

## Prä-analytische Qualitätskontrolle in Blutplasma Proben in Biobanken mittels Metabolomics Technologie

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Beate Kamlage, metanomics GmbH Oliver Schmitz, metanomics GmbH Philipp Schatz, Metanomics Health GmbH

Metanomics Health GmbH, Berlin a BASF Group Company Metanomics Health — a BASF Group Company



#### The Impact of the Pre-Analytical Phase on Sample Quality<sup>2</sup>

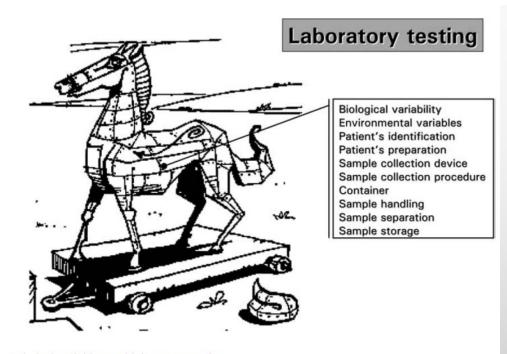


Figure 1 Preanalytical variables and laboratory testing.

Clin Chem Lab Med 2006;44(4):358-365 @ 2006 by Walter de Gruyter • Berlin • New York. DOI 10.1515/CCLM.2006.073

Review

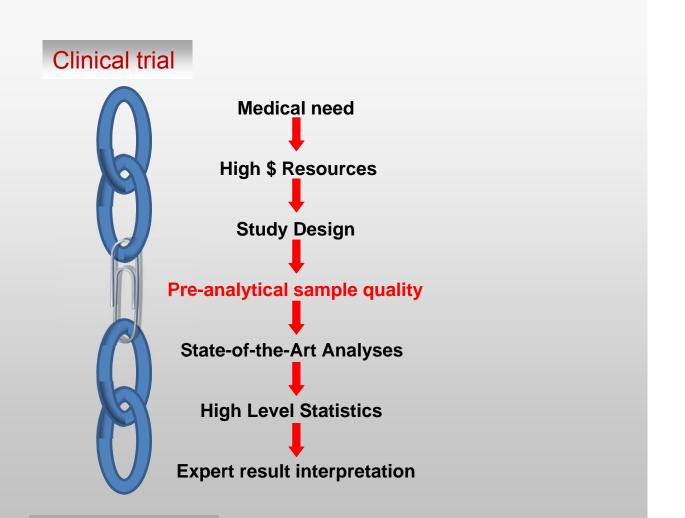
Preanalytical variability: the dark side of the moon in laboratory testing

Giuseppe Lippi<sup>1,\*</sup>, Gian Cesare Guidi<sup>1</sup>, Camilla Mattiuzzi<sup>2</sup> and Mario Plebani<sup>3</sup>

The medical error



#### -Analytics the Weakest Part of Your Research Pipeline?



### evance of Human Blood Sample Quality for Clinical narker Research

lood is easily accessible, collection is inexpensive and minimally invasive lideal for clinical assays

lood biomarkers are standard for diagnosis, prediction and monitoring of many seases (including cancer)

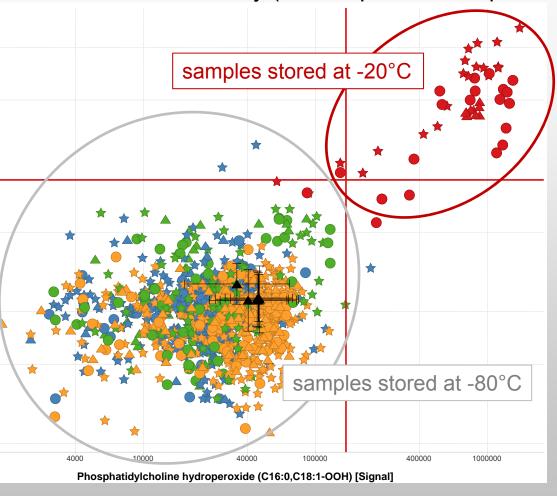
esults of targeted and -omics experimental approaches are influenced by prenalytical variations

he success of any biomarker study will depend in large part on the ality of the biospecimen analyzed"

