

Biobanking

Lecture: Master Molecular Medicine

SS 2023

Jena, 5th June 2023

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Please contact me if you have any questions regarding this lecture:

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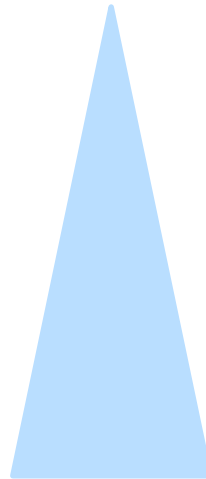
Biobanking? – More than just a Freezer!



Biobanking is more than just storing samples in a freezer and getting them out for a measurement!

Storage Complexity

1 sample
100 samples
10000 samples
100000 samples



Complexity of storage

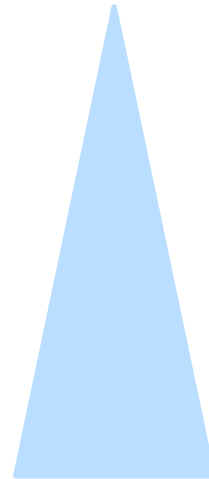
- Storage facilities
- Sample organization
- Data management

Biobanks offer professional sample storage and organization



Data Handling Complexity

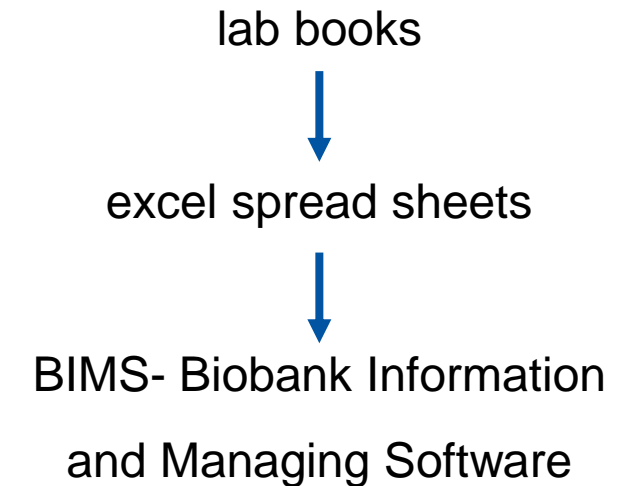
1 sample
100 samples
10000 samples
100000 samples



Complexity of storage

- Storage facilities
- Sample organization
- Data management

Professional data handling



Why Biobanking?



Guideline for quality- promoting aspects in medicine and biomedicine

Leitfaden für qualitätsfördernde Aspekte in der Medizin und Biomedizin

Erarbeitet von der Arbeitsgruppe „Qualität in der Klinischen
Forschung“ der Ständigen Senatskommission für Grundsatz-
fragen in der Klinischen Forschung der DFG

Published June 2021

https://www.dfg.de/foerderung/info_wissenschaft/2021/info_wissenschaft_21_54/index.html

Key Tasks for Biological Samples from Biobanks

Biomaterials from Biobanks should „reflect the biological or biochemical state of the donors at the time of sampling“.

- Even after long term storage

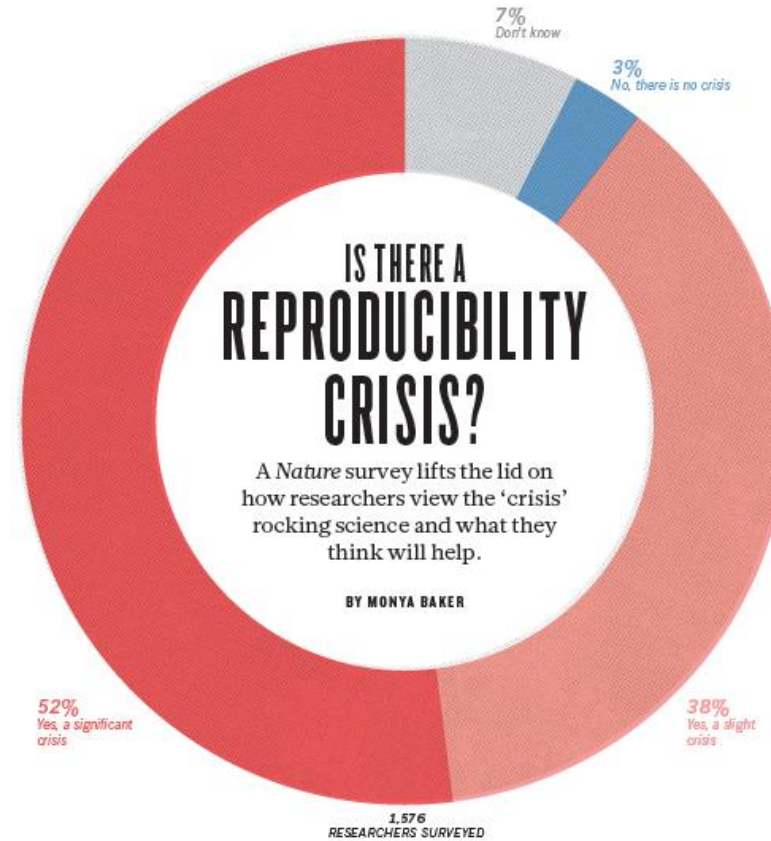
Sample integrity plays a major role for downstream analysis!

But:

There are multiple process- associated risks for biological samples endangering sample integrity!

