

NIRS uses for breeding purposes

- ⌘1) Breeding Purposes:
Targets and means.
- ⌘2) Laboratory Samples Analysis :
For Ground or Whole seeds
- ⌘3) On Line NIRS Plots Analysis :
For Fresh forage or fresh kernels.
- ⌘4) NIRS as a breeding tool, summary.

NIRS uses for breeding purposes: 1) Breeding Purposes: Targets and means.

⌘ 1.1) Summary of Main Breeding Crops Targets:

Breeding Purposes <i>Crops Targets</i> Summary:				
<i>Species</i>	<i>Fields Criteria</i>	<i>Maturity</i>	<i>Quality criteria</i>	<i>Process</i>
<i>Field Crops Grain:</i>				<i>Food & Feed</i>
Wheat (s)	Yield, diseases, lodging	%Water	Protein, Starch	Bakery
Barley (s)	Yield, diseases, lodging	%Water	Protein, Starch	Brewery
Peas	Yield, diseases, lodging	%Water	Protein, Starch,	Crushing
Maize	Yield, diseases, lodging	%Water	Starch, Protein, Oil	Dry & Wet Milling
Rapeseed	Yield, diseases, lodging	%Water	Oil, Protein, Gluco, F A	Crushing
Sunflower	Yield, diseases, lodging	%Water	Oil, Protein, Fatty Acids	Crushing
Soya	Yield, diseases, lodging	%Water	Oil, Protein	Crushing
<i>Field Crops Forages:</i>				<i>Feed</i>
Maize	Yield, diseases,	%HDM	Digest, NDF, Protein	Silage
Grass	Yield, diseases,	%HDM	Digest, NDF, Protein	Hay & Silage
Legume	Yield, diseases,	%HDM	Protein, Digest, NDF	Hay & Silage
<i>Vegetables Crops:</i>				<i>Food</i>
Roots	Yield, diseases	%Water	Sugars, colour, life	Fresh & Canned
Fruits	Yield, diseases	%Water	Sugars, colour, life	Fresh & Canned
Others	Yield, diseases	%Water	

NIRS uses for breeding purposes:
1) Breeding Purposes: Targets and means.

☼ 1.2) Summary of Main Breeding Means:

☼ Breeding means use either natural (Germplasm) or biotech tools...

Breeding Purposes <i>Crops Targets and means</i> Summary:				
<i>Fields Criteria</i>	<i>Estimation</i>	<i>Method</i>	<i>End use</i>	<i>Comments</i>
Yield	Ton/ha	Plot Weighting	Farmer Income	with an average yearly increase around 1%
Maturity	%HDM %Water	Fresh & Dry Weight Oven drying Or Electric Resistivity	Geographical planting & harvesting	Indicate maturity group area Earliness
Diseases	% Attack	Hand counting	Yield limitation	More pressure to resist to more pests (bacteria, viruses...) and insects (borer, worms...) attacks
Lodging	%Fall	Hand counting	Yield limitation	Resistance to stem breaking from wind storm
Quality criteria		a) Biochemistry		
Protein,	%DM	Kjeldahl,	Bakery Brewery Feed	Highest value for Wheat Lowest value for Barley Lysine complementation
Amino Acids	%DM	HPLC	FOOD	extraction, Omega 3 ratio Regulation Energy Intake
OIL, Fatty Acids Glucosinolates	%DM %DM %FA M. notes	SOXTEC GC HPLC	FEED	Energy intake Starch recovery, Starch functionality Taste, Energy recovery
Starch, Amylose Soluble Sugar	%DM %DM % Starch %DM	Ewers, Extract + Iodine Coloration	FEED Wet Milling ---	Celluloses Cell Wall contents
Fibre, NDF, ADF, ADL	%DM %DM	Weende, Van Soest	FEED	In Vitro Digestibility
Cellulasis IVDM	%DM	DeBoever, Aufrere	FEED	

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NIRS uses for breeding purposes:
1) Breeding Purposes: Targets and means.

☼ 1.3) Summary of Main Breeding Means:

☼ Process methods are time consuming !

Breeding Purposes <i>Crops Targets and means</i> Summary:				
<i>Quality criteria</i>	<i>Estimation</i>	<i>Method</i>	<i>Comments</i>	
d) Process Method		FOOD		
BREWERY	% Atls	MALT Extraction	Barley uses	
BAKERY: W, P, L or other European References ...	% Atls	Chopin Bread Making Sedimentation Tests	Wheat uses for different European bread needs...	
DRY Milling	% Atls	LAB Mini Mill Grinding	% Flour or Semolina on Wheat & Maize recovery	
Flour or Semolina quality	% Atls	Particules sizes distribution by sieves	Wheat & Maize final uses (Flakes/cakes)	
WET Milling	% Atls	Starch extract functionalities	Maize Starch industrial uses (coating, ...)	
CRUSHING	% Atls	LAB Oil Seeds Crusher	Sunflower oil extraction	
CANNED FOOD	% Atls	LAB process	Fresh pea industrial uses...	
e) Process Method		FEED		
In Vivo Digestibility	% DM	From Beef, Sheep	Silage & Hay % off efficient use by ruminants, On Maize, grass & legume	
In Vivo Energy	MI	Calorimetric bomb On before/after intakes	Metabolisable energies for Pig & poultry On Cereals Grains	
....				

