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# Heterogeneity of residual variances of milk fatty acids in dairy cattle

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# Collaborations

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## ■ CRA-W

– Frédéric Dehareng – Pierre Dardenne



## ■ Comité du Lait

– Didier Veselko – Emile Piraux



## ■ AWE

– Carlo Bertozzi – Laurent Laloux



# Context

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- Routine genetic evaluation for milk fatty acids is under development in the Walloon Region of Belgium
- The model used in genetic evaluations is a multi-trait random regression test-day model (RRTDM) (Auvray and Gengler, 2002)

## ■ Genetic evaluation model

$$y = X_t t + Q(W_t h + Z_t g + Z_t p) + e$$

### Fixed effects :

- herd \* test date
- stage of lactation
- gestation stage
- lactation stage\*calving season\*breed\*calving period regressed on age at calving

## ■ Genetic evaluation model

$$y = X_t t + Q(W_t h - Z_t g - Z_t p) + e$$

### Random regression effects :

- herd  $\times$  period of calving
- additive genetic
- permanent environmental

### Random residuals

## ■ Genetic evaluation model

$$y = X_t t + Q(W_t h + Z_t g + Z_t p) + e$$

### Matrices :

- **X, W, Z** are incidence matrices,
- **Q** is the covariate matrix for the second order Legendre polynomials.

# Context

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- ***Accuracy of estimated breeding values is one major component in designing breeding programs***
- In mixed models, it is often assumed that the residual variance is the same for all observations (Rönnegård et al., 2010)
- BUT: variation of the residual variances seems to be quite common
- BUT: Assuming a homogeneous residual variance could affect the genetic evaluation (Takma, 2009)
- → so, it could be important to include the heterogeneity of residual variance in the used model

# General objective

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*To test the heterogeneity of residual variances for*

- Milk, fat and protein yields
- Monounsaturated and saturated fatty acids

→ Indirectly, to study the goodness of fit of the model (average residuals)

# Materials & Methods

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## ■ Dataset

–First lactation

–Traits involved in the genetic evaluation :

Milk, fat and protein yields (kg/day)

(Milk, qFAT and qPROT)

→ +/- 6,687,000 records

–Traits not involved in the genetic evaluation : content of saturated and monounsaturated fatty acids in milk (g/dl of milk)

(SAT and MONO)

→ +/- 220,000 records

# Materials & Methods

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- Residuals:

- Observed values – Estimated values

- Squared residuals  $\approx$  residual variance

- Studied effects are:

- The month of test date
  - The calving month
  - The year of test date
  - The age at calving
  - And the stage of lactation

# Studied traits

Trait	N	Mean	SD
<b>MILK (kg/day)</b>	6,749,239	16.96	6.83
<b>qFAT (kg/day)</b>	6,746,993	0.68	0.29
<b>qPROT (kg/day)</b>	6,727,524	0.56	0.22
<b>SAT (%)</b>	220,397	2.79	0.49
<b>MONO (%)</b>	220,396	1.15	0.24

# Residuals

Trait	N	Mean	SD
MILK (kg/day)	6,749,239	0.00	1.35
qFAT (kg/day)	6,746,993	0.00	0.07
qPROT (kg/day)	6,727,524	0.00	0.05
SAT (%)	220,397	0.00	0.17
MONO (%)	220,396	0.00	0.09

# Results --- Squared residuals

Trait	N	Mean	SD
<b>MILK (kg<sup>2</sup>/day<sup>2</sup>)</b>	6,749,239	1.821	9.774
<b>qFAT (kg<sup>2</sup>/day<sup>2</sup>)</b>	6,746,993	0.0051	0.0208
<b>qPROT (kg<sup>2</sup>/day<sup>2</sup>)</b>	6,727,524	0.0023	0.0120
<b>SAT (%)<sup>2</sup></b>	220,397	0.0276	0.0615
<b>MONO (%)<sup>2</sup></b>	220,396	0.0089	0.0218

# Results --- Month of test-date

P value	Month of test date
Milk	***
qFAT	***
qPROT	***
SAT	***
MONO	***

The effect of month of test date on squared residuals is highly significant for all traits

# Results --- Month of test-date

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- Peak during the spring (April → June)
- Lowest values during the summer
- Similar trend for milk, fat and protein yields and for SAT **vs** MONO (smoother)

