

# Incorporating meat quality in sheep breeding programmes: *potential of non-invasive technologies*

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# Introduction

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- Genetic selection for lamb meat quality rare
- Difficult / expensive/ time consuming to measure
  - Direct tests:
    - post-mortem - on relatives, difficult to standardise
    - destructive - expensive, not possible on-line
  - Predictive tests:
    - mainly post-mortem; often destructive / invasive / slow
- Other potential hindrances:
  - Data feedback from abattoir; reliable traceability
  - On-line implementation

Species	Type	Country	Trait	Heritability	Reference
<i>Bos taurus</i>	Taurine	USA	Marbling	0.57 ± 0.13	(Wheeler <i>et al.</i> , 2001a)
			LT SF <sup>1</sup>	0.22 ± 0.12	
			Juiciness	0.09 ± 0.11	
			Flavour	0.07 ± 0.11	
			IMF	0.55 ± 0.14	
	Taurine	Australia	LT SF	0.11 ± 0.06 <sup>2</sup>	(Johnston <i>et al.</i> , 2003)
			Juiciness	0.15 ± 0.06	
			Flavour	0.05 ± 0.06	
	Zebu <sup>3</sup>	Australia	MQ4	0.13 ± 0.06	
			LT SF	0.31 ± 0.09	
Juiciness			0.20 ± 0.08 <sup>4</sup>		
Taurine	Australia	IMF	0.38 ± 0.04 <sup>5</sup>	(Reverter <i>et al.</i> , 2003)	
	Zebu	IMF	0.39 ± 0.03 <sup>5</sup>		
<i>Gallus gallus</i>	Broiler	France	Ultimate pH	0.49 ± 0.11	(Le Bihan-Duval <i>et al.</i> , 1999)
			Lightness	0.75 ± 0.08	
			redness	0.81 ± 0.04	
			yellowness	0.64 ± 0.06	
			IMF	0.08 ± 0.04	
<i>Ovis aries</i>	Merino	Australia	Meat pH	0.27 ± 0.09	(Fogarty <i>et al.</i> , 2003)
			Lightness	0.14 ± 0.07	
			redness	0.02 ± 0.06	
			yellowness	0.04 ± 0.06	
Composite	France	IMF	0.22	(Moreno <i>et al.</i> , 2001)	
<i>Sus scrofa</i>	Large White/Landrace	Australia	Meat pH	0.14 ± 0.04	(Hermesch <i>et al.</i> , 2000)
			Lightness	0.29 ± 0.06	
			Drip Loss	0.23 ± 0.05	
			IMF	0.35 ± 0.06	
	Duroc/Landrace	USA	Meat pH	0.14 ± 0.08	(Lo <i>et al.</i> , 1991)
			IMF	0.52 ± 0.13	
			Cooking loss	0.06 ± 0.06	
			Tenderness <sup>6</sup>	0.17 ± 0.08	
			Off flavour	0.03 ± 0.06	
			Consumer acceptance	0.34 ± 0.11	

<sup>1</sup> *Longissimus thoracis* shear force; <sup>2</sup> SE of heritability given as a range of 0.04–0.08 for the table see original reference; <sup>3</sup> Mixture of purebred zebu (e.g. Brahman) and breeds with some zebu ancestry; <sup>4</sup> SE of heritability given as a range of 0.07–0.09 for the table see original reference; <sup>5</sup> SE personal communication A. Reverter; <sup>6</sup> this is the objective measure of tenderness, for taste panel tenderness,  $h^2 = 0.45 \pm 0.12$ .

