

## Genetic selection as effective methane mitigation tool in livestock

Symposium on behalf of the PhD defence of Anouk van Breukelen

# Genomic and metagenomic tools for climate change mitigation in Spanish dairy cattle



Óscar Gonzalez Recio  
(INIA-CSIC, Madrid, Spain)



# Solutions to achieve net zero carbon emissions



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## COP26: US and EU announce global pledge to slash methane

2 November 2021

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Energy & Climate / Gas / The COP26 methane moment

**GAS**

### The COP26 methane moment

This is the second installment of the Topic of the Month: Decarbonising Gas Markets

European Commission English EN Search

Home > Press corner > Launch by US, EU and Partners of the Global Methane Pledge

Available languages: English

Statement | 2 November 2021 | Brussels

### Launch by United States, the European Union, and Partners of the Global Methane Pledge to Keep 1.5C Within Reach

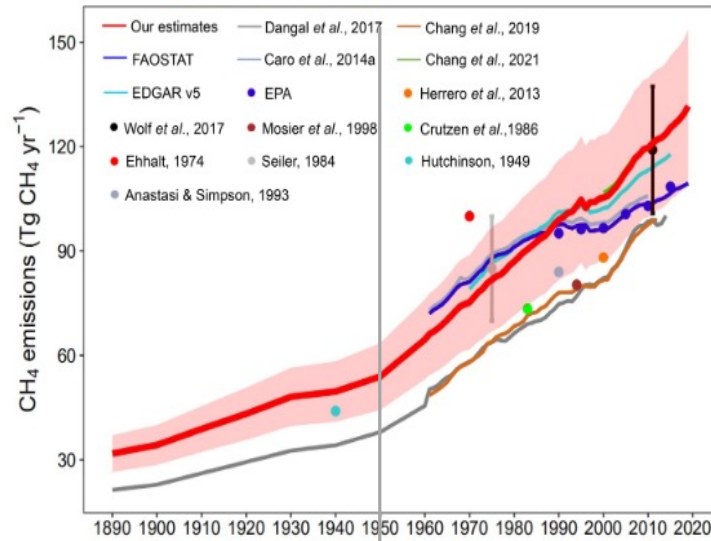
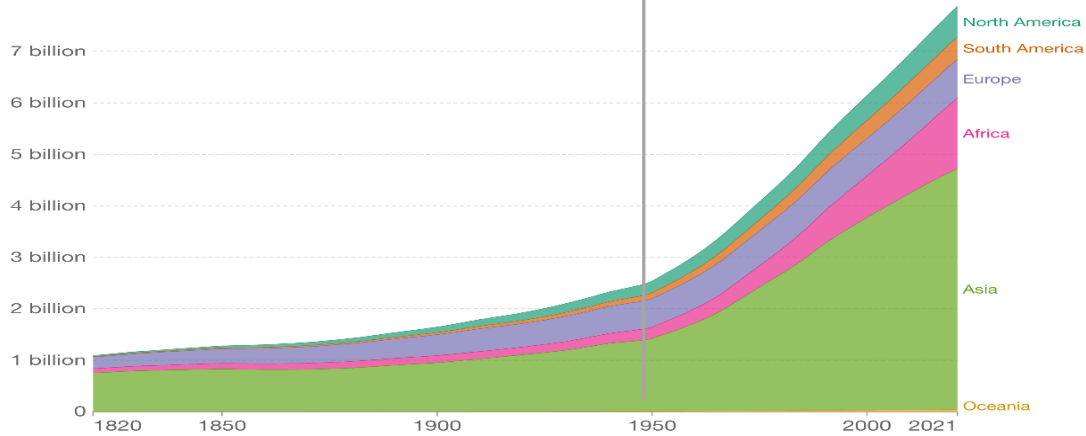


FIGURE 10 Estimates of CH<sub>4</sub> emissions from global livestock during 1890–2019 and comparisons with those reported in other inventories. The shaded area shows the 95% confidence interval of our estimates.

World population



Source: Gapminder (v6), HYDE (v3.2), UN (2019)

OurWorldInData.org/world-population-growth/ · CC BY



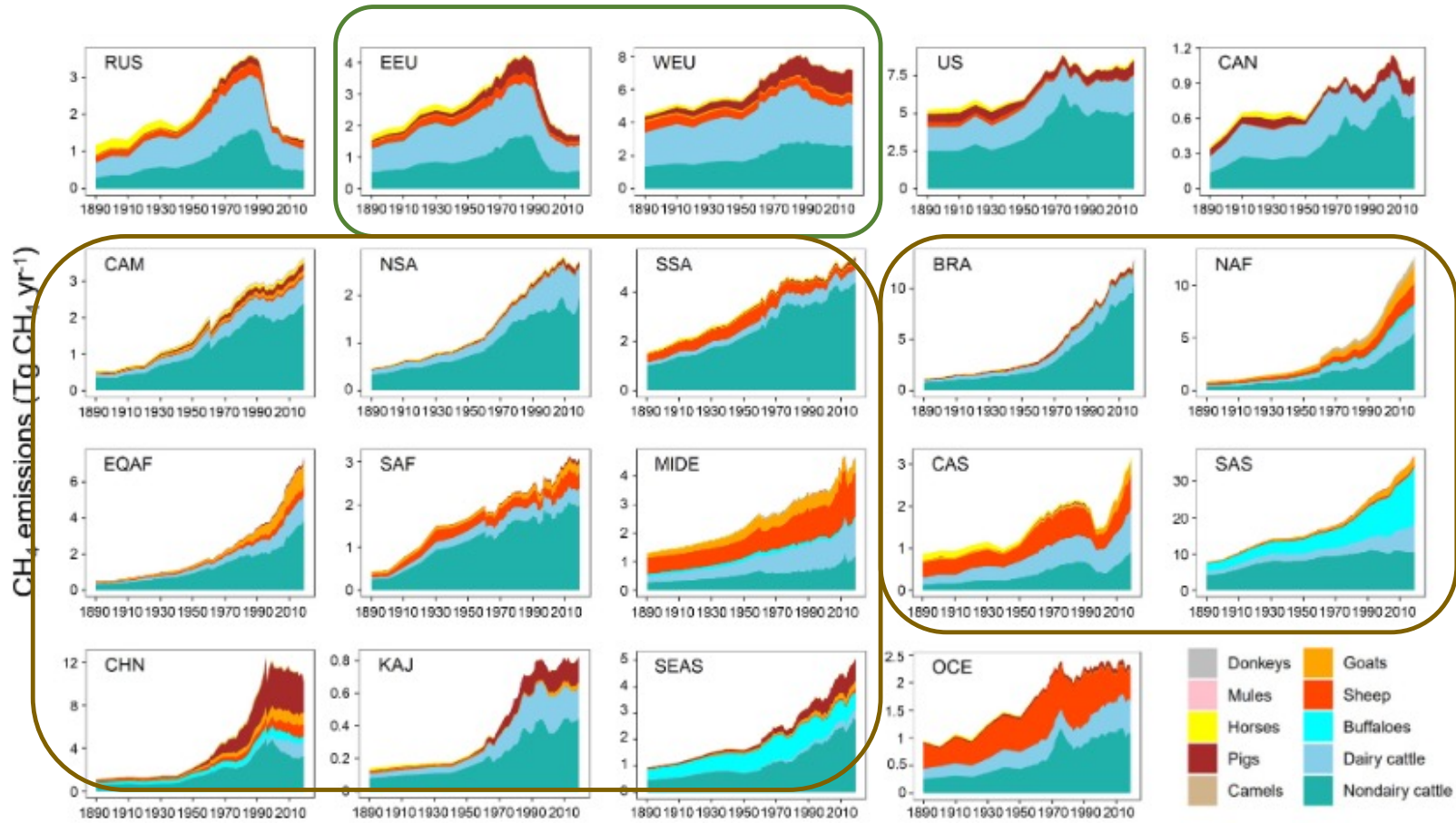
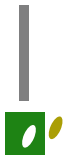
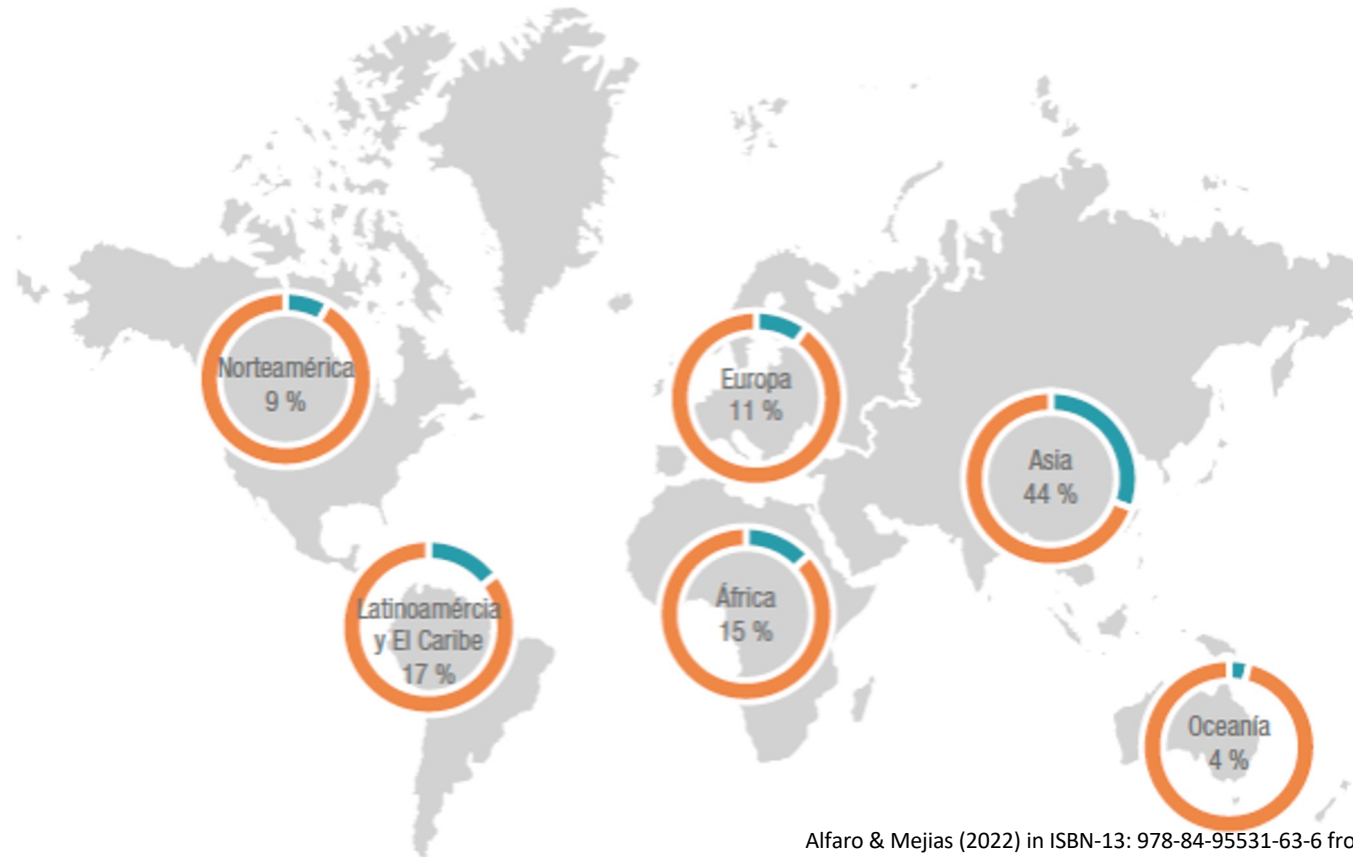


