

# Comparative proteomic profiling of two muscles from five divergent pig breeds using SELDI-TOF proteomics technology

*N. Mach*



ANIMAL SCIENCES GROUP  
WAGENINGEN UR

# Comparative proteomic profiling of two muscles from five divergent pig breeds using SELDI-TOF proteomics technology

## INTRODUCTION



1. SELDI-TOF increases understanding of the role of proteins expression in pig meat quality.
2. There is a lack of biomarkers to screen for divergent pig breeds and muscle types



# Comparative proteomic profiling of two muscles from five divergent pig breeds using SELDI-TOF proteomics technology

## OBJECTIVE



The specific objective of the present study was:

To identify biomarkers and to evaluate the influence of different breed types and muscle type on the expression of muscle proteins, as well as their interactions, using SELDI-TOF-MS techniques



# Comparative proteomic profiling of two muscles from five divergent pig breeds using SELDI-TOF proteomics technology

MATERIALS

AND

METHODS



126 pig males subjected to a 2 x 5 factorial design  
(2 muscles x 5 pure and divergent breeds)

Longissimus dorsi (LD)

Semimembranosus (SM)

43 Landrace

21 Duroc

43 Large White

13 Pietrain

6 Belgium Landrace



ANIMAL SCIENCES GROUP  
WAGENINGEN UR



# Comparative proteomic profiling of two muscles from five divergent pig breeds using SELDI-TOF proteomics technology

## MATERIALS AND METHODS



### --Protein expression-----

Isolation of water soluble fraction proteins

SELDI-TOF analysis

### --Statistic model -----

$$Y_{ijklmno} = \mu + A_i + B_j + M_h + (B \times M)_{jk} + \varepsilon_{ijh}$$

*A= pig; B= Breed type; M= Muscle type*



ANIMAL SCIENCES GROUP  
WAGENINGEN UR

RESULTS  
AND  
DISCUSSION



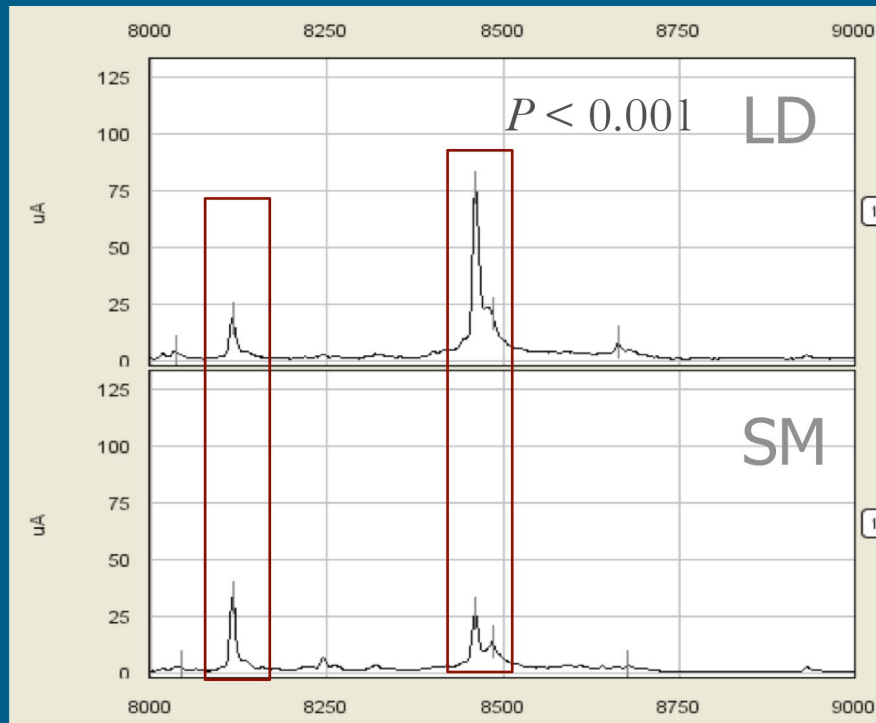
✓ The number of peaks per spectra ( $36.33 \pm 0.87$ ) was not affected by breed and muscle type.

✓ The intensity of peaks was strongly affected by breed, muscle type, as well as the interaction between them.



# Comparative proteomic profiling of two muscles from five divergent pig breeds using SELDI-TOF proteomics technology

## Muscle biomarkers intensity ( $\mu\text{A}$ )



Peak, <i>m/z</i>	LD ( $\mu\text{A}$ )	SM ( $\mu\text{A}$ )	SEM	<i>P</i> -Value
8,126	9.0	21.1	0.08	0.001
8,485	48.7	12.4	2.43	0.001
14,847	2.6	1.8	0.11	0.001

LD= *Longissimus dorsi* muscle

SM= *Semimembranosus* muscle



## Comparative proteomic profiling of two muscles from five divergent pig breeds using SELDI-TOF proteomics technology

### Breed biomarkers intensity ( $\mu\text{A}$ )

Peak ( $m/z$ )	DU	BL	LS	LW	PI	SEM	<i>P</i> -value
5,773	8.3	5.2	12.9	8.7	5.5	1.95	0.001
7,114	2.8	4.6	4.2	4.0	4.2	0.17	0.001

DU= duroc; BL = Belgian Landrace; LS= Landrace; LW= Large White; PI= Pietrain





## Comparative proteomic profiling of two muscles from five divergent pig breeds using SELDI-TOF proteomics technology

### Breed and muscle biomarkers intensity ( $\mu$ A)

Peak <i>m/z</i>	<i>Longissimus dorsi</i>					<i>Semimembranosus</i>					SEM	P- value
	DU	BL	LS	LW	PI	DU	BL	LS	LW	PI		
3,098	2.0	1.6	2.9	1.6	1.2	10.3	0.4	5.8	8.4	4.2	0.76	0.001
8,043	5.4	4.9	4.7	4.0	7.4	2.4	7.9	3.3	3.1	3.6	0.35	0.001

DU= duroc; BL = Belgian Landrace; LS= Landrace; LW= Large White; PI= Pietrain



## CONCLUSION



A total of 3 biomarkers to discriminate muscle type (the 8,126; 8,485 and 14,847  $m/z$ ), 2 to categorize breed types (5,773 and 7,114  $m/z$  peaks), and 2 to classify breed type according muscle type (3,098 and 8,043  $m/z$  peaks) were identified.

SELDI-TOF-MS is a highly efficient method for the identification of biomarkers and changes in muscle protein expression for divergent pure pig breeds and muscle type.



# Comparative proteomic profiling of two muscles from five divergent pig breeds using SELDI-TOF proteomics technology

## OBJECTIVE



## Acknowledgement

### Funding:

\*EU - Q-Porkchains

\*Dutch Agricultural Ministry Kennisbasis Funds

\*Departament d'Universitats, Recerca i Societat de la Informació de la Generalitat de Catalunya.



ANIMAL SCIENCES GROUP  
WAGENINGEN UR



Thank you

Questions



ANIMAL SCIENCES GROUP  
WAGENINGEN UR