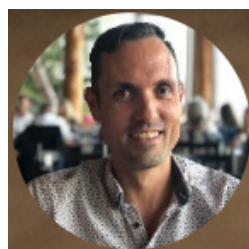
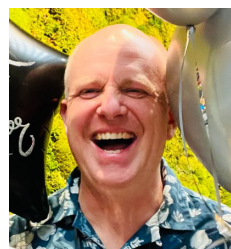


Species Classification Automation for Microfossil Photomicrograph Images (Scampi)

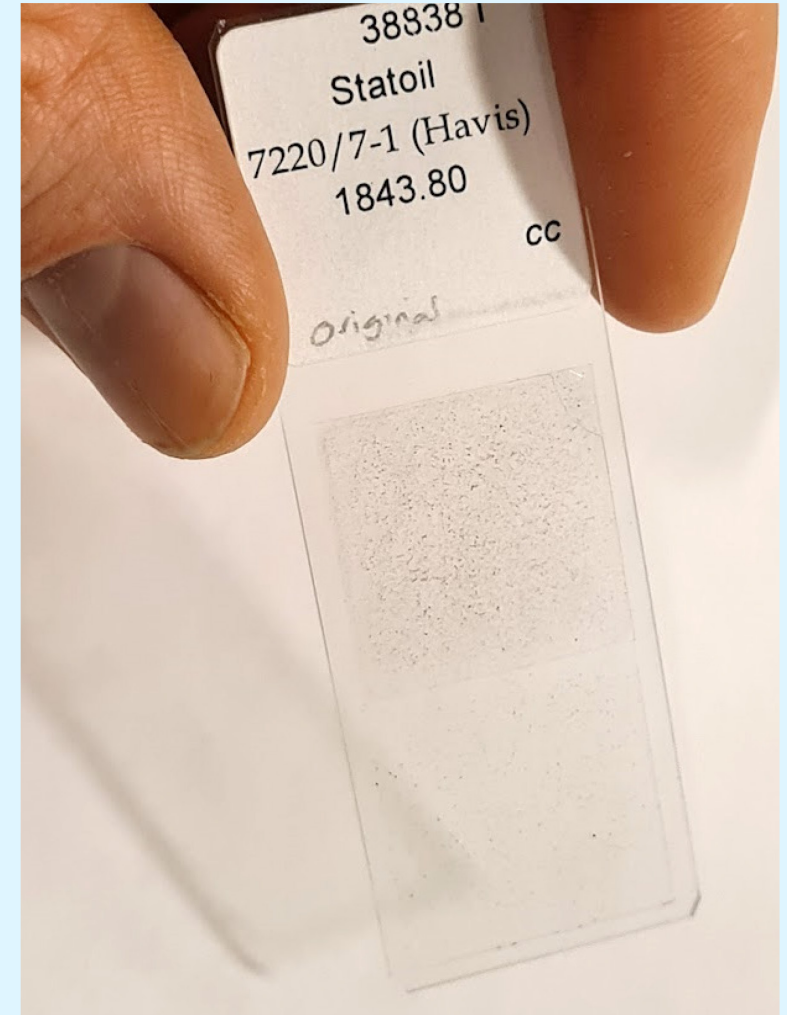
David Wade*, Sissa Stefanowicz, Alex Cullum & Erik Anthonissen



Palynology Slide Analysis

Traditional approach:

- Long hours with a microscope
- Make species counts
- Few practitioners



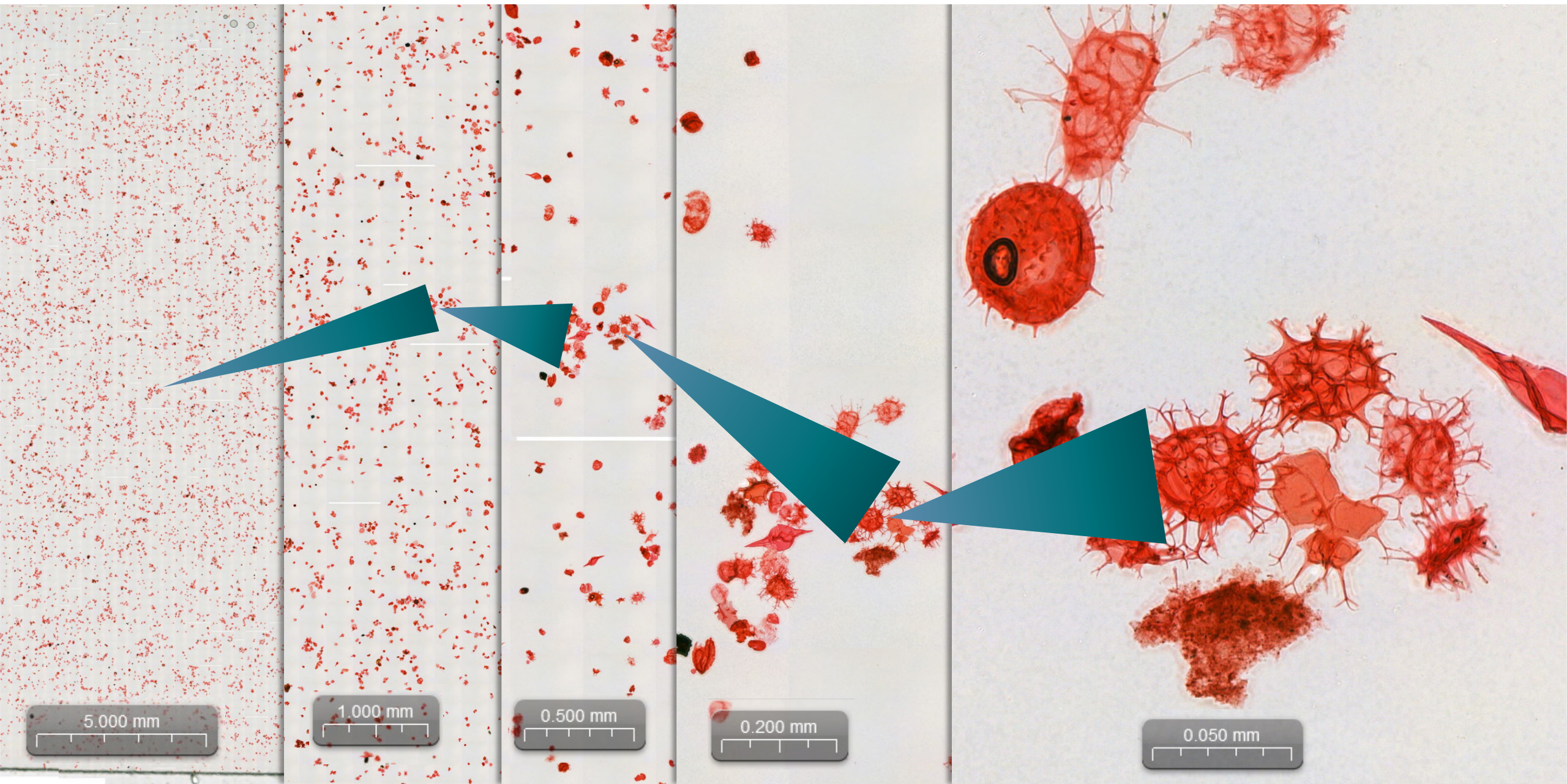


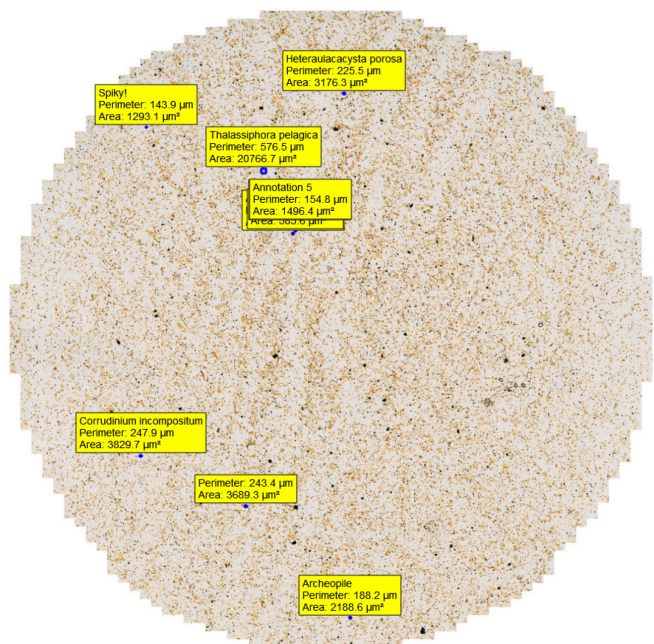
NORWEGIAN OFFSHORE
DIRECTORATE

Zoom : Whole slide

... to ...

