

Maternal and social genetic effects on growth of piglets from birth till weaning

A.C. Bouwman*, R. Bergsma†, N. Duijvesteijn† and P. Bijma*

* Wageningen University, Animal Breeding and Genomics Centre

† IPG, Institute for Pig Genetics

Aniek.Bouwman@wur.nl



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Animal Breeding &
Genomics Centre

Introduction

- Social interactions due to:
 - competition for limited resources
 - social behavior

(Muir, 1996 & 2005)

- Consequences for selection response

(Muir, 2005; Bijma et al. 2007)

- Heritable social effects found for growth of finishing pigs

(Arango et al. 2005; Bergsma et al. 2008; Chen et al. 2008; Chen et al. 2009)

Introduction

Social interactions between piglets in period from birth till weaning

- Milk is a limiting resource
- Teat order



(e.g. Hodge, 1974; Hartsock and Graves, 1976; Fraser 1980;
Fraser and Thompson 1986; Harrell et al. 1993;)

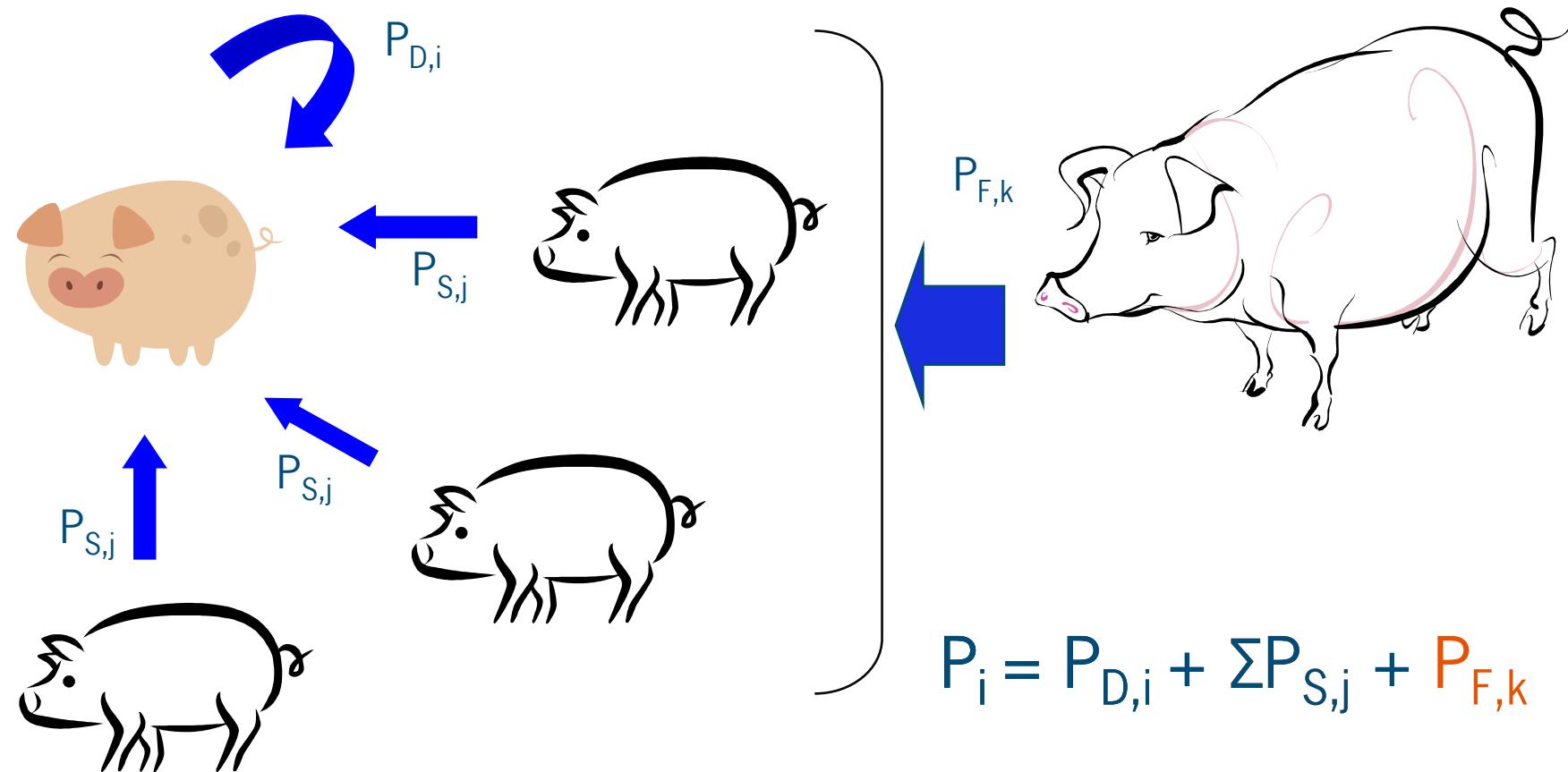
Aim

- Is there heritable social variation in growth of piglets?
- Extend the social interaction model with a maternal effect
- Compare models:
 - basic animal model
 - social model
 - maternal model
 - social-maternal model

Data

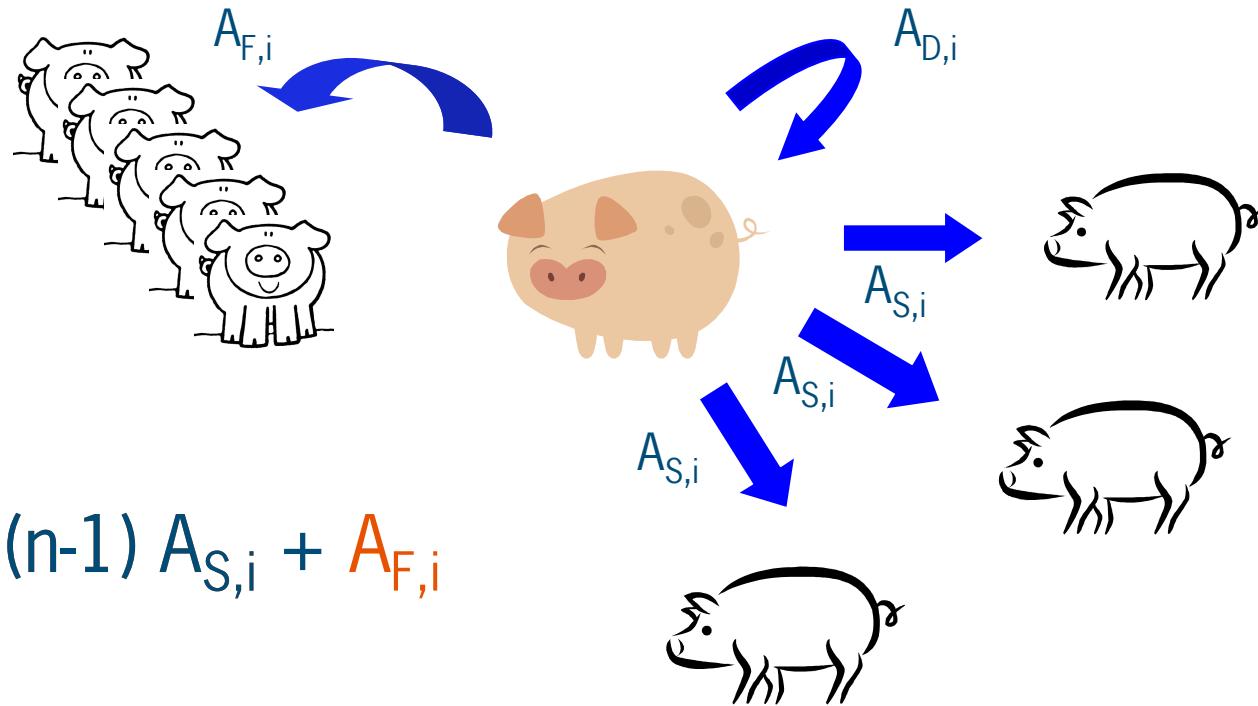
- # piglets: 17,053
- # (foster) groups: 1,604
- Pedigree: 49,645 (5 generations)
- Average group size: 10.8 (range 7-14)
- Cross-fostering %: 28.8%
- Average growth: 251 g/day (sd=61)

