



### Aims

Setup a routine genetic and genomic evaluation of gestation length (GL) Variance components estimation (VCE) for direct and maternal GL Investigation on the genetic relationship between GL and other economically important traits

	dGL1	dGL2+	mGL1
dGL1	0.43 (0.005)	0.92	-0.50
dGL2+		0.35 (0.004)	-0.29
mGL1			0.08 (0.003)
mGL2+			

Heritability on diagonal, genetic correlations above diagonal, posterior standard deviations in brackets. **d** = direct effect; **m** = maternal effect; **1** = first parities; **2+** = later parities.



Gestation length is suitable for genetic selection in both its components, direct and maternal. The identified statistical model is stable and suitable for routine genetic evaluation. A deeper analysis on the possible consequences of selecting for GL in Italian holstein is in progress.

# **Genetic evaluation of gestation length in Italian Holstein breed**

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**DATA:** 5,607,119 calvings from 2004 onwards; GL between 260-305 days; minimum observations per contemporary group: 10. Genetic evaluation: Multiple-trait (first and later parities) repeatability linear animal model. **Model validation:** LR method on first crop bulls.



# Conclusions

<u></u>-0,6

-0,8

Nilt

## Materials and methods

Correlation

Reliability

**0.957**(0.953,0.960) **0.944**(0.941,0.946) **0.617**(0.606,0.629)

**0.003**(-0.006,0.011) **0.902**(0.894,0.910) **0.784**(0.777,0.790) **0.186**(0.182,0.191)

LR validation results: bootstrap applied to 31 frist crop bulls. Confidence intervals in brackets.



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