Individual fossils





6
Gigapixels

=

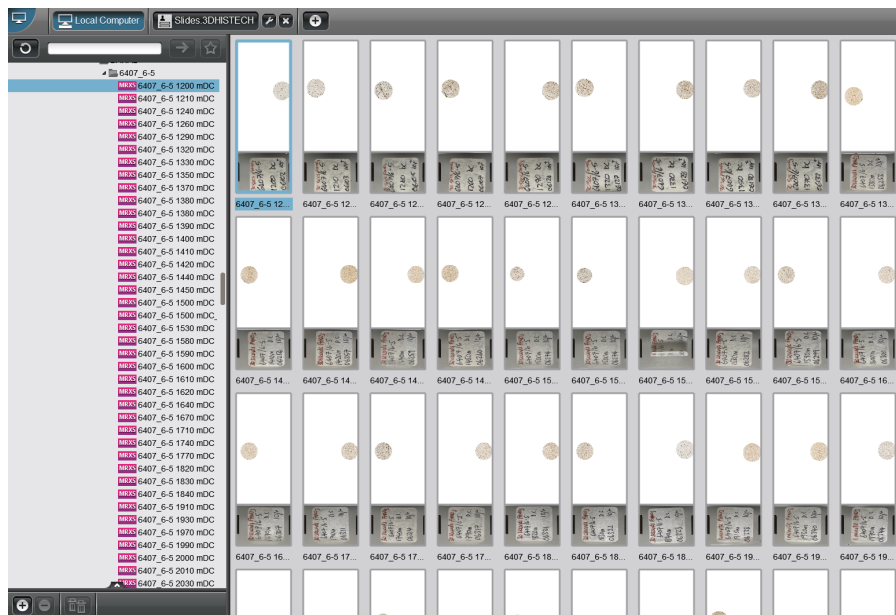
~25,000
Palynomorphs

x80
slides per well

x1800
wells

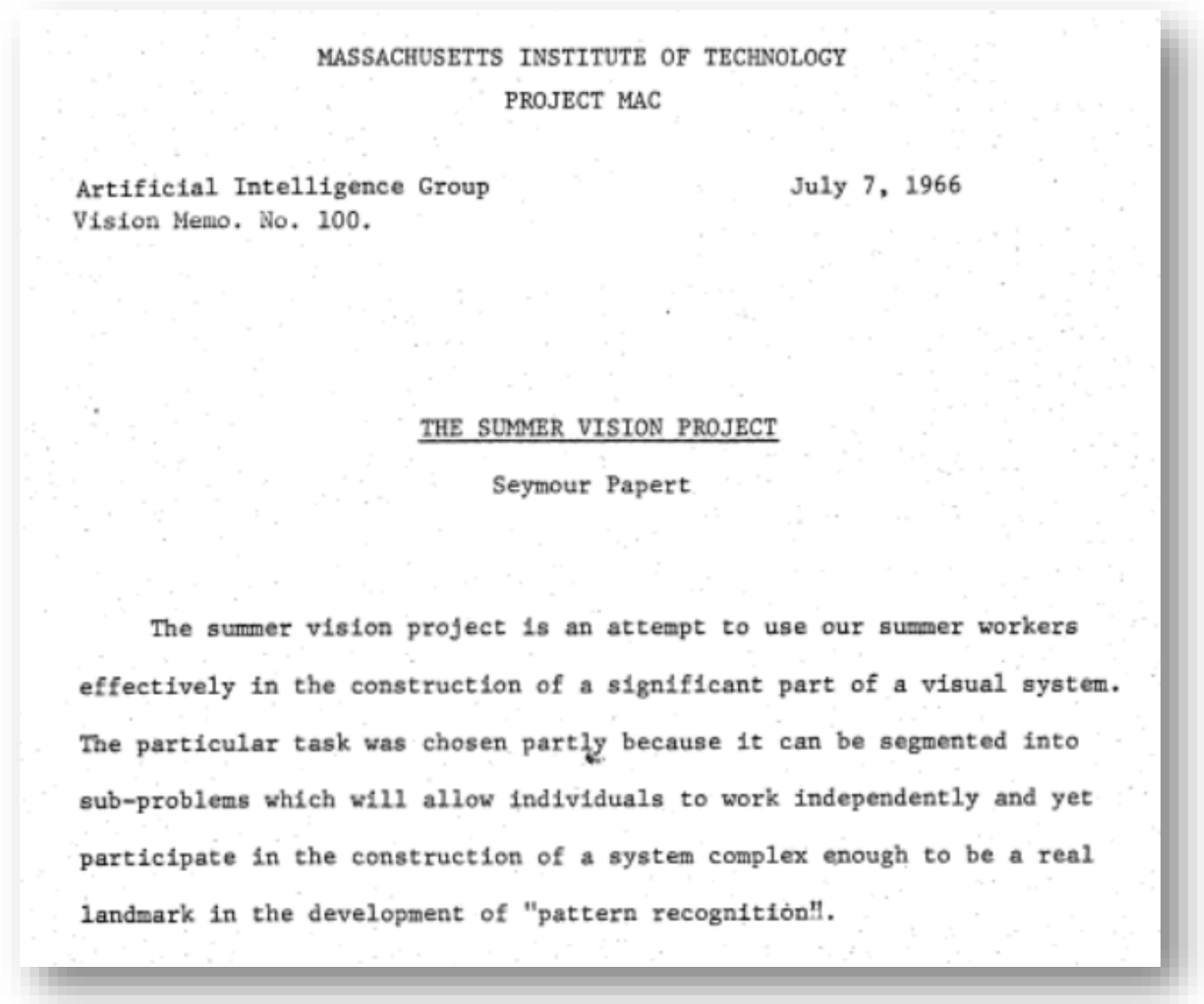
=

~3 billion
Palynomorphs
> 109 million
200TB



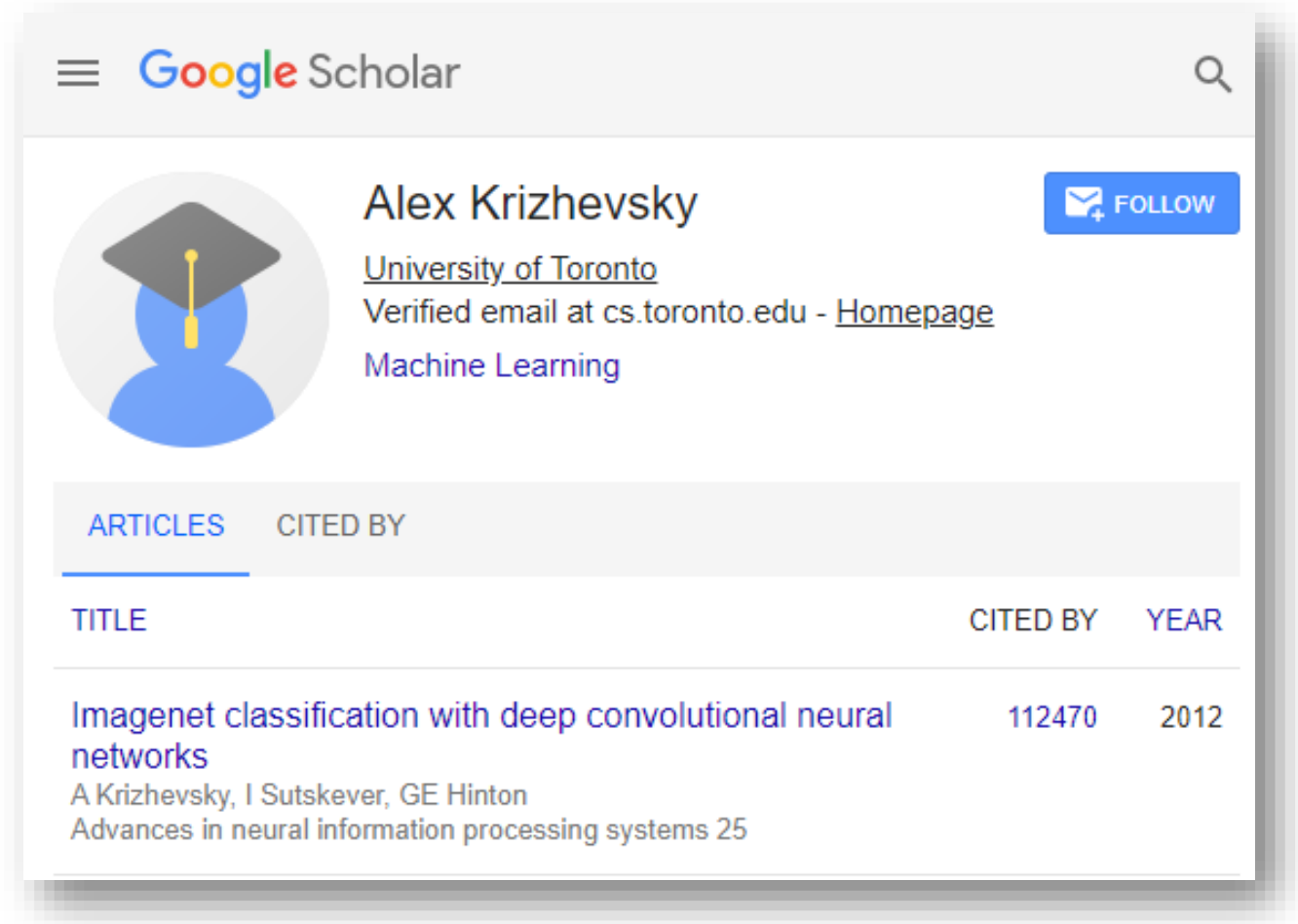
Computer-Vision Origins

- **1966** – ah, we should be able to crack this over the summer...



Computer-Vision Origins

- **1966** – ah, we should be able to crack this over the summer...
- **2012** – major breakthrough using DNNs to classify images

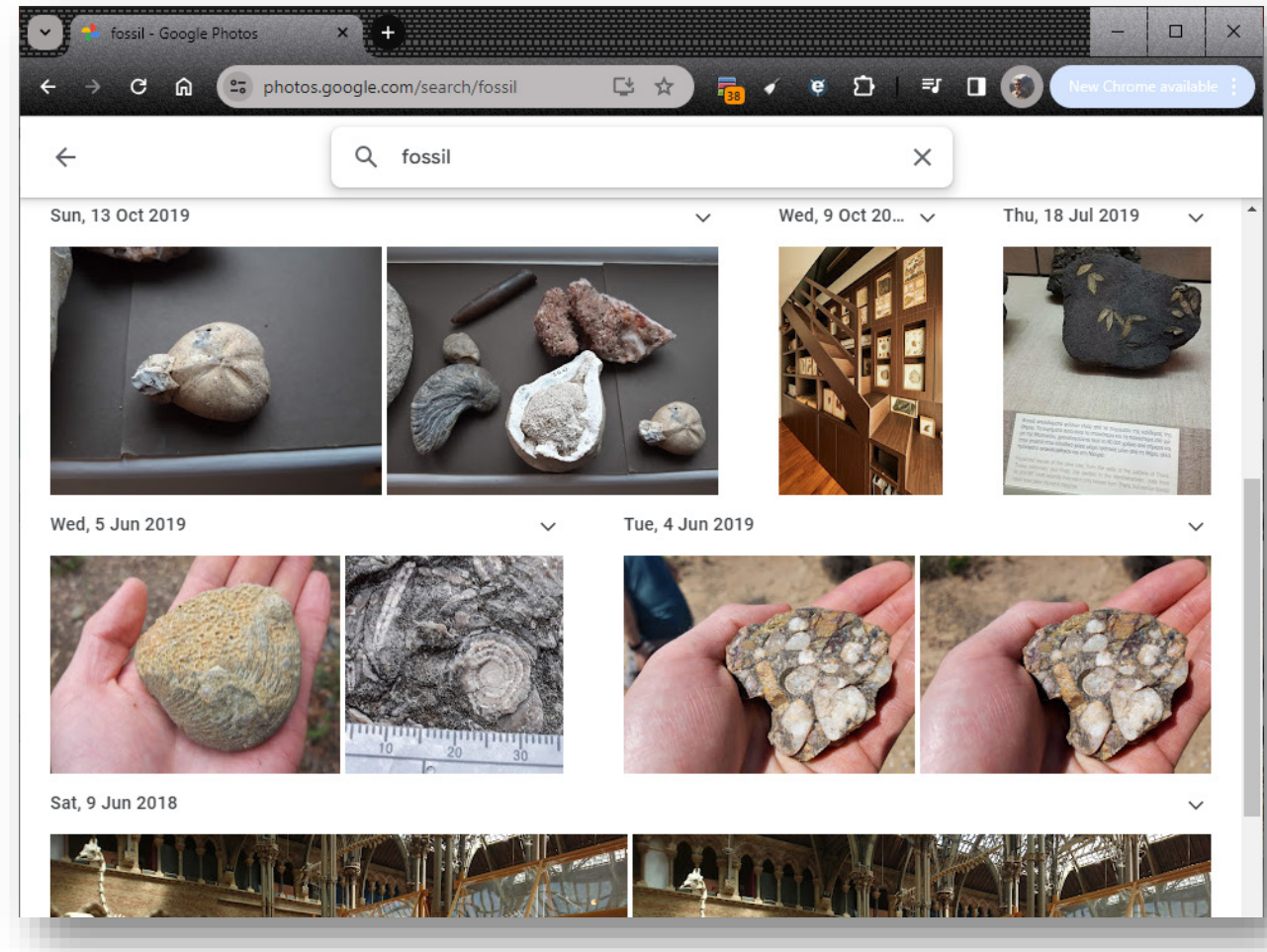


The screenshot displays the Google Scholar profile of Alex Krizhevsky. The profile includes a circular avatar with a graduation cap, his name, and a 'FOLLOW' button. Below this, it lists his affiliation as the University of Toronto, a verified email at cs.toronto.edu, and his research interests in Machine Learning. A tabbed interface shows 'ARTICLES' and 'CITED BY'. The 'ARTICLES' tab is active, displaying a table with the following data:

TITLE	CITED BY	YEAR
Imagenet classification with deep convolutional neural networks A Krizhevsky, I Sutskever, GE Hinton Advances in neural information processing systems 25	112470	2012

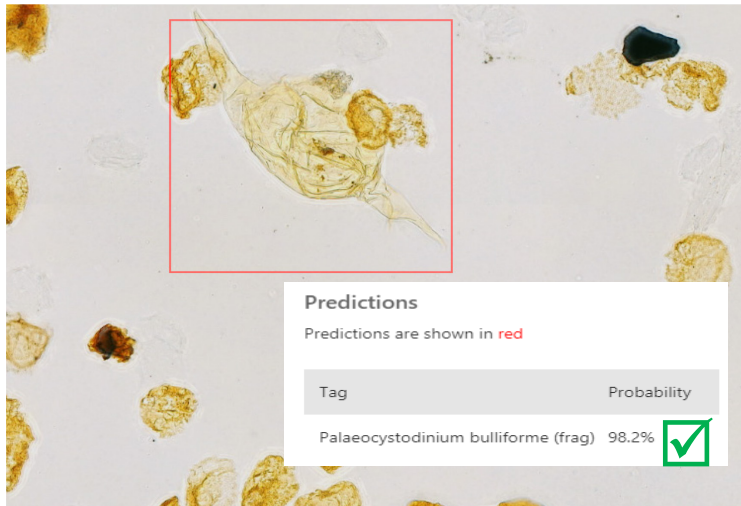
Computer-Vision Origins

- **1966** – ah, we should be able to crack this over the summer...
- **2012** – major breakthrough using DNNs to classify images
- **Now** – readily available commodity



Example results from supervised learning

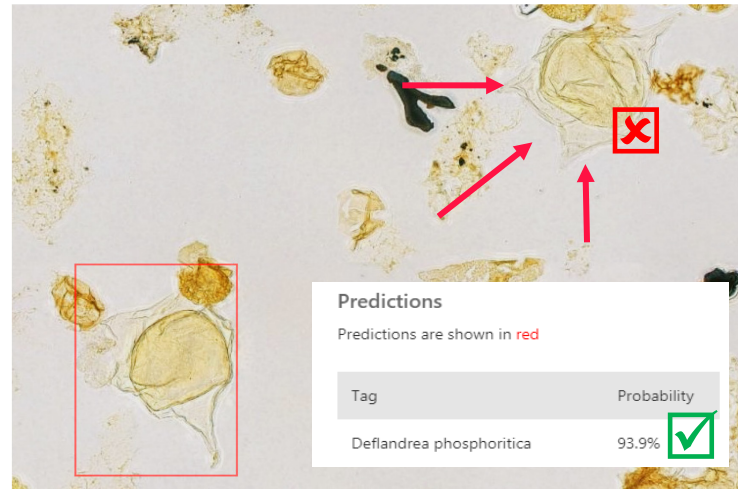
The Good



Correct prediction on image from outside training set

- ✓ *Palaeocystodinium* is here!
- ✓ No false positives

The Bad



Incomplete prediction on image from outside training set

- ✓ *Deflandrea phosphoritica* here
- ✓ Second *Deflandrea* missing?

The Ugly



Confident and wrong

- ✓ Not a *Deflandrea*
- ✓ Ignored pollen

10 |

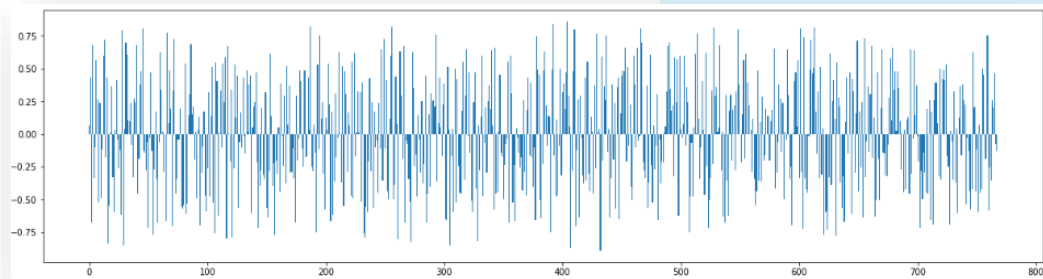
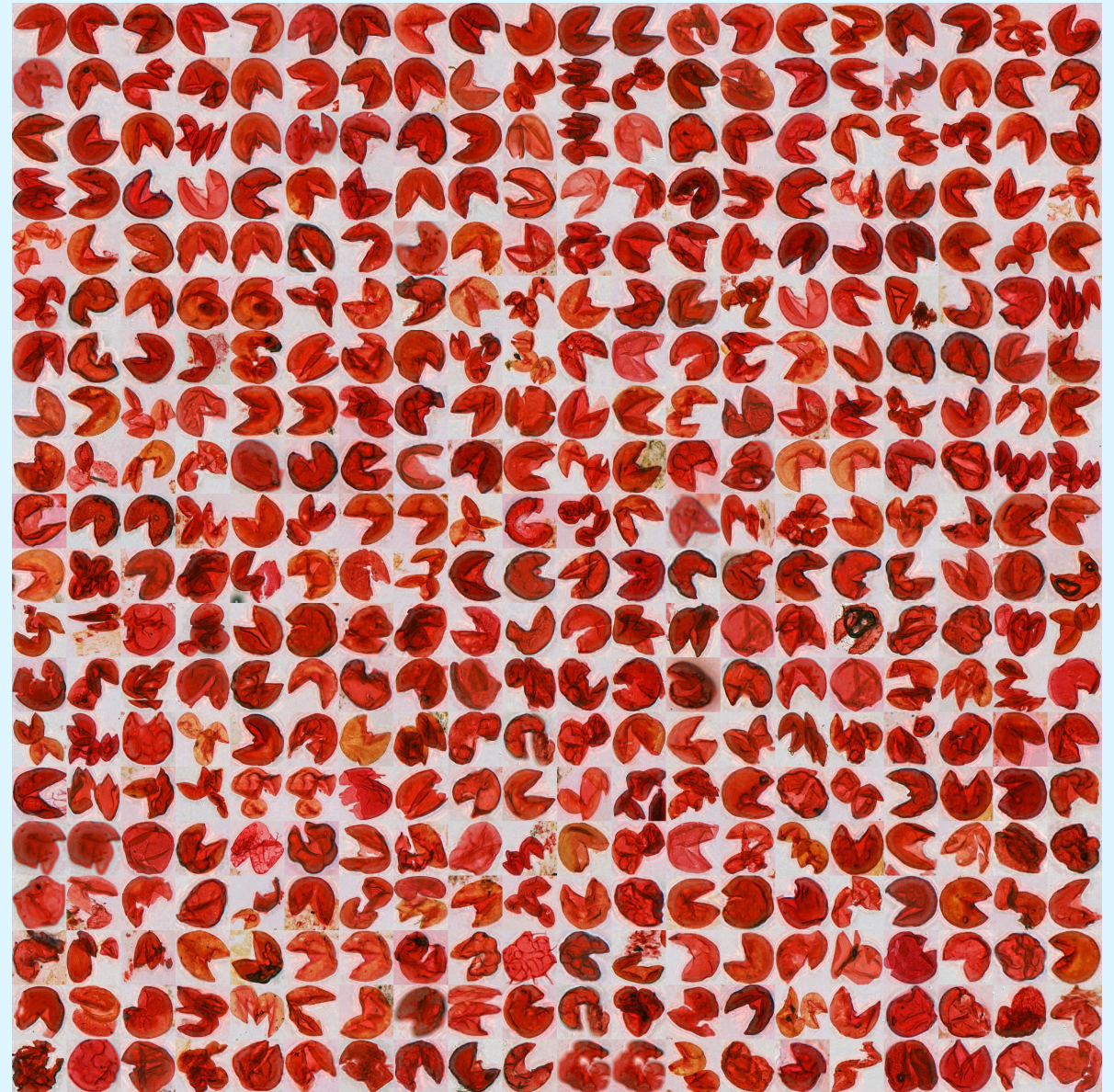


Content-Based Image Retrieval

Query :

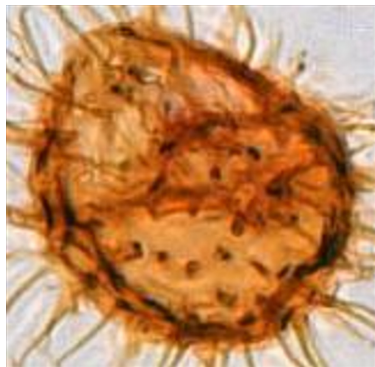


Result :

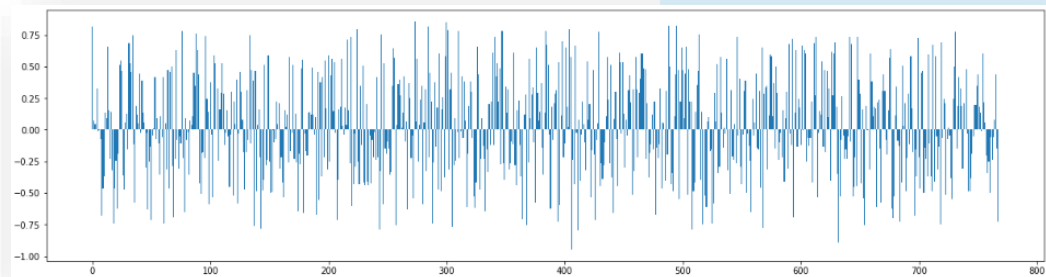


Content-Based Image Retrieval

Query :

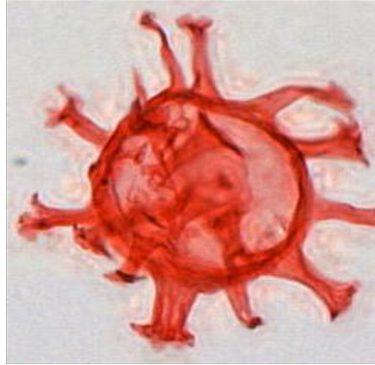


Result :