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**NIRS uses for breeding purposes:
1) Breeding Purposes: Targets and means.**

⌘ **1.4) Summary of Main Breeding Means:**

- ⌘ For all those criteria this means hundred of thousand of breeding lines & next to be varieties plots to be :
 - ⊗1) Planted,
 - ⊗2) Observed in many location with climatic stress,
 - ⊗3) Selfed or crossed, or open pollinated,
 - ⊗4) Harvested by hand (nurseries) or by harvesters/choppers (plots)
 - ⊗5) **Crops samples have to be collected, treated and analysed in a few weeks just after harvest.**
 - ⊗6) **Inbred seeds quality must be checked before next planting in winter (south hemispha) or spring...**
- ⌘ **All new varieties must fit end users needs for food & feed. Most process in formations are connected to biochemistry analysis**
- ⌘ **Quality criteria must be checked from very early stage of breeding process.**

**NIRS uses for breeding purposes:
1-5) Breeding Purposes: Breeding schema, Summary.**

Year	Source	E.Generation	I.%Purity	Nursery	Plots	Field Criteria	Quality Criteria	
1	GermPlasm R 0	R 0	0.1%	10		Diseases, Lodging, Maturity	Biochemistry	
2	GermPlasm R 1	R 1	0.1%	100		Diseases, Lodging, Maturity	Biochemistry	
3	GermPlasm R 2	R 2	0.1%	1 000		Diseases, Lodging, Maturity	Biochemistry	
1	Cross 1	C 1	C 1	50.0%	100			
	Cross 2	C 2	C 2	25.0%	50			
2	Cross 3	C 3	C 3	12.5%	25			
1	Self	S 1	S 1	50.0%	100			
2	Self	S 2	S 2	75.0%	50			
3	Self	S 3	S 3	97.5%	30	5 000	Yield, Diseases, Lodging, Maturity	Biochemistry
4	Self	S 4	S 4	98.0%	20	2 000	Yield, Diseases, Lodging, Maturity	Biochemistry
5	Self	S 5	S 5	98.4%	10	1 000	Yield, Diseases, Lodging, Maturity	Biochemistry
6	Self	S 6	S 6	98.8%	10	1 000	Yield, Diseases, Lodging, Maturity	Biochemistry
7	Self	S 7	S 7	99.2%	10	1 000	Yield, Diseases, Lodging, Maturity	Biochemistry
8	Self	S 8	S 8	99.6%	10	1 000	Yield, Diseases, Lodging, Maturity	Biochemistry, Process
	Hybrids	H 1	H 1	100%	100		Yield, Diseases, Lodging, Maturity	Biochemistry, Process
9	Test 1	Premium tests		80	5 000	Yield, Diseases, Lodging, Maturity	Biochemistry, Process	
10	Test 2	Final Tests		60	2 000	Yield, Diseases, Lodging, Maturity	Biochemistry, Process	
11	Test 3	Official Tests						
12	Test 4	Official Tests		10	1 000	Yield, Diseases, Lodging, Maturity	Biochemistry, Process	
1		Commercial use			100	Yield, Diseases, Lodging, Maturity	Biochemistry, Process	
2		Commercial use			1 000	Yield, Diseases, Lodging, Maturity	Biochemistry, Process	
3		Commercial use			2 000	Yield, Diseases, Lodging, Maturity	Biochemistry, Process	
4		Commercial use			1 000	Yield, Diseases, Lodging, Maturity	Biochemistry, Process	
5		Commercial use			100	Yield, Diseases, Lodging, Maturity	Biochemistry, Process	

NIRS uses for breeding purposes

- ⌘ 1) Breeding Purposes:
Targets and means.
- ⌘ 2) Laboratory Samples Analysis :
For Ground Dry Matter or
Intact Whole seeds (non destructive)



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NIRS uses for breeding purposes :

2-1) Powder Analysis for Forages & Ground Grain and for Small Grain Whole Seeds Analysis:

- ⌘ 2.11) Main fields crops: Silage Maize, Wheat, Barley, Rape seeds, Sunflower, Grass....



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NIRS uses for breeding purposes :
2-121) Powder Analysis for Forages & Ground Grain
and for Small Grain Whole Seeds Analysis:



- ⌘ **A) Forages & Grain Samples**
preparation: Drying & Grinding
with hammer mill (D. 1mm holes)



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NIRS uses for breeding purposes :
2-122) Powder Analysis for Forages & Ground Grain
and for Small Grain Whole Seeds Analysis:

- ⌘ **B) Scanning:** A volume of less than 5 cc of forage/grain powder or
small Whole Seed is enough for NIRS spectra with the Mini cell :



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**2-1) NIRSystem +Auto Sampler:
2.13) Powder Analysis & Small Whole Seeds Analysis
(Rape, Wheat, Barley...)**

- ⌘ An average of 300 cups or samples can be recorded per day !



