

Automating tropical pollen counts using convolutional neural nets: from image acquisition to identification

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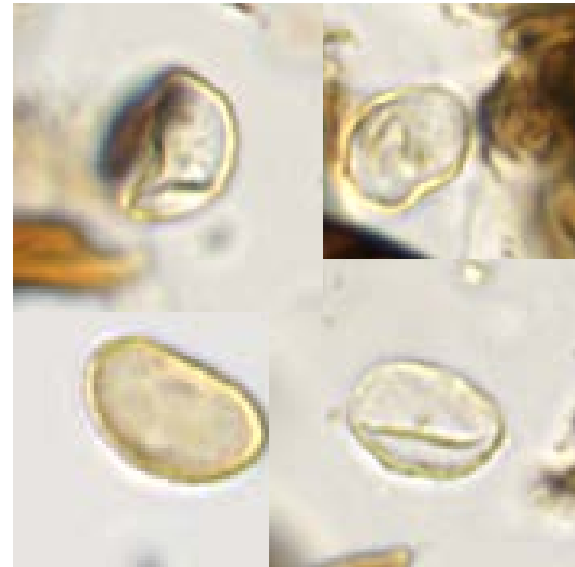
²*University of California at Irvine*

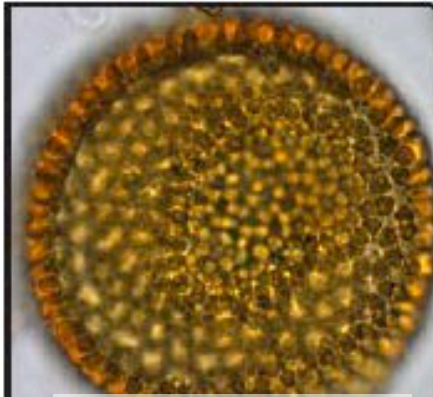
³*Smithsonian Tropical Research Institute, Panama City, Republic of Panama*

⁴*National Center for Supercomputing Applications,
University of Illinois at Urbana-Champaign*

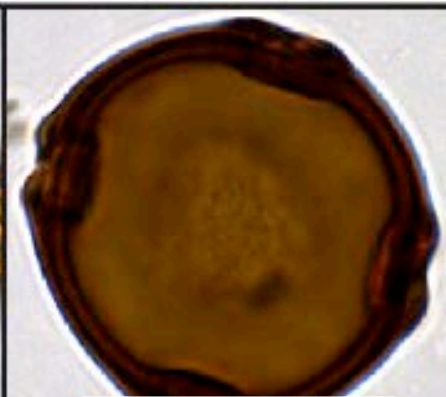


Cecropia (Urticaceae)

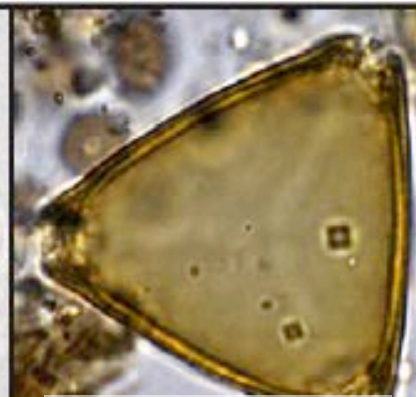




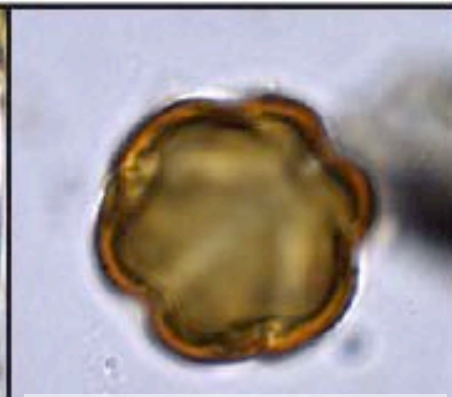
Croton (Euphorbiaceae)



Trichilia (Meliaceae)



Serjania (Sapindaceae)



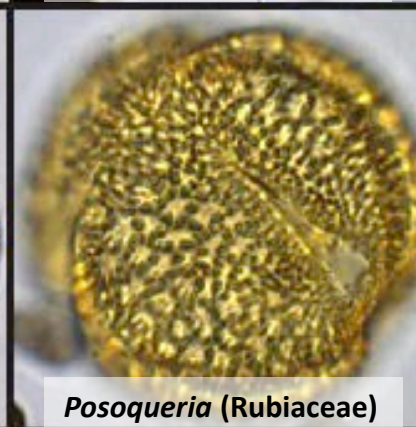
Combretum (Combretaceae)



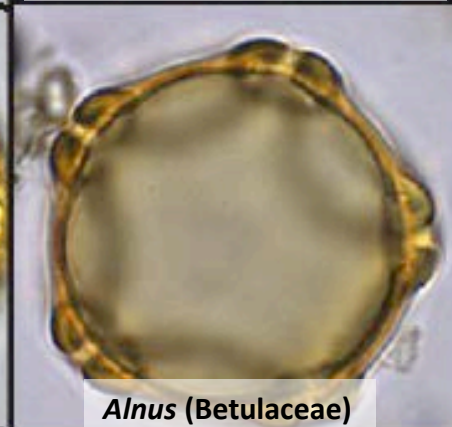
Quararibea (Malvaceae)



Pseudobombax (Malvaceae)



Posoqueria (Rubiaceae)



Alnus (Betulaceae)



