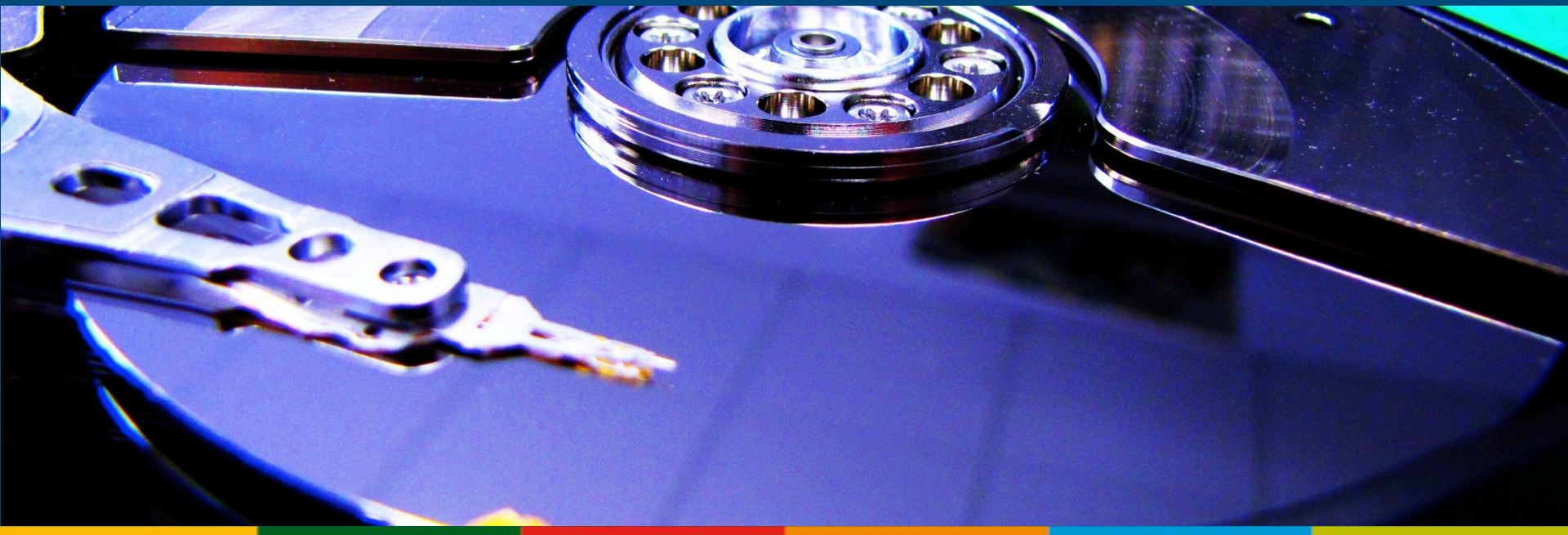


DLCL Key Technologies: Status + Future

**Rainer Stotzka, Nick Kepper,
Richard Grunzke, Volker Hartmann, Michael Hausmann, Jürgen
Hesser, Thomas Jejkal, Ralph Müller-Pfefferkorn, Halil Pasic,
Rasmus Schröder, Danah Tonne, Xiaoli Yang**



5 (+3) subprojects:

- ***Lightoptical Nanoscopy*** (Heidelberg, Mannheim)
- High Throughput Microscopy: ***Gen Scans***
(TU Dresden + MPI of Molecular Cell Biology and Genetics)
- High Throughput Microscopy: ***Selective Plane*** Illumination
Microscope (Karlsruhe)
- ***Electron*** Microscopy (Heidelberg)
- Material Research: ***Tomography*** Beam Line at ANKA (Karlsruhe)
- (Arts & Humanities): ESFRI ***DARIAH*** + BMBF ***eCodicology***
- *μ Tomography: in preparation (Institut für Angewandte Materialien,
Karlsruhe)*