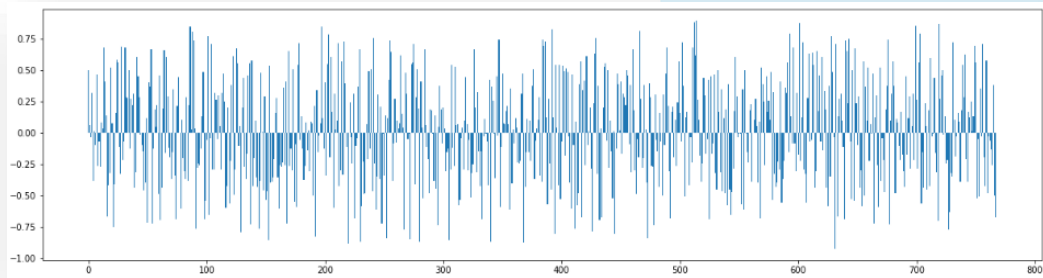
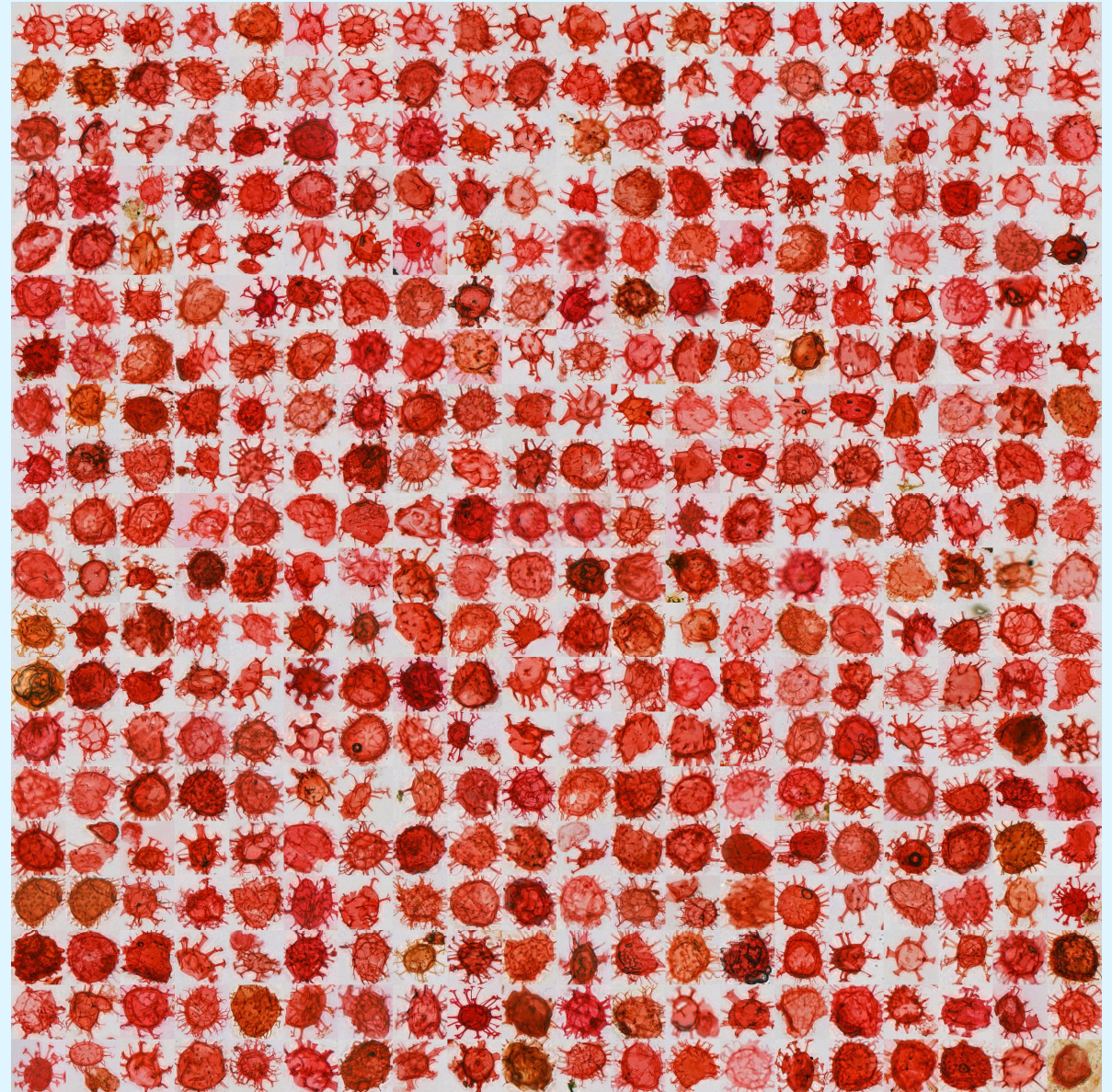


Content-Based Image Retrieval

Query :

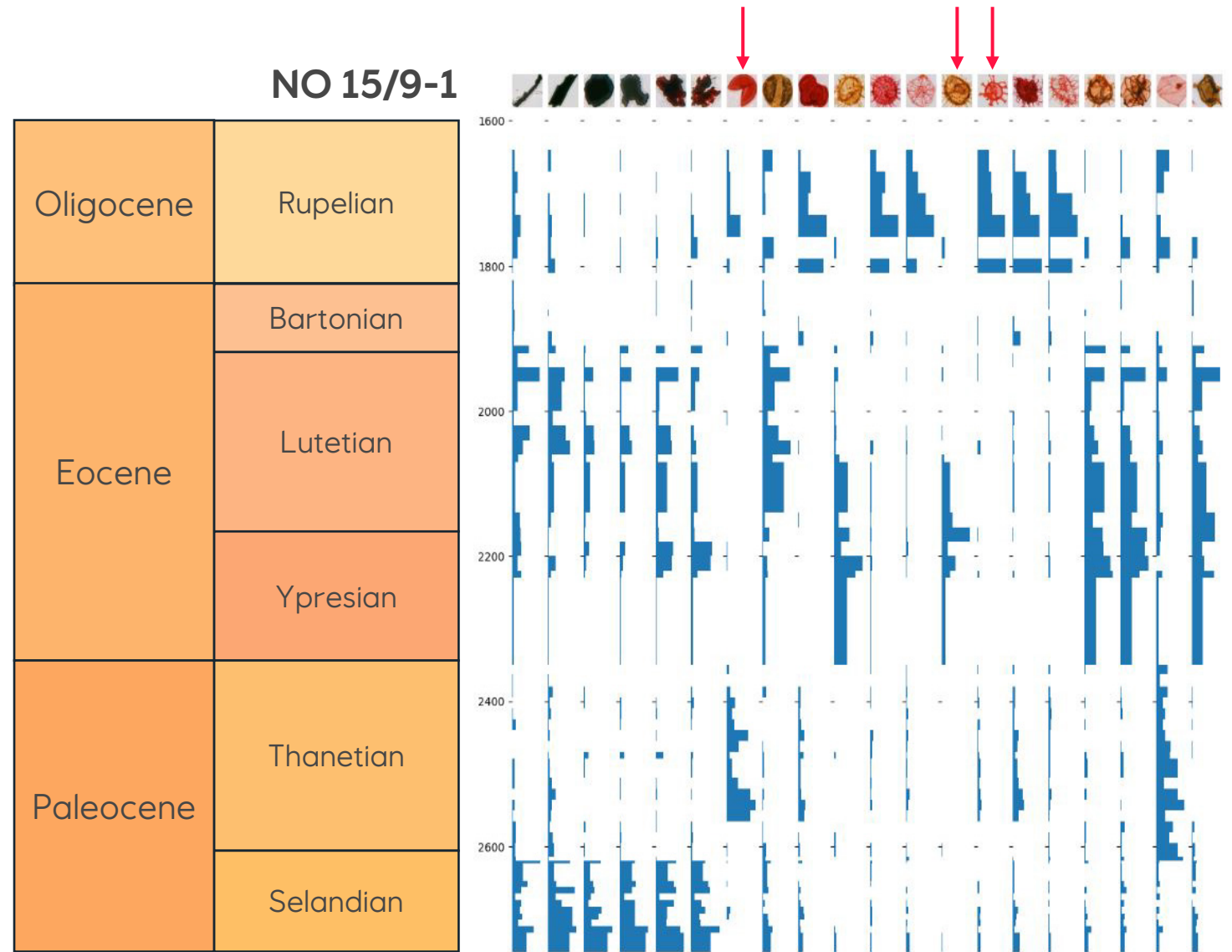


Result :



Chronostratigraphy

- Define multiple search crops
- Find depths of matching crops
- Plot histograms of matches
- Compare with well picks



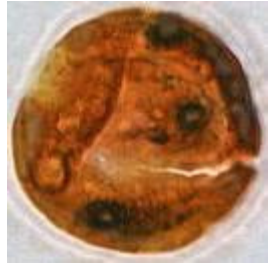
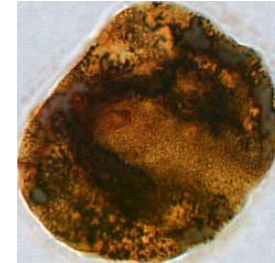
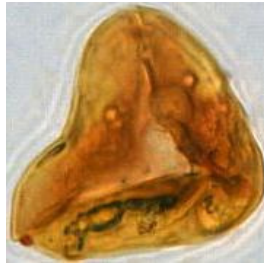
Use Case : Formation Zonation

Iteration #1 : Queries from not-a-palynologist

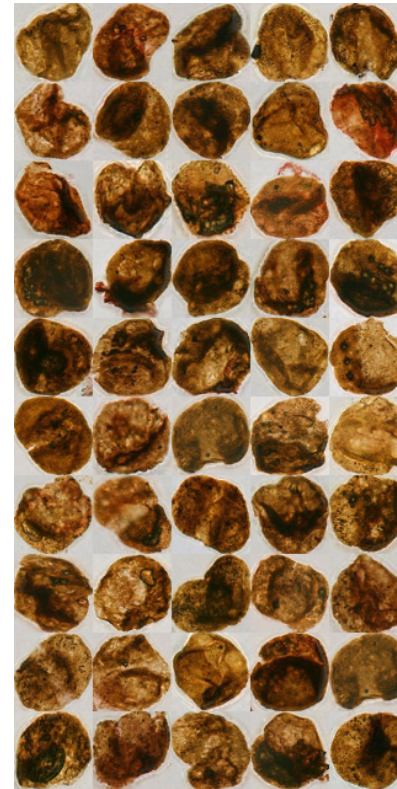
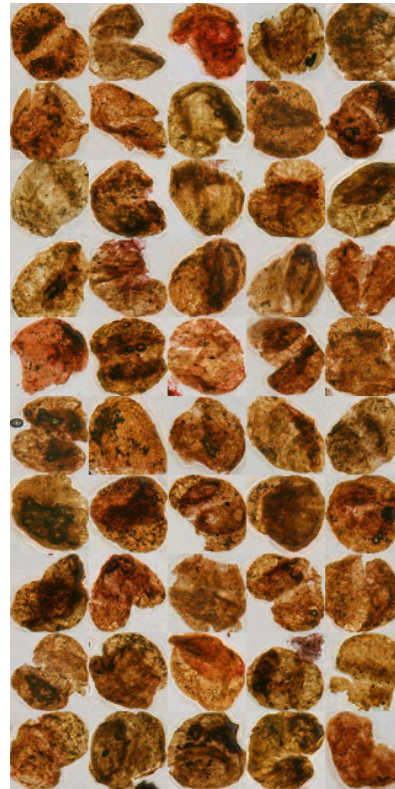
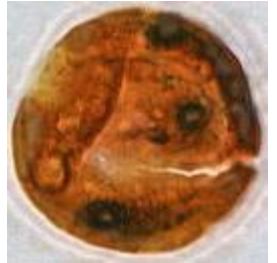
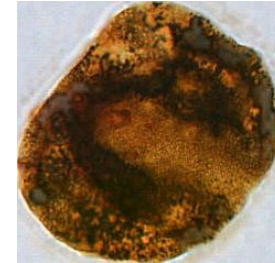
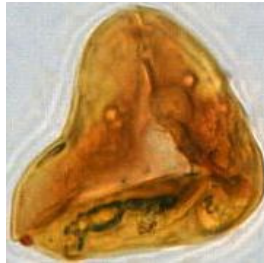
This works surprisingly well:

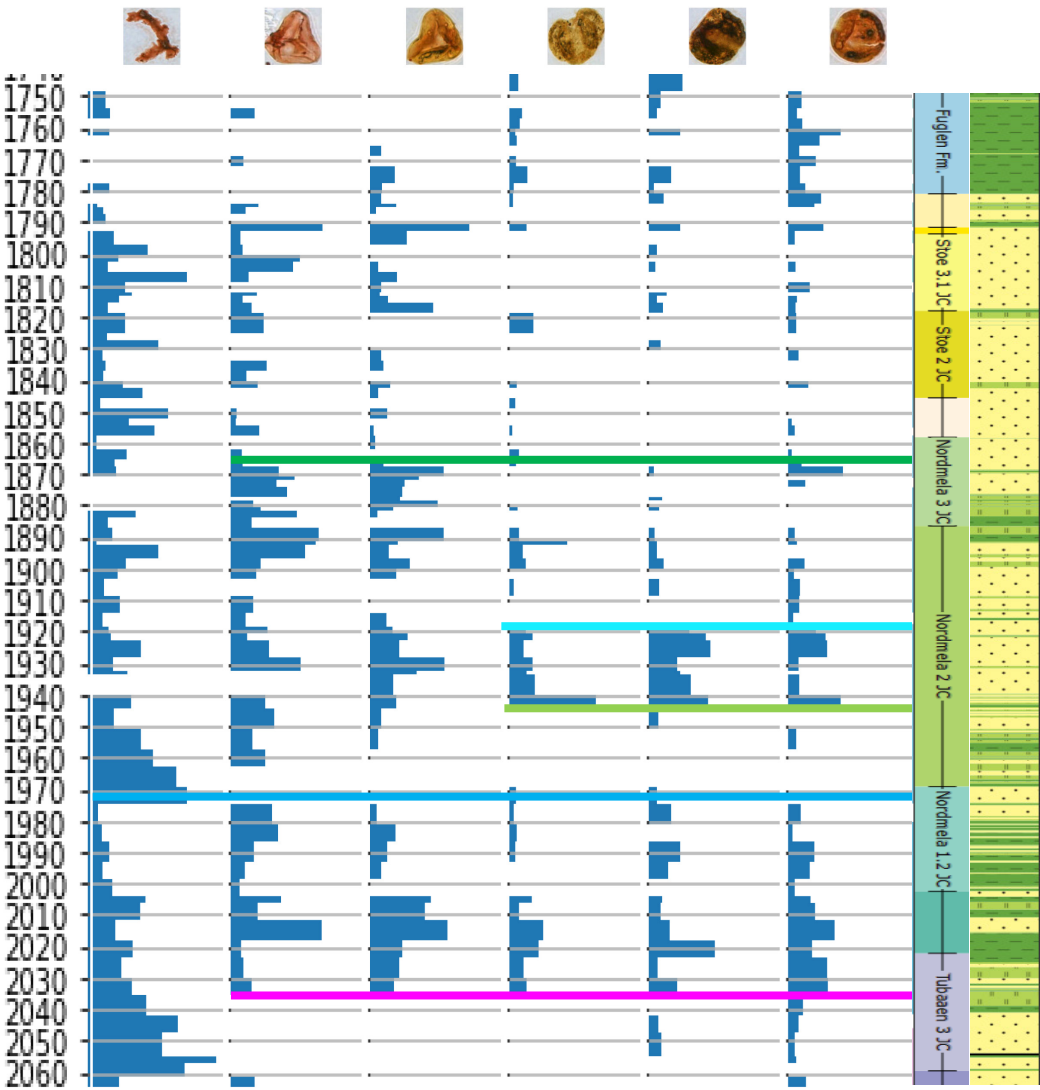
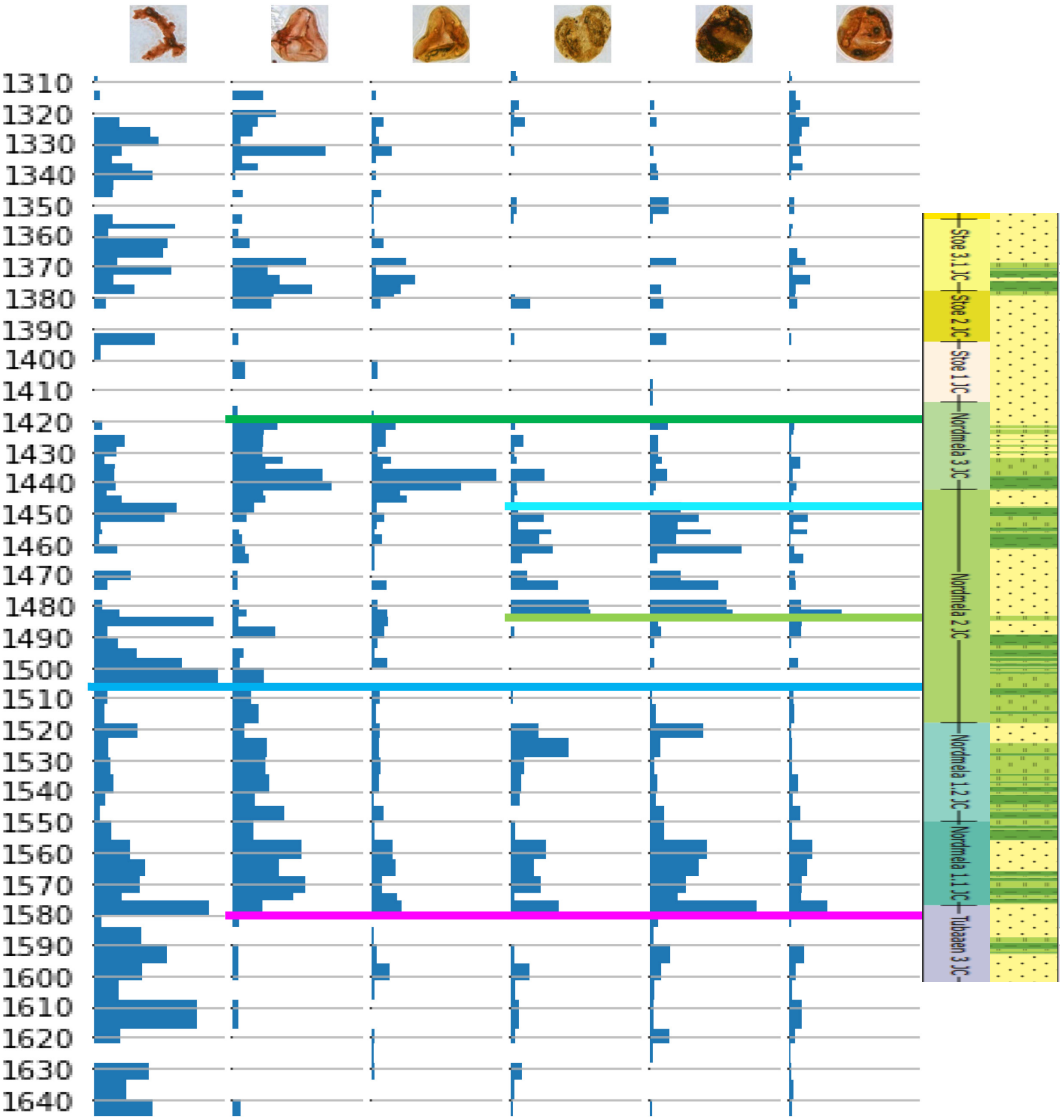
- ... finds things which look like the queries
- ... but what do the results mean?!

Well A – query results



Well B – query results





Use Case : Formation Zonation

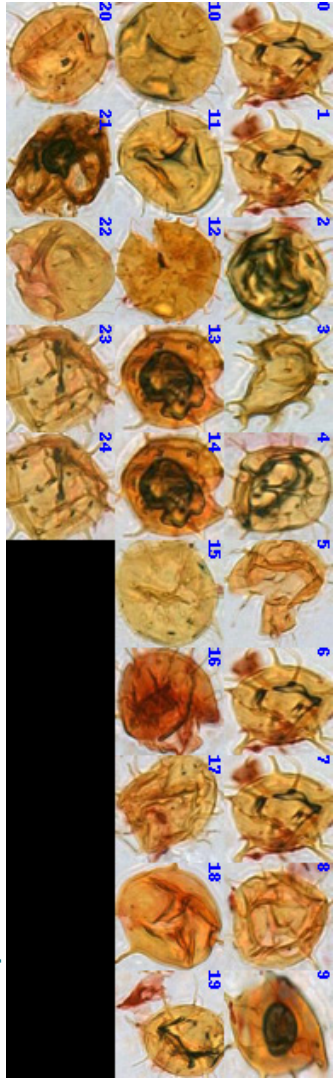
Iteration #2 : One real palynologist runs queries

Find *exactly* what we're looking for:

- ... interpretable results, fast!
- ... consistent patterns! (which *nearly* agree with official Equinor zonation)
- ... useful information for the asset!