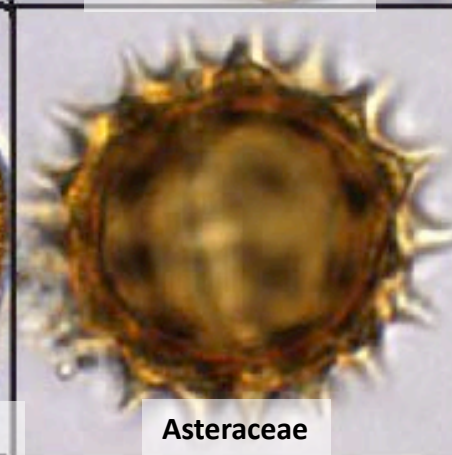
Quassia (Simaroubaceae)



Dalechampia (Euphorbiaceae)



Anacardium (Anacardiaceae)



Asteraceae

POLLEN AS “BIG DATA”



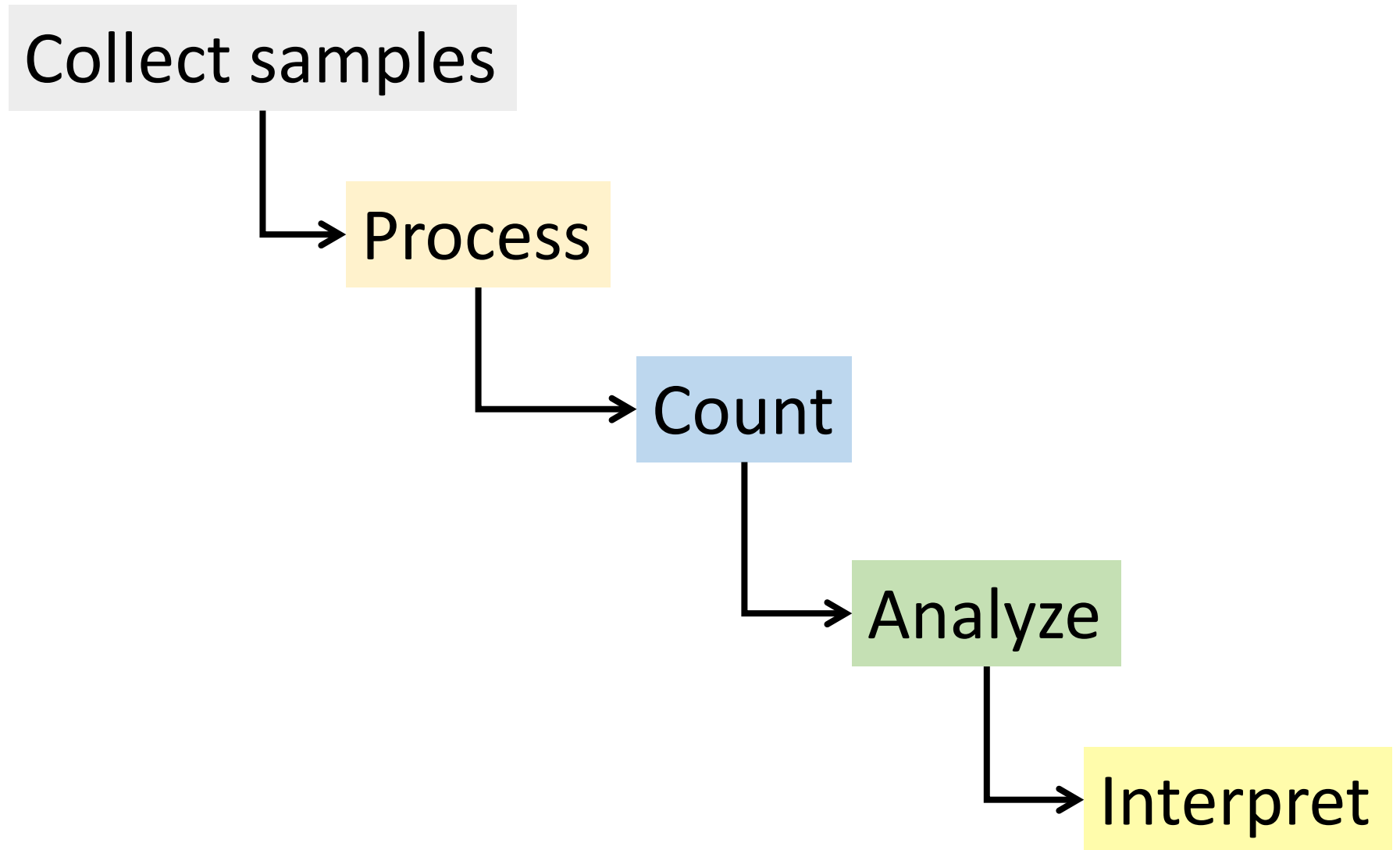
~470 million years of plant history

Billions of potential specimens

~Continuous deposition across a range of environments



REIMAGINING THE WORKFLOW



IS AUTOMATION THE ANSWER?