# Results --- Calving month

P value	Calving month
Milk	***
qFAT	***
qPROT	***
SAT	***
MONO	***

The effect of **calving month** on squared residuals is highly significant for all traits

# Results --- Month of test-date

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- Peak in August
- Lowest values during the spring (April → June)
  - Vs Month of calving effect
- Similar trend for milk, fat and protein yields **vs** for SAT and MONO (smoother)

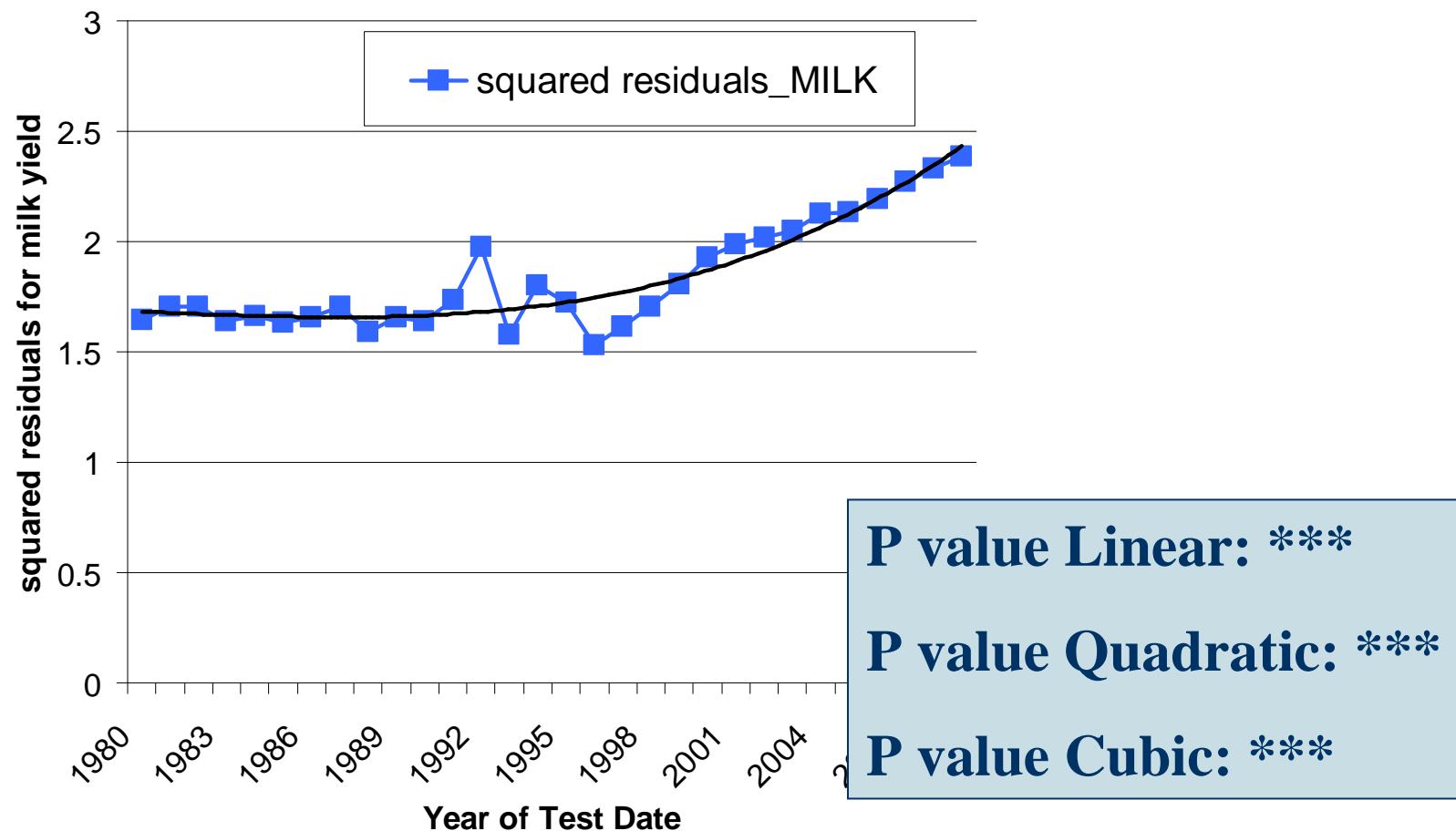
# Results --- Year of test date

P value	Year of test date
Milk	***
qFAT	***
qPROT	***

The effect of **year of test date** on squared residuals is highly significant for milk, fat and protein yields

