



Determination of fatty acid groups in intramuscular fat of pig by FT-NIRS within the project TREASURE

Bozzi R., Parrini S., Pugliese C., Bonelli A., Gasparini S., Karolyi D., Martins J.M., García-Casco J. M., Panella N., Nieto R., Petig M., Izquierdo M., Razmaite V., Djurkin Kušec I., Araújo J.P., Candek-Potokar M., Lebret B.



EAAP 2018

69th Annual Meeting of the European Federation of Animal Science

Dubrovnik, Croatia, 27th to 31st August 2018



Funded by European Union
Horizon 2020
Grant agreement No 634476

Task 3.2: Evaluation of sensory, healthy, technological and typical attributes of regional pork products from untapped breeds and new products issued from local chains

developed by partners within national or European research projects. In particular the fresh meat and pork products submitted to chemical analyses will be utilized to perform NIRS analysis in order to create a regression equation and implement the NIRS methodology as new method to predict meat quality, with special emphasis on prediction of lipid content and fatty acid composition (UNIFI, KIS).

Quantitative chemical analyses used for the determination of FA are:

a. costly b. time consuming

NIRS

- rapid and non-destructive: neither requiring reagents nor producing waste
- widely used **in research** for large-scale **meat quality evaluation**
- already applied to study the FA content in Iberian pig fat, intact pork loins and grounded beef.



Animals	Lipid	Fatty Acid Fat	IMF	Fatty Acid lean
512	270	372	452	197

**Present research
considered 165
fresh samples of:**

Partner	Breed	Sampled
KIS	Krskopolje	5
CICYTEX	Iberico	15
UNIFI	Cinta Senese	20
LUHS	Lithuanian White	24
LUHS	Lithuanian Wattle	9
CSIC	Iberico	5
UNIZG	Turopolje	5
PFOS	Crna Slawonska	16
IRTA	Negre Mallorqui	5
BESH	Schwabish Hallisches	16
UEVORA	Alentejana	10
UEVORA	Bisara	9
IPVC	Bisara	16
INIA	Iberico	5
INRA/IFIP	Gascon	5
Total		165

12 breeds in 8 different countries:



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Samples collection: *Longissimus dorsi*



Slices



Grounded



FT- NIRs
spectra
acquisition



Analysis with reference methods

From 3999 a 9999
wavenumber (cm^{-1})

- Lipid were extracted from intramuscular samples
- Fatty Acid Profile through Gas Chromatography



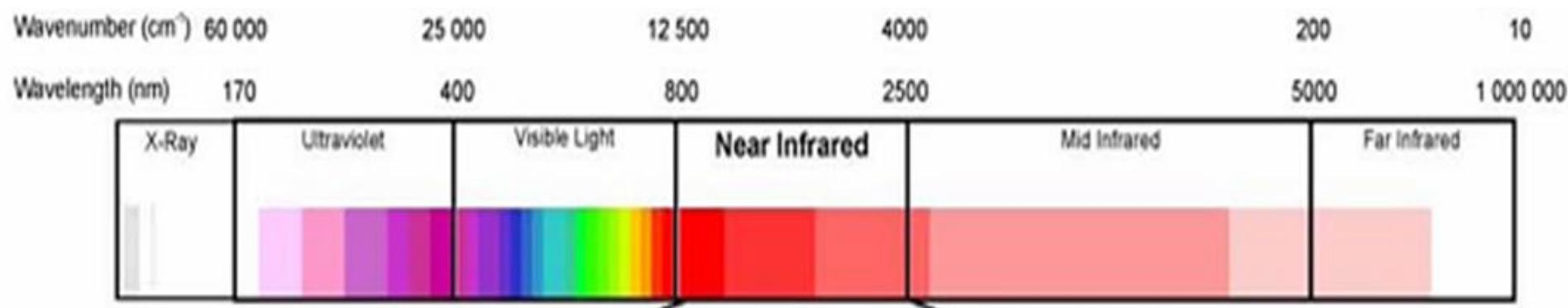
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Specific Aim



To evaluate the potential use of FT-NIRS for predicting:

- Intramuscular Fat IMF
 - fatty acids C18:1n9ct C18:3n3 C20:3 n6
 - FA groups SFA MUFA PUFA
 - n-6 PUFA n-3 PUFA n-4 PUFA n-1 PUFA
 - Ratios PUFA/SFA n-6/n-3