Quantity: increase the throughput of pollen analysis

Reproducibility: improve the consistency and accuracy of pollen identifications

Resolution: produce repeatable recognition of *species* from pollen for more precise biome reconstructions



TRAINING ON HYPERDIVERSE SAMPLES

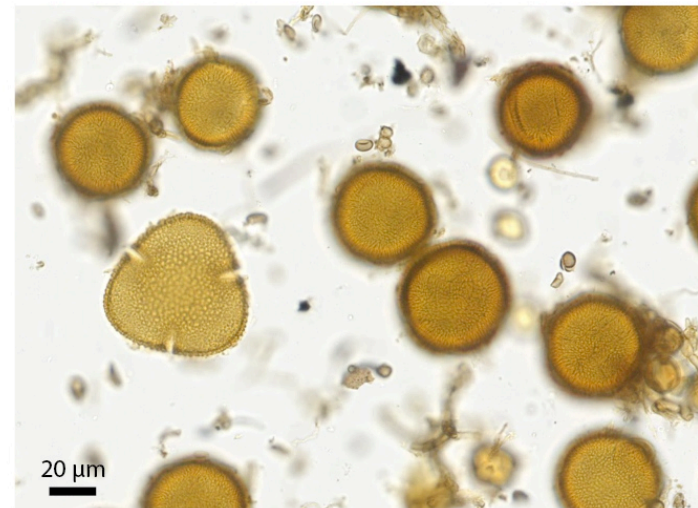
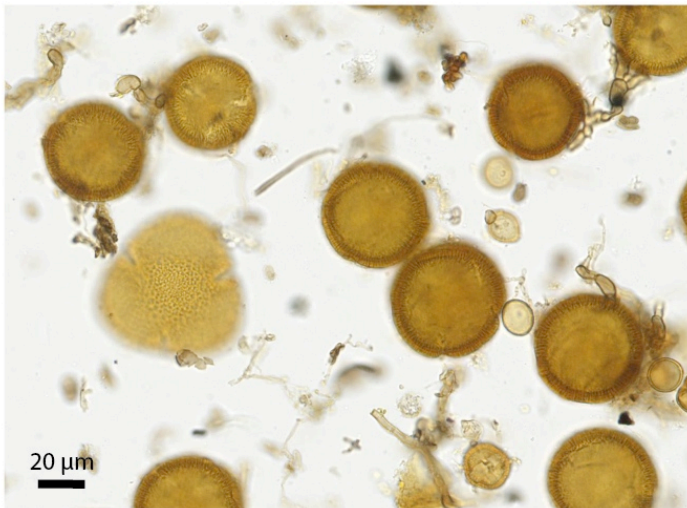
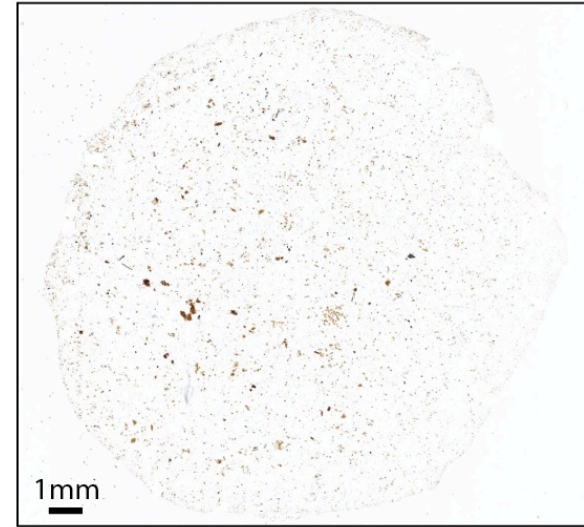
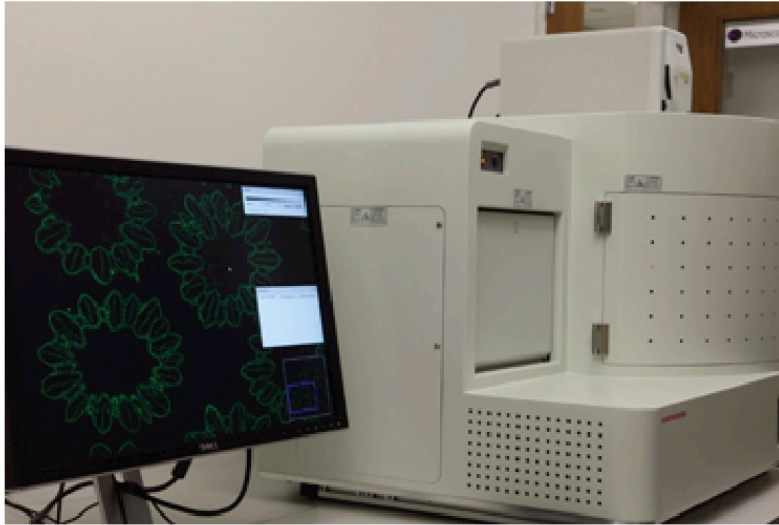
- A 15-year pollen rain record from Barro Colorado Island, Panama
 - Obtained from a series of 20, evenly spaced pollen traps along two parallel transect in the 50 ha CTFS plot
- A 10-year pollen record from the Lutz weather tower
 - Images to be analyzed this summer
- ~ 130 pollen morphotypes