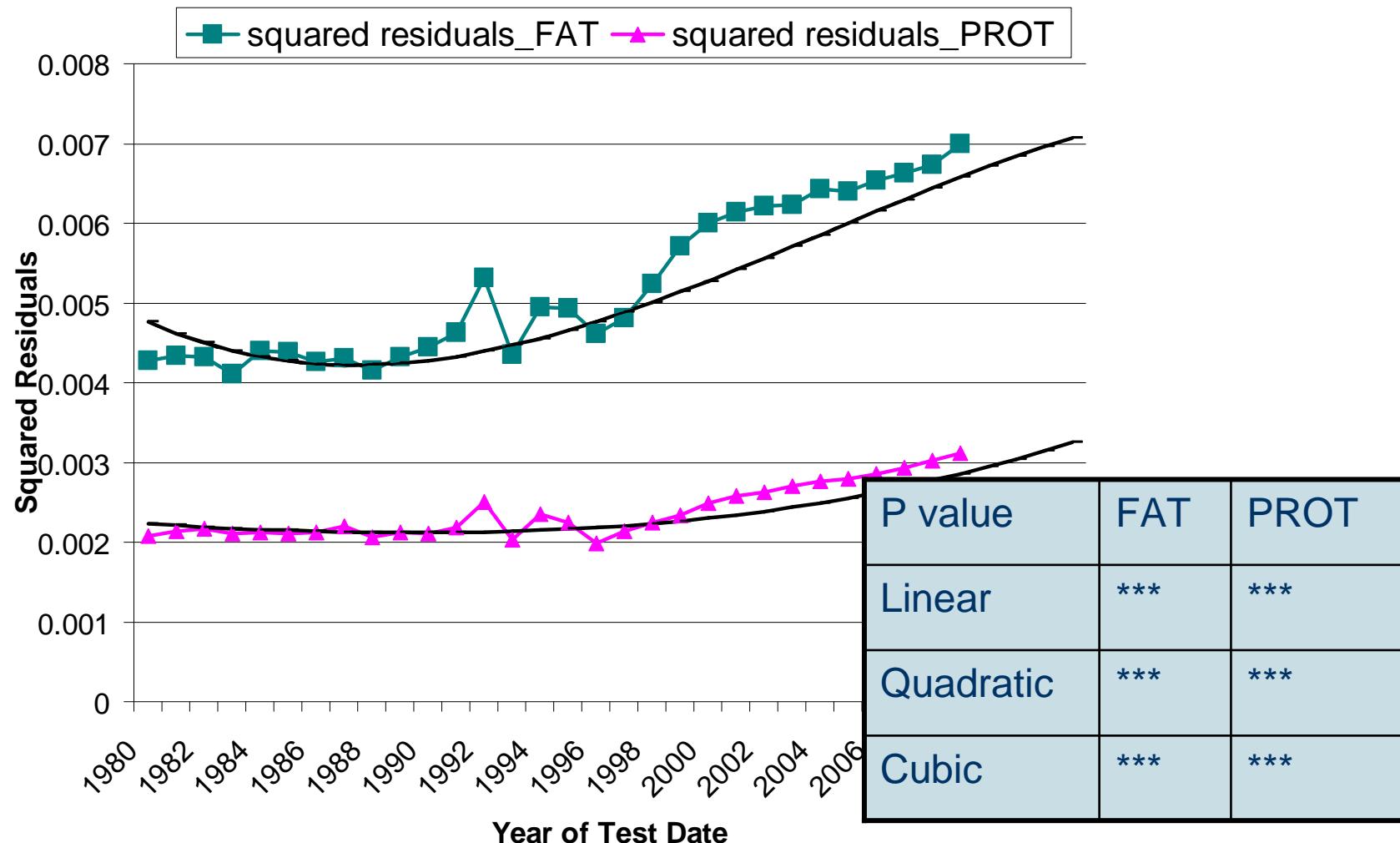
# Results --- Year of test date

Trend of squared residuals for milk yield according to year of test date



# Results --- Year of test date

Trend of squared residuals for fat and protein yields according to year of test date



# Results --- Age at calving

P value	Age at calving
Milk	NS
qFAT	NS
qPROT	NS
SAT	NS
MONO	NS

The effect of **age at calving** on squared residuals is not significant for any traits