Well A – query results

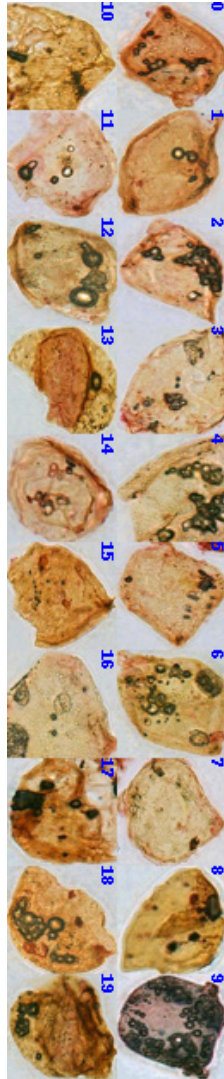
Micrhystridium



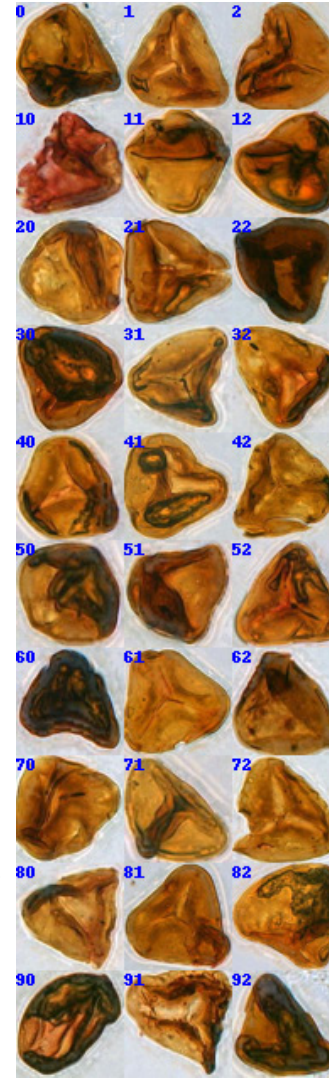
Limbicysta bjaerkei



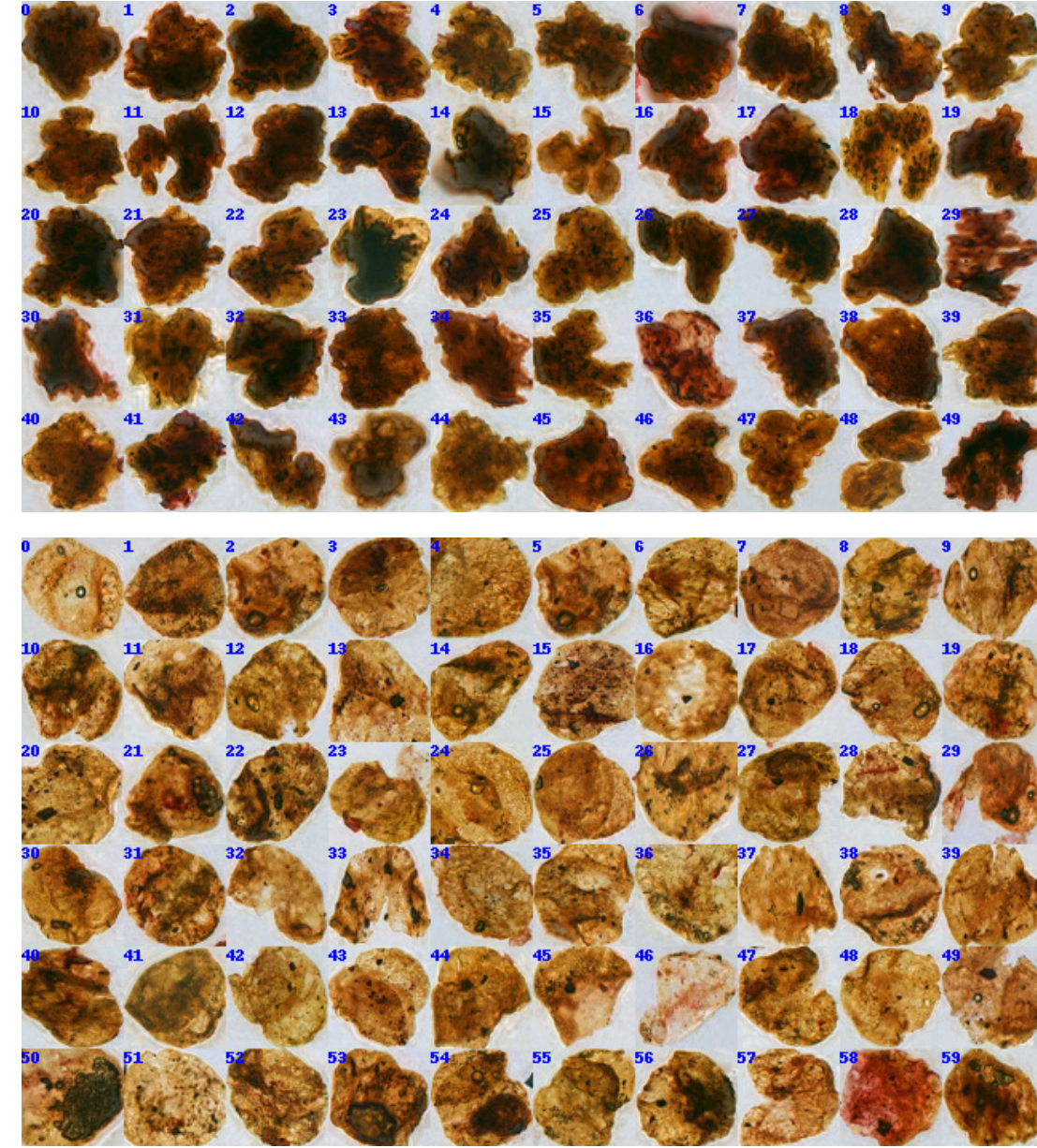
Nannoceratopsis spp.



Deltoidospora toralis

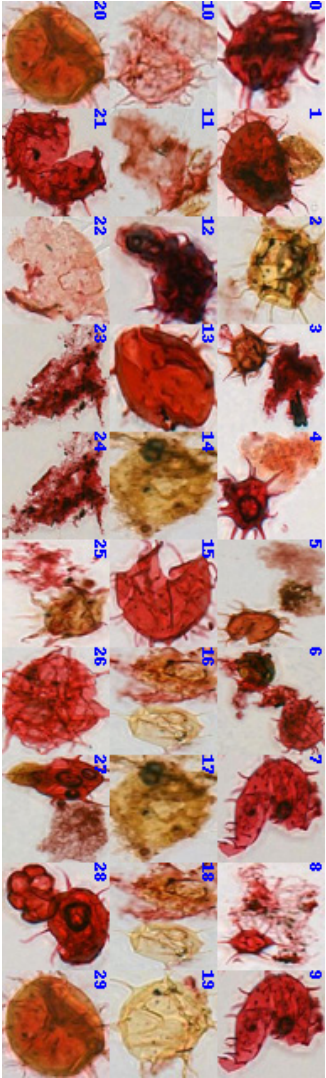


Botryococcus
Bisaccate

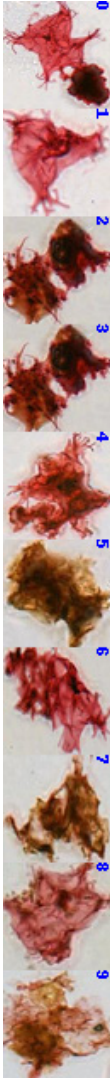


Well B – query results

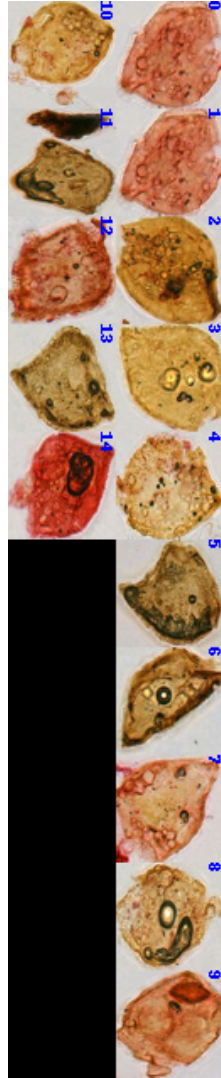
Michrystidium



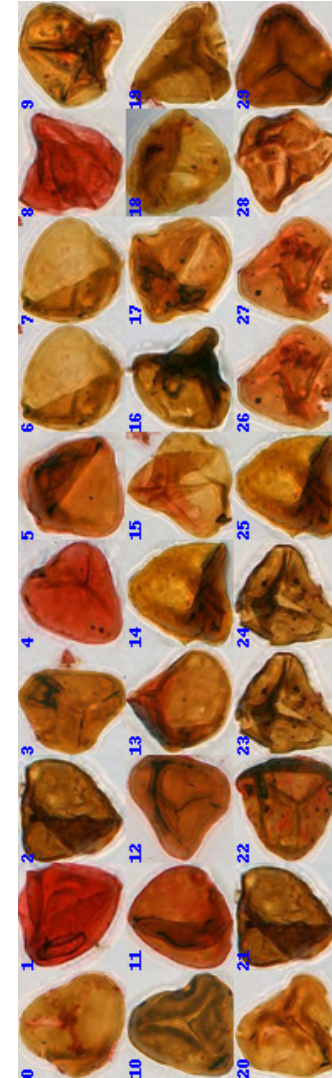
Limbicysta bjaerkei



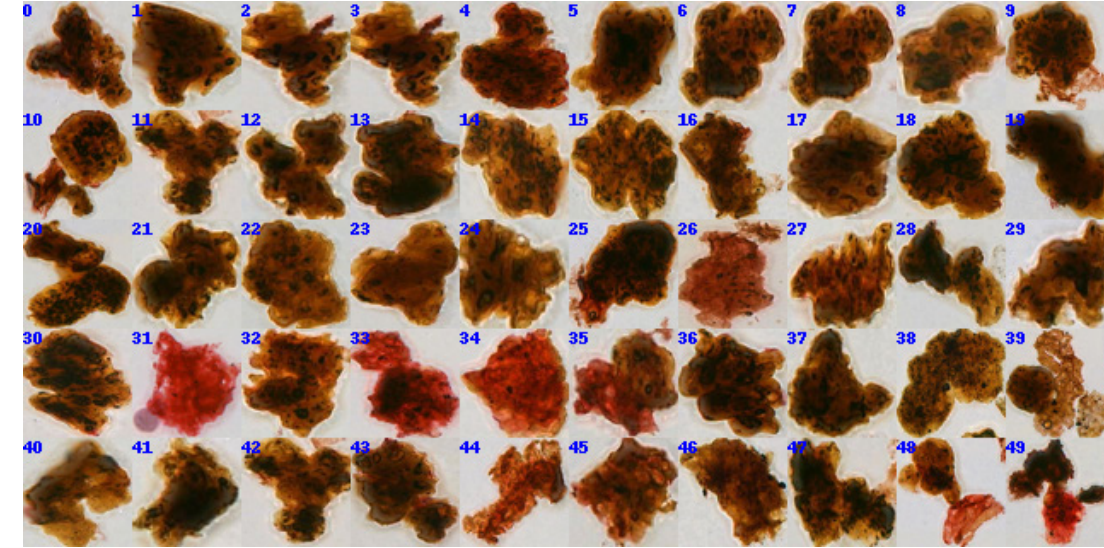
Nannoceratopsis spp.



