

CHALLENGE 2012

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Villeneuve d'ascq, 5-6 Décembre 2012

Polytech'Lille, Université Lille1, FRANCE

Identifiant

Mot de passe

Connexion

NAVIGATION

Accueil

Lieu de la conférence

Conférenciers invités

Liste de communications acceptées

Programme CHIMIOMETRIE 2012

Comités ▾

Résumé et inscription ▾

Publication dans CHEMOLAB

Challenge 2012

PRÉSENTATION

Le **Groupe Français de Chimiométrie (GFC)** sous l'égide de la **Société Française de Statistique (SFDS)** a le plaisir de vous annoncer la tenue de son congrès annuel **CHIMIOMETRIE 2012** qui se déroulera du **5 au 6 décembre 2012** à l'École Polytechnique Universitaire de Lille sur le campus de l'Université Lille 1.

Ce congrès vise à rassembler Universitaires et Industriels pour faire le point sur les avancées de la Chimiométrie, allant de la collecte de l'information aussi efficacement et précisément que possible à partir de plans d'expériences, à l'analyse et à la modélisation des données. Ce congrès, qui depuis de nombreuses années réunit théoriciens et expérimentateurs de tout domaine, a pour vocation de promouvoir l'utilisation de la Chimiométrie au-delà même de l'application des Statistiques dans le domaine de la Chimie. C'est un lieu privilégié de rencontres et d'échanges, qui donne la possibilité aux Universitaires et Industriels, sans restriction disciplinaire, de communiquer en faisant état de leurs expériences, de leurs préoccupations et/ou de leurs besoins actuels et futurs.

En cette année 2012, l'accent sera mis sur les disciplines et les thèmes suivants :



DATES IMPORTANTES

Date limite des soumissions :
22 octobre

Fin des inscriptions à tarif réduit :
20 octobre

Décision du comité scientifique,
notifications :



2004= MEAT (NIR) - Non lin

Data download from:

www.chimiometrie.fr

www.chimiometrie.org

dardenne@cra.wallonie.be

2005= STARCH (MIR) – Disc

2006= SOILS (NIR) – Large d

2007= WHEAT (NIR) - <<REF – Robustness

2008= none (CAC)

2009= RAPESEED (NIR) – Extrapolation

2010= BEERS (NIR-MIR-Raman) – Discrimination

2011= FEED (NIR) - Calibration Transfer

Challenge 2012



270 spectra in a CAL set with known concentration of a contaminant

276 spectra in a TEST set to be predicted

The goals:

Obtain the smallest RMSEP

Obtain the higher rate of good classifications;
not contaminated vs contaminated

Send the results including 2 vectors:

1. the quantification
2. the classification: 1 = contaminated 0 = clean

Send the xls file by **November 30** to **dardenne@cra.wallonie.be**

Add a DOC or PPT file with a summary of your approach

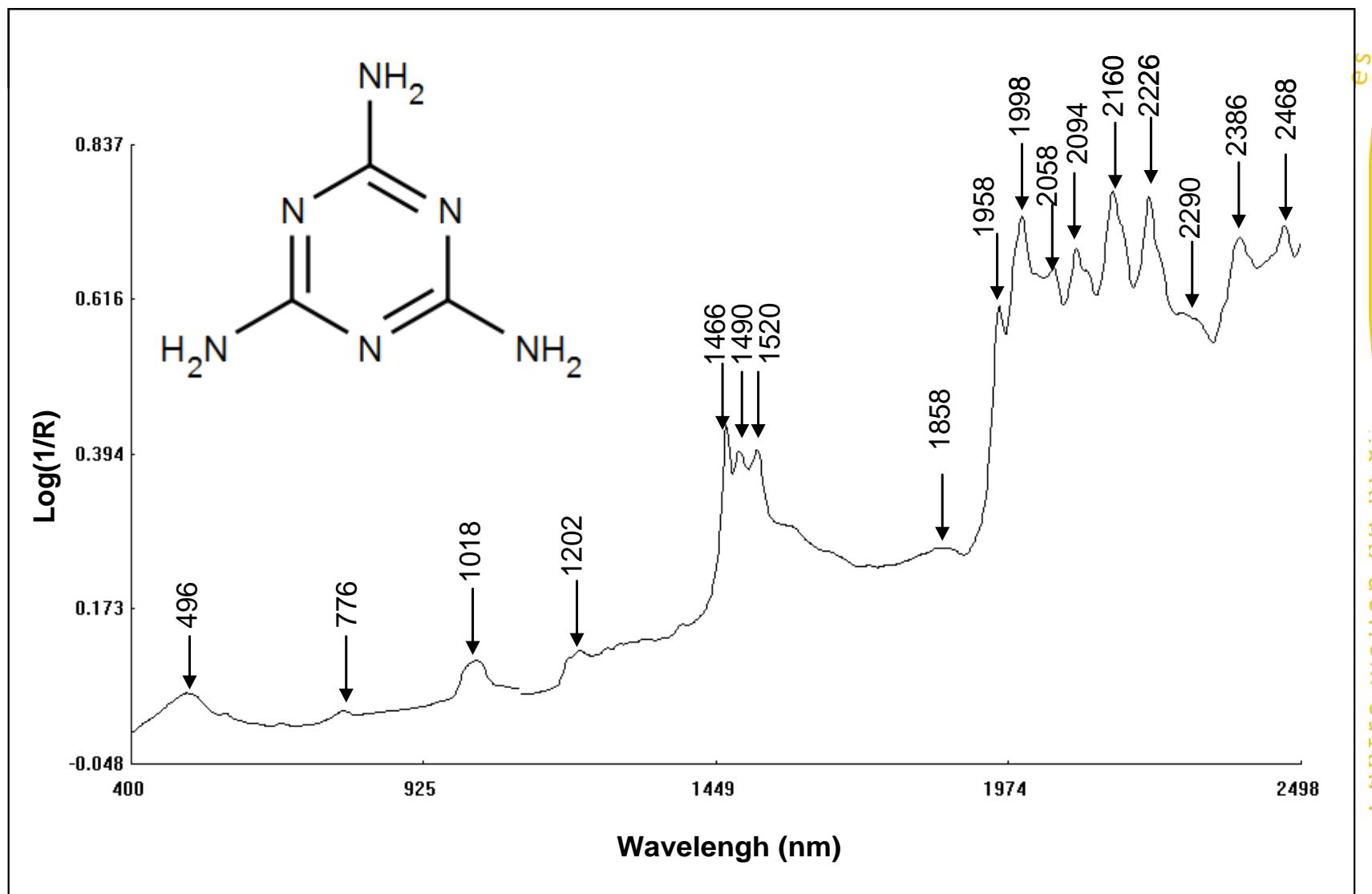
Participants

Florence Kussener, JMP (SAS)