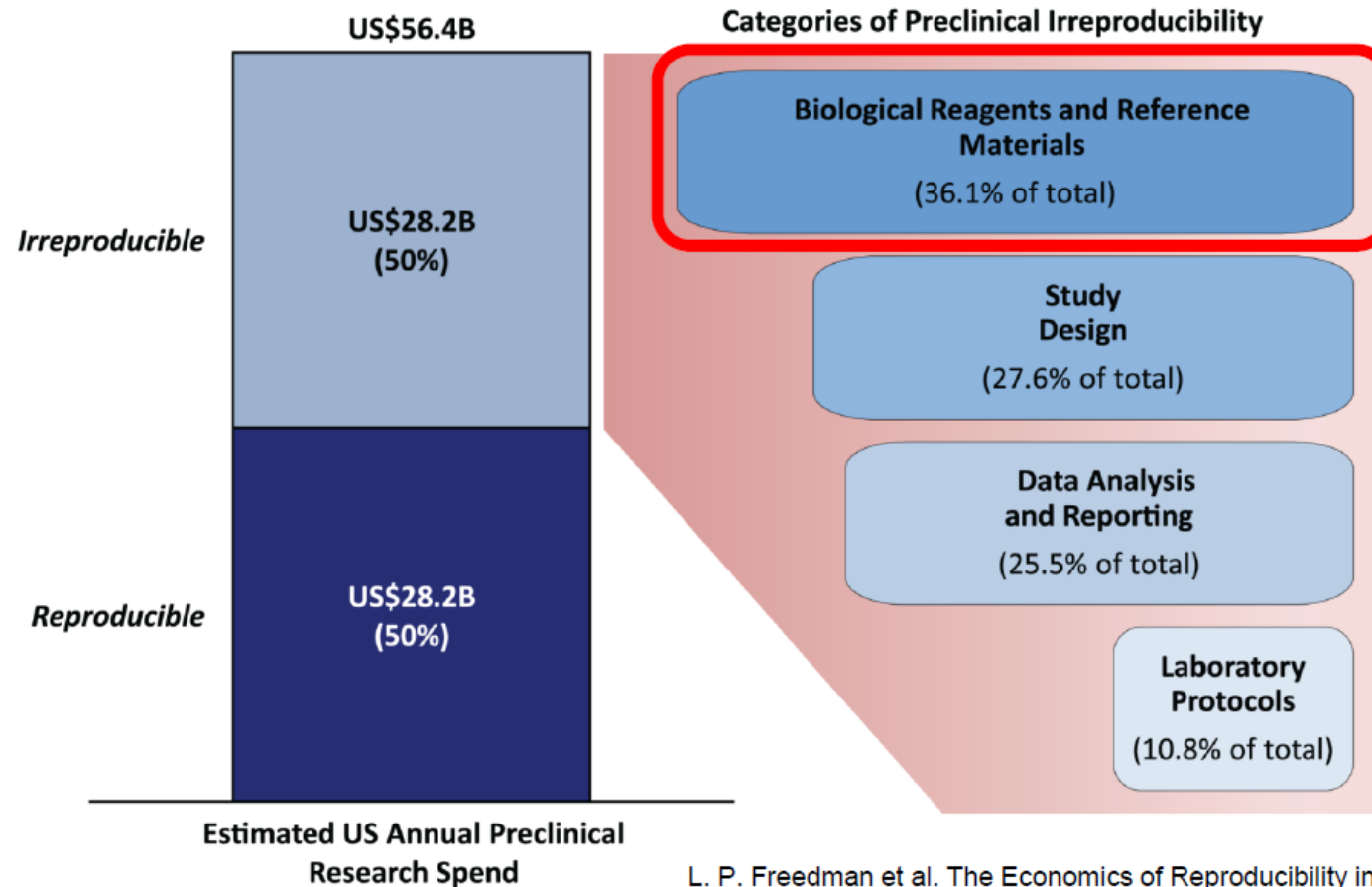
Why Biobanking?

**Biobanks help storing samples
at optimal conditions**



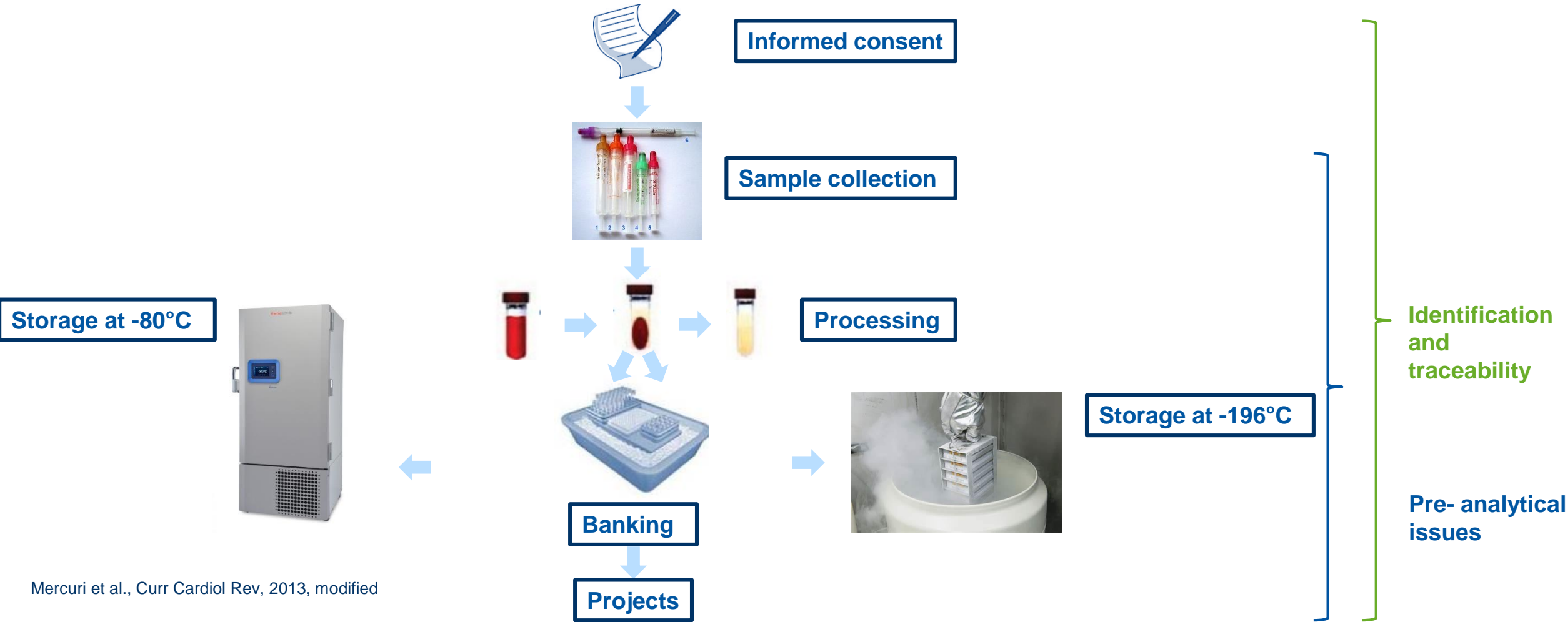
Nature (Vol.533, Issue 7604., May 2016)

Estimated US Preclinical Research Spend and Categories of Errors that Contribute to Irreproducibility



L. P. Freedman et al. The Economics of Reproducibility in Preclinical Research, *PLoS Biol* 13, e1002165–9 (2015)

Process- associated Risks for Biological Samples- Overview



Mercuri et al., Curr Cardiol Rev, 2013, modified

Pre- analytical Issues

Preenalytical phase

Biological/ environmental factors



Time of day/ Season



Coffee



Alcohol



Weight



Smoking

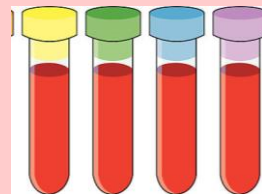


Medicine



Sunlight

Technical factors



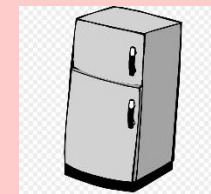
Collection



Processing



Transport



Storage

High risk for sample integrity

Analytical phase



Postanalytical phase



Ellervik & Vaught, Clin Chem, 2015, modified

Pre-analytical Problem- Processing

An MS-based metabolomics profiling on human EDTA plasma samples revealed that a high number of metabolites was significantly increased or decreased by preanalytical variation during blood and plasma processing

Short-term storage of blood either at room temperature or cooled on wet ice, hemolysis, and short-term storage of plasma for 5 or 16 h at room temperature resulted in statistically significant changes (4% to 19% of metabolites increased and 8% to 12% of metabolites decreased) of the human EDTA plasma metabolites.

