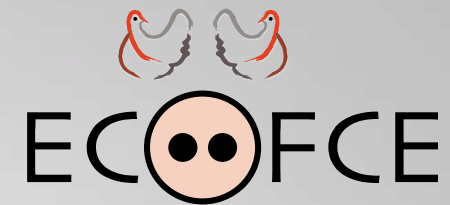


EFFICIENT & ECOLOGICALLY-FRIENDLY PIG AND POULTRY PRODUCTION.



A WHOLE-SYSTEMS APPROACH TO OPTIMISING FEED EFFICIENCY
AND REDUCING THE ECOLOGICAL FOOTPRINT OF MONOGASTRICS.



BASIC DATA

Funding:

EU-FP7
(€ 6 million)

Start date:

1 February 2013

Duration:

48 months
(2013 to 2017)



Optimising feed efficiency and reducing the ecological footprint of monogastrics (ECO-FCE)

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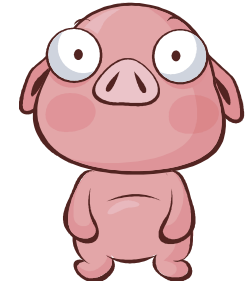
⁴University of Veterinary Medicine, Vienna, Austria;

⁵Hermitage Genetics, Kilkenny

ECO-FCE overview



- Feb 2013 – Feb 2017
- 17 partners
- Overall objectives
 - improve food security by optimising feed efficiency (FE) in pigs without negatively impacting on animal welfare or meat quality
 - reduce the ecological footprint of pig production
- Teagasc objective
 - to examine gut structure, function and microbiota in pigs divergent for FE



Expected benefits



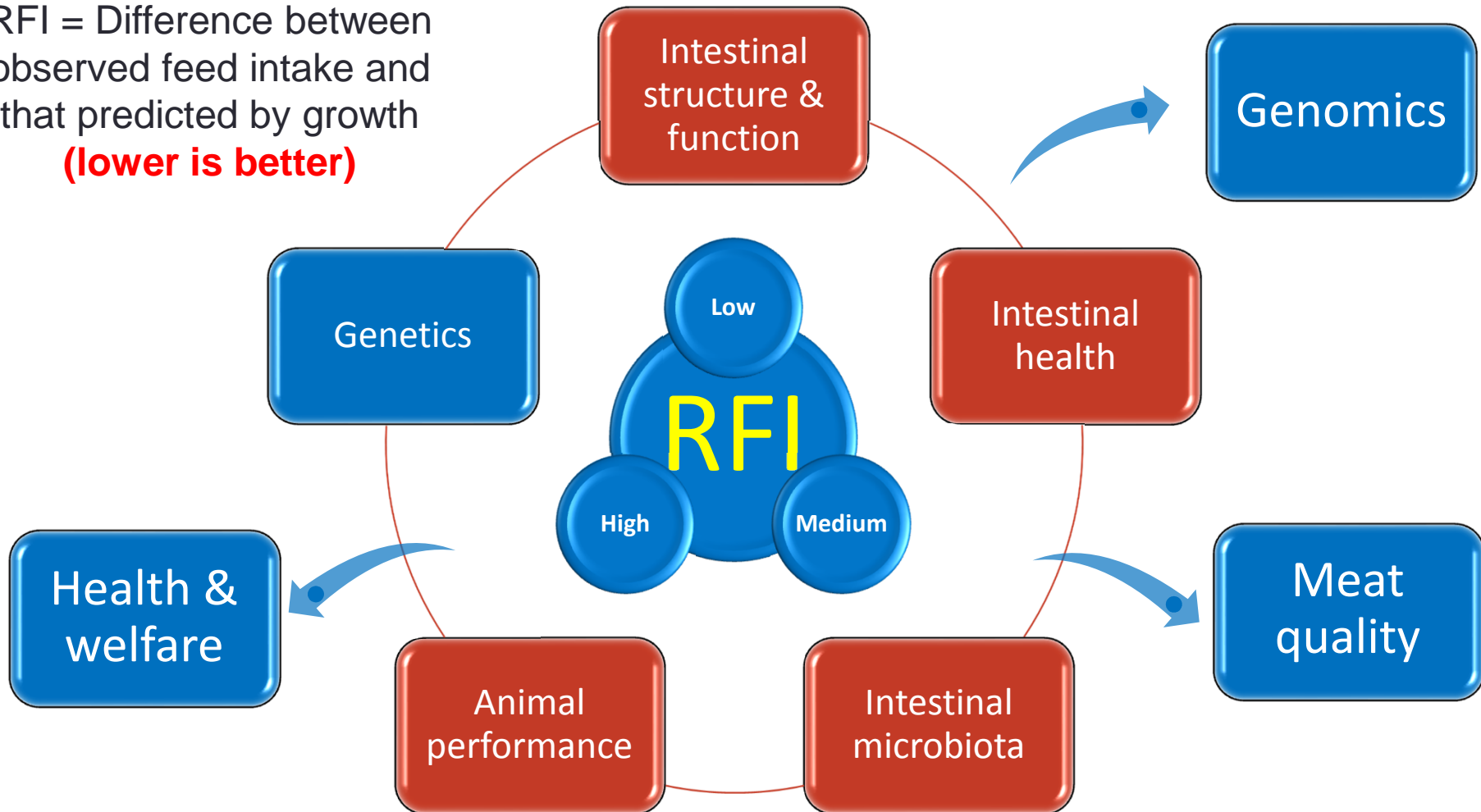
- **Improved FE (microbiota, feed additives & genetics)**
 - ↓ feeding costs
 - ↓ emissions
 - improve meat quality
 - improve animal welfare
- **Improved intestinal microbial profile**
 - ↓ costs associated with sick animals (growth check, medication, housing, labour, etc.)
 - better digestion and nutrient uptake
- **Implementation of an Internet ECO-FCE hub**
 - factors affecting feed efficiency
 - ecological calculators



