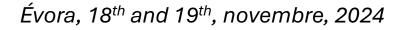


España - Portugal



Desarrollo de modelos predictivos mediante tecnología NIRS

Tecnologías avanzadas, innovadoras y digitales para el sector agroalimentario de la EUROACE





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Degree in food science and technology Meat Quality Area of CICYTEX <u>lucia.leon@juntaex.es</u>



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Index

- What is NIRS technology?
- The Unscrambler
 - The construction of the matrix
 - Quantitative models
 - Qualitative models
 - Pre-treatments



Index

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 - Quantitative models
 - Qualitative models
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What is NIRS technology?



What is NIRS technology?

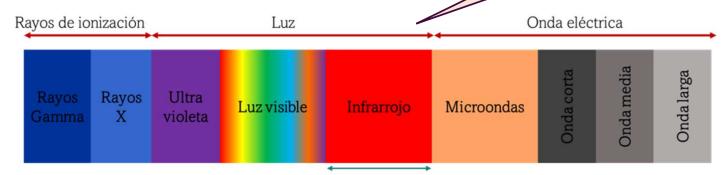
Near
InfraRed
Spectroscopy



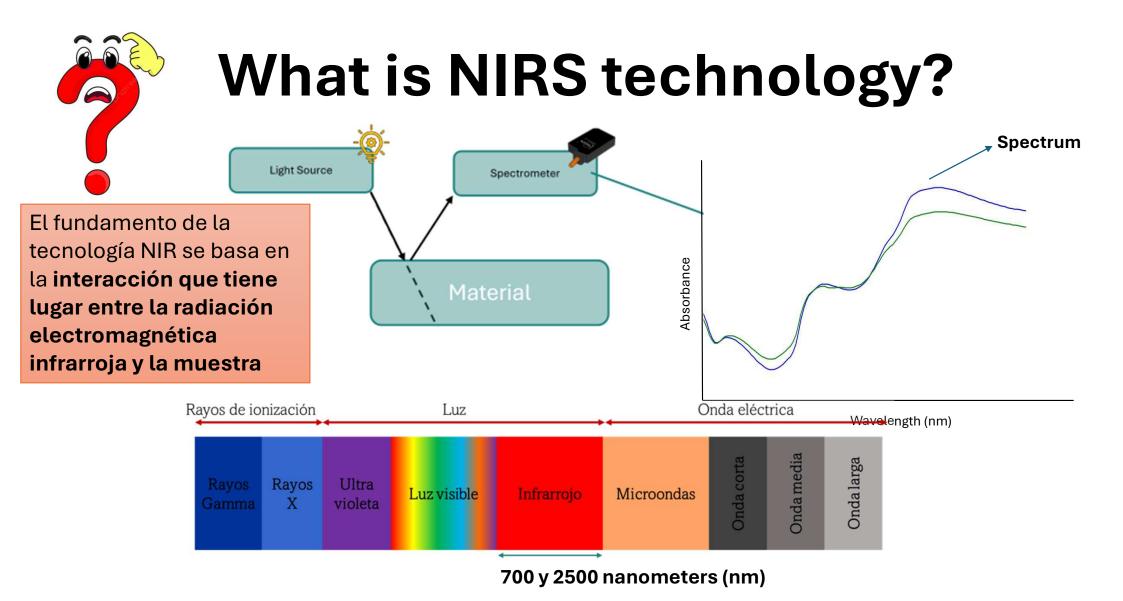
What is NIRS technology?

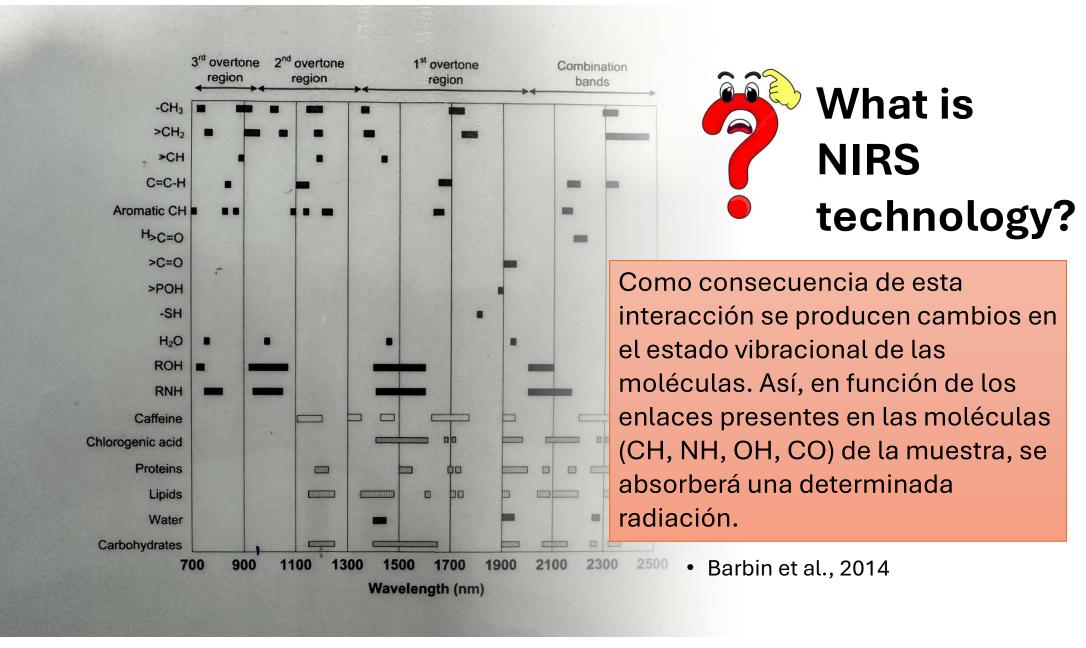
Near
InfraRed
Spectroscopy

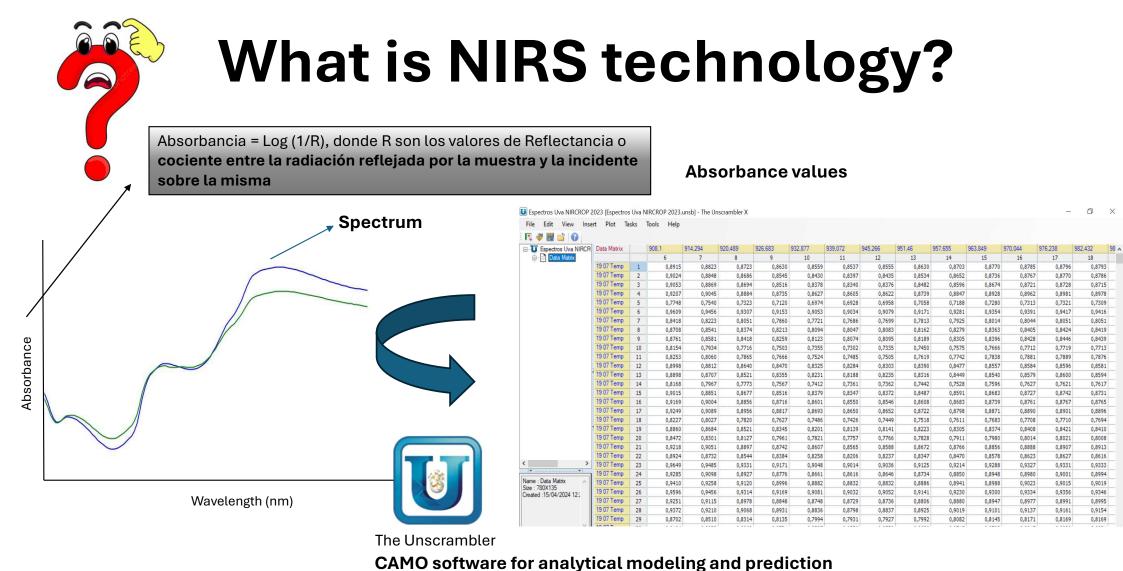
Near Infrared Spectroscopy (NIR) is an **analytical technique** that uses the **near infrared region** of the electromagnetic spectrum.

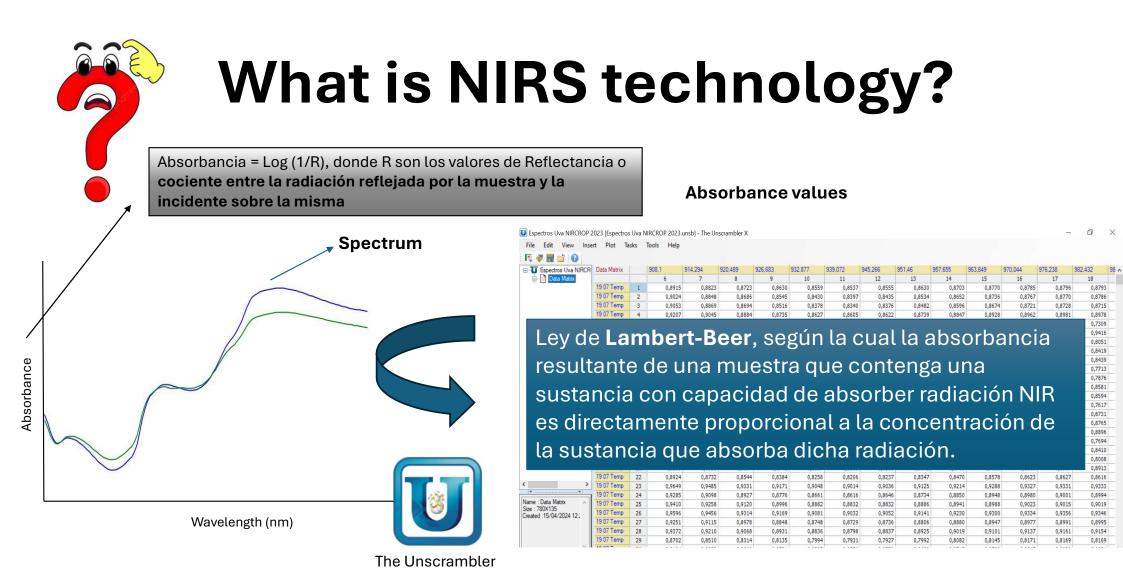


700 y 2500 nanometers (nm)

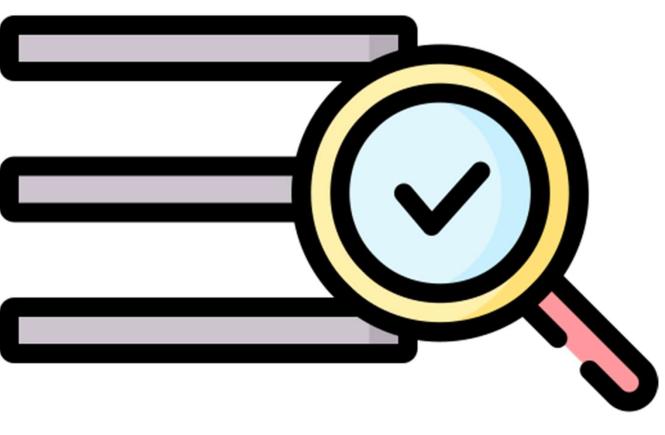








CAMO software for analytical modeling and prediction



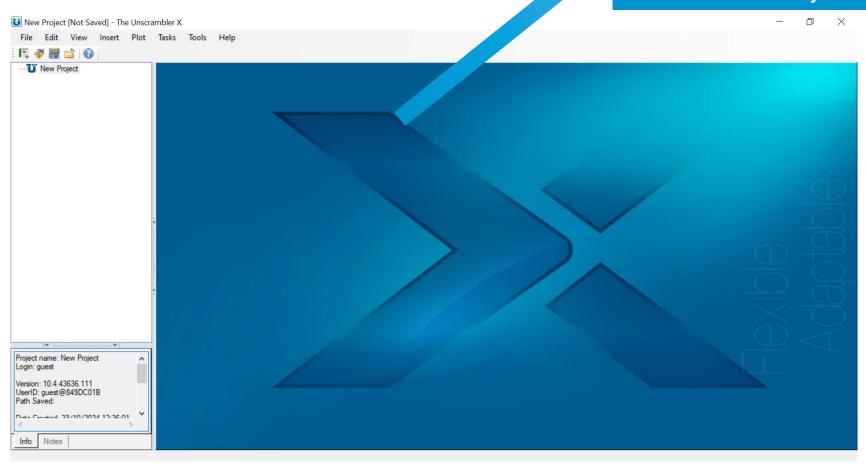
Index

- What is NIRS technology?
- The Unscrambler
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The Unscrambler [1]

Workspace

The Workspace occupies the largest area of the application window. It is a Viewer which displays results either graphically as plots or numerically as tables.

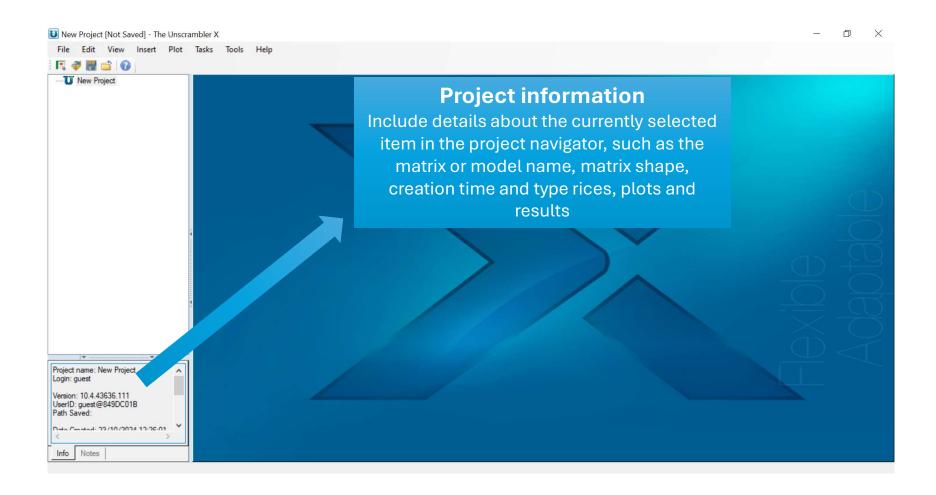


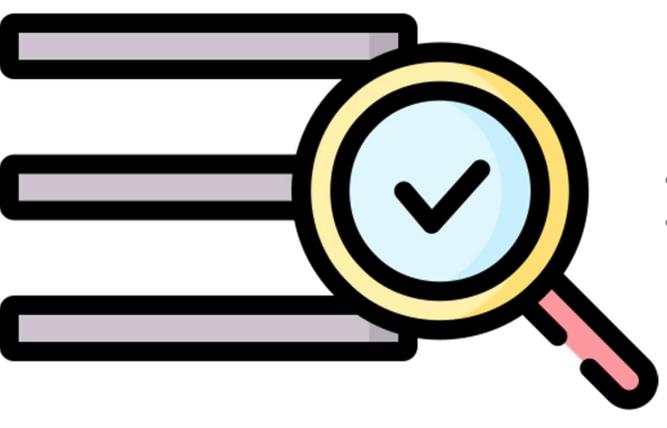
The Unscrambler [1]

Project navigator Is a tree-like structure consisting of data matrices and analysis results along with plots



The Unscrambler [1]





Index

- What is NIRS technology?
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 - Qualitative models
 - Pre-treatments

The construction of the matrix

- Data import
- Reduce (average)
- Insert row/column (to incorporate new samples or physical-chemical variables to predict)
- Insert Category variable
- Define range for the purpose of grouping rows and columns
- Plot line