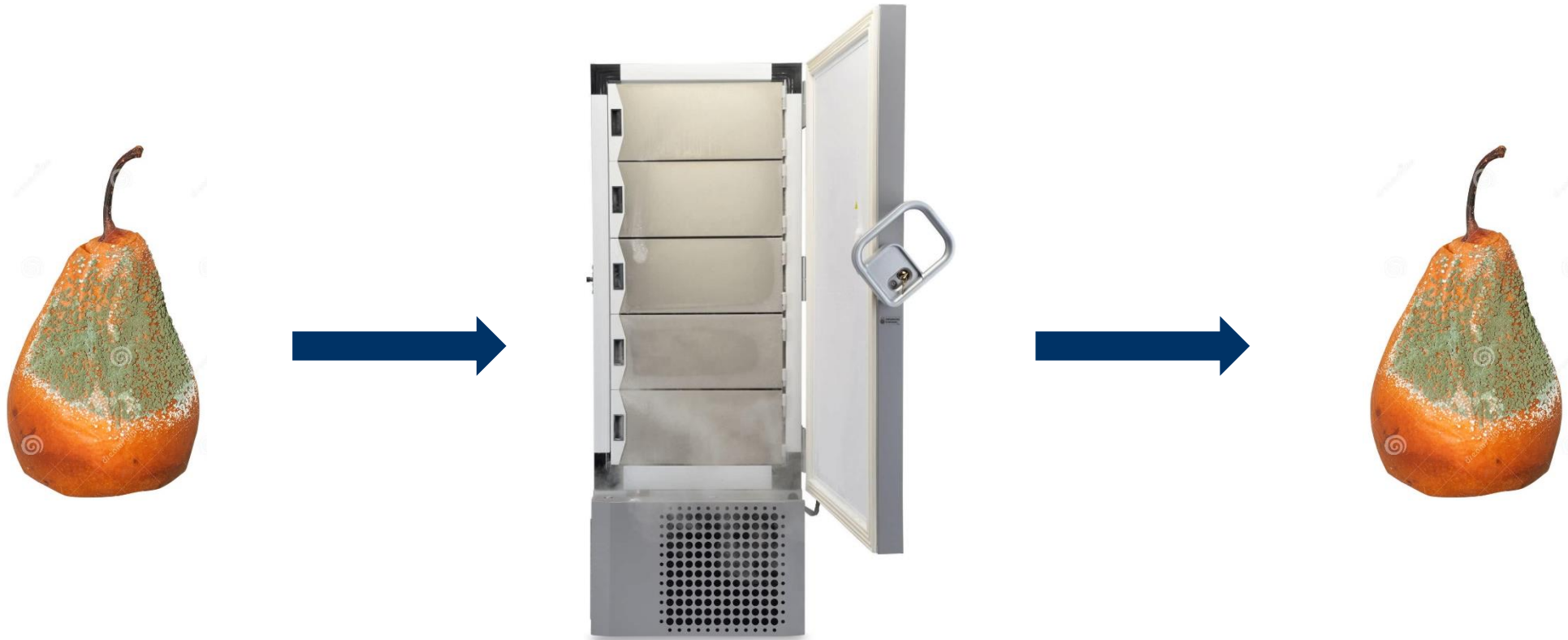
Table 1. Number and percentage of statistically significant metabolite changes after applying defined preanalytical confounders out of 267 metabolites for the blood and 262 metabolites for the plasma processing experiment.^a

Material	Preanalytical variable applied	Significantly changed metabolites (Increase/decrease) ^b	
		Number	Percent change
Blood	Microclotting	31 (3/28)	12 (1/10) ^c
	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) ^c
	Contamination with buffy layer, grade 1	0 (0/0)	0 (0/0)
EDTA plasma	Contamination with buffy layer, grade 2	8 (8/0)	3 (3/0)
	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) ^c
	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)	16 (10/6)
	Room temperature, 16 h	61 (41/20)	23 (16/8) ^c

^a Statistical analysis was done via ANOVA, the significance level was set to $P < 0.05$ and $FDR < 0.2$. Details are given in online Supplemental Tables 1 and 2.
^b Including free hemoglobin.
^c All 3 percentages are rounded to an integer, so the values in brackets do not necessarily add up to the total.

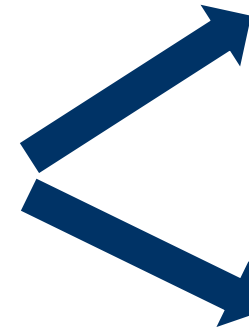
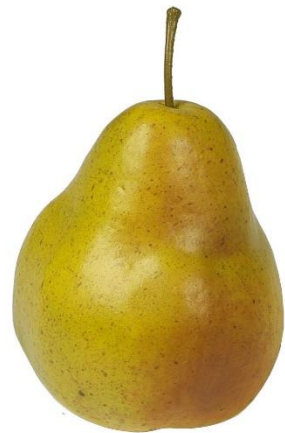
Kamlage et al., Clin Chem, 2014

