





ModWoodLife

NIR & WOOD – SOUNDS GOOD! #2

Monitoring thermally modified wood performance by NIR. Case of study: surface treatment

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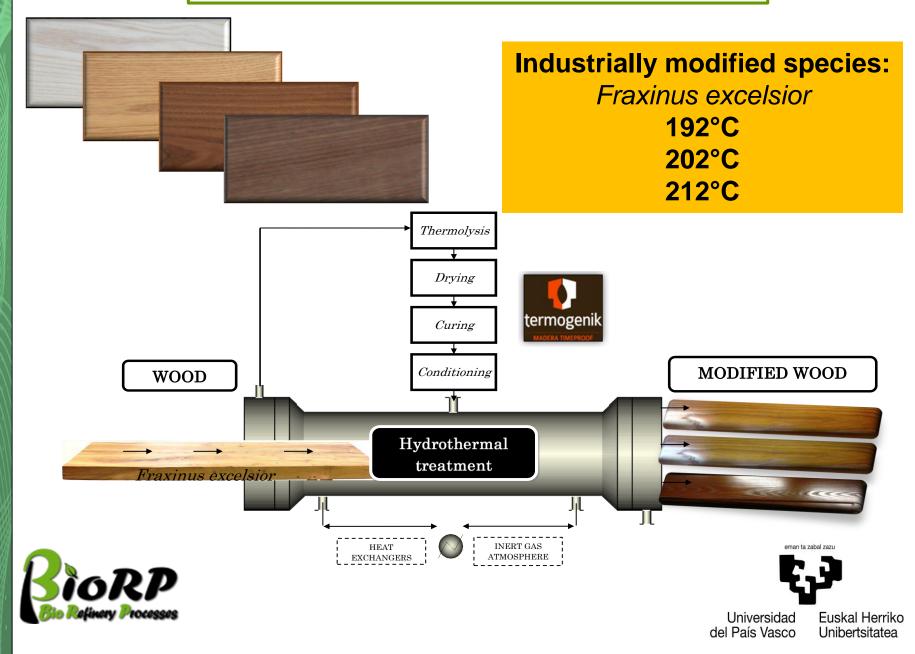




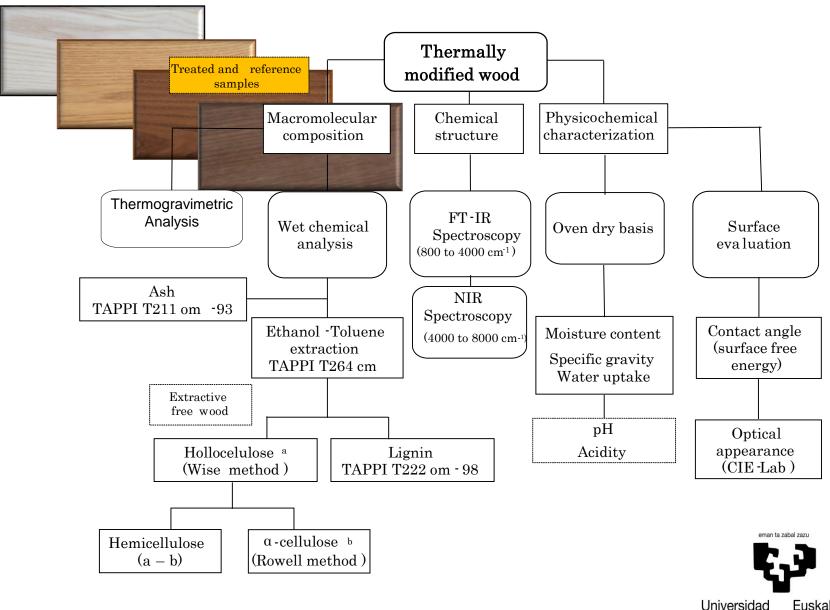


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Monitoring thermally modified wood