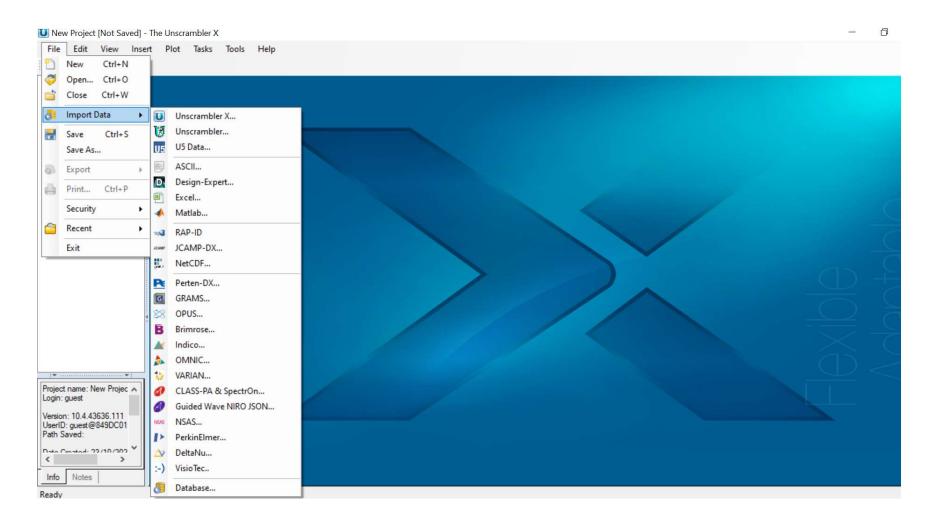


The construction of the matrix

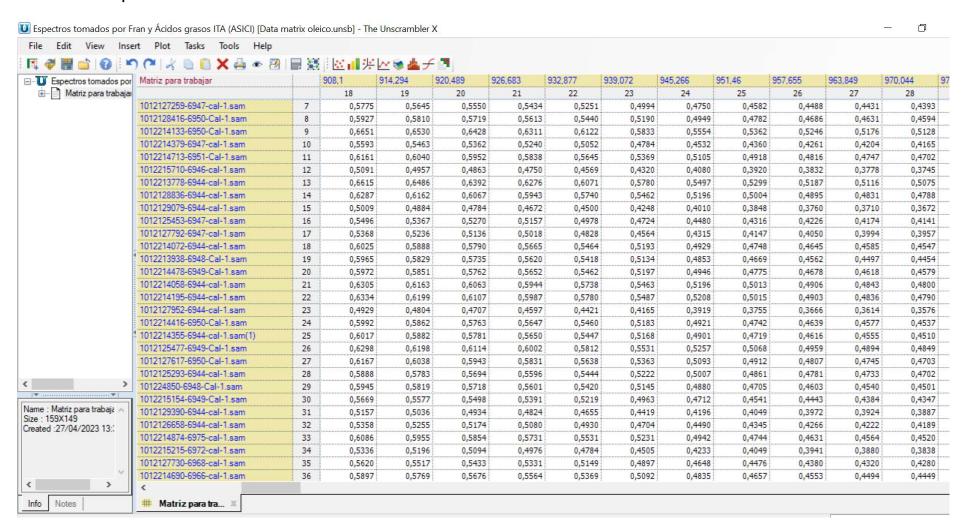
- Data import
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- Plot line



Data import



Data import



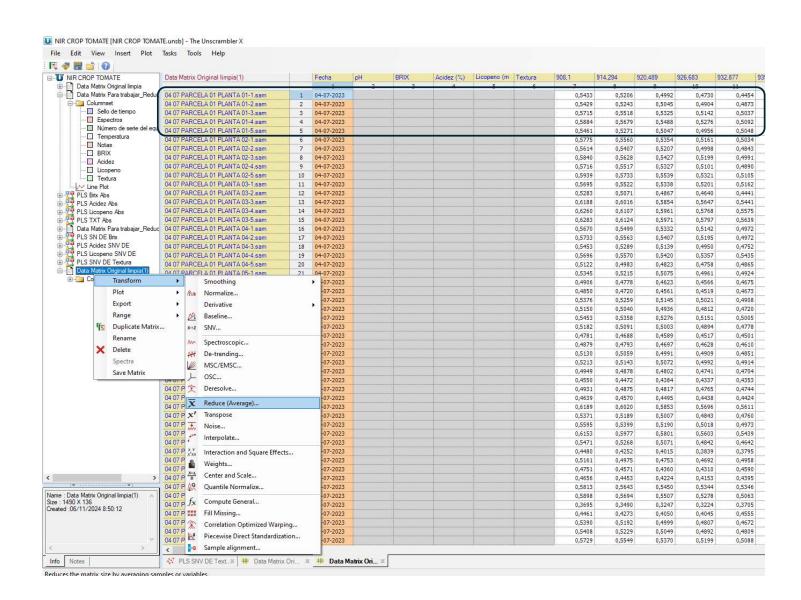
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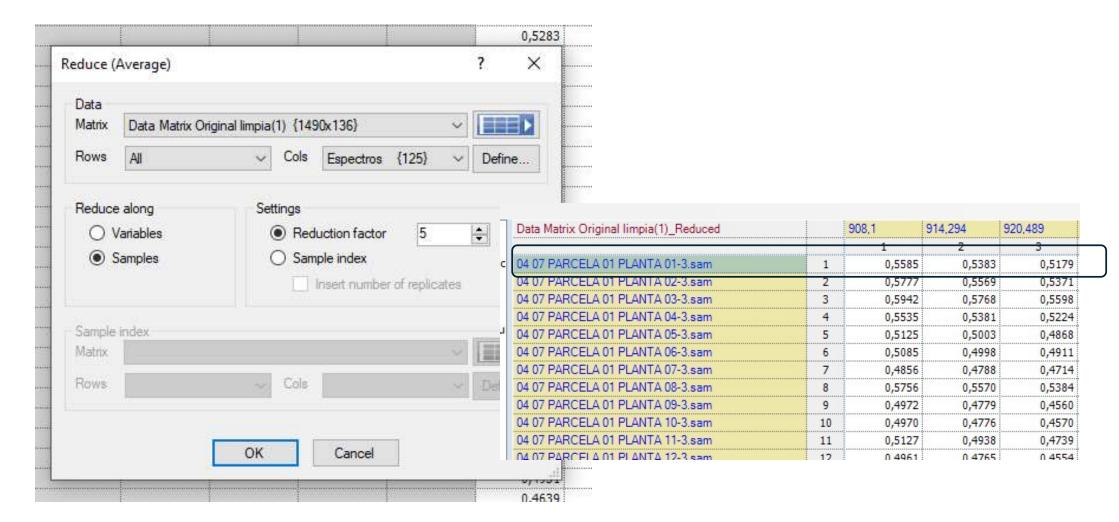


04 07 PARCELA 01 PLANTA 01-2.sam 04 07 PARCELA 01 PLANTA 01-3.sam	2 3	04-07-2023 04-07-2023	Reduce noise				
04 07 PARCELA 01 PLANTA 01-4.sam	4	04-07-2023	☐ Interpret the results more	o oscily			
04 07 PARCELA 01 PLANTA 01-5.sam	5	04-07-2023	interpret the results more	e easity			
04 07 PARCELA 01 PLANTA 02-1.sam	- 6	04-07-2023		0,5775	0,5560	0,5354	0,5161
04 07 PARCELA 01 PLANTA 02-2.sam	7	04-07-2023		0,5614	0,5407	0,5207	0,4998
04 07 PARCELA 01 PLANTA 02-3.sam	8	04-07-2023		0,5840	0,5628	0,5427	0,5199
04 07 PARCELA 01 PLANTA 02-4.sam	9	04-07-2023		0,5716	0,5517	0,5327	0,5101
04 07 PARCELA 01 PLANTA 02-5.sam	10	04-07-2023		0,5939	0,5733	0,5539	0,5321
04 07 PARCELA 01 PLANTA 03-1.sam	11	04-07-2023		0,5695	0,5522	0,5338	0,5201
04 07 PARCELA 01 PLANTA 03-2.sam	12	04-07-2023		0,5283	0,5071	0,4867	0,4640
04 07 PARCELA 01 PLANTA 03-3.sam	13	04-07-2023		0,6188	0,6016	0,5854	0,5647
04 07 PARCELA 01 PLANTA 03-4.sam	14	04-07-2023		0,6260	0,6107	0,5961	0,5768
04 07 PARCELA 01 PLANTA 03-5.sam	15	04-07-2023		0,6283	0,6124	0,5971	0,5797
04 07 PARCELA 01 PLANTA 04-1.sam	16	04-07-2023		0,5670	0,5499	0,5332	0,5142
04 07 PARCELA 01 PLANTA 04-2.sam	17	04-07-2023		0,5733	0,5563	0,5407	0,5195
04 07 PARCELA 01 PLANTA 04-3.sam	18	04-07-2023		0,5453	0,5289	0,5139	0,4950
04 07 PARCELA 01 PLANTA 04-4.sam	19	04-07-2023		0,5696	0,5570	0,5420	0,5357
04 07 PARCELA 01 PLANTA 04-5.sam	20	04-07-2023		0,5122	0,4983	0,4823	0,4758
04 07 PARCELA 01 PLANTA 05-1.sam	21	04-07-2023		0,5345	0,5215	0,5075	0,4961
04 07 PARCELA 01 PLANTA 05-2,sam	22	04-07-2023		0,4906	0,4778	0,4623	0,4566
04 07 PARCELA 01 PLANTA 05-3.sam	23	04-07-2023		0,4850	0,4720	0,4561	0,4519
04 07 PARCELA 01 PLANTA 05-4.sam	24	04-07-2023		0,5376	0,5259	0,5145	0,5021
04 07 PARCELA 01 PLANTA 05-5.sam	25	04-07-2023		0,5150	0,5040	0,4936	0,4812
OA OT DADCELA OF DIANTA OF 1 and	26	04 07 2022		U ENES	U ESEO	0 5776	0 6161

Reduce (Average)



Reduce (Average)

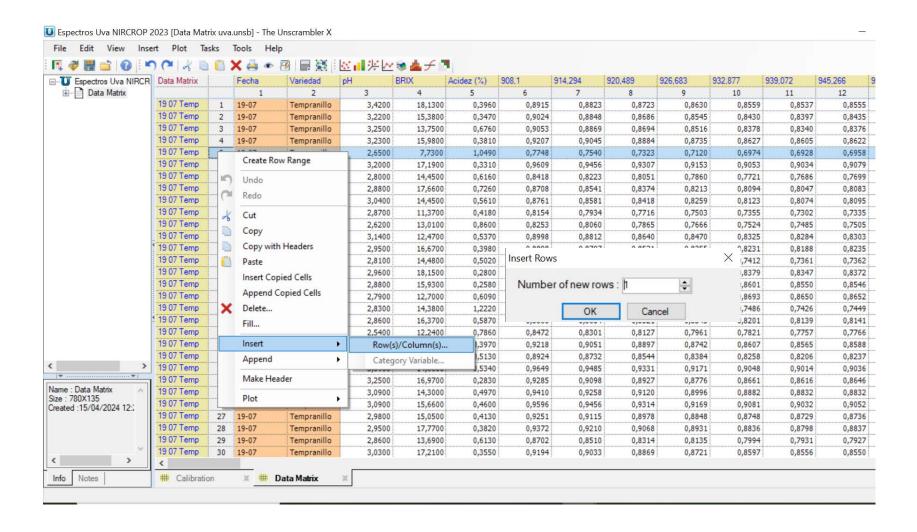


The construction of the matrix

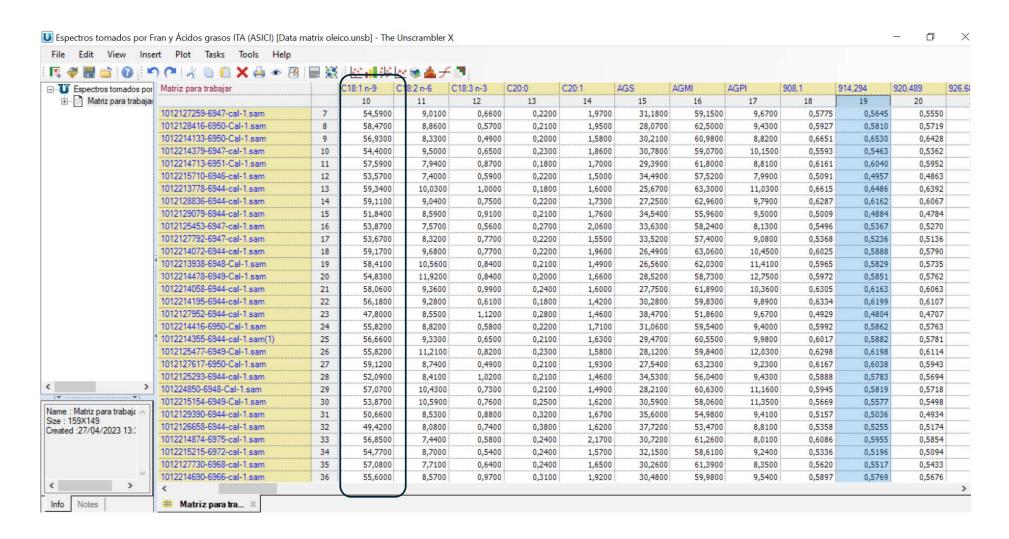
- Data import
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- Insert row/column (to incorporate new samples or physical-chemical variables to predict)
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- Define range for the purpose of grouping rows and columns
- Plot line



Insert row (to incorporate new samples)



Insert column (to incorporate physical-chemical variables to predict)