Pre-analytical Problem

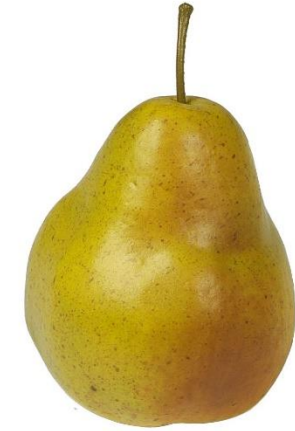


Pre-analytical Problem- Storage

Problem: Storage is a kind of a black box for sample quality



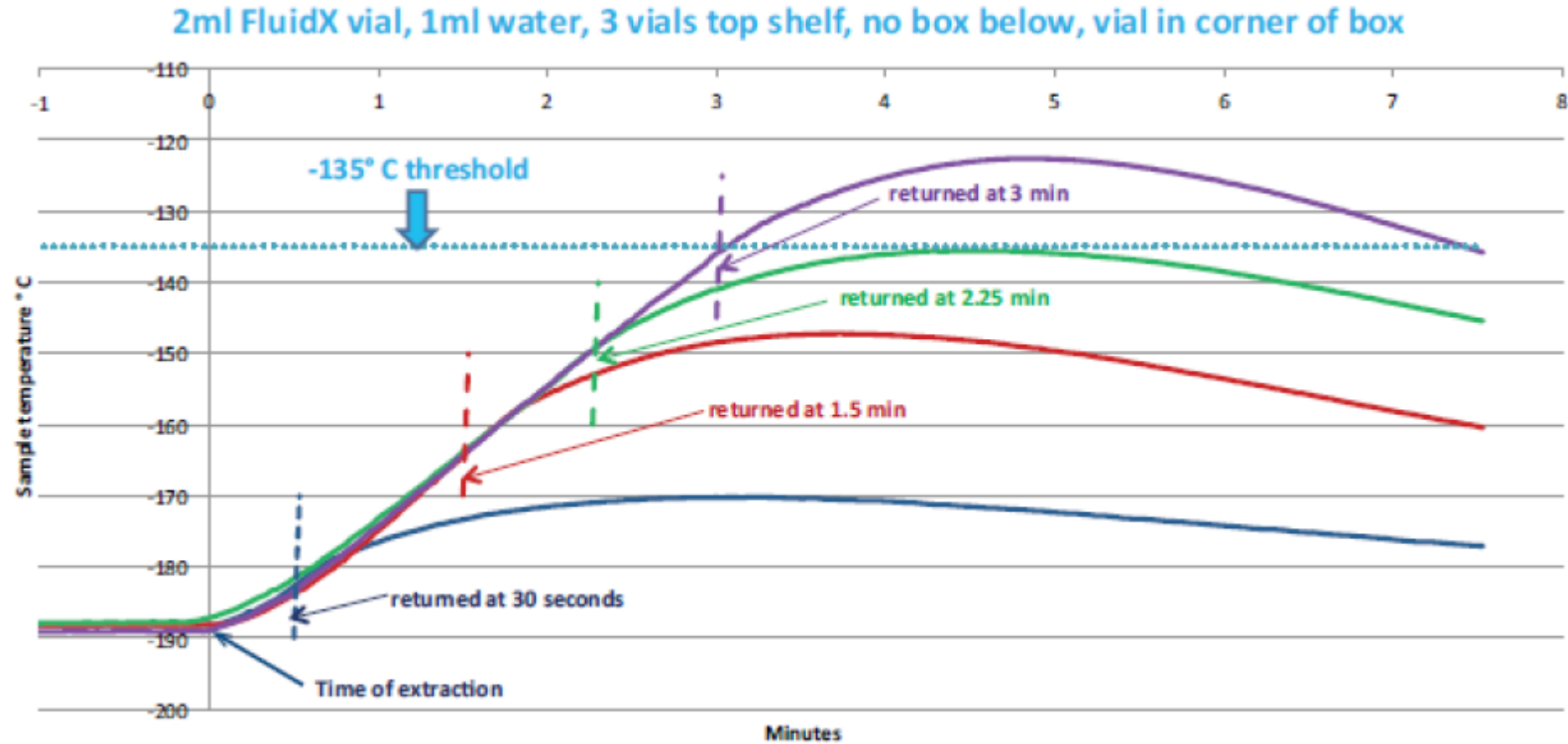
best case



worst case



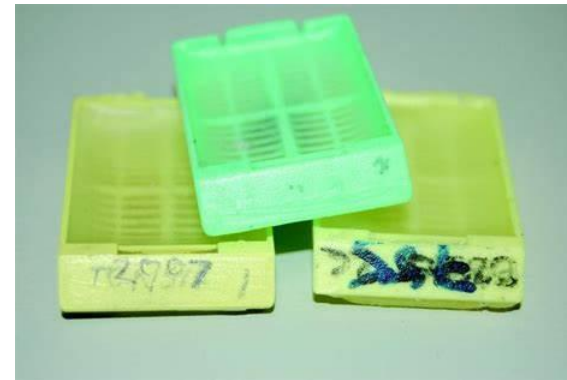
Pre-analytical Problem-Storage



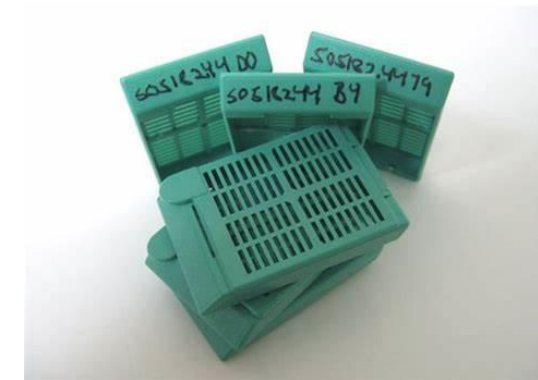
Warhurst et al., Conference Paper *in Cytotherapy*, 2015

Identification and Traceability

Identification of samples- Problems arising from today's archives



Poorly labeled samples will lead to confusion and possibly misidentification.



Ambiguous cassette labels „S05“, „50S“, „505“, „SOS“?

Identification of Samples

- Identification of samples is one of the key points in biobanking!
- It allows sample tracking and connection to sample and donor specific data.
- If a sample loses its identifier it is nearly impossible to save it.
- This has to be prevented as lots of work and money is needed to biobank samples!
- Also data protection plays an important role!
- Don't use donor identifying codes (names, address)

Identification of Samples- Examples from the IBBJ



Barcode and human readable label (optional)



Data matrix code

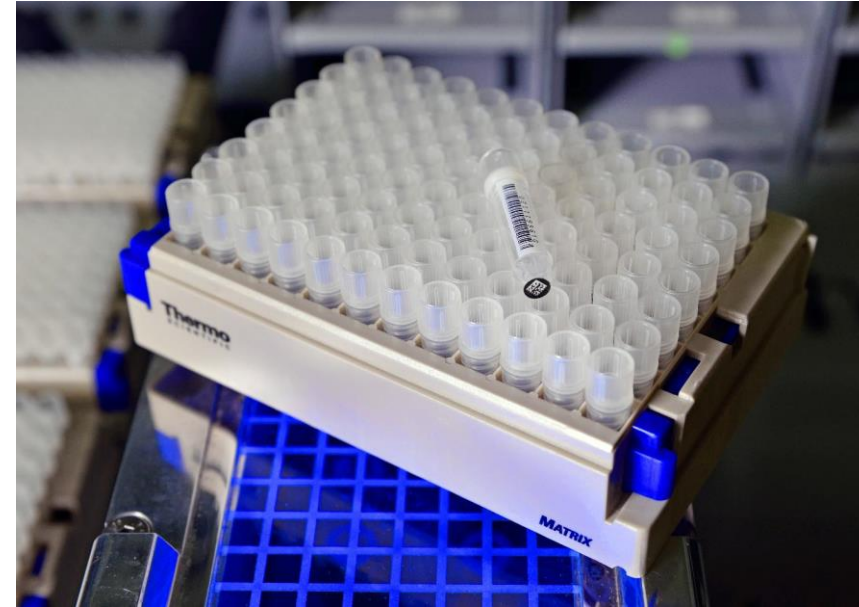
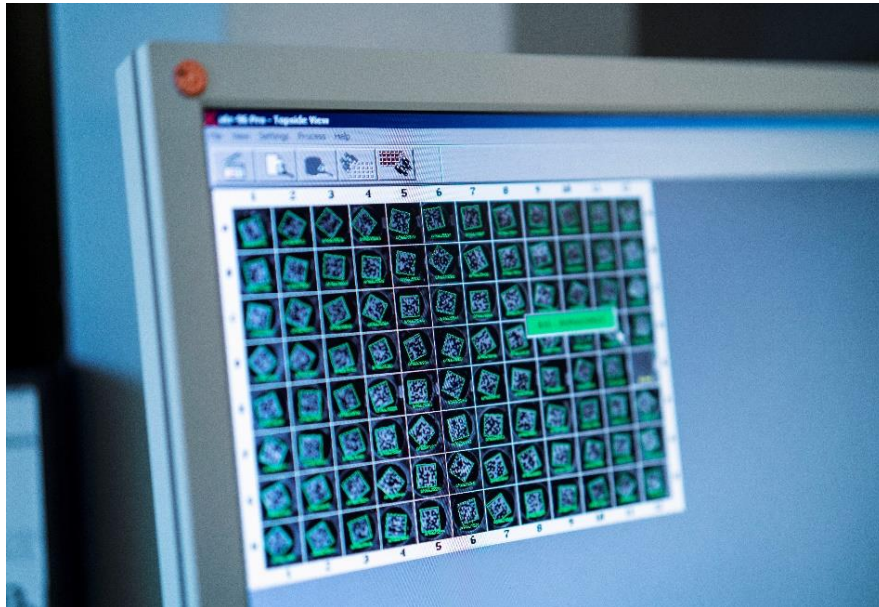