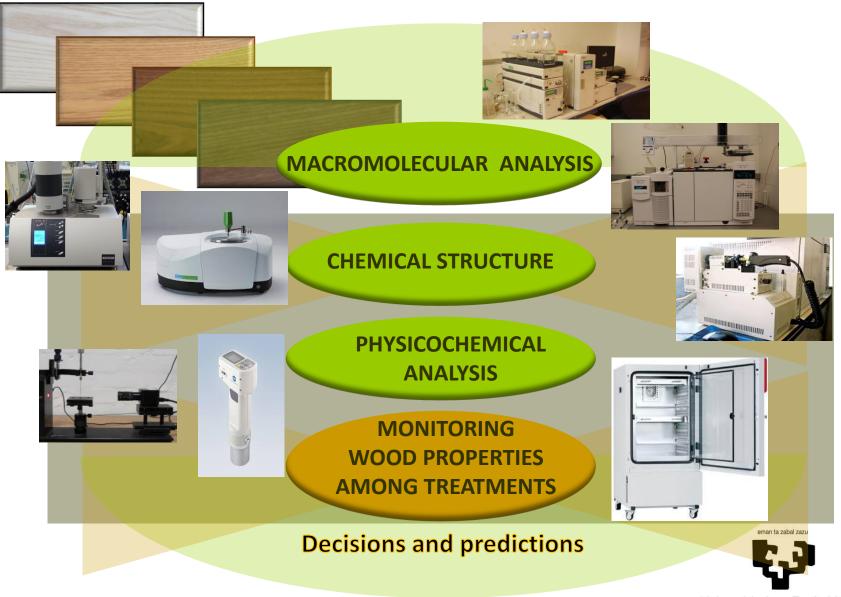


Monitoring thermally modified wood

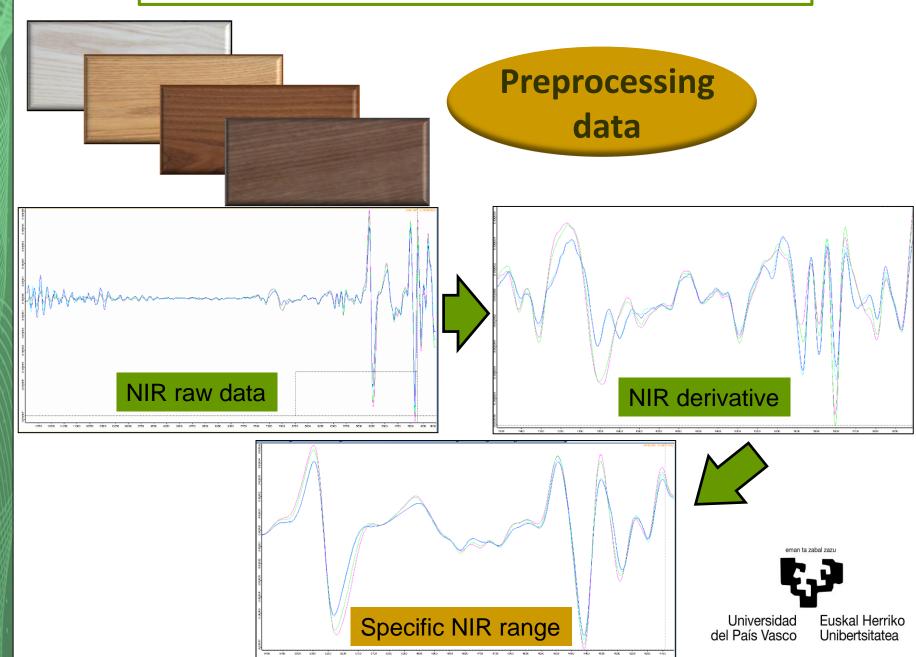


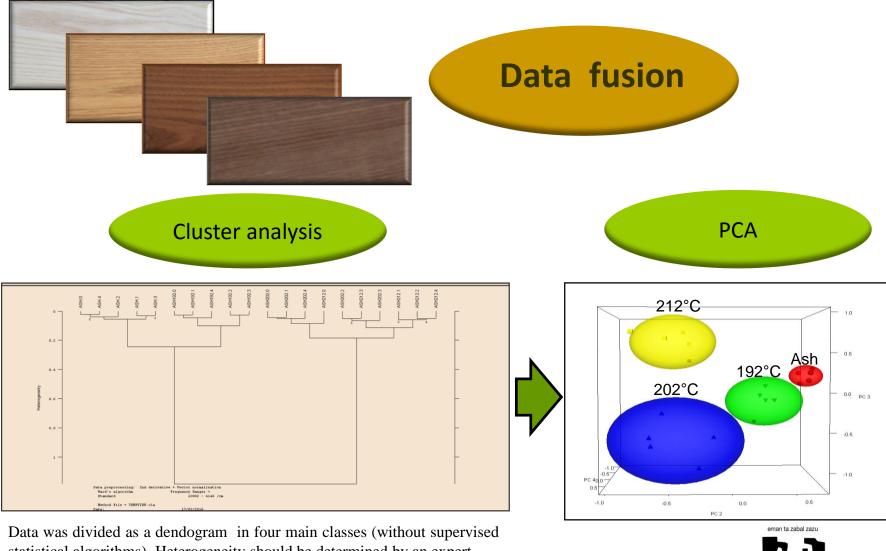
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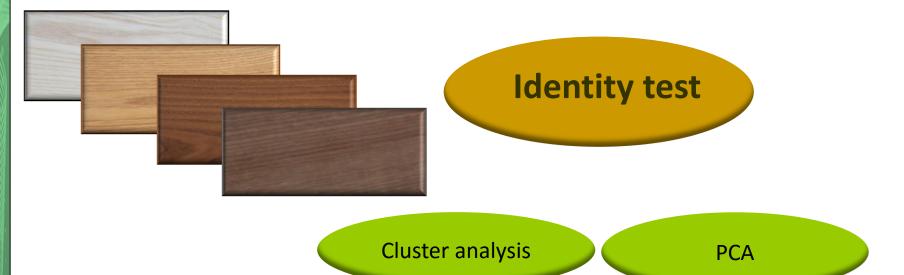




