

# PRELIMINARY RESULTS FROM A GENETIC ANALYSIS OF CLINICAL MASTITIS DATA FOR HOLSTEIN CATTLE IN CZECH REPUBLIC

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# Objective

- analyze data on **clinical mastitis (CM)**
  - Heritability
  - Genetic correlations with production and SCS
- determine their **suitability** for mastitis resistance **breeding value prediction**

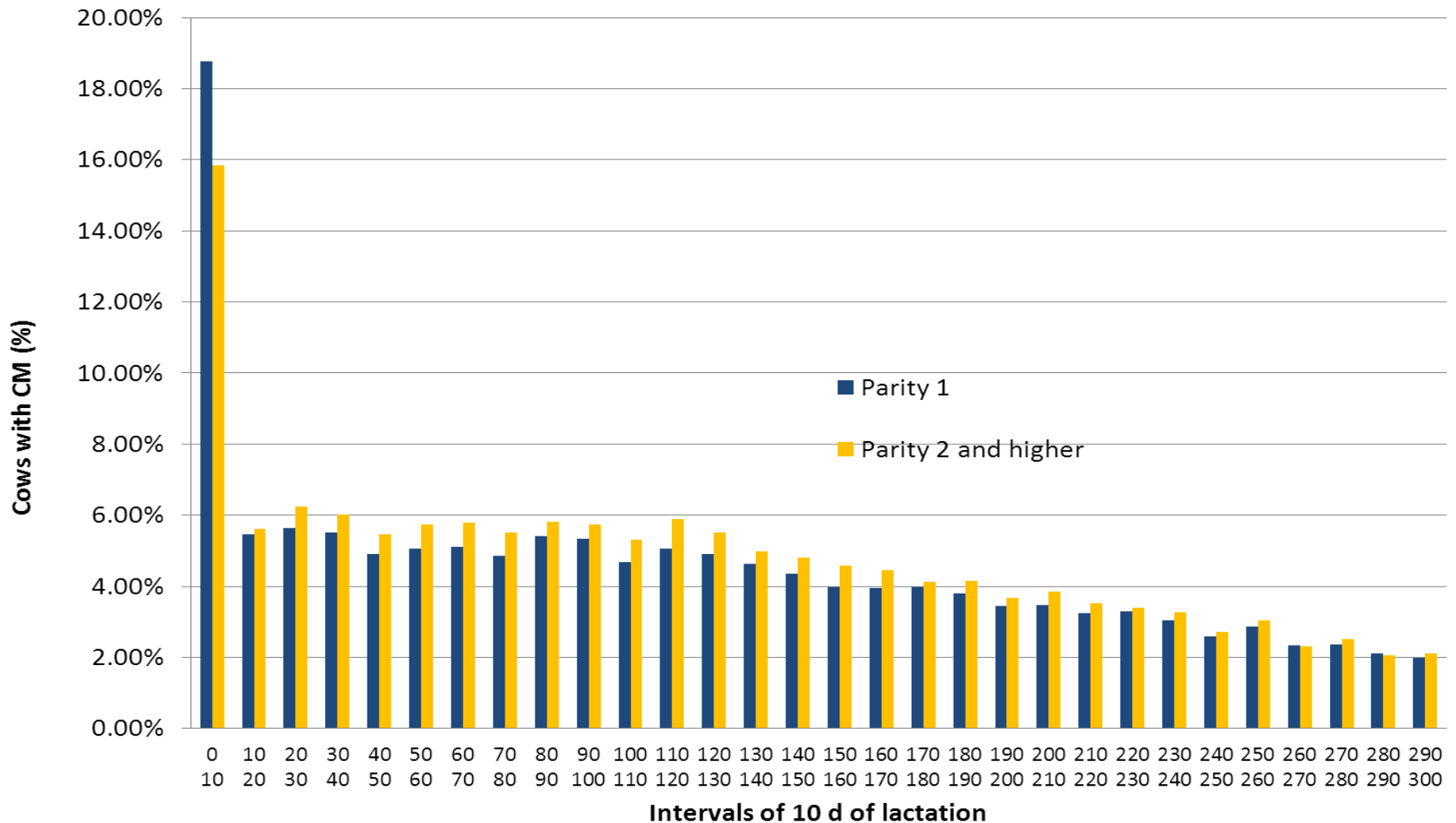
# Animals definition

- ✓ **10 294 cows**
- ✓ **22 812 lactation**
- ✓ **17 606 mastitis cases**
  
- ✓ **7 Holstein herds**
  - 150 (3) 200 (1) 500 (2) 900 (1)**
- ✓ **Time period 2000 and 2012**

