





## Modelling for the prediction of beef sensory quality

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In Europe, there is still no reliable tool to predict beef quality to deliver consistent quality beef to consumer.

Beef Quality depends on differences in muscle characteristics (muscle fibre types, collagen content, lipid content, etc.).

These differences are attributed to different factors: genetics, muscle type, breed and sex, etc.



aims to advance beef safety and quality across Europe

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#### Objective of this study: How muscle biochemical traits may explain variability in quality scores?

1. Trained consumers

2. Untrained consumers





we have compiled all biochemical data of the muscle tissue from a great number of experiments in a database called **BIF-Beef** (Integrated and Functional Biology of Beef)

→ The objective here is to perform meta-analyses in order to relate muscle biochemical data to meat quality.



1. Trained consumers





# We recorded sensory analysis according to the guidelines of **the Meat Standards Australia (MSA**) to relate MSA quality scores to muscle biochemical data system

Legrand et al., 2012, Animal, In Press

2. Untrained consumers







# Part 1 of this study



Parte 1 of this study:









Explain and predict variability in beef quality by muscle biochemical traits



#### Materials & methods



#### **4037 striploin** (M. longissimus thoracis)

samples from **YOUNG bulls** of similar age (15 months)

#### with a specific focus

21 Charolais young bulls which differed in age from 15 to 26 months

### **Trained panellists**

Relationship between IMF and Flavour (trained panellist consumers)

#### **Results**

On average, with 4037 striploin samples from mainly young bulls of similar age (15 months of age), the partial correlation between flavour and intramuscular fat level was low but significant (0.11\*\*\*). Thus, less than 2% of the variation in flavour could be explained by differences in intramuscular fat level with this homogenous population of young bulls.

> Hocquette et al., 2011 Animal Production Science, 2011, 51, 975–981

#### **Results of the specific analysis**

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In that particular case, we can explain up to 41% of the variation in flavour by differences in intramuscular fat content. Indeed, changes in age induce larger variation in intramuscular fat level and then large differences in flavour.





# Part 2 of this study

#### Part 2 of this study: We did sensory analysis according to MSA guidelines

Australia has developed the Meat Standards Australia (MSA) since 1996. MSA is a product grading scheme to predict beef quality by <u>untrained</u> consumers.

Score are allocated for each individual "muscle x cooking method" combination using various information on the corresponding animals and meats.





#### Materials & methods



108 cuts from 6 different muscles

[Outside (M. biceps femoris) Topside (M. semimembranosus) Striploin (M. longissimus thoracis), Rump (M. gluteus medius) Oyster blade (M. infraspinatus) Tenderloin (M. psoas major)]

sampled from 18 animals of different ages, breeds and sexes (3 young bulls +15 cows)

### **Untrained panellists**

### Relationship between IMF and Flavour MSA scores (untrained consumers)



Flavour MSA scores

Relationship between MSA scores and biochemical muscle data

#### Other significant correlations: R<sup>2</sup> (P<0.05)

Soluble / total collagen (Solubility indicator)

with <u>MSA tenderness</u> score: R<sup>2</sup> = 0.33

with <u>MSA overliking</u> score: R<sup>2</sup> = 0.29

with <u>MSA palatability</u> score: R<sup>2</sup> = 0.30

#### Palatability score

(0.3 Tenderness + 0.3 Flavour + 0.1 Juiciness + 0.3 overliking)





This is among the first studies which related biochemical parameters of the muscle tissue to quality scores determined by untrained consumers

With untrained consumers, this study confirmed the importance of intramuscular fat level for beef flavour and of collagen solubility for tenderness, which is in accordance with observations with trained panelists

It is possible to develop a predictive model of beef quality from muscle and biochemical traits (this study) combined with muscle structure and genomic biomarkers (not presented)



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# Thank you for your attention



