

HPA   

Genetic parameters for milk traits using fixed regression models for Pag sheep in Croatia

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Introduction


- Sheep breeding – autohtonous breeds
- Milk production
- Mediterranean area - extensive breeding
- Pag sheep



Pag sheep



- Breeding program since 2000
- Cheese processing
 - Great demand due to tourism
- Aim: Estimate genetic parameters for
 - Daily milk, fat and protein yields
 - Somatic cell score



Milk recording



- ICAR rules
- Croatian Agricultural Agency
- Central Laboratory for Milk Control
- AT4 method




Material

- Test-day records
- Pedigree information
- Central database of Croatian Agricultural Agency
- From January 2003 to March 2010



Data preparation

- Parity (1-6)
- Days in milk (6-180)
- Litter size (1, 2+)
- SCS $\rightarrow \log_{10} \text{SCS} = \log_{10} (\text{SCS}/100.000)$
- Data editing:
 - Animals without birth, lambing, or test-date
 - Age at lambing within parity (12-85)
 - Flock
 - Test-day (less than 10 test-days per flock deleted)



Descriptive statistics

Trait	n	\bar{x}	σ
DMY (kg)	37,916	0.786	0.342
DFY (kg)	37,880	0.058	1.347
DPY (kg)	37,857	0.047	0.020
SCS	34,753	7.08	1.87



Pedigree structure

	No.
Animals with records	4,449
Non base animals:	
- both parents known	1,895
- only sire known	18
- only dam known	288
Base animals (%)	58.2
Total number of animals	5,260



Models

Single-trait fixed regression repeatability test-day model

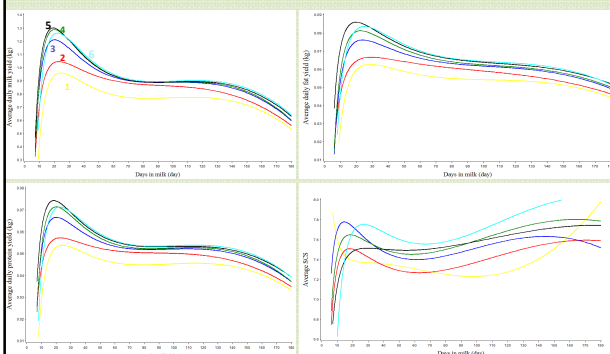
Trait	Fixed effects in the model					
	Parity	Litter size	Lambing season	Flock	Days in milk	Age at lambing
DMY, DFY, DPY	✓	✓	✓	✓	✓	✓
SCS	✓	✓	✓	✓	✓	
	Random effects in the model					
	Flock-test-date	Permanent environment	Additive genetic effect			
DMY, DFY, DPY, SCS	✓	✓	✓			

Method

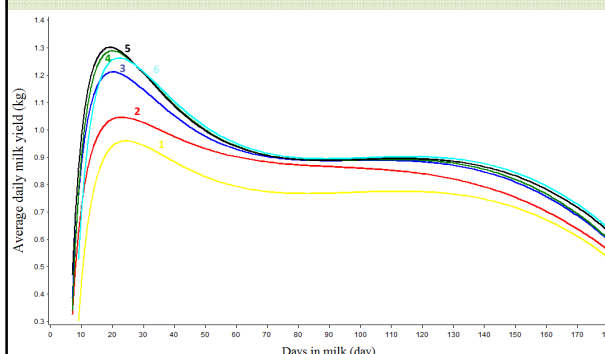
- SAS
 - SQL
 - GLM procedure (SAS/STAT)
 - Data coding
- Variance components estimation - method REML in VCE-6