Organization

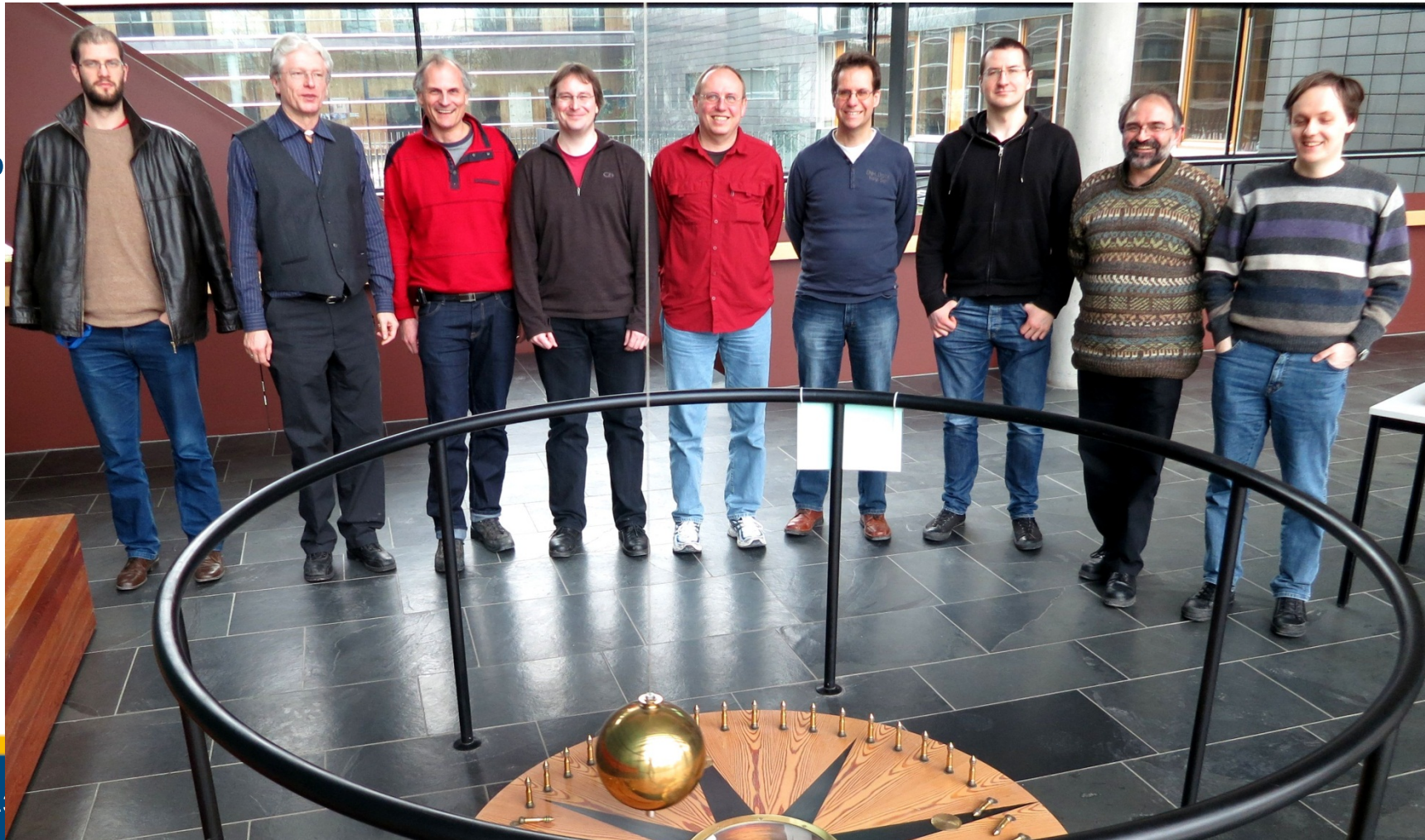


Speakers: Rainer Stotzka, Nick Kepper

Monthly phone conferences (each 2. Thursday): November 12, January 13

Face-2-Face meeting:

14. Feb. 2013, Heidelberg



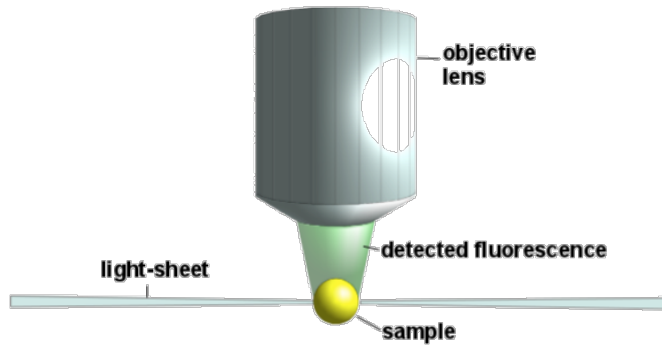
Publications (Paper/Book, Talk, LSDMA)