FIGURE 5 Temporal changes in regional CH<sub>4</sub> emissions from livestock during 1890-2019.

Zhang et al. (2022) in GCB

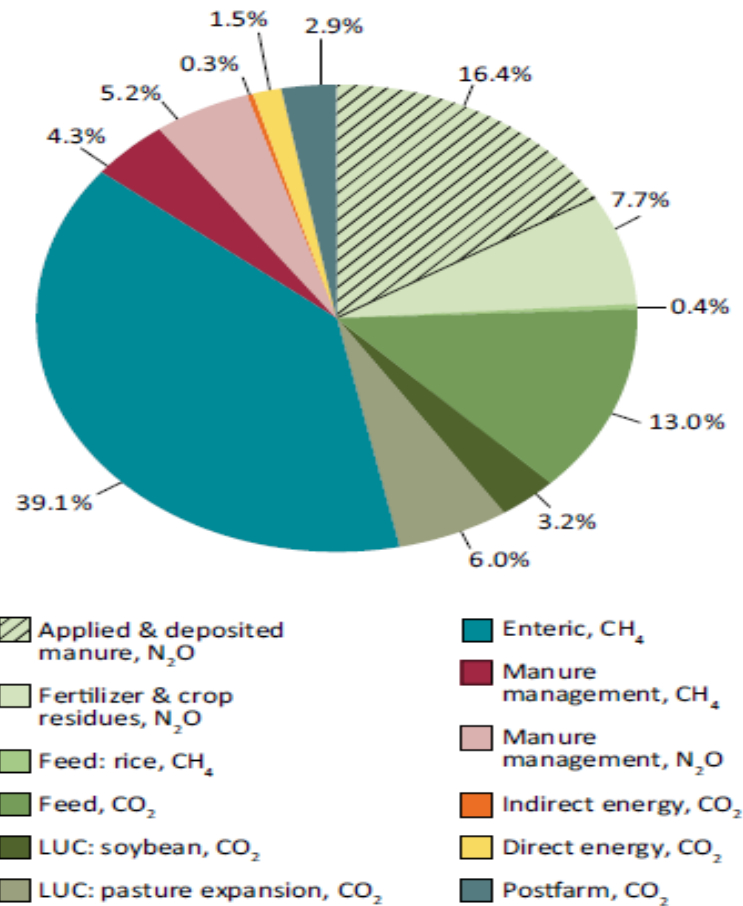


Alfaro & Mejias (2022) in ISBN-13: 978-84-95531-63-6 from FAO (2016)

Globally ~12% of total anthropogenic GHG emissions (direct and indirect) (FAO,2023)



**FIGURE 4.** Global emissions from livestock supply chains by category of emissions



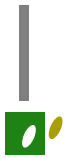
- Reduce Methane from enteric fermentation
- Reduce Feed associated CO<sub>2</sub> from improving Feed Efficiency



# | Solutions to achieve net zero carbon emissions



- **Contribution of livestock to society**
  - Healthy diet
  - Rural development and landscape maintenance
  - Negative consequences of eliminating livestock
  - Increase sustainability without compromising food security. New technologies and innovations are required.
- **Evaluate each case scenario**



**WA** WORLD ASSOCIATION  
**AP** for ANIMAL PRODUCTION  
www.waap.it

Number  
**26**  
2021  
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**The World Animal Science News**

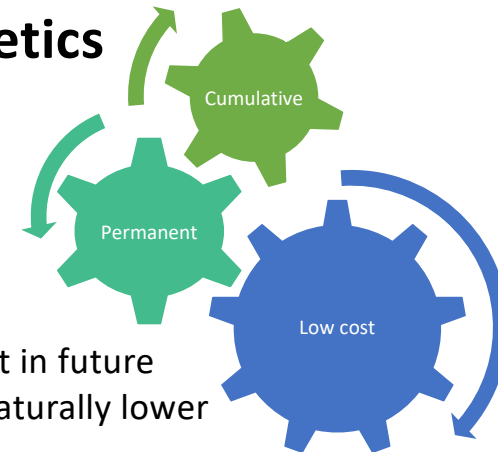
Main Topics

- From WAAP members
- News from Science
- News from Industry
- Job Offers
- Publications
- Meetings and Conferences

**EDITORIAL**

**When methane returns to the forefront of the climate scene, ruminants are in great danger!**

1. Implement proper calculation of GHG emissions \*
2. Nutrition
3. Technology: in-farm use of methane
4. **Genetics**



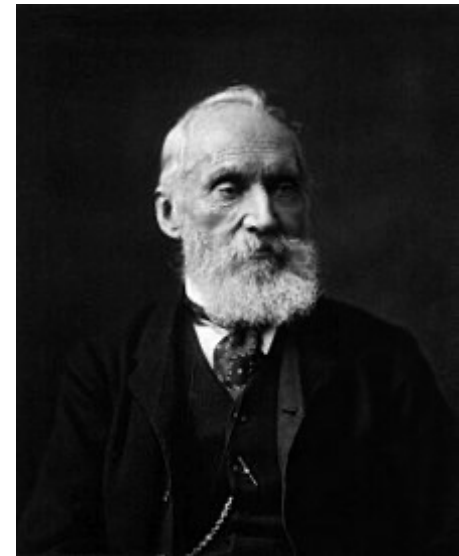
Targeted breeding can result in future generations of cattle with naturally lower emissions



# MEASURING METHANE

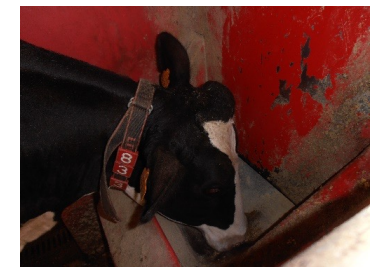
- If you **cannot measure it, you cannot improve it.**

“When you can measure what you are speaking about, and express it in numbers, you know something about it; but when you cannot measure it, when you cannot express it in numbers, your knowledge is of a meagre and unsatisfactory kind.”



