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**UNIVERSITY  
OF BERN**

# *u<sup>b</sup>* X-ray microtomography

## 9256-HS2024-0: Advanced Microscopy

**David Haberthür**

Institute of Anatomy, December 20, 2024

# *u<sup>b</sup>* Grüessech mitenang!

- David Haberthür
  - Physicist by trade
  - PhD in high resolution imaging of the lung, Institute of Anatomy, University of Bern, Switzerland
  - Post-Doc I: TOMCAT, Swiss Light Source, Paul Scherrer Institute, Switzerland
  - Post-Doc II: µCT group, Institute of Anatomy, University of Bern, Switzerland

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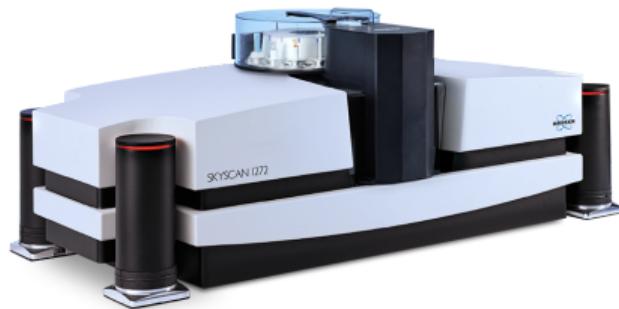
# Grüessech from the $\mu$ CT group



[David.Haberthuer@unibe.ch](mailto:David.Haberthuer@unibe.ch) [Ruslan.Hlushchuk@unibe.ch](mailto:Ruslan.Hlushchuk@unibe.ch) [Oleksiy.Khoma@unibe.ch](mailto:Oleksiy.Khoma@unibe.ch)

# $\mu$ CT-group

- microangioCT [1]
  - Angiogenesis: heart, musculature [2] and bones
  - Vasculature: (mouse) brain [3], (human) nerve scaffolds [4], (human) skin flaps [5] and tumors
- Zebrafish musculature and gills [6]
- (Lung) tumor detection and metastasis classification [7]
- Collaborations with museums [8] and scientist at UniBe [9], [10] to scan a wide range of specimens, from human hearing bones to meteorites
- Automate all the things! [11], [12]



[bruker.com/skyscan1272](http://bruker.com/skyscan1272)

*u*<sup>b</sup>

# Contents

Overview

    Imaging methods

Tomography

    History

    Tomography today

    Interaction of x-rays with matter

A scan, from *getting started* to *nice image*

Example: A study about teeth

    Overview

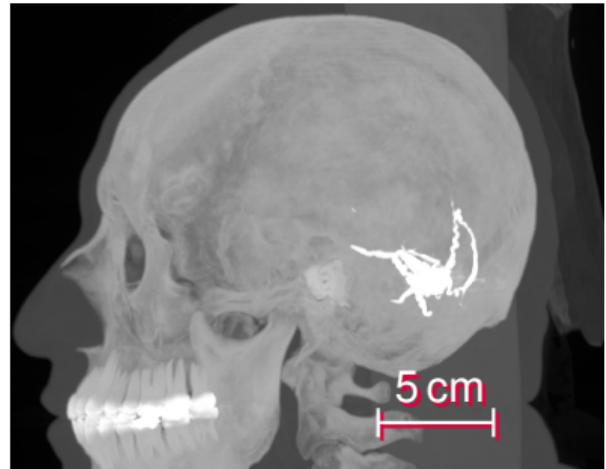
    Materials & Methods

    Results

*u<sup>b</sup>*

# micro-Computed tomography

- Allows for imaging dense and non-transparent samples
- Non-destructive imaging
- Results in three-dimensional images
- Covers a very large range of sample sizes

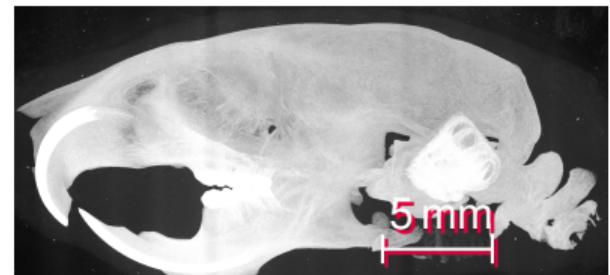


From [13], Subject C3L-02465

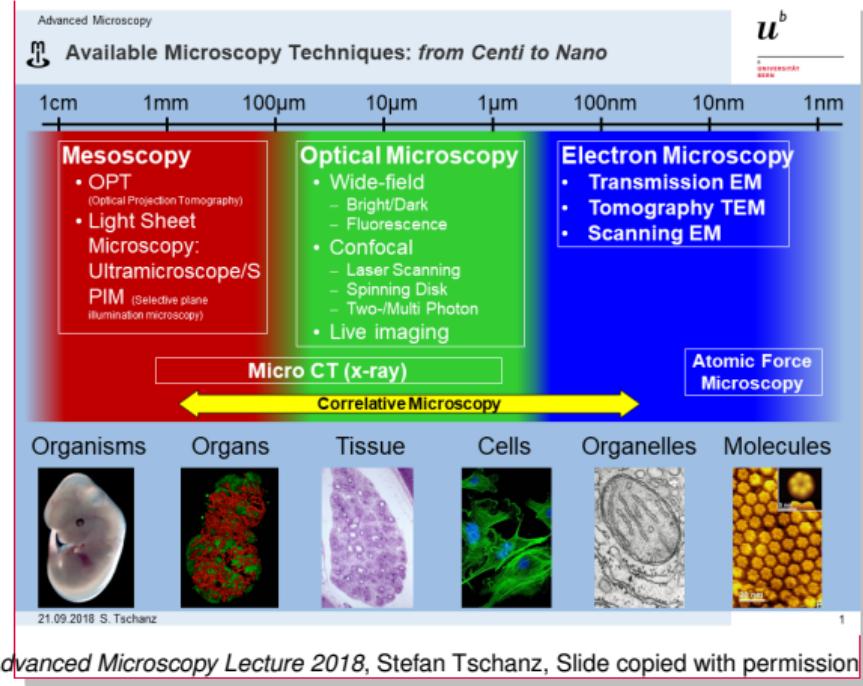
*u<sup>b</sup>*

# micro-Computed tomography

- Allows for imaging dense and non-transparent samples
- Non-destructive imaging, thus compatible with routine sample preparation
- Results in three-dimensional images with  $\mu\text{m}$  resolution
- Covers a very large range of sample sizes
- (Small) biological samples
- Enables correlative imaging pipelines, scanning of precious biological samples, as well as museum & collection material



# Imaging methods



- *Light Sheet Microscopy* by Nadia Mercader Huber
- X-ray imaging
- Electron microscopy
  - *Transmission Electron Microscopy* by Dimitri Vanhecke
  - *Scanning Electron Microscopy* by Sabine Kässmeyer and Ivana Jaric
  - *Cryoelectron Microscopy & Serial Block Face SEM* by Ioan Iacovache

Advanced Microscopy Lecture 2018, Stefan Tschanz, Slide copied with permission

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# CT-Scanner



[youtu.be/2CWpZKuy-NE](https://youtu.be/2CWpZKuy-NE)

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# CT History

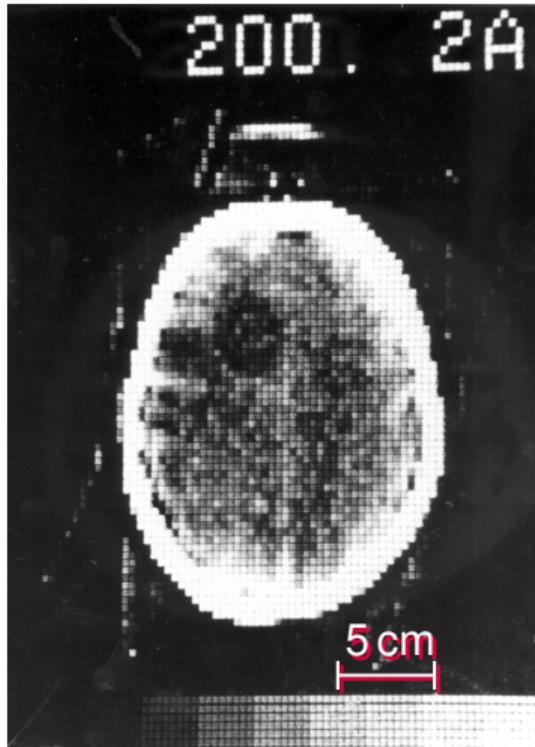
- 1895: Wilhelm Conrad Röntgen discovers X-rays



w.wiki/BHAN

# CT History

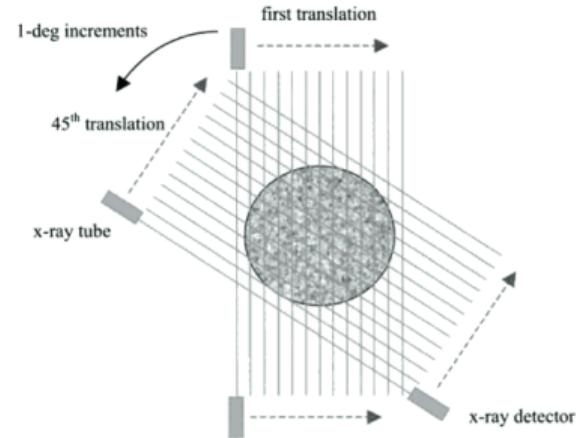
- 1895: Wilhelm Conrad Röntgen discovers X-rays
- 1963: Cormack used a collimated  $^{60}\text{Co}$  source and a Geiger counter as a detector [14]
- 1976: Hounsfield worked on first clinical scanner [15]