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**NIRS uses for breeding purposes :
2-14) Powder Analysis for Forages or Ground Seed,**

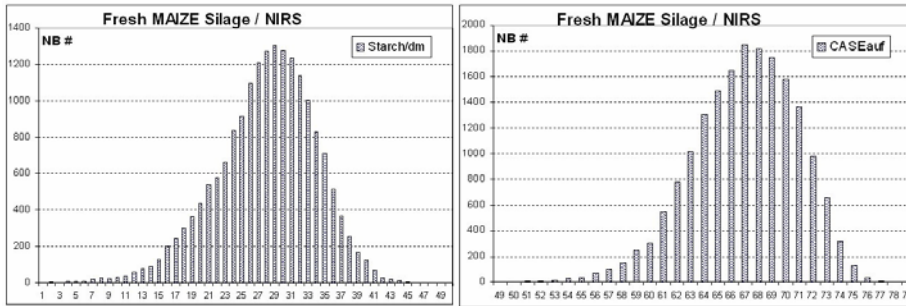
- ⌘ Powder NIRS analysis :
- * Need drying and grinding of fresh forage samples.
 - * Is a destructive method for all ground grain !
Only Homogeneous grain samples must be ground...
- ⌘ Mean spectra of Ground samples are done in one minute:
- * Sample preparation is similar to biochemistry process.
 - * NIRS precision (SECV) is very close to wet chemistry and slightly better than whole Seeds analysis.
- ⌘ Prediction of either biochemistry or process value on :
- * Plants parts such as Stover+leaves & ear or grain.
 - * Experimental open pollinated plot samples.
 - * Samples from Grain or Silage silo as final products (variety)

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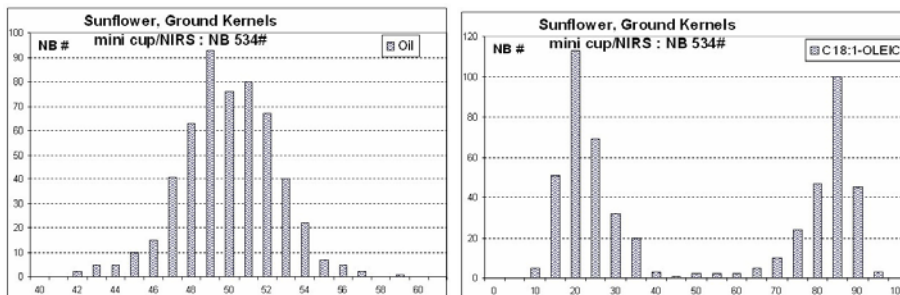
NIRS uses for breeding purposes :
2-15) Powder Analysis for Forages or Ground Seed,

- ⌘ Example #1: NIRS Prediction distribution for Fresh Silo Maize Samples (dry & ground) on 18 000 #
- ⌘ SECV : Starch/dm = 1.6 Cellulase IV/dm = 2.0



NIRS uses for breeding purposes :
2-16) Powder Analysis for Forages or Ground Seed,

- ⌘ Example #2: NIRS Prediction distribution for Ground Sunflower Samples on 534 #
- ⌘ SECV : Oil/dm = 1.10 Oleic/FA = 3.2



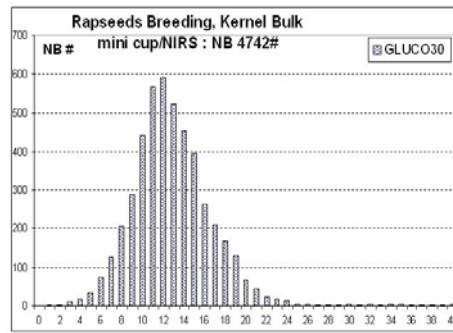
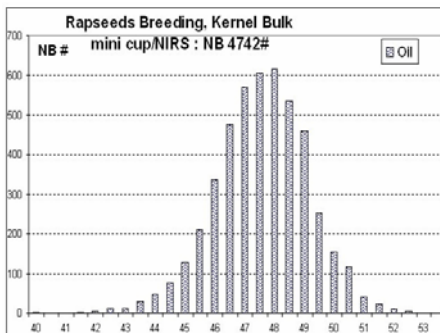
NIRS uses for breeding purposes :
2-2) Whole Seeds Analysis for Large Grain Crops
(Maize, Soy, Bean, Pea ...) :

- ⌘ NIRS analysis on whole grain is :
 - * a non destructive method !
 - All Selfed ear seed are send back to the breeder !*
 - * a very quick and cheap application.
 - ⌘ Mean spectra of Bulk analysis for:
 - * Small Seeds vial with 5 cc only
(=1 ear grain size on wheat or barley).
 - * Large Seeds vial with 15 to 25 cc.
 - They are recorded in 1' to 1'1/2.
 - ⌘ Prediction of either biochemistry or process value on :
 - * Selfed ear shelled grain,
 - * Experimental plot harvested grain Sample,
 - * Grain Silo or bin samples as end products.
- NB: Industrial Process value are only available for Whole seed!*