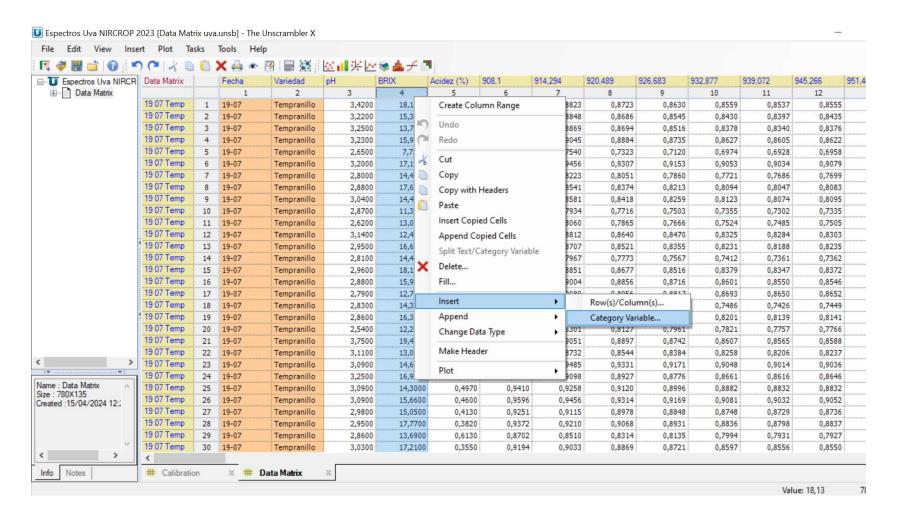


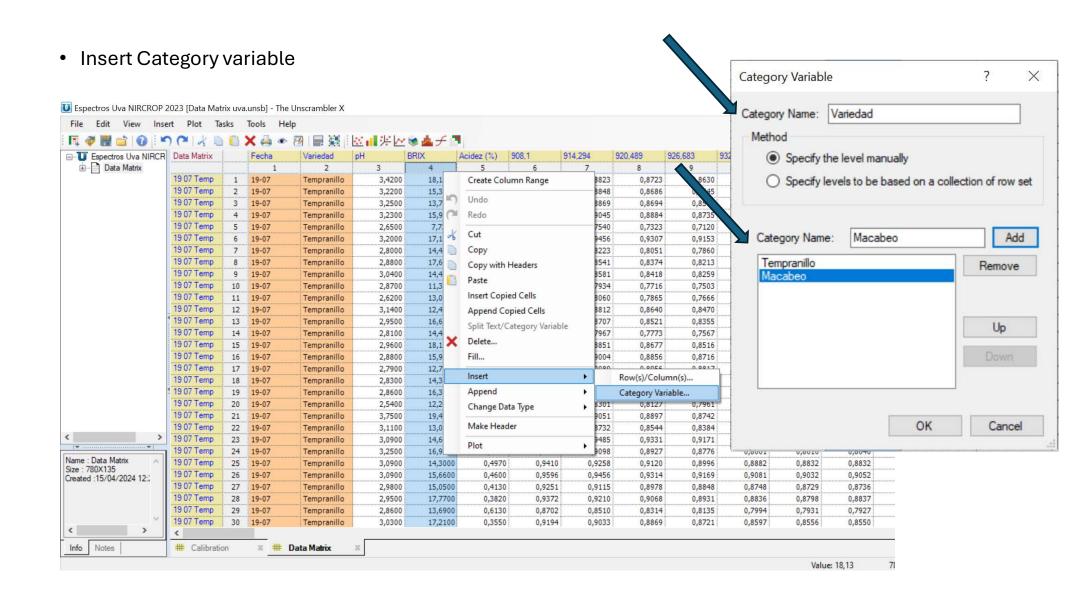
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- Data import
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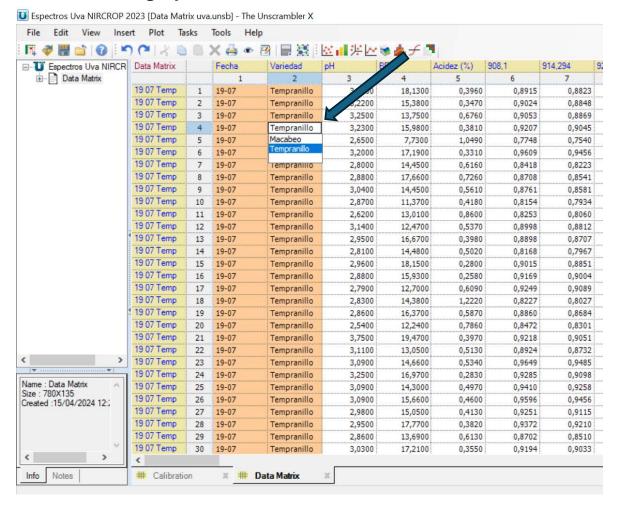


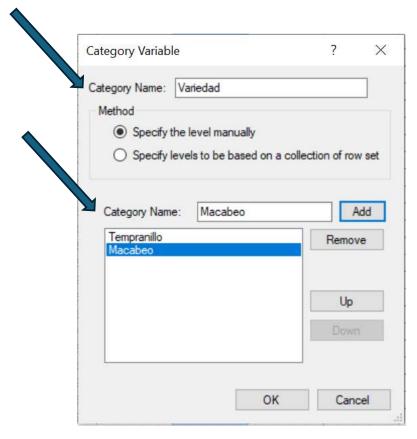
Insert Category variable





Insert Category variable





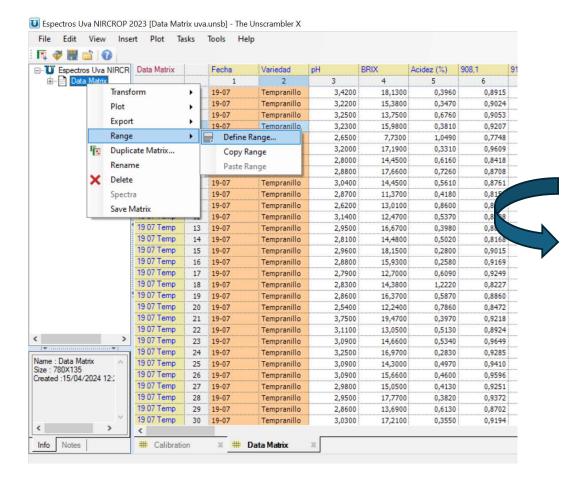
Right side/fill

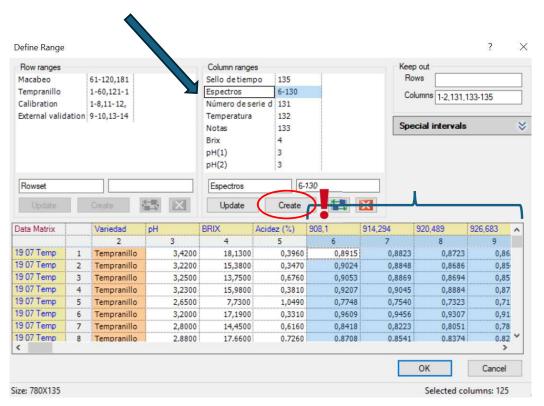
The construction of the matrix

- Data import
- Reduce (average)
- Insert row/column (to incorporate new samples or physical-chemical variables to predict)
- Insert Category variable
- Define range for the purpose of grouping rows and columns
- Plot line

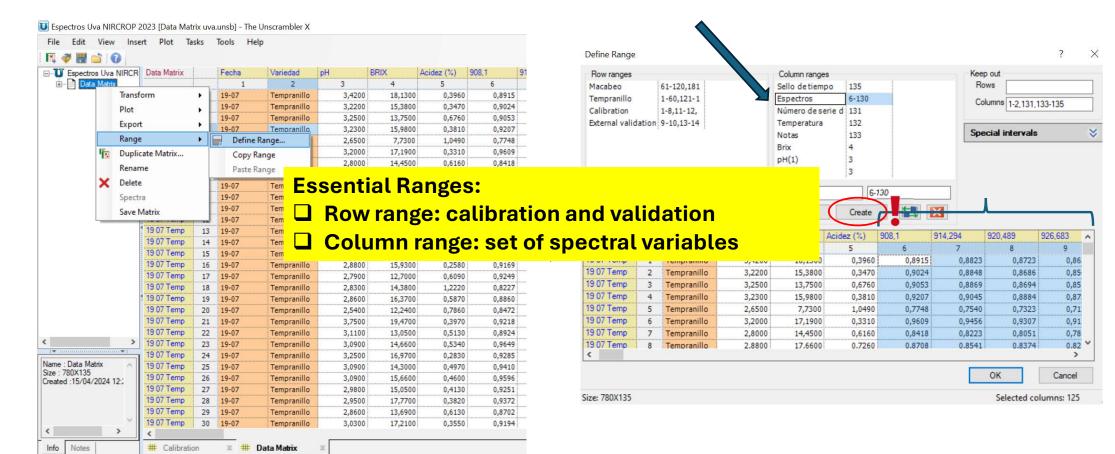


• Define range for the purpose of grouping rows and columns



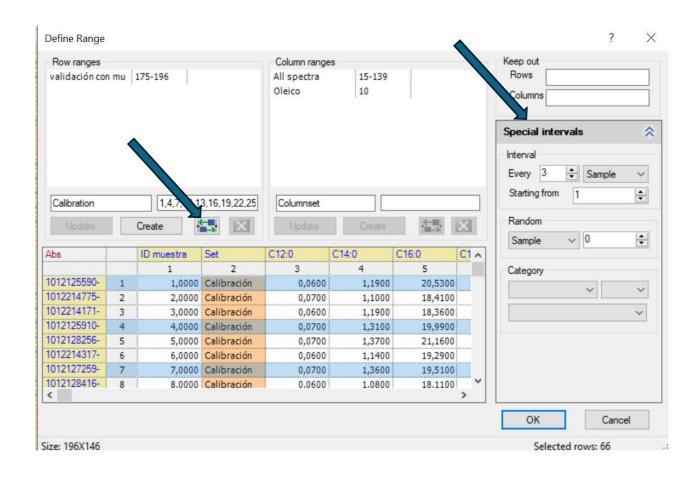


• Define range for the purpose of grouping rows and columns

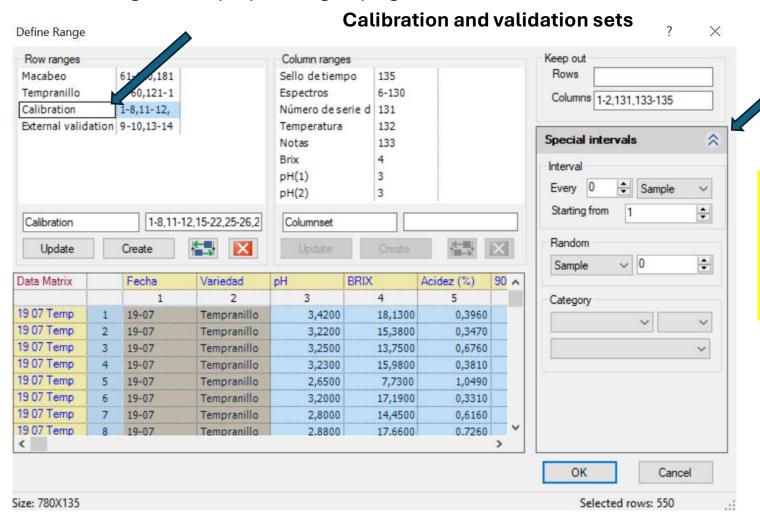


• Define range for the purpose of grouping rows and columns

Calibration and validation sets



Define range for the purpose of grouping rows and columns



It is also possible to define calibration and validation sets by means of a category variable

The construction of the matrix

- Data import
- Reduce (average)
- Insert row/column (to incorporate new samples or physical-chemical variables to predict)
- Insert Category variable
- Define range for the purpose of grouping rows and columns
- Sets (Calibration and Validation)
- Plot line

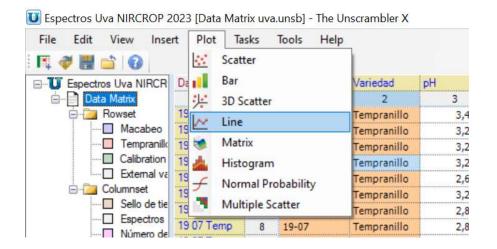


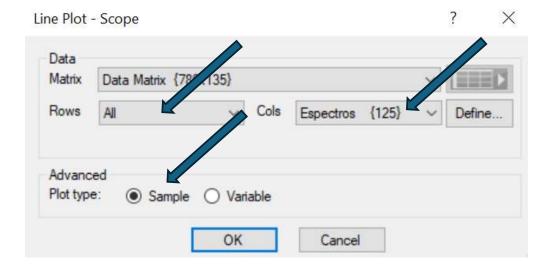
Plot line



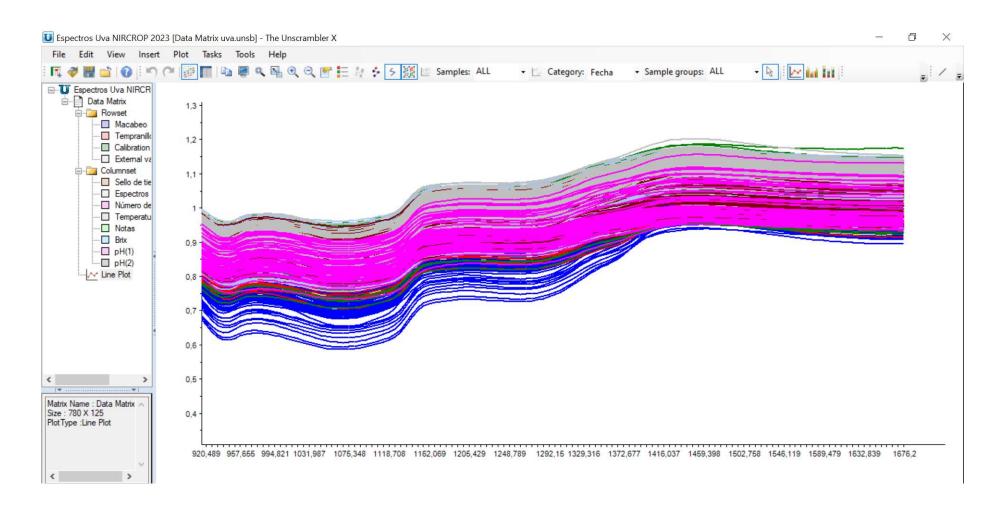
In order to visualize the spectra and detect possible anomalies

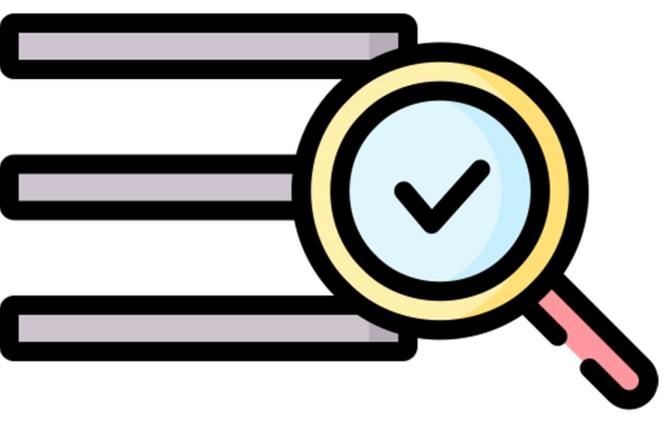
Plot/line





Plot line





Index

- What is NIRS technology?
- The Unscrambler
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 - Quantitative models
 - Qualitative models
 - Pre-treatments



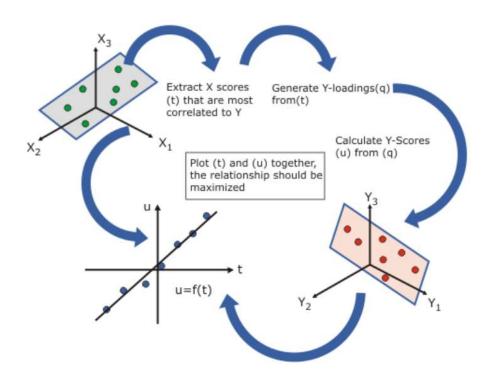
Quantitative models

- Construction of the model by means of Partial least squares regression (PLSR)
- Methods available for validation
- Plots
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- Predict
- Outliers

Construction of the model by means of Partial least squares regression (PLSR)

Partial Least Squares regression models both the X- and Y-matrices simultaneously **to find the latent variables in X that will best predict the latent variables in Y.** These PLSR components are similar to principal components; however, they are referred to as *factors*.

(It finds a linear regression model by projecting the predicted variables and the observable variables to a new space).



Construction of the model by means of Partial least squares regression (PLSR)

Partial Least Squares regression is a multivariate analysis, i.e. it uses multiple variables - spectral variables - to estimate a given physicochemical parameter.