- [1] P. Kilpatrick, P. Milligan, R. Stotzka (eds.), Proceedings of the 21st International Euromicro Conference on Parallel, Distributed and Network-based Processing, IEEE Conference Publishing Services, DOI 10.1109/PDP.2013, 2013
- [2]+[3] Danah Tonne, **Jedrzej Rybicki**, Stefan E. Funk, and Peter Gietz, Access to the DARIAH Bit Preservation Service for Humanities Research Data, 21st Euromicro International Conference on Parallel, Distributed, and Network-Based Processing, pp. 9-15, DOI 10.1109/PDP.2013.12, 2013
- [4]+[5] Xiaoli Yang, Thomas Jejkal, Halil Pasic, Rainer Stotzka, Achim Streit, Jos van Wezel, and Tomy dos Santos Rolo, Data Intensive Computing of X-Ray Computed Tomography: 21st Euromicro International Conference on Parallel, Distributed, and Network-Based Processing, pp. 86-93, DOI 10.1109/PDP.2013.21, 2013
- [6] Danah Tonne, Rainer Stotzka, Integration von eCodicology in die DARIAH-Dienstewelt, Tagung Digitale Rekonstruktion Mittelalterlicher Bibliotheken, 18.1.2013, eingeladener Vortrag, Trier
- [7] Haas, D.; Mexner, W.; Spangenberg, T.; Cecilia, A.; Kopmann, A.; Balzer, M.; Vogelgesang, M.; Pasic, H.; Chilingaryan, S., Status of the ultra fast tomography experiments control at ANKA. 9th Internat.Workshop on Personal Computers and Particle Accelerator Controls (PCaPAC-2012), 2012
- [8] Halil Pasic, Managing Large Scale Scientific Data at ANKA, Data Rate Initiative of the HGF Programme PNI, 2.3.2013
- [9] Xiaoli Yang, Thomas Jejkal, Halil Pasic, Rainer Stotzka, Achim Streit, Jos van Wezel, Tomy dos Santos Rolo, Algebraic Reconstruction of X-ray Tomography on parallel computing Architecture of LSDF, Data Rate Initiative of the HGF Programme PNI, 2.3.2013
- [10] Rainer Stotzka, Marcus Hardt, Facilitating Data Upload, Access and Deployment, eIRG, Amsterdam, 2012
- [11] Sutter, M.; Hartmann, V.; Götter, M.; Wezel, J.; Trunov, A.; Jejkal, T. & Stotzka, R. File systems and access technologies for the large scale data facility Remote Instrumentation for eScience and Related Aspects, Springer, 2012, 239-256
- [12] Vanscheidt, P.; Rapp, A. & Tonne, D. (2012), 'Storage Infrastructure of the Virtual Scriptorium St. Matthias"Digital Humanities 2012', Poster, 2012
- [13] **DLCL-Key Aktivitäten Quartal 3/2012, October 2012**
- [14] **DLCL-Key Aktivitäten Quartal 4/2012, February 2013**

- [15] Grunzke et al., A Data Driven Science Gateway for Computational Workflows, in: UNICORE Summit 2012 Proceedings, pages 35-49, 2012
- [16] Gesing et al., A Science Gateway Getting Ready for Serving the International Molecular Simulation Community, in: Proceedings of Science, 2012
- [17] Gesing, Grunzke, et al, A Single Sign-On Infrastructure for Science Gateways on a Use Case for Structural Bioinformatics (2012), in: Journal of Grid Computing, 10:4(769-790)
- [18] Herres-Pawlis et al., Workflow-enhanced conformational analysis of guanidine zinc complexes via a science gateway, Studies in Health Technology and Informatics, 175:142-151, IOS Press., 2012
- [19] Pyatykh, Stanislav, Zheng, Lei and Hesser, Jürgen (2012), "Efficient method of pixel neighborhood traversal", Journal of Visual Communication and Image Representation, 23, 1: 719-728, 2012.
- [20] Teuber, T., Remmele, S., Hesser, J. and Steidl, G. (2012), "Denoising by second order statistics", Signal Processing, 92, 12: 2837-2847, 2012.

Single Plane Illumination Microscope



Novel microscope to observe zebra fish embryos in vivo:

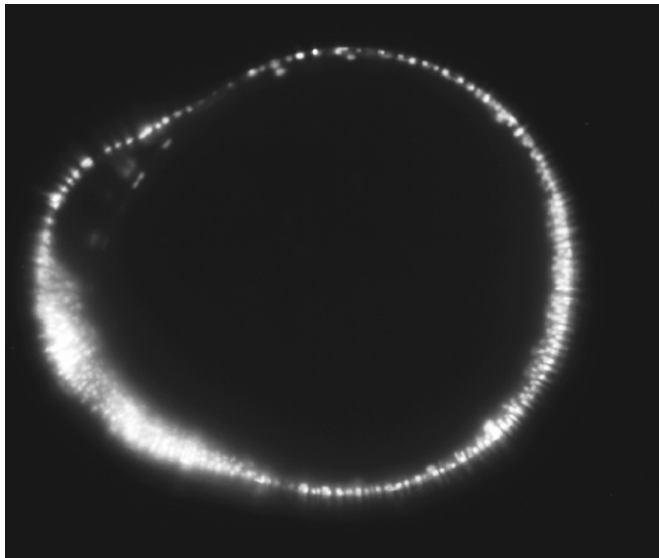
- Very high 3D resolution
- Short data acquisition time

Each day one observation of the development of an embryo:

Data: 7 TB/10 h → Σ 300 TB

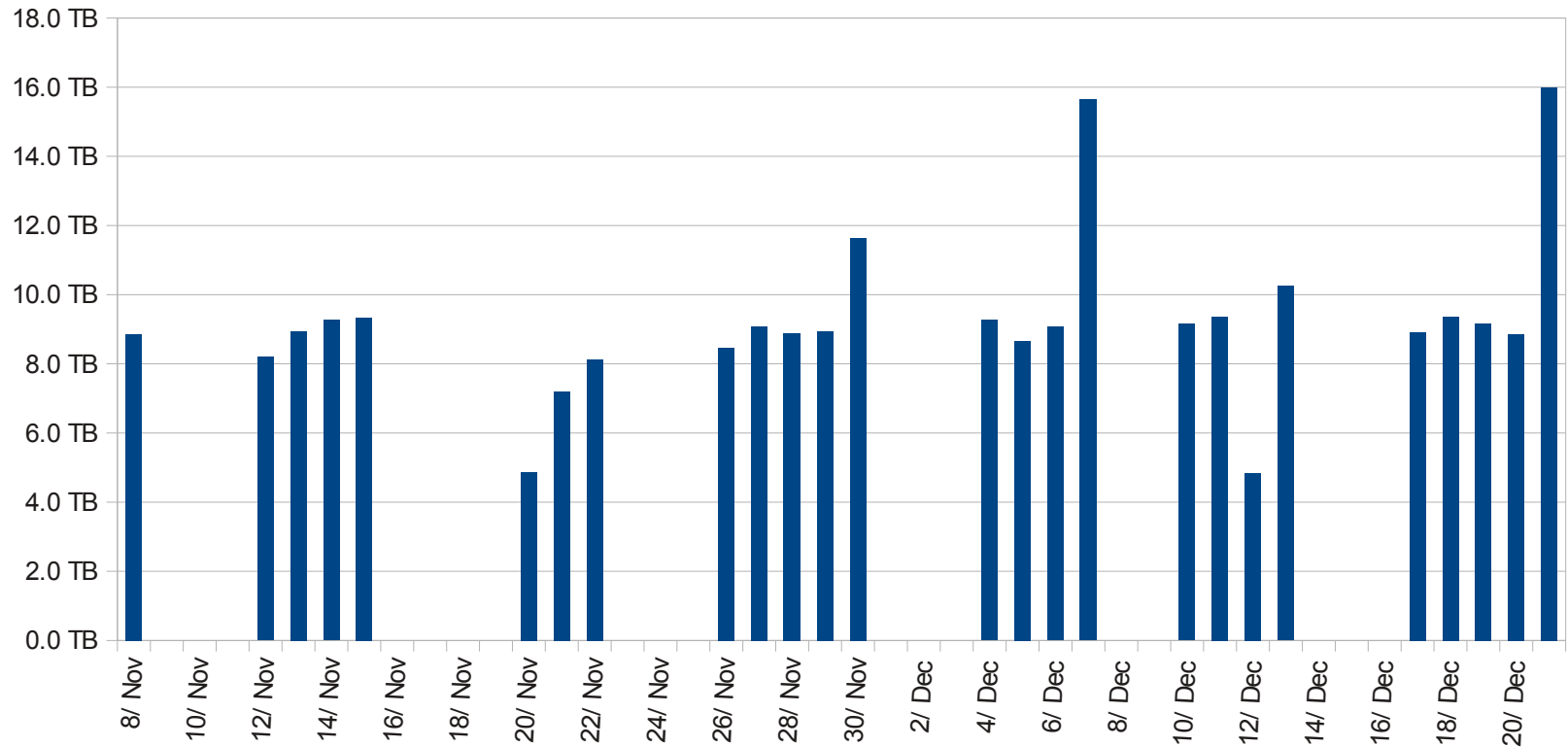
Challenges:

- Meta data extraction
- **Data transfer** source to LSDF > 400 MB/s
- Data intensive **analysis** < 24 h:
registration, fusion, segmentation, tracking, statistics



Status SPIM

Datatransfer SPIM from CS to CN
8/Nov - 21/Dec



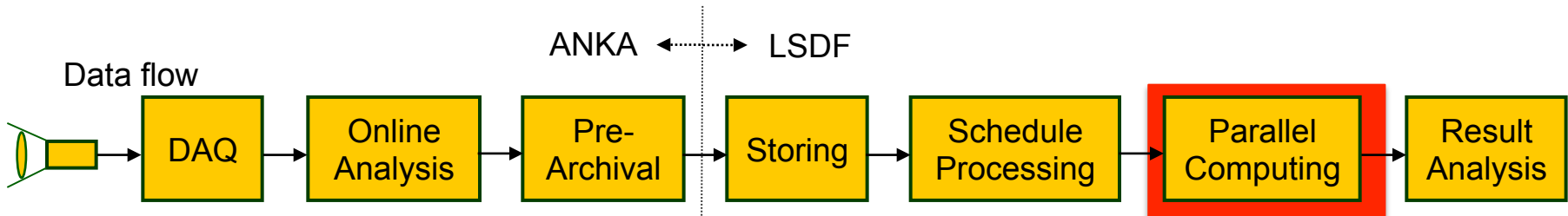
Transfer:

Total amount: 253 TB
Files: 53,333
Total transfer time: 7d 15h

Transfer rate:

Maximum: 442 MB/s
Minimum: 366 MB/s
Average: 403 MB/s

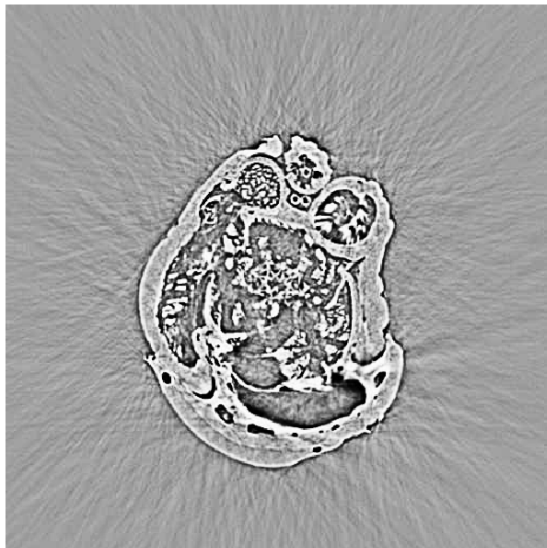
Material Research: Tomography Beam Line at ANKA



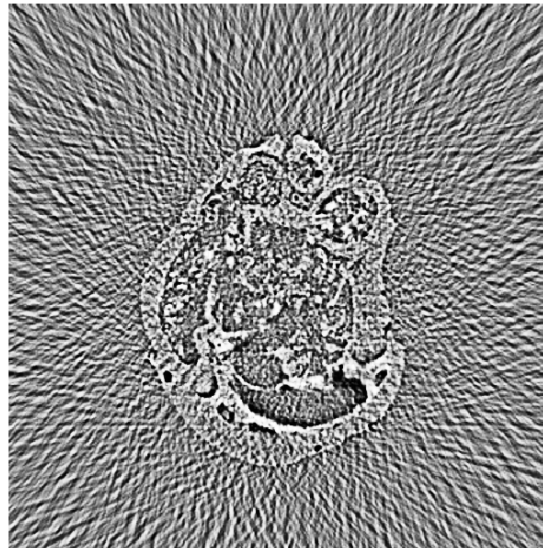
Novel algorithms enabling the world fastest tomography beamline

- Compressive Sampling Algebraic Reconstruction (**CS-ART**)

Standard: 1500 projections



Standard: 60 projections



CS-ART: 60 projections



Material Research: Tomography Beam Line at ANKA

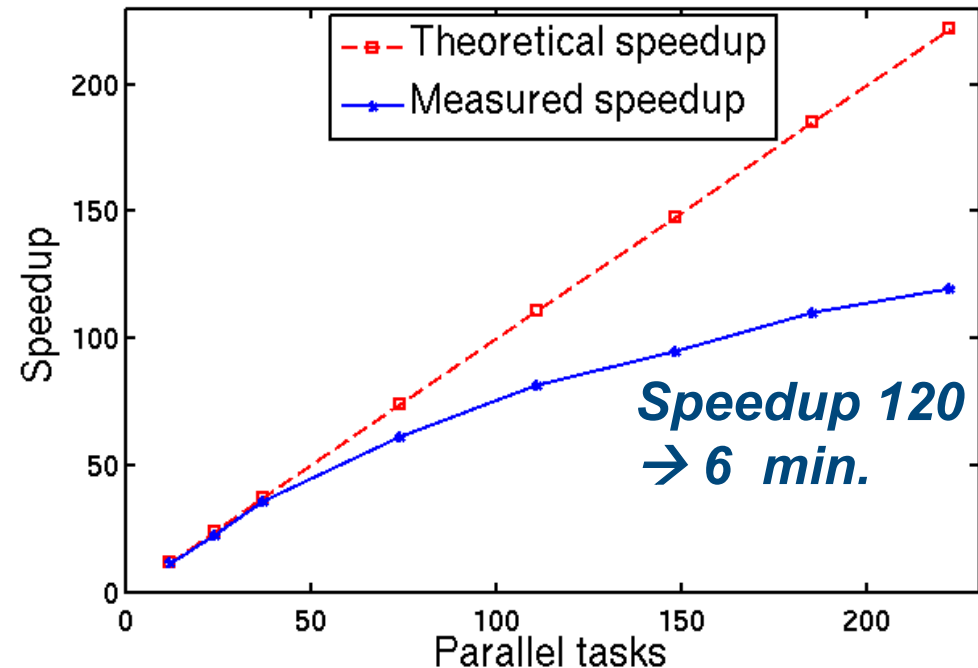


- CS-ART implemented in Matlab
- Computational expensive

→ **11 h !!!!** >> **1 min.**

- Parallelized using DSIT DIC, speedup of 120 

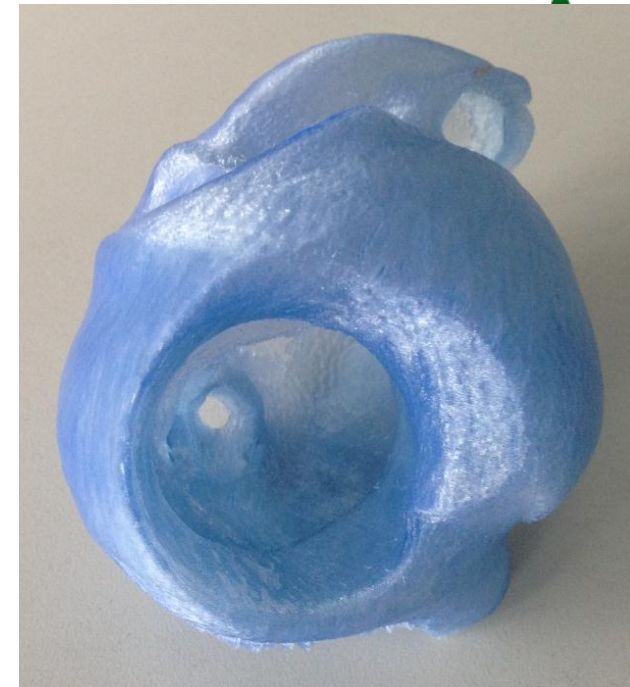
- CS-ART: **better** images
- **Automatic** DIC workflow 2014
- **Enables** world's fastest tomography beamline
- **General**, not limited to ANKA