Barcoded rack showing data matrix orientation code on underside



Identification of Samples- Examples from the IBBJ

Key to efficient sample management is automatisation:
All tubes (samples) are tracked by barcode scanning



- Standardized racks with tubes on a barcode scanner
- Scan position of tube on rack and storage in central data base
- Automated cryo storage: Cherry picking of single tubes and output

Requirements for Automatisations

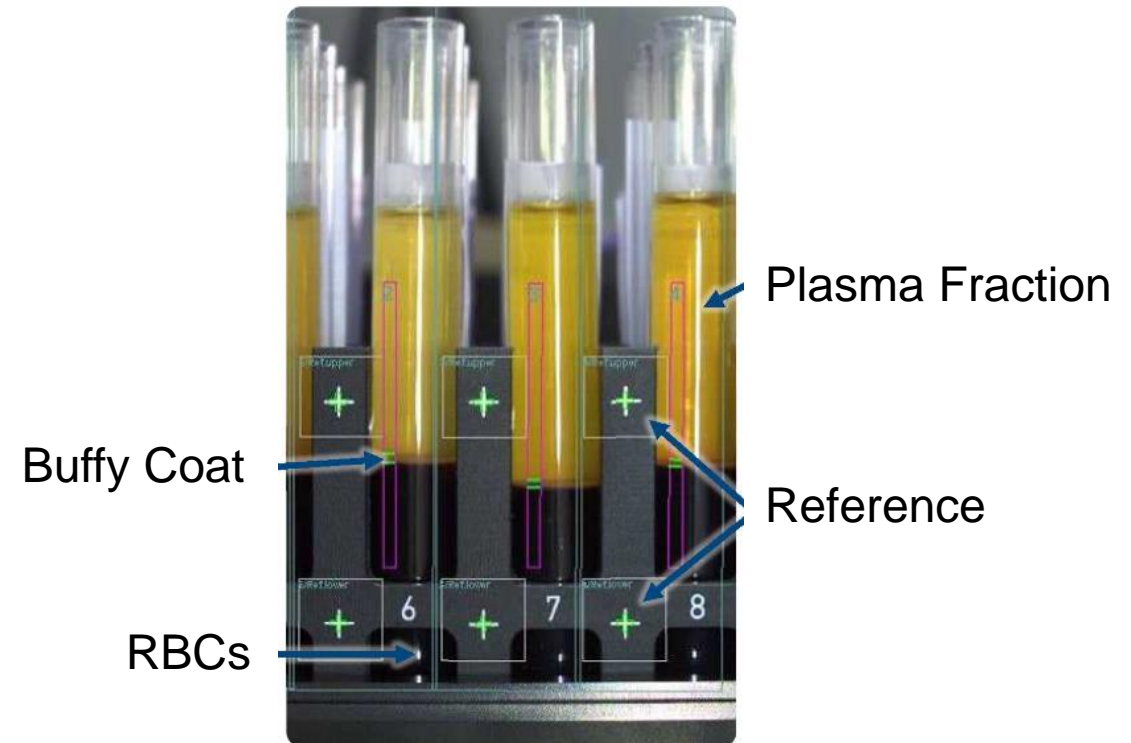
- Automated tube handling
- Tubes must be made for automatisations (opening/ closing)
- As much standardisation as possible
- Barcode identification
- Sample ID is automatically read out by a scanner and is stored together with the sample position in the managing software

Automated Biobank Work Flow



- Fraction analysis
- Aliquoting in 2D- coded target tubes without cap
- Capping target tubes
- Transport in -20°C freezing unit for short term storage

Fully automated blood fractionation



Traceability- Automated Storage Systems



IBBJ 2.0: Automated -80°C Storage for 1.5 Mio tubes

Traceability- Automated Storage Systems



Running aliquoted
samples into the
automated storage

Cap/ Decap tubes

Aliquot samples

Traceability- Automated Storage Systems

IBBJ: Perspectives 2023

- Implementation of a fully automated cryostorage system for storage of > 360.000 vials at -150°C N_2 vapor phase for newly founded **Leibniz Center for Photonics in Infection Research (LPI)**

