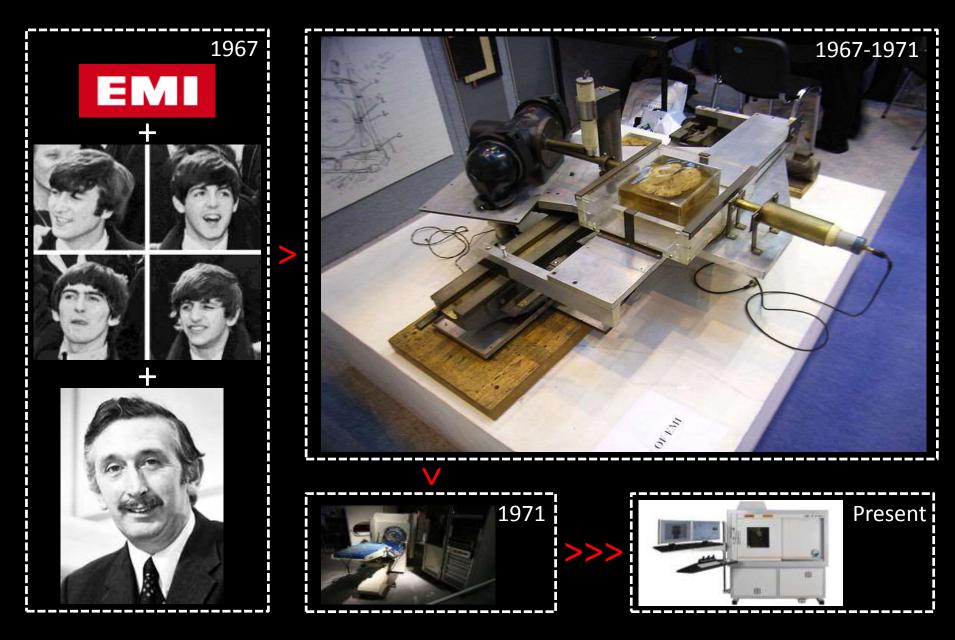
Imperial College London

Royal School of Mines Earth Science & Engineering Imperial College London

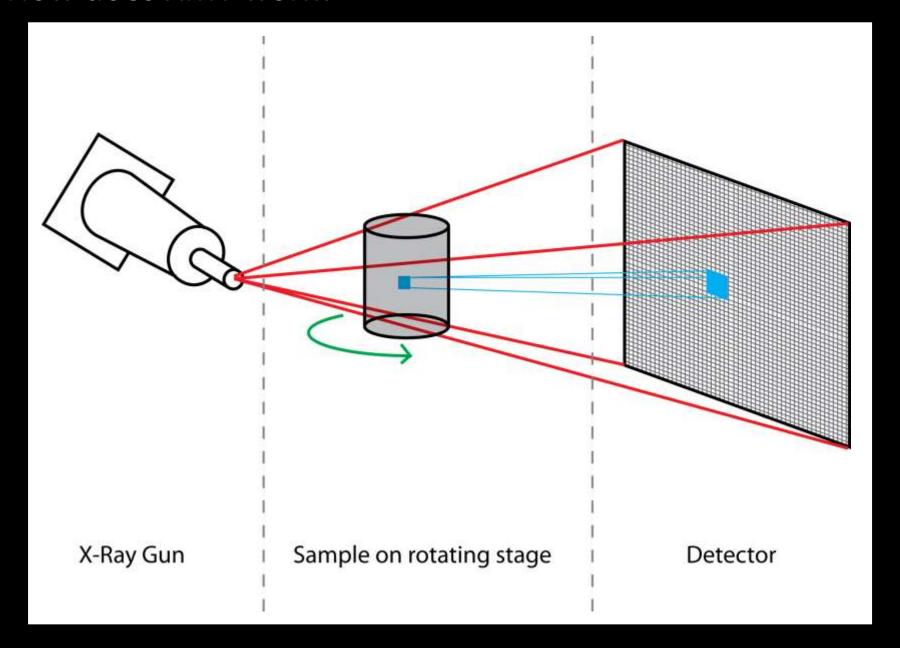
3D X-RAY MICROTOMOGRAPHY RECONSTRUCTIONS: AN IMPORTANT PRELIMINARY TOOL FOR PALAEOBOTANISTS