statistical algorithms). Heterogeneity should be determined by an expert.

Principal component analysis decomposed highly correlated data in 4 plotted PC

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	ID	Group1	Group2	IP-Level	s	Threshold1
-	1	ash	ash192	IP1: New	0.951993	0.141386
Γ		"	ash202	IP1: New	1.906599	"
		"	ash212	IP1: New	2.327233	
-	2	ash192	ash	IP1: New	0.952004	0.375173
		"	ash202	IP1: New	0.977602	
		"	ash212	IP1: New	1.648072	"
 -	3	ash202	ash192	IP1: New	0.977605	0.586747
Γ		"	ash212	IP1: New	1.383708	"
		"	ash	IP1: New	1.906604	"
	4	ash212	ash202	IP1: New	1.383700	0.390028
		"	ash192	IP1: New	1.648064	"
Ì		"	ash	IP1: New	2.327225	

It is possible to discover trends when playing with sensor data and validating clusters within threshold (S>1) PCA and CA can be useful for differentiation of samples and classified unmeasured samples into defined groups

Sample name: wo Sample: D:NIR Re Data and time ima		5 IGMT+1			
	Rene only ash 5 spectra.faa	5 (GMT+1)			
Hit no.	Sample name	Hit qual.	Threshold	Group	
1 wood		0.07973	0.14139	ash	
2 wood		0.42296	0.37517	ash192	
3 wood		1.19121	0,39003	ash212	
4 wood		1.30935	0.58675	ash202	
IDENTIFIED AS	ash				
100 C	OK				