Photo: STRI



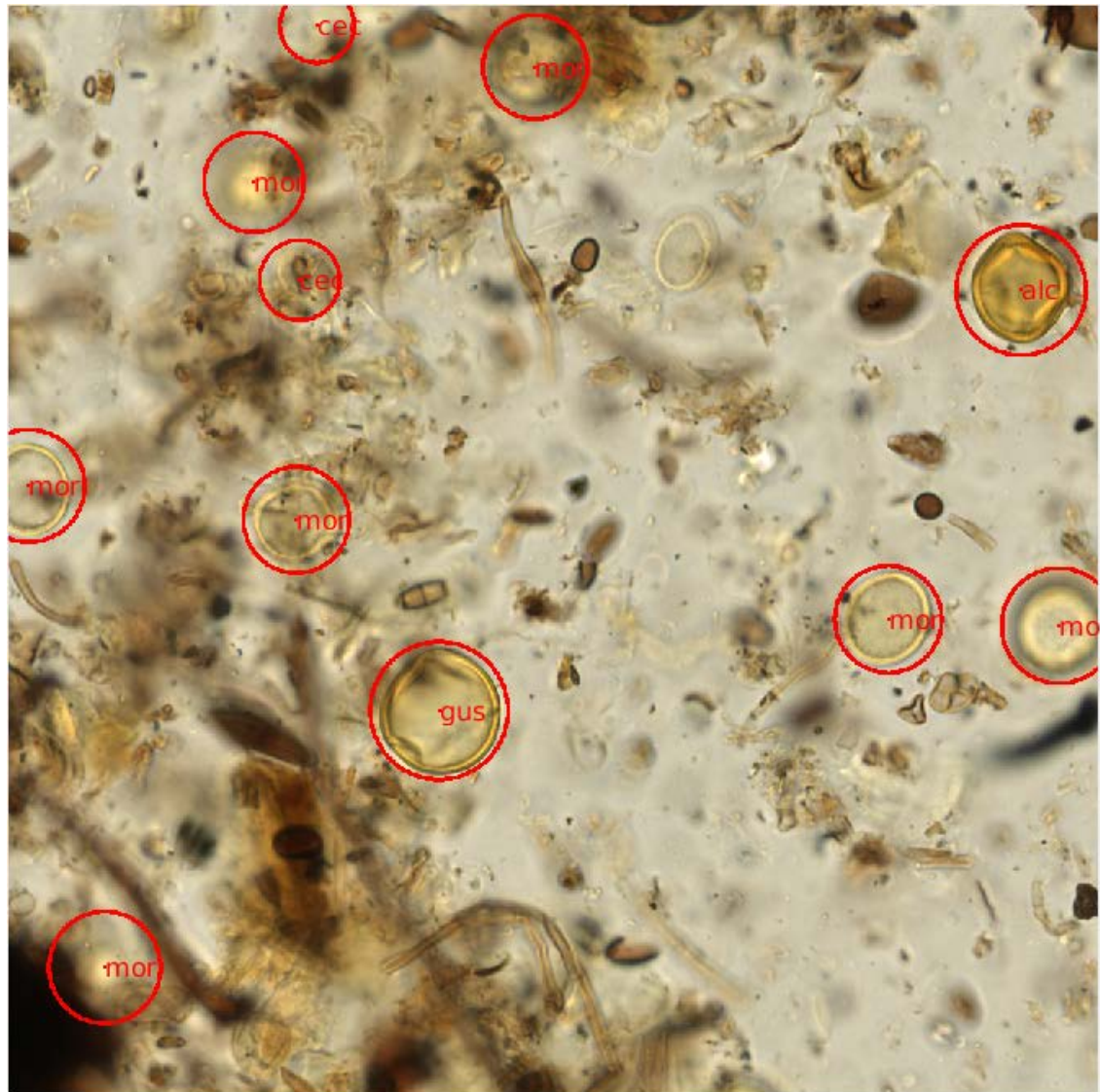
IMAGING POLLEN SLIDES FOR COUNTING



400x, 0.23 μm/pixel

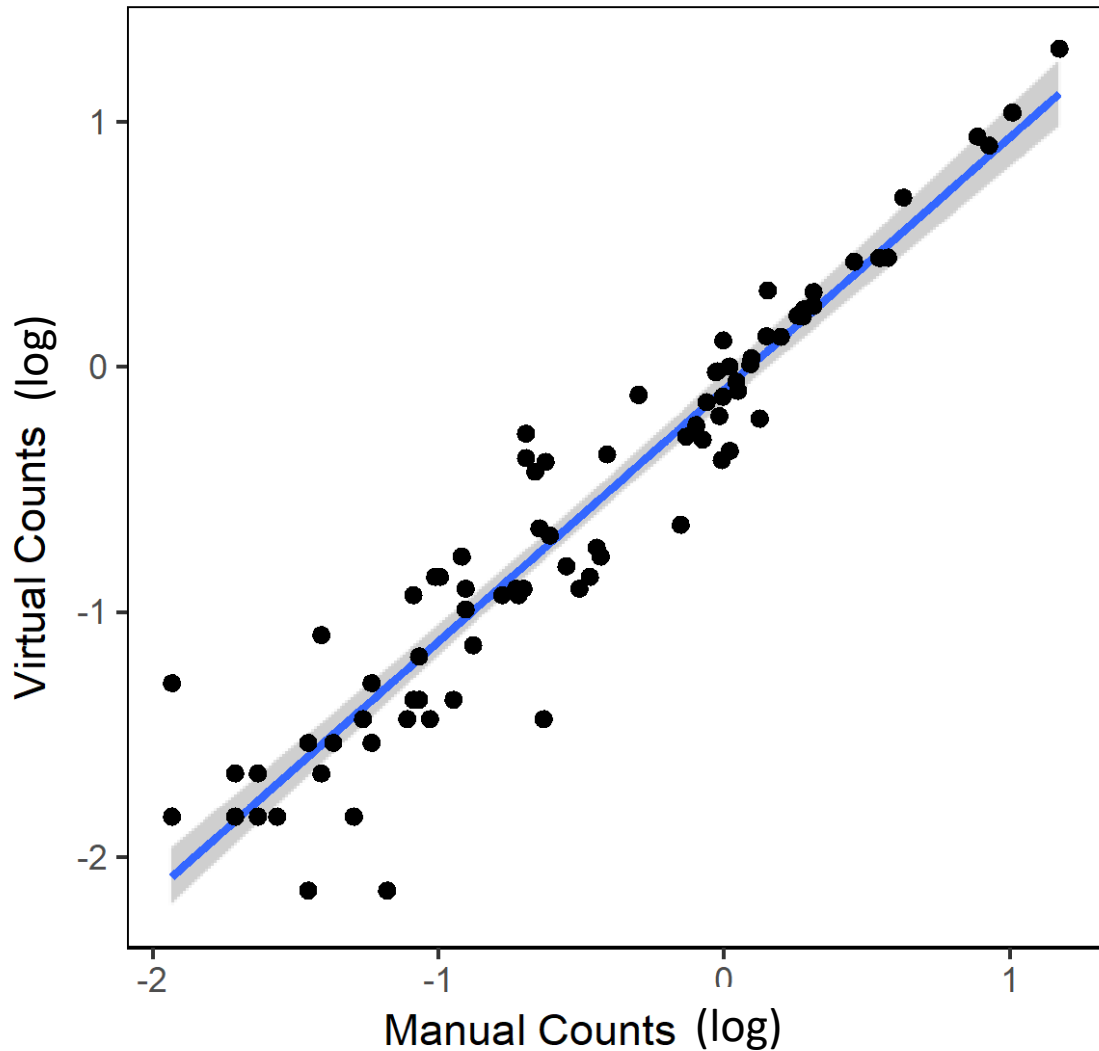
One sample (41 @ 1 μm axial planes) = ~400 GB