Material Research: Data to Finger

Printing reconstructed volumes in 3D

HECTOR Seminar: highly skilled MINT students



Projects considered:

- Changes density and packaging in hetero- and euchromatin by radiation damage via high resolution SPDM images
- Analysis genome architecture of breast cancer marker Her2/neu as marker for the tumorigenesis
- Micro- and nanoarchitecture of Her2/neu receptor clusters before and after amplification of genes in tumor cells

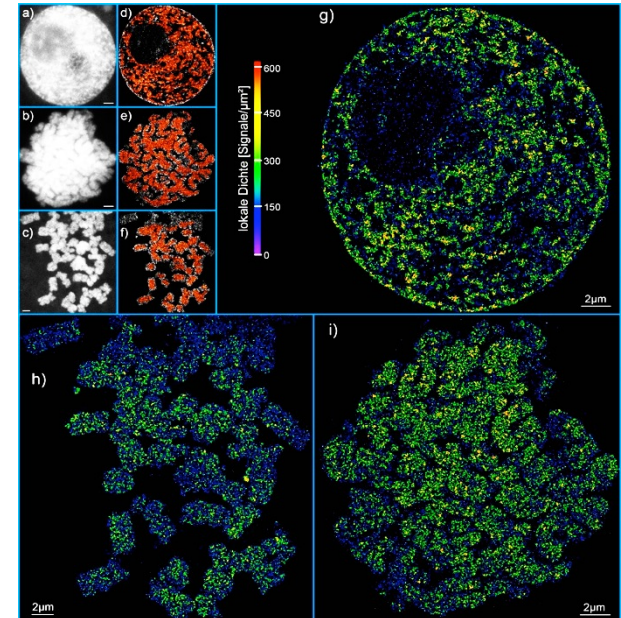
Tools:

- **Automatic analysis** with parallelized MatLab programs: localization, distribution, and classification

Visualization:

- Scientific **visualization** with ParaView

SPDM of nucleosomes



Important requirements:

- Data storage facilities
- Data life cycle definition
- Application dependent evaluation
- Meta data handling

Electron Microscopy

To do:

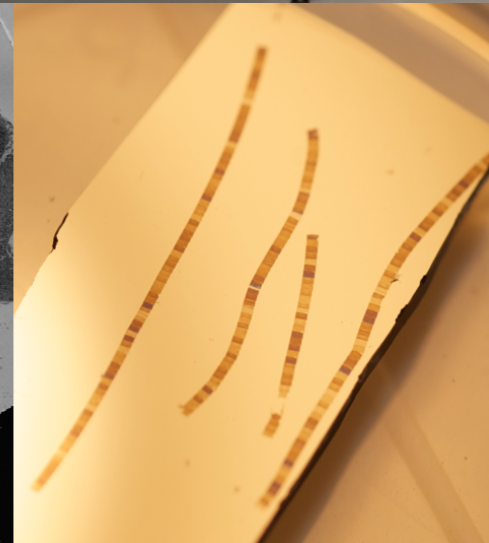
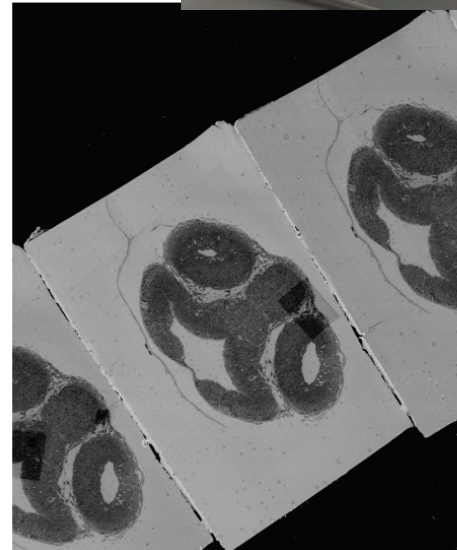
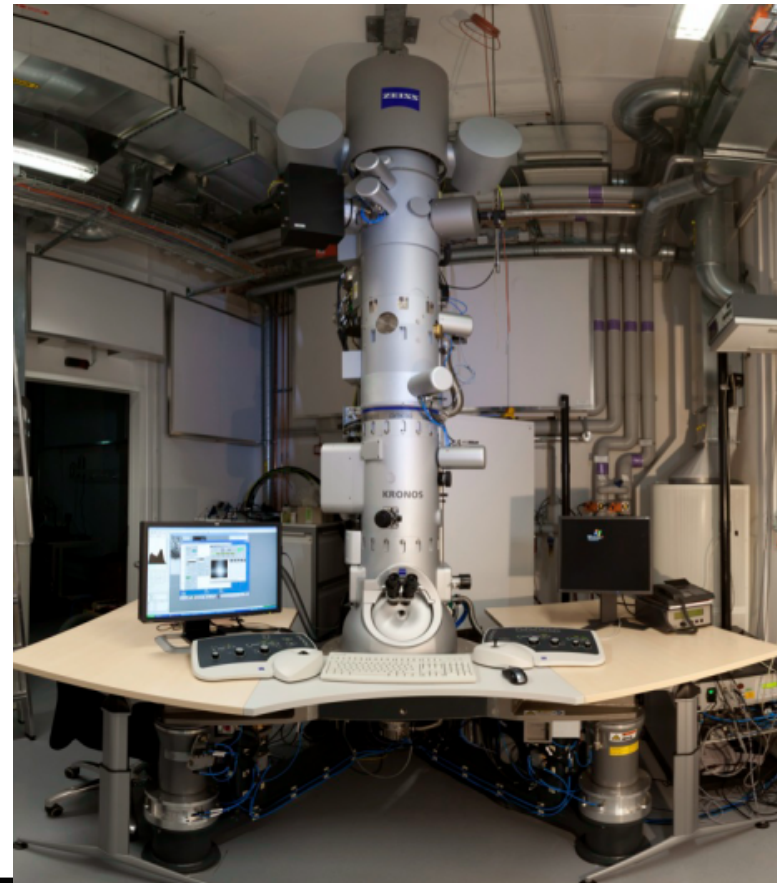
- Acquiring data: automation, workflow, electronic lab book (EMEN2)
- Image data data base structure (LEGINON)
- Storage (local for processing?)
- Processing pipeline (registration, segmentation) needs to be developed
- Data browsers, 3D-Viewers? Data reduction?
- Archiving?