Lord Kelvin (1824 – 1907)

# METHANE PHENOTYPING



Precision farming



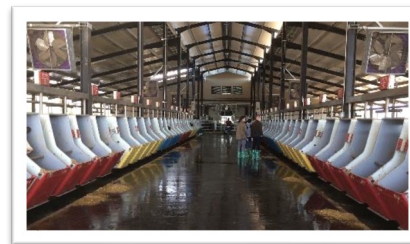
Genotypes (50k SNPs)

2017\_09\_21.093427



# METHANE PHENOTYPING

## 1. Recording methane in commercial and experimental farms



**Samples= 3281 dairy cows in 34 farms**

# METHANE PHENOTYPING

## 1. Recording methane and DMI in commercial and experimental farms



## 2. Genomic selection (genotyping, phenotyping). International collaborations (AUS, CAN, DNK, ESP, GER, NDL, SWI, USA)



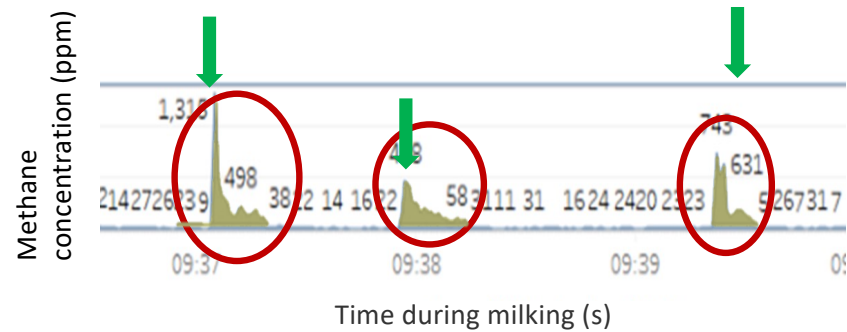
# METHANE PHENOTYPING

- **Phenotype definition**

1. Mean CH<sub>4</sub> (by second and every 5 s)
2. Sum of peaks CH<sub>4</sub> (by second and every 5 s)
3. Sum of max peaks CH<sub>4</sub>
4. Area under the curve (AUC CH<sub>4</sub>)
5. Ratio of (mean) CH<sub>4</sub>/CO<sub>2</sub>
6. CH<sub>4</sub> grams per day (Madsen et al., 2010)

$$\text{Prod CH}_4\left(\frac{g}{d}\right) = 0.714 * \text{ratio}(\text{ppm}) * 180 * 24 * 0.001 * (5.6 * \text{kg body mass}^{0.75}) + 22 * \text{ECM} + 1.6 * 10^{-5} * \text{days in pregnancy}$$

7. CH<sub>4</sub> grams per day (in-house\*)



Weekly averages were used for all phenotypes

# Heritabilities

| Trait                     | $h^2$             | $h^2$ (van Breukelen) | $r^2$            |
|---------------------------|-------------------|-----------------------|------------------|
| Mean CH4 (ppm)            | 0.08 (0.05-0.11)  |                       | 0.54 (0.52;0.56) |
| Mean CH4 5 s(ppm)         | 0.08 (0.05-0.11)  | 0.21-0.27             | 0.54 (0.53;0.56) |
| Sum of peaks CH4 (ppm)    | 0.09 (0.06;0.12)  |                       | 0.55 (0.53;0.57) |
| Sum of peaks CH4 5s (ppm) | 0.10 (0.06;0.13)  |                       | 0.55 (0.53;0.57) |
| Sum of max peaks (ppm)    | 0.08 (0.05;0.11)  |                       | 0.52 (0.50;0.54) |
| AUC CH4 (ppm)             | 0.10 (0.07;0.13)  |                       | 0.55 (0.53;0.57) |
| CO <sub>2</sub> (ppm)     | 0.02 (0.004;0.04) | 0.23-0.44             | 0.58 (0.56;0.60) |
| CO <sub>2</sub> (L/d)*    | 0.02 (0.004;0.05) |                       | 0.61 (0.59;0.63) |
| Ratio CH4/CO <sub>2</sub> | 0.10 (0.05;0.16)  | 0.02                  | 0.42 (0.39;0.45) |
| MeP (g/d) (Madsen eq)     | 0.12 (0.06;0.17)  | 0.33 (GF)             | 0.51 (0.48;0.54) |
| MeP (g/d)*                | 0.10 (0.04;0.15)  |                       | 0.55 (0.53;0.58) |



Low to moderate heritabilities and large repeatability

# Results: Genetic correlations between methane traits

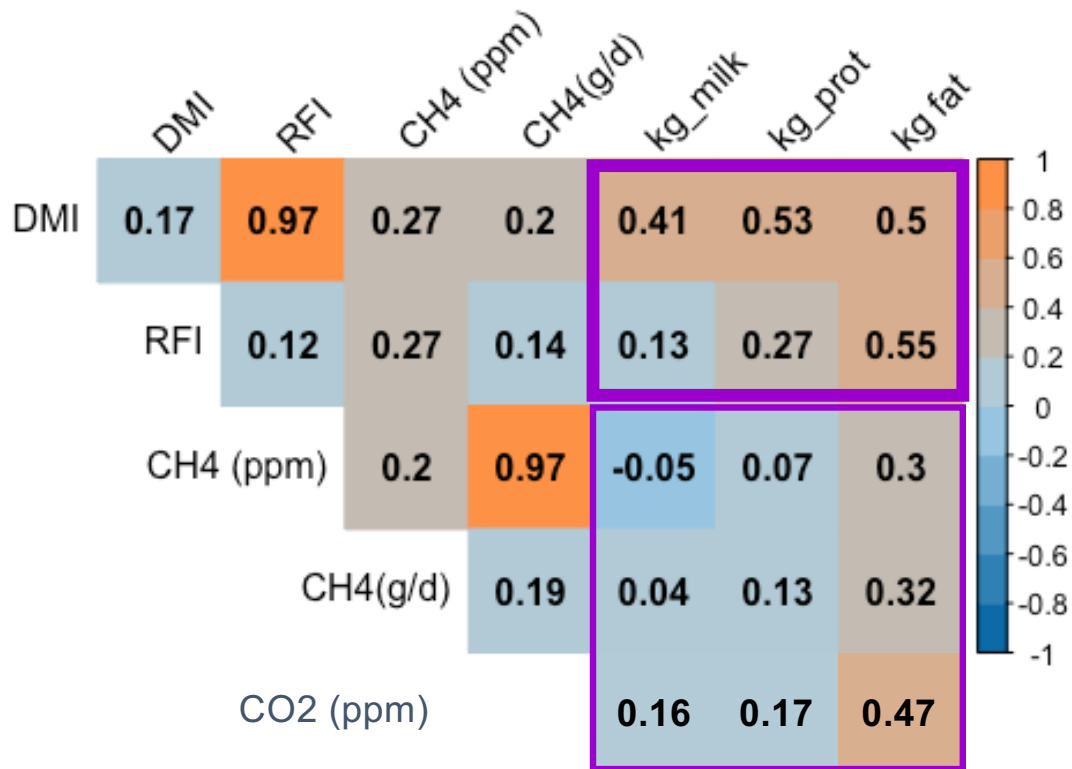
| Traits                           | Mean CH <sub>4</sub> | Mean CH <sub>4</sub> 5 s | MeP (g/d)* | Sum of peaks CH <sub>4</sub> | Sum of peaks CH <sub>4</sub> 5s | Sum of max peaks | AUC CH <sub>4</sub> |
|----------------------------------|----------------------|--------------------------|------------|------------------------------|---------------------------------|------------------|---------------------|
| Mean CH <sub>4</sub>             | 0.08                 | 0.99                     | 0.99       | 0.82 (0.74;0.90)             | 0.82 (0.75;0.90)                | 0.77 (0.67;0.85) | 0.83 (0.74;0.89)    |
| Mean CH <sub>4</sub> 5 s         |                      | 0.08                     | -          | 0.82 (0.73;0.89)             | 0.84 (0.76;0.92)                | 0.72 (0.57;0.85) | 0.84 (0.76;0.90)    |
| MeP (g/d)*                       |                      |                          | 0.10       | 0.52                         | -                               | 0.63             | 0.52                |
| Sum of peaks CH <sub>4</sub>     |                      |                          |            | 0.09                         | 0.99                            | 0.99             | 0.99                |
| Sum of peaks CH <sub>4</sub> 5 s |                      |                          |            |                              | 0.10                            | 0.99             | 0.99                |
| Sum of max peaks                 |                      |                          |            |                              |                                 | 0.08             | 0.99                |
| AUC CH <sub>4</sub>              |                      |                          |            |                              |                                 |                  | 0.10                |