VIRTUAL MICROSCOPE POLLEN IDENTIFICATIONS



- PNG images subsampled using a java script
- Slide images read from Matlab script
- Image metadata recorded for each individual pollen grain
 - Slide ID
 - Pollen coordinate
 - Pollen radius
 - 3-letter ID
 - Confidence level (0-9)
- Images and metadata then shared with UC-Irvine Computer Vision Collaborators

COMPARING VIRTUAL AND LIGHT MICROSCOPY COUNTS



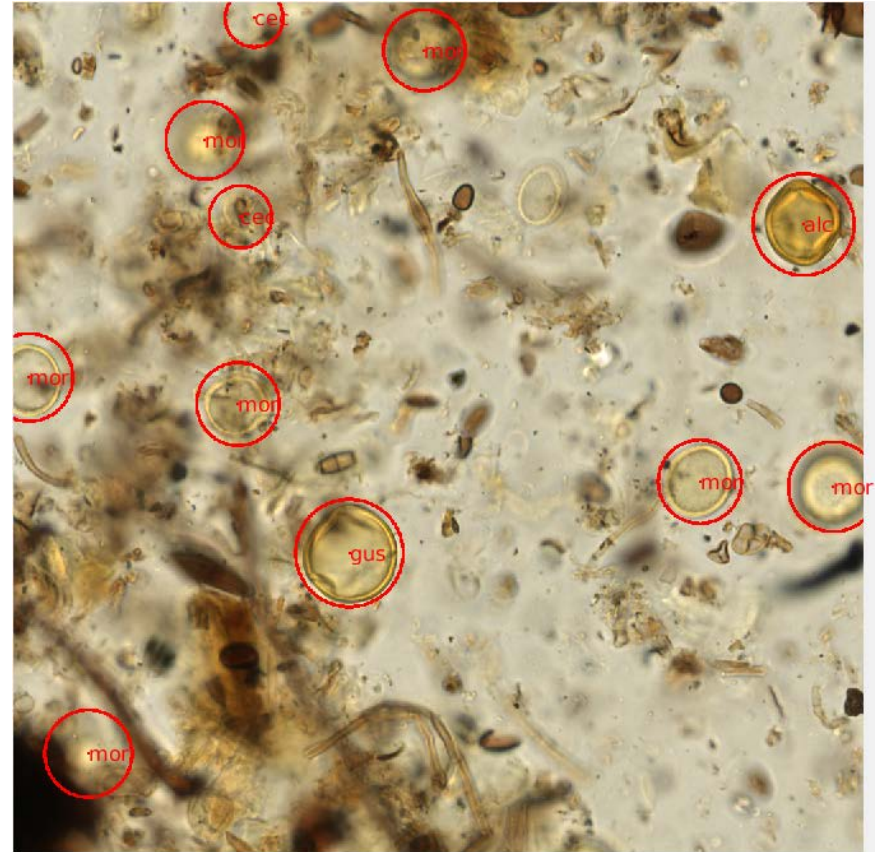
- Testing the fidelity of the virtual microscope using the 10-year Lutz tower record
- Can pollen can be identified at the same taxonomic resolution and frequency?
- $R^2 = 0.97$
- Observed differences are likely reflective of differences in counting strategy:
 - Manual: slide transects
 - Virtual: randomized images

(log)