From [16], Figure 5

# CT History

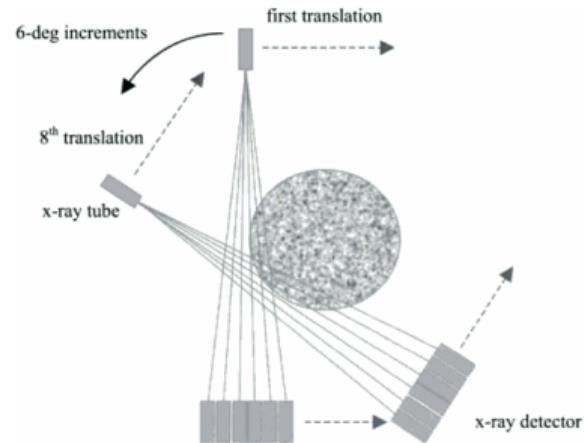
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- 1976: Hounsfield worked on first clinical scanner [15]
- CT scanner generations
  - First generation



From [17], Figure 1.12

# CT History

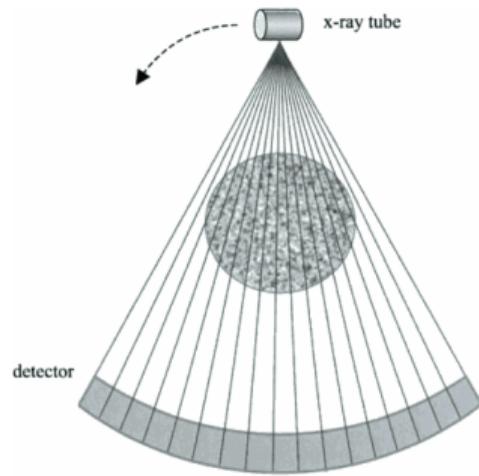
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- CT scanner generations
  - First generation
  - Second generation



From [17], Figure 1.13

# CT History

- 1895: Wilhelm Conrad Röntgen discovers X-rays
- 1963: Cormack used a collimated  $^{60}\text{Co}$  source and a Geiger counter as a detector [14]
- 1976: Hounsfield worked on first clinical scanner [15]
- CT scanner generations
  - First generation
  - Second generation
  - Third generation



From [17], Figure 1.14

# $\mu$ CT History I

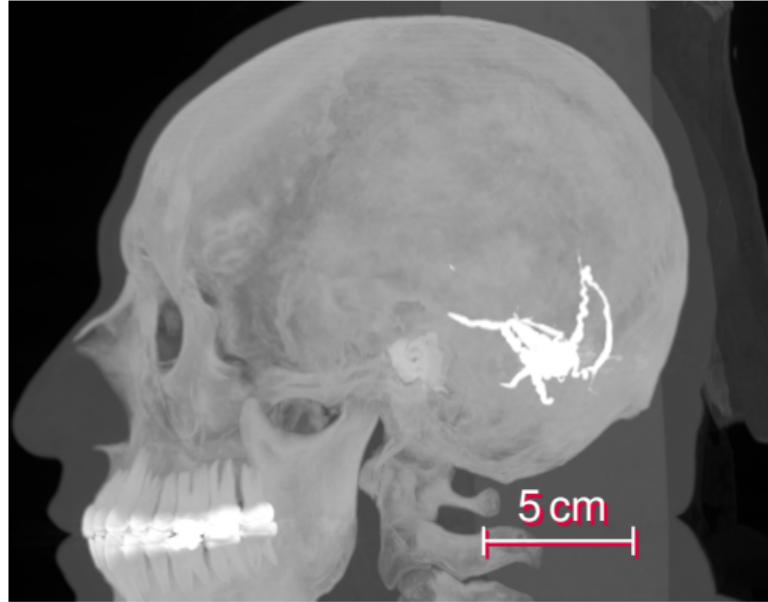
- X-ray computed tomography began to replace analog focal plane tomography in the early 1970s [18]
- Non-medical use in the late 1970s, for detection of internal defects in fabricated parts and equipment
- Lee Feldkamp [19] developed one of the early laboratory microCT systems by assembling a micro-focus cone beam x-ray source, specimen holder and stages, and an image intensifier at Ford Motor Company's Scientific Research Laboratory to nondestructively detect damage in ceramic manufactured automobile parts
- Feldkamp met with scientists at Henry Ford Hospital and University of Michigan interested in understanding the relationship between the microstructure and biomechanical function of trabecular bone to study osteoporotic fractures [20]

# $\mu$ CT History II

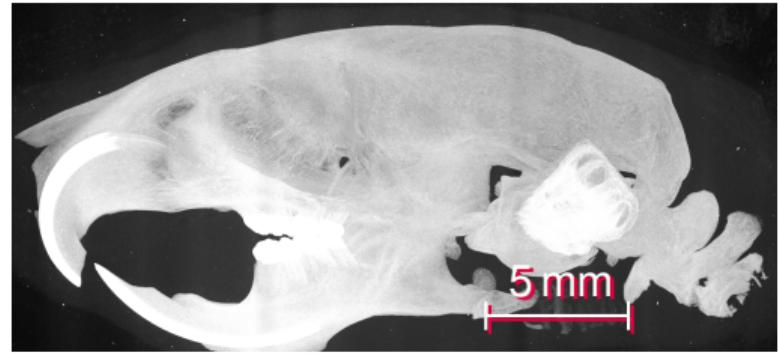
- $\mu$ CT was first reported in the 1980s, for scanning gemstones to cut out the largest possible one
- Today: Nondestructive imaging for quantifying the (micro)structure of organic materials
  - Mineralized bone tissue and the relationships between the mechanical behavior of bone to its structural and compositional properties
  - Teeth and their internal details
  - Soft tissues and vasculature using radio-opaque contrast agents
  - Characterization of anatomical details in high resolution
- $\approx$ 2500  $\mu$ CT systems are in use worldwide with over 1000 publications annually

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# Why $\mu$ CT?

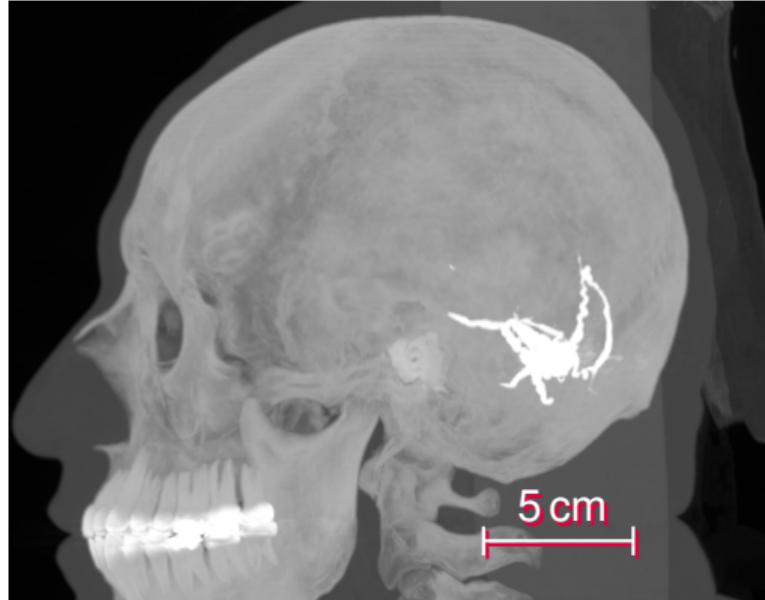


From [13], Subject C3L-02465



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# Why $\mu$ CT?