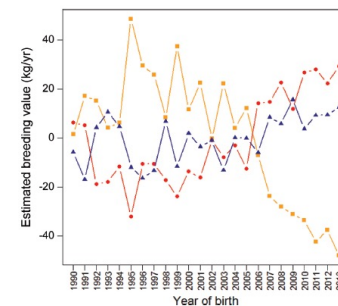
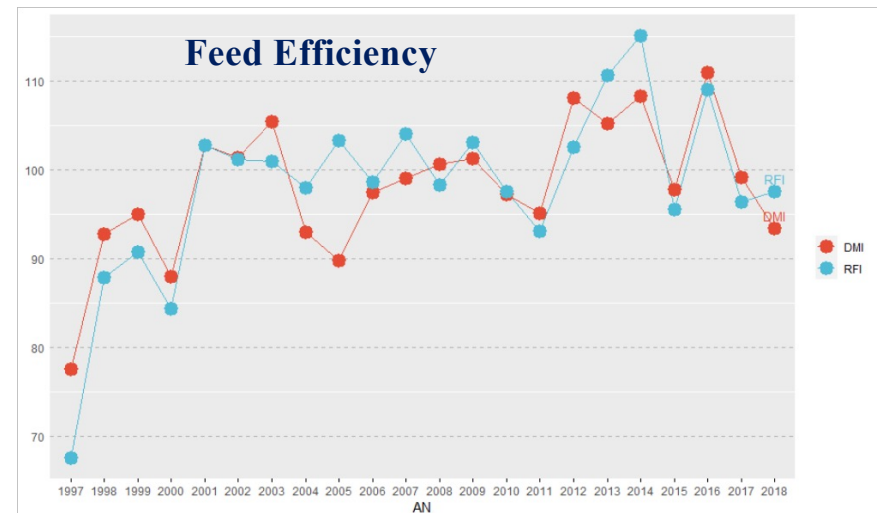
High correlations between phenotypes



# Genetic correlations



## Genetic trendsx



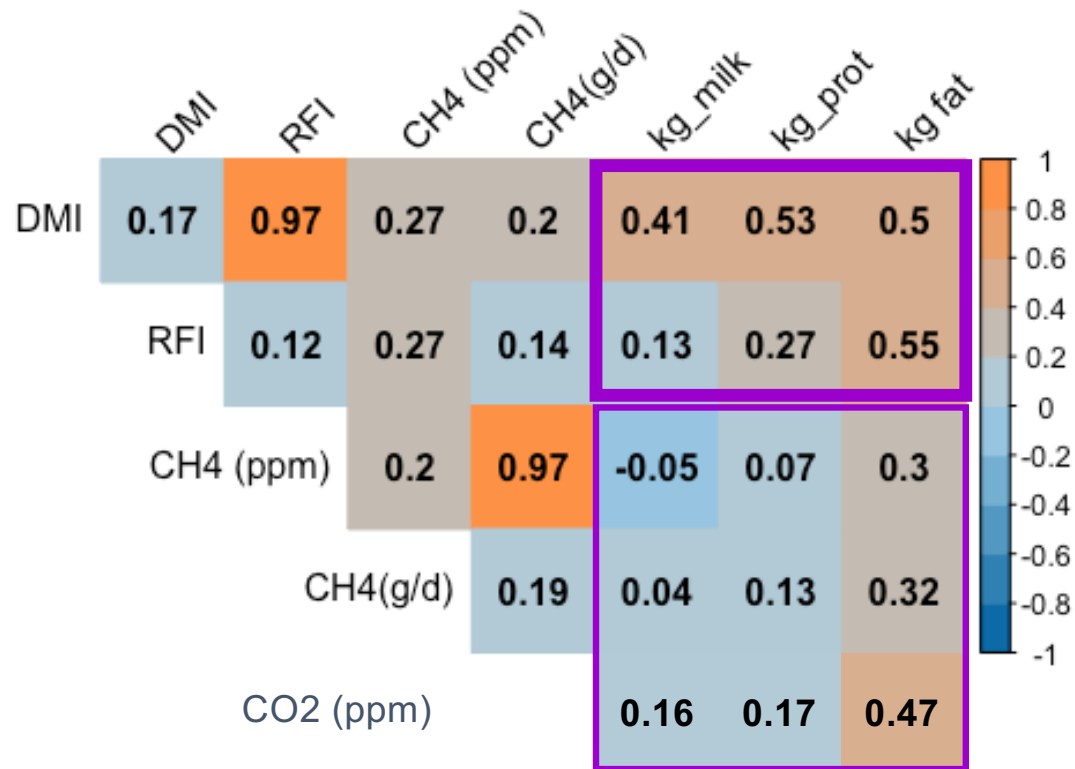
J. Dairy Sci. 98:7340–7350  
<http://dx.doi.org/10.3168/jds.2015-9621>  
 © American Dairy Science Association®, 2015.

**Hot topic: Definition and implementation of a breeding value for feed efficiency in dairy cows**

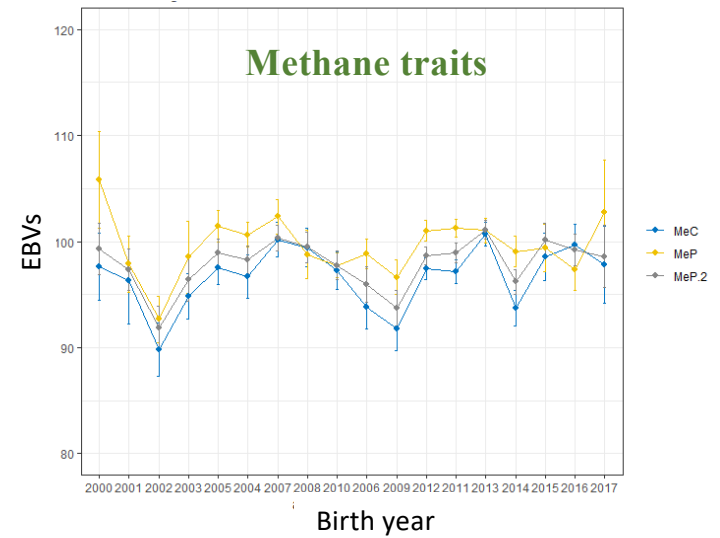
J. E. Pryce,<sup>1\*</sup> O. Gonzalez-Recio,<sup>1</sup> G. Nieuwhof,<sup>2</sup> W. J. Wales,<sup>3</sup> M. P. Coffey,<sup>4</sup> B. J. Hayes,<sup>1†</sup> and M. E. Goddard<sup>1‡</sup>



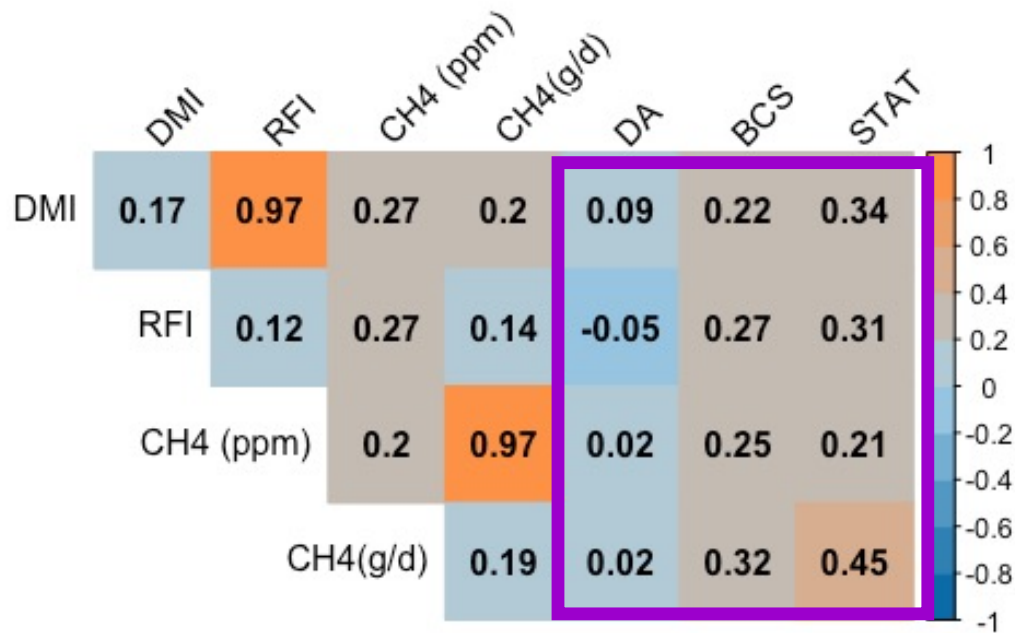
# Genetic correlations



Genetic trends



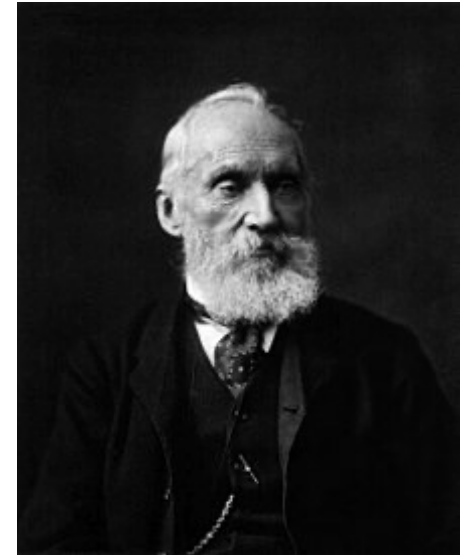
# Genetic correlations



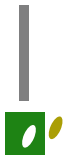
- Bigger cows tend to be less efficient and produce more methane

# Challenges at methane phenotyping

- Large scale phenotyping – sniffers, which can only be installed in AMS (yet)
- Rely on a reference population and genomic selection
- Other options to increase large scale phenotyping?