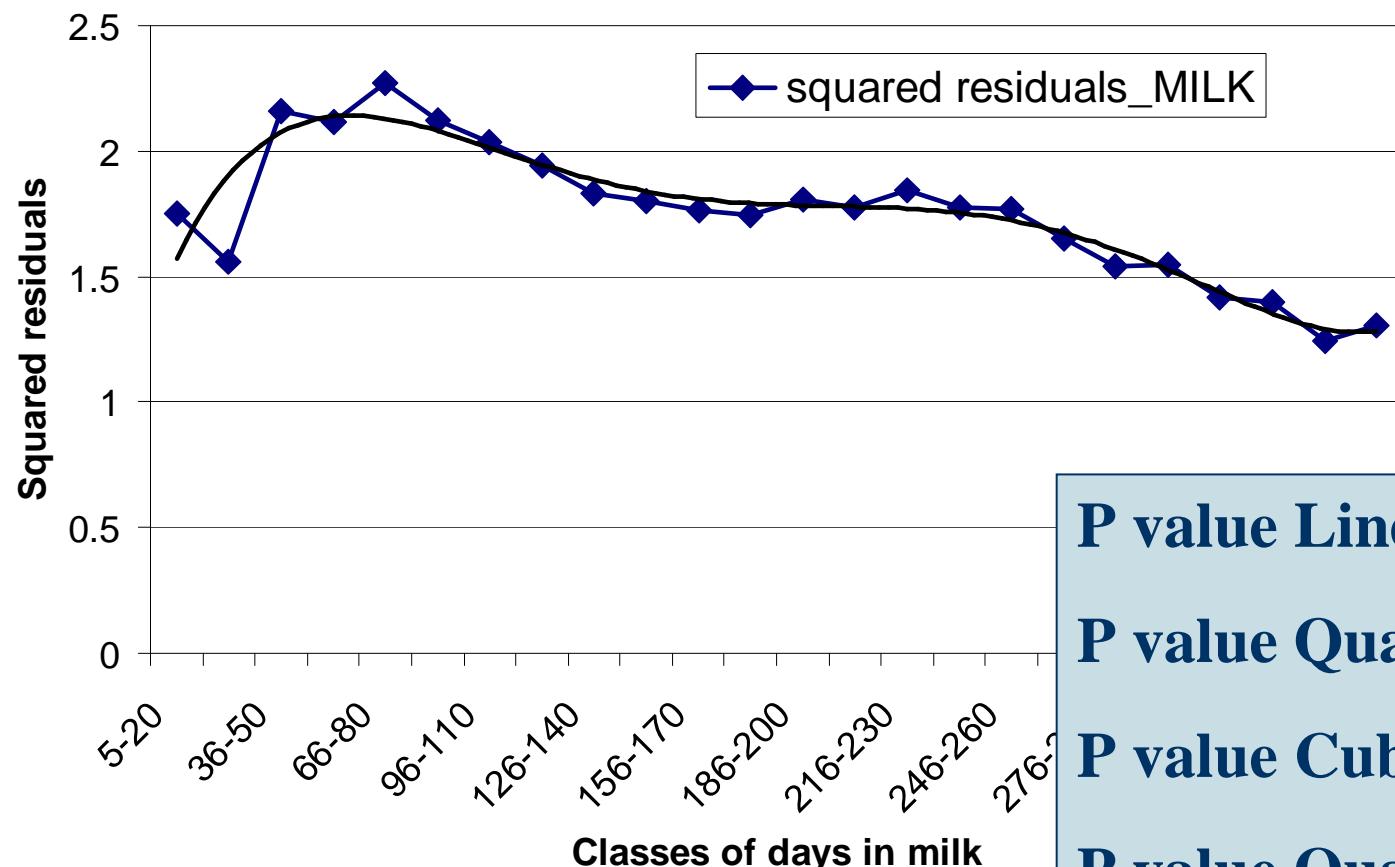
# Results --- Stage of lactation

P value	DIM
Milk	***
qFAT	***
qPROT	***
SAT	***
MONO	***

The effect of **stage of lactation** on squared residuals is highly significant for all traits