Alan R.T. Spencer & Mark D. Sutton

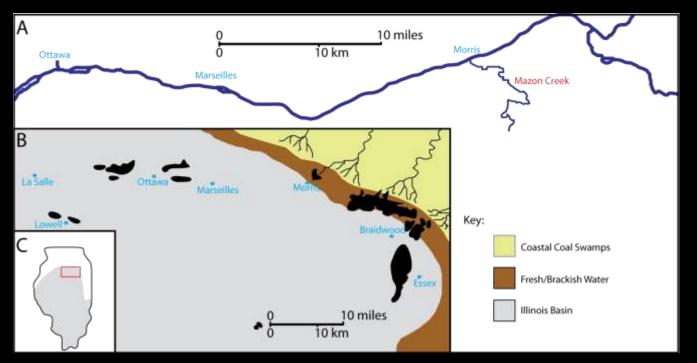
CT/CAT Scanning History (in brief):



How does XMT work?



Specimen P30420





Location: Francis Creek Shale, Mazon Creek, II, USA

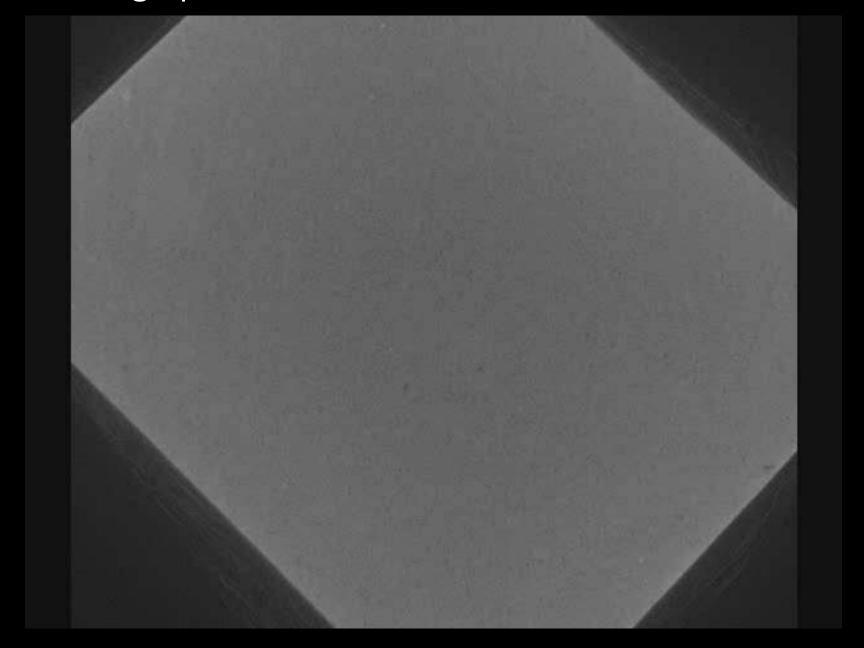
Age: Carboniferous, Pennsylvanian (318-299 Ma)

Preservation: Siderite Nodule (FeCO₃)

Owner: The Field Museum, Chicago, USA

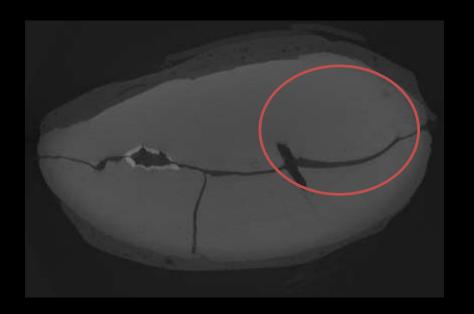
Description: Stephanospermum Seed

2D Tomographic Data Set:

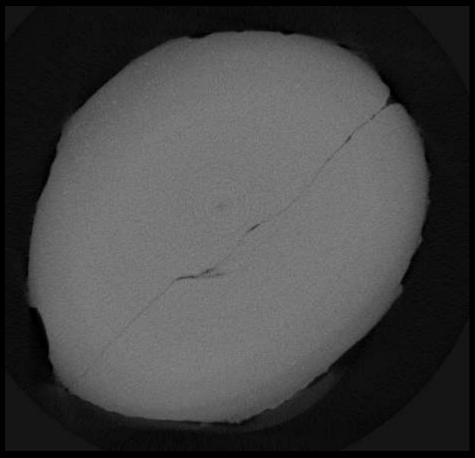


Problems with XMT Data collection:

Low phase contrast



<u>Artefacts</u>



Data Set Size (storage)

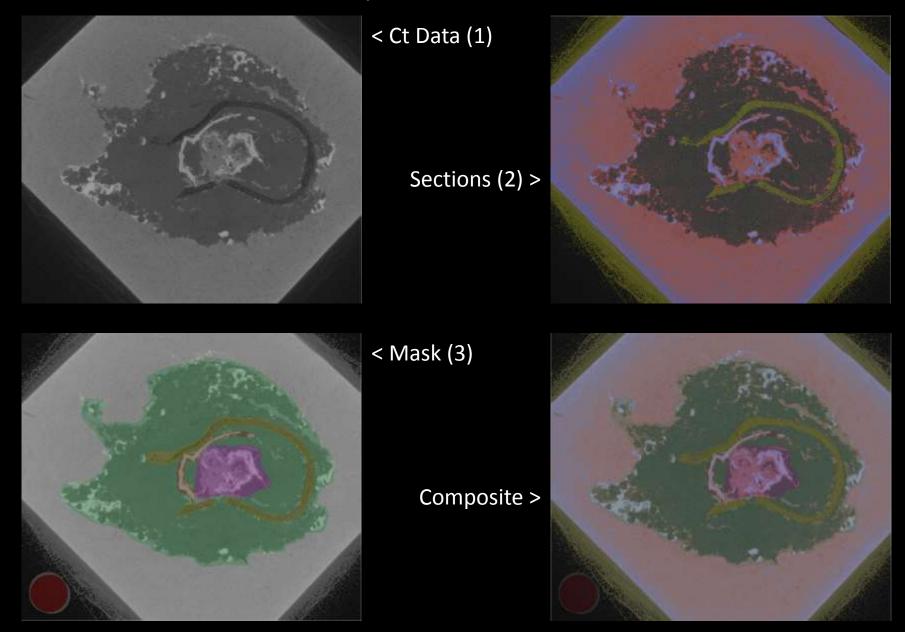
Direct from Scanner = 2.02 GB

After conversion to BMP = 1.01 GB

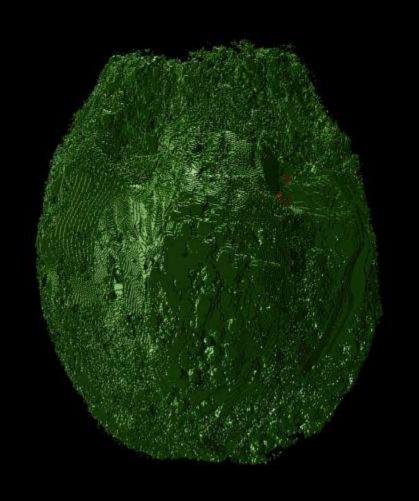
After initial conversion for Modelling = 6.10 GB (depending on complexity)

TOTAL: = 9.11 GB (before anything useable is outputted)