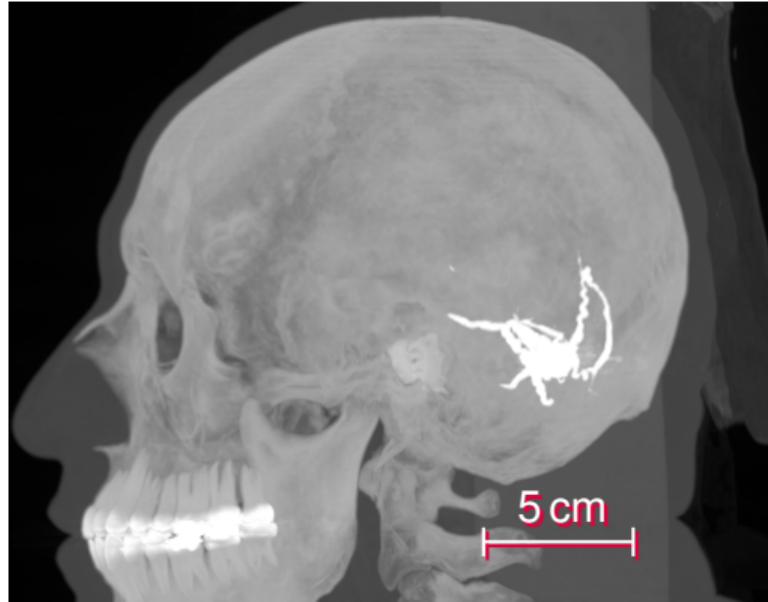


From [13], Subject C3L-02465

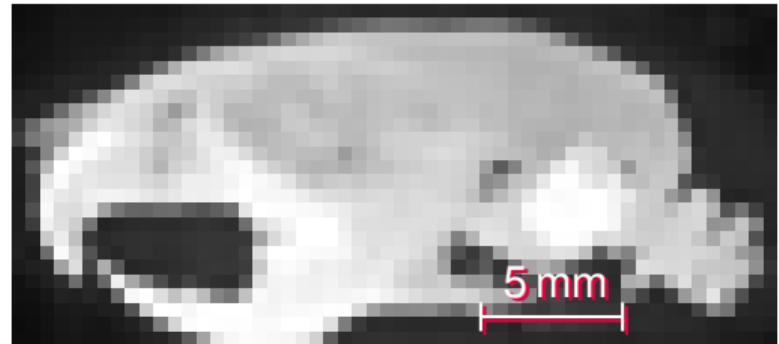


$u^b$

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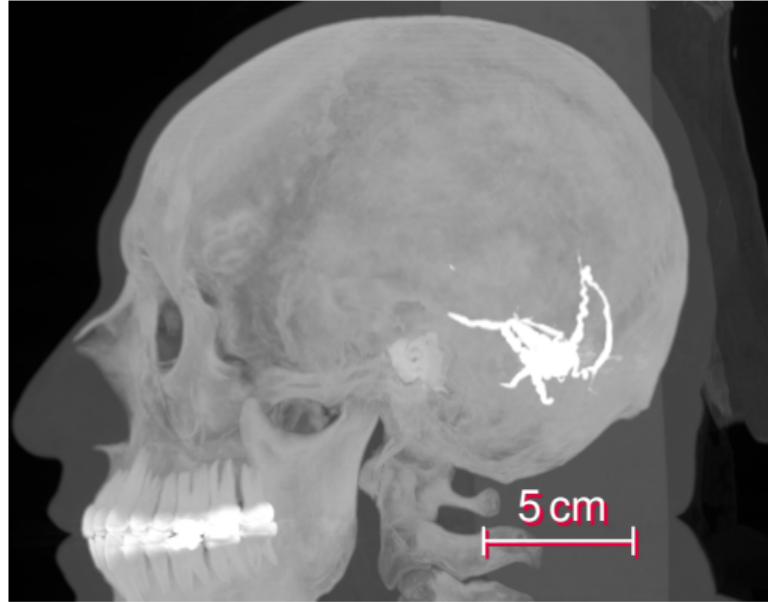


From [13], Subject C3L-02465

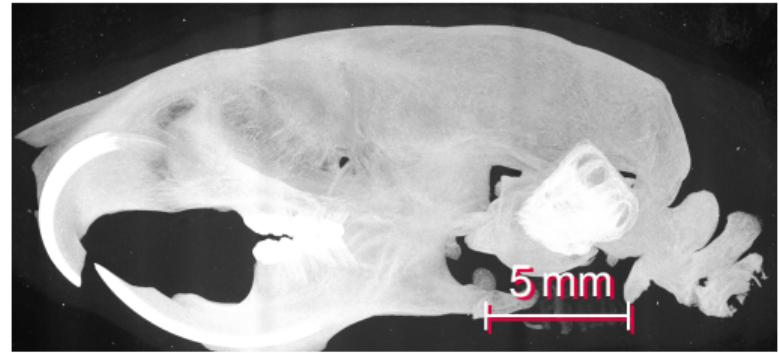


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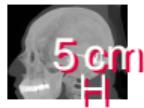


From [13], Subject C3L-02465

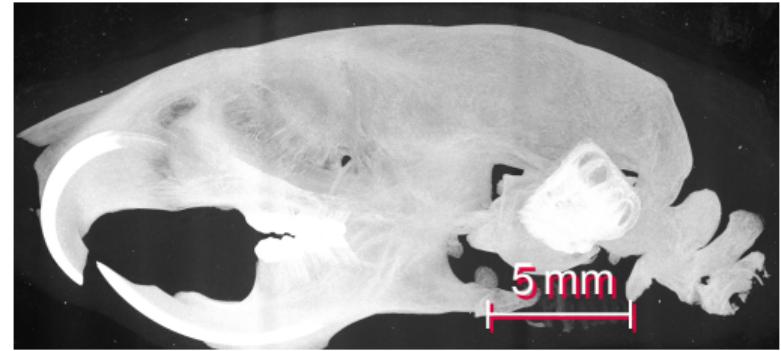


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# Why $\mu$ CT?



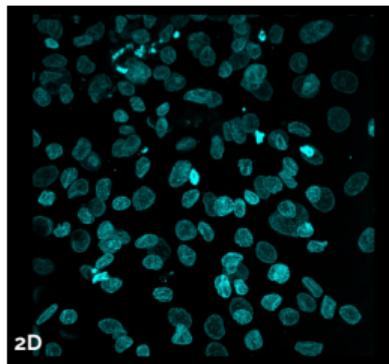
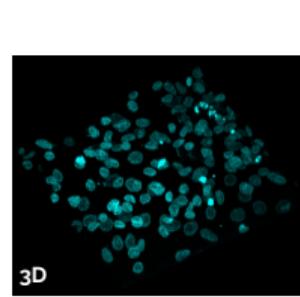
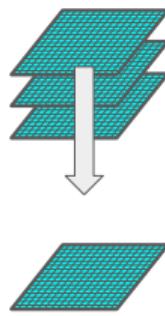
From [13], Subject C3L-02465



# Maximum intensity projection

## Projections

Reducing the dimensions of a dataset. For example projecting a volume (3D) to a surface by taking the maximum value across planes for each pixel.



*Fundamentals of Digital Image Processing*, Guillaume Witz, Slide 23

# Machinery

- Hospital CT
  - Voxel size around 0.5 mm
- Lab/Desktop CT
  - Voxel size around 7  $\mu\text{m}$  (*in vivo*)
  - Voxel size around 0.5  $\mu\text{m}$  (*ex vivo*)
- Synchrotron CT
  - Voxel size down to 160 nm



flic.kr/p/D4rbom

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[flic.kr/p/fpTrGu](https://flic.kr/p/fpTrGu)

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[bruker.com/skyscan1272](http://bruker.com/skyscan1272)

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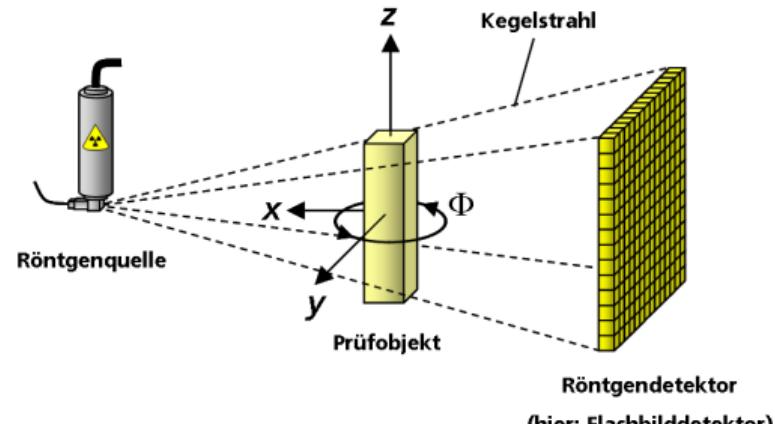
flic.kr/p/7Xhk2Y © ⓘ ⓘ

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# What is happening?