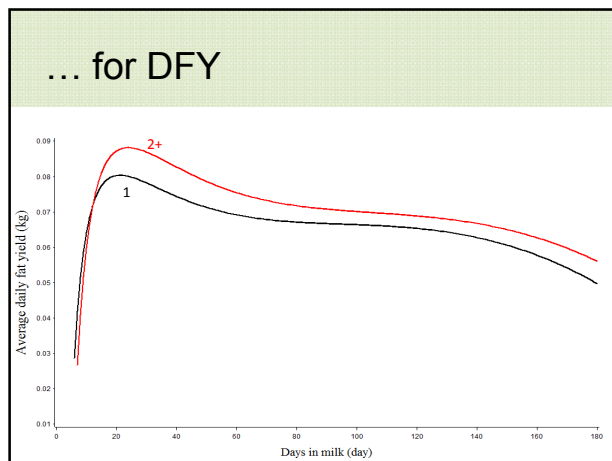
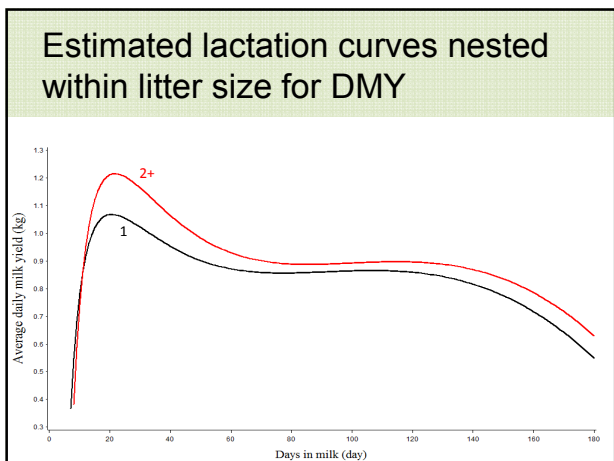
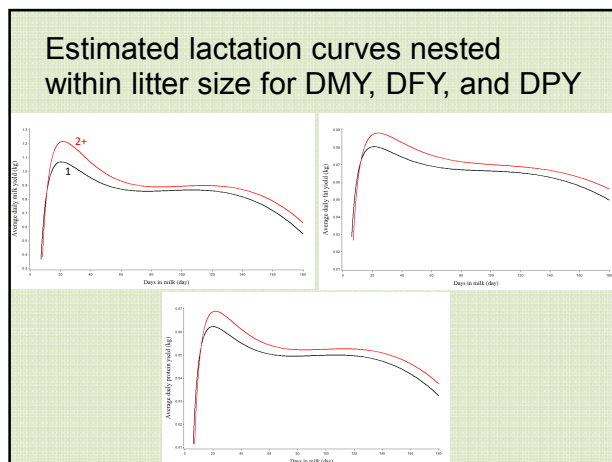
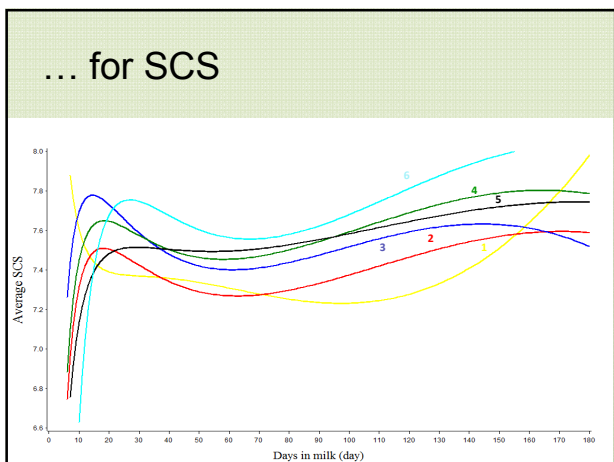
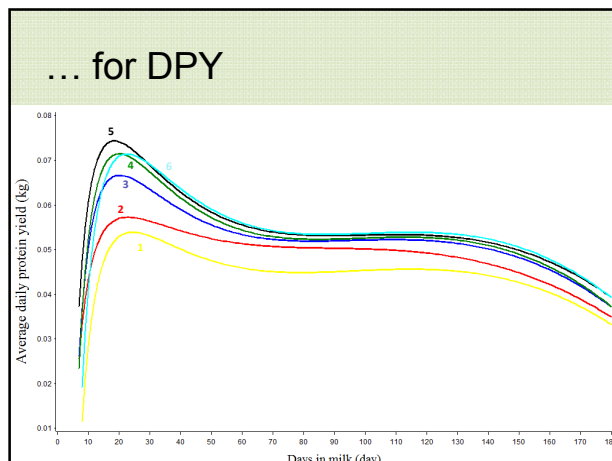
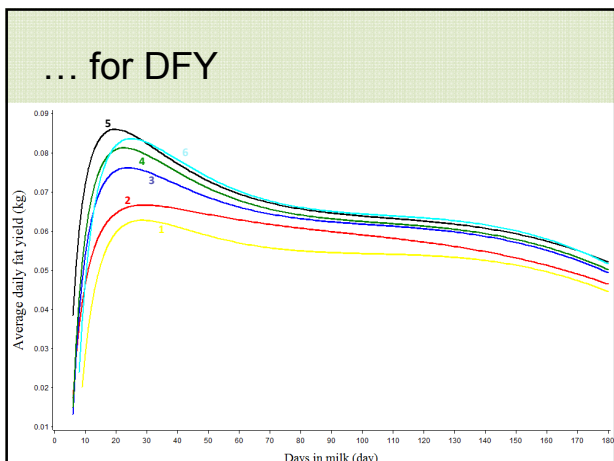