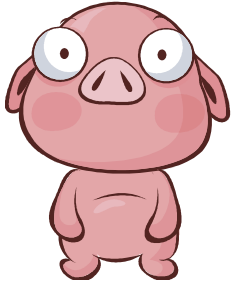
ECO-FCE overview



RFI = Difference between observed feed intake and that predicted by growth
(lower is better)

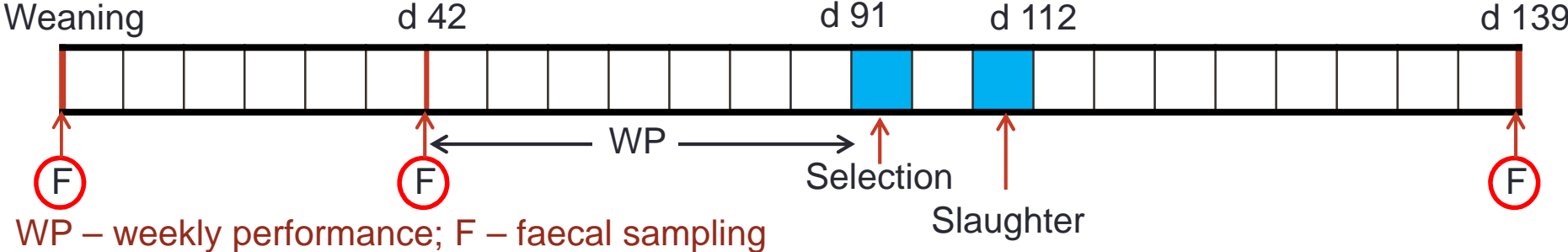
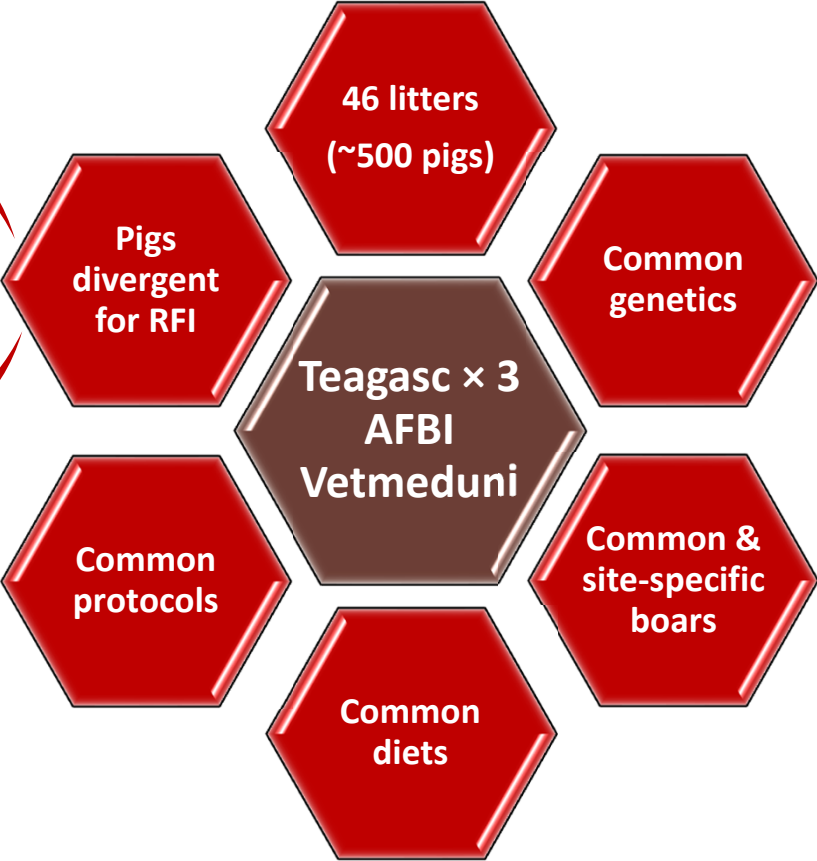