Jean-Claude Boulet, INRA

Nadège Brun, TOTAL

Pure melamine



480 SM spectra with contamination (0.1, 0.25, 0.5, 1, 2, 3, 4, 5%)



233 SM spectra clean

CAL SET (only soya meal)

240 with 4 concentrations (0.10, 0.25, 0.50, 1.00%)

+30 randomly selected from the clean set

270 spectra

+ Noise on Y
0.15 noise
 $N(0, 0.15)$
 $M=0.41 \rightarrow 36\%$

TEST SET

T1 = 40 randomly selected from the 240 with high contamination (2,3,4,5%)

T2 = 50 randomly selected from the clean set

T3 = 43 soyameal spl from another origin + another instrument

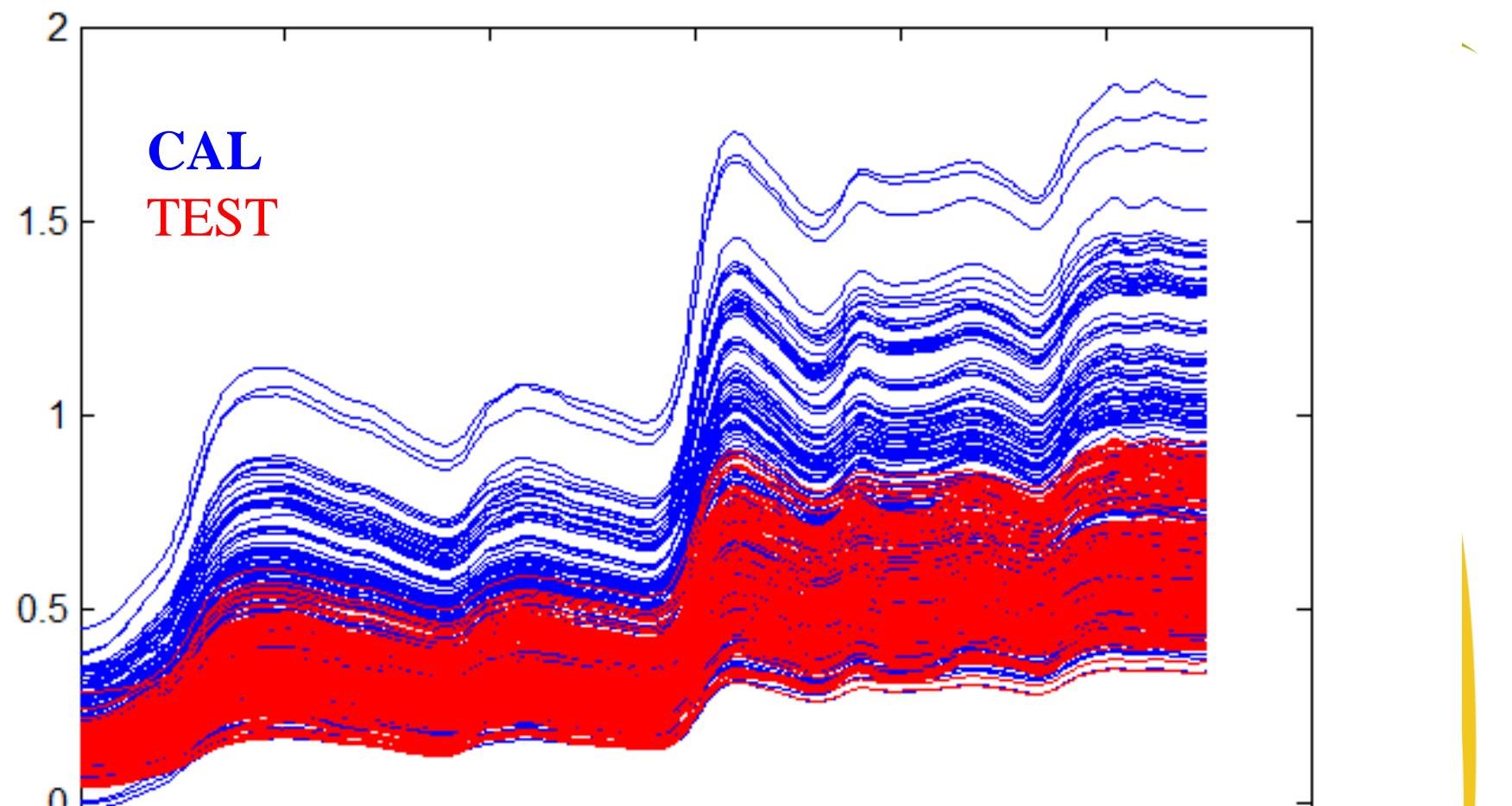
T4 = 78 wheat gluten (39 in duplicates)