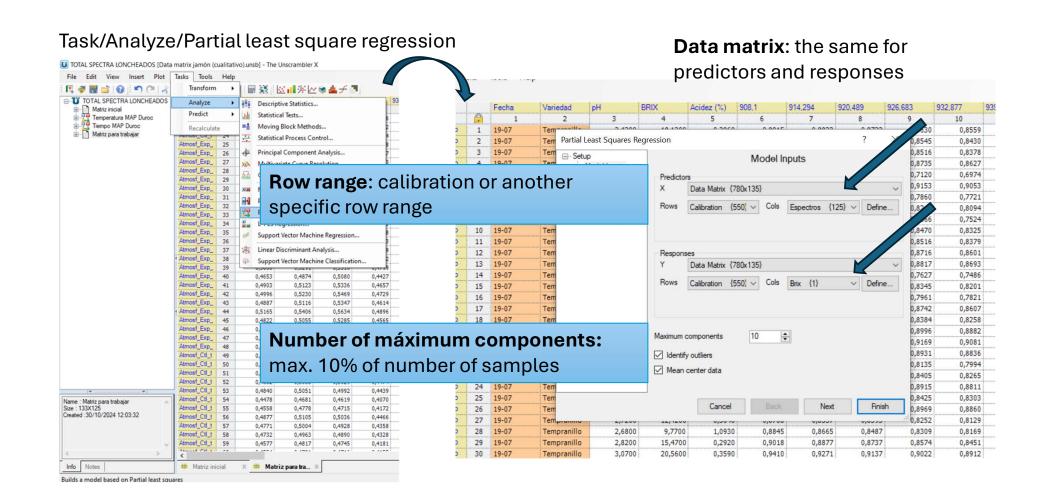
Why cannot a univariate approach be performed in NIRS?

- Overlapping absorption bands of different sample constituents
- Large amount of data
- ☐ High redundant information (collinearity)

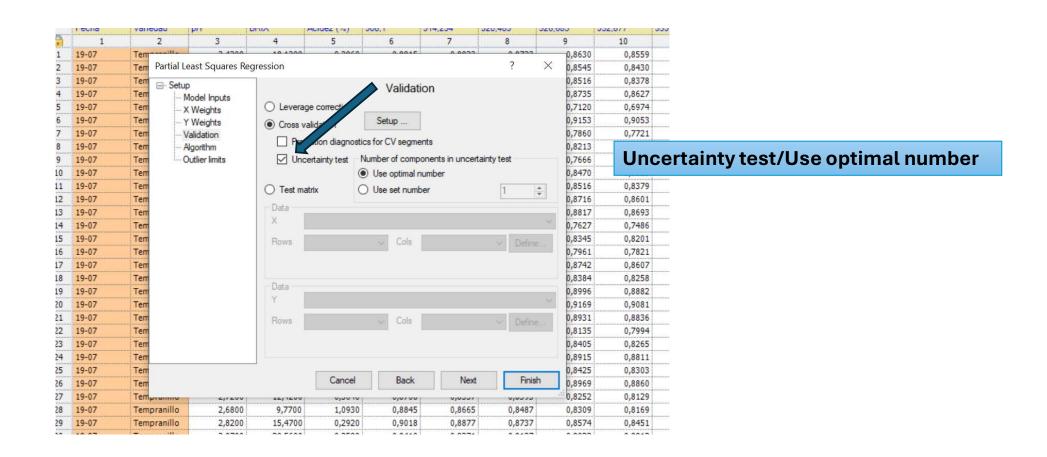


Causes low selectivity of NIR data

Construction of the model by means of Partial least squares regression (PLSR)



• Construction of the model by means of Partial least squares regression (PLSR)

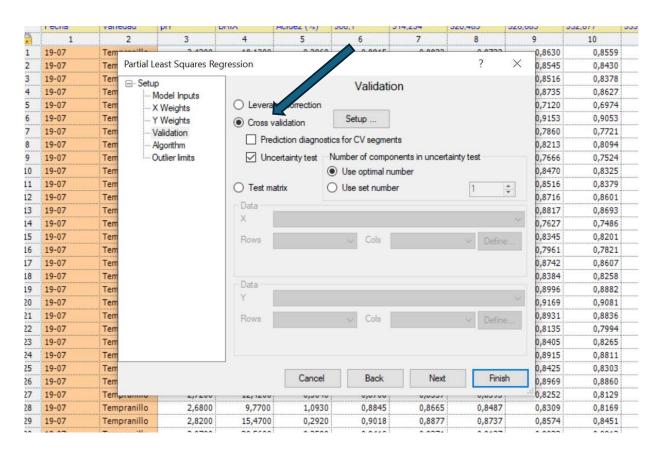




Quantitative models

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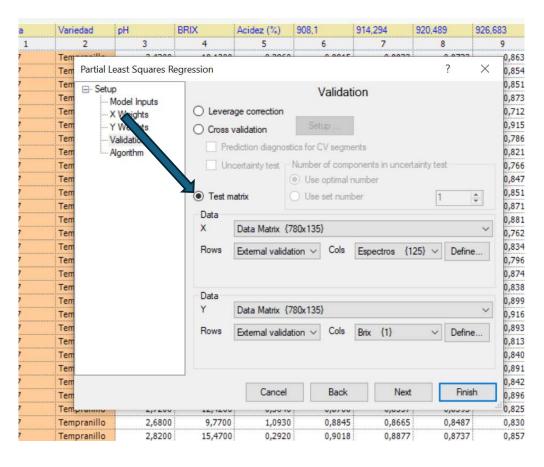
Methods available for validation



Cross-Validation

This method is used when either there are not enough samples available to make a separate test set, or for simulating the effects of different validation test cases, e.g. systematically leaving samples out vs. randomly leaving samples out, etc.

Methods available for validation



Test matrix

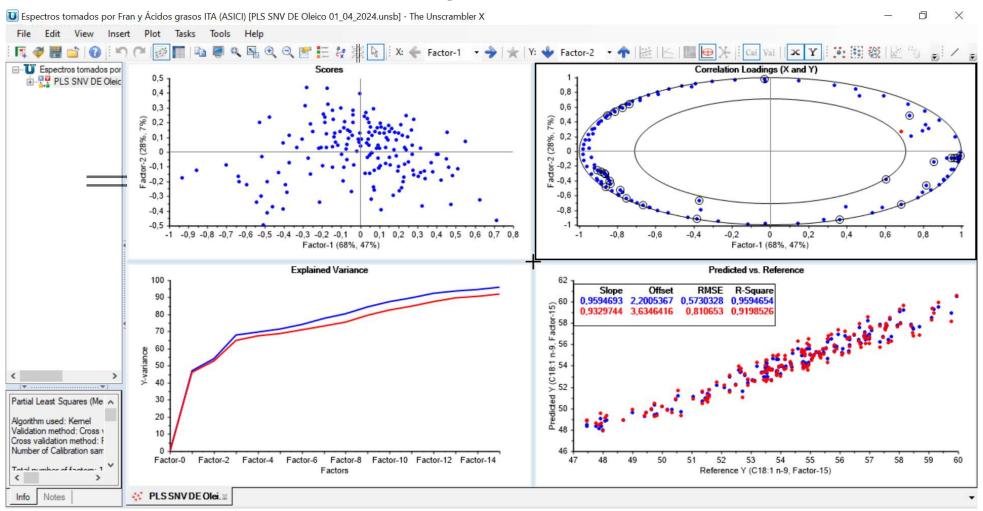
This is also known as *Test Set Validation* and uses independent samples that have not taken part in the calibration for validation. Both X- and Y-matrices need to be defined in this case. This is the preferred method for validation and should be aimed for.



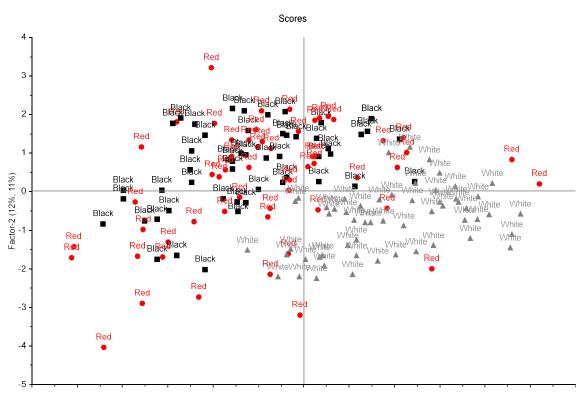
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Regression Overview



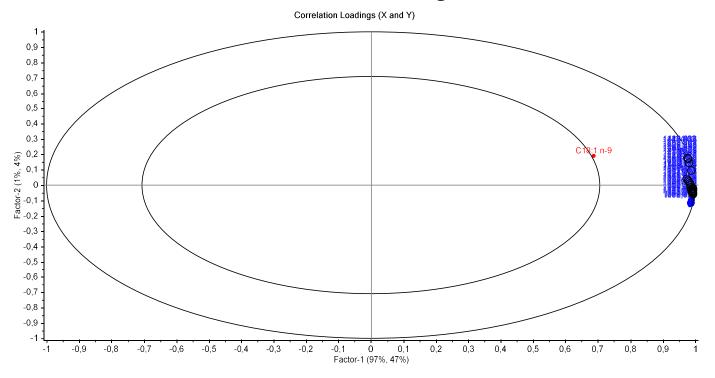
Scores plot



This is a two-dimensional scatter plot (or map) of scores for two specified factors (latent variables or PCs) from PLS regression.

The plot gives information about patterns in the samples. The scores plot for (factor 1,factor 2) is especially useful, since these two components summarize more variation in the data than any other pair of components. The closer the samples are in the scores plot, the more similar they are with respect to the two components concerned. Conversely, samples far away from each other are different from each other, so the plot can be used to interpret differences and similarities among samples. It allows to study sample distribution or even detect an outlier.

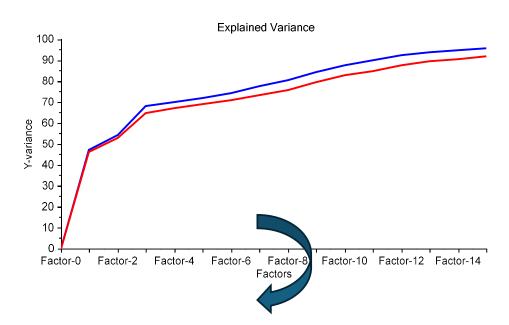
Correlation loadings



A 2-D scatter plot of X- and Y-loadings for two specified components (factors) from PLS is a good way to detect important variables and relationships between variables. The two ellipses help to check how much variance is taken into account. The outer ellipse indicates 100% explained variance. The inner ellipse indicates 50% of explained variance.

- ➤ Predictors (X) projected in roughly the same direction from the center as a response, are positively linked to that response.
- Predictors projected in the opposite direction have a negative link.
- > Predictors projected close to the center, are not well represented in that model and cannot be interpreted.

Explained/Residual variance variance

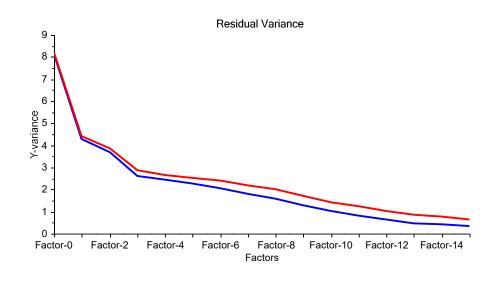


Important!!!

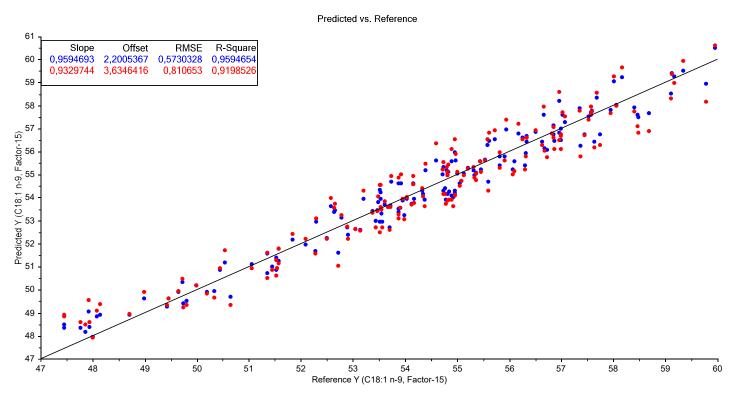
Compare the two variances (calibration and validation): if they differ significantly, there is good reason to question whether either the calibration data or the test data are truly representative!!

This plot illustrates how much of the variation in the response is described by each different component. It is the percentage of the original variance in the data that is taken into account by the model

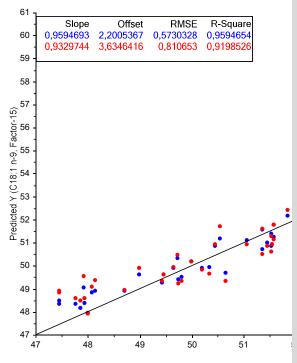
Residual variance



Predicted vs. Reference



The predicted Y-value from the model is plotted against the measured Y-value. This is a good way to check the quality of the regression model. If the model gives a good fit, the plot will show points close to a straight line through the origin and with slope equal to 1



The predicted Y-value from the check the quality of the regrectors to a straight line through

To determine the quality of the fit, the following statistics are available,

Slope

The closer the slope is to 1, the data are better modelled.

Offset

This is the intercept of the line with the Y-axis when the X-axis is set to zero (**Note**: It is not a necessity that this value is zero!)

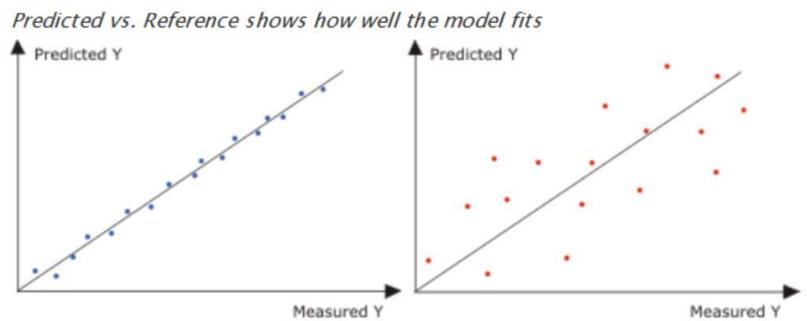
RMSE

The first one (in blue) is the Calibration error. This is a measure of the dispersion of the calibration samples about the regression line. The second one (in red) is the expected Prediction error, depending on the validation method used. Both are expressed in the same unit as the response variable Y.

R-squared

It tells how good a fit can be expected for future predictions for a defined number of components. The first one (in blue) is the calibration, the second one (in red) for the validation set.

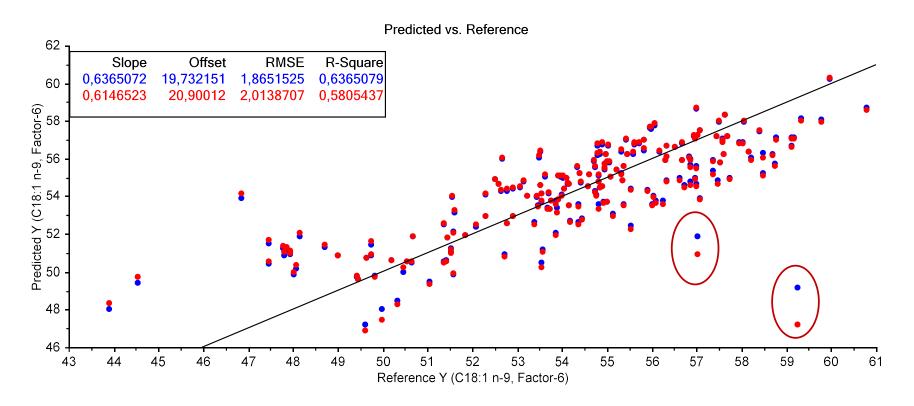
Predicted vs. Reference



Left: Good fit. Right: Poor fit.