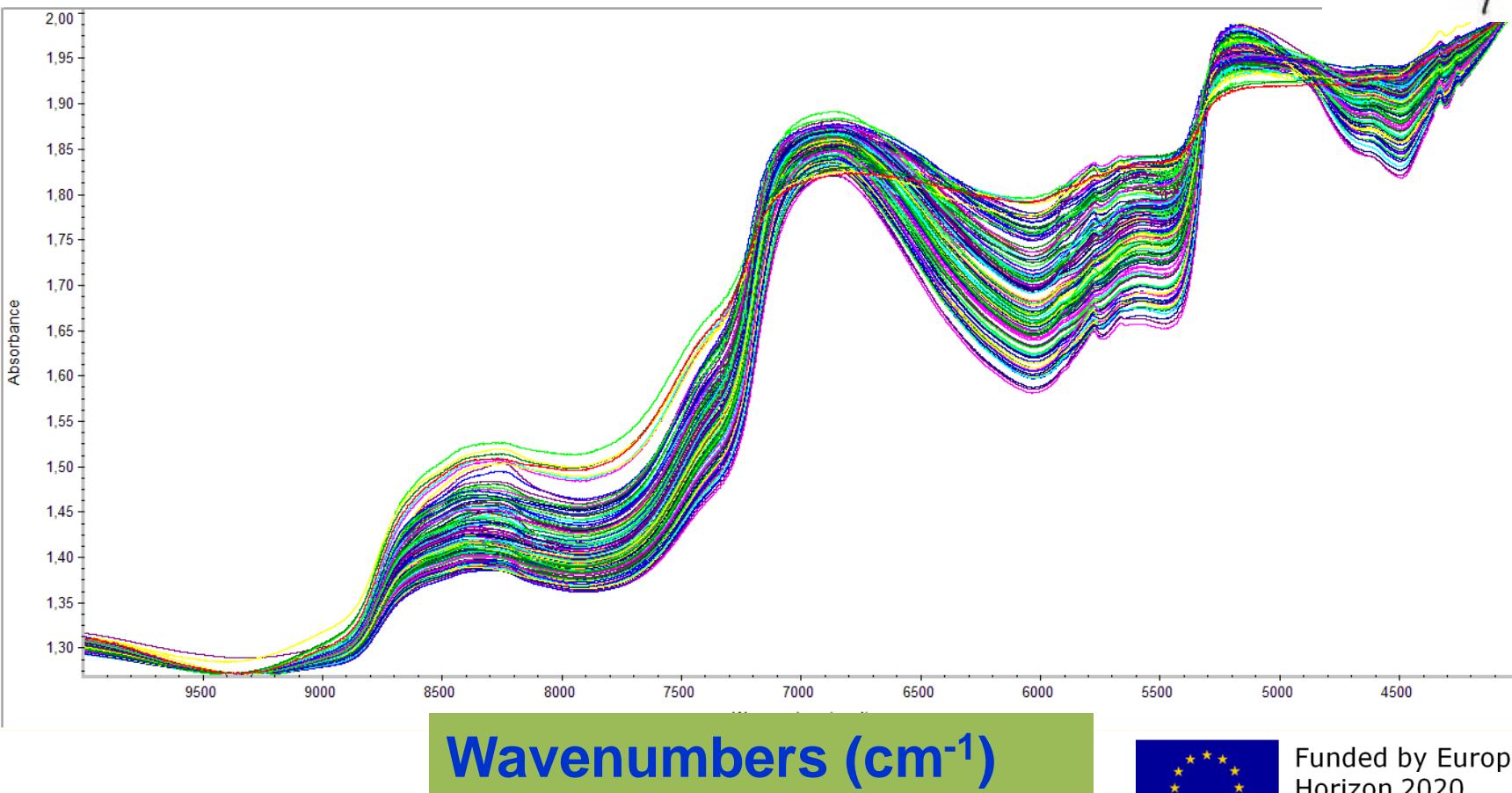
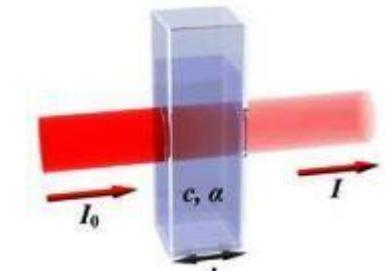
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Scan of samples



FT- NIRS (Antaris II):

- 2 scan for each sample (32 scans 16^{-1} cm)
- Absorbance mode



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Mathematical pre-treatments

- Multiplicative Scatter Correction
 - 1st and 2nd derivative
- Smoothing: Savitzky-Golay Filter
 - outliers' spectra