T5 = 65 maize gluten

Total : 276

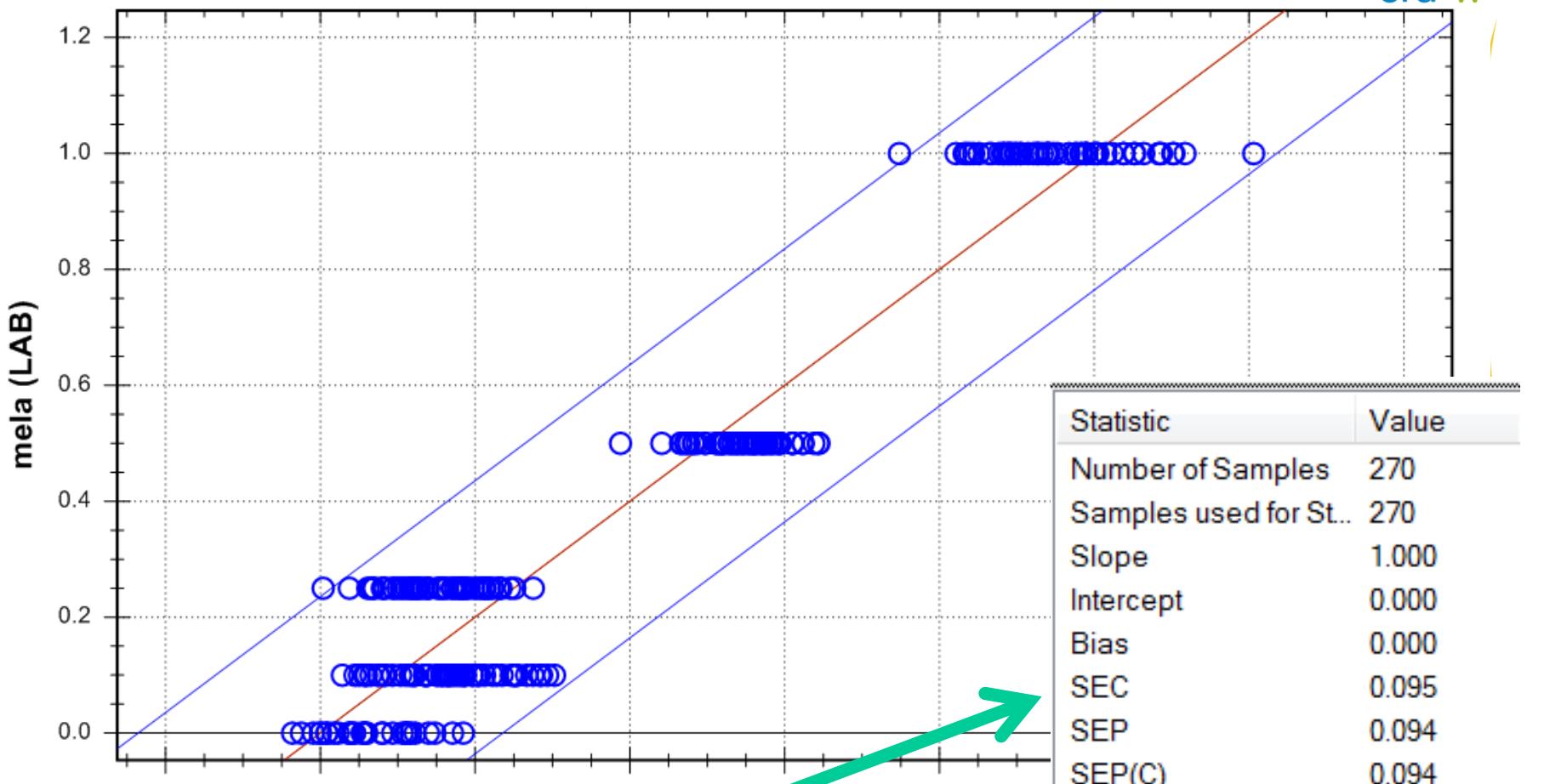
Foss XDS

Foss 6500
Clean & Conta.
Melamine
Cyanuric Acid



	PROT as is		CF as is		FAT as is	
	MEAN	SD	MEAN	SD	MEAN	SD
Soya meal	46,60	1,97	4,37	1,29	1,88	1,59
Wheat gluten	80,54	2,23	-	-	-	-
maize gluten meal	61,64	2,99	1,53	0,72	1,85	0,83