TRAINING CONVOLUTIONAL NEURAL NETS (CNN) FOR POLLEN IDENTIFICATIONS FROM ANNOTATED IMAGE DATA

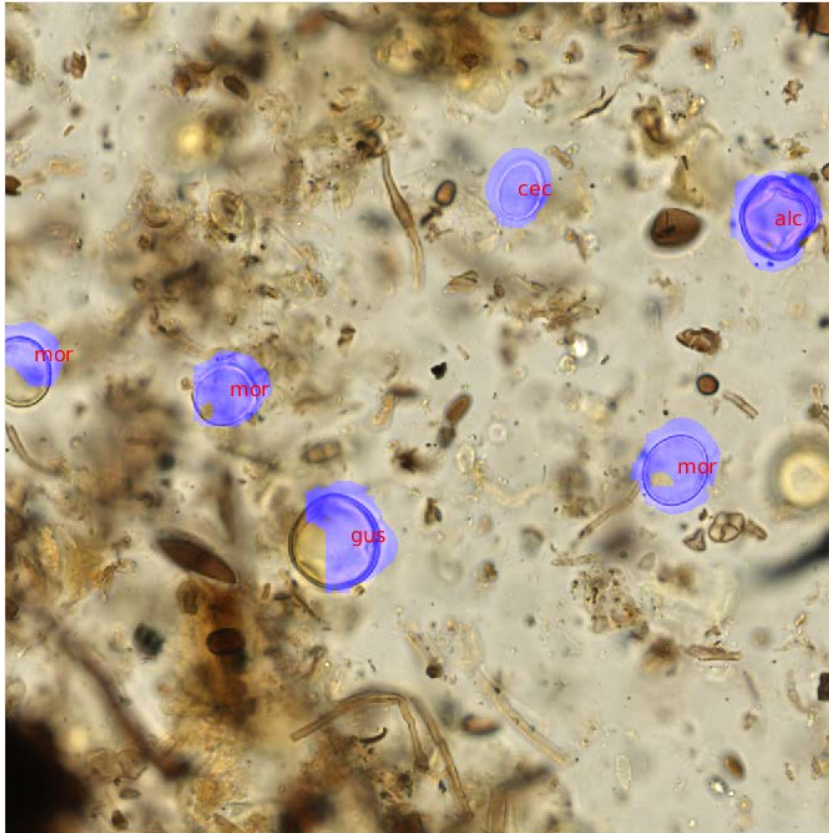
- Images were randomly split into training and testing sets
- Annotated training (ground truth) pollen image examples included the pollen id, location coordinate, and pollen grain radius
- CNN searches each image for patterns corresponding to each pollen id morphology
- Non-maximum suppression was used to identify pollen grains according to pollen ornamentation

