



Fortschritte für die Standardisierung der Präanalytik



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Möglichkeiten der Ergebnisverwertung aus Forschungsprojekten

Publikation

Journal of proteome research

Delayed Times to Tissue Fixation Result in Unpredictable Global Phosphoproteome Changes

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Supporting Information

ABSTRACT: Protein phosphorylation controls the acuity of signal transduction pathways regulated by kinases and phosphatases. Little is known, however, about the impact of protein phosphorylation changes on the levels of phosphopeptides in biological tissues at time points after tissue fixation. The aim of this study was to characterize the potential effects of delayed tissue preservation (cold ischemia) on the levels of phosphopeptides using targeted and unspecific approaches. We analyzed rat liver tissue under different cold ischemia conditions with cold ischemic conditions ranging from 10 to 360 min prior to cryopreservation. The phosphoproteome was analyzed using several proteome array (RPPA) technology and phosphopeptide enrichment followed by LC-MS/MS analysis. In addition, we analyzed the analysis of rat liver tissues with long (up to 360 min) cold ischemia times did not reveal statistically significant alterations of specific phosphorylation events brought nonphosphorylated cytoskeletal proteins to a higher level of phosphorylation. However, freezing the samples in ice prior to cryopreservation prevented this effect. LC-MS/MS-based quantification of 1684 phosphorylation sites in rat liver tissues showed changes of their distribution compared to time point zero. After reaching statistical significance for individual phosphopeptides. Similarly, RPPA analysis of rat liver tissue with short (60 min) cold ischemia times did not reveal direct or predictable changes of protein and phosphopeptide levels. Using LC-MS/MS and quantification of 791 phosphorylation sites, we found that the distribution of ratios compared to time point zero broadens with prolonged ischemia times, but there were rather undirected and diffuse changes of protein phosphorylation. In conclusion, our results show that both RPPA and LC-MS/MS analysis of rat and mouse liver tissues, we conclude that prolonged cold ischemia results in unspecific phosphoproteome changes that cannot be predicted nor assigned to individual proteins. On the other hand, we identified a number of proteins that are significantly affected by cold ischemia and, therefore, may be used as general reference markers for future companion diagnostics for kidney substitutes.

KEYWORDS: tissue, proteomic, proteomics, Novartis, Novartis, RPPA, phosphoproteins, mass spectrometry

INTRODUCTION

Phosphorylation and dephosphorylation are key mechanisms of intra- and intercellular signal transduction and reflect the activation status of a cell. The alteration of gene phosphorylation patterns are often used to predict biological processes involved against deregulated signaling pathways in cancer patients. However, knowledge of the impact of preanalytical variations such as delayed time to tissue fixation or freezing, on global phosphoproteome changes is limited.

Modulations of cellular signaling cascades and accompanying changes of phosphoprotein levels in tissue samples have to be investigated in much more detail, and the most critical pre-analytical parameters affecting protein profiling have to be identified. Otherwise the information gained by proteomic studies is a complete distortion of what is really happening. Consequently, researchers are realizing that preanalytical variations influence sample quality and integrity and thus may affect phosphoprotein levels. Recently, it was shown in an elegant and highly recognized study that tissues after surgical resection are still viable.

Received: May 13, 2013
Published: August 29, 2013

DOI: 10.1021/acs.jproteome Res. 2013.12.4024

Published online in Wiley Online Library (wileyonlinelibrary.com) on May 13, 2013; DOI: 10.1021/acs.jproteome Res. 2013.12.4024

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4024

Published online in Wiley Online Library (wileyonlinelibrary.com) on May 13, 2013; DOI: 10.1021/acs.jproteome Res. 2013.12.4024

Printed by Jove, ISSN 1541-429X

Patent

EP 1 356 302 B1

EUROPEAN PATENT SPECIFICATION

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Bild 1

(11) EP 1 356 302 B1

(12) Date of publication and mention of the grant of the patent: 29.01.2008 Bulletin 2008/04

(21) Application number: 01991160.9

(22) Date of filing: 07.11.2001

(37) International publication number: WO 2002059006 (18.07.2002 Gazette 2002/25)