No matter what kind of machine, the basic principle is always

- an x-ray source
- a sample
- a detector



w.wiki/7g3 @①②

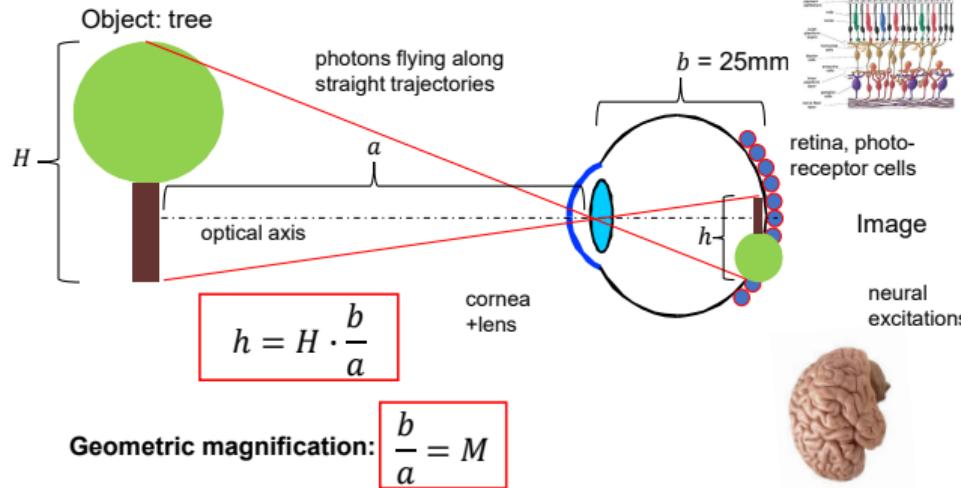
$u^b$

# Magnification

## Introduction – why do we need microscopes?

4

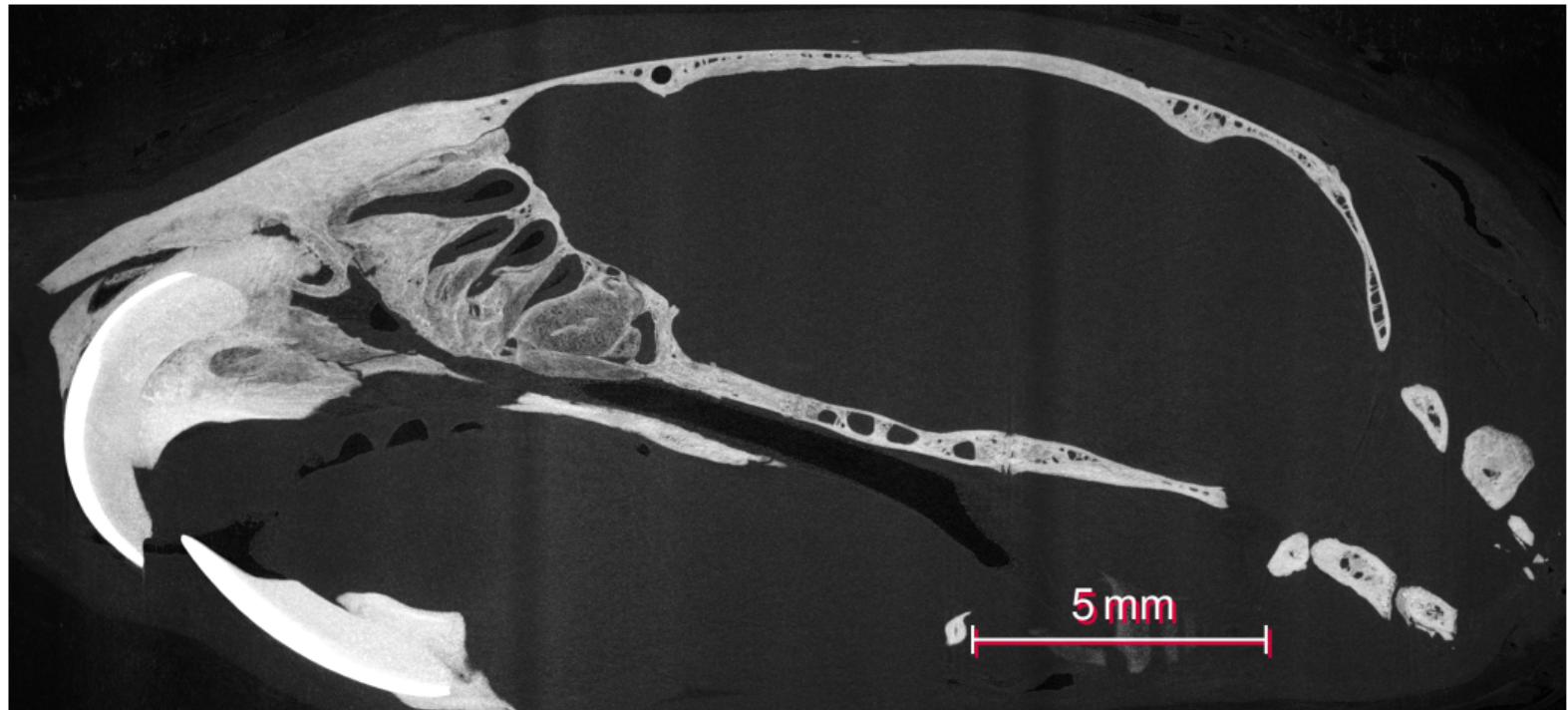
- Basic geometric optics: how does the eye form an image?



Contrast, Magnification and Resolution, Michael Jaeger, Slide 4

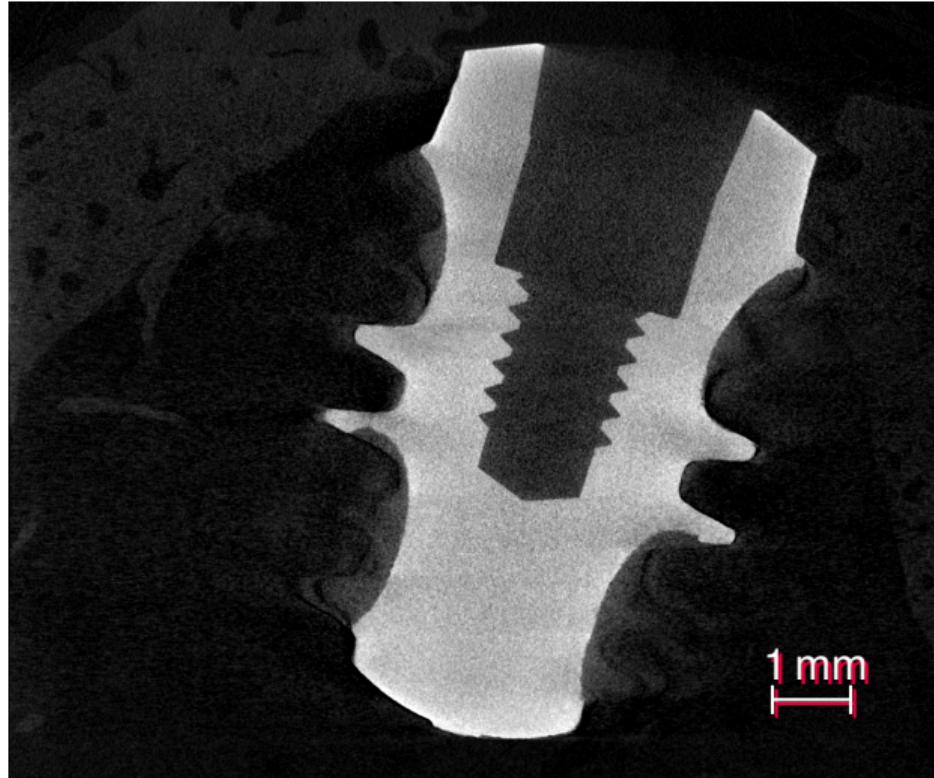
*u*<sup>b</sup>

# Examples



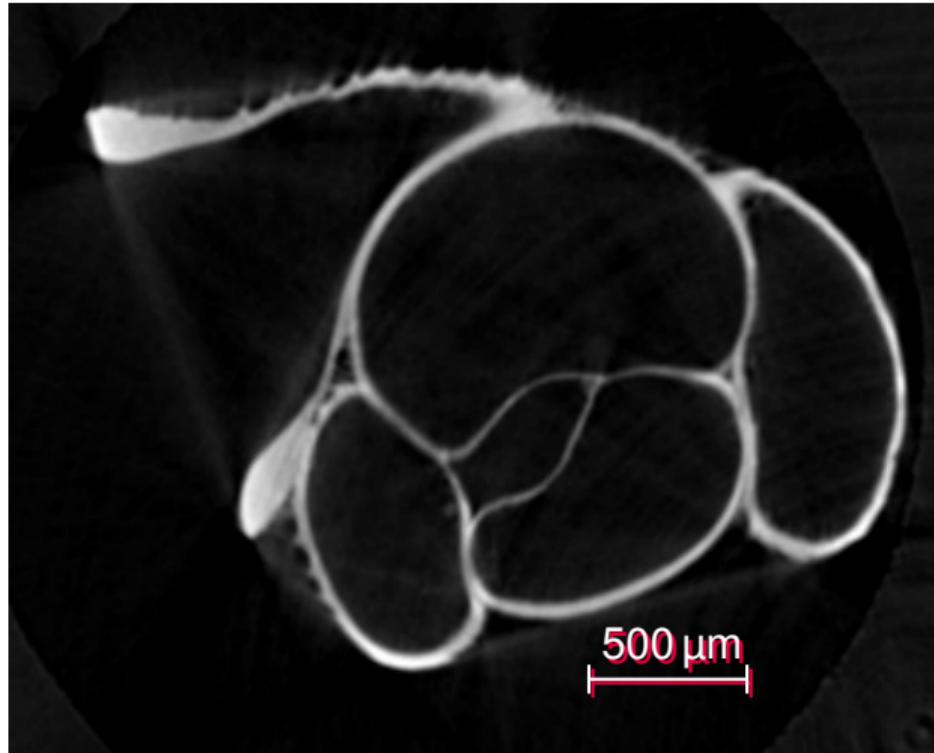
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# Examples



*u<sup>b</sup>*

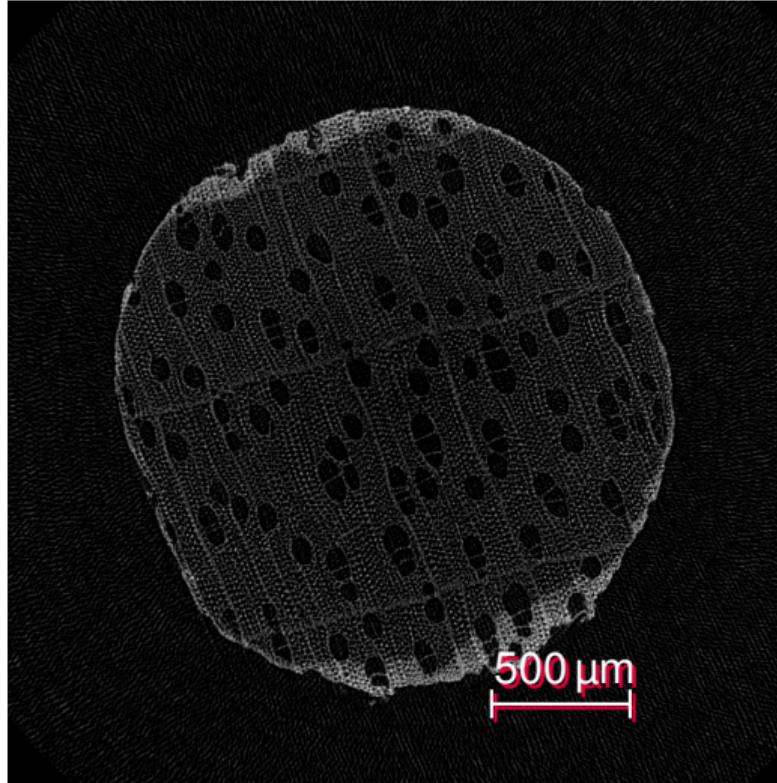
# Examples



From [8], *Diancta phoenix*

$u^b$

# Examples



*u*<sup>b</sup>

# Examples



# X-ray interaction

- “X-rays interact with tissue in 2 main ways: photoelectric effect and Compton scatter. To a first approximation, the photoelectric effect contributes to contrast while the Compton effect contributes to noise. Both contribute to dose.”  
**(xrayphysics)**
  - Photoelectric absorption ( $\tau$ ) is strongly dependent on the atomic number  $Z$  of the absorbing material:  $\tau \propto \frac{Z^4}{E^{3.5}}$
  - Compton scattering is one of the principle forms of photon interaction and is directly proportional to the (electron & physical) density of the material. It does *not* depend on the atomic number:  $\lambda' - \lambda = \frac{h}{m_e c} (1 - \cos \theta)$
- Lowering x-ray energy increases contrast
- X-ray penetration decreases exponentially with sample thickness **wiki:beer-lambert**:  $I(t) = I_0 e^{-\alpha z}$