“If you cannot measure it, you cannot improve it”



#### ROLE OF MICROBIOME IN

- *Feed digestion*
- *Feed efficiency (DMI)*
- *Production traits*

#### ROLE OF MICROBIOME IN

- *Methane emissions*

#### ROLE OF MICROBIOME IN

- *Overall health*
- *Dysbiosis*
- *Pathogens*



#### ROLE OF HOST

- *Symbiosis*
- *Microbiome composition*
- *Heritability, and genetic correlation*

# Previous findings (summary)



- Rumen eukaryotes are the main phenotypic risk factors for larger methane emissions in dairy cattle (Saborío-Montero et al., 2022).
- Certain microbial groups are likely causal effects of larger/lower methane production in the rumen (Saborío-Montero et al. 2019; López-García et al. 2022).
- The cow genome partially controls the rumen microbiome composition (Saborío-Montero et al., 2019; González-Recio et al. 2023)

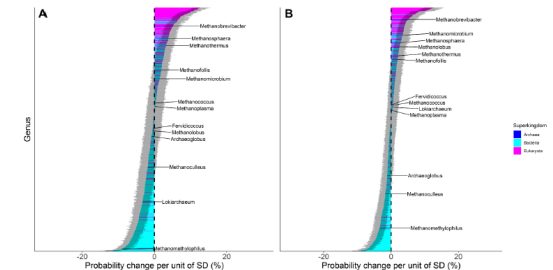
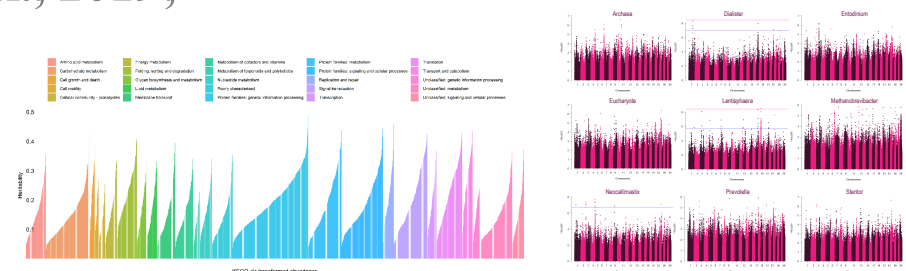
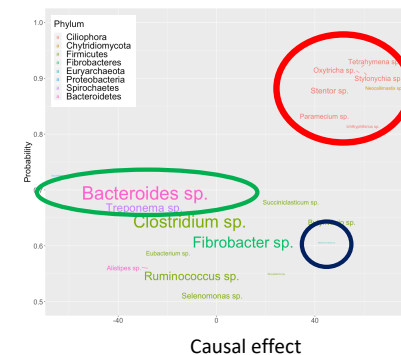


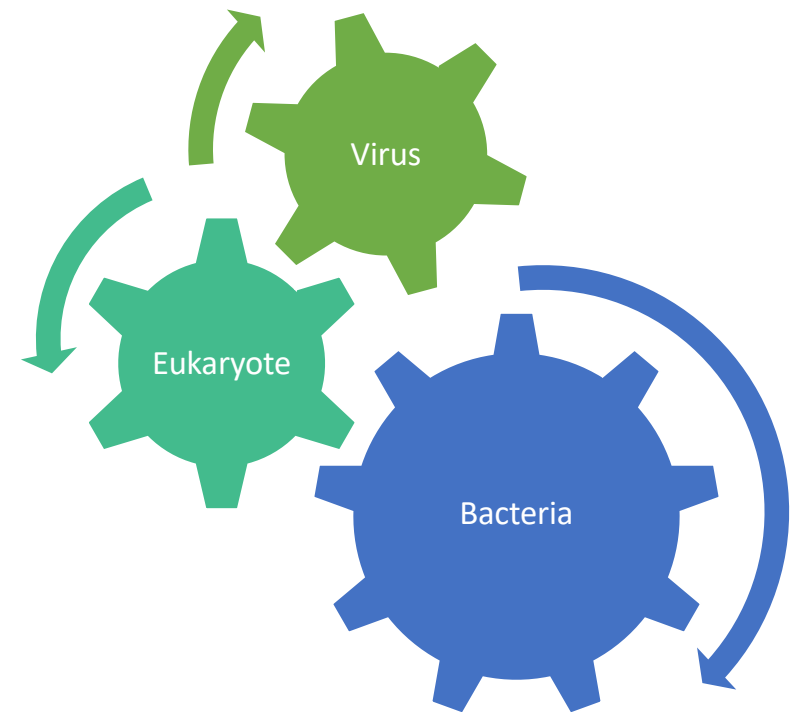
Fig. 3. Change in the probability of being classified in the upper quartile for (A) methane concentration (ppm CH<sub>4</sub>) and (B) methane intensity (ppm CH<sub>4</sub>/kg milk) per unit of standard deviation for relative abundance (%) of 1240 genera colored by superkingdom. Black dashed line indicates the baseline probability of being classified in the upper quartiles without any genus effect. All the archaea genera are explicitly indicated. Probability intervals based on posterior standard deviations are depicted in gray for all genera.



# Rumen microbiome community

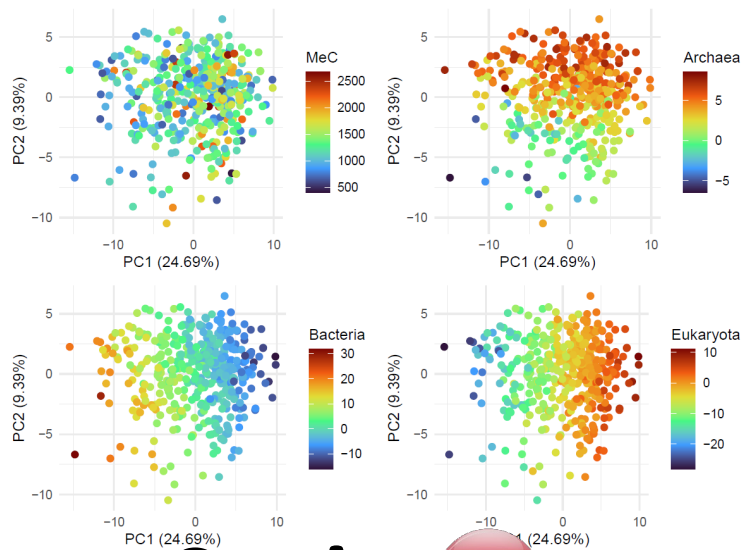


- Large dimensionality, redundancy and interplay
- Should focus on one or few microbial taxa/genes as there will be changes in other groups
- Develop strategies to modulate the whole microbiome towards low emitting, efficient and healthy microbiomes.



# HOW TO ACCOUNT FOR MICROBIOME COMPLEXITY

- Aggregated variables capture a relevant portion of microbial variability



Paves the way to international collaborations to create multi country reference population for metagenome composition

**Spain**



33% total microbiome variability



*Similar microbiome composition and relationship with methane under 2 different production systems (Australia and Spain)*