# Genetic control of meat quality



From: J.P. Kerry and David Ledward:  
Improving the Sensory and Nutritional Quality of Fresh Meat  
Elsevier, 2009

# Introduction

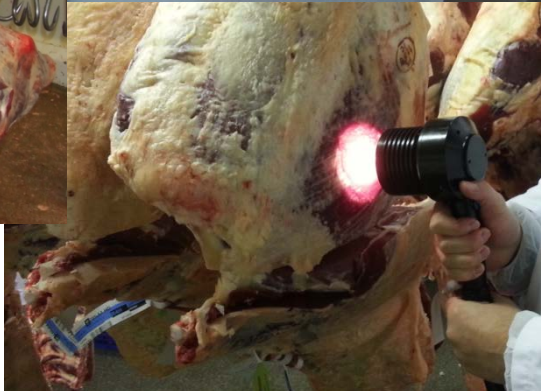
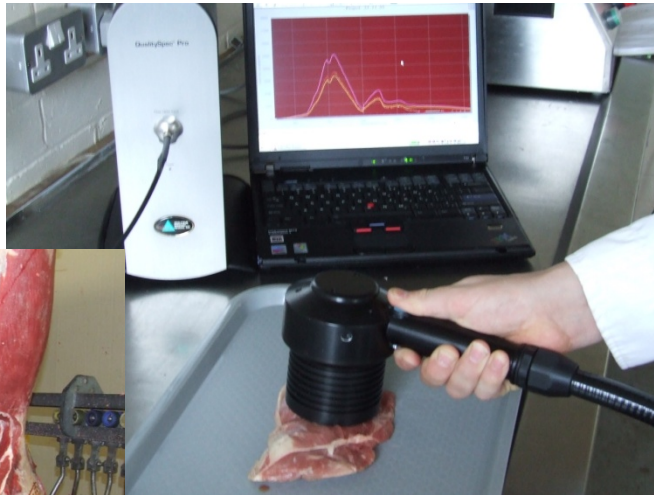
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- Few examples of commercial implementation
  - large scale progeny tests (NZ, Australia)
  - genomics
  - (SRUC “More taste, less waste” industry-led project)
- Accurate phenotypes are key
  - rapid, routine, non-destructive, non-invasive, cost-effective
  - Imaging technologies?



# Non-invasive *post-mortem* predictors

- Visible and Near Infra-red spectroscopy (VISNIR)



## Predicts:

- Colour
- Cooking loss
- Composition
- IMF; fatty acids
- Mechanical tenderness
- Sensory traits

## Pros:

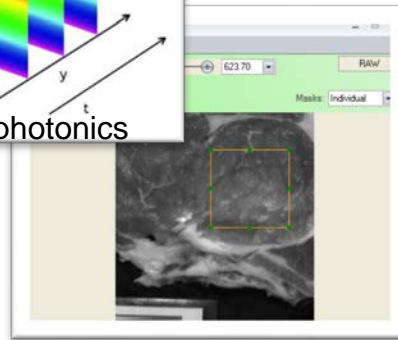
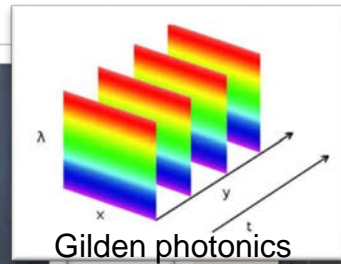
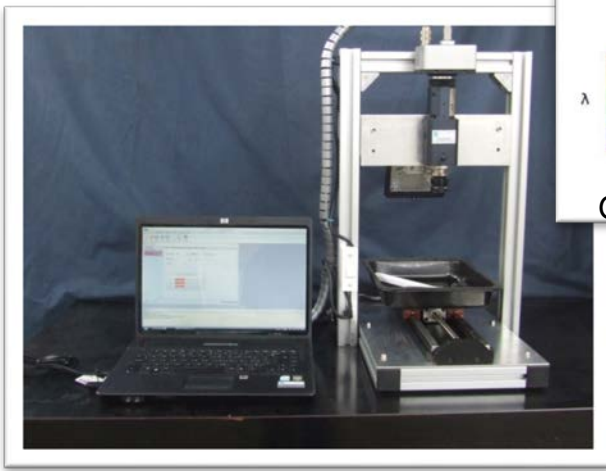
- Fast, non-invasive, cost-effective, on-line
- High  $R^2$  for colour & composition

## Cons:

- $R^2 \ll 1$  for technological/ sensory traits (Prieto et al., '09)
- predictions complex

# Non-invasive *post-mortem* predictors

- Hyperspectral imaging



## Predict:

- Colour
- Cooking loss
- Mechanical tenderness
- Composition; IMF
- Fatty acid composition
- Sensory traits

- Raman spectroscopy



## Pros:

- non-invasive, cost-effective
- wealth of information
- $R^2 > 0.8$  for several traits<sup>1</sup>

## Cons:

- practicality in plant
- predictions complex
- price?