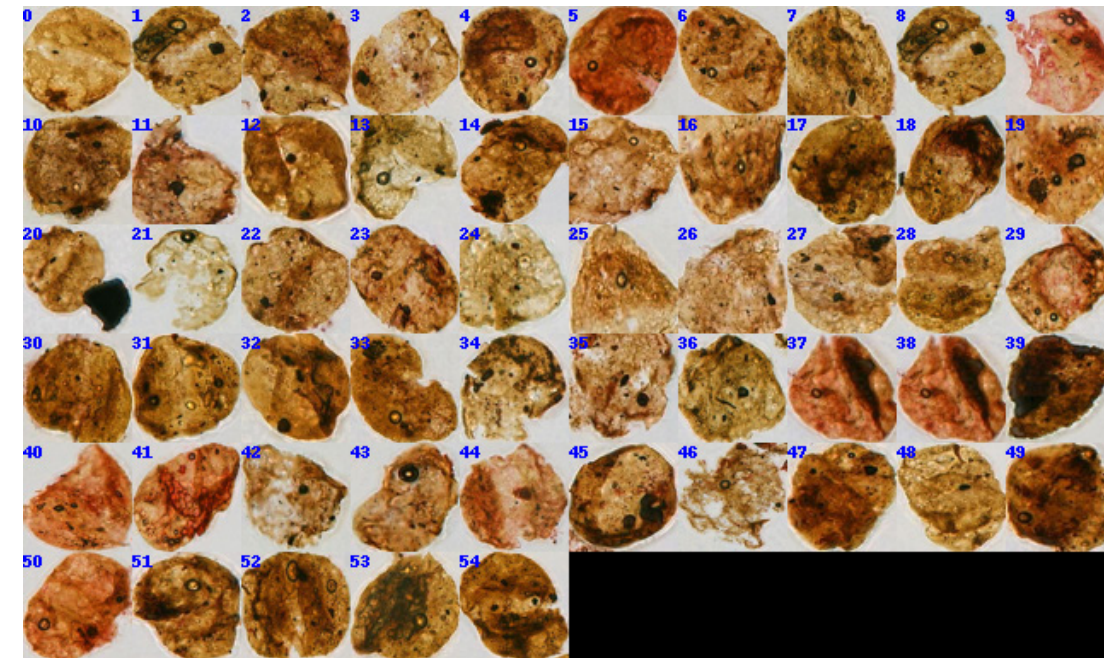
Deltoidospora toralis



Botryococcus



Bisaccate

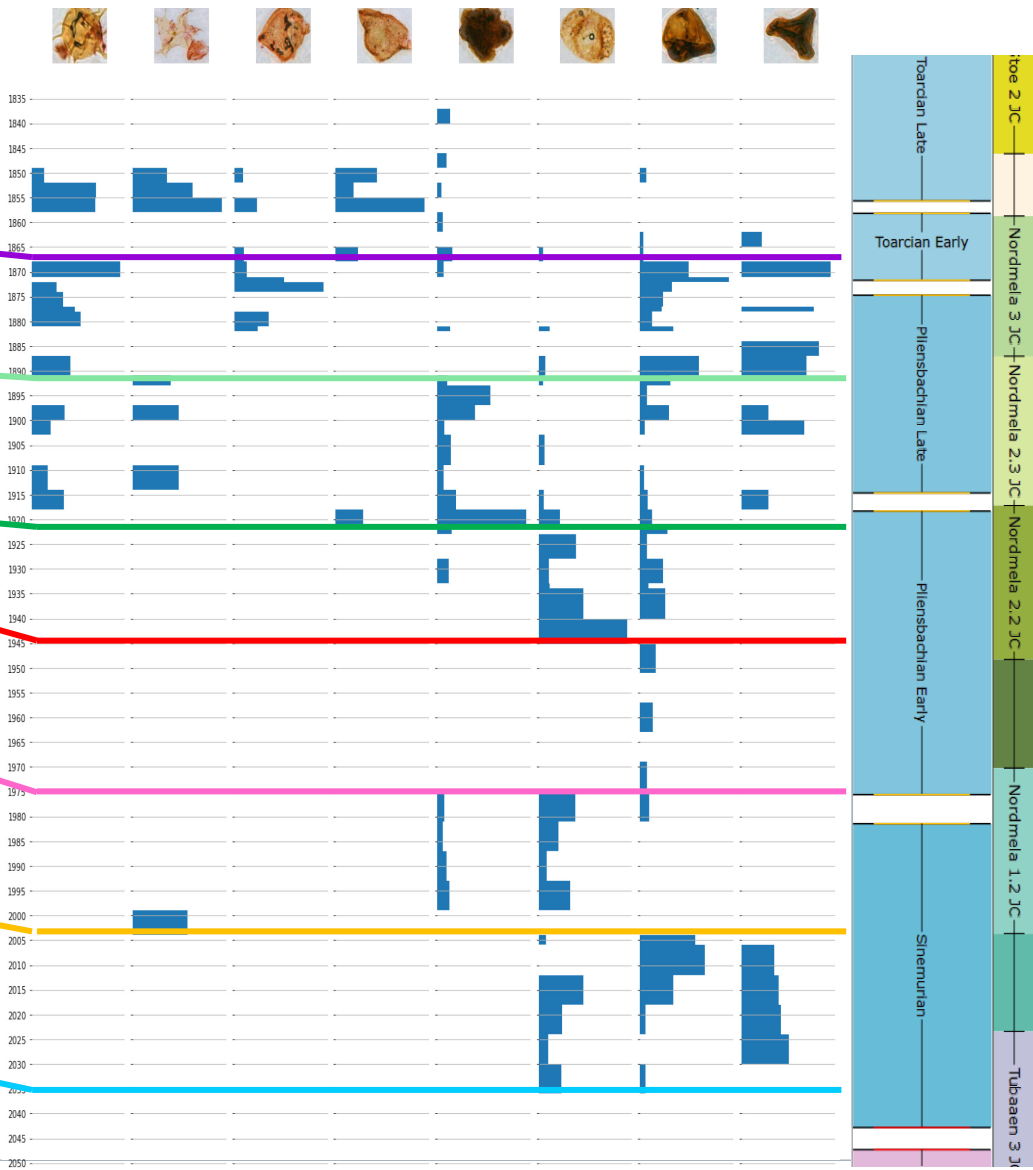


Scampi charts

Well A

Palynologists finds some interesting images to query with
- **and** refines queries **and** QCs results,

Well B



Implementation

One webapp for all:

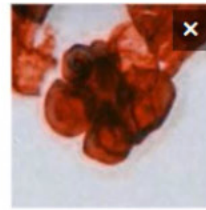
- Logging contractors
- Equinor biostrat team in Shared Services
- Equinor generalists

Choose a file

Drop your file here

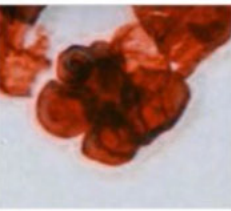
Search

Choose a file



Selected file:
foram_lining.JPG

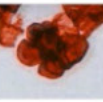
Search



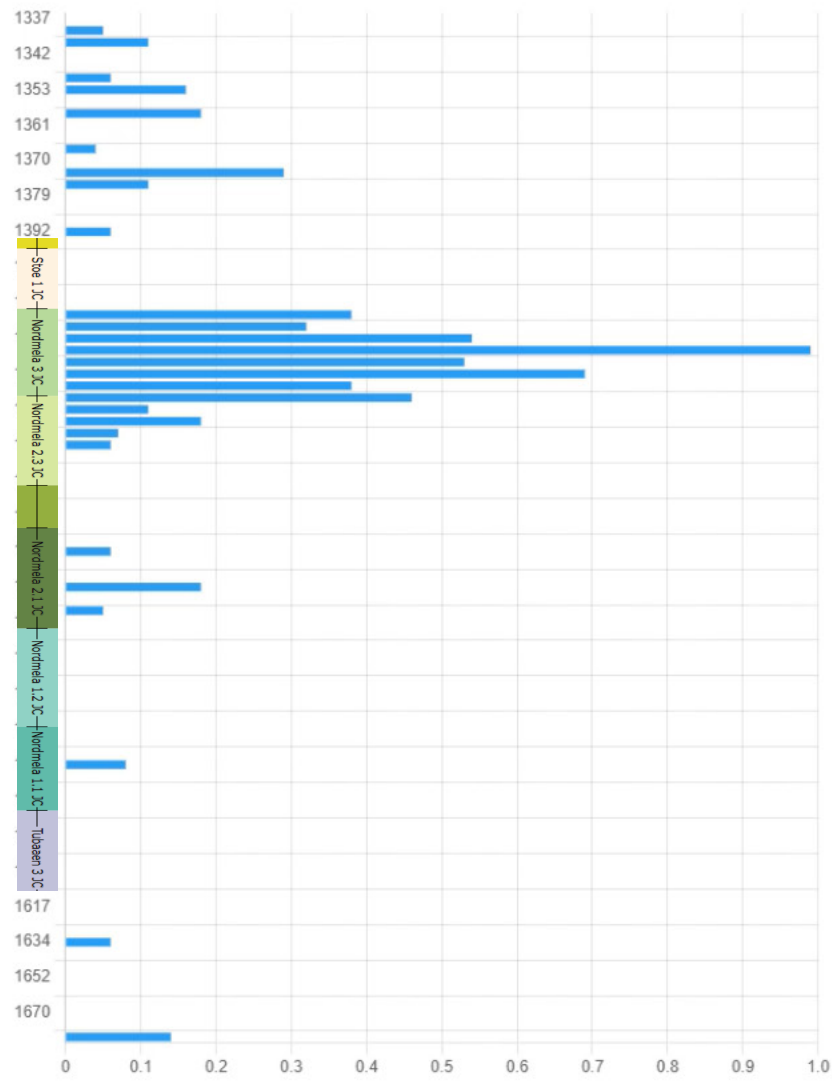
number of result slides to show is 100



Number of crops loaded: 100/100



Relative Counts



Select query crop

No well selected

NO 16/2-10

NO 16/2-12

NO 33/9-6

NO 34/10-30

NO 34/2-2

NO 34/2-3

Select query crop

Library

MultiQuerySearch and LineProjection

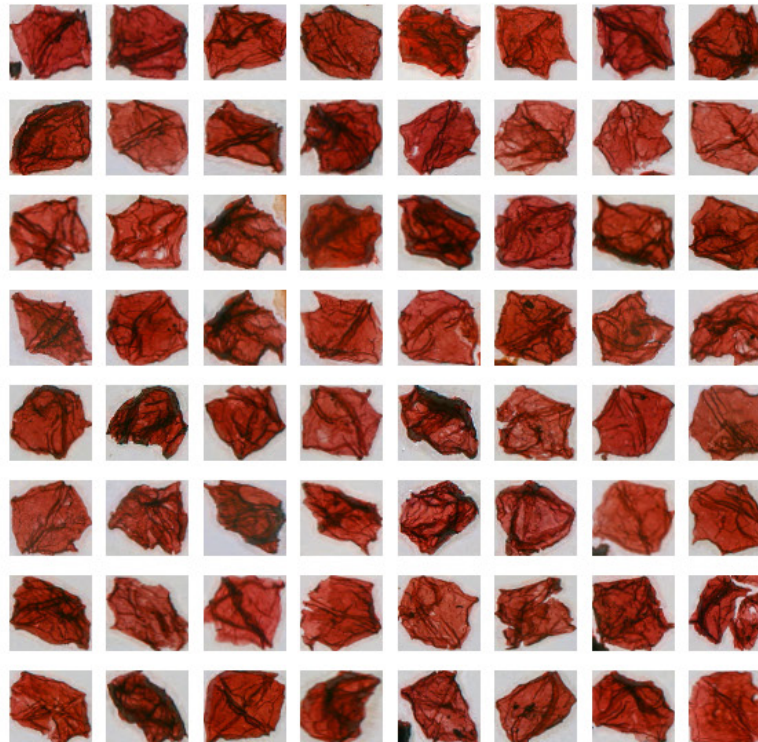
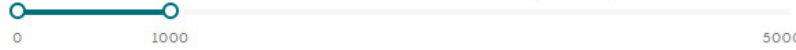
Refine search