30 GB / image

Depth in 3rd dimension $\rightarrow 10^5$ images

1 eye: 3 PB

Image sources: Rasmus R. Schröder
CellNetworks, BioQuant, CAM, Universität Heidelberg



DARIAH + eCodicology



DARIAH-DE

Digital Research Infrastructure
for the Arts and Humanities

ESFRI project,
preservation
infrastructure

eCodicology: BMBF joint project (Karlsruhe,
Trier, Darmstadt), PhD students + **code camps**

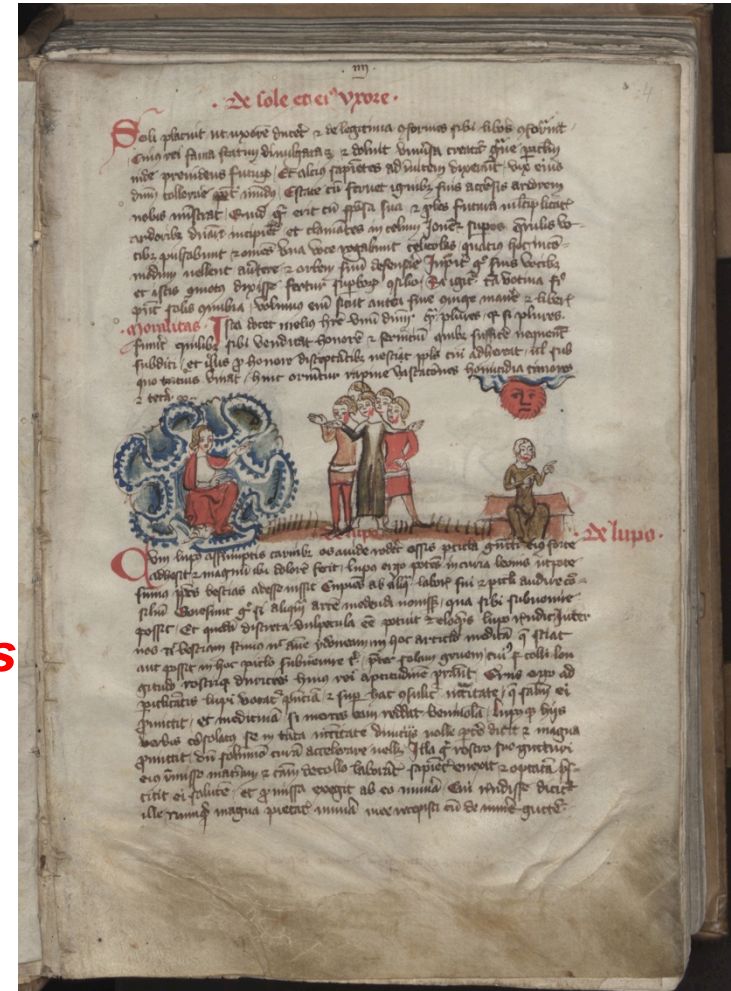
- Automatic **analysis** of 500 medieval
manuscripts, 150,000 pages