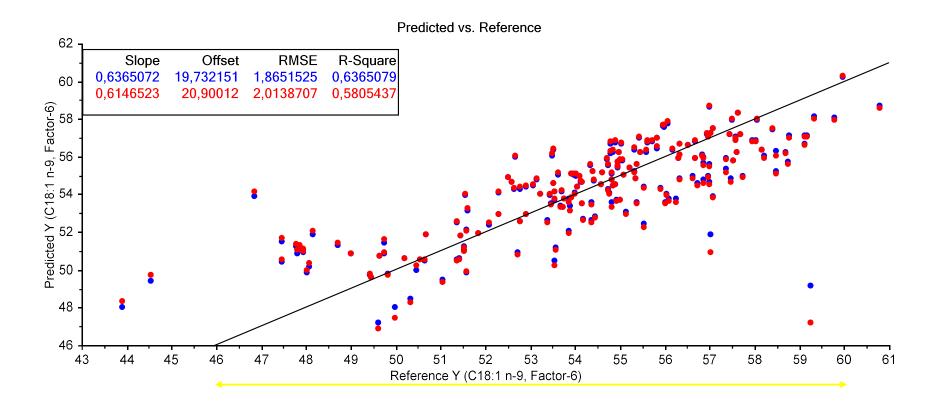
Predicted vs. Reference

If there are large differences between the calibration and validation results (in cross-validation), the model cannot be trusted.



Predicted vs. Reference

To achieve a good (robust, reliable) quantitative model, it is necessary to have a wide range of the variable to be predicted.





Quantitative models

- Construction of the model by means of Partial least squares regression (PLSR)
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- Predict
- Outliers

• Statistics for assessing a quantitative model

Coeficiente de determinación en validación cruzada (1-VR)	Error en validación cruzada
 1-VR ≥ 0,90. Excellent calibration 0,89 > 1-VR > 0,70 Good calibration 0,69 > 1-VR > 0,50 The model could be used to discriminate between low, medium and high values. 	Should be as little as possible
Desidual Duadiation Deviation (DDD)	Daniela Finan Datia (DED)
Residual Prediction Deviation (RPD)	Range Error Ratio (RER)
Residual Prediction Deviation (RPD) RPD = SD/RMSE	RER = Range/RMSE



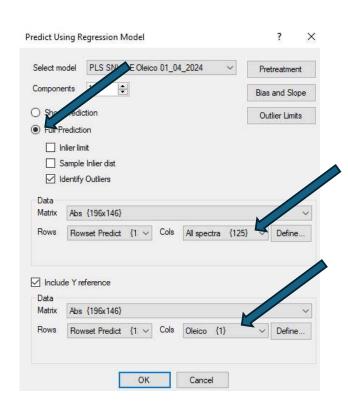
Quantitative models

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Predict

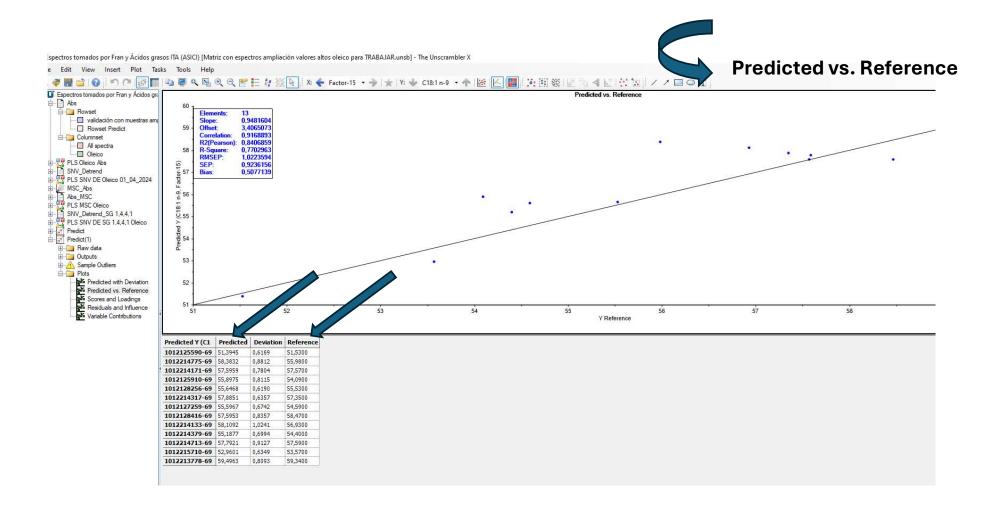
Use an existing regression model to predict response values for new samples.

Tasks/Predict/Regression



- Full Prediction uses a projection on the latent space in the calculation. (It will provide comprehensive results such as plots and additional matrices for increased data interpretation and outlier diagnostics).
- ☐ Short Prediction uses only the extracted Regression (Beta) coefficients. There are no plots associated with this type of prediction.

Predict





Quantitative models

- Construction of the model by means of Partial least squares regression (PLSR)
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Outliers

An outlier is an <u>object which deviates from the other objects</u> in a model and may not belong to the same population as the majority and therefore <u>can disturb the model</u>.

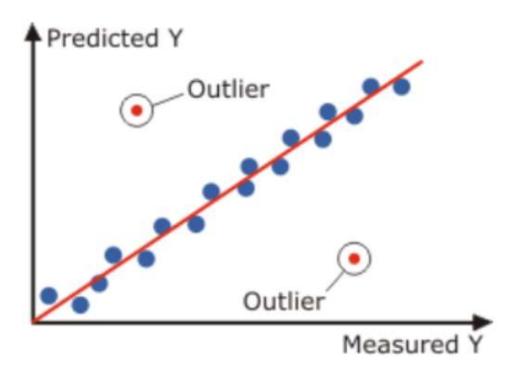
The cause of outliers could be one or more of the following:

- Measurement error
- Wrong labeling
- Deviating products / processes
- Noise
- **Extreme / interesting sample**

For projection methods like PLSR, outliers can be detected using scores plots, residuals, leverages and influence plots

• Outliers. How to detect them?

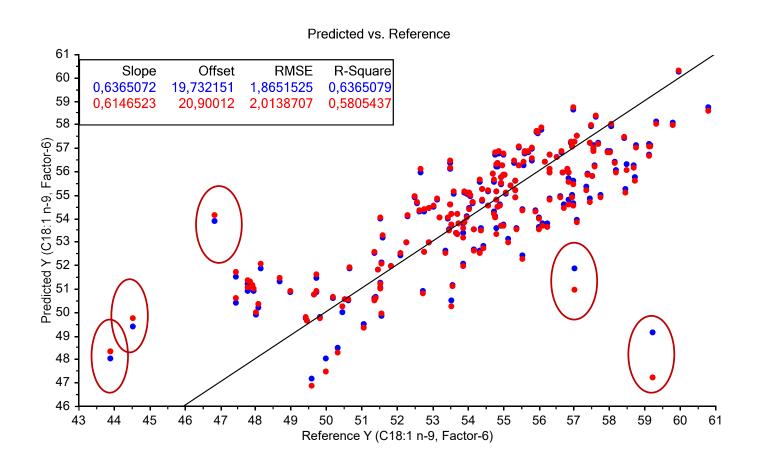
Predicted vs. Reference



One may also see cases where the majority of the samples lie close to the line while a few of them are further away. This may indicate good fit of the model to the majority of the data, but with a few outliers present

• Outliers. How to detect them?

Predicted vs. Reference



Outliers. How to detect them?

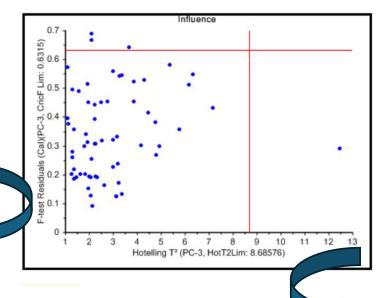
Influence plot with Hotelling's T² statistic

Representa dos tipos de outliers

Influence plot with Hotelling's T² on the abscissa and F-residuals on the ordinate

El eje de ordenadas describe la distancia de la muestra al modelo.

Las muestras con una varianza residual elevada están mal descritas por el modelo.



Las muestras con un valor elevado en el eje de abscisa (a la derecha) están bien descritas por el modelo. Sin embargo, las puntuaciones de las muestras pueden tener valores muy altos o bajos para algunos componentes en comparación con el resto de las muestras. Estas muestras son peligrosas en la fase de calibración porque modelo. influyen Una muestra suficientemente extrema puede abarcar por sí sola un componente entero, en cuyo caso el modelo dejará de ser fiable. Si la varianza descrita por la muestra es importante pero única, hay que intentar obtener más muestras del mismo tipo para estabilizar el modelo. De lo contrario, la muestra debe descartarse como valor atípico.

Outliers. How to detect them?

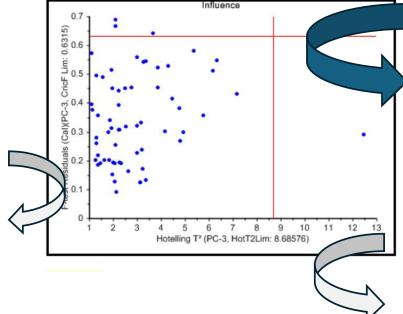
Influence plot with Hotelling's T² statistic

Representa dos tipos de outliers

Influence plot with Hotelling's T2 on the abscissa and F-residuals on the ordinate

El eje de ordenadas describe la distancia de la muestra al modelo.

Las muestras con una varianza residual elevada están mal descritas por el modelo.

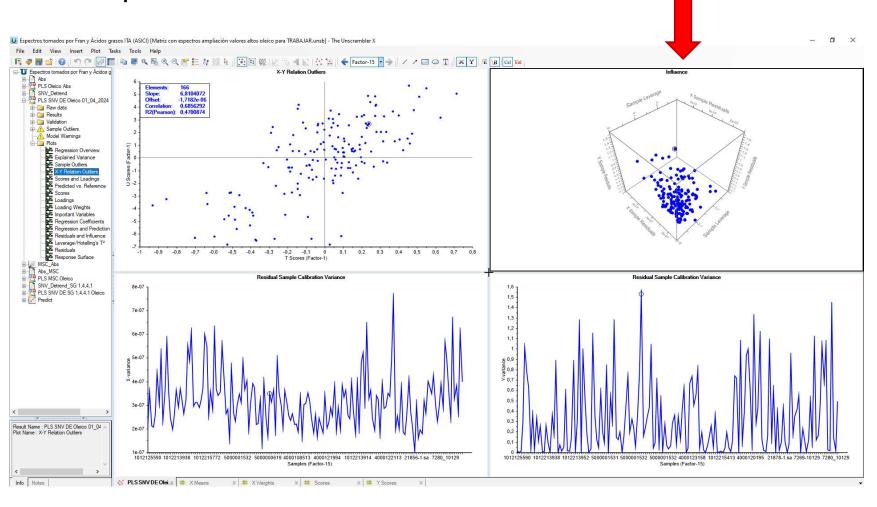


Obsérvese que una muestra con una varianza residual y un valor de Hotelling elevados es el valor atípico más peligroso. No sólo está mal descrita por el modelo, sino que además es influyente.

muestras. Estas muestras son peligrosas en la fase de calibración porque influyen en el modelo. Una muestra suficientemente extrema puede abarcar por sí sola un componente entero, en cuyo caso el modelo dejará de ser fiable. Si la varianza descrita por la muestra es importante pero única, hay que intentar obtener más muestras del mismo tipo para estabilizar el modelo. De lo contrario, la muestra debe descartarse como valor atípico.

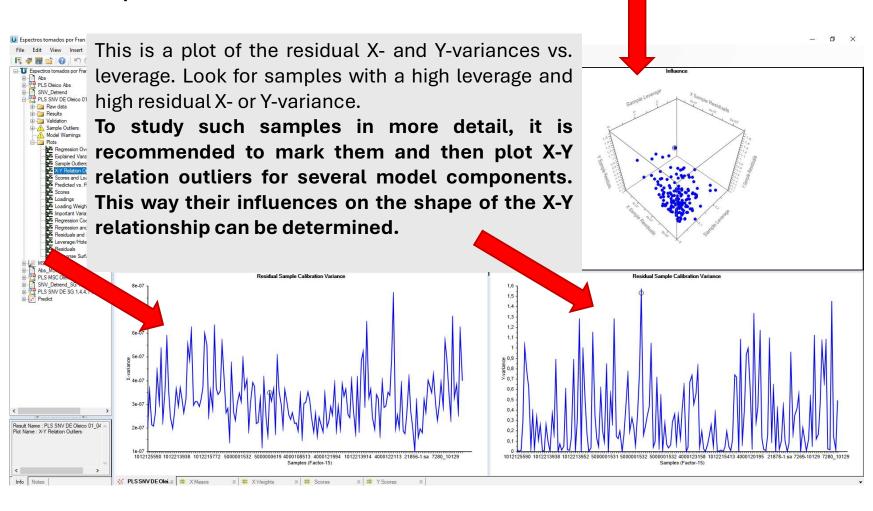
• Outliers. How to detect them?

Influence plot



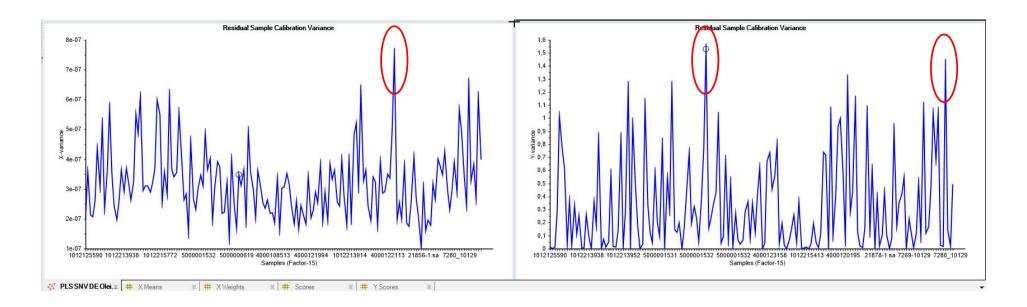
• Outliers. How to detect them?

Influence plot

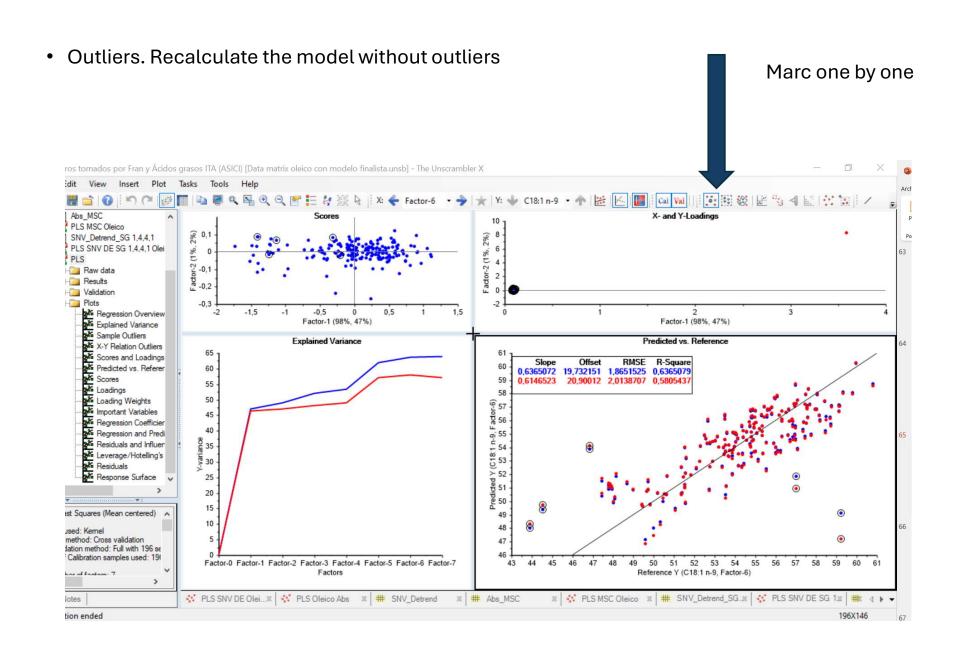


Outliers. How to detect them?