<sup>1</sup>review by Xiong et al., 2014

# Non-invasive *post-mortem* predictors

- X-ray computed tomography (CT)



## Predicts:

- IMF
  - beef ( $R^2=0.71-0.76$ )<sup>1</sup>
  - pork ( $R^2 = 0.63-0.83$ )<sup>2</sup>
  - lamb ( $R^2= 0.36$ )<sup>3</sup>
- fatty acid profile ( $R^2=0.61-0.75$ )<sup>1</sup>
- low accuracy for tenderness and sensory traits

## Pros:

- fast; non-invasive; packaged meat
- simultaneously predicts composition

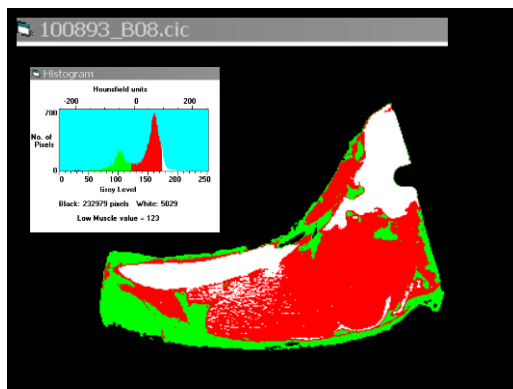
## Cons:

- $R^2 \ll 1$
- portability
- price

<sup>1</sup>Prieto et al., 2010

<sup>2</sup>Font-i-Furnols et al., 2013

<sup>3</sup>Lambe et al., 2009

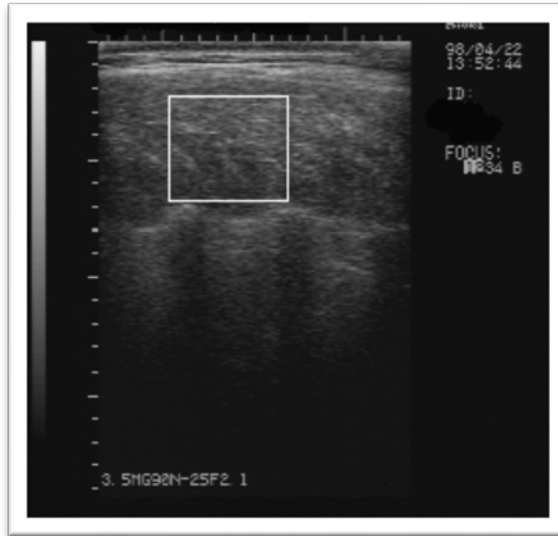


# Non-invasive *in-vivo* predictors

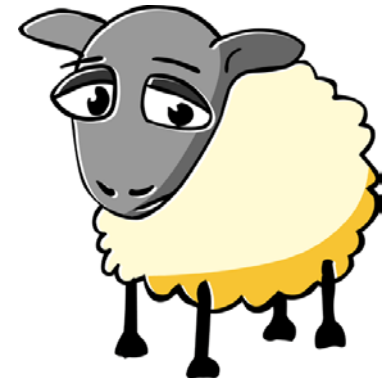
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- Ultrasound

- predicts IMF in pigs and beef cattle with mod-high accuracy (Newcom et al. '02; Aass et al., '06,'09)



