



TOWARDS IMPROVEMENT OF RUMINANT BREEDING
THROUGH GENOMIC AND EPIGENOMIC APPROACHES

Meta-analysis for heat stress tolerance traits in Holstein in France, the Netherlands and Spain

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RUMIGEN – Introduction

- In the context of climate change, animals will be more and more exposed to heat-stress conditions
- Challenge : ensure that animals maintain a good production and remain healthy
- **European project RUMIGEN**
Evaluate the impact of heat-stress conditions on performances and estimate the genetic variability for heat-stress tolerance in Holstein dairy cattle breed
 - **Objective : Meta-analysis to estimate genetic correlations between countries for heat stress tolerance traits in Holstein**

RUMIGEN – Datasets

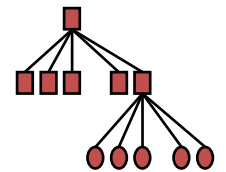
- **Countries:** the Netherlands (WUR), Spain (INIA, IRIAF), France (INRAE, IDELE)
- **Breed:** Holstein – common to all 3 countries



- **Existing large scale national datasets:**

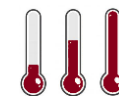
- **Phenotypes** : national datasets for

- production : milk yield (MY), fat yield (FY), protein yield (PY)
 - udder health : somatic cell score (SCS)



- **Pedigrees**

- **Weather** : daily records provided by national Weather Agencies associated to each herd through zip code





RUMIGEN – Heat stress indicator

- **Temperature-Humidity Index (THI)**

calculated from the average daily temperature (T in °C) and relative humidity (RH)

$$\text{THI} = (1.8 * T + 32) - (0.55 - 0.0055 * RH) * (1.8 * T - 26)$$

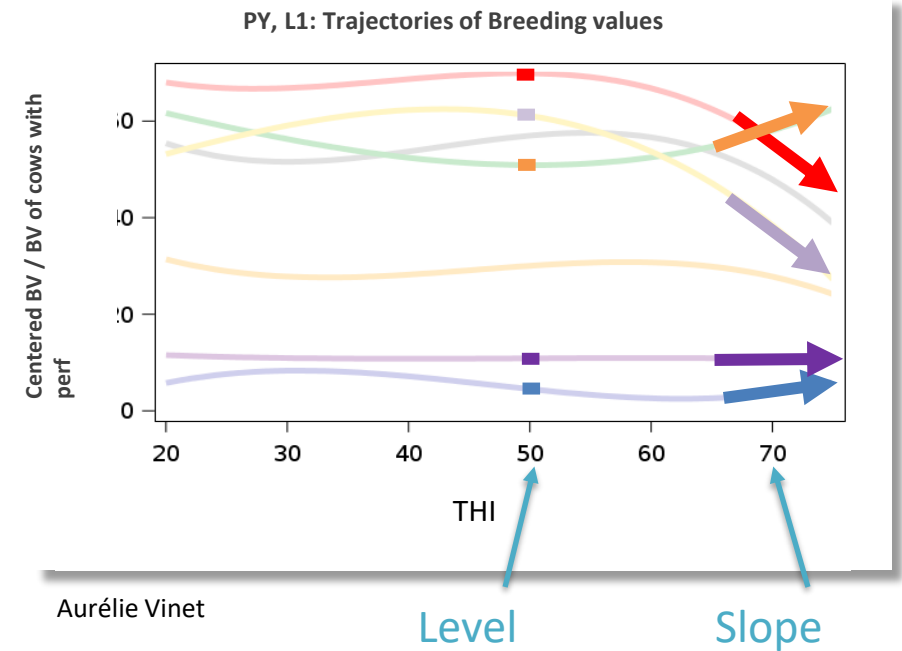
(National Research Council. A guide to environmental research on animals. Washington: National Academy of Sciences; 1971.)