Background



Total breeding value

$$TBV_i = A_{D,i} + (n-1) A_{S,i} + A_{F,i}$$



Total heritable impact an individual can have on the population

Models

- 1_{basic}: $\mathbf{y} = \mathbf{Xb} + \mathbf{Z}_D \mathbf{a}_D + \mathbf{Wc} + \mathbf{Vg} + \mathbf{Upe} + \mathbf{e}$
- 2_{social}: $\mathbf{y} = \mathbf{Xb} + \mathbf{Z}_D \mathbf{a}_D + \mathbf{Z}_S \mathbf{a}_S + \mathbf{Wc} + \mathbf{Vg} + \mathbf{Upe} + \mathbf{e}$
- 3_{maternal}: $\mathbf{y} = \mathbf{Xb} + \mathbf{Z}_D \mathbf{a}_D + \mathbf{Z}_F \mathbf{a}_F + \mathbf{Wc} + \mathbf{Vg} + \mathbf{Upe} + \mathbf{e}$
- 4_{social-maternal}: $\mathbf{y} = \mathbf{Xb} + \mathbf{Z}_D \mathbf{a}_D + \mathbf{Z}_F \mathbf{a}_F + \mathbf{Z}_S \mathbf{a}_S + \mathbf{Wc} + \mathbf{Vg} + \mathbf{Upe} + \mathbf{e}$

c: common birth litter = common env. in uterus of biological dam

g: group = common postnatal env. due to being in same (foster) litter

pe: perm. env. from foster dam = 75% fostered 2/more litters

Results

Model	h_D^2	h_S^2	h_F^2	c^2	g^2	pe^2	T^2
1 _{basic}	0.07	-	-	0.05	0.10	0.09	0.07
2 _{social}	0.07	0.0009 _{0.0005}	-	0.05	0.09	0.08	0.11
3 _{maternal}	0.07	-	0.06	0.06	0.09	0.04	0.10
4 _{social-maternal}	0.07	0.0007 _{0.0005}	0.06	0.05	0.09	0.03	0.15

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Model	h_D^2	h_S^2	h_F^2	c^2	g^2	pe^2	T^2
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Results

$$T^2 = \sigma_{TBV}^2 / \sigma_P^2$$

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Results

Log Likelihood

Model	LogL
1 _{basic}	-8600
2 _{social}	-8597
3 _{maternal}	-8592
4 _{social-maternal}	-8590

Likelihood-ratio tests (Chi²- distribution)

Models	P-value
1 _{basic} vs 2 _{social}	0.074
1 _{basic} vs 3 _{maternal}	<0.001
2 _{social} vs 4 _{social-maternal}	0.004
3 _{maternal} vs 4 _{social-maternal}	0.385

Conclusion model 3_{maternal} performs best

Remarks

- Foster dam as maternal effect
 - Both biological and foster dam as genetic effect
 - unsuccessful in convergence → not separable
 - Genetic effect foster dam > genetic effect biological dam

- Non genetic terms
 - Without **common**: **direct** genetic effect increased (+social)
 - Without **group**: genetic **foster + social** effect increased
 - Without **perm. env.**: genetic **foster** effect increased (+social)

Conclusion

- Separation of heritable direct, social & foster effect is possible when substantial cross-fostering is applied
- No evidence that growth in the period from birth till weaning was affected by genetic social interactions
- Maternal model performed best

Acknowledgement



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