Calibration

Partial Least Square Regression
80% of total samples

Validation

20% of total samples
(randomly selected)

R^2

RMSEC

Coefficient of determination
Root mean square error

R_p^2

RMSECP



Intramuscular Fat and Fatty Acid composition (all units in %)

Parameter	n= 165	Mean	SD	Minimum	Maximum
IMF		4.30	2.33	0.56	14.15
SFA		38.71	2.44	31.87	45.61
MUFA		52.48	3.22	43.23	59.45
PUFA		8.26	2.72	3.31	19.50
n-6 PUFA		8.01	2.61	3.24	18.40
n-3 PUFA		0.26	0.14	0.07	1.10
n-4 PUFA		0.01	0.01	0.00	0.08
n-1 PUFA		0.20	0.14	0.03	1.04
C18:1n9ct		42.30	2.45	34.47	48.19
C18:3n3		0.31	0.13	0.11	0.78
C20:3n6		0.16	0.08	0.04	0.58
<i>Ratios</i>					
PUFA/SFA		0.22	0.08	0.08	0.55
n-6/n-3		35.23	11.84	16.69	61.35



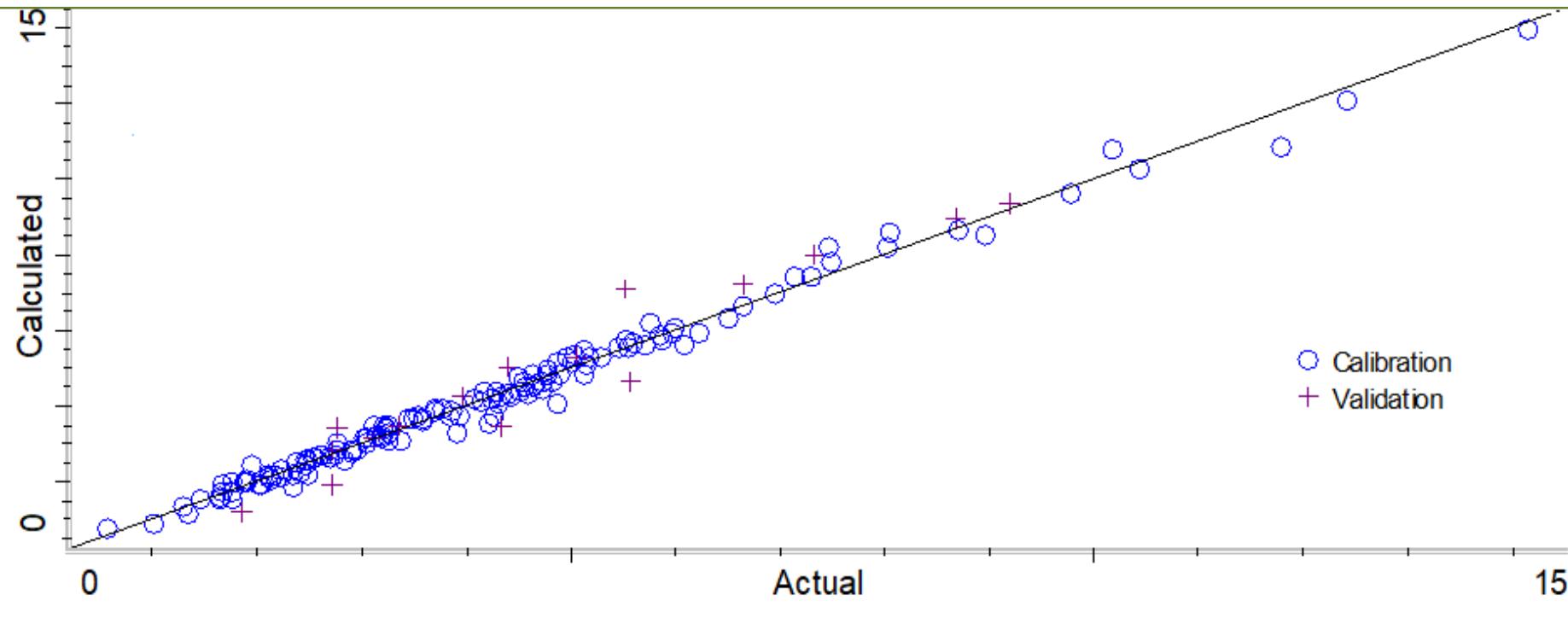
NIRS spectra of grounded meat



IMF

R^2_c 0.994
RMSEC 0.245

R^2_p **0.966**
RMSEP 0.664



Prevolnik Povše et al. 2017

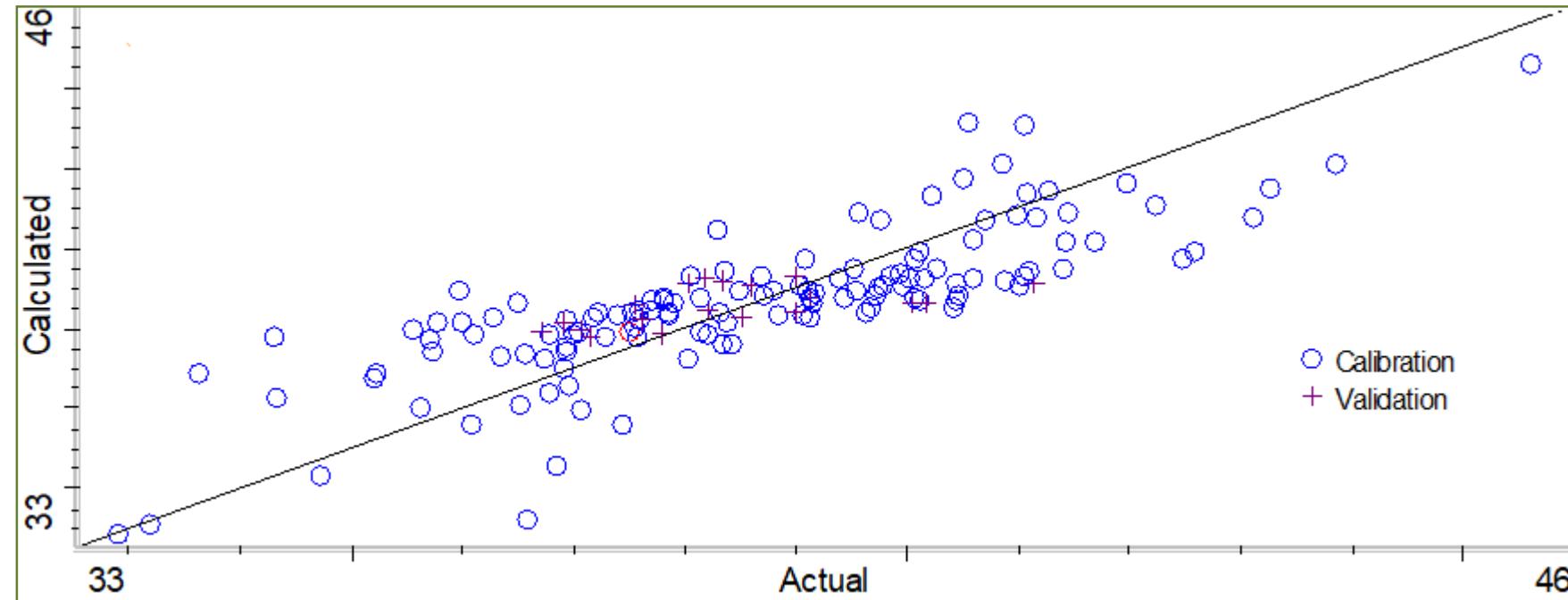
Pork fresh meat LD muscle ($n = 75$)