(54) METHOD AND DEVICE FOR COLLECTING AND STABILIZING A BIOLOGICAL SAMPLE
VERFAHREN UND VORRICHTUNG ZUM SAMMELN UND ZUM STABILISIEREN EINER BIOLÓGISCHEN PROBE
MÉTHODE ET DISPOSITIF DE PRÉLEVEMENT ET DE STABILISATION D'UN ÉCHANTILLON BIOLÓGIQUE

(56) Designated Contracting States: AT BE CY CH DE DK ES FI FR GB GR IE IT LU MC NL PT SE TR

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Postfach 10 22 41
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(58) Reference cited: WO-A-9418155 DE-A-59 031 236
US-A- 6 015 055

(59) Priority: 01.11.2000 US 2002/04
30.11.2001 US 99/4058

(60) Date of publication of application: 29.11.2001 Bulletin 2008/04

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(75) Description:
The explosive human immunodeficiency virus type I (HIV-1) antigen, in essence a C virus of restricted genovariety, "JOURNAL OF VIROLOGY", vol. 68, no. 3, pp. 1146-1157, XN002372729 (USN: 6052-584K)
The HIV-1 antigen, "JOURNAL OF VIROLOGY", vol. 68, no. 3, pp. 1146-1157, XN002372729 (USN: 6052-584K)
"Direct Detection of Herpesvirus (HSV) DNA in Human Peripheral Blood and Comparison With HSV RNA in Plasma and Peripheral Blood Monocyte Cells," JOURNAL OF MEDICAL VIROLOGY, vol. 47, no. 2, 1995, pages 153-160, L1900X0003 01040005

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Printed by Jove, ISSN 1541-429X

Standard

CEN/TC 140
Date: 2014-06
TC 140/WI 00140098
CEN/TC 140
Secretariat: DIN

Molecular in-vitro diagnostic examinations — Specifications for pre-examination processes for frozen tissue — Extracted proteins

Molekuläranalytische in-vitro diagnostische Verfahren — Spezifikationen für präanalytische Prozesse für gefrorene Gewebepräparate — Extrahierte Proteine

Élément introductif — Élément central — Élément complémentaire

ICS:
Description:

Document type: Technical Specification
Document subtype: Document
Document stage: Publication
Document language: E

C:\Users\kth\Desktop\protein_cryo_2014-07-04\kth.doc STD Version 2.5a

Normen und Spezifikationen als Beitrag zur Innovationsförderung

Hightech-Strategie 2020 für Deutschland

- „**Normung und Standardisierung werden in Deutschland zunehmend integraler Bestandteil des Forschungs- und Innovationsprozesses**, denn frühzeitig eingeleitet fördern sie den Transfer von Forschungsergebnissen in marktfähige Produkte und Dienstleistungen und den schnellen Marktzugang von Innovationen“
- „**Eine aktive Beteiligung an Normungs- und Standardisierungsaktivitäten** verschafft der deutschen Wirtschaft zudem globale Wettbewerbsvorteile.“