# Composition of biological tissues

Tissue: content by mass percentage

| Element<br>Atomic number | H<br>1 | C<br>6 | N<br>7 | O<br>8 | Na<br>11 | P<br>15 | S<br>16 | Cl<br>17 | K<br>19 | Ca<br>20 |
|--------------------------|--------|--------|--------|--------|----------|---------|---------|----------|---------|----------|
| Fat                      | 11.4   | 59.8   | 0.7    | 27.8   | 0.1      |         | 0.1     | 0.1      |         |          |
| Water                    | 11.2   |        |        | 88.8   |          |         |         |          |         |          |
| Blood                    | 10.2   | 11     | 3.3    | 74.5   | 0.1      | 0.1     | 0.2     | 0.3      | 0.2     |          |
| Liver                    | 10.2   | 13.9   | 3      | 71.6   | 0.3      | 0.2     | 0.3     | 0.2      | 0.3     |          |
| Brain                    | 10.7   | 14.5   | 2.2    | 71.2   | 0.2      | 0.4     | 0.2     | 0.3      | 0.3     |          |
| Bone                     | 3.4    | 15.5   | 4.2    | 43.5   | 0.1      | 10.3    | 0.3     |          |         | 22.5     |

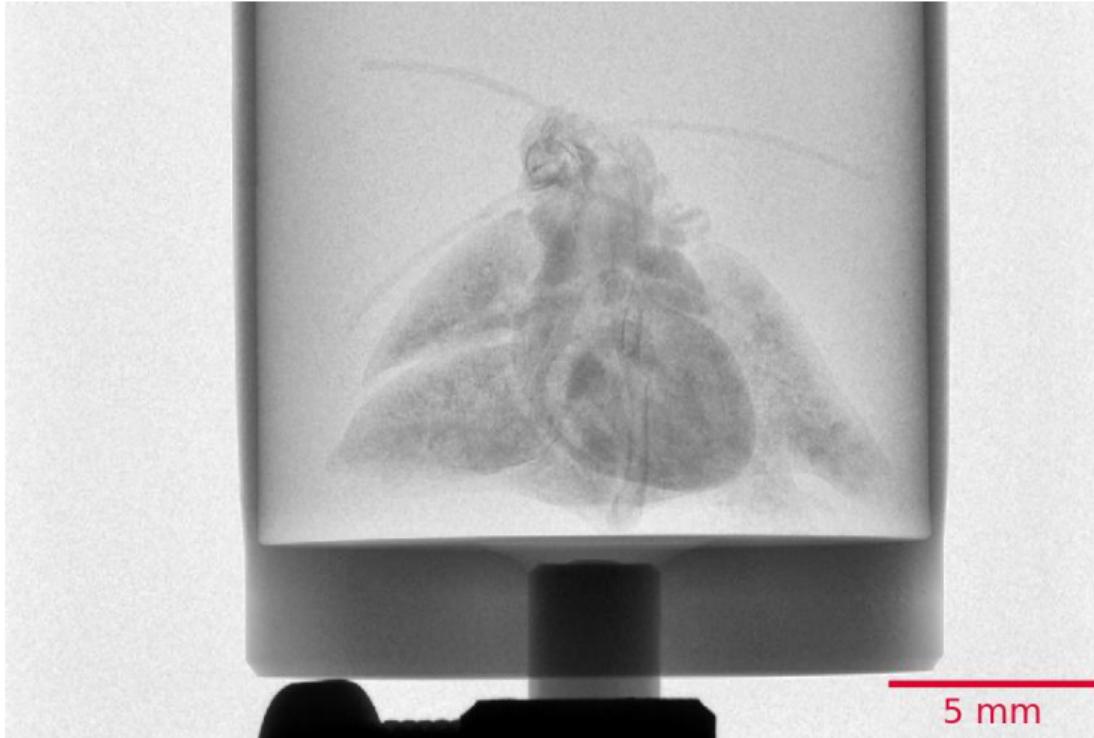
*u<sup>b</sup>*

# Preparation

- Study design
- Sample preparation

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# Projections



# Projections

- A (micro-focus) x-ray source illuminates the object
- The x-rays penetrate the sample and are attenuated
- A scintillator converts the x-rays to visible light
- A (planar) x-ray detector collects (magnified) projection images.
- The projections are recorded on disk

*u*<sup>b</sup>

# Reconstructions



# Reconstructions

- Based on hundreds of angular views acquired while the object rotates, a computer synthesizes a stack of virtual cross section slices through the object.
- Radon Transformation
- Filtered back projection
- Cone beam reconstruction [19]
- Corrections (beam hardening, etc.)
- Writing to stack

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# Visualization



# Visualization

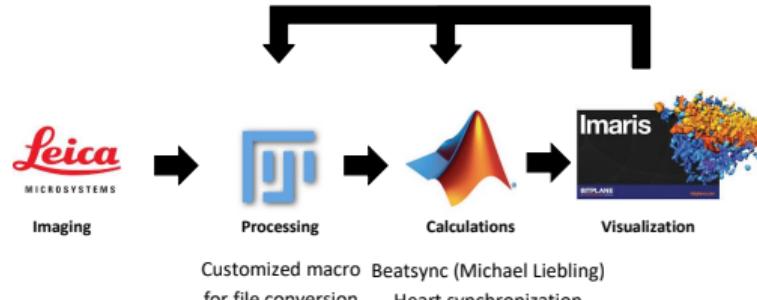
- Based on the calculated reconstructions, a computer synthesizes a (three-dimensional) visualization of the scanned sample

*u*<sup>b</sup>

# Imaging

CEM 2024-Light Sheet Microscopy

How to deal with large multidimensional data?



CEM 2024-Light Sheet Microscopy

*Light Sheet Microscopy*, Nadia Mercader, Slide 43

*u*<sup>b</sup>

# What to use?

- ImageJ/Fiji [21]
- Also see *Fundamentals of Digital Image Processing* by Guillaume Witz
- Reproducible research
  -  in Jupyter [22]
  - **git**
  - Script all your things!
  - Data repositories; i. e. sharing is caring!

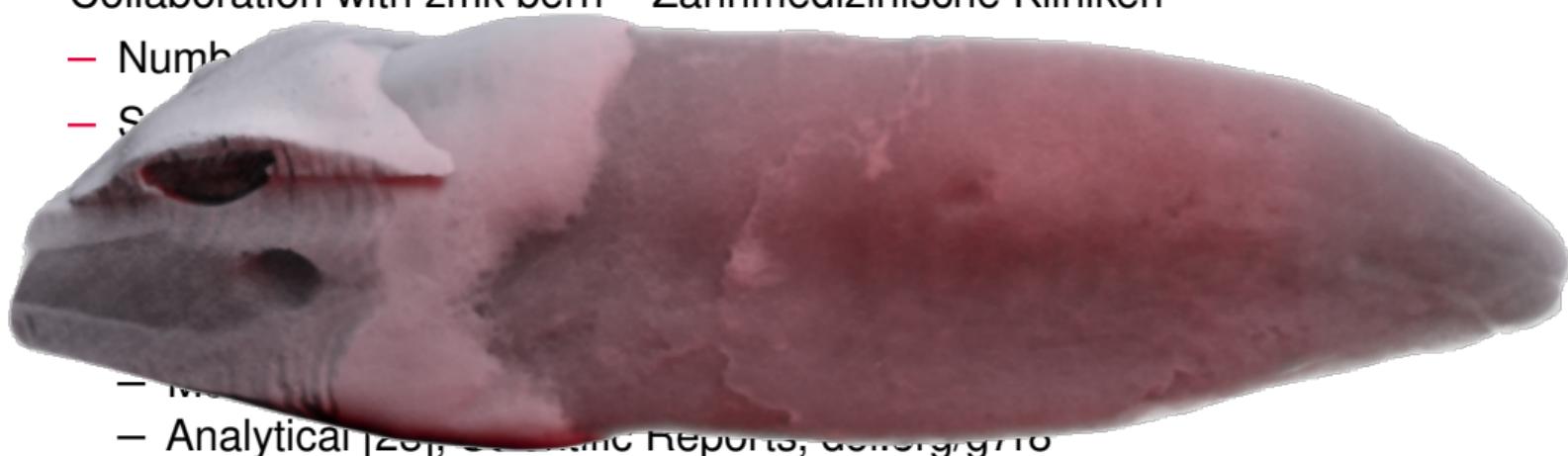
# Quantitative data

- Pretty images are nice, but we need quantitative numbers
- Segmentation
- Characterization