	R^2_c	SE_c	R^2_{cv}	SE_{cv}
IMF	0.99	0.14	0.95	0.25



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R^2_c **0.817**
 RMSEC 1.30

 R^2_p **0.771**
 RMSEP 0.975

 Gonzalez Martins et al. (2003) in **subcutaneous fat of Iberian pig**

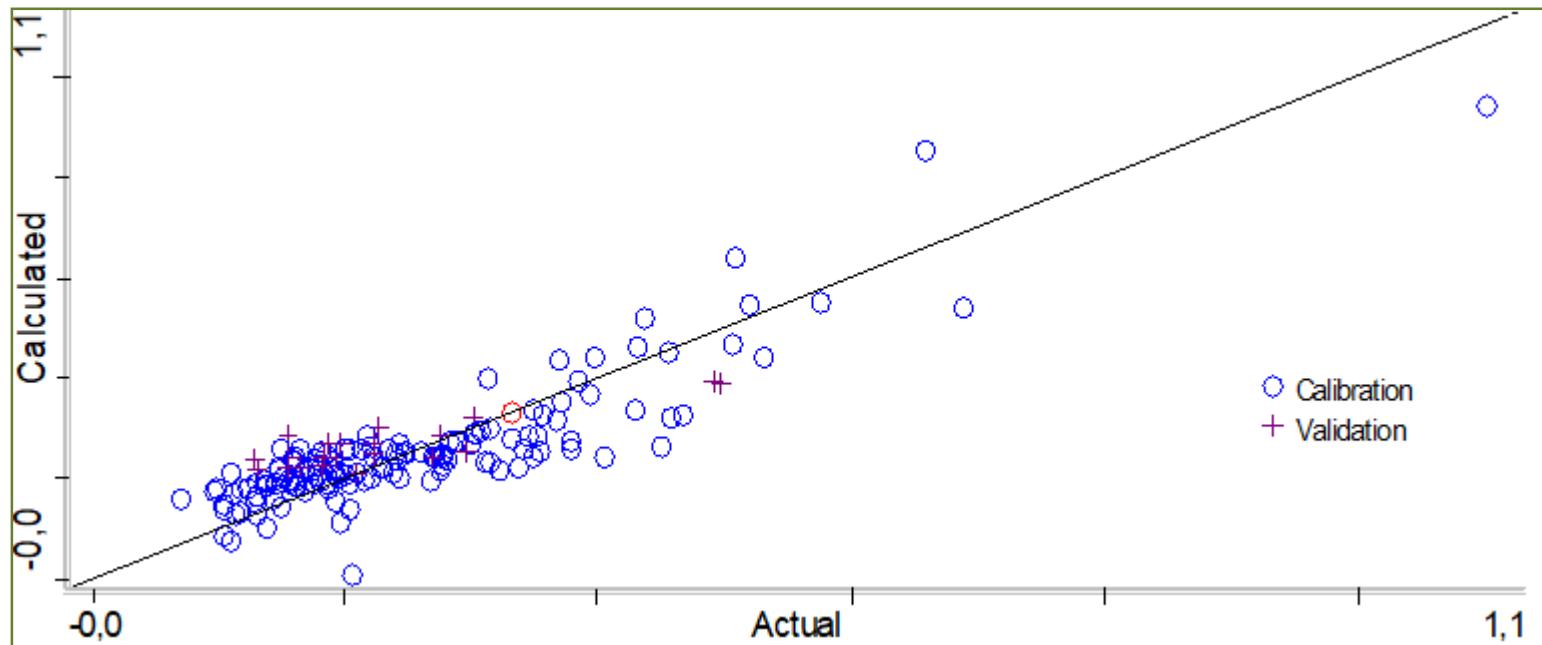
	RSQ	Se_c or SE_p	SE_{cv}	SEP_c	Range	n	N. PC
Saturated							
CALIBRATION	0.96	0.86	1.10	1.10	22.09-47.31	157	12
EXT. VALIDATION	0.90	1.18	1.18	1.18	29.26-40.75	23	


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PUFA

R^2_c 0.881
RMSEC 0.066

R^2_p 0.872
RMSEP 0.070



Gonzalez Martins et al. (2003) in subcutaneous fat of Iberian pig

	RSQ	Se_c or Se_p	SE_{cv}	SEP_c	Range	n	N. PC
Polyunsaturated							
CALIBRATION	0.95	0.47	0.60	0.60	2.31-14.82	157	12
EXT. VALIDATION	0.88	0.76	0.78	0.78	5.44-13.31	23	