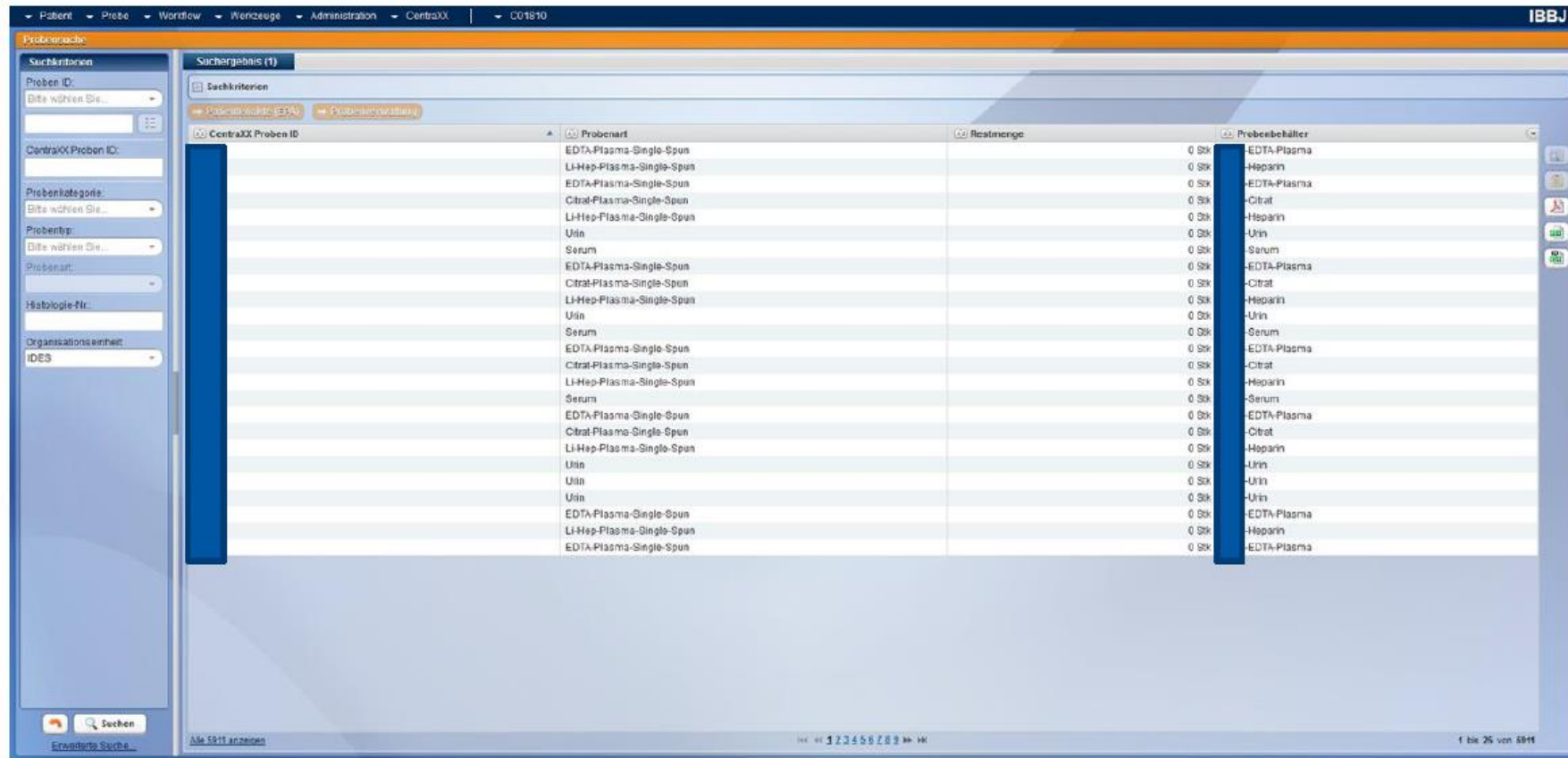


 LEIBNIZ-ZENTRUM für
PHOTONIK in der
INFEKTIONSFORSCHUNG



The IBBJ BIMS *CentraXX*

The Biobanking Information and Management System (BIMS) is especially designed for sampling data handling



The IBBJ BIMS CentraXX

The Biobanking Information and Management System (BIMS) is especially designed for sampe data handling

Sample life cycle

Description	ID	Code / Type	Remaining Amount	Date
▼ Liquid Sample (Urin)	[REDACTED]	Urin	0 pc	04/29/2022 12:44
▼ Liquid Sample (Urin)			5 / 5 Aliquots	
▲ Liquid Sample (Urin)	A1972426	Urin	400.00 µl	05/02/2022 12:06
▲ Liquid Sample (Urin)	A1972428	Urin	400.00 µl	05/02/2022 12:06
▲ Liquid Sample (Urin)	A1972430	Urin	400.00 µl	05/02/2022 12:06
▲ Liquid Sample (Urin)	A1972432	Urin	400.00 µl	05/02/2022 12:06
▲ Liquid Sample (Urin)	A1972417	Urin	400.00 µl	05/02/2022 12:07
▼ Liquid Sample (Urinsediment)			1 / 1 Aliquots	
▲ Liquid Sample (Urinsediment)	0356892403	Urinsediment	400.00 µl	05/02/2022 14:55

Details

Extraction date: 04/29/2022 12:44
 Storage date: 05/02/2022 12:11
 Date of receipt: 05/02/2022 09:06
 Date of distribution: 05/02/2022 12:06
 Date of first storage: 05/02/2022 12:11
 Stocked by: [REDACTED]
 Organization unit: [REDACTED]
 CentraXX Sample ID: 1619933
 Proben ID: A1972426
 Sample status:
 Master sample (Sample ID):
 Diagnosis:
 Histology No.:
 Order No.: [REDACTED]
 Patient consent:
 Age at extraction time:
 Use SPREC: Yes (Code URN - ZZZ - A - B - N - B - S)
 Sample type: Liquid Sample
 Sample kind: Urin
 Remaining amount: 400.00 µl
 Concentration:
 Primary container: Other (ZZZ)
 Pre-Centrifugation delay: RT < 2h (A)
 Post-Centrifugation delay: RT < 1h (B)
 1st centrifugation: RT 10 - 15 min < 3,000 g mit Bremse (B)
 2nd centrifugation: Keine Zentrifugation (N)
 Sample container: Matrix50
 Stock location: UKJ => IBBJ => U2.025 => LICONIC Store NEU (BioLix) => TS01244879 (A5)

Primary organ:
 Site of sample:
 Episode: CentraXX ID: 96239 ([REDACTED])
 Study:
 Project:
 Cooperation partner:
 Appointment number:
 Passage cell line:
 Freezer thaw cycle:
 SOP deviation: No
 Note:
 CentraXX Patient ID: 49821
 MPI: [REDACTED]

Stock location

	1	2	3	4	5	6	7	8	9	10	11	12
A												
B												
C												
D												
E												
F												
G												
H												

The IBBJ BIMS *CentraXX*

All processes from sample arrival to aliquotation, storage and shipping are documented

Data is automatically updated when a sample barcode is scanned

The database can be searched for:

- Sample category
- Sample name
- Storage place
- Associated study
- Every other data attached to the sample

- ✓ For every sample information about the aliquotation status and the remaining number of aliquots is available
- ✓ Detailed history for the sample including its former storage places is available

Data Protection

Peter Meyer

Date of birth 12.05.1965, Address Jena, Sample ID 125656, Diagnosis
Diabetes II, ... Freezer 3 Box 37.

Accumulation of data is convenient but not always legal!

Need for Data Protection

Peter Meyer

Date of birth 12.05.1965, Address Jena, Sample ID 125656, Diagnosis Diabetes II, ... Freezer 3 Box 37.

Examples of types of data:

- **MDTA:** medical or research data
- **OrgDat:** place of storage, freezing date
- **ProbDat:** sample data: results of assays, measurements
- **IDAT:** patient identifying data (name, health insurance number)
- ❖ It is not allowed to store IDAT in the Biobank
- ❖ Every employee should always only see the data he needs to know
- ❖ Pseudonymisation of samples: no sample is stored under patients name but a sample number.

MDTA: Material and Data Transfer Agreement



 UNIVERSITÄTS
KLINIKUM
jena

MATERIAL AND DATA TRANSFER AGREEMENT (MDTA)
(Academic Relations)

Supplier: Integrated Biobank Jena (IBBJ)
University Hospital Jena
Am Klinikum 1
07747 Jena
Germany

 iiIBBJ
Integrated Biobank Jena

Represented by: PD Dr. Dr. Michael Kiehntopf

Recipient:

Represented by:

Material: Human blood serum and plasma
(according to the attached sample- and data description)

Use: Analysis of XYZ quality indicators in the joined research project:
XXX

Every time someone requests samples from the biobank a MDTA is needed to:

- Ensure that the recipient complies to data protection laws
- Specify the allowed usage of the material (an ethics committees vote is always needed)
- Agree on the usage of produced data (publication, patents)
- ...