- **Heat stress indicator**

Average THI over 3 days: 2 days before test day + test day

RUMIGEN - Traits definition

- At the **country level**, **variance components** and **breeding values** were estimated:
 - under **thermo-neutral conditions**
 - **level of production** at DIM 150 and **THI 50**
 - under **heat-stress conditions**
 - **level of production** at a given THI
 - **slope** using the **first derivative** at a given THI



RUMIGEN - Traits definition

- **Under heat-stress, traits were defined at different THI in each country**

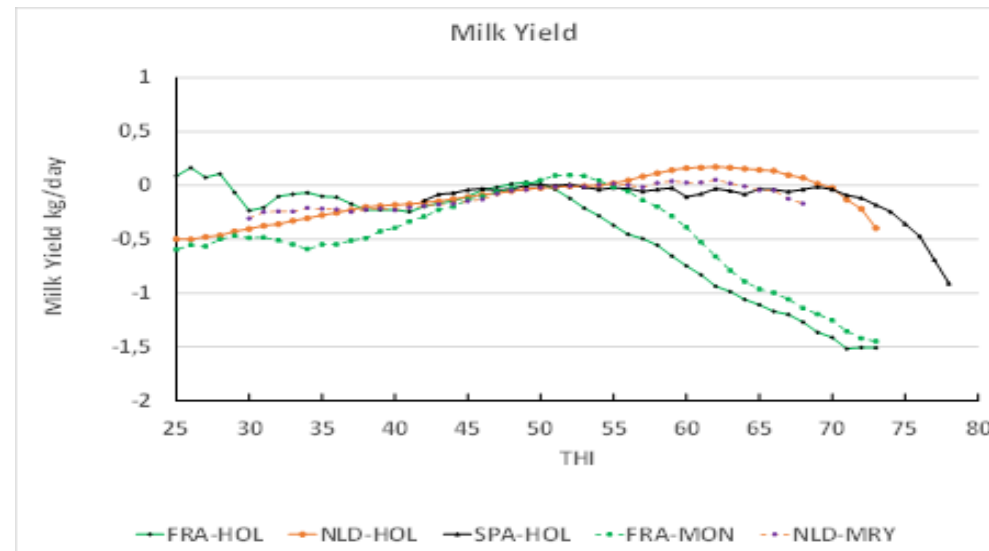
Different break points at the population level in each country

→ Adapt to the situation of each country

FRA : THI 65

NLD : THI 68

SPA : THI 77





RUMIGEN - Traits definition

- Under heat-stress, traits were different THI in each country

Different break

→ Adap

FRA :

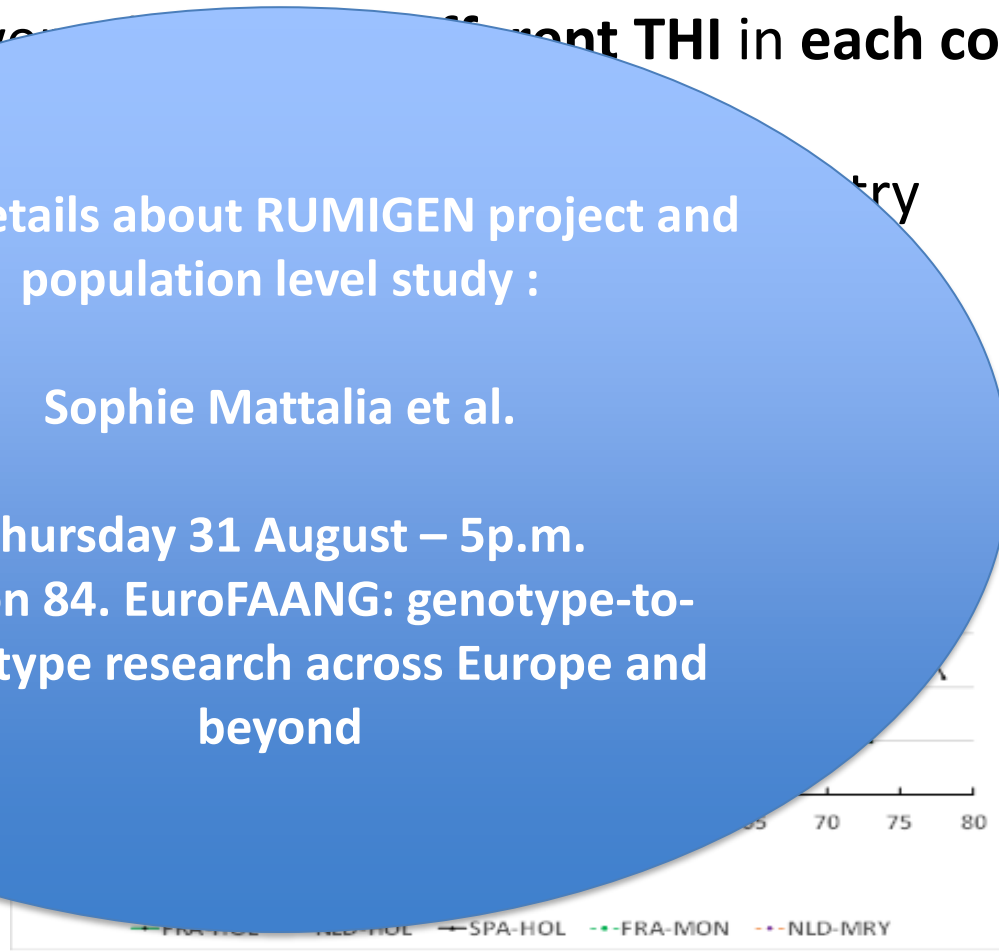
NLD :