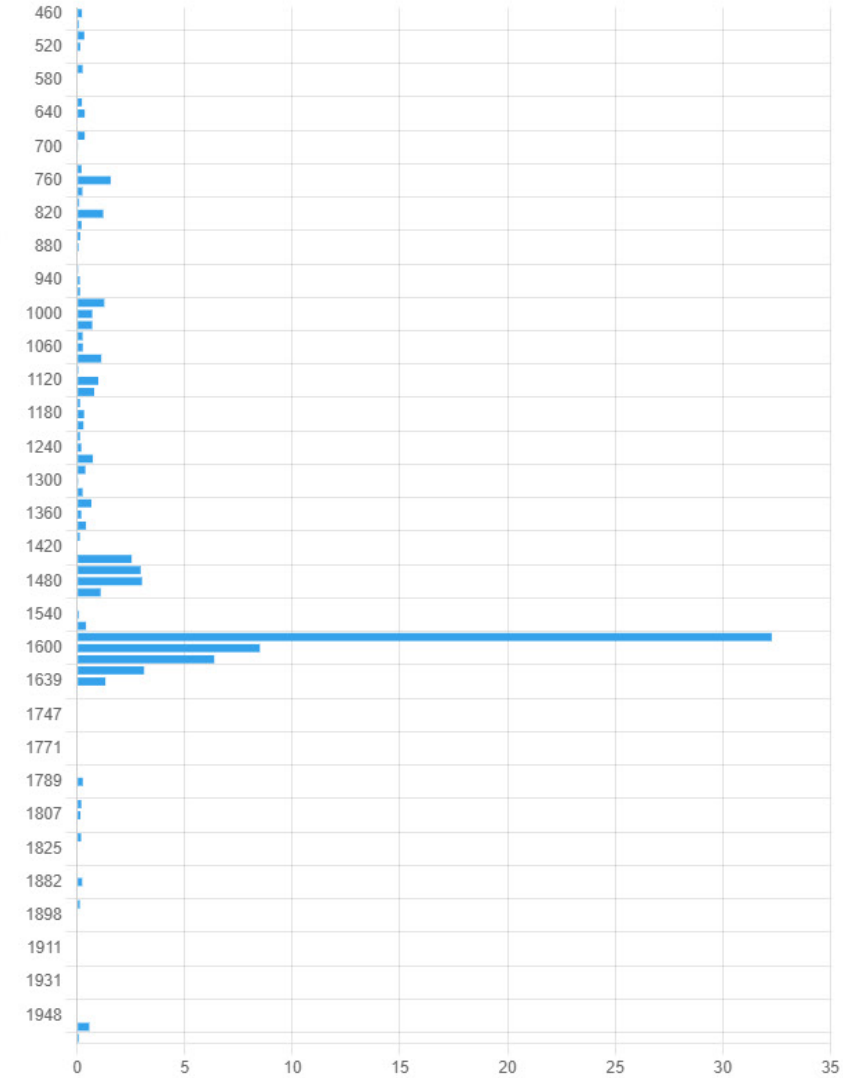
Relative Counts



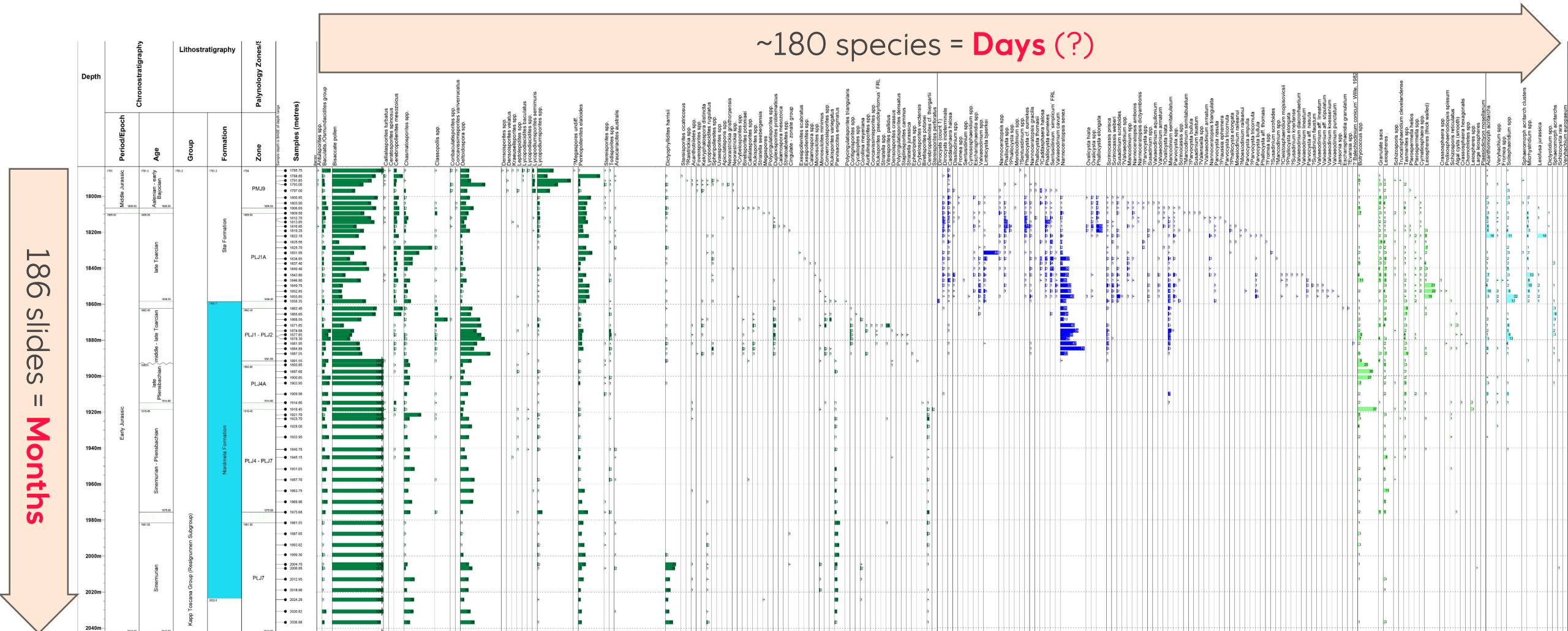
number of result slides to show is 1000



Number of crops loaded: 100/1000

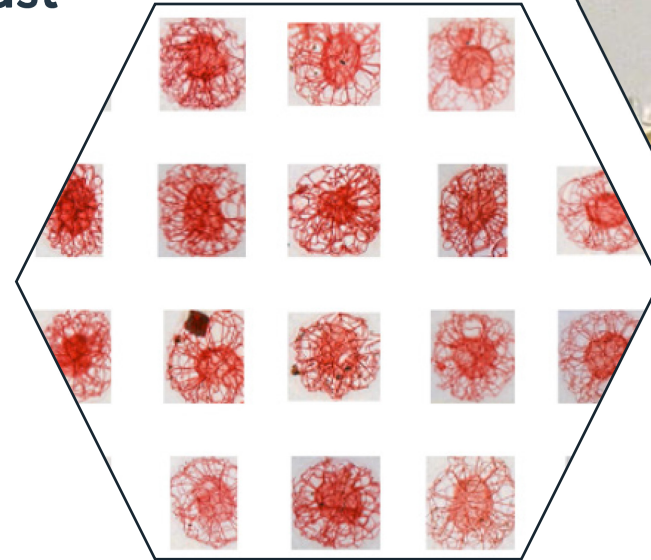
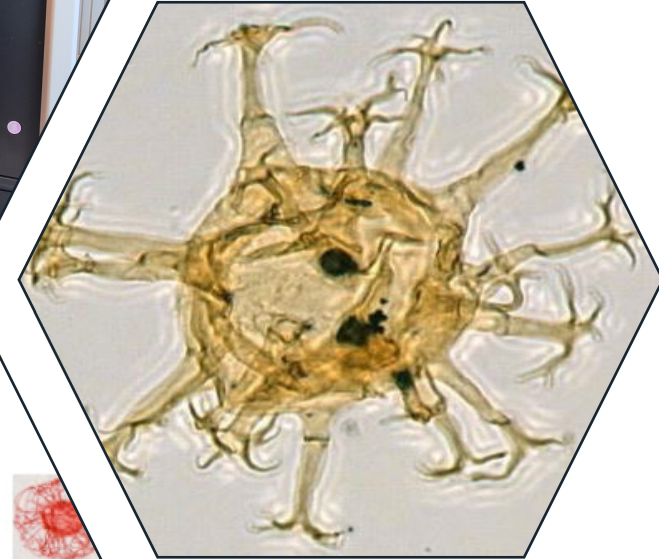
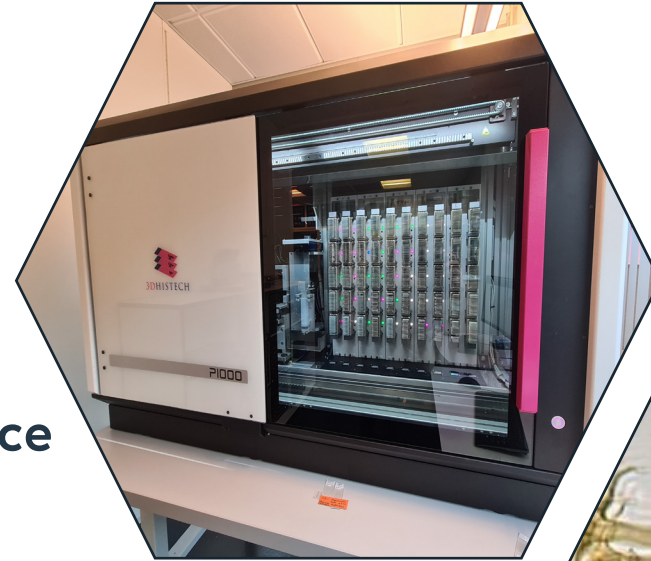


Imagine creating a distribution chart for a new well...



Summary

- NOD's Digital Palynology is an amazing resource
- Scampi enables us to maximize value from it, fast
- Totally new way to view and use biostrat data
- Improves communication with non-specialists



Species Classification Automation for Microfossil Photomicrograph Images (Scampi)

David Wade*, Sissa Stefanowicz, Alex Cullum & Erik Anthonissen
Muhammad Gibran Alfarizi, Håkon Ruud,
Thomas Elvestad†, Einar Salomonsen†, Thomas Karlsson†
† Bouvet ASA