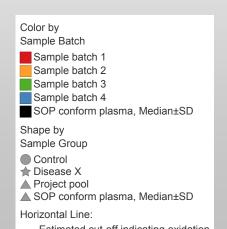
uilar-Mahecha et al. 2012, PLoS ONE 7, 1-10

#### mple of a Quality Issue: Lipid Peroxidation

narker identification study (human plasma samples from an academic partner)

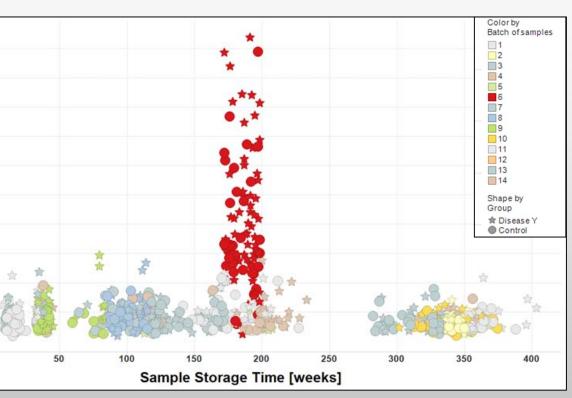


- Metabolite profiling of >1200 human plasma samples
- Batch 1 of samples presented lipohydroperoxides above the estimated thresholds
- Oxidation most likely due to inadequate storage temperature of -20°C



#### ample of a Quality Issue: Batch-to-Batch Variability

narker identification study (human plasma samples from a biobank)



- Human plasma samples stored in a biobank were thawn, split into aliquots and sent in batches to metanomics GmbH
- A quality marker is increased in a single batch (#6) of samples
- Independent of disease and storage time

ntification of pre-analytical failure in one batch of a major biomarker study

#### etanomics Health: Because Quality Matters Initiative

**Quality Assurance (QA) Service**: standardization of preanalytical processing steps, addressing key confounders for metabolomics research

→ SOP guidance, sample repository quality assessment, consulting

**Quality Control (QC) Service:** effectively assess the quality of biobank samples, e.g. pre-testing for quality markers or multivariate identification and exclusion of outliers

→ QC biomarkers, sample quality report, outlier elimination



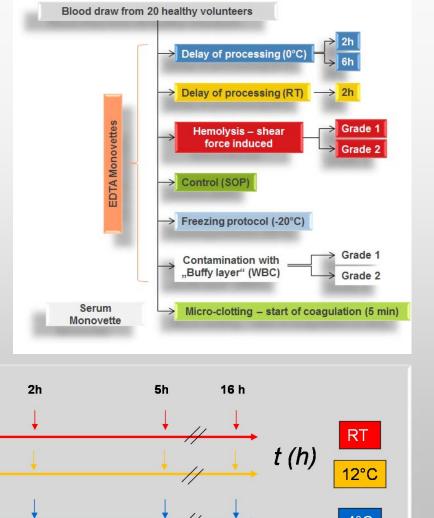


#### signs of Framework Studies – Quality Markers

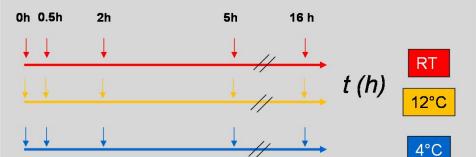
erimental designs analyzing ct of pre-analytical variables ne plasma metabolome ood processing asma processing abolomics analysis xP™ Broad Profiling xP™ Catecholamines

xP™ Eicosanoids

xP™ Lipids (sphingoid fraction)



Human plasma (EDTA, pooled)



#### sults: Percentage of Metabolome Changes

Clinical Chemistry 60:2 Other Areas of Clinical Chemistry 000-000 (2014)