Calibration (SEC) with unmodified Y values



$0.1 = 1000 \text{ ppm}$

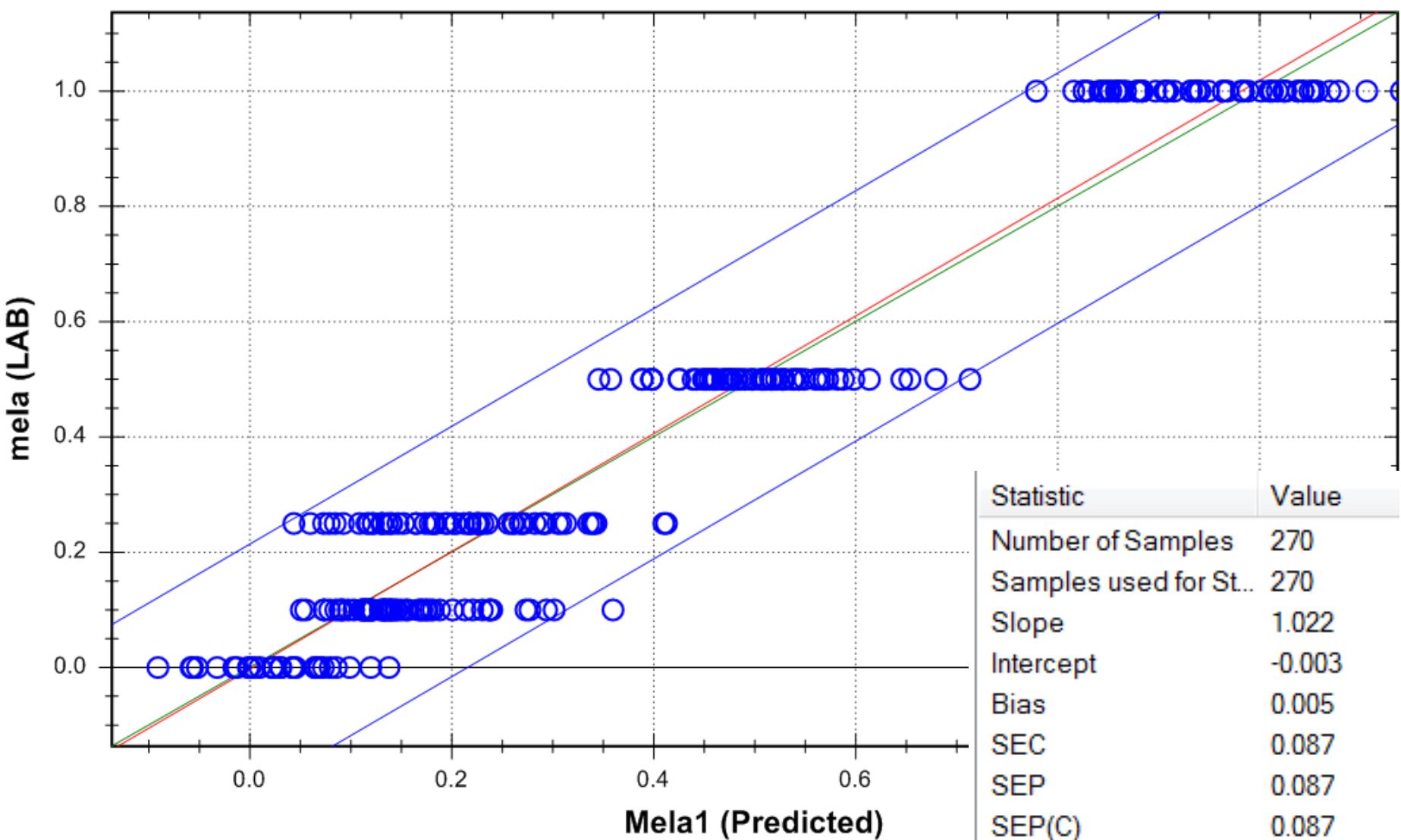
Multiple Linear Regression – Manual Step up

Segment 1 1300 - 2398, 2

Constituent	N	Mean	SD	SEC	1-VR	SECV	1-VR
mela	270	0.4111	0.3540	0.0828	0.9442	0.0835	0.9442

5dp → PLS instead of MLR

Mean	0.411	Range		0.00 - 1.00
SEC	RSQ	F	SECV	1-VR
1	0.348	0.031	9.62	0.348
2	0.203	0.670	518.94	0.204
3	0.087	0.940	1209.55	0.087