Bundesministerium
für Bildung
und Forschung



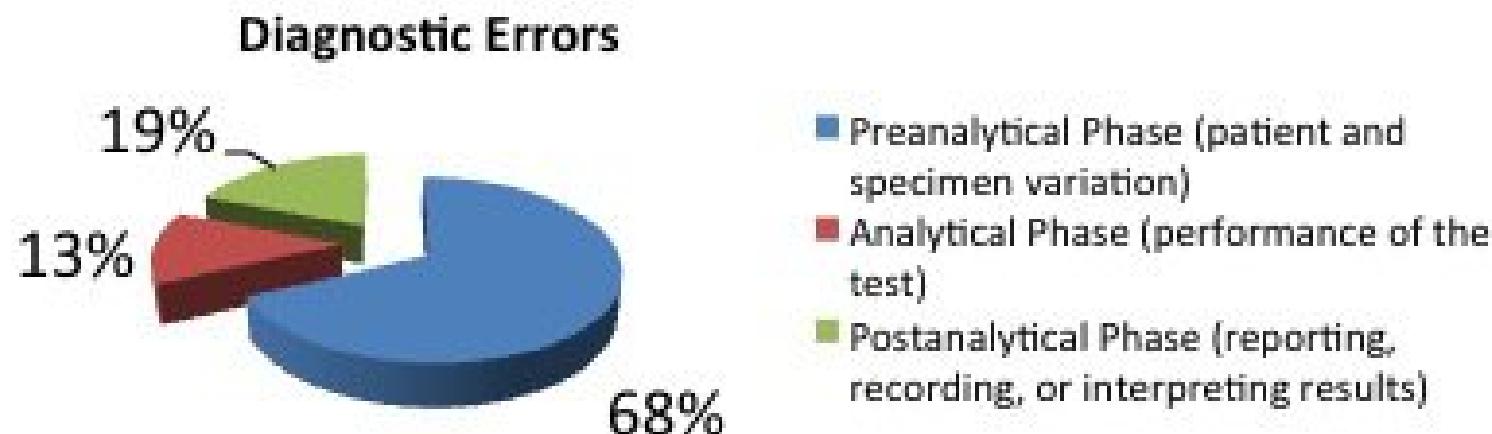
Ideen. Innovation. Wachstum

Hightech-Strategie 2020 für Deutschland

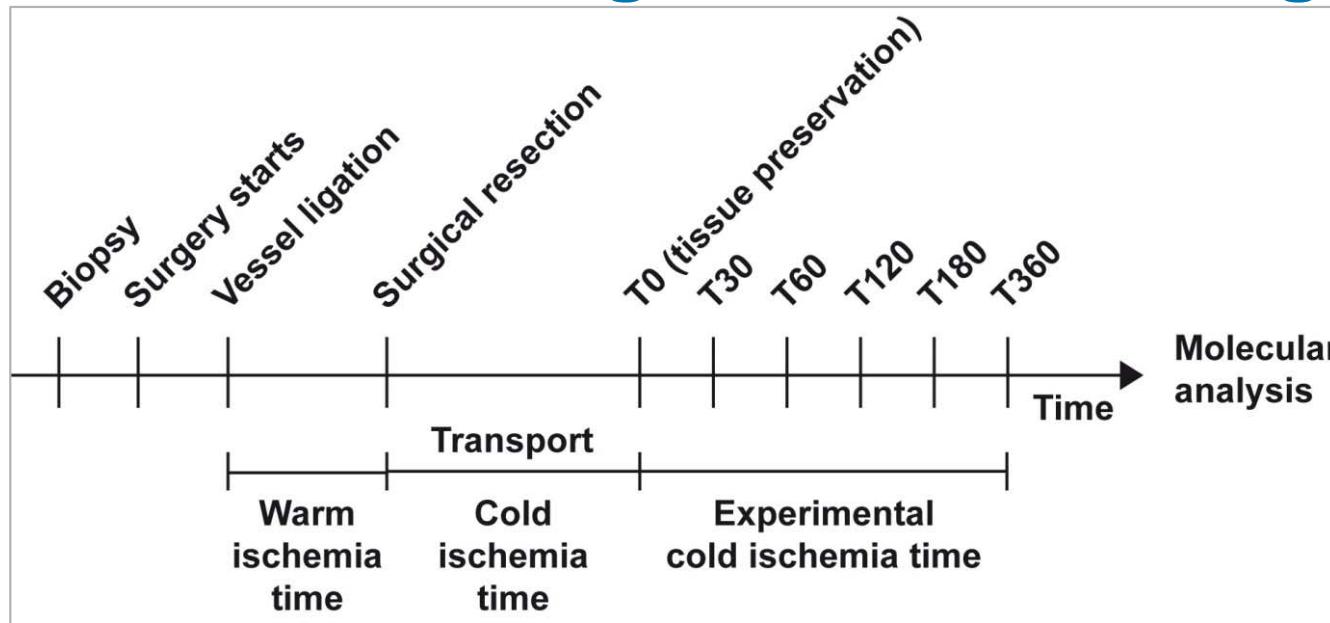


Quelle: Hightech-Strategie 2020,
BMBF 2010 (Hrsg.), S. 10

Entwicklung von Standards für die Bioproben sammlung und *in vitro* Diagnostik



Entwicklung von Standards für die Bioprobensammlung und *in vitro* Diagnostik



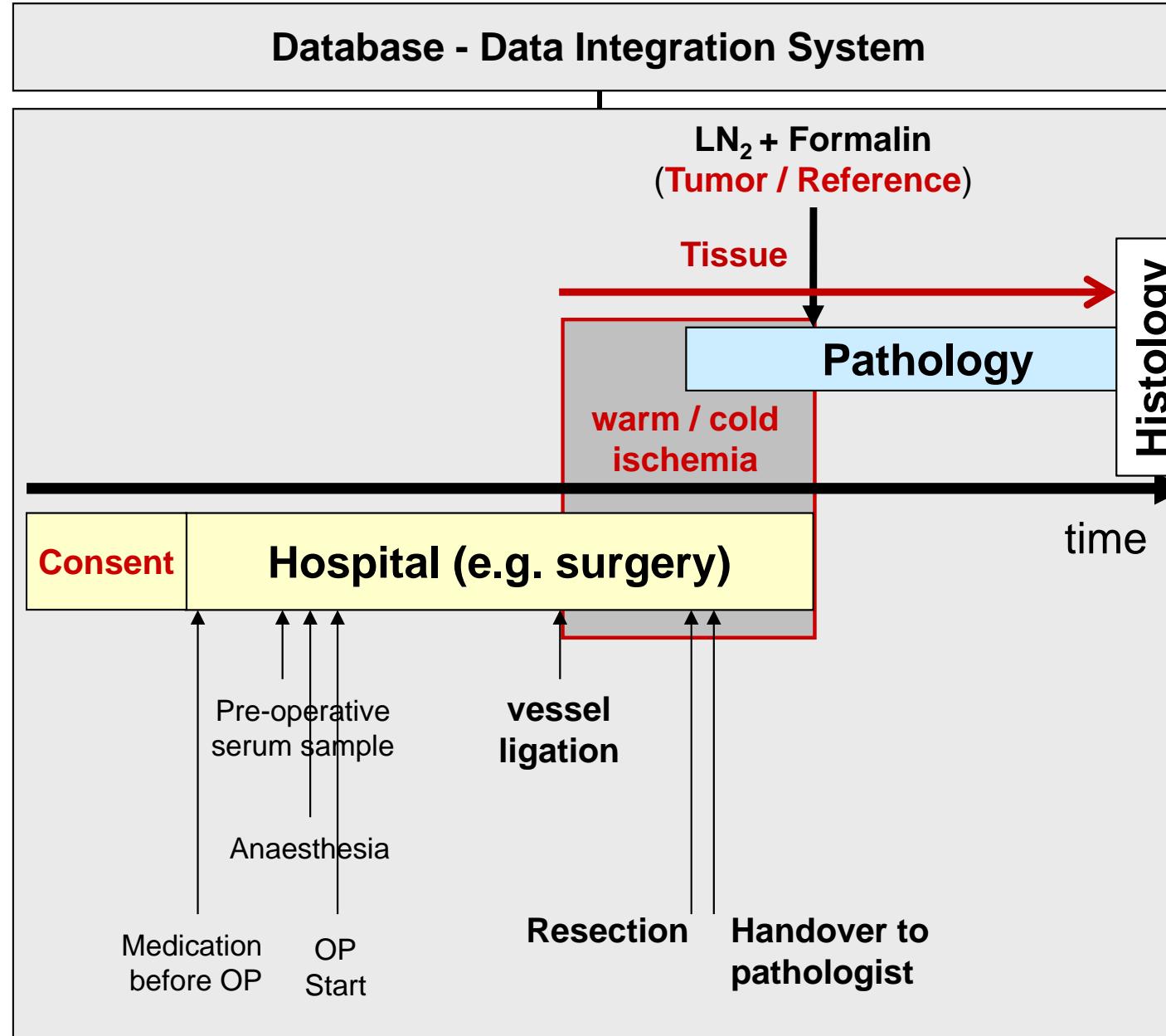
RNA analysis

Protein analysis

Metabolome analysis

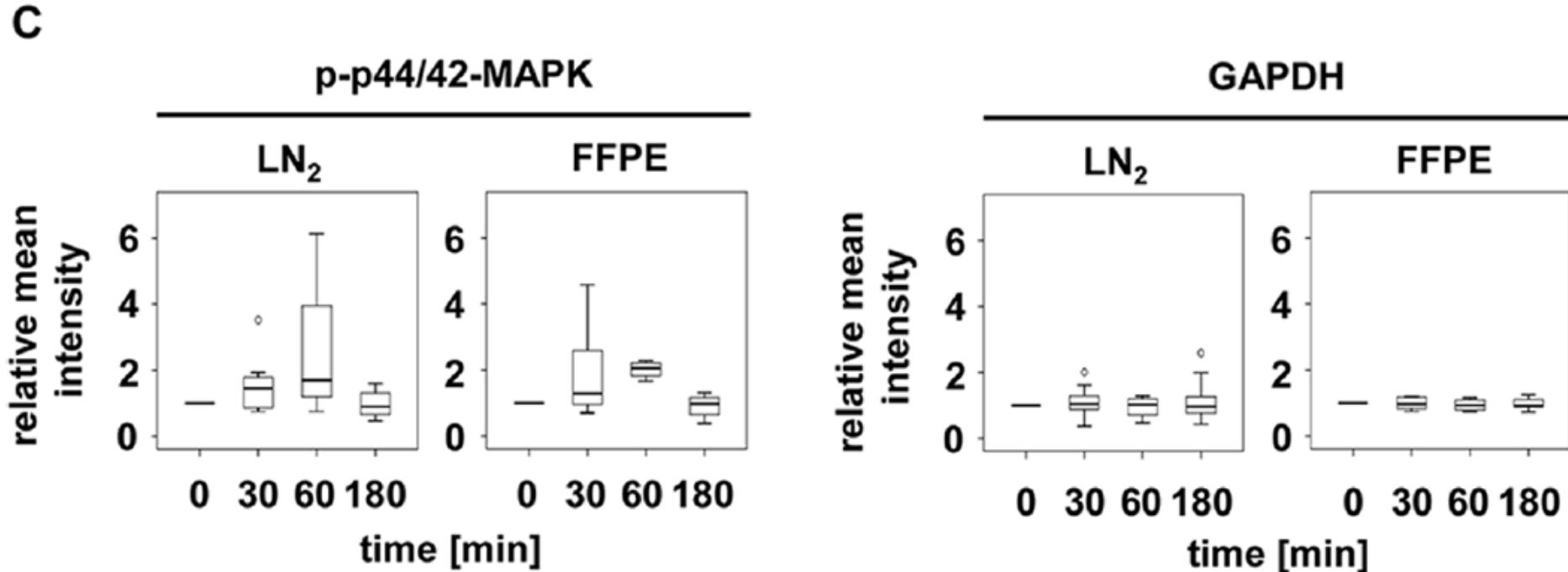
European Committee for Standardization (CEN)

Collection of tumor tissue samples exposed to different ischemic conditions



1411
samples
from 128
patients

Impact of cold ischemia duration on protein levels



Gündisch et al. 2012, J Proteome Res
Gündisch et al. 2013, J Proteome Res

Example of a European Standard for the preanalytical phase