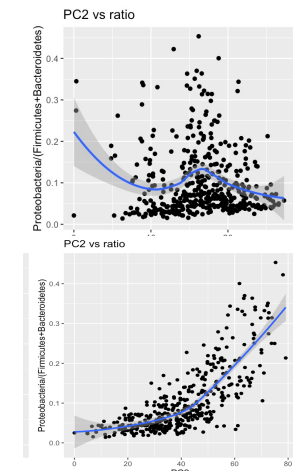
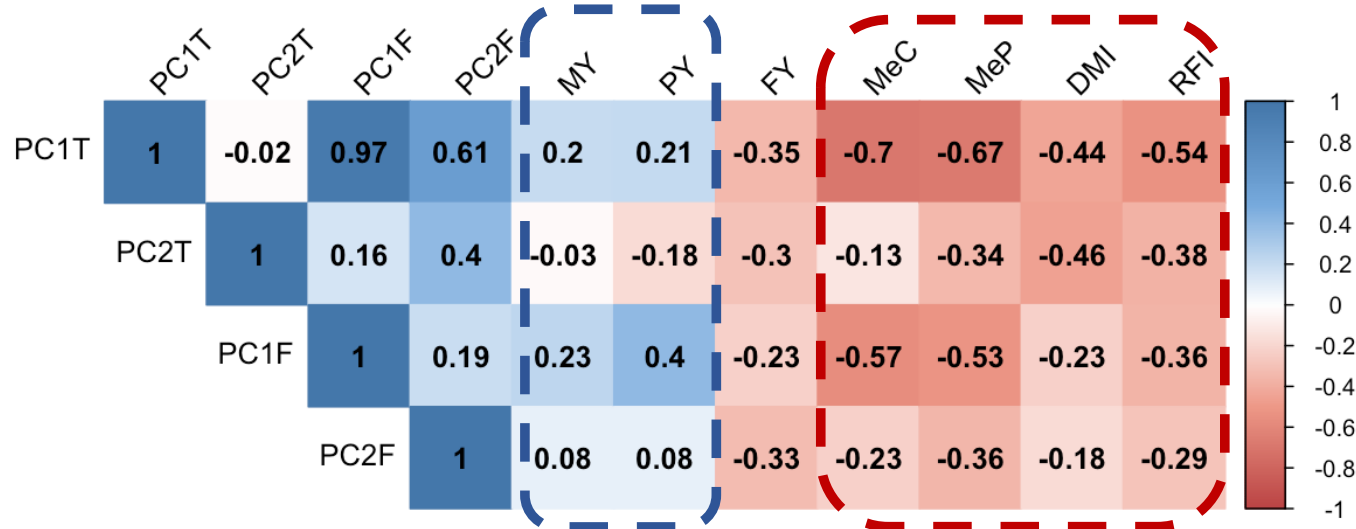
J. Dairy Sci. 104:8135–8151  
<https://doi.org/10.3168/jds.2020-20005>  
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**A dimensional reduction approach to modulate the core ruminal microbiome associated with methane emissions via selective breeding**

Alejandro Saborio-Montero,<sup>1,2</sup> Adrian López-García,<sup>1</sup> Mónica Gutiérrez-Rivas,<sup>1</sup> Raquel Atxaerandio,<sup>1</sup> Idoia Goiri,<sup>2</sup> Aser García-Rodríguez,<sup>2</sup> José A. Jiménez-Montero,<sup>2</sup> Carmen González,<sup>1</sup> Javier Tamames,<sup>2</sup> Fernando Puente-Sánchez,<sup>2</sup> Luis Varona,<sup>1</sup> Magdalena Serrano,<sup>1</sup> Cristina Ovilo,<sup>1</sup> and Oscar González-Reco<sup>1,2\*</sup>

# Genetic correlations: Rumen microbiome community

## Genetic correlation between PC and other traits



- Aggregated microbial variables have heritabilities 0.20-0.30.
- Aggregated microbial variables are favorably genetically correlated with milk, protein, methane emissions and feed efficiency (unfavorable with fat)
- Breeding for lower methane or feed efficiency may impact microbiota composition (dysbiosis)
- What are the expected consequences?



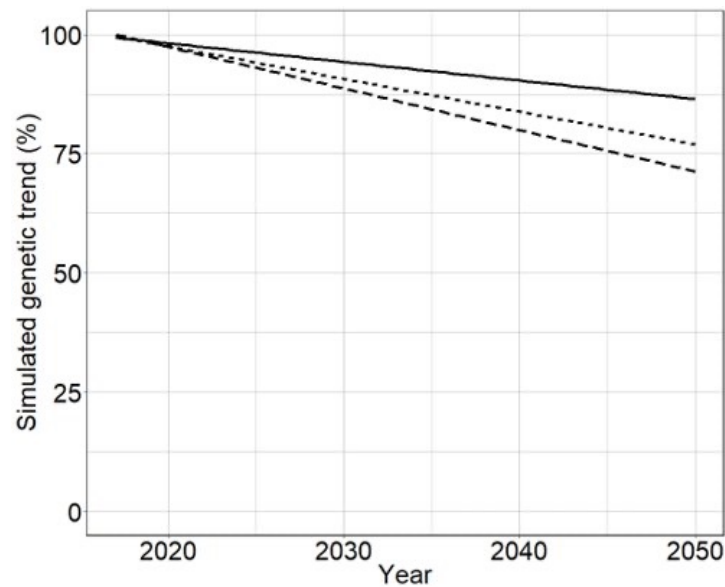


# Methane emissions in breeding goals and selection indices

# Animal breeding as a mitigation strategy

## Methane intensity (g/kg milk) & methane emissions (g/d)

- Current trend
- Combined selection for CH<sub>4</sub> and other traits
- Theoretical maximum (exclusively focusing on methane)



<https://doi.org/10.1016/j.animal.2021.100294>  
(de Haas *et al.* 2021)

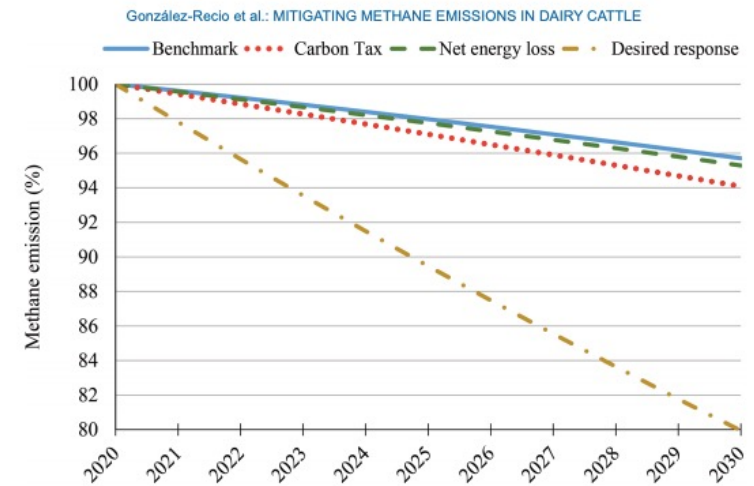
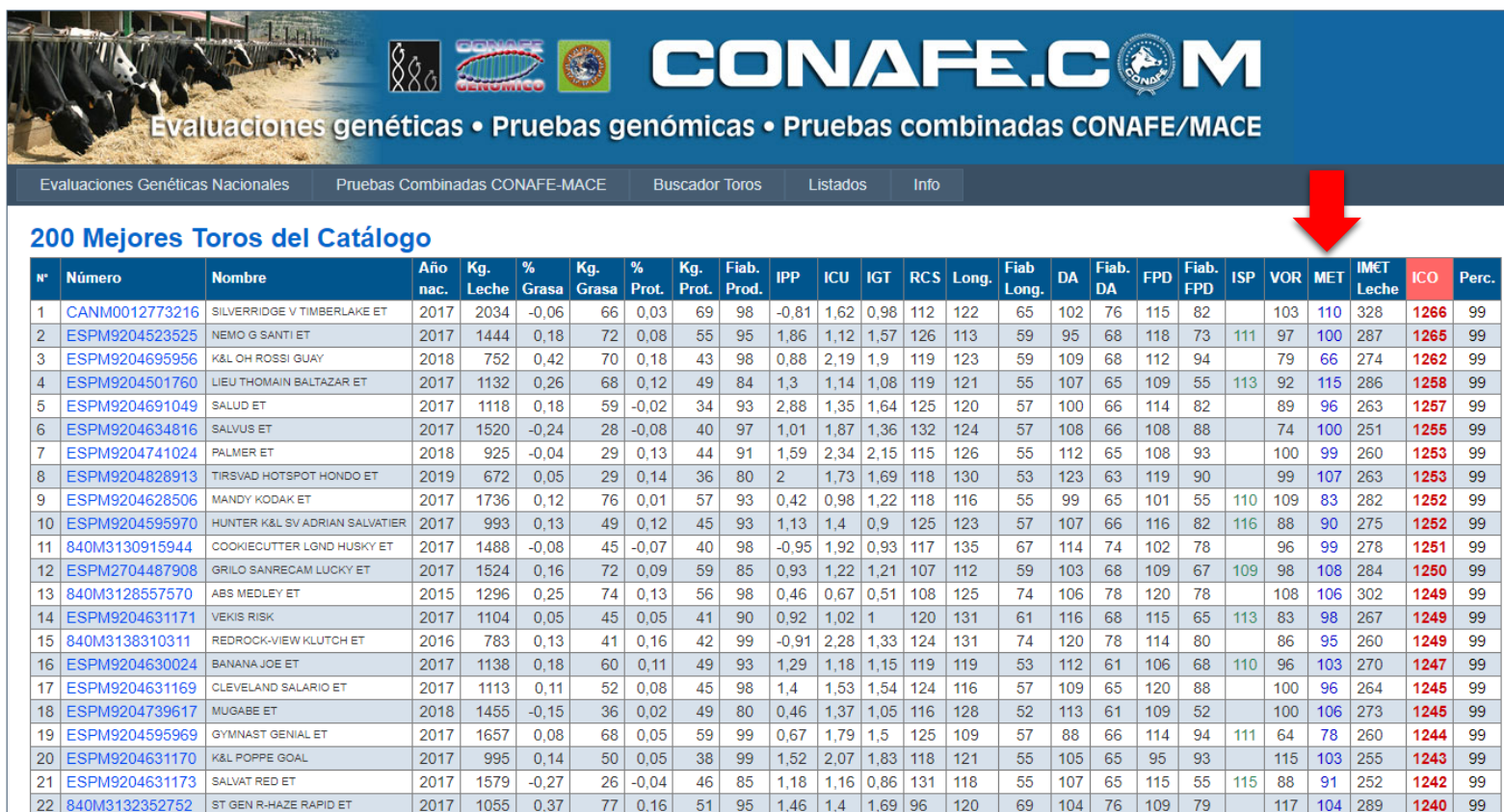


Figure 2. Expected reduction in percentage from current levels in methane emissions produced by Holstein cows in Spain based on genetic gain in methane emissions (MET; t/yr) under the 4 scenarios: benchmark, carbon tax, net energy loss, and desired response (i.e., number of cows  $\times$  MET genetic gain  $\times$  time/1,000). A decrease of 1.5% in the number of dairy cows was considered each year, following census data from the Spanish Holstein association: ([http://www.conafe.com/VisorDocs.aspx?pdf=estadisticas\\_CENSO\\_DE\\_ANIMALES.pdf](http://www.conafe.com/VisorDocs.aspx?pdf=estadisticas_CENSO_DE_ANIMALES.pdf)).