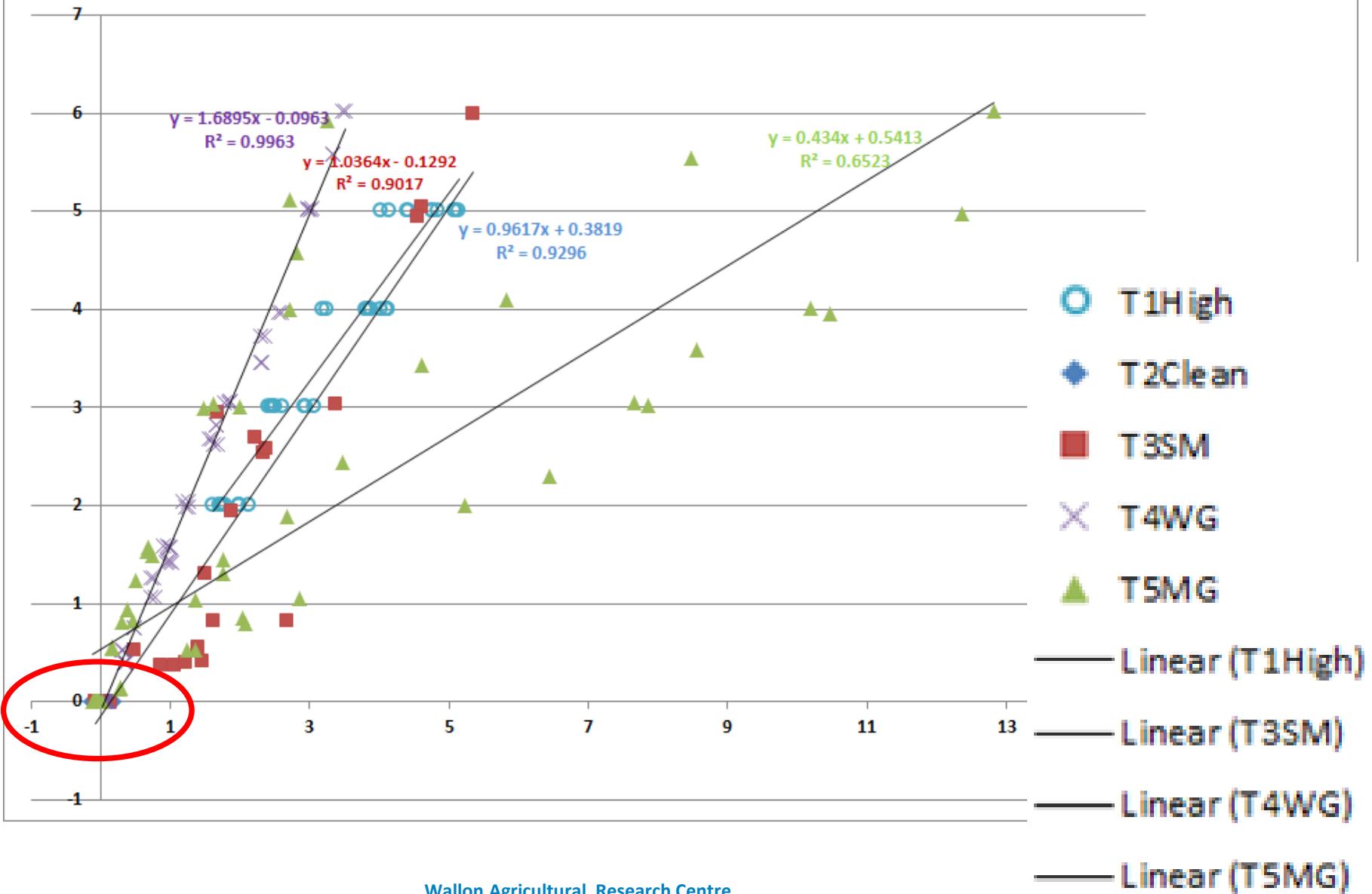
>Calibration set = 0.1 % = 1000 ppm

Ref values without noise

Raw Data 5 WL

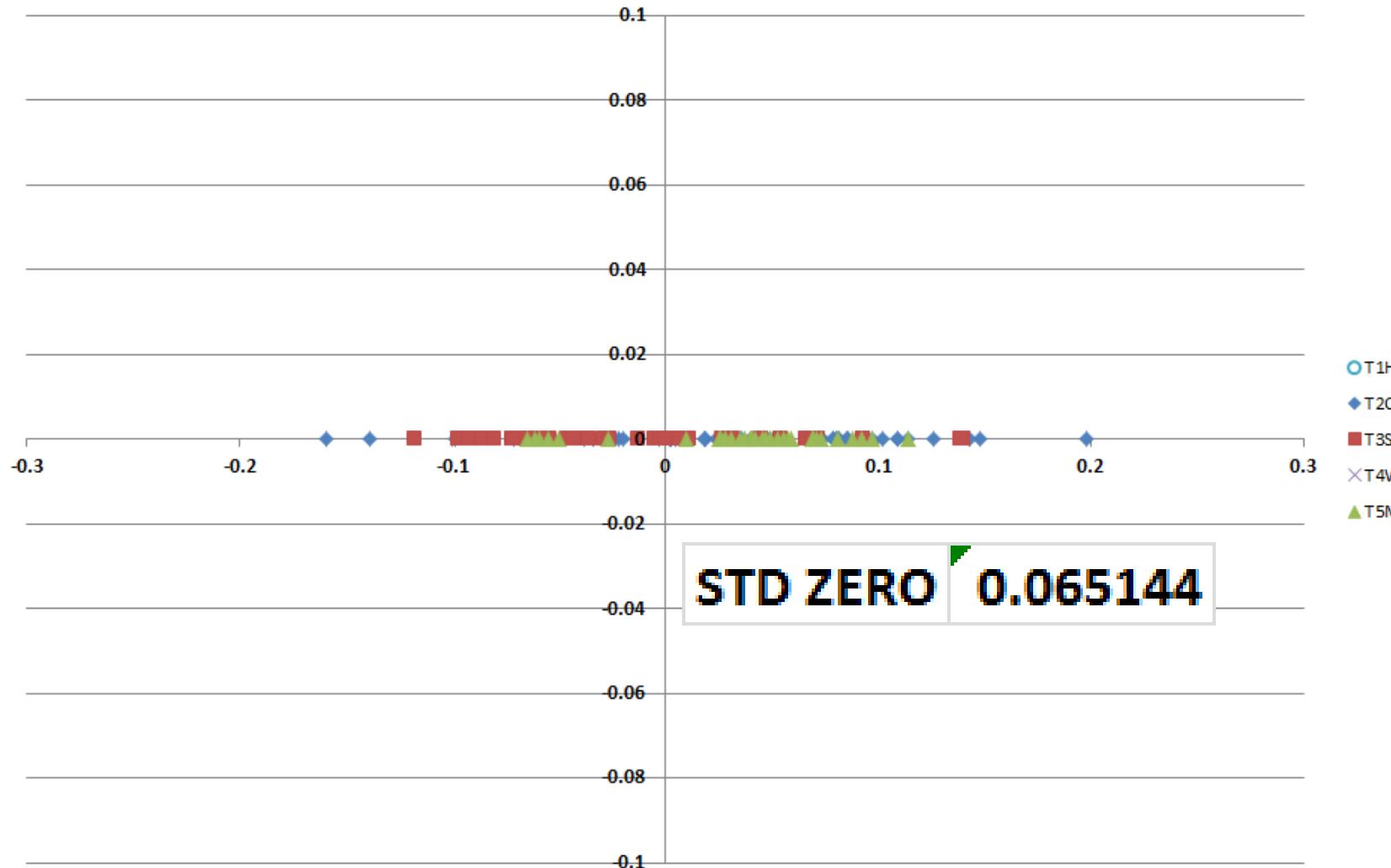
3 PLS

RMSEP=1.22



Raw Data 5 WL

3 PLS

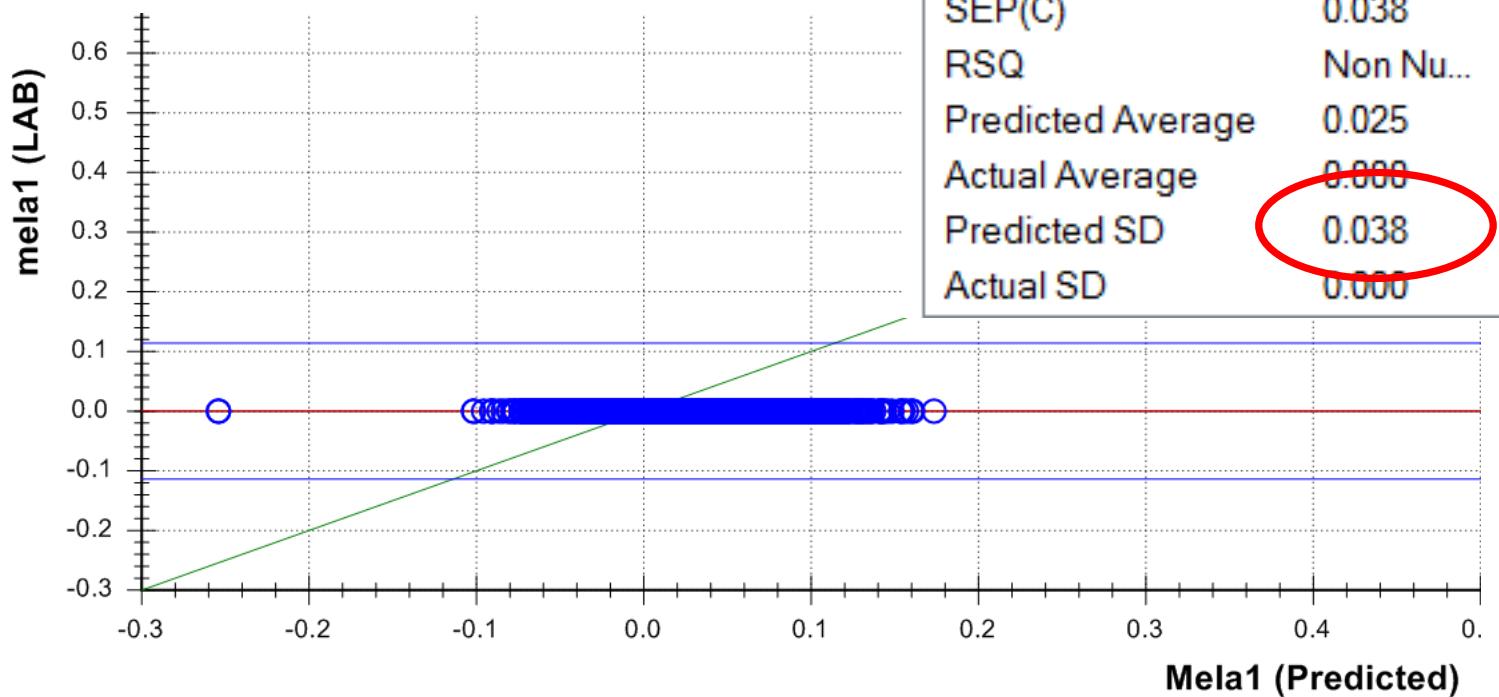


65 Mais Gluten predicted + GH based on 30 Clean # : 5 dp Log(1/R)



		>7		>4				>7		>4				>7		>4	
Posit	Sample Numbe	REF	mela	GH-mela	NH-mela	Positic	Sample Numb	REF	mela	GH-mela	NH-mela	Positi	Sample Num	REF	mela	GH-mela	NH-mela
1	T5MG_094	0.00	-0.04	2.58	0.48	26	T5MG_119	3.02	8.06	7376.12	7180.28	51	T5MG_144	1.23	0.49	30.08	17.86
2	T5MG_095	2.00	5.37	3253.86	3124.86	27	T5MG_120	0.00	-0.08	2.01	0.96	52	T5MG_145	3.03	1.64	316.29	275.44
3	T5MG_096	3.05	7.86	7043.32	6851.92	28	T5MG_121	1.49	0.73	66.34	47.81	53	T5MG_146	0.00	0.00	0.41	0.06
4	T5MG_097	3.95	10.75	13228.21	12964.88	29	T5MG_122	3.01	2.04	489.63	438.66	54	T5MG_147	1.04	1.37	205.73	174.43
5	T5MG_098	6.02	13.17	19905.54	19581.54	30	T5MG_123	3.99	2.77	900.07	830.86	55	T5MG_148	2.44	3.54	1424.05	1338.43
6	T5MG_099	0.00	-0.05	3.13	0.83	31	T5MG_124	4.58	2.87	968.33	896.57	56	T5MG_149	3.43	4.73	2544.45	2429.50
7	T5MG_100	0.00	-0.06	3.06	0.68	32	T5MG_125	0.00	-0.09	2.23	1.25	57	T5MG_150	5.54	8.70	8670.86	8457.39
8	T5MG_101	0.00	-0.11	4.15	1.09	33	T5MG_126	0.00	-0.09	2.17	1.21	58	T5MG_151	0.00	-0.04	0.89	0.02
9	T5MG_102	0.00	-0.13	4.48	1.40	34	T5MG_127	0.00	-0.09	2.21	1.19	59	T5MG_152	0.00	-0.06	1.31	0.15