TECHNICAL SPECIFICATION
SPÉCIFICATION TECHNIQUE
TECHNISCHE SPEZIFIKATION

FINAL DRAFT
FprCEN/TS 16827-1

March 2015

ICS 11.100.10

English Version

Molecular in vitro diagnostic examinations - Specifications for pre-examination processes for FFPE tissue - Part 1: Isolated RNA

Tests de diagnostic moléculaire in vitro - Spécifications relatives aux processus préanalytiques pour les tissus FFPE - Partie 1: ARN extrait

Molekularanalytische in-vitro-diagnostische Verfahren - Spezifikationen für präanalytische Prozesse für FFPE-Gewebe - Teil 1: Isolierte RNAs

This draft Technical Specification is submitted to CEN members for formal vote. It has been drawn up by the Technical Committee CEN/TC 140.

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Recipients of this draft are invited to submit, with their comments, notification of any relevant patent rights of which they are aware and to provide supporting documentation.

Warning: This document is not a Technical Specification. It is distributed for review and comments. It is subject to change without notice and shall not be referred to as a Technical Specification.

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COMITÉ EUROPÉEN DE NORMALISATION
EUROPAISCHES KOMITEE FÜR NORMUNG

CEN-CENELEC Management Centre: Avenue Marnix 17, B-1000 Brussels

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Ref. No. FprCEN/TS 16827-1:2015 E

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6/9 Technischen Spezifikationen sind Ende 2015 von CEN publiziert worden*



Molecular in-vitro diagnostic examinations — Specifications for pre-examination processes for **blood**

- Part 1: cellular RNA
- Part 2: genomic DNA
- Part 3: cell free circulating DNA

Molecular in-vitro diagnostic examinations — Specifications for pre-examination processes for **FFPE tissue**