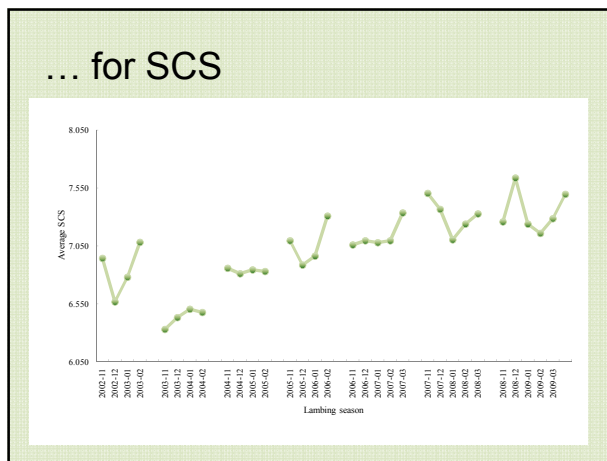
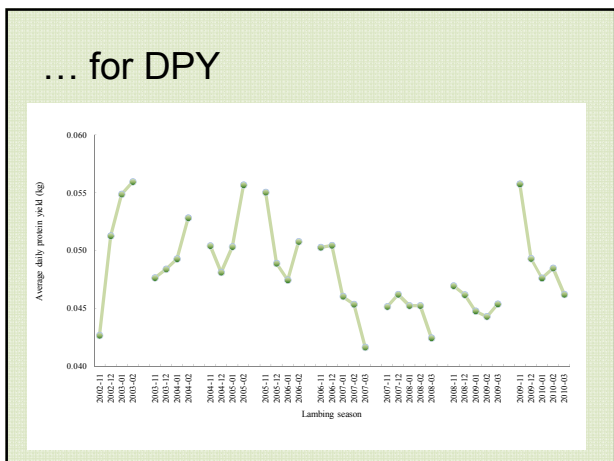
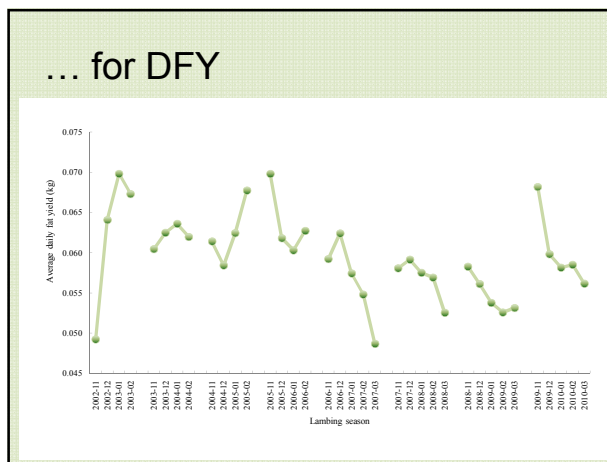
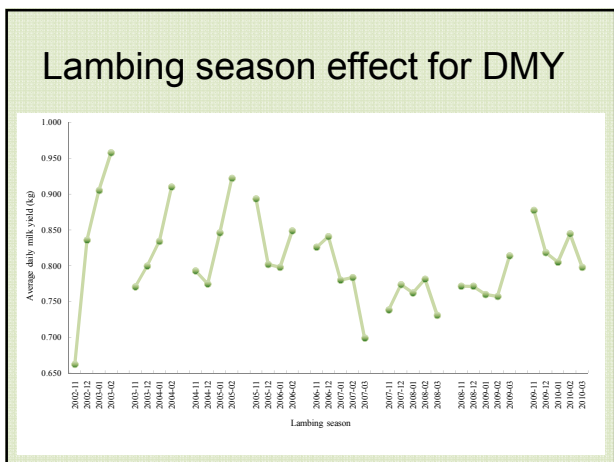
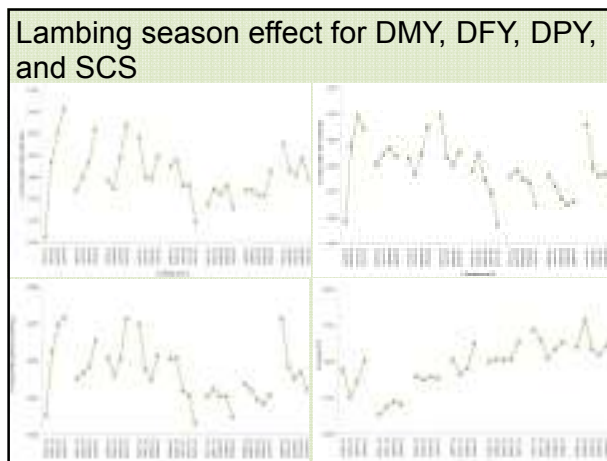
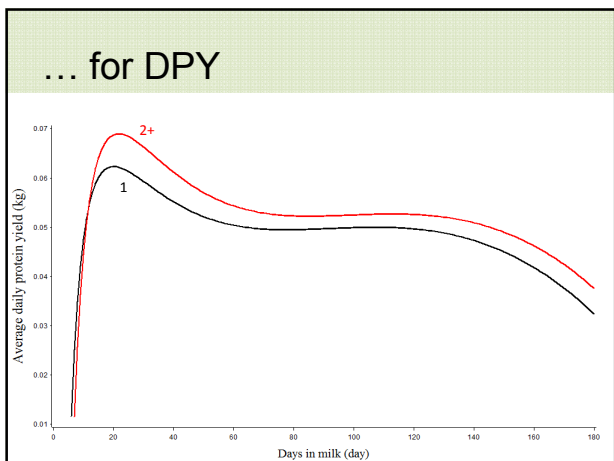
Estimated lactation curves nested within parity for DMY, DFY, DPY, and SCS