Pig studies



Selection within litter & gender

More accurate FE indicator

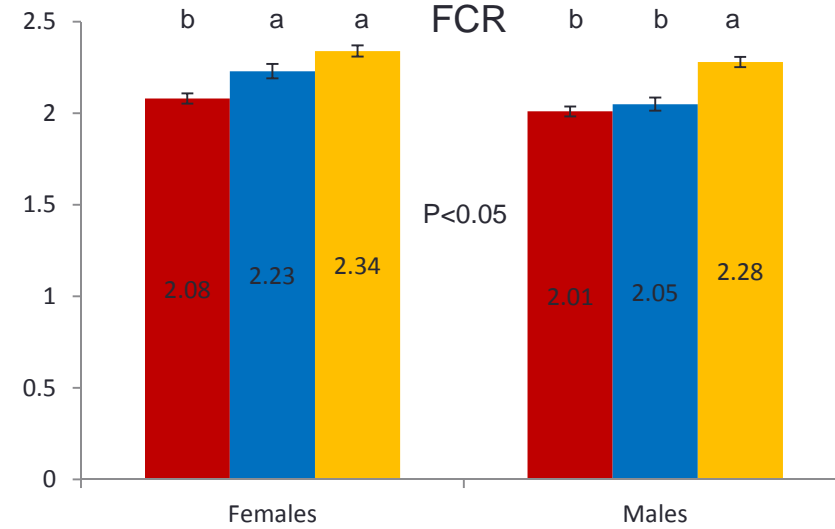
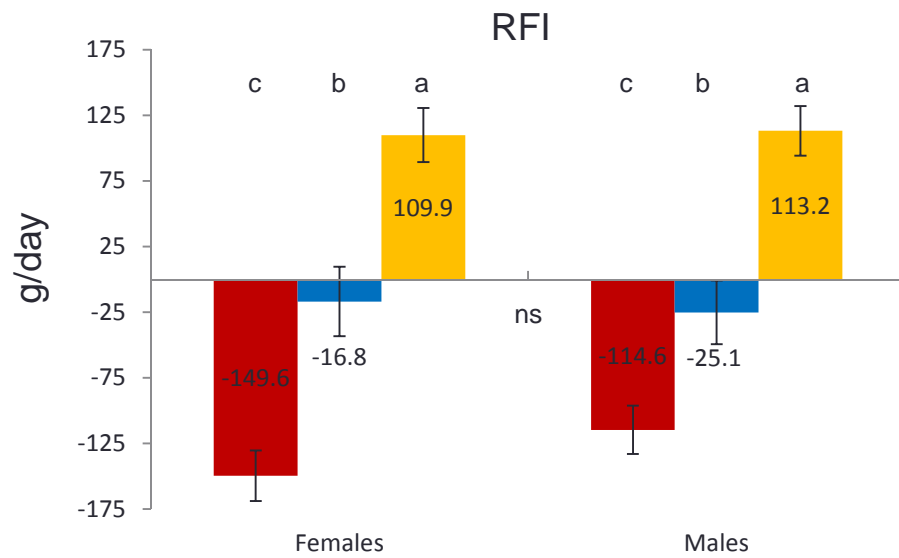