# Traits definition

- ✓ **CM1** - the **number of CM cases** per lactation
- ✓ **CM2** - the **number of *days* of CM** per lactation
- ✓ **CM3** - CM considered as an **all-or-none trait** with values of 0 (no CM case) and 1 (at least 1 CM case)
- **305-d milk yield (MY305)**
- **average lactation somatic cell count (SCC)**

# Frequency of cows with clinical mastitis (CM) as a function of the 10 d interval of days of lactation



# Linear animal model

- ✓ Herd – 7
- ✓ Parity – 1, 2, 3,  $\geq 4$
- ✓ Year of calving - 2000-2012
- ✓ Season of calving – 4
  
- ✓ *Permanent environmental effect of cow* 10 294
- ✓ *Additive genetic effect* 25 359

Data were analyzed using the DMU package (Madsen and Jensen, 2010).

# Variance components and heritabilities

	CM1	CM2	CM3
Variances			
Additive	0.19	8.1	0.019
PE	0.13	6.4	0.012
Residual	1.49	68.2	0.193
		Heritability	
	0.10	0.10	0.09

**CM1 – no CM cases**

**CM2 – days of CM**

**CM3 – 0 or 1**

# Variance components and heritabilities

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# Additive genetic correlations

Trait	CM1	CM2	CM3
CM1		0.90 **	0.97 **
CM2			0.93 **
MY(305)	0.23 **	0.24 **	0.29 **
SCS(305)	0.80 **	0.79 **	0.83 **

**CM1 – no CM cases**

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# Breeding value and ranking of sires

- ✓ equal equation
- ✓ linear animal model
- ✓ the same dataset
  
- ✓ univariate models
  - CM1, CM2, CM3 and SCS(305)
- ✓ bivariate models
  - CM trait and SCS(305)

# Breeding value and ranking of sires

- ✓ Spearman **rank correlations** between breeding values
- ✓ for **139** sires
- ✓ with reliability of breeding value over 50% for all estimated breeding values.

# Spearman rank correlations between breeding values

CM1	x	CM2	0.87
CM1	x	CM3	0.84
CM2	x	CM3	0.81
CM1scs	x	CM2scs	0.92
CM1scs	x	CM3scs	0.88
CM2scs	x	CM3scs	0.88

**CM1 – no CM cases**

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**CM1 – no CM cases**

**CM2 – days of CM**

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# Spearman rank correlations between breeding values

CM1	x	SCS(305)	0.53
CM2	x	SCS(305)	0.55
CM3	x	SCS(305)	0.59
CM1scs	x	SCS(305)	0.70
CM2scs	x	SCS(305)	0.71
CM3scs	x	SCS(305)	0.77

**CM1 – no CM cases**

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CM3scs	x	SCS(305)	0.77

**CM1 – no CM cases**

**CM2 – days of CM**

**CM3 – 0 or 1**



# Range of breeding values

	Absolute value	SD
CM1	0.34	4.99
CM2	2.13 d	5.85
CM3	0.10	4.95
CM1scs	0.34	5.04
CM2scs	2.14 d	5.64
CM3scs	0.10	5.11

**CM1 – no CM cases**

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**CM1 – no CM cases**

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# Conclusion

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- ✓ suitable for genetic evaluation
- ✓ linear animal model including the permanent environmental effect of the cow
- ✓ Traits
  - CM1 – no. CM cases
  - CM2 – days of CM
  - CM3 – 0 or 1

# Acknowledgments

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