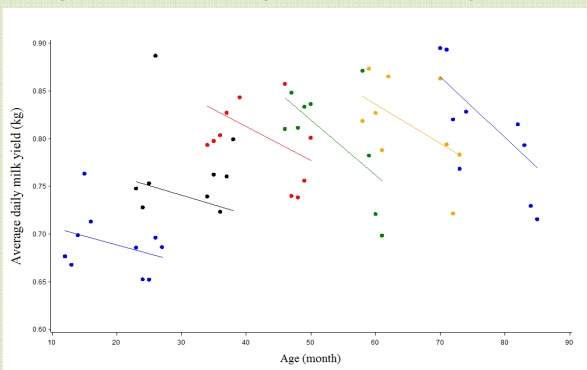
Estimated lactation curves nested within parity for DMY



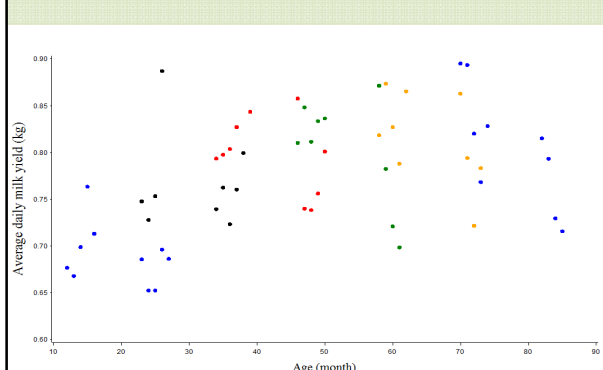




Age at lambing within parity



Age at lambing within parity



Estimated ratios

Trait	σ_a	σ_{hid}	σ_{perm}	σ_{perm}
DMY (kg)	0.26±0.01	0.24±0.01	0.22±0.01	0.30±0.02
DFY (kg)	0.25±0.01	0.16±0.01	0.23±0.01	0.30±0.02
DPY (kg)	0.24±0.01	0.26±0.01	0.21±0.01	0.29±0.02
SCS	0.19±0.01	0.06±0.01	0.19±0.01	0.23±0.02

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Bayesian analysis

- Bayesian analysis (the same priors as in REML)
- Methods
 - Markov chain Monte Carlo (MCMC)
 - Integrated Nested Laplace Approximation (INLA)



Conclusions

- Heritability for milk traits were in expected range
- Results indicate the possibility of using test-day records for genetic evaluation of the Pag sheep in Croatia



Thank you for
attention!