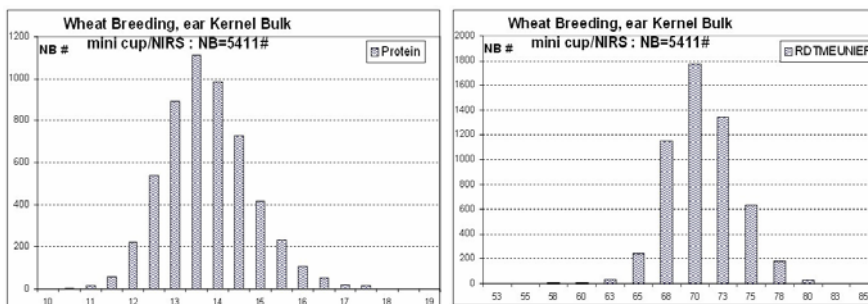
NIRS uses for breeding purposes :
2-21) Whole Seeds analysis on small grain

- ⌘ **Example #3:** NIRS Prediction distribution for Rape Seeds Samples on 4 742 #
- ⌘ **SECV :** Oil/dm = 1.10 Glucosinolates M.moles = 3.2



**NIRS uses for breeding purposes :
2-22) Whole Seeds analysis on small grain**

- ⌘ Example #4: NIRS Prediction distribution for Wheat Grain breeding on 5 411#
- ⌘ SECV : Protein/dm = 0.35 Mill Recovery = 3.3



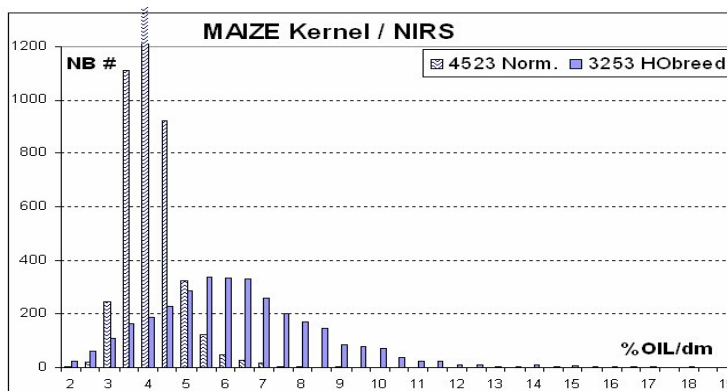
**NIRS uses for breeding purposes :
2-23) NIRSystem +Transport Module:
Whole Seeds Analysis for Large Grain Crops (Maize, Soy, Bean, Pea ...) :**

- ⌘ An average of 250 Maize samples can be recorded per day with the large cell.



NIRS uses for breeding purposes :
2-24) Whole Seeds Analysis for Large Grain Crops

- ⌘ **Example #5: NIRS Prediction distribution for Maize Grain Samples on Whole Seed bulks:**
- ⌘ left = 4 523 normal field samples (grey line),
- ⌘ right= 3 253 ear to row "HO breed" (blue lines) SECV : Oil/dm = 0.50



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NIRS uses for breeding purposes :
2-3) NIRS Single Seeds Analysis



For S.S.A. on large size grain crops two apparatus are available, depending on :

- ⌘ Low Number of kernels per vial (<300K):

-> NIRSystem + auto sampler + Mini cell.

An average of 350 hand sorted kernels in 8 hours for:

- * Oil, protein, starch, Amylose...
 - * Or others fine biochemistry contents.
- All steps are done by hand...*

- ⌘ For High Number of kernels per vial

(>300K to 3 kg) we must use the
 -> BRIMROSE SEED MEISTER



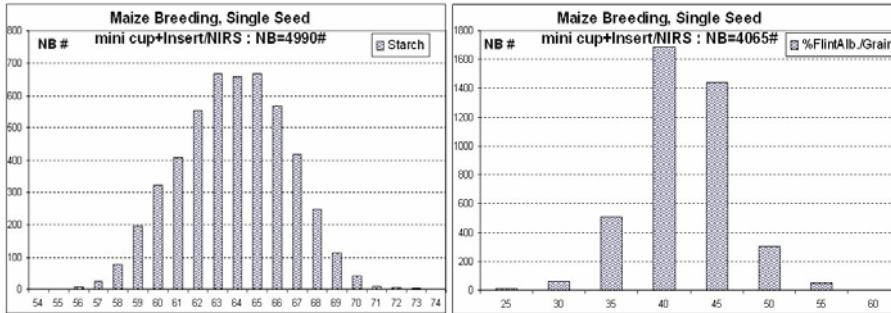
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NIRS uses for breeding purposes
2-31) NIRS Single Seeds Analysis