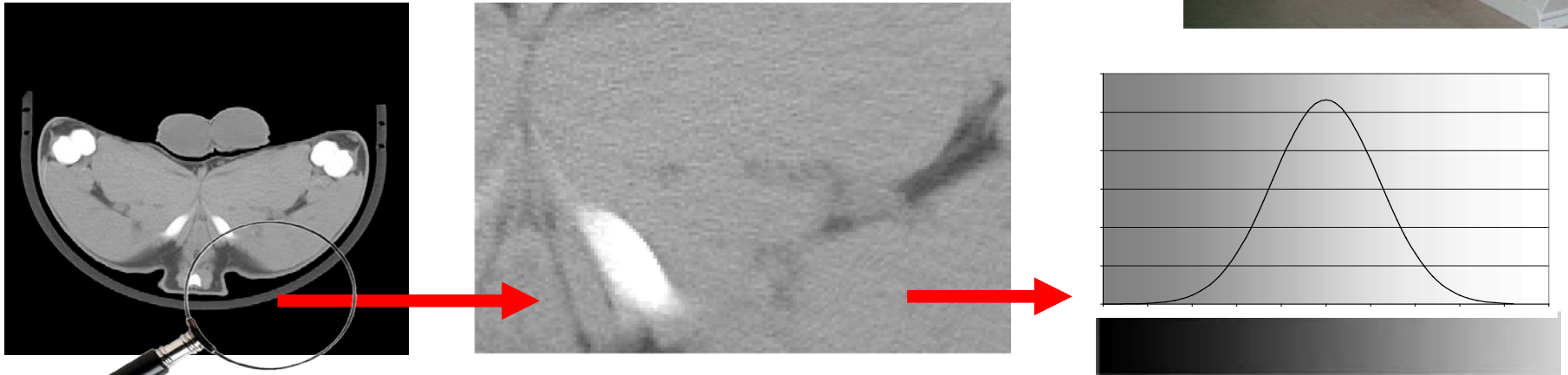
- not successful in sheep





# Non-invasive *in-vivo* predictors

- X-ray computed tomography (CT)



- CT tissue density distributions reflect IMF levels in live lambs ( $R^2 > 0.6$ )
- Does not accurately predict mechanical tenderness or taste panel traits



# Previous research: lamb IMF vs MQ

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- Acceptable levels for IMF (loin)
  - > 2-3% grilled red meat / lamb<sup>1</sup>
  - > 5% for “better than every day” eating quality<sup>2</sup>
  - SRUC slaughter lamb mean IMF:
    - Texel 1.4-1.6%
    - Texel X Mule 2.2%
    - Scottish Blackface 2.3%
- Concerns about fat reduction for eating quality



<sup>1</sup>Savell and Cross, 1988; Heylen et al., 1998; <sup>2</sup> Hopkins et al., 2006

# Genetic control of CT-IMF



- Data set from UK terminal sire breeding programme
  - ~2000 Texel ram lambs over 12 years
  - CT and performance records:  
2-stage selection for carcass composition
- Genetic analysis of CT-predicted IMF (ASReml):
  - heritability = 0.31 (s.e. 0.07)
  - genetic correlation with total carcass fat = 0.68 (s.e. 0.08)

# More taste, less waste

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Industry led research project with SRUC as lead research partner



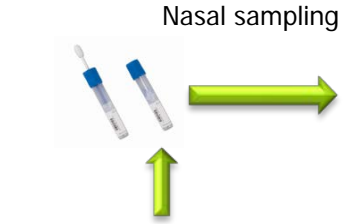
# More taste, less waste project



Terminal sire rams  
CT scanned



Mated to Mule ewes



Nasal sampling



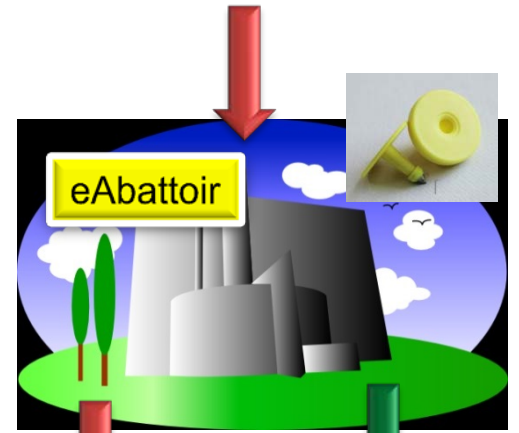
N= 5000  
crossbred  
lambs



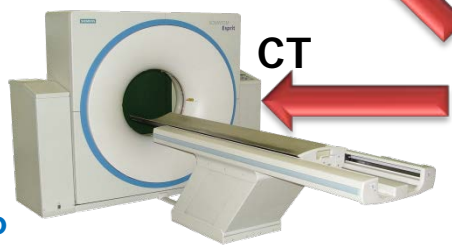
EBVs based on  
meat and carcass quality  
of crossbred lambs