Results



Growth performance



Feed efficiency

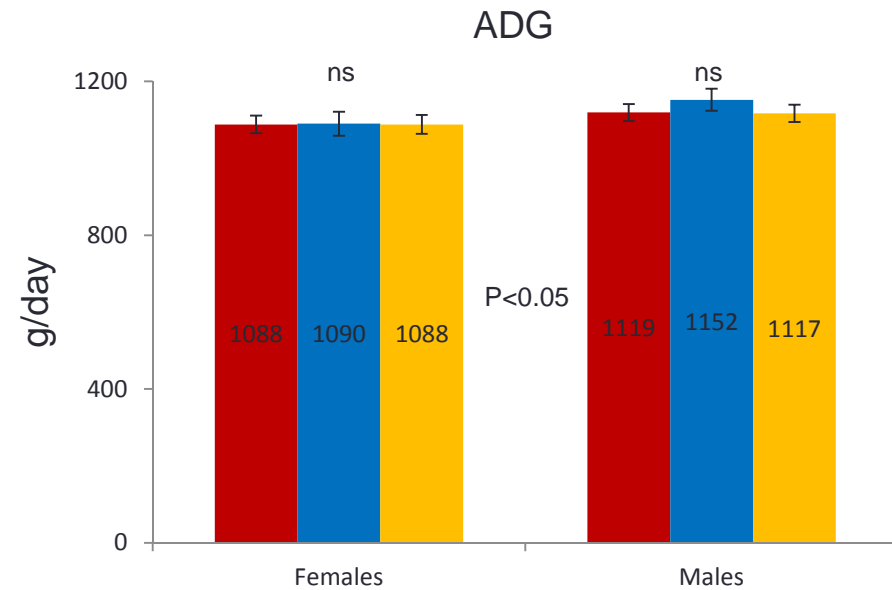
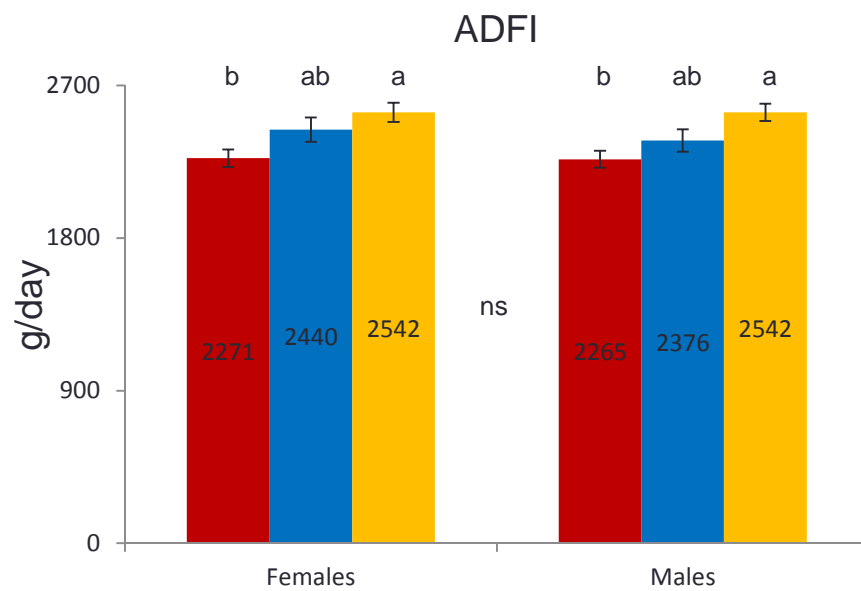
■ Good ■ Medium ■ Poor

a,b,c $P \leq 0.05$; ns – $P > 0.05$;

This project has received funding from the European Union's Seventh Framework Programme for research, technological development and demonstration under grant agreement No. 311794.