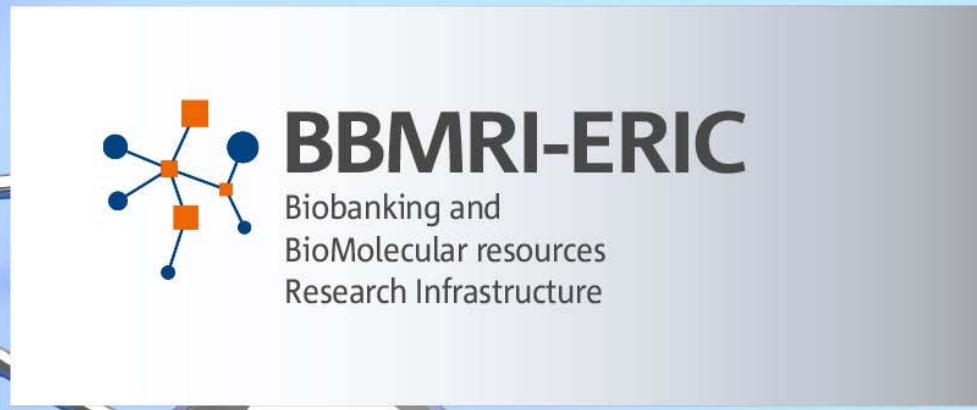
- Part 1: RNA
- Part 2: Proteins
- Part 3: DNA

Molecular in-vitro diagnostic examinations — Specifications for pre-examination processes for **snap frozen tissue**

- Part 1: RNA
- Part 2: Proteins

Molecular in-vitro diagnostic examinations — Specifications for pre-examination processes for **metabolomics in urine, serum and plasma**

Relevant für *in vitro* Diagnostiklabore und deren Kunden, Hersteller von *in vitro* Diagnostika, Molekulare Pathologie, Biobanken, Akkreditierungsorganisationen, akademische und industrielle medizinische Forschung, ...



slides provided to Prof. Karl-Friedrich Becker

Andrea Wutte

26 November 2016 (for presentation purpose – call for technical experts BBMRI-ERIC)

BBMRI-ERIC Work Programme 2016 Quality

Work Stream 2.1 CEN/TC 140 / ISO 212 **Quality of the sample**



- Set up Expert-WG for Evaluation of Pre-examination processes
- **Evaluation of the pre-examination processes of the 9 published CEN/TC 140 Standards by the BBMRI-ERIC experts of the field**
- Definition of Evaluation Tasks (Criteria, Documentation, Outcome and Timeline)
- Nomination of Experts for specific Evaluation task
- Evaluation process
- WG meetings (TC, Webinars, meetings) to discuss and evaluate developments
- Documentation for Self-assessment (**deliver main criteria for Self-Assessment-Tool and Documentation for community use**)

Der Weg zu einem ISO-Standard



- 2017: Publication of ISO Standards
- 2014: 8 new projects for ISO Standards approved in ISO/TC 212 „Clinical laboratory testing and *in vitro* diagnostic test systems“