# Results --- Stage of lactation

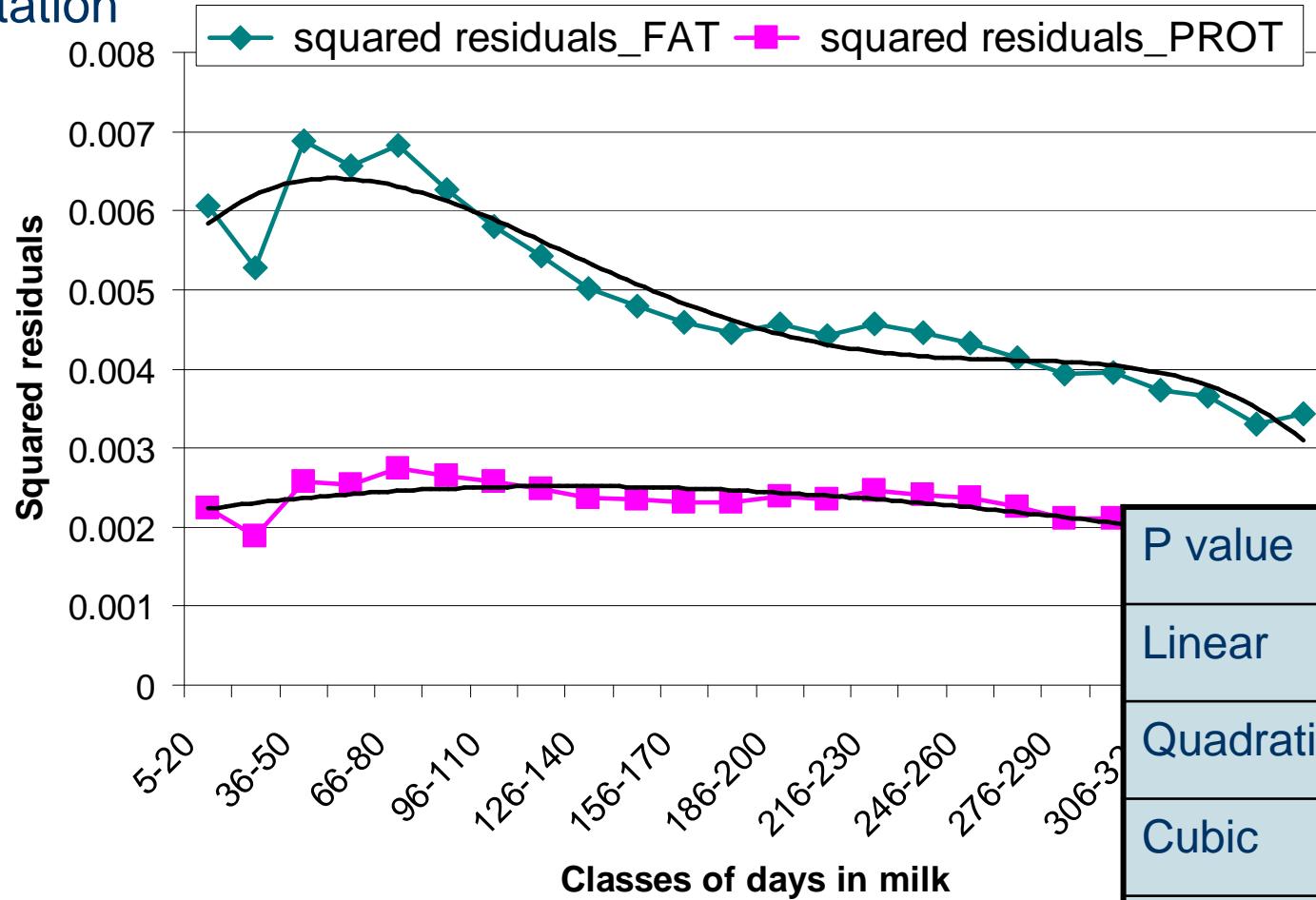
Trend of squared residuals for milk yield according to the stage of lactation