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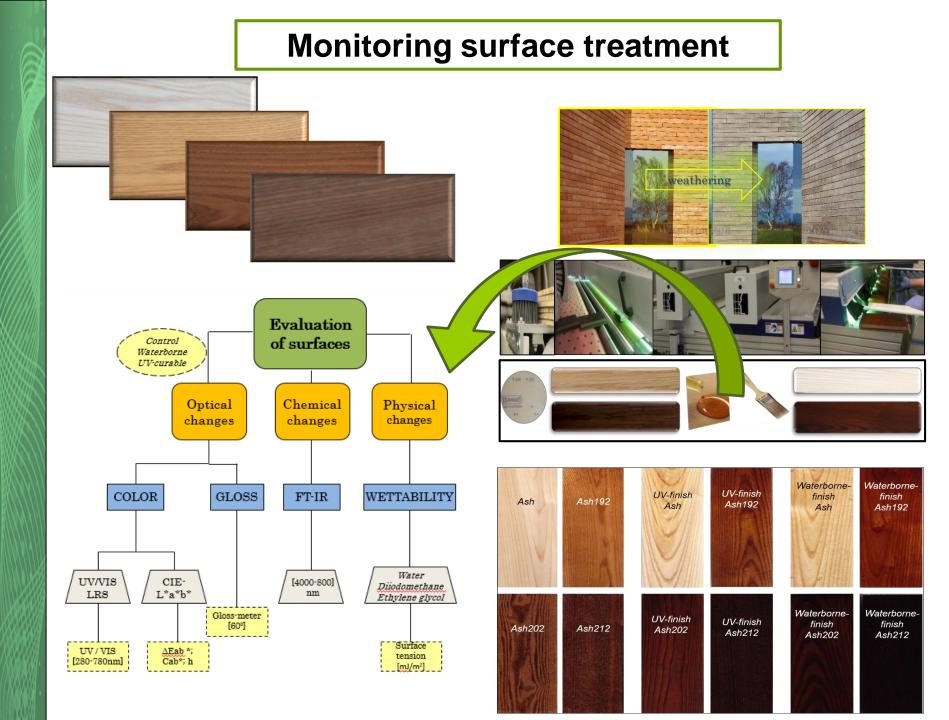
									ial Least uares		
Calibration data set									PLS		
Analysis	Lignin c	x-Cellulose	Hemice	llulose	Extracts	R2 93.59 RMSECV: 0.202	Prediction vs True / liquin [5] / Cross Validation	R2 98.99 RMSECV: 0.339	Prediction vs True / hemicelillose [5] / Cross Validation	112 SELIS PIMSEEV 0.335	Prediction vs True / extractives [5] / Cross Validation
Ash	28.88	45.38	28.	88	4.69	RPD: 15.7 Bias: -0.0018	34 33 32 31	8PD: 3.33 Biac: 0.00775		10100 5-46 Bioc 	0.6
	29.13	43.81	29.13 35.12		6.11	E Live		E Line		E Live	
	35.12	43.93			8.06	FT Color	28 28 5 29 29 5 30 30 5 31 31 5 32 32 5 33 33 5 34 34 5 35 36 5 36 Prediction vs. Ima / cellulose 151 / Gross Validation	E Color	10 11 12 13 14 15 16 17 18 19 20 Prediction vs True / acidity (meg)g) / Cross Validation	E Edor	Prediction of True / contact ands / Cross Validation
	Ash212 35.54 38.14 35.54 9.49 Ash212 Density Asidity rdl Mark Contac					HIS RHSECV. 1.30 HITC: 2.17 Bion HITC: 2.17 Bion HITC: 2.17 Bion HITC: 2.17		BILTS PARSETV CO0254 HPD: 7.39 Bias: -0.000108 Con-		IS 26 RHSECV. 0.001 Pro- 203 Bioc -0.00105 Filme Filme	
	Density	Acidity	рН	WWA	t angle	R2. 99.96	Prediction vs True / WWA [5] / Cross Validation	R2. 95.44	Prediction vs True / density [q/cm3] / Cross Validation	R2 81.84	Prediction vs True / pH / Cross Validation
Ash	0.69	6.74 10 ⁻²	4.55	54.65	61.31	PMSECV: 0.143 RPD: 63.0	535 515 485 485	RMSECV: 0.00509 RPD: 4.7	0.6865	RMSECV: 0.134 RPD: 2.25	
Ash192	0.68	4.35 10 ⁻²	4.55	34.59	99.96	Bias: 0.00046	48.5 43.5 99.6	Bias: 0.000438	0.0000	8ias: -0.00524	49 40 47 46
Ash202	0.65	2.79 10 ⁻²	4.75	33.13	99.19	E Line	35.5 33.5 31.5 31.5 33.5 35.5 37.5 39.5 41.5 43.5 45.5 47.5 49.5 51.5 53.5 65.5	E Line	0.6365	E Line	45 44 43 451 461 471 481 491 501 511 521 531
Ash212 0.63 1.59 10 ⁻² 5.32 33.03 97.22 Models PLS Depending on the number of available samples two emanta zabal zazu										eman ta zabal zazu	
The development of Partial Least Squares									ssible: cross-validation		
	(PLS) model starts with computation of PCs					test-set validation.				1 27	

(PLS) model starts with computation of PCs on the base of calibration dataset.

The PLS model has to be validated after calibration.

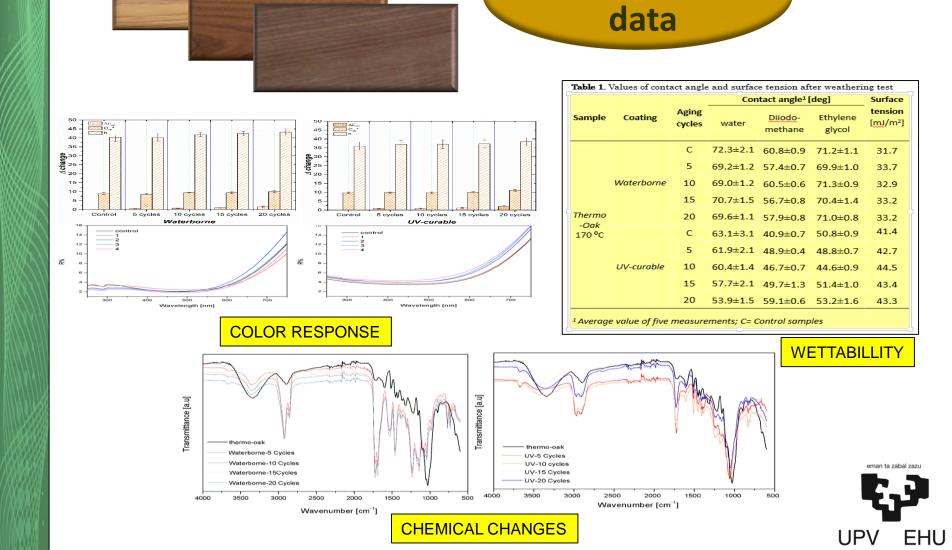
The high value of R2 and small value of RMSEP indicates excellent PLS models.