From 2D to 3D: Sections, Masks & Models

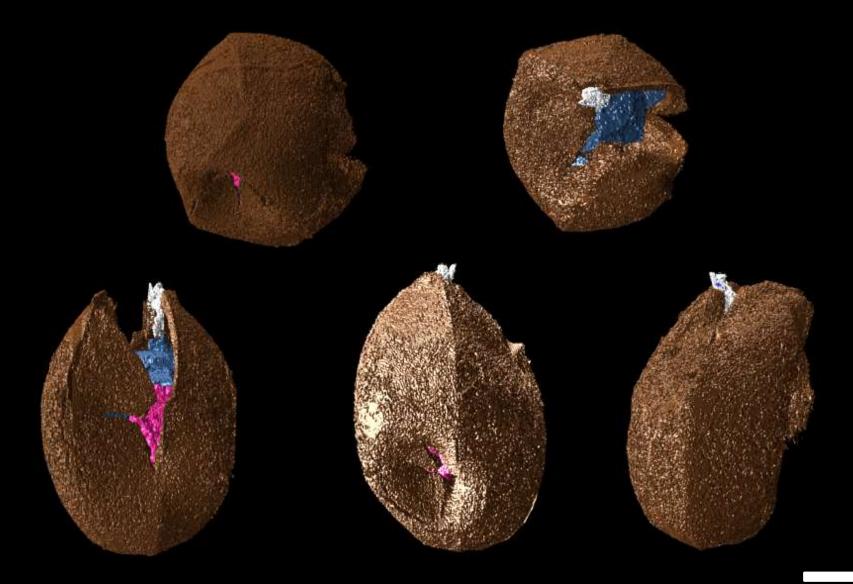


3D Model:

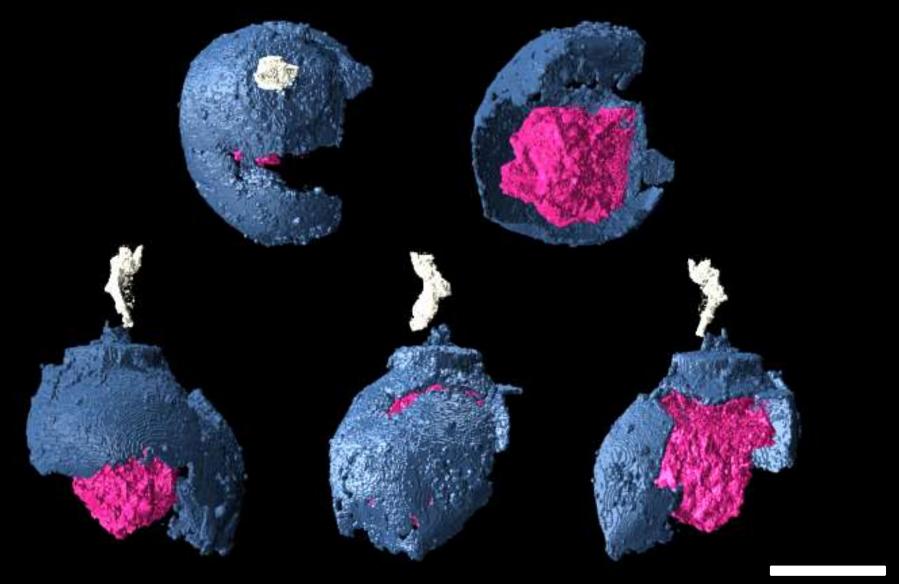


Sarcotesta:

Sclerotesta:



Nucellus, Micropyle, Pollen Chamber, & Megaspore:



Real World vs Virtual:



Conclusion:

<u>PROS</u>

- •the ability to virtually dissect fossils
- recovery of full morphological data
- all within a non-destructive environment
- models can be used to guide the application of traditional destructive techniques

CONS

- resolution required to render detail, for example the nucellus membrane, is compromised by the need to reconstruct an entire fossil
- only fossil material that has a high enough phase contrast under x-ray can produce a scan of sufficient quality for successful 3D reconstruction
- time required to produce detailed model

The combination of XMT with traditional techniques provides a powerful approach to the study of three-dimensionally preserved palaeobotanical specimens by allowing the traditional approaches to perform to their maximal potential.

The End

(...runs away to hide from zombies)