# Genetic evaluations

Published as an individual trait in the catalog.



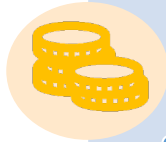
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### 200 Mejores Toros del Catálogo

| Nº | Número         | Nombre                         | Año nac. | Kg. Leche | % Grasa | Kg. Grasa | % Prot. | Kg. Prot. | Fiab. Prod. | IPP   | ICU  | IGT  | RCS | Long. | Fiab. Long. | DA  | Fiab. DA | FPD | Fiab. FPD | ISP | VOR | MET | IMET Leche | ICO  | Perc. |
|----|----------------|--------------------------------|----------|-----------|---------|-----------|---------|-----------|-------------|-------|------|------|-----|-------|-------------|-----|----------|-----|-----------|-----|-----|-----|------------|------|-------|
| 1  | CANM0012773216 | SILVERRIDGE V TIMBERLAKE ET    | 2017     | 2034      | -0,06   | 66        | 0,03    | 69        | 98          | -0,81 | 1,62 | 0,98 | 112 | 122   | 65          | 102 | 76       | 115 | 82        |     | 103 | 110 | 328        | 1266 | 99    |
| 2  | ESPM9204523525 | NEMO G SANTI ET                | 2017     | 1444      | 0,18    | 72        | 0,08    | 55        | 95          | 1,86  | 1,12 | 1,57 | 126 | 113   | 59          | 95  | 68       | 118 | 73        | 111 | 97  | 100 | 287        | 1265 | 99    |
| 3  | ESPM9204695956 | K&L OH ROSSI GUAY              | 2018     | 752       | 0,42    | 70        | 0,18    | 43        | 98          | 0,88  | 2,19 | 1,9  | 119 | 123   | 59          | 109 | 68       | 112 | 94        |     | 79  | 66  | 274        | 1262 | 99    |
| 4  | ESPM9204501760 | LIEU THOMAIN BALTAZAR ET       | 2017     | 1132      | 0,26    | 68        | 0,12    | 49        | 84          | 1,3   | 1,14 | 1,08 | 119 | 121   | 55          | 107 | 65       | 109 | 55        | 113 | 92  | 115 | 286        | 1258 | 99    |
| 5  | ESPM9204691049 | SALUD ET                       | 2017     | 1118      | 0,18    | 59        | -0,02   | 34        | 93          | 2,88  | 1,35 | 1,64 | 125 | 120   | 57          | 100 | 66       | 114 | 82        |     | 89  | 96  | 263        | 1257 | 99    |
| 6  | ESPM9204634816 | SALVUS ET                      | 2017     | 1520      | -0,24   | 28        | -0,08   | 40        | 97          | 1,01  | 1,87 | 1,36 | 132 | 124   | 57          | 108 | 66       | 108 | 88        |     | 74  | 100 | 251        | 1255 | 99    |
| 7  | ESPM9204741024 | PALMER ET                      | 2018     | 925       | -0,04   | 29        | 0,13    | 44        | 91          | 1,59  | 2,34 | 2,15 | 115 | 126   | 55          | 112 | 65       | 108 | 93        |     | 100 | 99  | 260        | 1253 | 99    |
| 8  | ESPM9204828913 | TIRSVAD HOTSPOT HONDO ET       | 2019     | 672       | 0,05    | 29        | 0,14    | 36        | 80          | 2     | 1,73 | 1,69 | 118 | 130   | 53          | 123 | 63       | 119 | 90        |     | 99  | 107 | 263        | 1253 | 99    |
| 9  | ESPM9204628506 | MANDY KODAK ET                 | 2017     | 1736      | 0,12    | 76        | 0,01    | 57        | 93          | 0,42  | 0,98 | 1,22 | 118 | 116   | 55          | 99  | 65       | 101 | 55        | 110 | 109 | 83  | 282        | 1252 | 99    |
| 10 | ESPM9204595970 | HUNTER K&L SV ADRIAN SALVATIER | 2017     | 993       | 0,13    | 49        | 0,12    | 45        | 93          | 1,13  | 1,4  | 0,9  | 125 | 123   | 57          | 107 | 66       | 116 | 82        | 116 | 88  | 90  | 275        | 1252 | 99    |
| 11 | 840M3130915944 | COOKIECUTTER LGND HUSKY ET     | 2017     | 1488      | -0,08   | 45        | -0,07   | 40        | 98          | -0,95 | 1,92 | 0,93 | 117 | 135   | 67          | 114 | 74       | 102 | 78        |     | 96  | 99  | 278        | 1251 | 99    |
| 12 | ESPM2704487908 | GRILO SANRECAM LUCKY ET        | 2017     | 1524      | 0,16    | 72        | 0,09    | 59        | 85          | 0,93  | 1,22 | 1,21 | 107 | 112   | 59          | 103 | 68       | 109 | 67        | 109 | 98  | 108 | 284        | 1250 | 99    |
| 13 | 840M3128557570 | ABS MEDLEY ET                  | 2015     | 1296      | 0,25    | 74        | 0,13    | 56        | 98          | 0,46  | 0,67 | 0,51 | 108 | 125   | 74          | 106 | 78       | 120 | 78        |     | 108 | 106 | 302        | 1249 | 99    |
| 14 | ESPM9204631171 | VEKIS RISK                     | 2017     | 1104      | 0,05    | 45        | 0,05    | 41        | 90          | 0,92  | 1,02 | 1    | 120 | 131   | 61          | 116 | 68       | 115 | 65        | 113 | 83  | 98  | 267        | 1249 | 99    |
| 15 | 840M3138310311 | REDROCK-VIEW KLUTCH ET         | 2016     | 783       | 0,13    | 41        | 0,16    | 42        | 99          | -0,91 | 2,28 | 1,33 | 124 | 131   | 74          | 120 | 78       | 114 | 80        |     | 86  | 95  | 260        | 1249 | 99    |
| 16 | ESPM9204630024 | BANANAJOE ET                   | 2017     | 1138      | 0,18    | 60        | 0,11    | 49        | 93          | 1,29  | 1,18 | 1,15 | 119 | 119   | 53          | 112 | 61       | 106 | 68        | 110 | 96  | 103 | 270        | 1247 | 99    |
| 17 | ESPM9204631169 | CLEVELAND SALARIO ET           | 2017     | 1113      | 0,11    | 52        | 0,08    | 45        | 98          | 1,4   | 1,53 | 1,54 | 124 | 116   | 57          | 109 | 65       | 120 | 88        |     | 100 | 96  | 264        | 1245 | 99    |
| 18 | ESPM9204739617 | MUGABE ET                      | 2018     | 1455      | -0,15   | 36        | 0,02    | 49        | 80          | 0,46  | 1,37 | 1,05 | 116 | 128   | 52          | 113 | 61       | 109 | 52        |     | 100 | 106 | 273        | 1245 | 99    |
| 19 | ESPM9204595969 | GYMNAST GENIAL ET              | 2017     | 1657      | 0,08    | 68        | 0,05    | 59        | 99          | 0,67  | 1,79 | 1,5  | 125 | 109   | 57          | 88  | 66       | 114 | 94        | 111 | 64  | 78  | 260        | 1244 | 99    |
| 20 | ESPM9204631170 | K&L POPPE GOAL                 | 2017     | 995       | 0,14    | 50        | 0,05    | 38        | 99          | 1,52  | 2,07 | 1,83 | 118 | 121   | 55          | 105 | 65       | 95  | 93        |     | 115 | 103 | 255        | 1243 | 99    |
| 21 | ESPM9204631173 | SALVAT RED ET                  | 2017     | 1579      | -0,27   | 26        | -0,04   | 46        | 85          | 1,18  | 1,16 | 0,86 | 131 | 118   | 55          | 107 | 65       | 115 | 55        | 115 | 88  | 91  | 252        | 1242 | 99    |
| 22 | 840M3132352752 | ST GEN R-HAZE RAPID ET         | 2017     | 1055      | 0,37    | 77        | 0,16    | 51        | 95          | 1,46  | 1,4  | 1,69 | 96  | 120   | 69          | 104 | 76       | 109 | 79        |     | 117 | 104 | 289        | 1240 | 99    |