The IBBJ



IBBJ

... is part of the research infrastructure of the University Hospital Jena.

Its mission is to support medical research through the quality-assured, standardized and fiduciary collection, storage and provision of liquid and non-liquid high-quality biomaterials and corresponding clinical context data.



Biobank Competence Network
Sepsis



Biobank Centre for Innovation
Competence (ZIK) Septomics



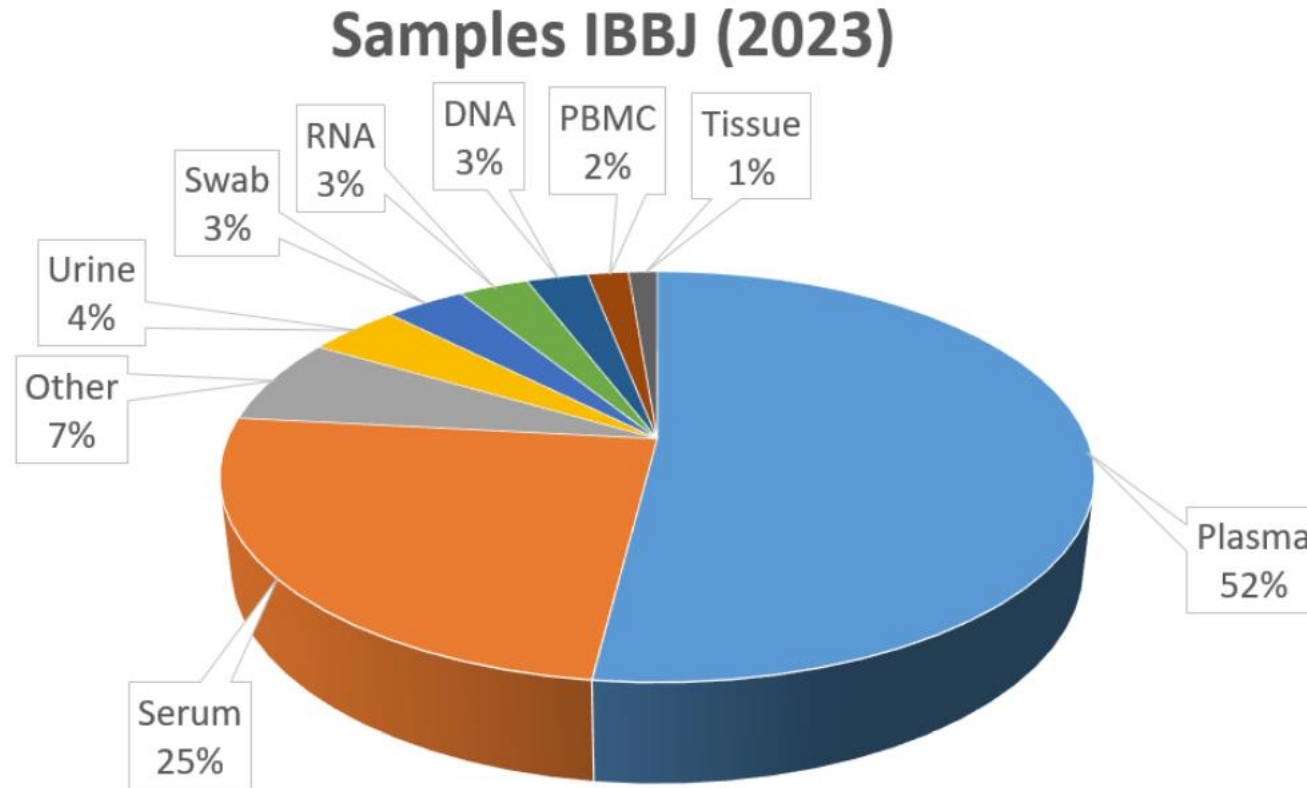
Biobank Center for Sepsis Control
and Care

The Integrated Biobank Jena (IBBJ)- Expansion to the UKJ Biobank Infrastructure



Sample Numbers and Capacities*

Liconic: 1.5 Mio*
 UTKs 21 = 1.88 Mio*
 LN₂ 8 = 0.55 Mio*
 ASKION = 0.38 Mio*
 * 500 µL Tubes