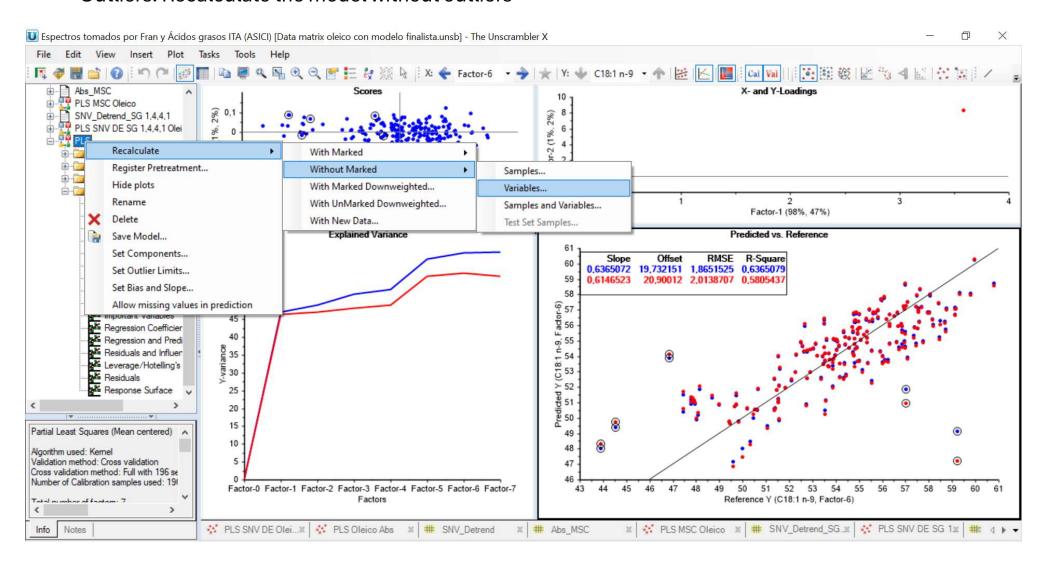
Residual sample variance

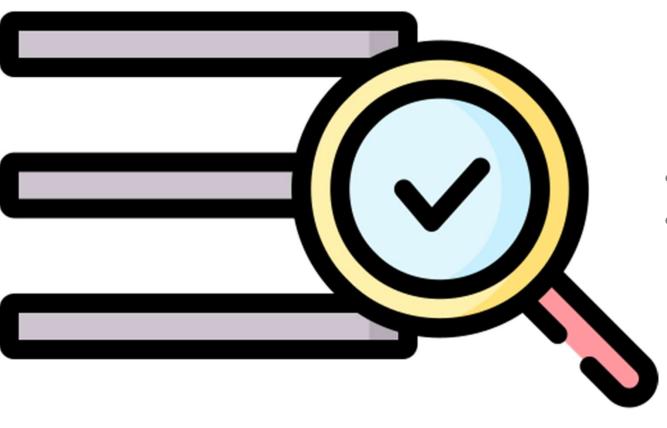


High residuals indicate an outlier. Incorporating more components can sometimes model outliers; avoid doing so since it will reduce the prediction ability of the model.



Outliers. Recalculate the model without outliers





Index

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 - Quantitative models
 - Qualitative models
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Qualitative models

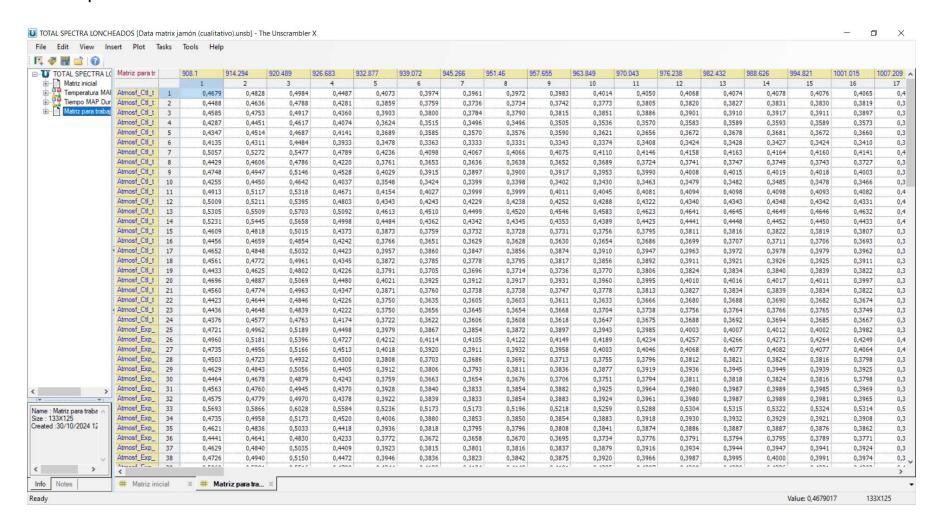
- Data import
- Insert row/column (to incorporate new samples or physical-chemical variables to predict)
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Data import

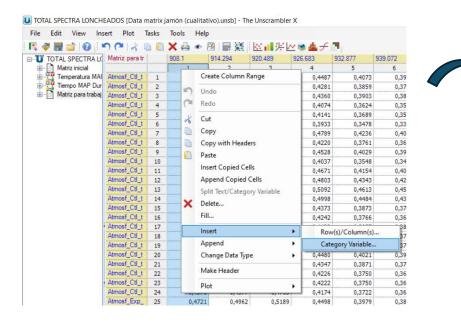


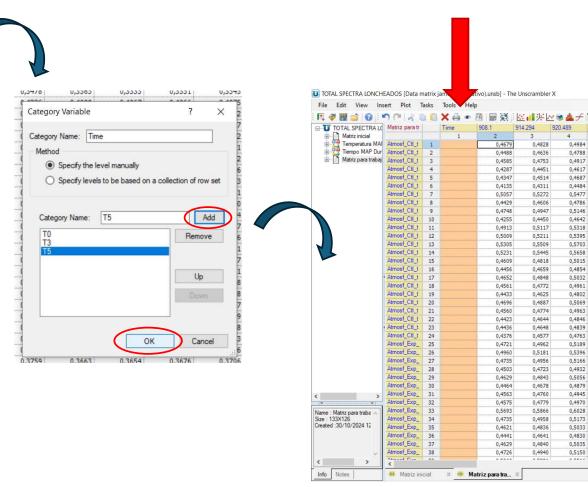


Qualitative models

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- Insert row/column (to incorporate new samples or physical-chemical variables to predict)
- Insert Category variable





• Insert Category variable

Assignment of dummy classes

01011 11010 01011

 $1 \rightarrow$ Assigned to the sample belonging to the study class

 $0 \rightarrow$ Assigned to the sample that does **NOT** belong to the study class.

	T3	T5	Time
4	5	6	
1,0000	0,0000	0,0000	T0
1,0000	0,0000	0,0000	T0
1,0000	0,0000	0,0000	T0
1,0000	0,0000	0,0000	TO
1,0000	0,0000	0,0000	T0
1,0000	0,0000	0,0000	T0
1,0000	0,0000	0,0000	T0
1,0000	0,0000	0,0000	T0
1,0000	0,0000	0,0000	T0
1,0000	0,0000	0,0000	T0
1,0000	0,0000	0,0000	T0
1,0000	0,0000	0,0000	T0
1,0000	0,0000	0,0000	T0
1,0000	0,0000	0,0000	T0
1,0000	0,0000	0,0000	T0
1,0000	0,0000	0,0000	T0
1,0000	0,0000	0,0000	T0
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1,0000	0,0000	0,0000	T0
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1,0000	0,0000	0,0000	T0
1,0000	0,0000	0,0000	T0
1,0000	0,0000	0,0000	T0
1,0000	0,0000	0,0000	T0
1,0000	0,0000	0,0000	TO
	1,0000 1,0000	4 5 1,0000 0,0000 1,0000 <td< td=""><td>4 5 6 1,0000 0,0000 0,0000 1,0000 0,0000</td></td<>	4 5 6 1,0000 0,0000 0,0000 1,0000 0,0000

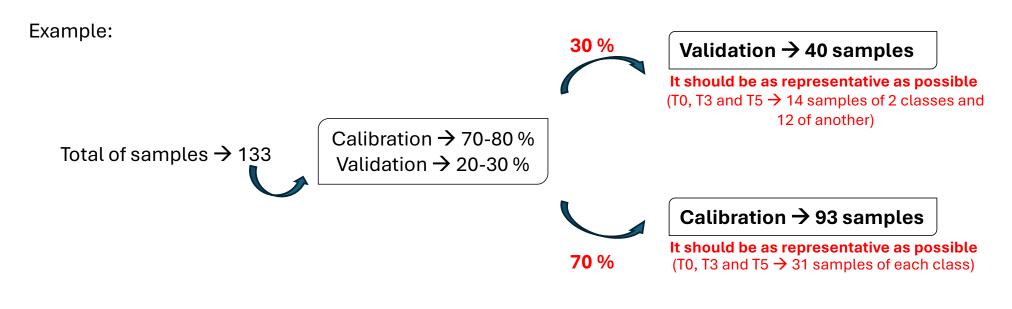
Split Category Variable

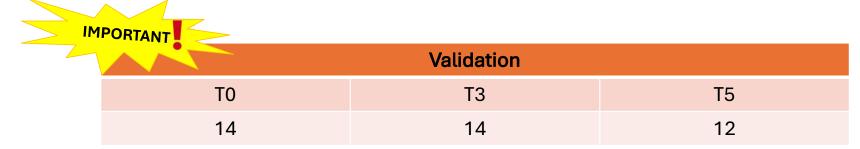


Qualitative models

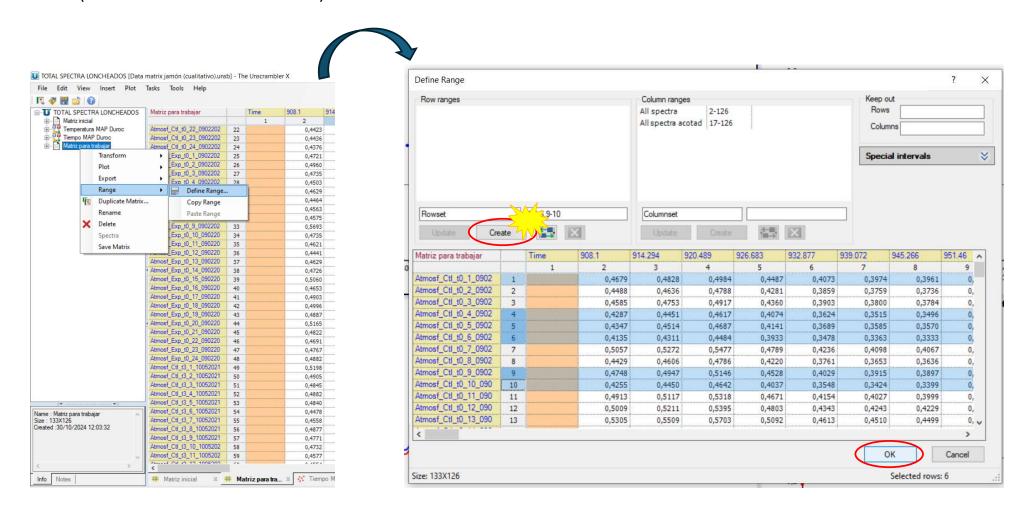
- Data import
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- Define range for the purpose of grouping rows and columns
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- Define range for the purpose of grouping rows and columns
- Sets (Calibration and Validation)





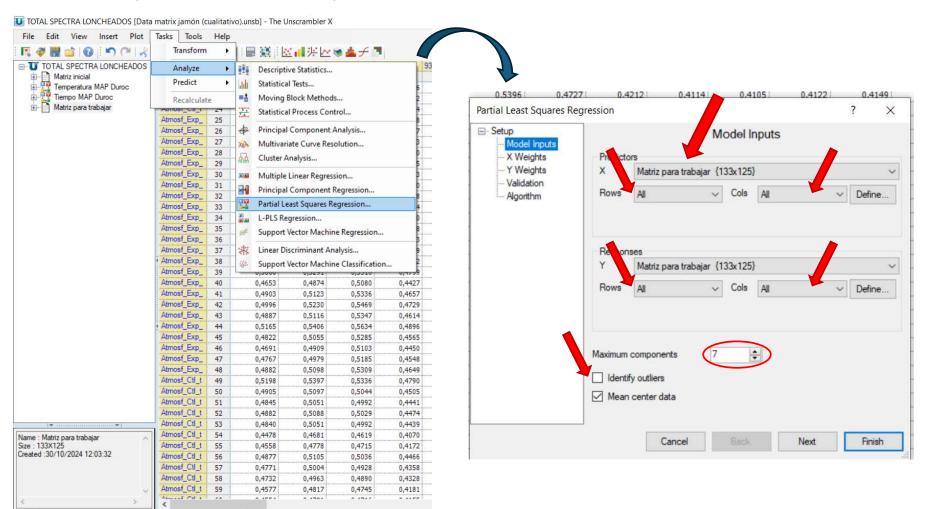
Qualitative models

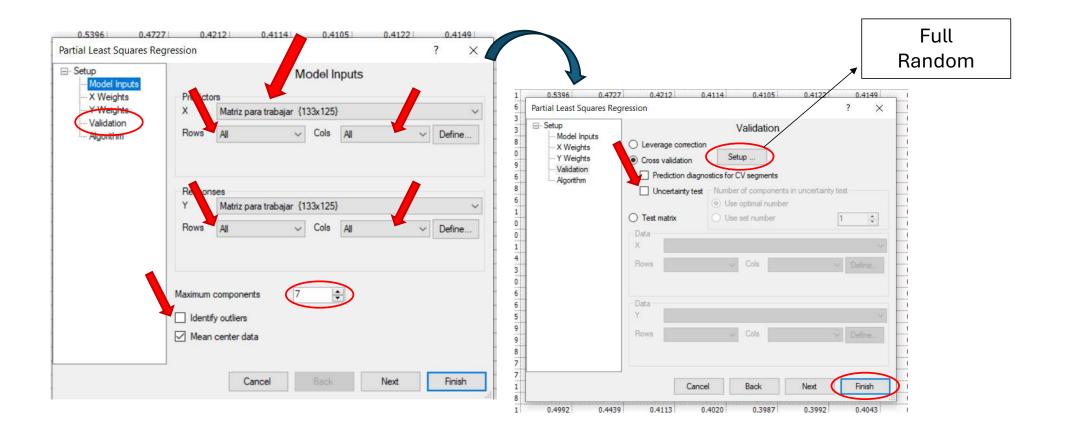
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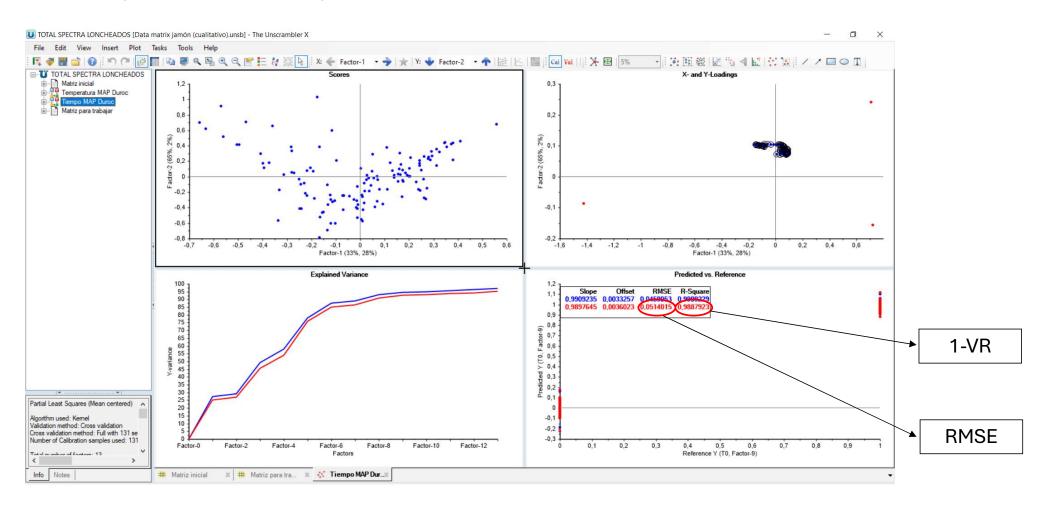
Matriz inicial