Growth performance



Feed efficiency

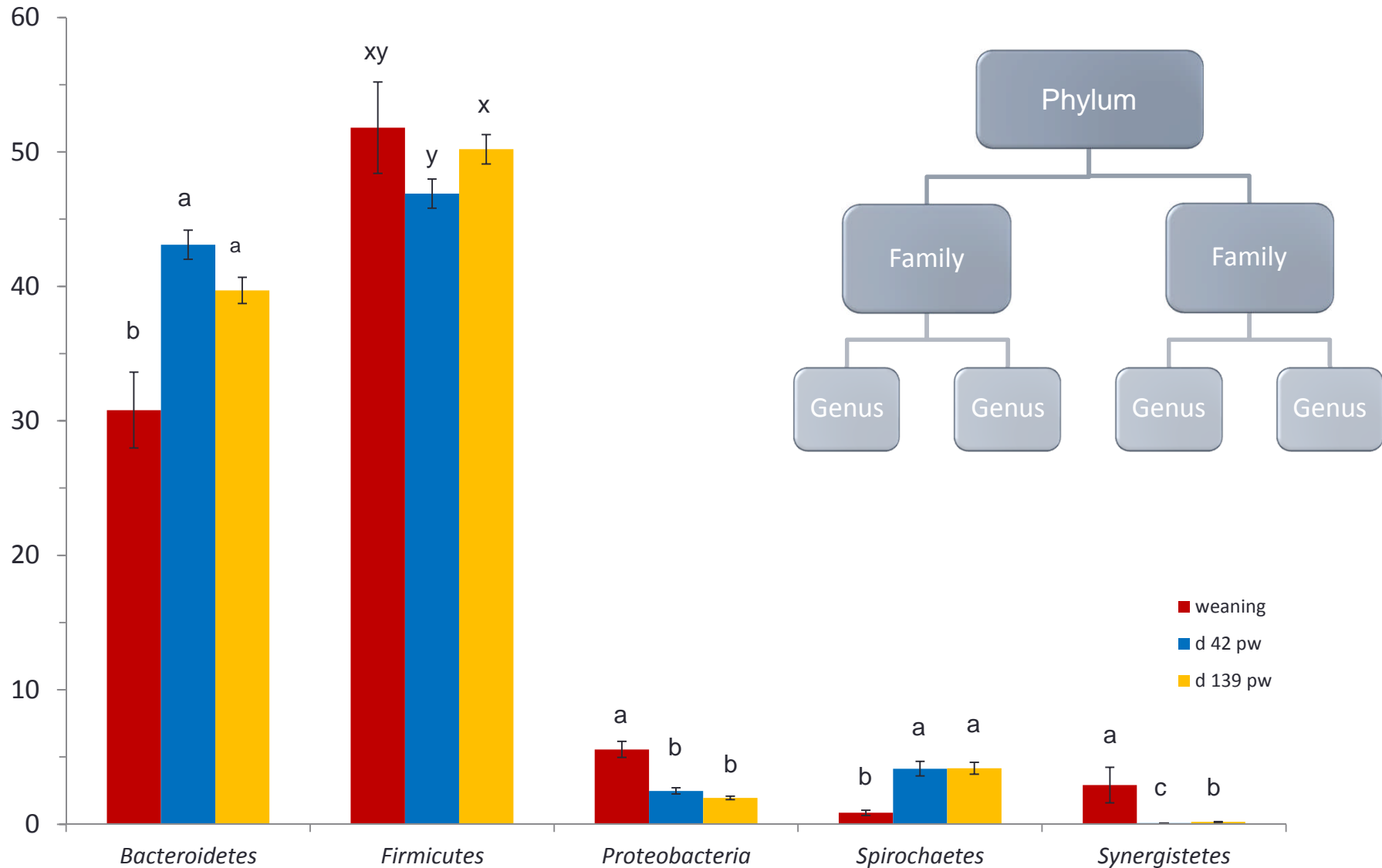
■ Good ■ Medium ■ Poor

a,b,c $P \leq 0.05$; ns – $P > 0.05$;

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Phylum – level differences in faecal microbiota over time



Columns within a group that don't share a common letter are significantly different ($a,b,c P \leq 0.05$; $x,y 0.05 < P < 0.10$)
 Standard errors represented by error bars

Genus – level differences in faecal microbiota

Taxon	Feed efficiency			95% CI/5-95 th percentiles	P value	Role in the intestine
	Good	Medium	Poor			
<i>Streptococcus</i>	0.80 ^b	1.00 ^{ab}	2.09 ^a	0.470-3.077	0.01	Some species pathogenic
<i>Campylobacter</i>	0.63 ^x	0.40 ^{xy}	0.36 ^y	0.236-0.812	0.04	Potentially pathogenic
<i>Pseudobutyrvibrio</i>	0.09 ^y	0.09 ^y	0.17 ^x	0.061-0.225	0.03	Beneficial
<i>Adlercreutzia</i>	0.000021 ^a	0.000017 ^a	0.000004 ^b	0-0.000123	0.04	Beneficial

^{a,b} $P \leq 0.05$; ^{x,y} $0.05 < P < 0.10$; Values within a row with a different superscript are different ($P < 0.05$).

Better FE = better intestinal microbial profile?

- relevance to be investigated further
- analysis of additional samples on-going - **confirmation**



Conclusions



- RFI ranking
 - differences in ADFI and FCR
 - gender differences for ADG and FCR
 - requires extensive data manipulation
 - **FCR may be more practical in most circumstances**
 - **RFI – more useful than FCR to standardise FE ranking across sites**
- Major effects of **geographical site & boar** on growth performance
 - despite diets being formulated to common specifications
- Intestinal microbiota
 - 4/333 taxa differed by feed efficiency rank
 - functionality to be investigated further
 - major litter and time effects



See poster



Acknowledgements



- Pig Development Department farm staff
- Teagasc Walsh Fellowship Scheme
- EU 7th Framework Programme
- Waterford Institute of Technology



Waterford Institute of Technology

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Thank you