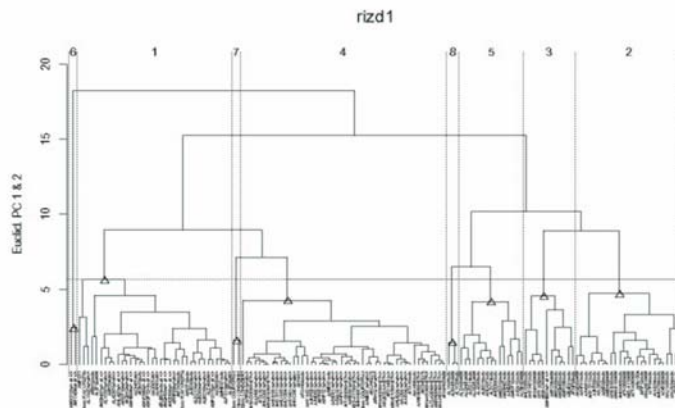
⌘ **Example #6: Maize Kernels, Single Seeds Analysis**
Left to Right : Dent, Dent x Flint, Flint germ plasm

⌘ SECV : Starch/dm = 1.9 %Flint Albumen/WS = 6.0



NIRS uses for breeding purposes
2-32) NIRS Single Seeds Analysis

⌘ **Example #7: NIRS Rice kernels Single Seeds Factorial Discriminant Analysis (cluster)**



NIRS uses for breeding purposes

2-4) High Kernels Number Per Vial (< 3 Kg):



- ⌘ **BRIMROSE Seed Meister :**
An average of 300 automated sorted kernels *per hour* !



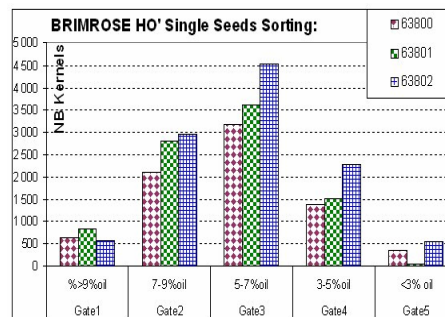
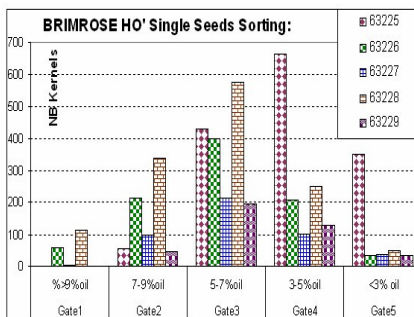
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NIRS uses for breeding purposes

2-41) High Kernels Number Per Vial (< 3 Kg):

- ⌘ **Example #8: NIRS Prediction & Oil sorting for Single Seeds**
on Maize Grain for High Oil Breeding

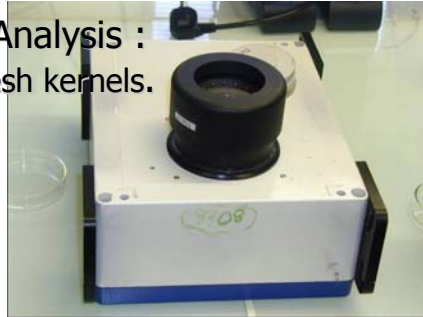


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NIRS uses for breeding purposes

- ⌘ 1) Breeding Purposes:
Targets and means.
- ⌘ 2) Laboratory Samples Analysis :
For Ground or Whole seeds
- ⌘ 3) On Line NIRS Plots Analysis :
For Fresh forage or fresh kernels.



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3) On Line NIRS Plots Analysis :

3-101) NIRS ON LINE for Research Plots Harvester;

ZEISS CORONA Outside Laboratory Analysis:

- ⌘ I) At Line Analysis:
 - * Used for dedicated application on collected samples, when no harvester or chopper are available...
- ⌘ II) One Line Analysis :
 - a) Harvest plot" flow measurement for chopped whole plant or threshed grain :
 - b) Each harvest plot flow is multi scanned & spectra averaged *in less than 3 seconds*, with no added delay while harvesting silage or grain crops...
 - c) Crop Spectra are treated and stored on special on board enforced computer,
 - d) NIRS prediction are done on the spot and used by breeder team the same day, like fresh weight & moisture information.

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3) On Line NIRS Plots Analysis :

3-102) NIRS ON LINE for Research Plots Harvester ZEISS CORONA;

- ⌘ On & At line prediction available for :
 - ☒ *Dry matter* for grass, legume and maize fresh silage on whole plants or Stover.
or *Water contents* for fresh cereals grain (Wheat, Barley, Maize...) and Rapeseeds grain,
 - ☒ *Main biochemistry* variable for fresh silage & cereals grain : protein, oil, starch, fibre...
 - ☒ *Process information* for end users :
 - + Some industrial food quality information for dry milling process on Maize Grain & Wheat.
 - + Some Animal feed process variable for Maize Grain & Wheat.
- ⌘ Reference samples are picked up by automated device on board on request in order to check & update NIRS calibration.
- ⌘ *High Moisture crops PLS NIRS prediction are more difficult to build than on low moisture crops...*

3) At Line NIRS samples Analysis with ZEISS CORONA :