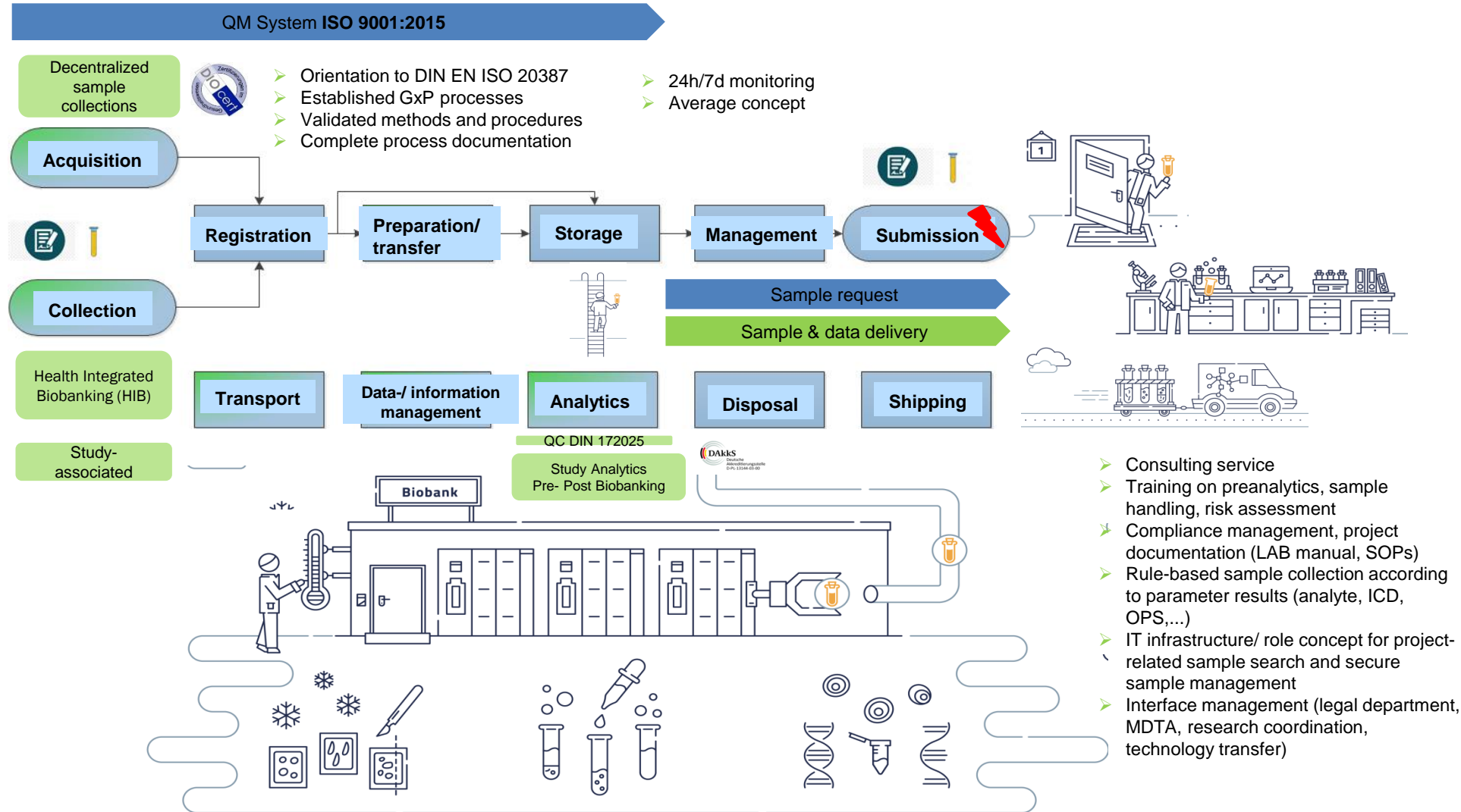
91 collections

Total sample size:
860.000

Core Services

The IBBJ:

- It is partner for the conduction of studies and collections
- It is being run by a Steering Committee and sample access will be organised via a Use and Access Committee



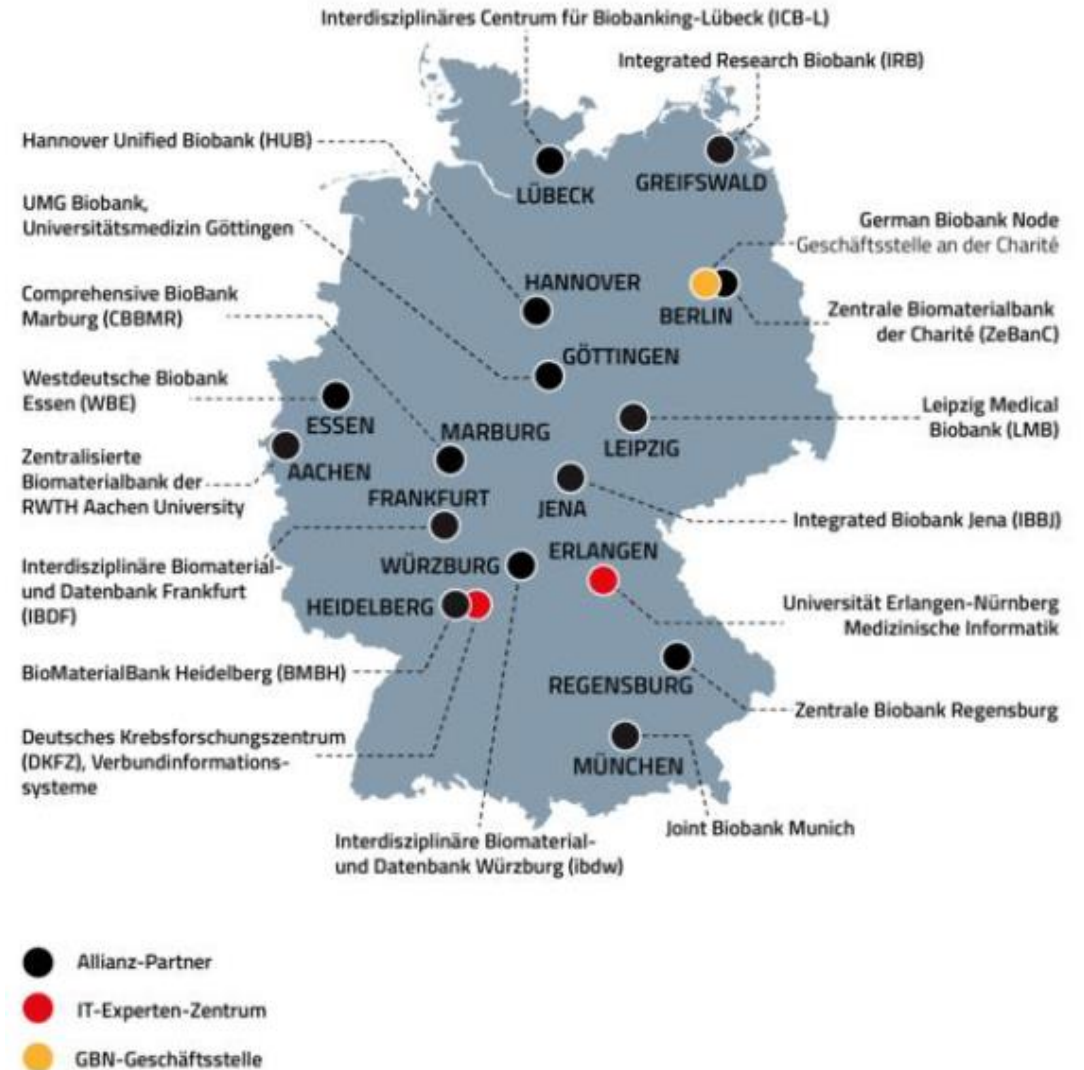
Summary

- Biobanked samples are important for clinical research and medical progress
- Sample integrity plays a major role for downstream analysis and data interpretation as time point of sampling is linked to data acquisition
- Sample integrity can be impaired by an interplay of pre- analytical factors and loss of identification and traceability
- How to solve the pre- analytical problem:
 - ❖ Standardized working with standard operating procedures
 - ❖ High grade of automation
 - ❖ Documentation of processing times and all deviations from protocol

Further Information

German Biobanks are organised in the German Biobank Alliance (GBA) and part of the European Network
BBMRI-ERIC
ABIDE

These Biobanks offer a centralised sample search and aim for a harmonization and standardization of biobanking processes and protocols.



Biobanking offers access to samples to other researchers

- Biobanking enhances the visibility of sample collections by displaying them on their websites and national or international sample search sites like the BBMRI-ERIC Directory:
<https://directory.bbmri-eric.eu/menu/main/app-molgenis-app-biobank-explorer/biobankexplorer>
- Using preexisting sample collections can be more cost effective and much faster than setting up your own collection. In addition, it reduces the overall amount of samples obtained from patients.

Movies and Links to some large Biobanks

Movie about Biobanks produced by the German Biobank Alliance
(showcasing the ZeBank at the Charité)

<https://youtu.be/njmeLYZujZo>

Germany NaKo Gesundheitsstudie (Nationale Kohorte; NaKo)

<https://nako.de/infomationen-auf-englisch/>

UK Biobank

<https://www.ukbiobank.ac.uk/>

Thank you for your attention!

Daniel.Barth@med.uni-jena.de

IBBJ Tour