10	T5MG_103	0.14	0.27	5.53	2.75	35	T5MG_128	0.00	-0.09	2.26	1.29	60	T5MG_153	0.00	-0.07	1.56	0.21
11	T5MG_104	0.85	2.08	471.47	424.09	36	T5MG_129	0.83	0.45	26.41	14.98	61	T5MG_154	0.00	-0.08	1.68	0.24
12	T5MG_105	0.80	2.13	492.93	444.22	37	T5MG_130	0.81	0.28	11.88	4.55	62	T5MG_155	1.89	2.74	845.92	780.51
13	T5MG_106	4.01	10.47	12532.79	12276.61	38	T5MG_131	0.94	0.36	18.23	8.89	63	T5MG_156	1.45	1.79	359.53	317.32
14	T5MG_107	0.00	-0.03	2.73	0.67	39	T5MG_132	2.99	1.51	270.33	232.54	64	T5MG_157	1.31	1.79	355.09	313.28
15	T5MG_108	0.52	1.39	201.07	171.20	40	T5MG_133	0.00	-0.06	1.86	0.75	65	T5MG_158	4.10	5.97	4091.09	3944.82
16	T5MG_109	1.05	2.91	932.71	865.25	41	T5MG_134	0.55	0.16	4.96	0.79						
17	T5MG_110	3.59	8.79	8808.25	8594.03	42	T5MG_135	1.53	0.66	54.61	37.84						
18	T5MG_111	4.97	12.70	18482.90	18171.29	43	T5MG_136	5.12	2.77	898.71	829.56						
19	T5MG_112	0.00	-0.06	3.21	0.73	44	T5MG_137	5.93	3.32	1289.58	1206.78						
20	T5MG_113	0.00	-0.07	3.77	0.94	45	T5MG_138	0.00	-0.08	2.06	1.15						
21	T5MG_114	0.00	-0.09	3.95	1.00	46	T5MG_139	0.00	-0.06	1.90	0.75						
22	T5MG_115	0.00	-0.10	4.52	1.29	47	T5MG_140	0.00	-0.08	2.07	1.02						
23	T5MG_116	0.13	0.27	5.26	2.27	48	T5MG_141	0.00	-0.07	2.11	1.02						
24	T5MG_117	0.52	1.27	167.21	140.38	49	T5MG_142	0.54	0.15	4.73	0.71						
25	T5MG_118	2.30	6.61	4950.30	4790.38	50	T5MG_143	1.58	0.68	58.53	41.13						