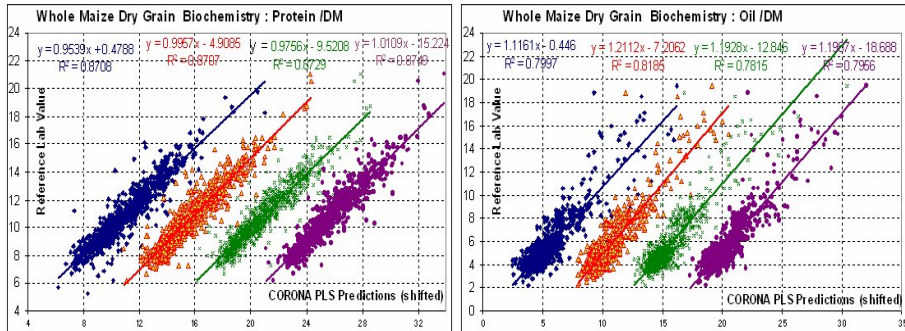
3.111) At Line Applications (near by the oven) for Fresh forages samples or any else!



3) At Line NIRS samples Analysis with ZEISS CORONA::

3.112) Maize Dry Grain same PLS calibration checks on 4 #:

Validation #1: Corona reproducibility, Protein & Oil prediction /4#

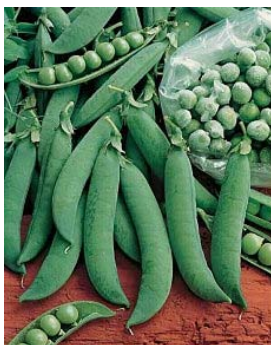


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3) At Line NIRS samples Analysis with ZEISS CORONA::

3.121) Fresh Peas harvests for canned food with At Line application:

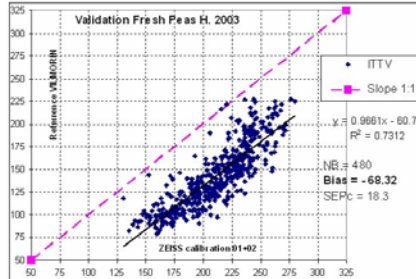
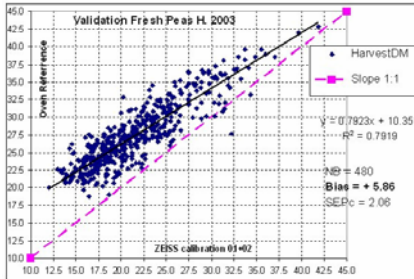


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3) At Line NIRS samples Analysis with ZEISS CORONA::
 3.122) *Fresh Peas harvests for canned food At Line application:*

Validation #2: Validation Results from Harvest 2003:
Dry Matter (Left) & Tenderness (Right)



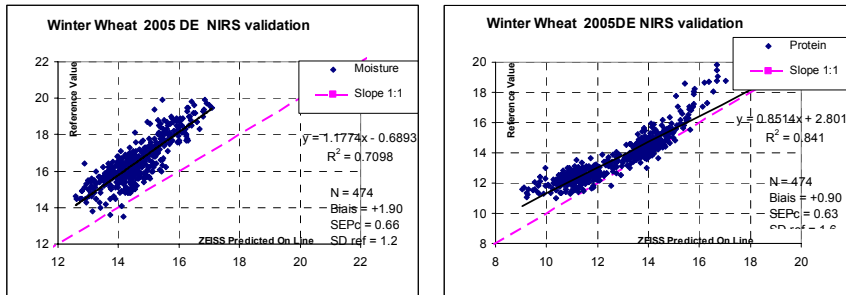
3) At & On Line NIRS samples Analysis with ZEISS CORONA:
 3.131) *For Wheat, Barley, Rape Seeds Samples :*

At Line = Calibration (Left) & On Line = monitoring (Right):



**3) At Line NIRS samples Analysis with ZEISS CORONA:
3.132) Wheat Samples At Line for On Line monitoring:**

Validation #3: Wheat Samples Moisture & Protein prediction for harvest 2005



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3) On Line NIRS Plots Analysis ZEISS CORONA :

3.201) On Line Applications @DLF FR :

Fresh forages plots at DLF Trifolium FR location

=> canopy measurement since 2004 harvests:

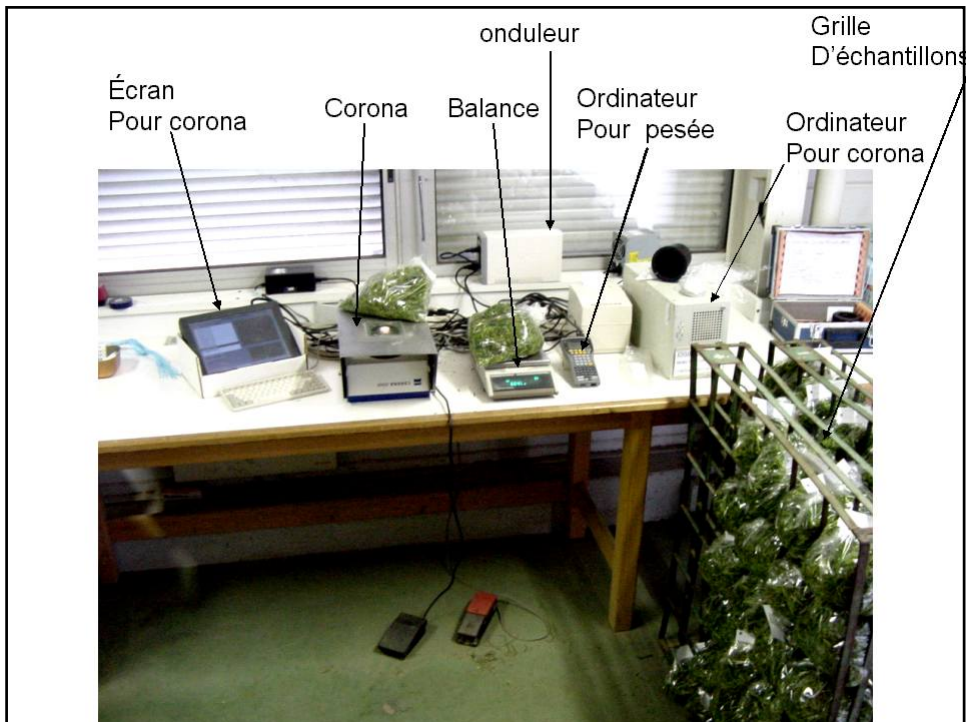
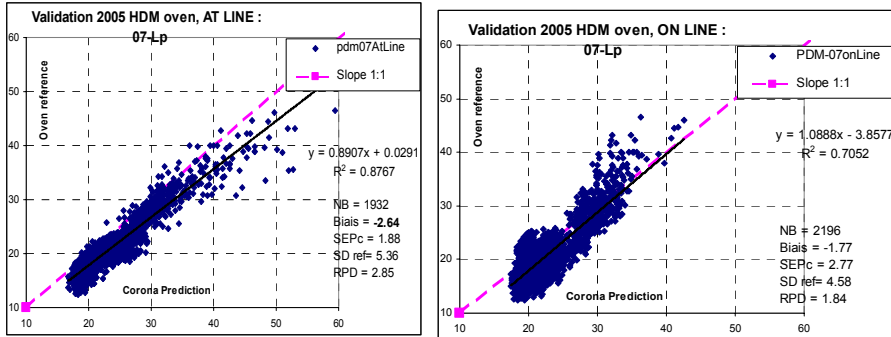


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3) On Line NIRS Plots Analysis ZEISS CORONA :
3.202) DLF FR Harvests 2002-2005, HDM Oven value:

Validation #4 On & At Line HDM validation for Harvest 2005:
 At Line (left) & On Line (right)



3) On Line NIRS Plots Analysis :

3-301) NIRS ON LINE for Research Plots Silage Harvester ZEISS CORONA;

- ⌘ Flow measurement of chopped maize whole plant.



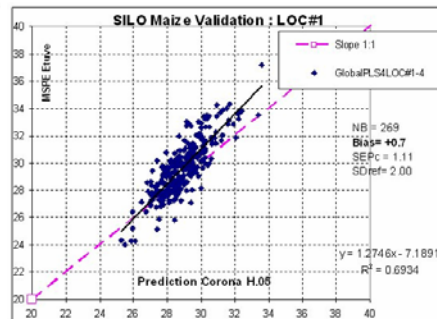
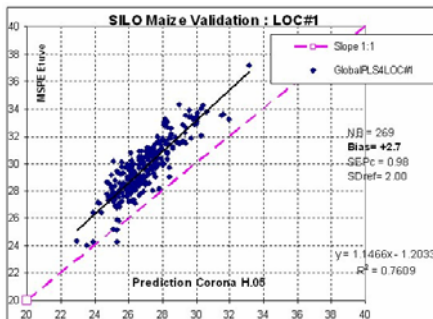
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3) On Line NIRS Plots Analysis ZEISS CORONA :

3-302) Harvest Dry Matter prediction validation

- ⌘ Validation #5: On Fresh Chopped Maize Whole Plants :
- ⌘ Two PLS models "local #1" & "global #2", validation on an Independent set of 270#



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3) On Line NIRS Plots Analysis :

3-40) NIRS ON LINE for Research Plots Grain Harvester **ZEISS CORONA**;



⌘ Flow measurement of whole plot grain flow.

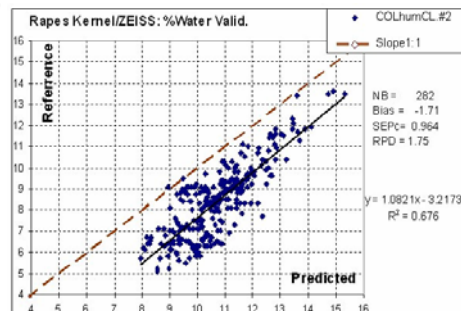
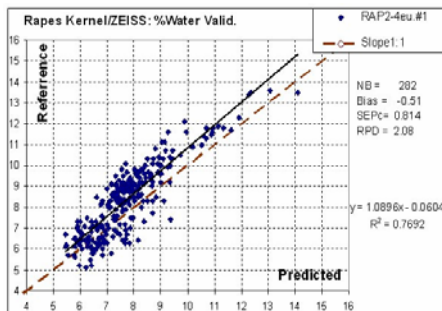