SPA : THI

More details about RUMIGEN project and population level study :

Sophie Mattalia et al.

Thursday 31 August – 5p.m.
 Session 84. EuroFAANG: genotype-to-phenotype research across Europe and beyond



RUMIGEN – Genetic evaluation

- Estimation of the effect of THI at the individual level:

Reaction norm models

$$y = \sum \text{fixed effects} + f(\mathbf{GxTHI}) + f(\mathbf{pxTHI}) + e$$

- y: performances
- f(GxTHI): random additive genetic effects - Legendre polynomials
- f(pxTHI) : random permanent environment effect
- e: residual

➔ **Models specific to each trait and harmonised as far as possible between countries**

- **Genetic evaluation:**

Each partner provided :

- **EBVs** and **reliabilities** estimated for sires with at least 20 daughters
- Pedigree file

RUMIGEN - Meta-analysis methodology

- **Deregression:**

Estimation of **deregressed proofs** (phenotypes) and **equivalent record contributions** (weights)

- **Single trait animal model** – deregression by country and by trait
- **Heritabilities:**
 - **estimated** for the **levels**
 - **assumed** to be equal to **0.10** for MY, FY, PY and **0.03** for SCS slopes
- **Minimum reliability of 0.25**

- **Genetic correlations among countries :**

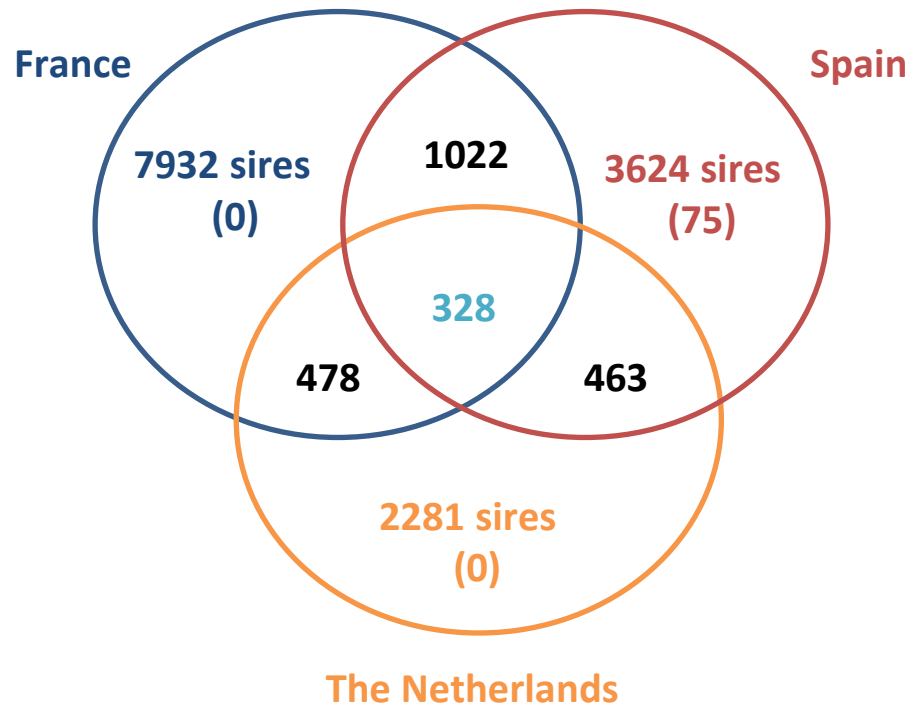
Similar to MACE approach

Multiple trait model: $\mathbf{y} = \text{country} + \text{bull genetic effect} + \mathbf{e}$

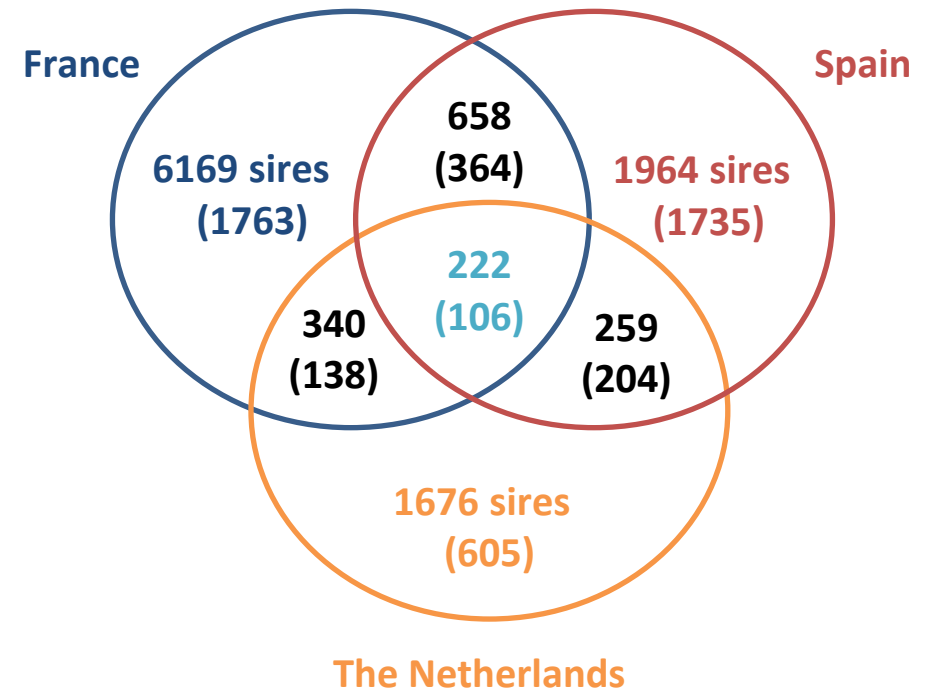
Blupf90 (Misztal *et al.*)

RUMIGEN - Common sires to all three countries

- Milk productions traits - Levels



- Slopes for SCS



() : number of sires deleted of the dataset because of low reliability

- Elimination of sires with **low reliability** had **more impact on slopes** than levels

RUMIGEN - Genetic correlations for the levels under thermo-neutral conditions

- Levels at THI 50

Milk yield	FRA	NLD	SPA
FRA	0.18 ± 0.01	0.94 ± 0.02	0.96 ± 0.01
NLD		0.25 ± 0.02	0.92 ± 0.02
SPA			0.19 ± 0.01

SCS	FRA	NLD	SPA
FRA	0.34 ± 0.01	0.88 ± 0.02	0.95 ± 0.02
NLD		0.15 ± 0.01	0.89 ± 0.03
SPA			0.08 ± 0.01

Fat yield	FRA	NLD	SPA
FRA	0.25 ± 0.01	0.90 ± 0.02	0.97 ± 0.01
NLD		0.21 ± 0.01	0.89 ± 0.02
SPA			0.13 ± 0.01

Protein Yield	FRA	NLD	SPA
FRA	0.15 ± 0.01	0.89 ± 0.02	0.96 ± 0.01
NLD		0.20 ± 0.01	0.90 ± 0.02
SPA			0.12 ± 0.01

- Heritabilities were mostly consistent with the national estimations.
- Genetic correlations** between countries estimated for the **levels of production at THI 50** were **high and consistent with Interbull estimates**.