Wheat flour data set :



Statistic	Value
Number of Samples	5269
Samples used for St..	5269
Slope	0.000
Intercept	0.000
Bias	-0.025
SEC	0.000
SEP	0.046
SEP(C)	0.038
RSQ	Non Nu...
Predicted Average	0.025
Actual Average	0.000
Predicted SD	0.038
Actual SD	0.000



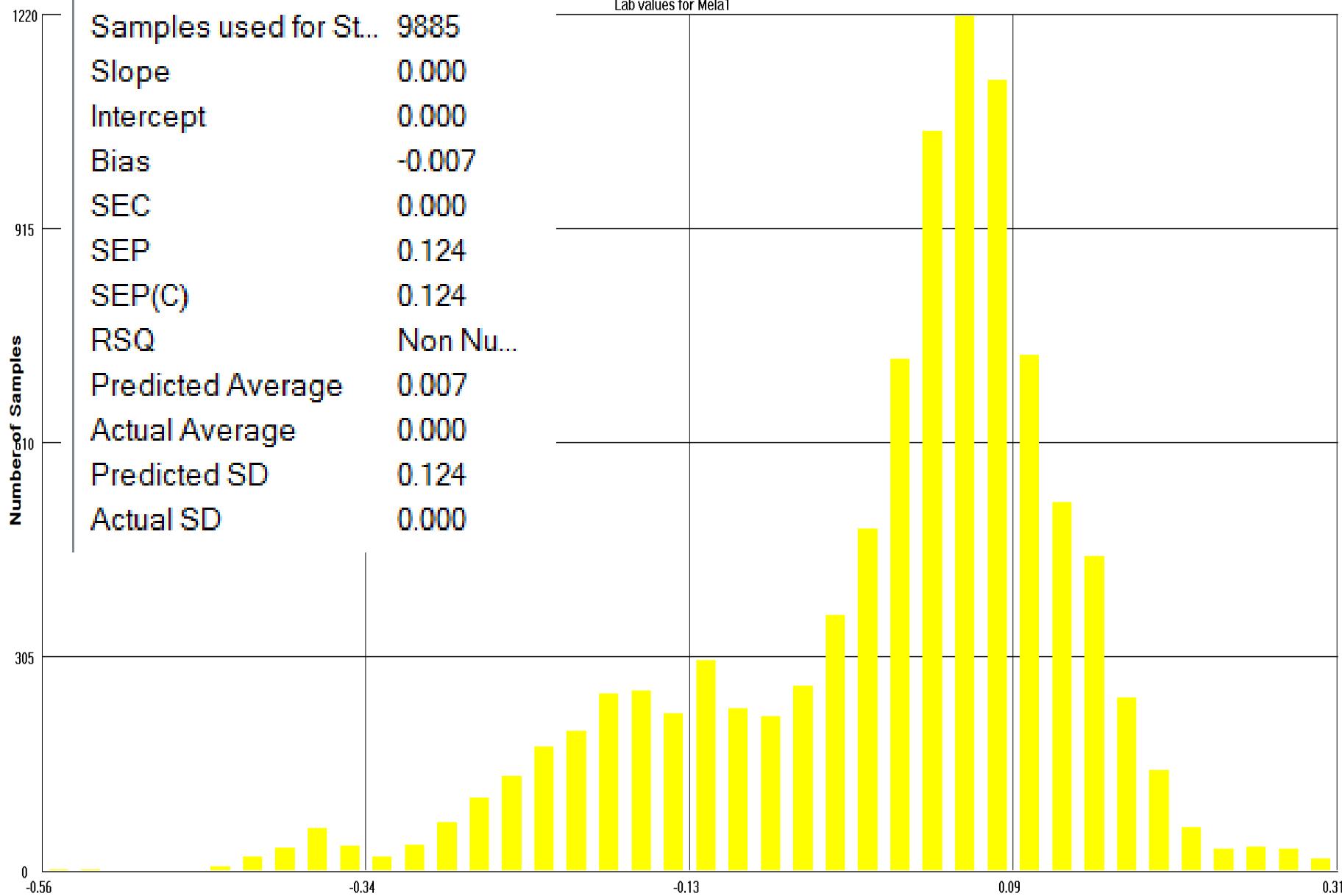
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HIGH PROTEIN FEED INGREDIENTS