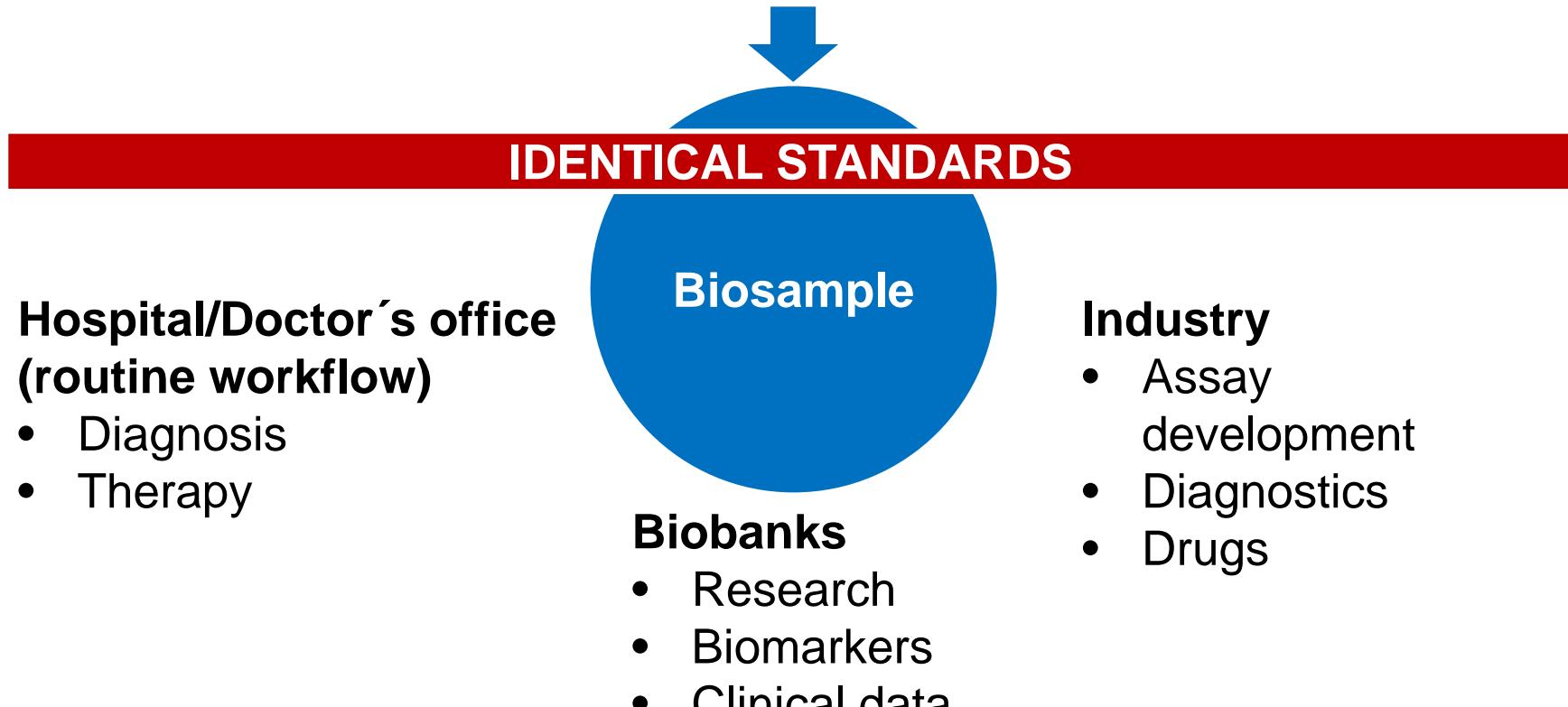
CLIA



- 2015: 9 CEN Technical Specifications are being published
- 2013: 9 new projects approved in CEN/TC 140 „In vitro diagnostic medical devices“
- 2010: Start of standardization work

Aim: One Standard for the same workflow

Sample donor/Patient



- ISO/TC 140 (In vitro Diagnostics)
- ISO/TC 276 (Biotechnology -> Biobanks)
- FDA, CLIA, EDMA, CANCER ID, ESP,

...

Zusammenfassung

- Verwertung von Forschungsergebnissen als Standard
- Prä-analytische Phase zur Probensammlung muss verbessert werden
- 6/9 CEN Standards zur Prä-Analytik wurden 2015 publiziert
- ISO Standards zur Prä-Analytik werden derzeit erstellt
- Publikation der ISO Standards geplant für 2017
- Relevant für
 - *in vitro* Diagnostik Labore und Kunden dieser Labore
 - Hersteller von Diagnostika
 - Biobanken
 - Industrielle und akademische medizinische Forschung
 - Akkreditierungsorganisationen, Zulassungsbehörden
 - ...
- International gleiche Standards für die Bioprobengewinnung im Krankenhaus und Arztpraxen, für Biobanken und Industrie

Thanks

Pathology TUM/HMGU

- Christa Schott
- Christian Beese
- Sibylle Gündisch
- Julia Slotta-Huspenina
- Heinz Höfler



Klinikum rechts der Isar, Munich

- Surgery
 - Klaus-Peter Janssen
- Institut für Medizinische Statistik und Epidemiologie
 - Klaus Kuhn
 - Rainer Blaser
 - Andreas Lehmann

Qiagen GmbH

- Uwe Oelmueller
- Daniel Grölz

DIN German Institute for Standardization

- Lena Krieger
- Margit Heinrich

Medical University of Graz (MUG)

- Kurt Zatloukal
- Christian Viertler

Erasmus Medical Center Rotterdam (EMC)

- P. Riegman
- M. Kap

Medical Sciences Department - University of Trieste

- Giorgio Stanta