Info Notes

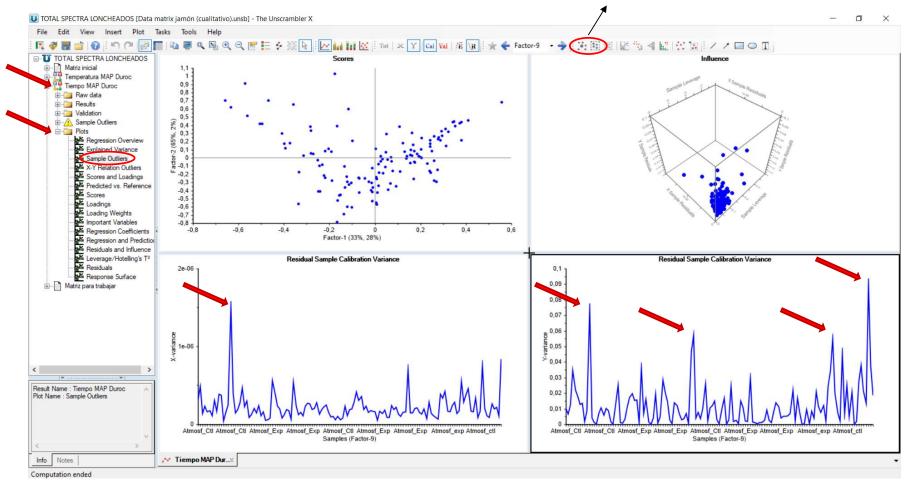
Builds a model based on Partial least squares







Click to select the outliers we have identified





Qualitative models

- Data import
- Insert row/column (to incorporate new samples or physical-chemical variables to predict)
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STATISTICS

SENSITIVITY (SE)

Which denotes the percentage of samples belonging to an established class that the studied model has recognized as belonging to that class

$$SE \ (\%) = \frac{TP}{TP + FN}$$

TP = True positive FN = False negative

SPECIFICITY (SP)

Which refers to the percentage of samples that do not belong to the target class and that the model has correctly rejected

$$SP\ (\%) = \frac{TN}{TN + FP}$$

TN = True negative FP = False positive

STATISTICS

ACCURACY

Defined as the percentage of samples correctly classified during external validation

$$Accuracy (\%) = \frac{TN + TP}{TN + TP + FN + FP}$$

TP = True positive

TN = True negative

FP = False positive

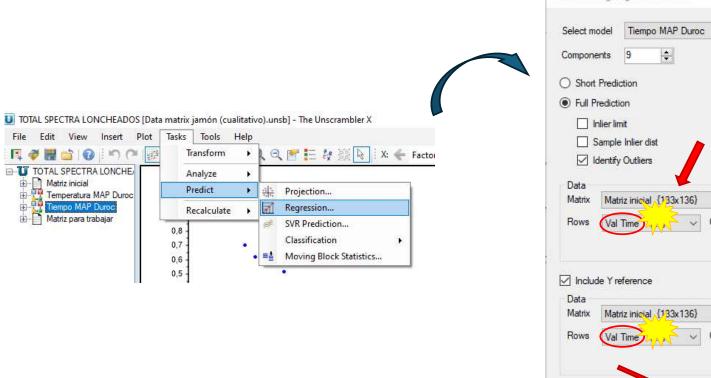
FN = False negative

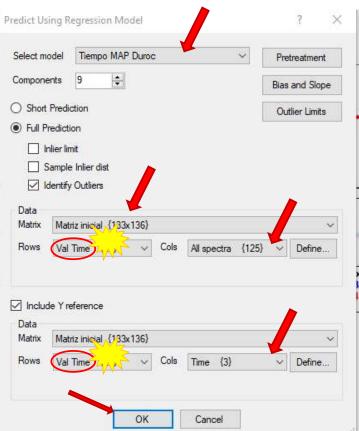
MATTHEWS CORRELATION COEFFICIENT (MCC)

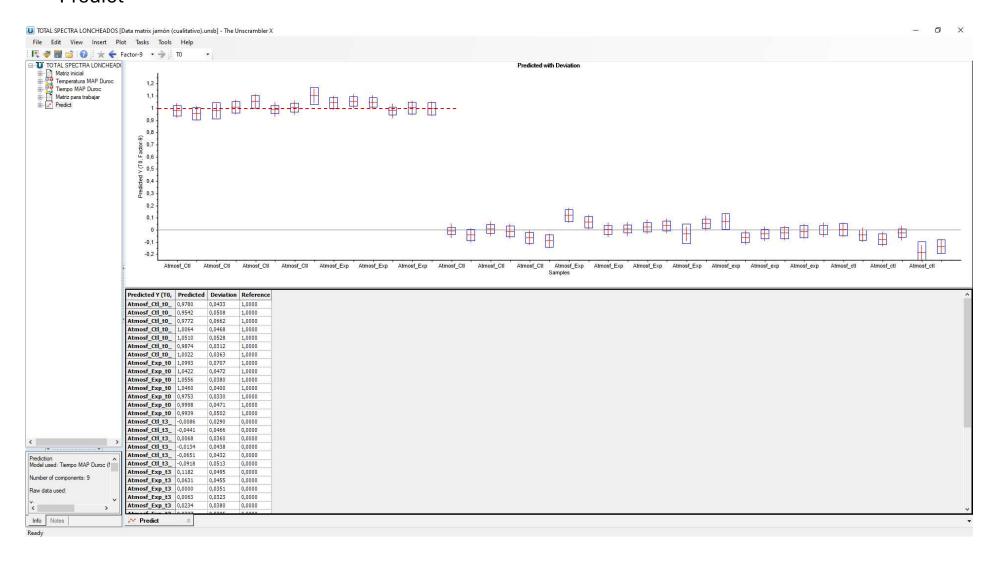
Which is suitable for assessing the performance of the model when the number of samples in the various classes is not the same

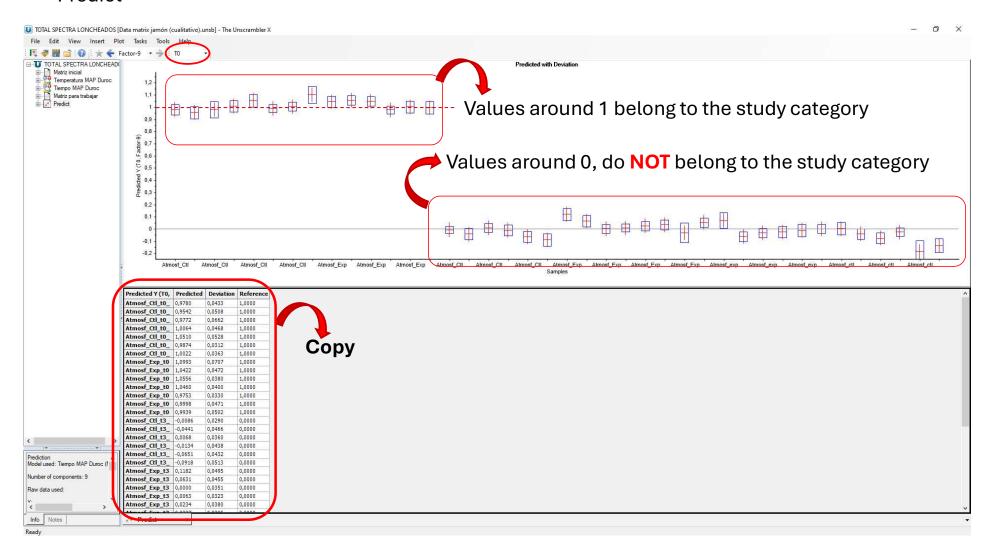
$$MCC = \frac{TP \times TN - TP \times FN}{\sqrt{(TP+FN)(TP+FN)(TN FN)(TN+TP)}}$$

Value of 1 indicates a perfect prediction; -1 represents a total disagreement between the prediction and the actual values; and 0 means "no better than a random prediction"

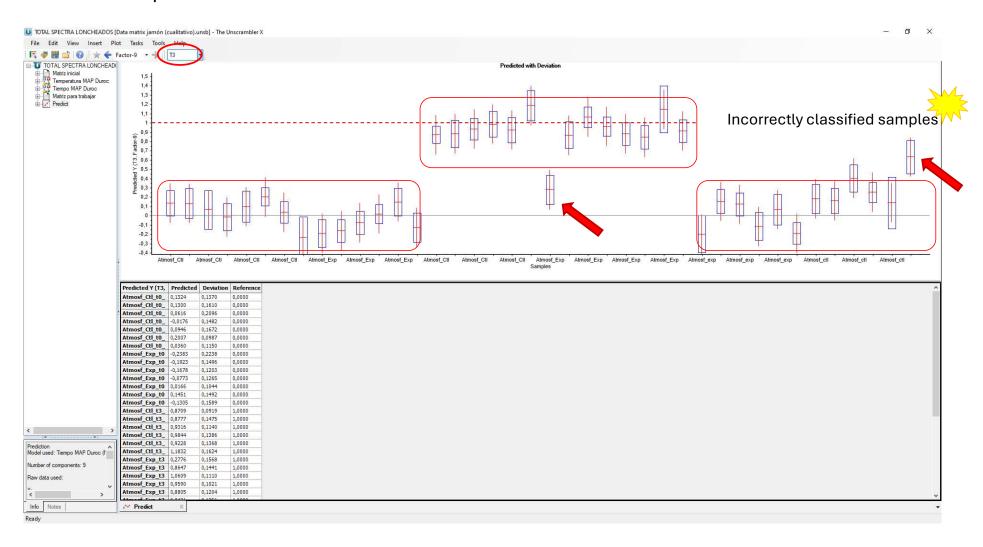






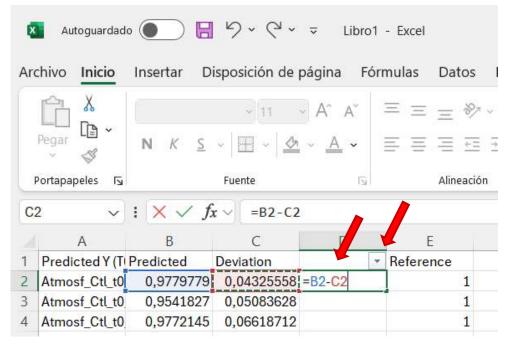


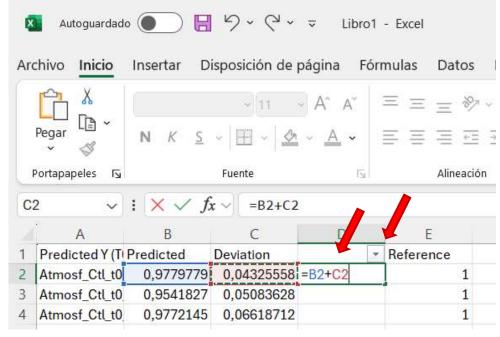
Another example:



Predicted value - deviation $\geq 0.5 \rightarrow$ **SENSITIVITY**. Predicted value + deviation $< 0.5 \rightarrow$ **SPECIFICITY**.

We calculate the most unfavorable case





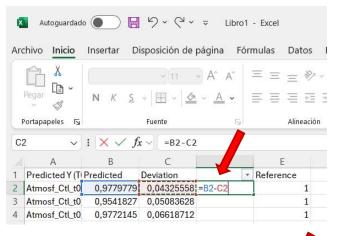
SENSITIVITY

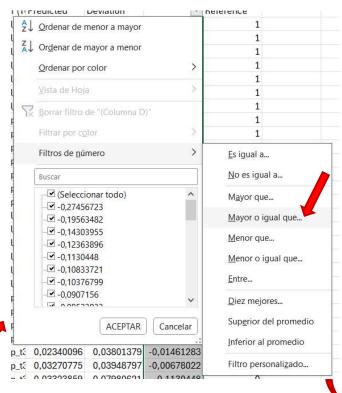
SPECIFICITY

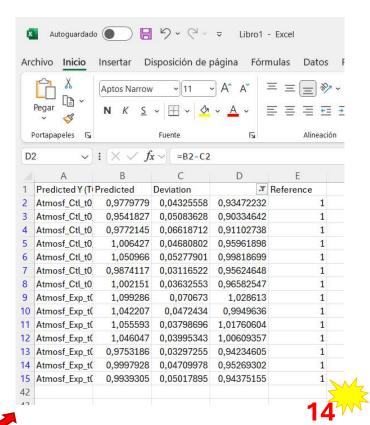
Predicted value - deviation $\geq 0.5 \rightarrow$ **SENSITIVITY**. Predicted value + deviation $< 0.5 \rightarrow$ **SPECIFICITY**.

We calculate the most unfavorable case

SENSITIVITY







True positive

Validation				
ТО	Т3	T5		
14	14	12		



$$SE (\%) = \frac{TP}{TP + FN}$$

$$SE\ (\%) = \frac{14}{14+0}$$



TP = True positive

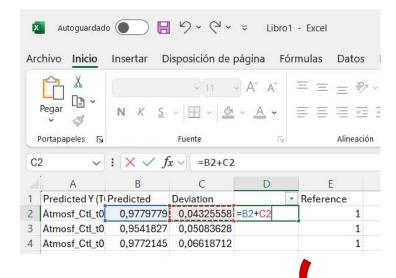
FN = False negative

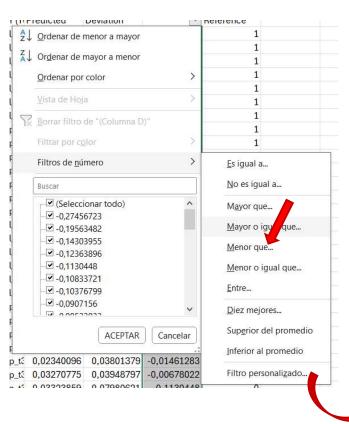
FN = Total class samples - TP

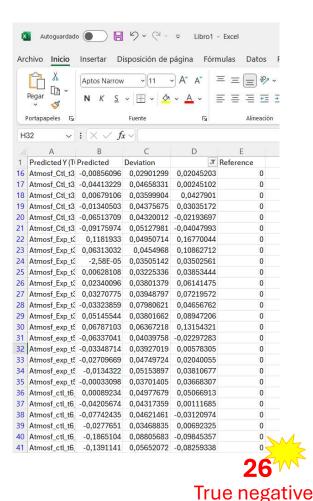
Predicted value - deviation $\geq 0.5 \rightarrow$ **SENSITIVITY**. Predicted value + deviation $< 0.5 \rightarrow$ **SPECIFICITY**.