Most important requirements:

- Meta data handling
- Reproducibility
- Performant data access
- Data intensive workflows

DSIT Services

- ❖ Federated Data Access
- ❖ Meta Data Catalogues and Repositories
- ❖ Data Intensive Computing



Some GENERAL Results

Similarities found in nearly all sub-projects:

- ***Meta data is important !***
- Automatic meta data extraction required
File format → standard meta data format

Experiment view

- EMEN2
- „by hand“
- proposal DB

Instrument view

- OME
- Tango
- Labview
- LEGINON

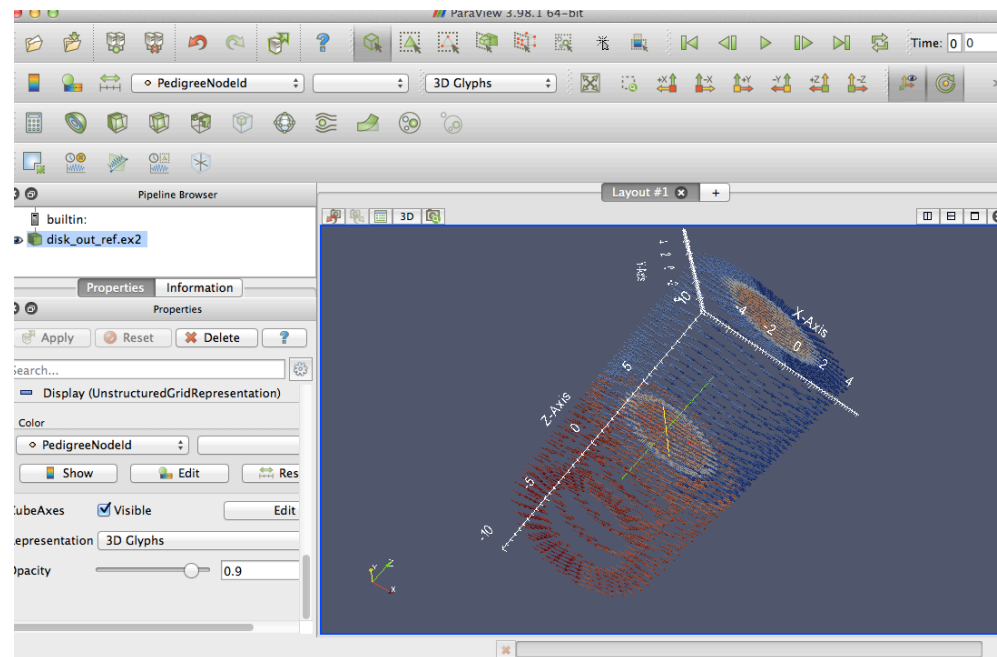
- Data provenance
- Where to store the data?
→ KIT LSDF as the platform for tool development & testing
→ TU Dresden

Some GENERAL Results

Established knowledge exchange

Common tools defined:

- **Data transfer client** requires federated AAI !!!
- **Meta data** management tools
- Data **analysis** tools:
 - Digital image processing lib. consisting of basic functions, e.g. denoising
- **Visualization**



Additional Funding



- **eCodicology**: 1 (+2) PhD positions
Relevance to LSDMA:
 - Data life cycle and data sharing
 - Meta data
 - Image analysis
 - Data intensive computing
- 1 PhD student **scholarship** (4 years)
Relevance to LSDMA:
 - Algorithms for volume reconstruction & analysis
 - Data intensive computing
- **Hector** Seminar: 1 year's students project
Relevance to LSDMA:
 - Novel data representations
- **MoSGrid** (→ ER-flow : 2 years)
Relevance to LSDMA:
 - Science gateway
 - Workflow management & sharing
 - Data interoperability in workflows
- **MoSGrid** (→ SCI-BUS: 3 years)
Relevance to LSDMA:
 - Science gateway
 - Meta data repository
 - Result visualization
- BMBF Hausmann: „Fokussierte Fluoreszenzmarkierung (COMBO-FISH)“
- BMU Hausmann: „3D/4D Architektur von chromosomalen Bruchpunktregionen in Zellkernen nach Bestrahlung von Krebs- und nicht-Krebszellen“

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- J. Lux, ..., M. Hausmann, Reconstruction and optical sequencing of a DNA repeat expansion unit by location nanoscopy.
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- Halil Pasic et al., Managing Large Scale Scientific Data at ANKA
- Grunzke et al., Standard-based Metadata Management for Molecular Simulations, 2013
- Krüger, Gesing, Grunzke, et al. , The MoSGrid Science Gateway, 2013
- Herres-Pawlis et al., User-friendly Workflows in Quantum Chemistry, 2013
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Conclusions

- a. DLCL-Key: broad **spectrum** of scientific projects
- b. Heterogeneous team, highly motivated
- c. DSIT services** will be required
- d. Results:** established knowledge exchange
→ common **tools** defined

Next steps:

- Establish software exchange, common tool development and integration