# Economic Selection indices proposed in Spain (IM€T)

## Breeding goals



- Production
- Mastitis & Lameness
- Feed Efficiency
- Fertility
- Longevity
- Live Weight
- Calving Ease
- Methane production (g/d)

### Economic value Feed intake:

- Cost of MFU per kg DM

### Economic value methane:

- Carbon price voluntary market for traditional systems → -0.14 € (g/d/cow)
- Carbon price for organic systems → -0.72 € (g/d/cow)

## Selection index

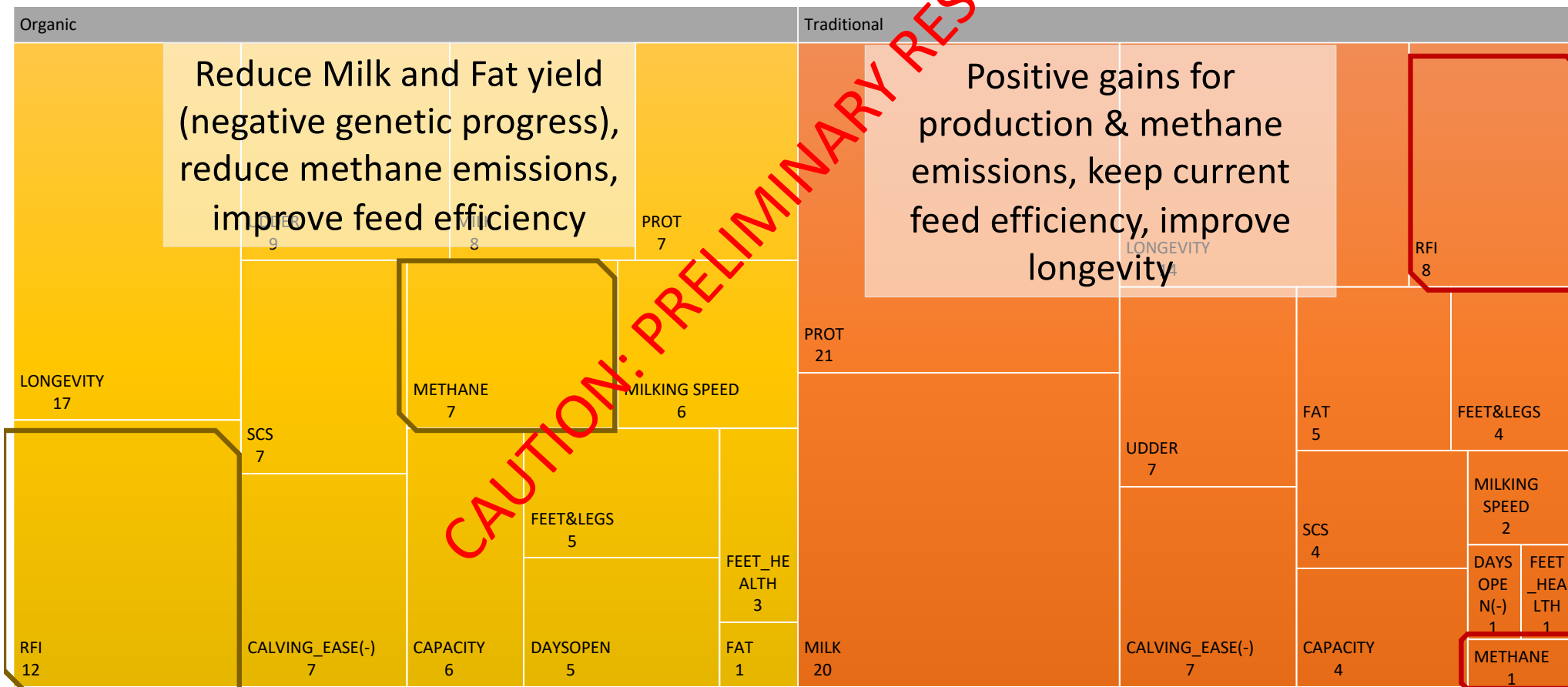
**Traditional**, cheese focused, pasture, **organic**

- kg milk
- kg fat
- kg prot
- RCS
- Feet & Legs
- Days Open
- Longevity
- Milking speed
- Maternal calving ease
- Capacity
- Residual Feed Intake
- Methane (ppm)

# Example of implementation

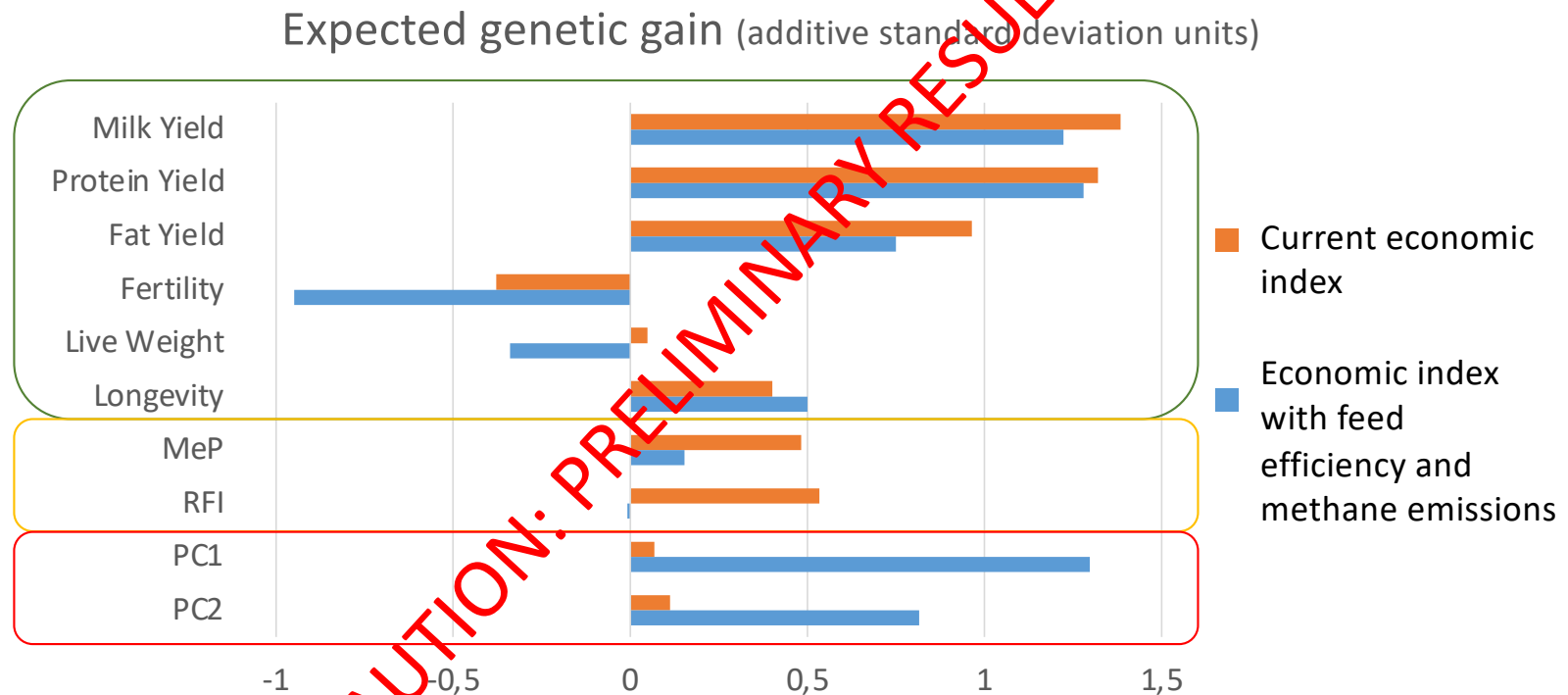


- Efficiency and sustainability index in Spanish Dairy cattle



# Example of implementation

- Efficiency and sustainability index in Spanish Dairy cattle
  - Subindex (will be integrated in the total merit index)



# TAKE HOME MESSAGE

**01**

**Measure, measure, measure**

Also in beef and small ruminants

**02**

**Reducing methane emission via selective breeding can have a great impact if, and only if, farmers are encouraged to breed for lower emissions**

**03**

**Rumen microbiome information is a suitable complementary phenotype for reducing methane emissions**

Selecting for lower emissions and better feed efficiency may impact rumen microbiome health, and it should be controlled

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