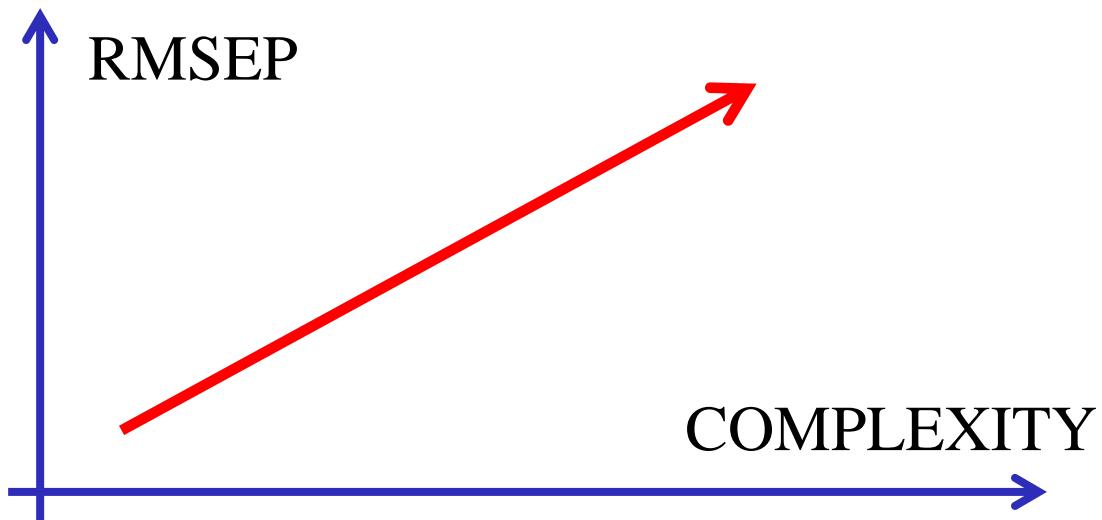


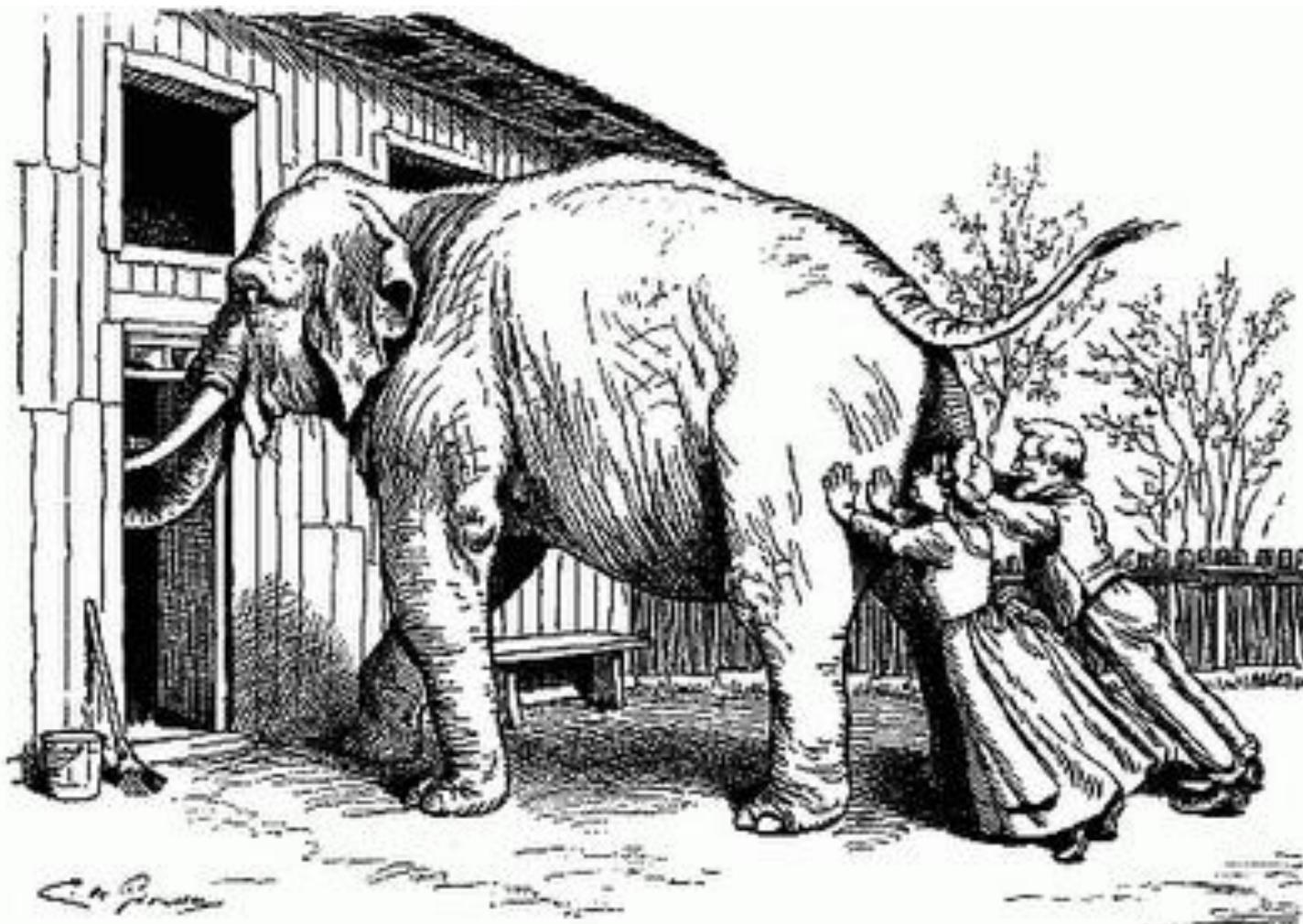
Number of Samples	9885
Samples used for St...	9885
Slope	0.000
Intercept	0.000
Bias	-0.007
SEC	0.000
SEP	0.124
SEP(C)	0.124
RSQ	Non Nu...
Predicted Average	0.007
Actual Average	0.000
Predicted SD	0.124
Actual SD	0.000

Lab values for Mela1



		RMSEP	% err.
Florence	ANN	2.36	40.2
Jean-Claude	Independant Interference Reduction risques 1er et 2ieme espèce	1.81	16.3
Nadege	Classical PLS - SG2 - Windows (IPLS)	1.41	5.4
Pierre	MLR (best 3) -> PLS 3F with 5 Log(1/R)	1.22	0.7





SOMETIMES IT IS OBVIOUS THAT SHEER
BRUTE FORCE IS NOT GOING TO WORK!

Next events



NIR platform detection

27 March 2013

We are pleased
platform worksh

Location :Gembil



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Information supplied by : Peter FLINN
11 MARCH 2013

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