Human

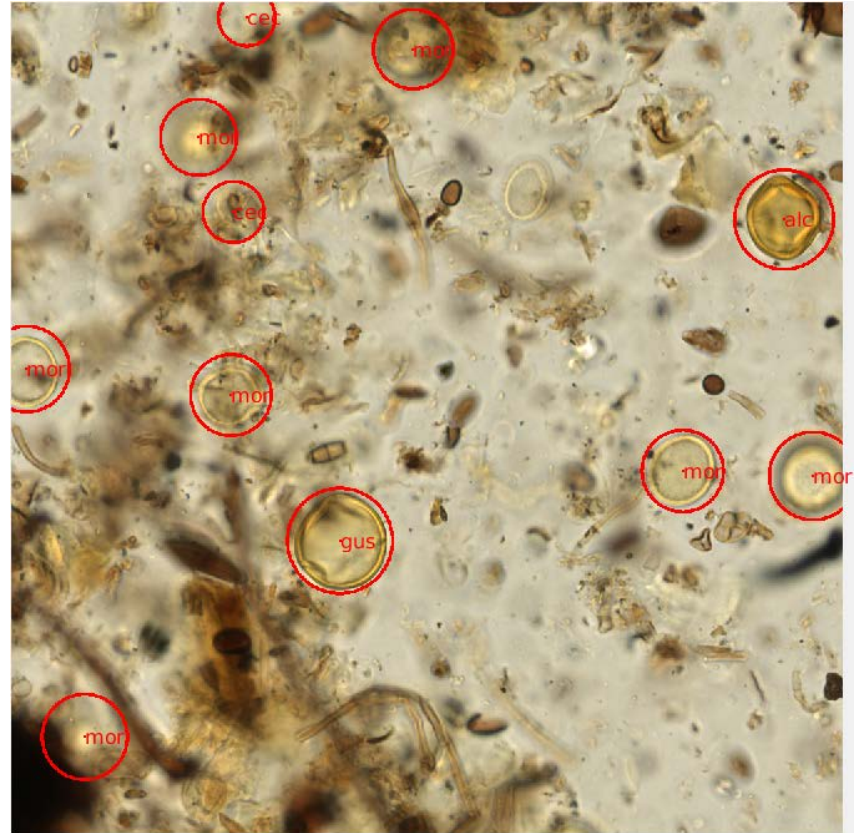


SIMULTANEOUS POLLEN SEGMENTATION AND IDENTIFICATION

Machine



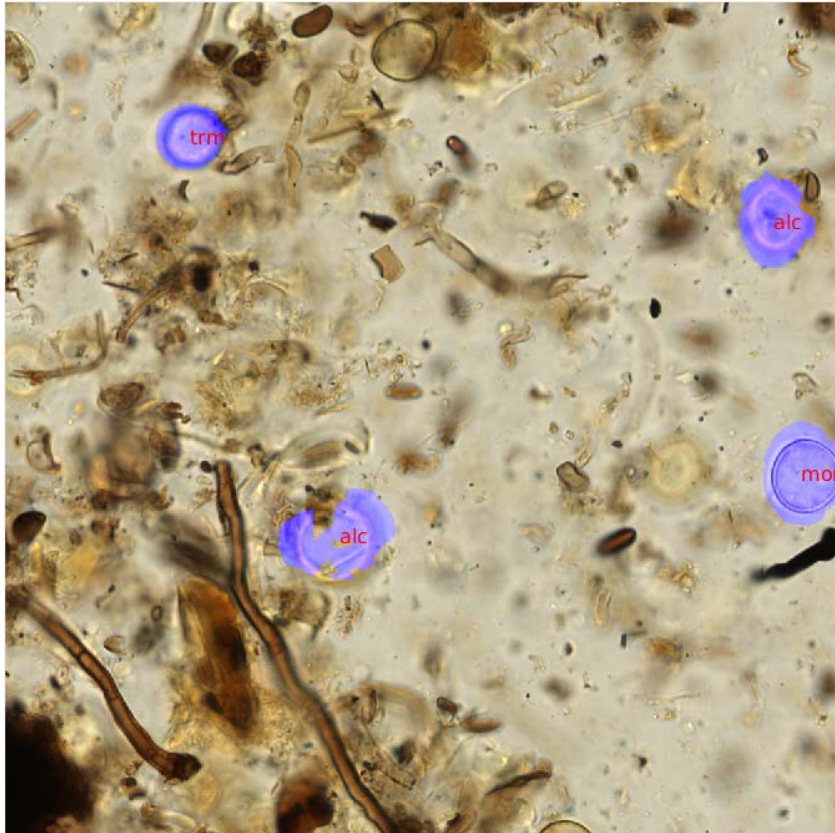
Human



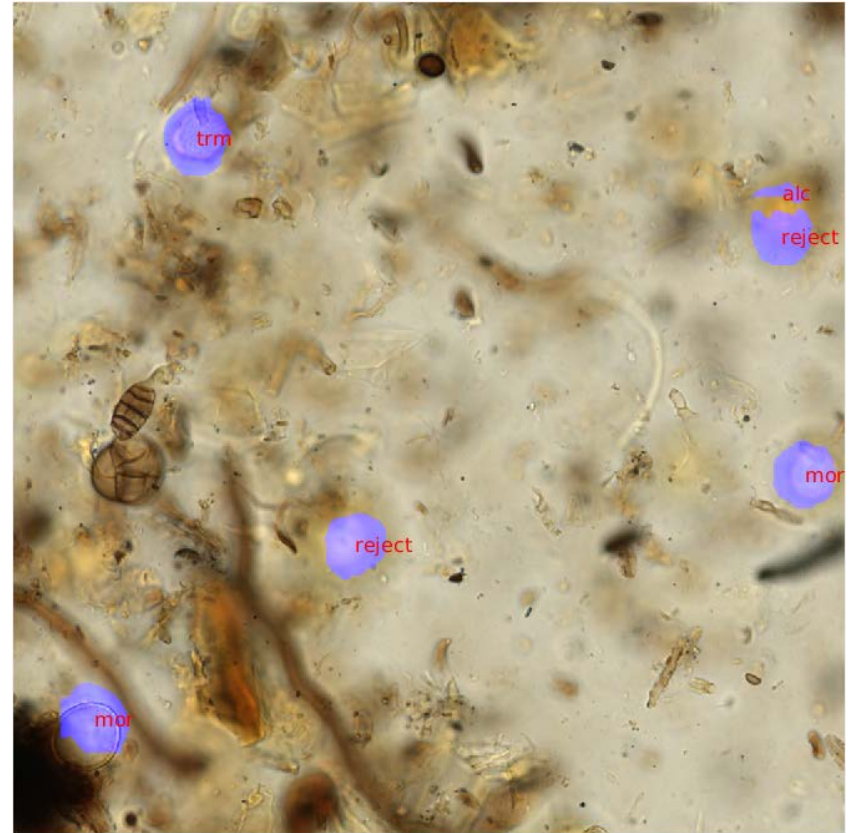
- 48-way classification matrices were constructed using the 47 most abundant pollen types and an additional category called “reject” comprised of pollen types not included in the 47 most abundant

SIMULTANEOUS POLLEN SEGMENTATION AND IDENTIFICATION

Machine



Machine



- 48-way classification matrices were constructed using the 47 most abundant pollen types and an additional category called “reject” comprised of pollen types not included in the 47 most abundant

COMPLETE AUTOMATION: CONFUSION MATRICES

~70% accurate on full 47 pollen type training set, 87.25% on 25 most accurate types