# Internal morphology of human teeth

Collaboration with zmk bern – Zahnmedizinische Kliniken

- Number of teeth
- Structure of teeth



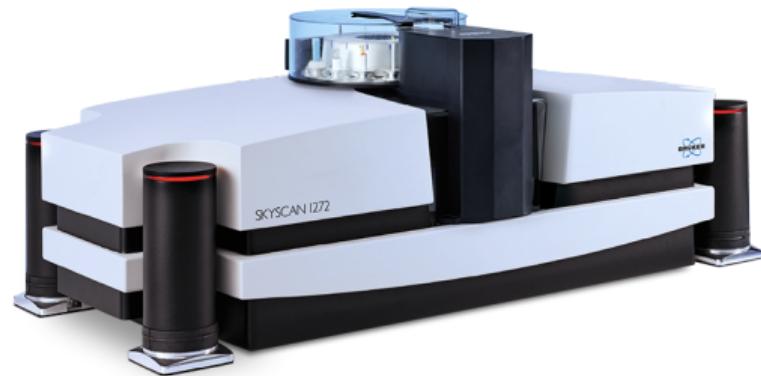
- Morphology [20], [Scientific Reports, doi.org/9710](https://www.ncbi.nlm.nih.gov/pmc/articles/PMC9710)
- Analytical [20], [bioRxiv preprint doi: https://doi.org/10.1101/2023.09.21.554710](https://doi.org/10.1101/2023.09.21.554710)

# How?

- 104 extracted human permanent mandibular canines
- $\mu$ CT imaging
- Root canal configuration, according to Briseño-Marroquín *et al.* [24]
- *Reproducible* analysis [25], e. g. you can click a button to double-check or recalculate the results yourself!



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[bruker.com/skyscan1272](http://bruker.com/skyscan1272)

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```
Scanner=SkyScan1272
Instrument S/N=15G09089-B
Software Version=1.1.19
Filename Prefix=Tooth045~00
Number Of Files= 482
Number Of Rows= 1092
Number Of Columns= 1632
Source Voltage (kV)= 80
Source Current (uA)= 125
Image Pixel Size (um)=9.999986
Exposure (ms)=950
Rotation Step (deg)=0.400
Frame Averaging=ON (3)
Filter=Al 1mm
Study Date and Time=02 Jul 2020 08h:23m:34s
Scan duration=0h:39m:51s
```

- 104 extracted human permanent mandibular canines
- $\mu$ CT imaging
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- *Reproducible* analysis [25], e. g. you can click a button to double-check or recalculate the results yourself!

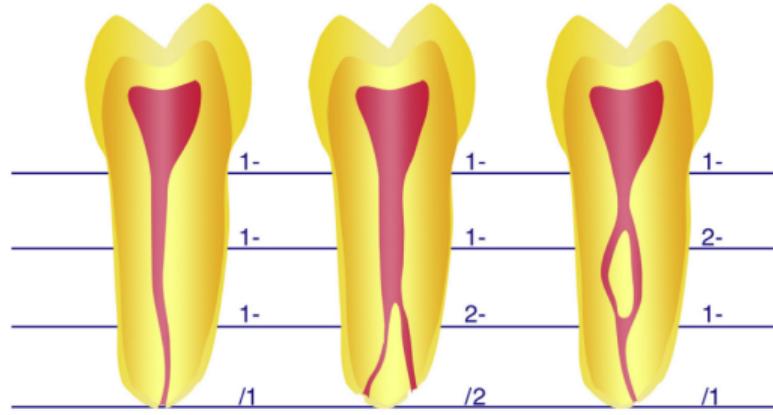
*Sample changer* on the SkyScan 1272

In total:

- 13 days of *continuous*  $\mu$ CT scanning
- 819 GB of raw data
- 230 648 TIFF projections
- 326 GB data as input for analysis
- 282 062 PNG reconstructions

# How?

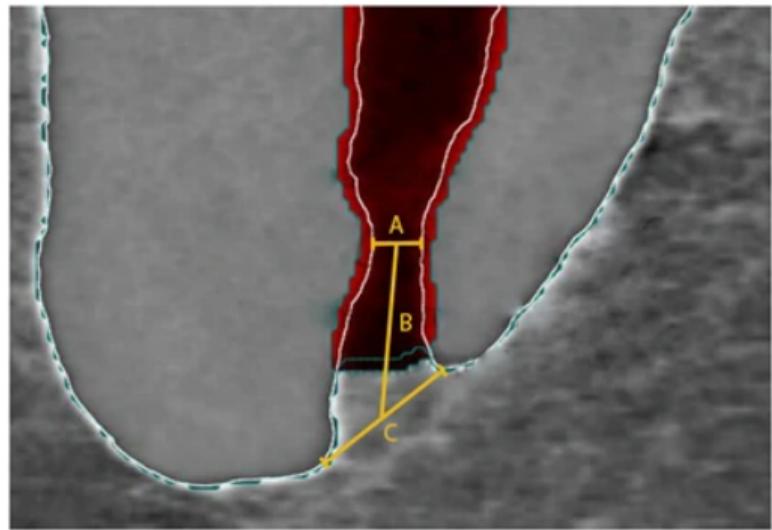
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From [24], Fig. 2

# How?

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From [26], Fig. 1

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[gph.is/2nqkple](https://gph.is/2nqkple)

*u*<sup>b</sup>

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The screenshot shows a GitHub repository interface. At the top, there are buttons for 'master' (with a dropdown arrow), '1 branch' (with a dropdown arrow), '1 tag' (with a dropdown arrow), 'Go to file', 'Add file', and 'Code'. Below this is a list of commits from a user named 'habi'. The commits are:

| File                       | Description                                        | Date         |
|----------------------------|----------------------------------------------------|--------------|
| .github/workflows          | Update actions file                                | 20 days ago  |
| .gitignore                 | Only 'mode' changes                                | 2 months ago |
| DownloadFromOSF.ipynb      | Clean run of download script                       | 22 days ago  |
| README.md                  | Typo in Binder badge & link to full repo on Binder | 22 days ago  |
| Tooth.Border.jpg           | Only 'mode' changes                                | 2 months ago |
| Tooth.Characterization.jpg | Only 'mode' changes                                | 2 months ago |
| ToothAnalysis.ipynb        | Only select a subset if we actually have data:wq   | 22 days ago  |
| ToothAxisSize.ipynb        | Clean run of notebook                              | 22 days ago  |
| ToothDisplay.ipynb         | Display Tooth045 for manuscript                    | 22 days ago  |
| requirements.txt           | We also need this                                  | 2 months ago |
| treebeard.yaml             | Add treebeard configuration                        | 20 days ago  |

Below the commits is a preview of the 'README.md' file. It contains:

```
README.md
DOI: 10.5281/zenodo.3999402 treebeard.yml failing
launch binder
```

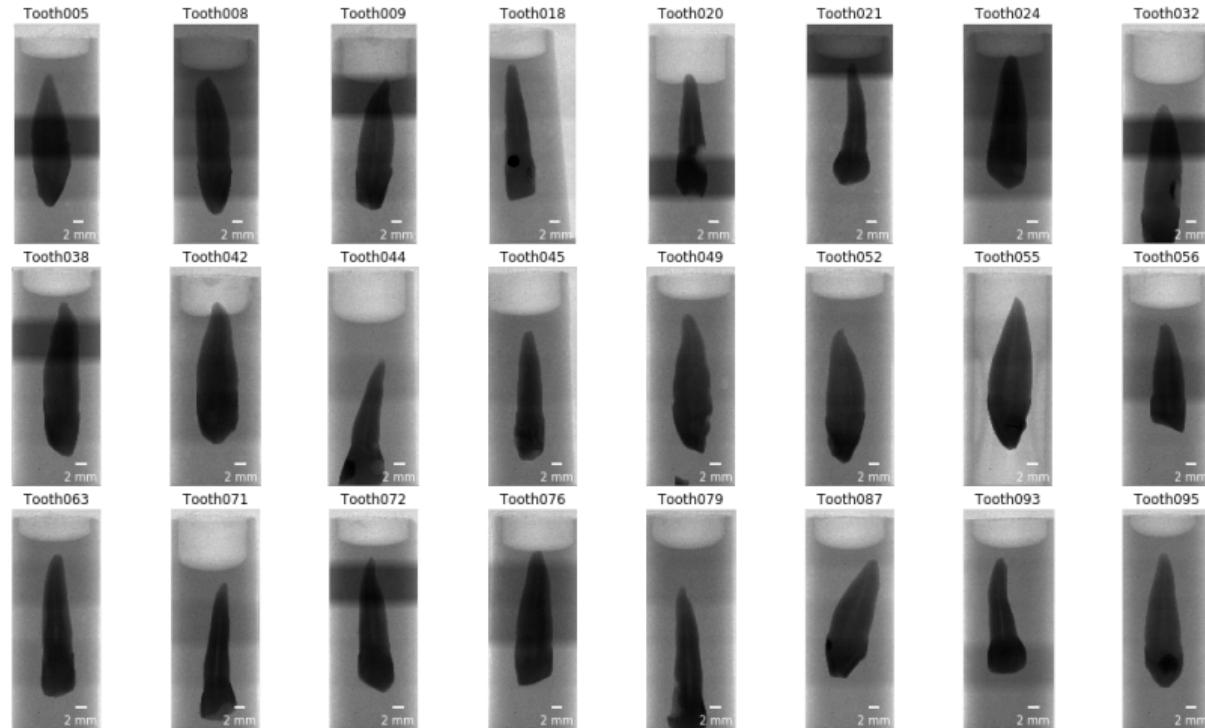
**A big tooth cohort**

We scanned a *big* bunch of teeth for a team of the dental clinic of the University of Bern.

To get an overview of the samples while we scanned the whole tooth cohort we generated a [preview](#) and [analysis notebook](#). The analysis notebook (with download possibility for two of the +100 teeth) can be started in your browser by clicking on the 'Binder' badge above, without installation of any software. If you'd like to start a Binder instance with the full repository, you can click [here](#).

$\mu$ b

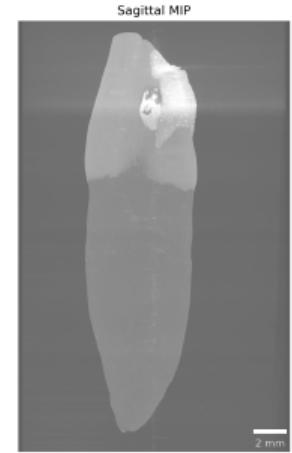
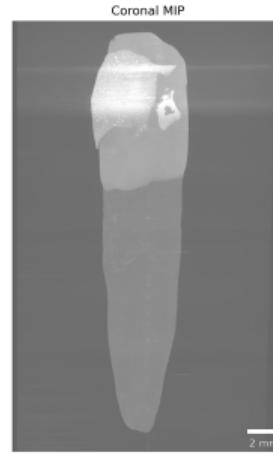
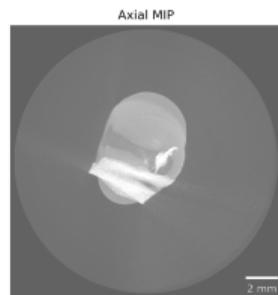
# $\mu$ CT imaging



*u*<sup>b</sup>

# Dataset cropping

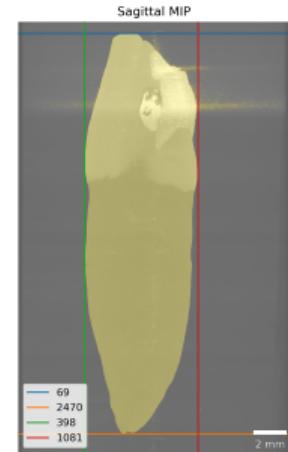
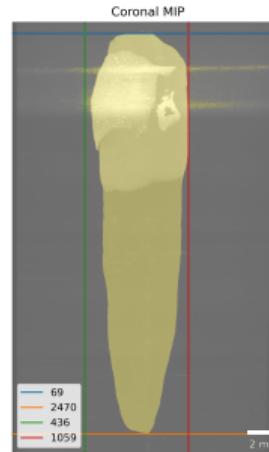
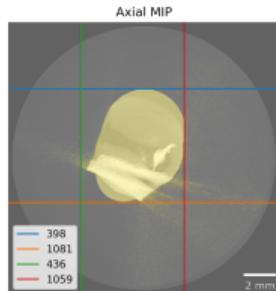
- Full datasets: 326 GB
- Cropped datasets: 115 GB



*u*<sup>b</sup>

# Dataset cropping

- Full datasets: 326 GB
- Cropped datasets: 115 GB



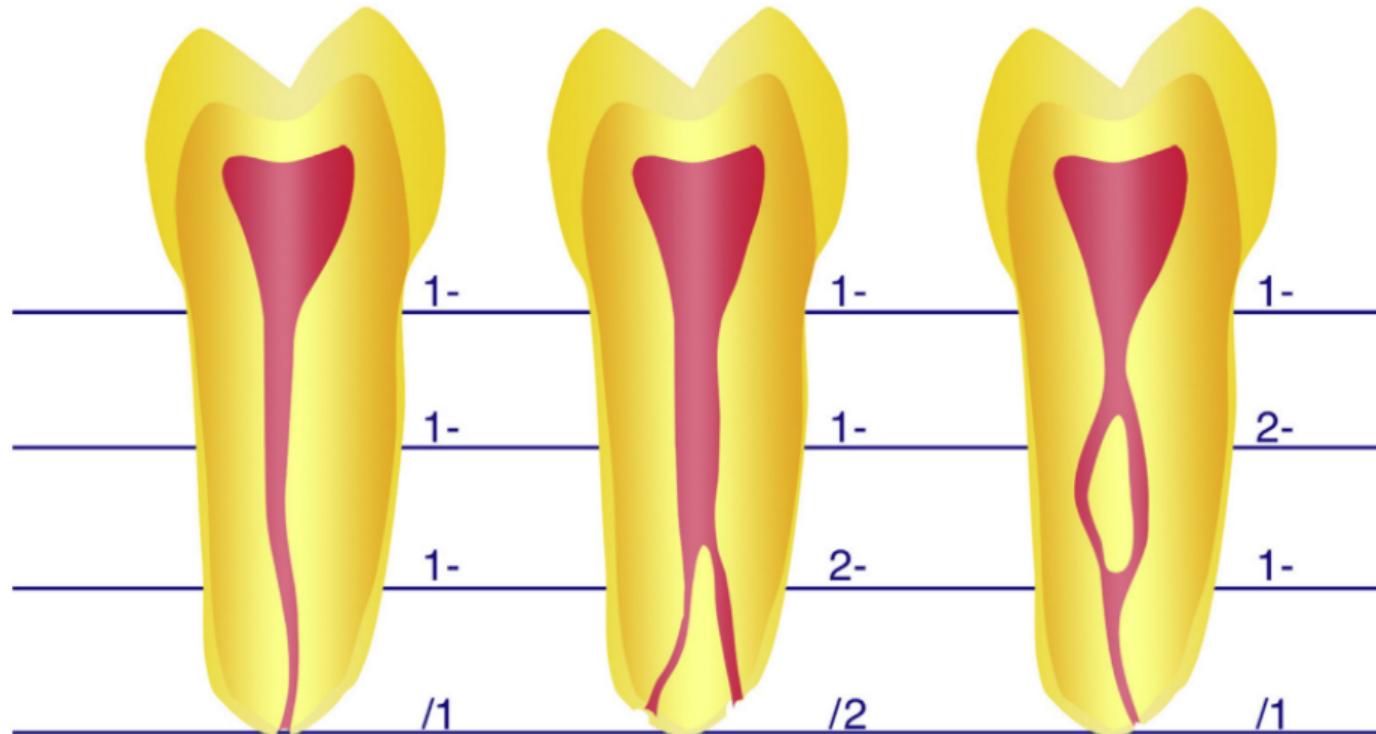
*u<sup>b</sup>*

# Tooth morphology



*u<sup>b</sup>*

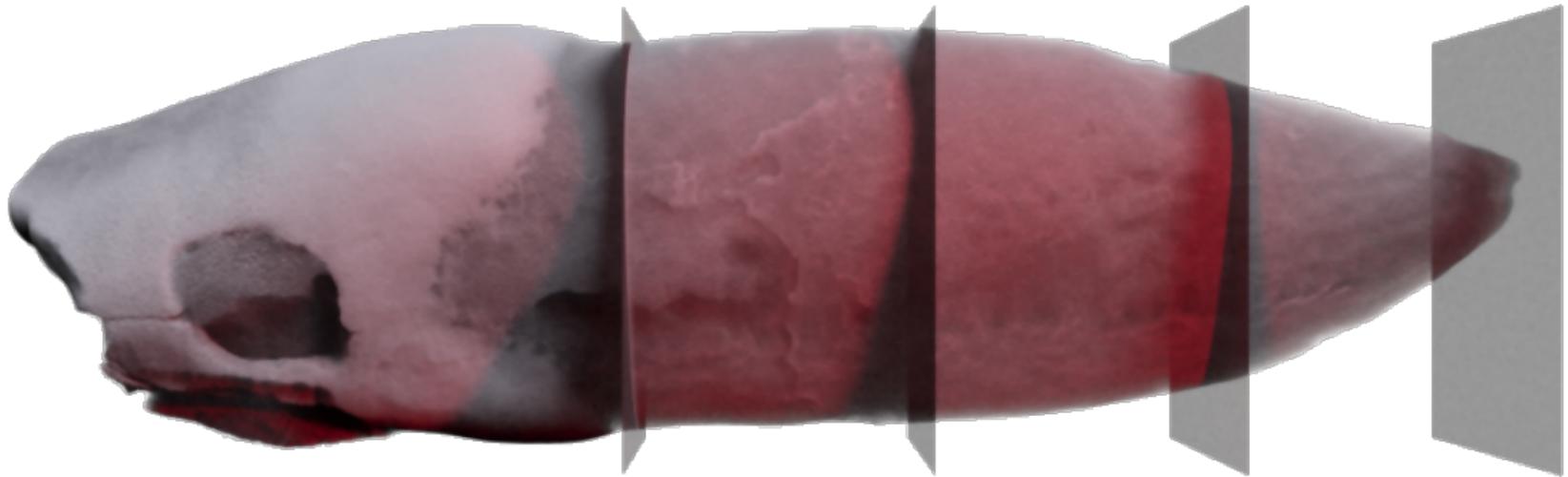
# Tooth morphology



From [24], Fig. 2

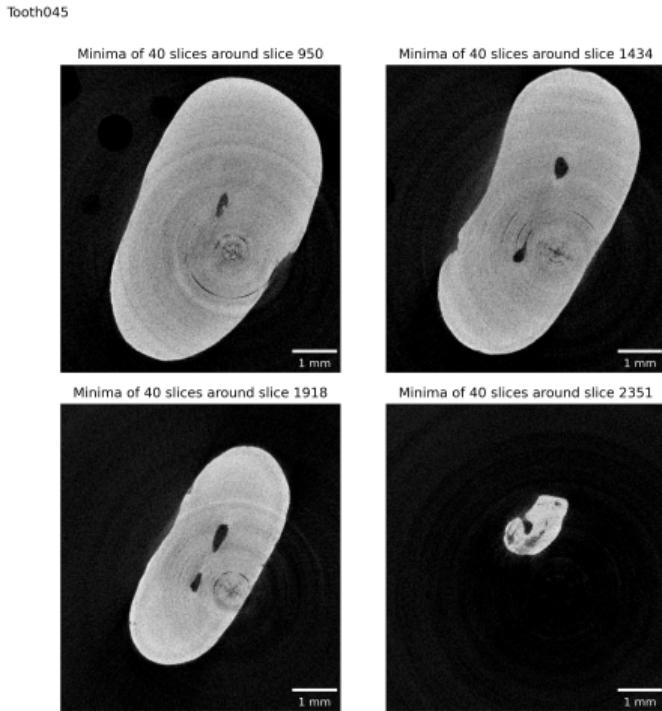
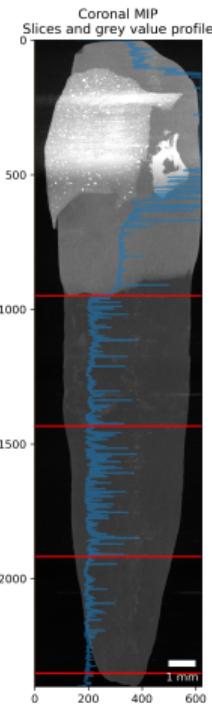
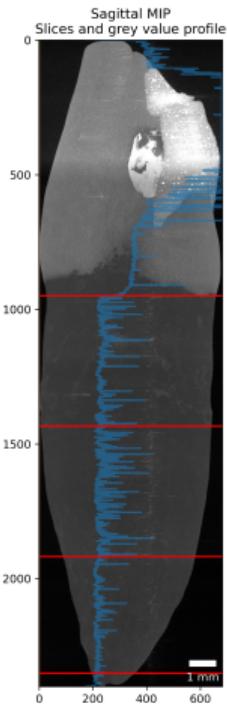
*u<sup>b</sup>*

# Tooth morphology



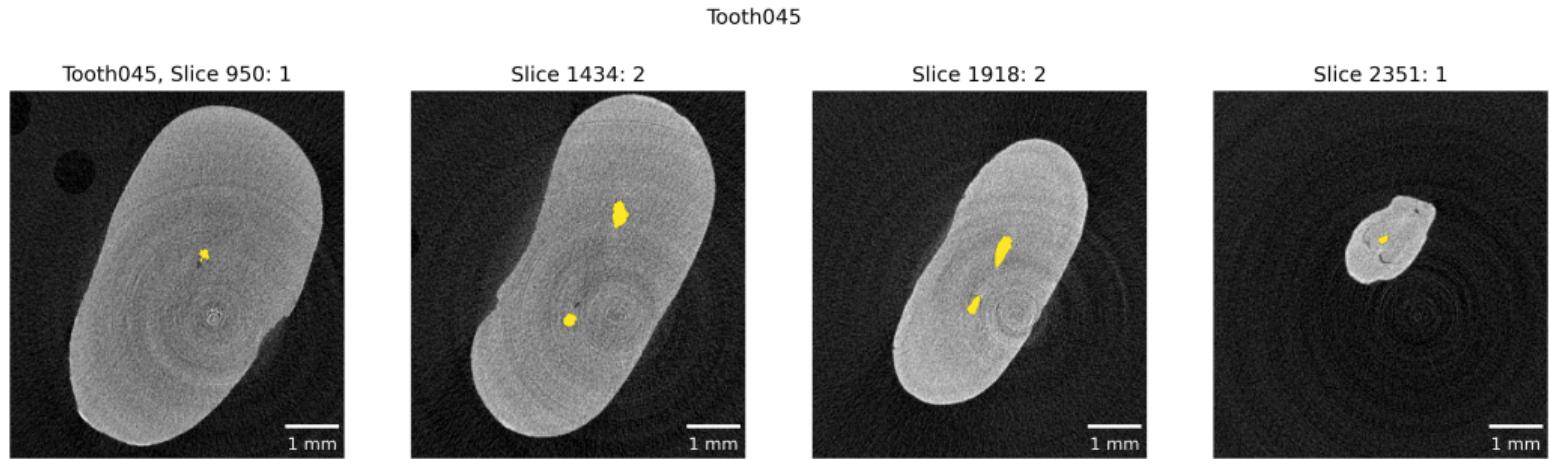
*u<sup>b</sup>*

# Detection of enamel-dentin border



*u*<sup>b</sup>

# Detection of enamel-dentin border



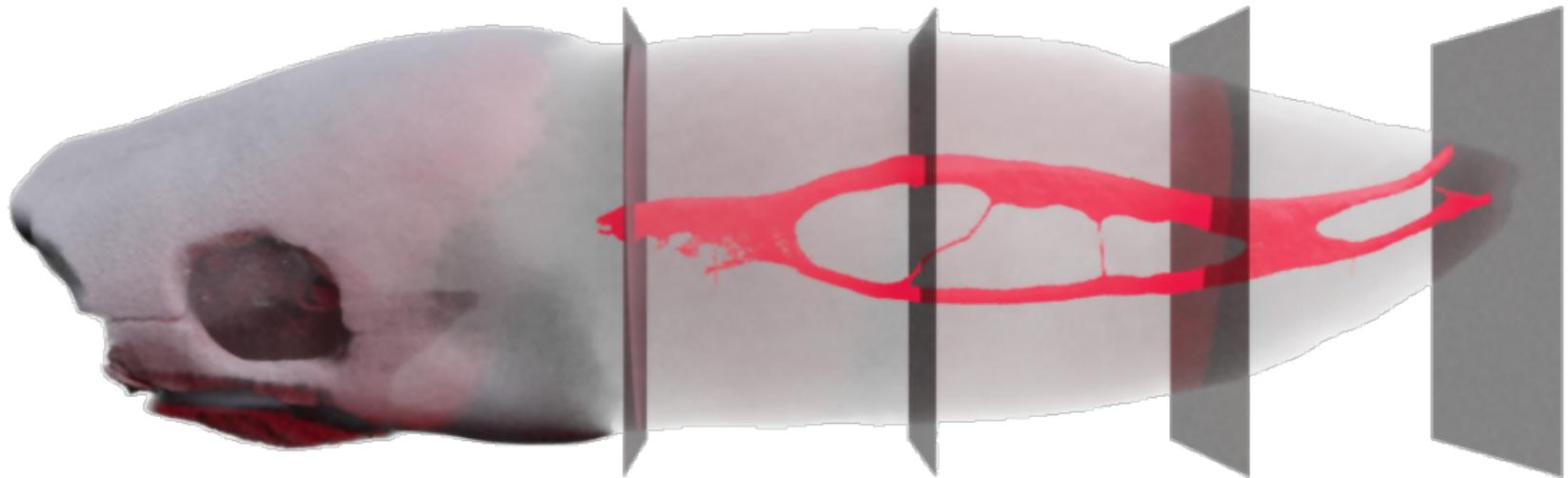
*u<sup>b</sup>*

# Outcome root canal configuration classification

| Roots         | RCC     | #       | %    |