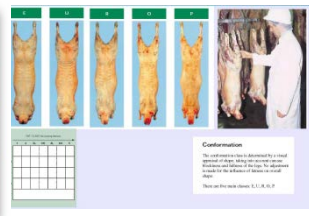
VISNIR



eAbattoir



CT



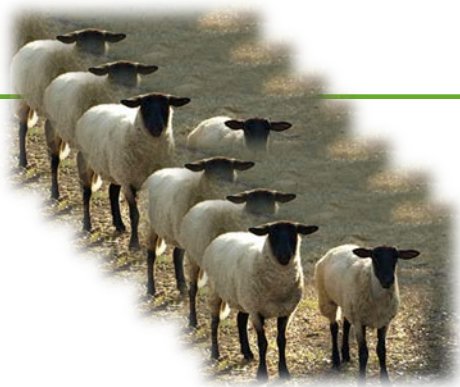
Loin info

Carcass info

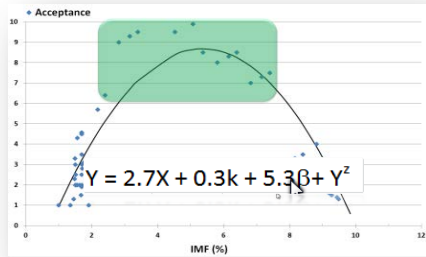
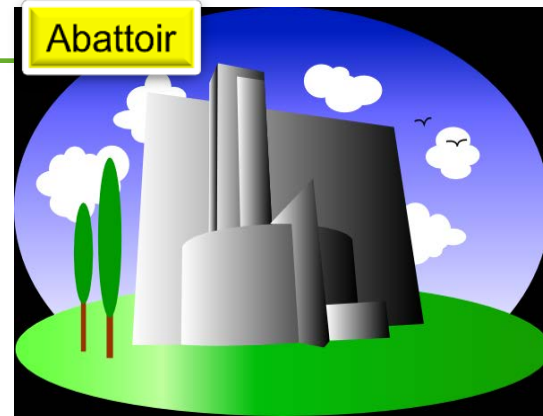


SRUC/EGENES

# More taste, less waste project – WP1



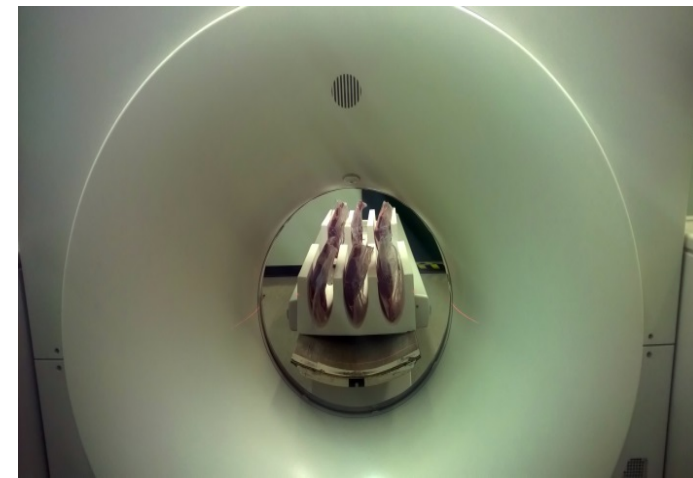
N= 300 across all specs



Prediction equations & correlations;  
windows of acceptability



# (Multiplex) CT to predict meat quality in lamb meat cuts



Trait	Accuracy of prediction (R <sup>2</sup> )
IMF	0.36
Shear Force	0.03
Texture (TP)	0.08
Flavour (TP)	0.09
Juiciness (TP)	0.06
Liking (TP)	0.10

Prediction equations combining CT traits and weights of loin & carcass

Best single CT predictor of all traits = % fat in sample (estimated by CT)

