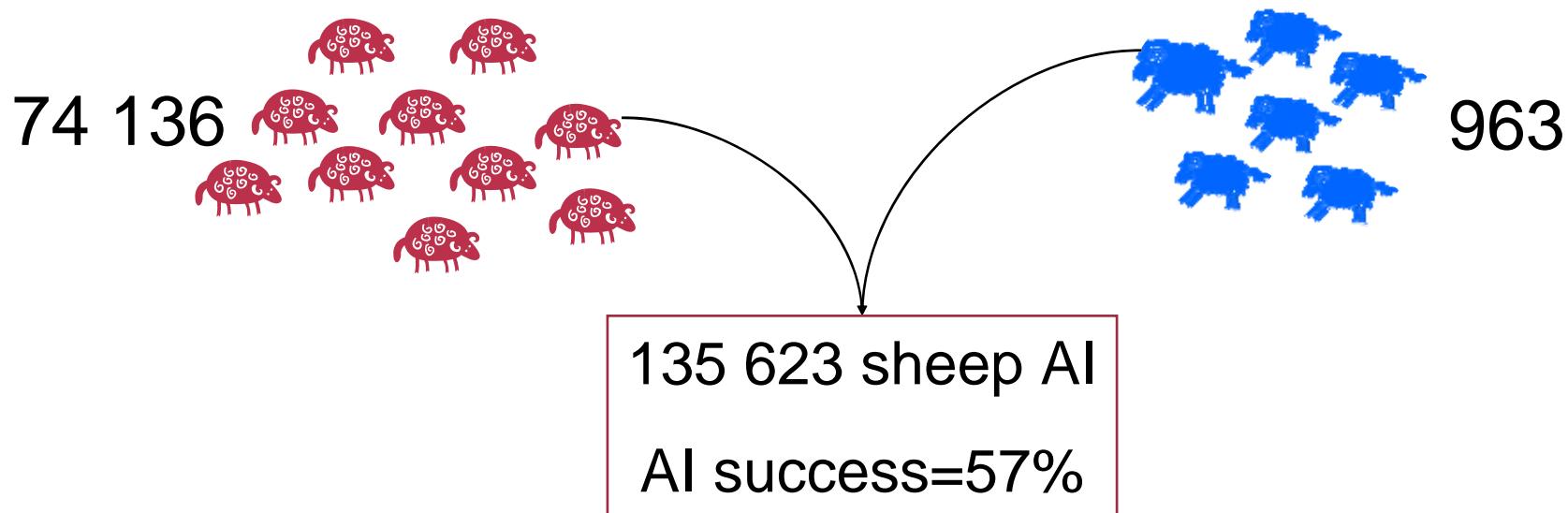
	Simulated	Result	
Genetic Male fecundancy variance	0.25	0.25	[0.21,0.29]
Genetic Female fertility variance	0.36	0.36	[0.33,0.41]
Genetic correlation	0.30	0.21	[0.03,0.41]
year effect	Male	-1.50	-1.46 [-1.52,-1.41]
		0.50	0.53 [0.47,0.60]
year effect	female	1.50	1.54 [1.47,1.63]
		1.00	1.00 [0.97,1.04]
Male age effect	2.00	1.99	[1.94,2.03]

Good correlation between BV

<i>Correlation EBV</i>	EBV
Male fecundancy	0.97
Real BV	
Female fertility	0.94



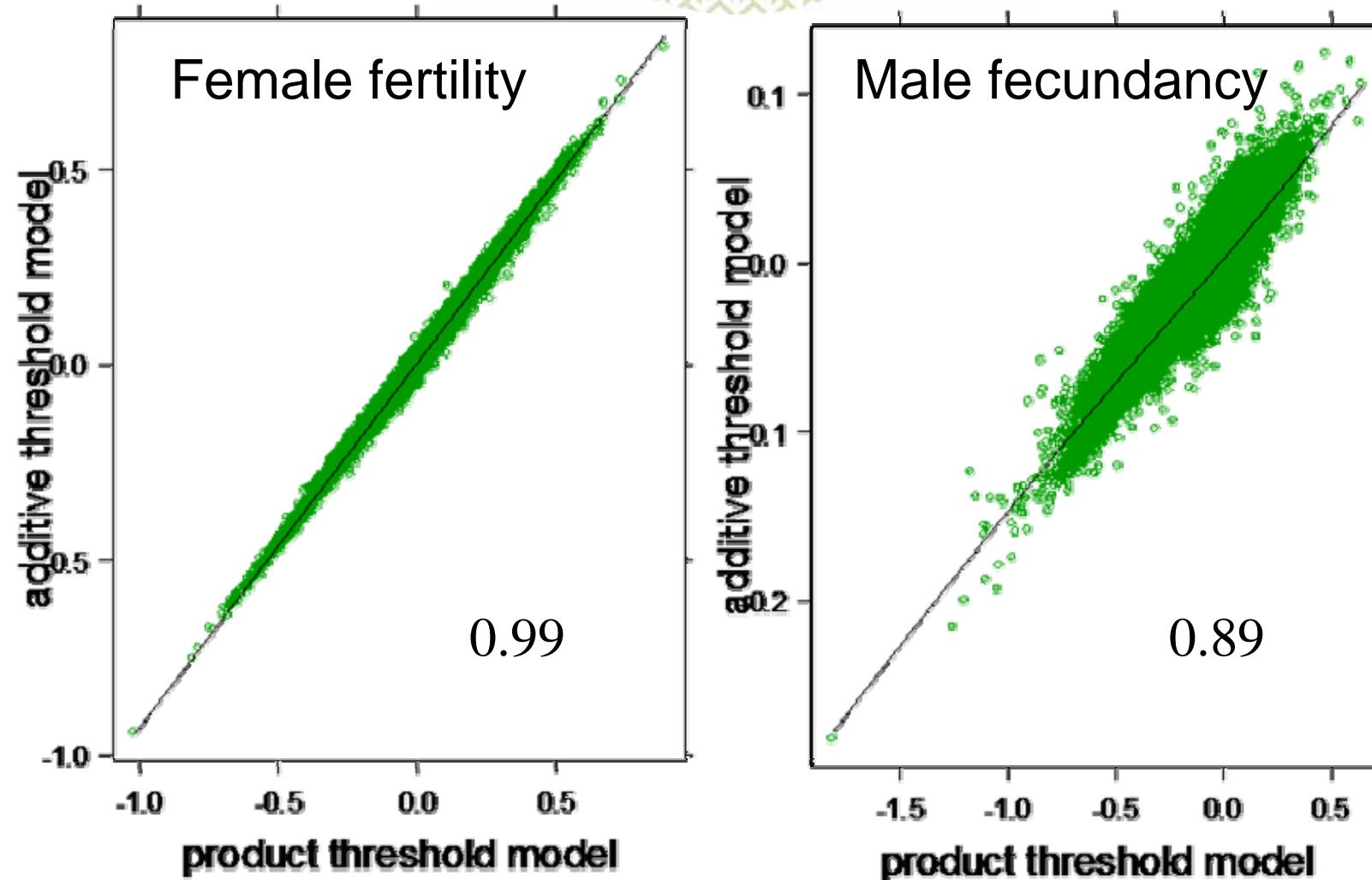
Field data



models

- Product threshold model
- Additive threshold model

Field data: preliminary results



Discussion / conclusion

Advantage

- In accordance with biology
- Easy to implement
- Estimate effects for male and female separately

Disadvantage

- Estimate parameters for the unobserved phenotypes

$$0.75=0.87*0.87$$

Convergence limit:
extreme case problem