|---------------|---------|---------|------|
| Single (N=98) | 1-1-1/1 | 73      | 74.5 |
|               | 1-1-1/2 | 14      | 14.3 |
|               | 1-1-1/3 | 1       | 1.0  |
|               | 1-1-1/4 | 2       | 2.1  |
|               | 1-1-2/1 | 1       | 1.0  |
|               | 1-2-1/1 | 4       | 4.1  |
|               | 1-2-1/2 | 1       | 1.0  |
|               | 1-2-2/2 | 1       | 1.0  |
|               | 2-3-1/1 | 1       | 1.0  |
|               |         |         |      |
| Double (N=3)  | Buccal  | 1-1-1/1 | 2    |
|               |         | 1-2-1/1 | 1    |
|               | Lingual | 1-1-1/1 | 2    |
|               |         | 1-1-1/2 | 1    |

*u*<sup>b</sup>

# Extraction of root canal space



# Conclusion ZMK

- Efficient use of time, e. g. more teeth does not mean more (human) work
- Reproducible analysis with *free and open-source* software, usable by *anyone*
- Objective analysis, e. g. no operator bias

*u*<sup>b</sup>

# Thanks!

- Thanks for listening to me!
- What questions do you have for me?

# Colophon

- This BEAMER presentation was crafted in  $\text{\LaTeX}$  with the (slightly adapted) template from *Corporate Design und Vorlagen* of the University of Bern.
  - Complete source code: [git.io/fjpP7](https://git.io/fjpP7)
  - The  $\text{\LaTeX}$  code is automatically compiled with a GitHub action to a (handout) PDF which you can access here: [git.io/JeQxO](https://git.io/JeQxO)
- Did you spot an error?
  - File an issue: [git.io/fjpPb](https://git.io/fjpPb)
  - Submit a pull request: [git.io/fjpPN](https://git.io/fjpPN)
  - Send me an email: [david.haberthuer@unibe.ch](mailto:david.haberthuer@unibe.ch)