RUMIGEN - Genetic correlations for the levels under heat-stress conditions

FRA : THI65
 NLD : THI68
 SPA : THI77

- Levels at high THI

Milk yield	FRA	NLD	SPA
FRA	0.18 ± 0.01	0.89 ± 0.02	0.92 ± 0.01
NLD		0.25 ± 0.02	0.86 ± 0.02
SPA			0.25 ± 0.01

SCS	FRA	NLD	SPA
FRA	0.30 ± 0.01	0.88 ± 0.02	0.96 ± 0.02
NLD		0.16 ± 0.01	0.88 ± 0.03
SPA			0.09 ± 0.01

Fat yield	FRA	NLD	SPA
FRA	0.25 ± 0.01	0.87 ± 0.03	0.97 ± 0.02
NLD		0.20 ± 0.02	0.85 ± 0.04
SPA			0.13 ± 0.01

Protein yield	FRA	NLD	SPA
FRA	0.17 ± 0.01	0.81 ± 0.02	0.89 ± 0.01
NLD		0.21 ± 0.01	0.83 ± 0.02
SPA			0.17 ± 0.01

- Genetic correlations between countries estimated for the levels of production at high THI were high...

RUMIGEN – Comparison of genetic correlations under thermo-neutral conditions and under heat-stress conditions

- Levels at THI 50

Milk yield	FRA	NLD	SPA
FRA	0.18 ± 0.01	0.94 ± 0.02	0.96 ± 0.01
NLD		0.25 ± 0.02	0.92 ± 0.02
SPA			0.19 ± 0.01

- Levels at high THI

Milk yield	FRA	NLD	SPA
FRA	0.18 ± 0.01	0.89 ± 0.02	0.92 ± 0.01
NLD		0.25 ± 0.02	0.86 ± 0.02
SPA			0.25 ± 0.01

- ... even though they were lower than at THI 50

RUMIGEN - Genetic correlations for the slopes under heat-stress conditions

Example of milk yield

- **Inconsistent genetic variances from one country to another**
 - France : 0.5 ± 0.01
 - The Netherlands : 2.7 ± 0.05
 - Spain : 0.01 ± 0.001
- **Different variance estimates at different THI**
 - Residual variances of 4.9 ± 0.17 at THI68 vs 8.6 ± 0.66 at THI70 for the Netherlands
- **h^2 estimates much higher than expected (expected value ≈ 0.10)**
 - France : 0.21 ± 0.01
 - The Netherlands : 0.44 ± 0.06
 - Spain : 0.09 ± 0.01

→ **Unreliable results**

RUMIGEN – Discussion

- **Levels:**

- Our approach **was validated**
- **High genetic correlations** were found between the 3 countries
 - **Consistent with Interbull references at THI50**
 - **Lower correlations under heat stress** than under thermo-neutral conditions

- **Slopes:**

Difficulties in correctly estimating variances

- Estimated as **derivative at a given THI** → sensitivity to potential error

- **Deregression**

Few data at high THI on which to base estimates of the slopes

→ **indirect prediction** of the slopes based on the performances at low THI

It leads to several issues for the deregression :

- low reliabilities
- the hypothesis of a single trait model for the deregression is probably too strong

- **Slopes:**
 - give an indication of the **animals' ability to maintain production despite high temperatures**

BUT

 - were **difficult to use in practice** with our dataset (few data at high THI)
- **Levels:**
 - give an indication of **the level of production we can expect in the future**
 - **there is an interest in working on levels at high THI**
 - there is an interest in developing **collaborations** between countries that are already facing high temperatures and countries that will face high temperatures in the future

RUMIGEN PARTNERS



Thank you for your attention

www.rumigen.eu



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