#### Quality Markers Addressing Preanalytical Variations of Blood and Plasma Processing Identified by Broad and Targeted Metabolite Profiling

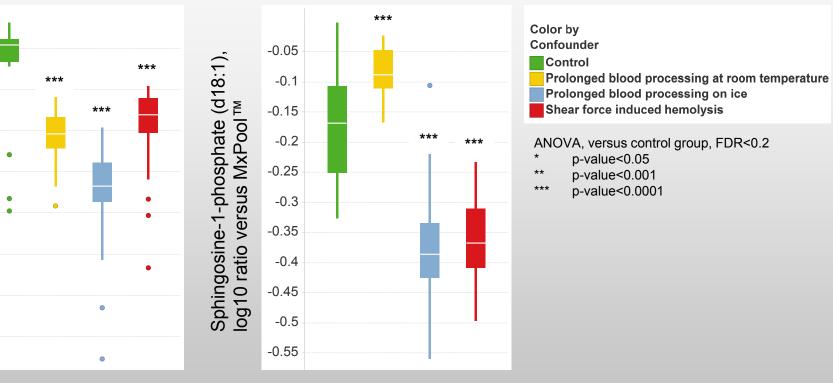
Beate Kamlage, 1\* Sandra González Maldonado, 1 Bianca Bethan, 1 Erik Peter, 1 Oliver Schmitz, 1 Volker Liebenberg, 2 and Philipp Schatz 2

Table 1. Number and percentage of statistically significant metabolite changes after applying defined preanalytical confounders out of 267 metabolites for the blood and 262 metabolites for the plasma processing experiment.<sup>a</sup>

Material	Preanalytical variable applied	Significantly changed metabolites (increase/decrease) <sup>b</sup>	
		Number	Percent change
Blood	Microclotting	31 (3/28)	12 (1/10) <sup>c</sup>
	Room temperature, 2 h	59 (27/32)	22 (10/12)
	Wet ice, 2 h	44 (12/32)	16 (4/12)
	Wet ice, 6 h	46 (17/29)	17 (6/11)
	Hemolysis, grade 1	47 (15/32)	18 (6/12)
	Hemolysis, grade 2	81 (50/31)	30 (19/12) <sup>c</sup>
	Contamination with buffy layer, grade 1	0 (0/0)	0 (0/0)
	Contamination with buffy layer, grade 2	8 (8/0)	3 (3/0)
EDTA plasma	4 °C, 0.5 h	0 (0/0)	0 (0/0)
	4 °C, 2 h	7 (7/0)	3 (3/0)
	4 °C, 5 h	16 (12/4)	6 (5/2) <sup>c</sup>
	4 °C, 16 h	30 (24/6)	11 (9/2)
	12 °C, 0.5 h	1 (1/0)	0 (0/0)
	12 °C, 2 h	7 (7/0)	3 (3/0)
	12 °C, 5 h	14 (11/3)	5 (4/1)
	12 °C, 16 h	37 (29/8)	14 (11/3)
	Room temperature, 0.5 h	4 (4/0)	2 (2/0)
	Room temperature, 2 h	28 (25/3)	11 (10/1)
	Room temperature, 5 h	47 (27/20)	18 (10/8)
	Room tomporature 10 h	(1 (41/20)	22 /1/2/0\c