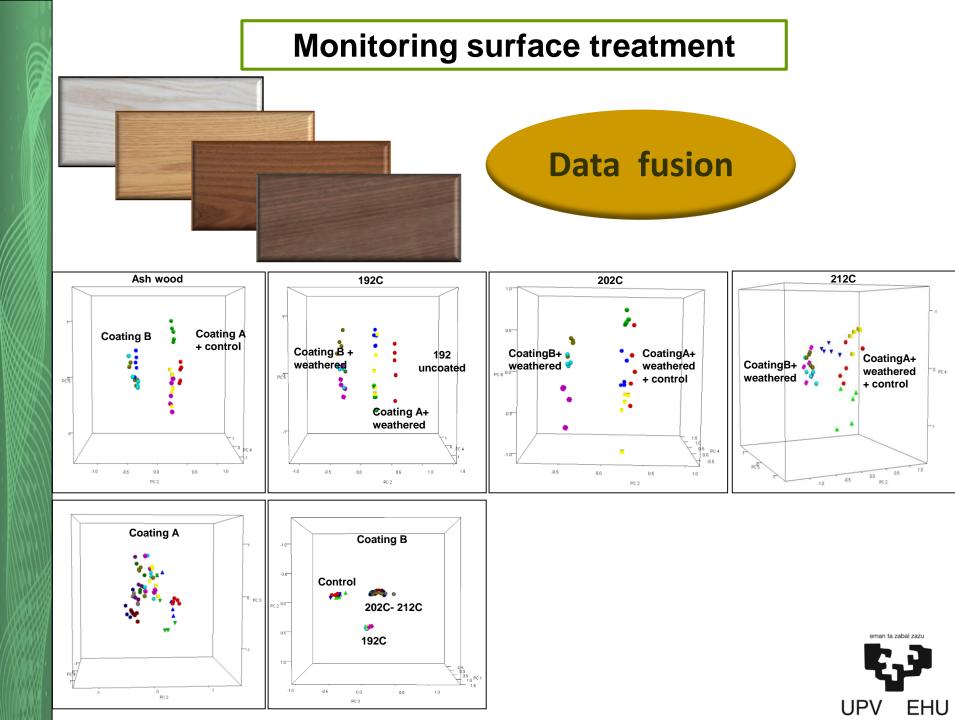
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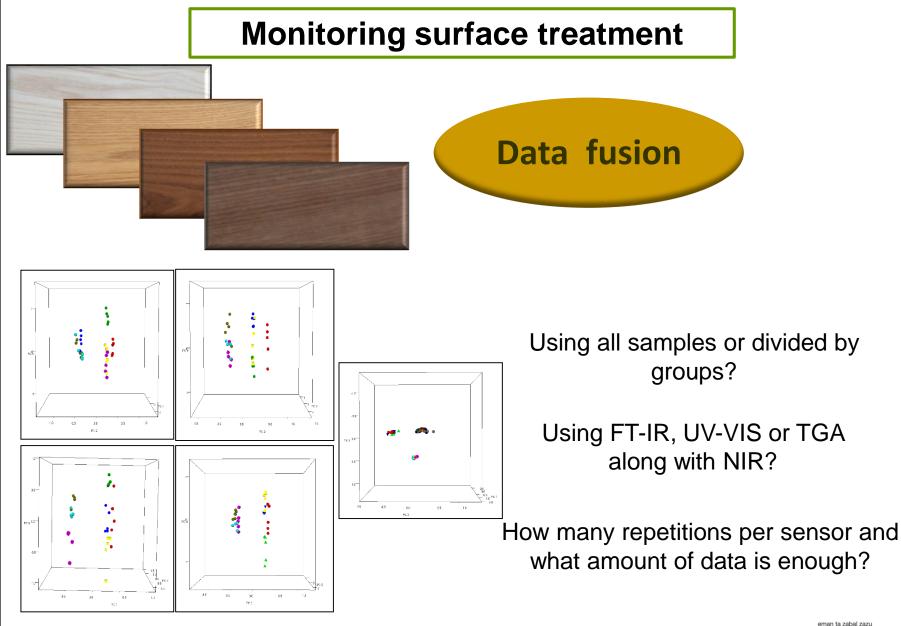


Monitoring surface treatment

Preprocessing







TO BE CONTINUED...









THANK YOU FOR YOUR ATTENTION!