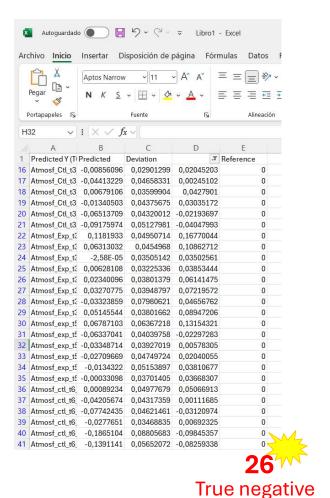
We calculate the most unfavorable case

SPECIFICITY









	Validation	
T0	Т3	T5
14	14	12
26		

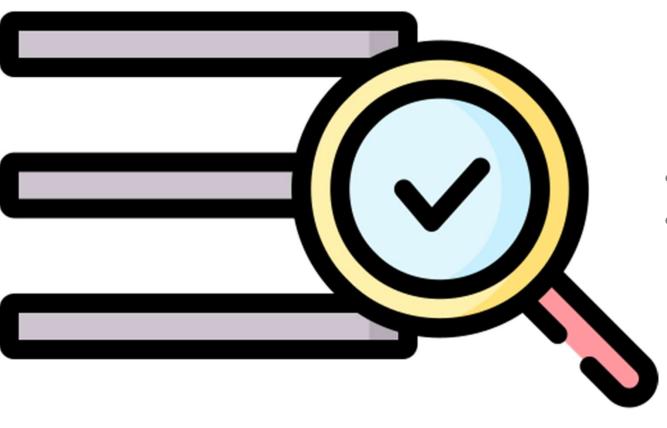
$$SP\ (\%) = \frac{TN}{TN + FP}$$

$$SP\ (\%) = \frac{26}{26+0}$$



TN = True negative FP = False positive

FP = Total of samples not belonging to the class - TN



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Los **efectos no deseados presentes en las señales** obtenidas en espectroscopia se denominan comúnmente **ruido**.

Este ruido puede tener diferentes causas u orígenes.

- ☐ Puede haber ruido debido a los componentes de la instrumentación utilizada para el registro del espectro (ruido instrumental) (variaciones de temperatura, humedad u otras)
- o bien variaciones en la señal debidas a la propia naturaleza de la muestra



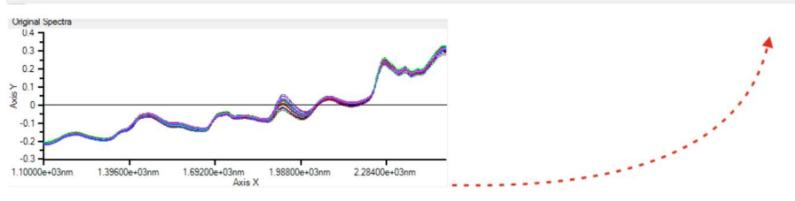
Pre-treatments

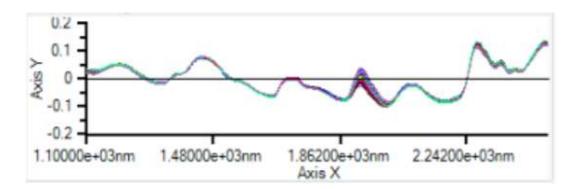
Tratamientos espectrales para reducir el ruido

Pre-treatments	
□ Scattering effect refers to the dispersion of it (texture, size and geometry of the particles the variations in the spectral data.	incident radiation due to physical phenomena at make up the sample). This causes unwanted Scatter correction treatments Detrending (DT) Standard Normal Variate (SNV) Multiplicative Scatter Correction (MSC)
☐ Average Averaging several analytical signals of the same noise ratio.	sample decreases noise and improves the signal-to-
□ Spectral smoothing to mathematically redu	uce random noise and increase signal-to-noise ratio Derivatives

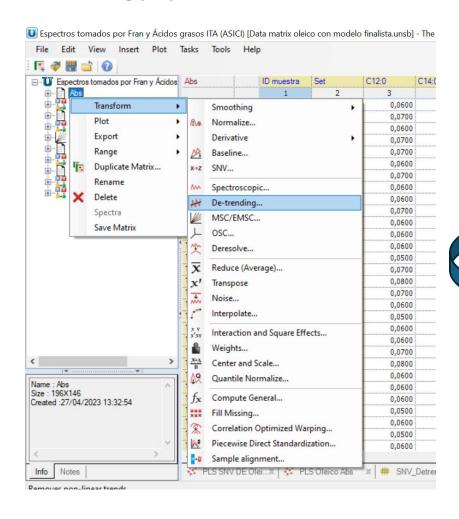
Detrending (DT)

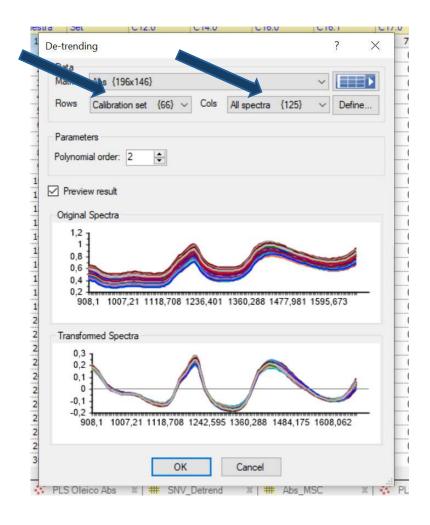
Detrending (DT) is a transformation which seeks to remove nonlinear trends in spectroscopic data, normally caused by scatter effect.





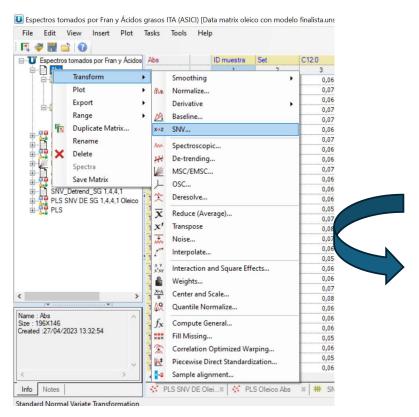
Detrending (DT)

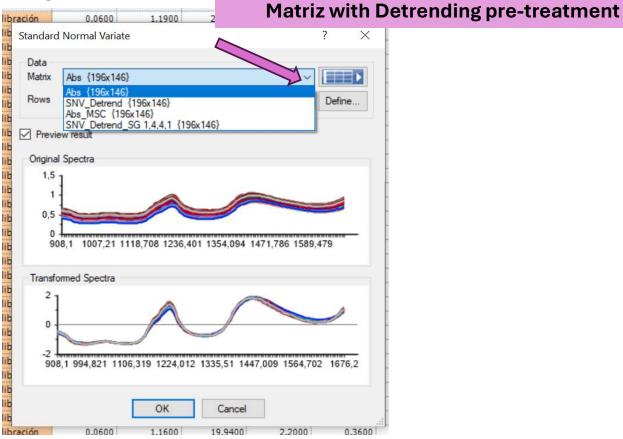




Standard Normal Variate (SNV)

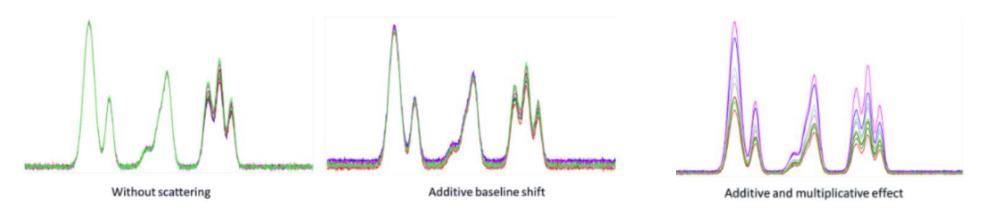
It is usually applied together with the DT. The SNV is mainly used to correct baseline variations caused by scattering.



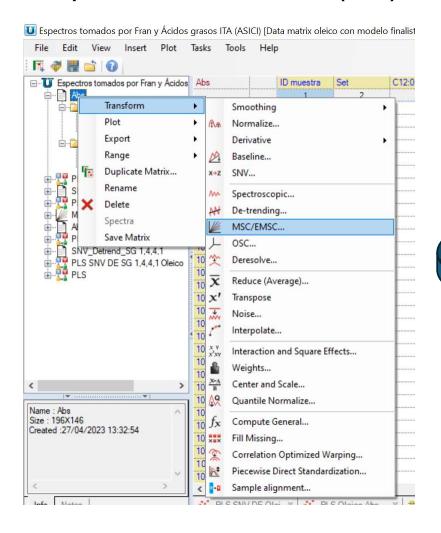


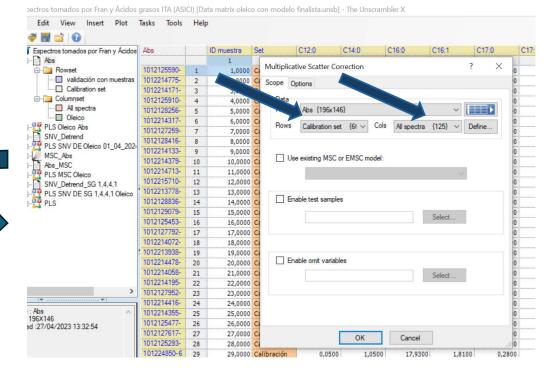
Multiplicative Scatter Correction (MSC)

Multiplicative Scatter Correction (MSC) is a transformation method used **to compensate for pure additive and/or multiplicative effects** in spectral data.



Multiplicative Scatter Correction (MSC)





Derivatives

Derivatives are applied to correct for baseline effects (baseline shifts) in spectra for the purpose of removing nonchemical effects. Derivatives may also aid in resolving overlapped bands which can provide a better understanding of the data, emphasizing small spectral variations not evident in the raw data.

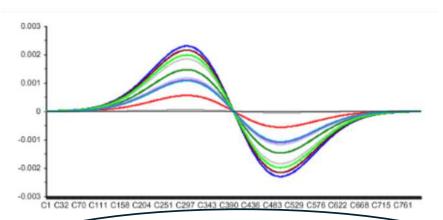
Raw data

0.9 0.8 0.7 0.6 0.5 0.4 0.3 0.2 0.1 0 C1 C32 C68 C108 C153 C198 C243 C288 C333 C378 C423 C468 C513 C558 C603 C648 C693 C738 C783

The baseline offset has been removed under derivatization

The peak maxima in the raw data has now become a zero point in the derivative (The zero point can be explained by the fact that at a peak maxima (minima), the derivative is zero).

First derivate

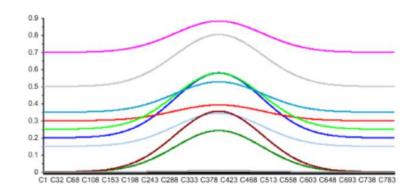


The first derivative eliminates CONSTANT baseline shifts

Derivatives

Derivatives are applied to correct for baseline effects (baseline shifts) in spectra for the purpose of removing nonchemical effects. Derivatives may also aid in resolving overlapped bands which can provide a better understanding of the data, emphasizing small spectral variations not evident in the raw data.

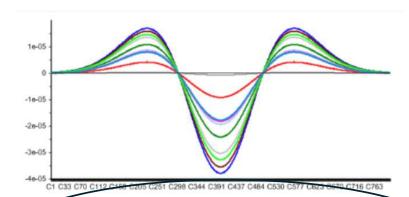
Raw data



The second derivative is a measure of the change in the slope of the curve. In addition to removing pure additive offset, it is not affected by any linear "tilt" that may exist in the data.

The second derivative can help resolve nearby peaks and sharpen spectral features

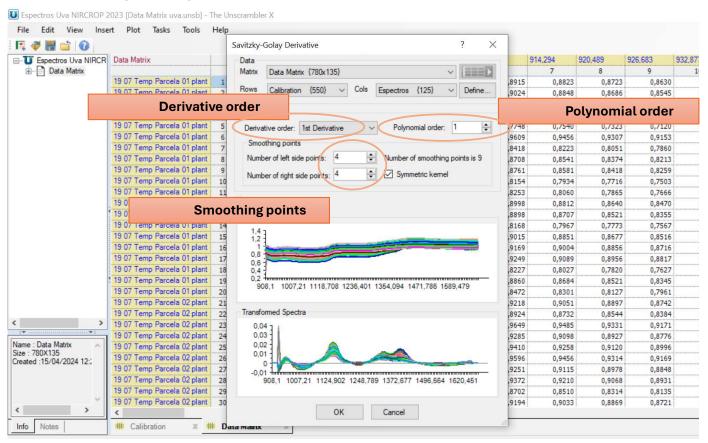
Second derivate

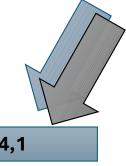


The **second derivative** eliminates **shifts** that **vary linearly with** wavelength

Derivatives

The Savitzky-Golay method

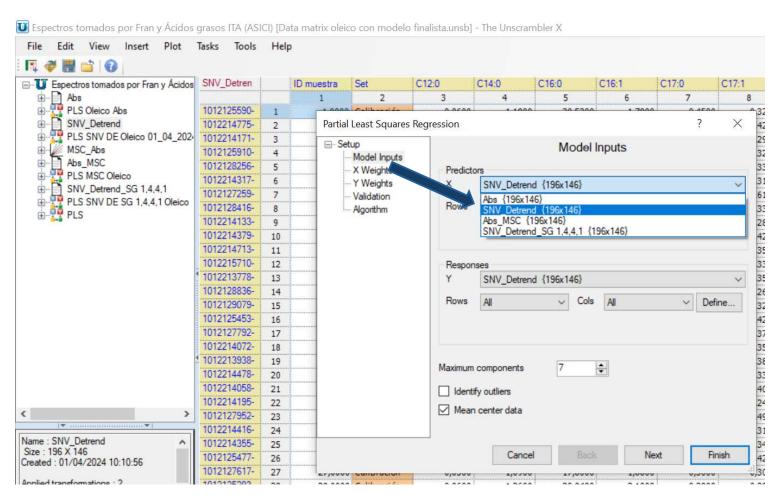




1,4,4,1

2,5,5,2

Remake the model on the pretreated matrix



Pretreatments are only applied to spectral variables

Pretreatments are applied to the calibration set

The selection of the most appropriate one is purely experimental



Archivos de Zootecnia

Comparación de herramientas quimiométricas de clasificación para la identificación de grasa perirrenal en corderos

Agudo, B.1; Delgado, J.V.2; López, M.M.3 y Rodríguez, P.L.40

¹Ovino del Suroeste Soc. Coop. Ltda. (OVISO), Villanueva de la Serena, Badajoz, España.

²Departamento de Producción Animal, Facultad de Veterinaria, Universidad de Córdoba, España.

³Departamento de Producción Animal, CYCITEX, Finca la orden, Gobierno de Extremadura, España.

**Oepartamento de Producción Animal y Ciencia de los Alimentos, Facultad de Veterinaria de Cáceres, España.



España - Portugal



Desarrollo de modelos predictivos mediante tecnología NIRS

Tecnologías avanzadas, innovadoras y digitales para el sector agroalimentario de la EUROACE

Évora, 18th and 19th, novembre, 2024