#### sults: Examples of Blood Processing

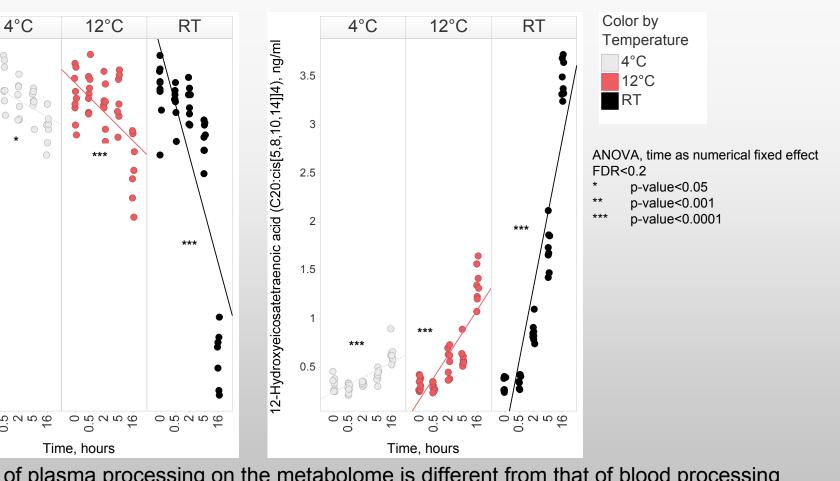
tonin and sphingosine-1-phospate levels in plasma are influenced by preytical blood processing



onin concentration in plasma is dependent on pre-analytics [Brand and Anderson, 2011, Clin Chem 57:10, 1376-86] pecial regard to platelet metabolism

gosine-1-phosphate changes are probably related to platelet metabolism (Hammad et al. 2010, Julio

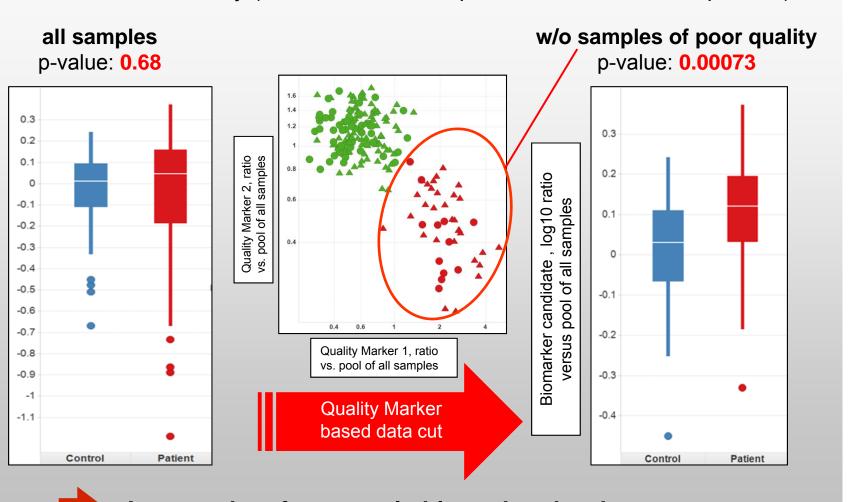
#### sults: Examples of Plasma Processing



of plasma processing on the metabolome is different from that of blood processing ical-oxidative processes and remaining enzyme activities in plasma change the plasma

#### nefit of Quality Control

rker identification study (human serum samples from an academic partner)



lity Control of Clinical Trial and Biobank Samples

### Clinical trial

Medical need



**High \$ resources** 



Pre-analytical sample quality



State-of-the-Art analyses



**Expert interpretation** 

- Biomedical research demands highquality samples (QA, QC)
- Pre-analytical processes impact plasma sample quality
- MxP™ quality control assay currently developed by Metanomics Health **GmbH** 
  - Checks sample type (EDTA -, citrate-, heparin-plasma; serum)
  - Controls for blood processing
  - Controls for plasma processing, storage and shipping
- Clinical use cases
  - Monitoring clinical sample quality
  - Selection of high quality samples from biobanks

#### Sincere thanks to ....



- Our academic and clinical partners
- Our pharma and nutrition partners
- Our dedicated and enthusiastic staff at Metanomics Health and metanomics



### E-BASF The Owning Conyany

#### Headquarters:

Metanomics Health GmbH Tegeler Weg 33 10589 Berlin Germany

Phone: +49 30 3480 7400

#### **US Offices:**

BASF Corporation 100 Park Avenue Florham Park, NJ 07932, USA Dr. Hajo Schiewe, Senior Manager North America

Email: hajo.schiewe@basf.com

Phone: +1 612 605 7971

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The Chemical Company