**P value Linear: \*\*\***  
**P value Quadratic: \*\*\***  
**P value Cubic: \*\*\***  
**P value Quartic: \*\*\***

# Results --- Stage of lactation

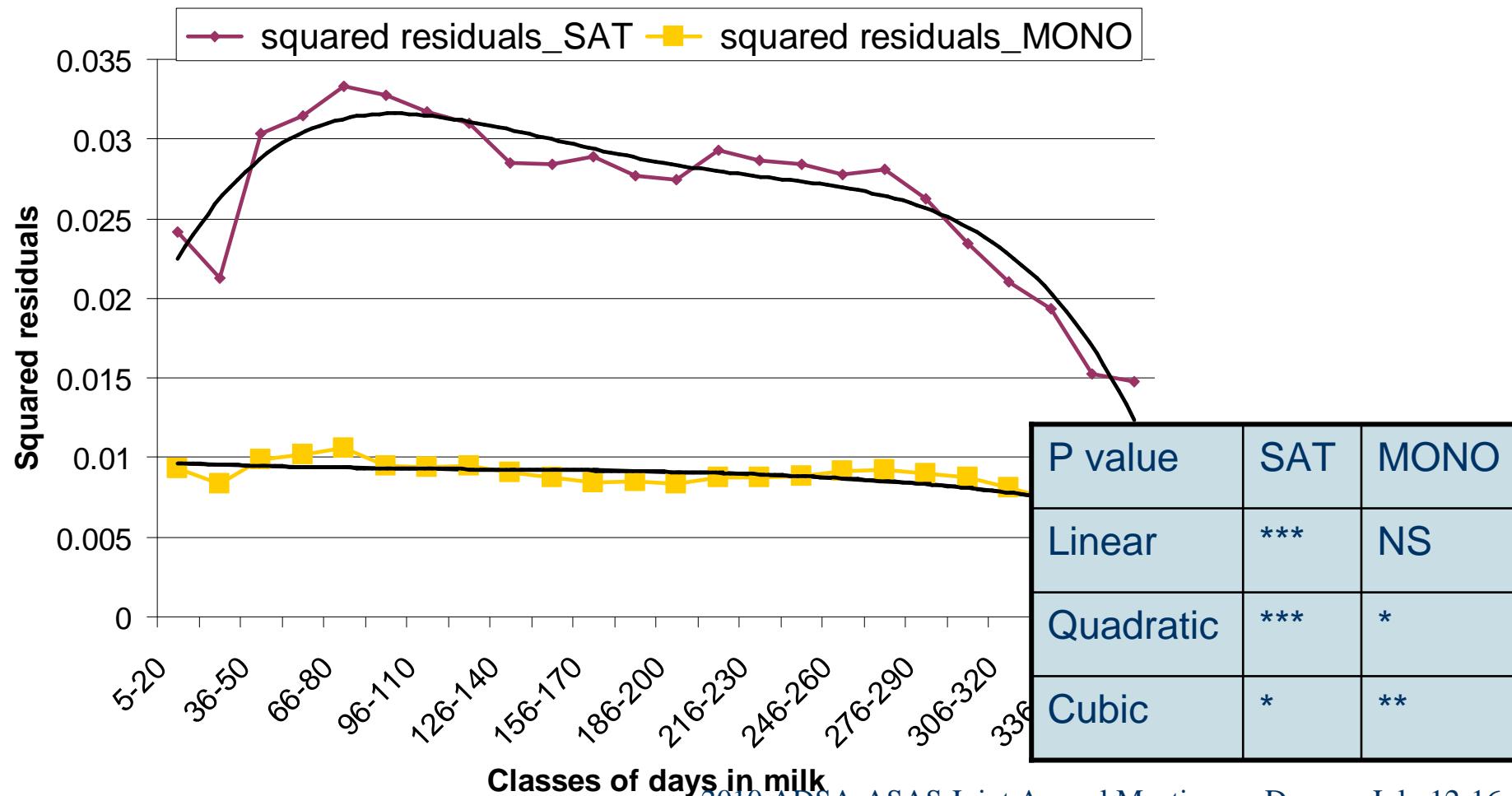
Trend of squared residuals for fat and protein yields according to the stage of lactation



P value	FAT	PROT
Linear	***	***
Quadratic	***	***
Cubic	***	***
Quartic	***	***

# Results --- Stage of lactation

Trend of squared residuals for SAT and MONO according to the stage of lactation



# Conclusions

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- Means of residuals were stable and close to zero for all traits
- Trends of squared residuals: differences between milk, fat and protein yields and SAT and MONO
- → Introduction of heterogeneous residual variance could be interesting for the accurate model definition

# Conclusions

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- How to introduce this heterogeneity?
- Suggestion : introduction of correction of variability of squared variances according to stages of lactation

# Acknowledgment

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