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3) On Line NIRS Plots Analysis :

3-51) Water prediction validation **ZEISS CORONA**

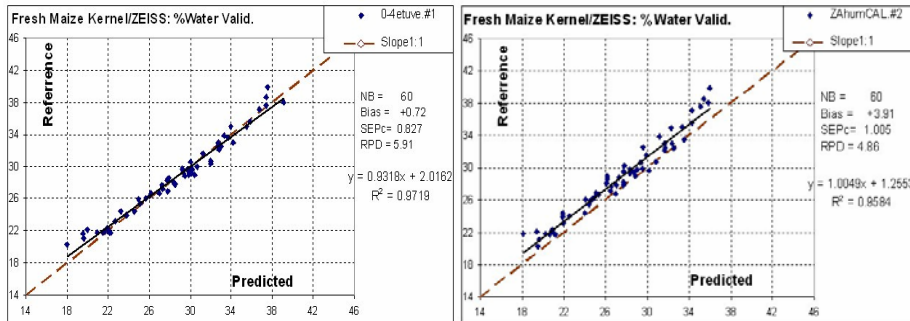
- ⌘ A) Validation #6, On Fresh Rapeseeds :
- ⌘ Two independent PLS models ("DB #1" & "DB #2") validation on an Independent set of 280#.



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3) On Line NIRS Plots Analysis : 3-52) Water prediction validation ZEISS CORONA

- ⌘ B) Validation #7, On Fresh Maize Grain :
- ⌘ Two independent PLS models ("DB #1" & "DB #2") validation on an Independent set of 60#.



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Targets and means.
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For Fresh forage or fresh kernels.
- ⌘ 4) NIRS as a breeding tool, summary.

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4) NIRS uses for breeding purposes, summary.

4-1) *They are many possible Ways to help Breeders!*

⌘ **A) "Plant to plant" analysis:**

- * Fresh chopped dry & ground single maize plant upper parts for digestibility traits before Selfed ear harvest.
- * Nursery Selfed ear grain after harvest for biochemistry content.
- * Single Seeds sorting before planting for biochemistry traits.

⌘ **B) "Plants parts" samples:**

- * Grasses Stem+leaves for cell wall digestibility & husked maize ear,
- * All field crops dry grain, *even ground sunflower!*
- * Vegetable roots or fresh fruits, *carrots, peas, melon, tomatoes...*



4) NIRS uses for breeding purposes, summary.

4-2) *They are many possible Ways to help Breeders!*

⌘ **C) "Plot bulks" or Seed production vials:**

- * Germ Plasm (parents) stored seed bags,
- * Control trials networks samples from harvesters.
- * Whole plot On line analysis from choppers.
- * Vials homogeneity by flow analysis.

⌘ **D) What could be Next, *maybe?***

- * *Single plant (or spotted) analysis in nursery, greenhouse...*



4) NIRS uses for breeding purposes, summary.

⌘ E) Basic rules :

NIRS application can work only if we can bring together :

- **Large Germ Plasm variability on parents for food & feed end use.**
- **Good reference value from laboratory or suitable process.**
- **NIRS cheap & representative spectra on provided samples or flow plots.**
- **Non destructive method for whole or single seeds only.**
- **Efficient NIRS calibration uses and updates for breeding works.**
- **Very quick feed back quality information to breeders for all NIRS routine analysis data (to Breeders' Computer Data Bases).**
- ***NIRS is a team work !***

- *Many thanks to :*
Dr P. DARDENNE CRAW,

a) NIRS ON LINE presentation & publication:

- GEMBLoux, Belgium, November 2004,
- BRAUNSchWEIG, Germany, Marsh 2003,
- Diode array near infrared instrument to analyse fresh forages on a harvest machine.
Dardenne P., Féménias N., 1999, In: Davies A.M.C. and Giangiacomo R. (eds),

Near Infrared Spectroscopy : Proceeding of the 9th Conference. NIR Publications, UK, 121.

•b) Others NIRS publication:

- Prediction of the metabolisable energy value of maize in adult cockerels.
Lessire, M.; Hallouis, J.M.; Barrier-Guillot, B.; Champion, M.; Femenias N.,
In British Poultry Science, 44:813-814 N. 2003..

- Interaction entre la teneur en acides aminés et la... (Les Journées de La recherche Avicole)
_Lessire Michel, Hallouis Jean Marc, Barrier-Guillot Bruno, Orlando D., Champion M., Féménias N.
p253....

- Composition and nutritive value of whole maize plants fed fresh to sheep. I: Factors of variation,
ANDRIEU J.; DEMARQUILLY C.; DARDENNE P.; BARRIERE Y.; LILA M.; MAUPETIT P.; RIVIERE F. ;
FEMENIAS N. In Annales de zootechnie 1993, vol. 42, no3, pp. 221-249;

- Use of NIRS dermination of quality in a silage maize breeding program. Femenias, N., and T.
Ronsin. 1990.. In Proc. Third Intern. Conf. on Near Infrared Spectroscopy. Brussels.

THANK YOU !

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