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MUFA

R^2_c 0.894

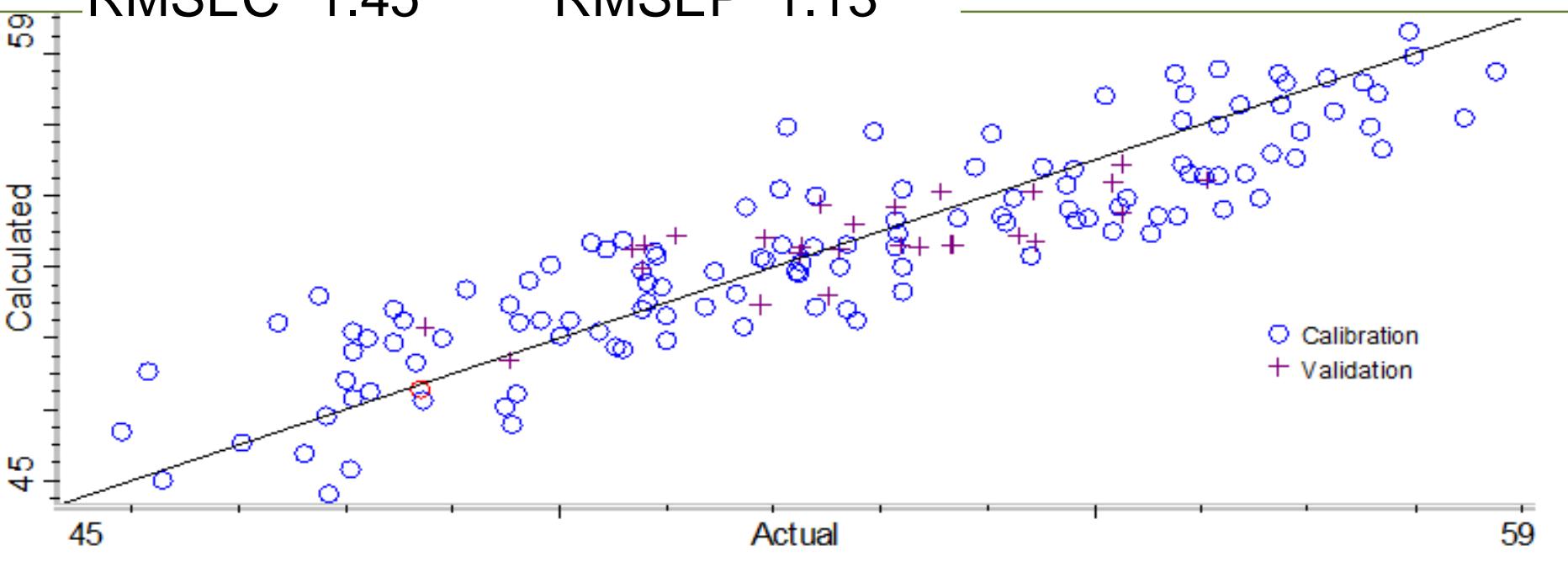
RMSEC 1.45

R^2_p 0.770

RMSEP 1.13



TREASURE



Gonzalez Martins et al. (2003) in subcutaneous fat of Iberian pig

	RSQ	Se_c or SE_p	SE_{CV}	SEP_c	Range	n	N. PC
Monounsaturated							
CALIBRATION	0.89	0.98	1.50	1.50	47.37-65.62	157	12
EXT. VALIDATION	0.74	1.21	1.23	1.23	51.43-60.70	23	



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PLS calibration and validation models

Trait	Calibration		Validation		Math treat
	R ² _c	RMSEC	R ² _p	RMSEP	
IMF	0.994	0.24	0.966	0.66	1;2
SFA	0.817	1.30	0.771	0.97	1;3;4
MUFA	0.894	1.45	0.770	1.13	1;3;4
PUFA	0.881	0.66	0.872	0.70	1;3

Math treatment:

- | | |
|--|---|
| 1. Multiplicative Scatter Correction
3. Second Derivative | 2. First Derivative
4. Savitzky-Golay filter |
|--|---|



PLS calibration and validation models

Trait	Calibration		Validation		Math treat
	R ² _c	RMSEC	R ² _p	RMSEP	
n-6 PUFA	0.754	1.43	0.694	1.52	1;3
C18:1 n9ct	0.768	1.46	0.687	1.19	1;3;4
<i>Ratios</i>					
n-6/n-3	0.824	6.77	0.801	5.94	1;3

Math treatment:

- 1. Multiplicative Scatter Correction
- 2. First Derivative
- 3. Second Derivative
- 4. Savitzky-Golay filter



PLS calibration and validation models

Trait	Calibration		Validation		Math treat
	R ² _c	RMSEC	R ² _p	RMSEP	
n-3 PUFA	0.756	0.07	0.701	0.06	1
n-4 PUFA	0.721	0.00	0.671	0.00	1;3;4
n-1 PUFA	0.834	0.06	0.648	0.05	1;2
C18:3 n3	0.812	0.07	0.768	0.07	1;2;4
C20:3 n6	0.740	0.05	0.721	0.04	1;2;4
<i>Ratios</i>					
PUFA/SFA	0.704	0.44	0.686	0.37	1;3

Math treatment:

- 1. Multiplicative Scatter Correction
- 2. First Derivative
- 3. Second Derivative
- 4. Savitzky-Golay filter



Resuming

- Developed model applicable for IMF quality control
- Promising results for FA groups
- Good results even when low proportion of FA
- Single fatty acid even if highly represented gave poor result
- Mathematical pre-treatments needed
- Possible application for Nutritional Label
- Additional work expected





Thanks for your attention



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