IMF % band	CT-predicted IMF converted to % band				Total
	1-2%	2-3%	3-4%	4-5%	
1-2%	5	17			22
2-3%	3	70	55		128
3-4%		34	83	1	118
4-5%		2	23	1	26
>5%		1	2		3
<b>Total</b>	8	124	163	2	297

54% samples – band correct

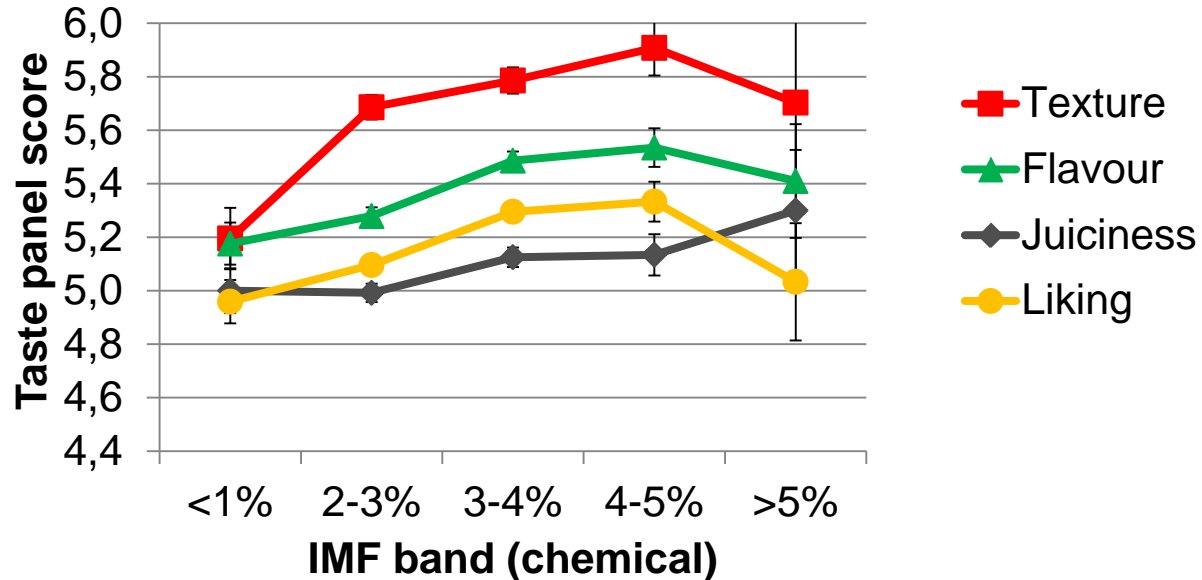
63% samples with IMF < 3% = < CT band 3-4%

25% samples with IMF > 3% = < CT 3-4%

# IMF influences sensory traits

Sensory traits significantly affected by IMF level:

- Assessed by chemical IMF extraction

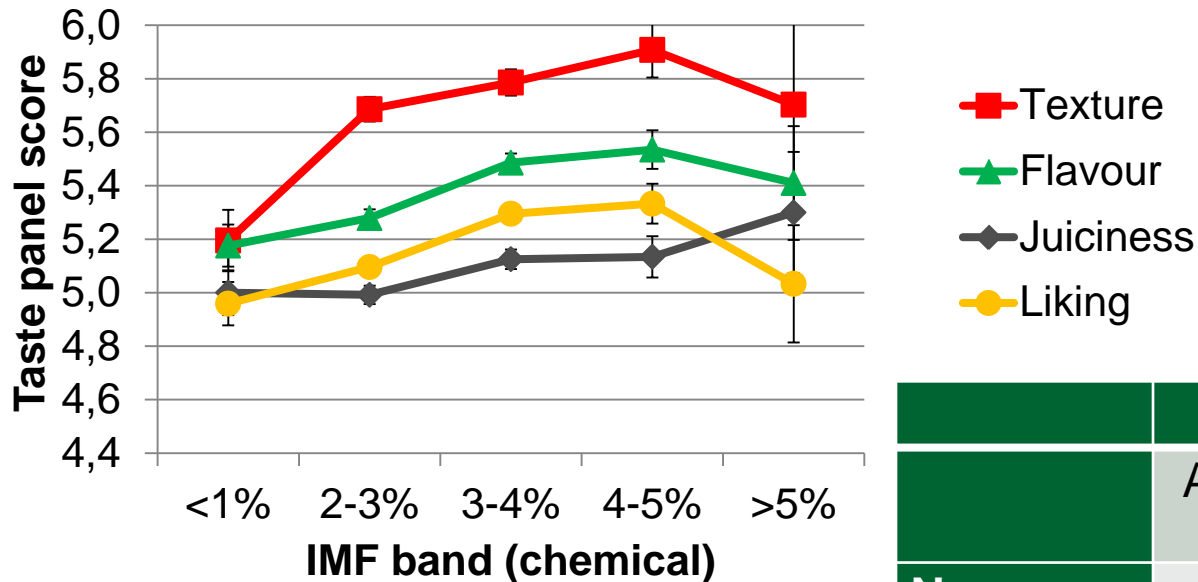




# IMF influences sensory traits

Sensory traits significantly affected by IMF level:

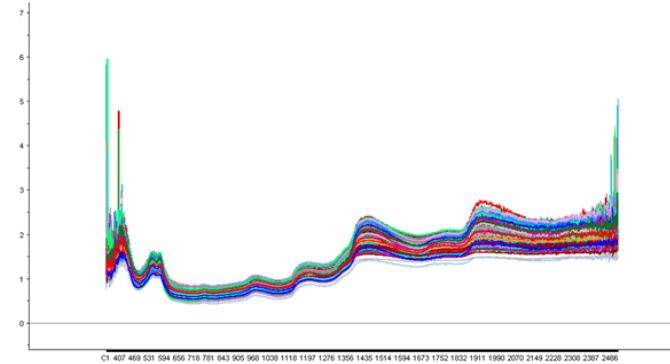
- Assessed by chemical IMF extraction OR predicted by CT



	CT-predicted IMF band			
	Adj-R <sup>2</sup>	<3%	>3%	P value
<b>N</b>		132	165	
<b>Texture</b>	7.0	5.55	5.85	<0.001
<b>Flavour</b>	3.8	5.29	5.45	<0.001
<b>Juiciness</b>	4.4	4.98	5.15	<0.001
<b>Liking</b>	5.7	5.08	5.28	<0.001

# VISNIR to predict MQ in lamb meat cuts

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- Spectra from 500-2400 nm used in analysis
- Median spectra of 10 replicates used
- Unscrambler (v10.3) multivariate analysis software

# VISNIR to predict MQ in lamb meat cuts

	Unpackaged		Vacuum-packed	
	$R^2_{Cal}$	$R^2_{Val}$	$R^2_{Cal}$	$R^2_{Val}$
<b>IMF</b>	0.35	0.23	0.23	0.18
<b>ShF</b>	0.03	0.01	0.11	0.04
<b>Texture</b>	0.03	0.01	0.07	0.06
<b>Flavour</b>	0.02	0.01	0.05	0.02
<b>Juiciness</b>	0.01	0.01	0.01	0.003
<b>Overall liking</b>	0.008	NA	0.001	NA

<sup>3</sup> $R^2_{Cal}$ =Coefficient of determination of calibration.

<sup>4</sup> $R^2_{Val}$ =Coefficient of determination of validation.

# Discussion – More taste, less waste

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- Can we increase accuracies to predict IMF *post-mortem*?
  - VISNIR on fresh cut meat; analysis method
  - CT on whole carcasses
- Project has produced:
  - high accuracy *in-vivo* phenotypes for IMF
  - moderate accuracy *post-mortem* phenotypes for IMF
  - data set to develop SNP-keys for genomic selection
- A combination of *in-vivo*, *post-mortem* and genomic predictors could be used to develop a sustainable breeding programme including lamb meat quality traits

# General discussion

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- Clear breeding goals required
  - MQ and other traits – multi-trait selection index
  - genomic selection + phenotyping
- Need to overcome the barriers to practical implementation and routine phenotyping
- Move from R&D to commercial implementation



# Acknowledgements



Supportive funding of the “More taste, less waste” project came from Innovate UK

Innovate UK  
Technology Strategy Board

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- Ian Richardson and team, University of Bristol



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