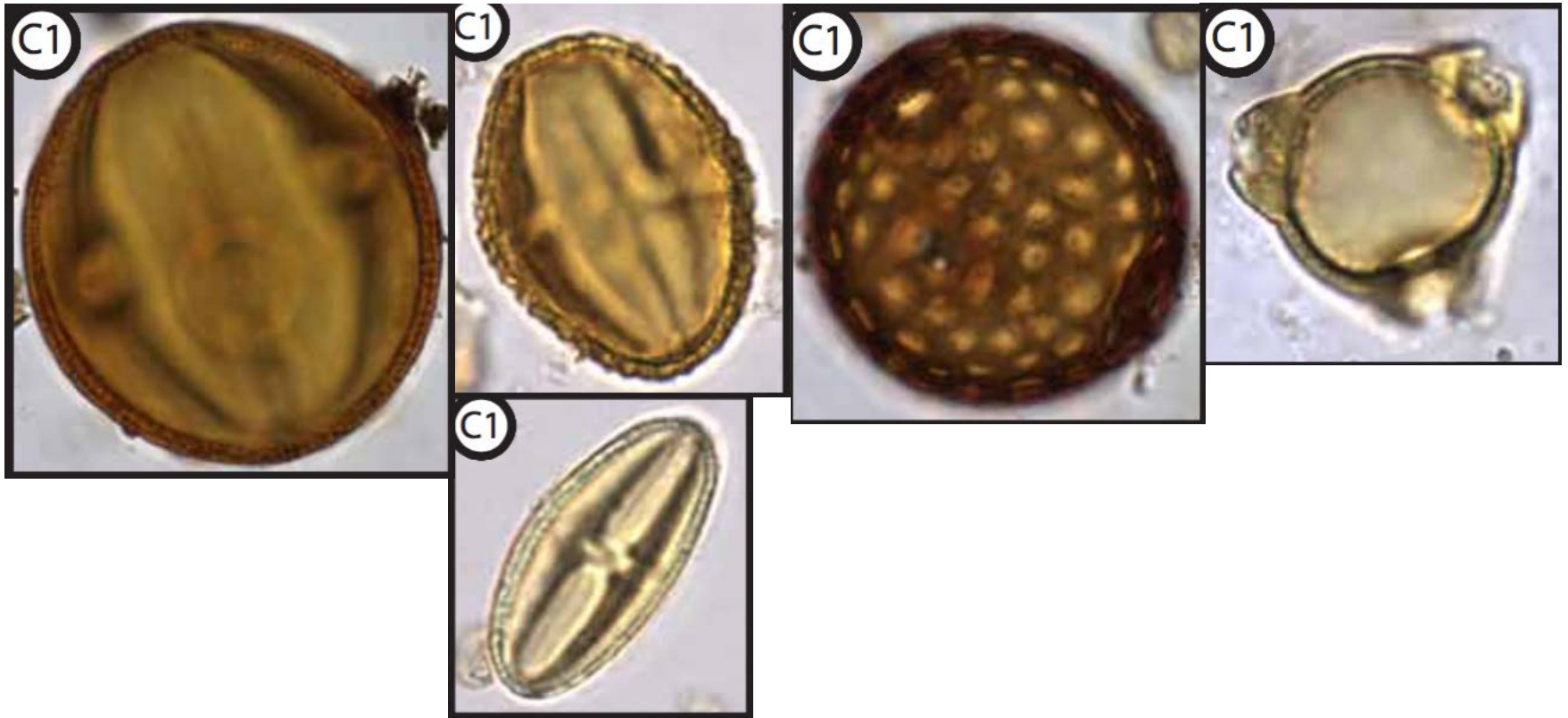
confusion matrix on test set (acc=87.25%)

ground-truth label	als	ant	cas	cec	cor	fic	fra	hir	hyr	lae	lue	lyc	mic	mor	oen	ply	qua	sim	slo	tab	tch	unc	vir	vra	zna	
als	0.77	0.01	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.19
ant	0.08	0.62	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.08
cas	0.00	0.00	0.75	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.04
cec	0.00	0.00	0.00	0.86	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
cor	0.00	0.00	0.00	0.00	0.89	0.00	0.05	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.02
fic	0.00	0.00	0.00	0.05	0.00	0.71	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
fra	0.00	0.00	0.00	0.00	0.00	0.00	0.94	0.00	0.03	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
hir	0.00	0.00	0.00	0.00	0.03	0.00	0.00	0.79	0.03	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
hyr	0.00	0.00	0.01	0.00	0.01	0.00	0.00	0.00	0.95	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.01
lae	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	1.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
lue	0.00	0.00	0.00	0.00	0.07	0.00	0.02	0.00	0.00	0.00	0.91	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
lyc	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.01	0.00	0.01	0.97	0.00	0.00	0.00	0.00	0.00	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.00	0.00
mic	0.00	0.00	0.02	0.01	0.00	0.01	0.00	0.00	0.03	0.00	0.00	0.00	0.83	0.11	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.01	0.00	0.00	0.00
mor	0.00	0.00	0.00	0.10	0.00	0.01	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.88	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
oen	0.00	0.00	0.00	0.00	0.00	0.00	0.01	0.00	0.01	0.00	0.00	0.00	0.00	0.00	0.03	0.87	0.06	0.01	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
ply	0.00	0.00	0.00	0.00	0.00	0.00	0.01	0.00	0.00	0.03	0.01	0.01	0.00	0.00	0.00	0.07	0.87	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
qua	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.01	0.00	0.00	0.00	0.00	0.00	0.99	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
sim	0.02	0.02	0.00	0.02	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.88	0.00	0.00	0.00	0.00	0.00	0.05
slo	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.14	0.00	0.00	0.00	0.00	0.79	0.00	0.00	0.07	0.00	0.00
tab	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.06	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.94	0.00	0.00	0.00	0.00
tch	0.00	0.00	0.15	0.00	0.00	0.00	0.00	0.00	0.02	0.00	0.00	0.00	0.00	0.00	0.00	0.02	0.00	0.00	0.00	0.00	0.00	0.00	0.80	0.00	0.00	0.00
unc	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.06	0.11	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.83	0.00	0.00
vir	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.04	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
vra	0.00	0.00	0.00	0.00	0.00	0.00	0.04	0.00	0.08	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
zna	0.06	0.01	0.01	0.00	0.01	0.00	0.00	0.00	0.03	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00

predicted label

COMPLETE AUTOMATION: CONFUSION MATRICES

>90% accuracy



COMPLETE AUTOMATION: CONFUSION MATRICES

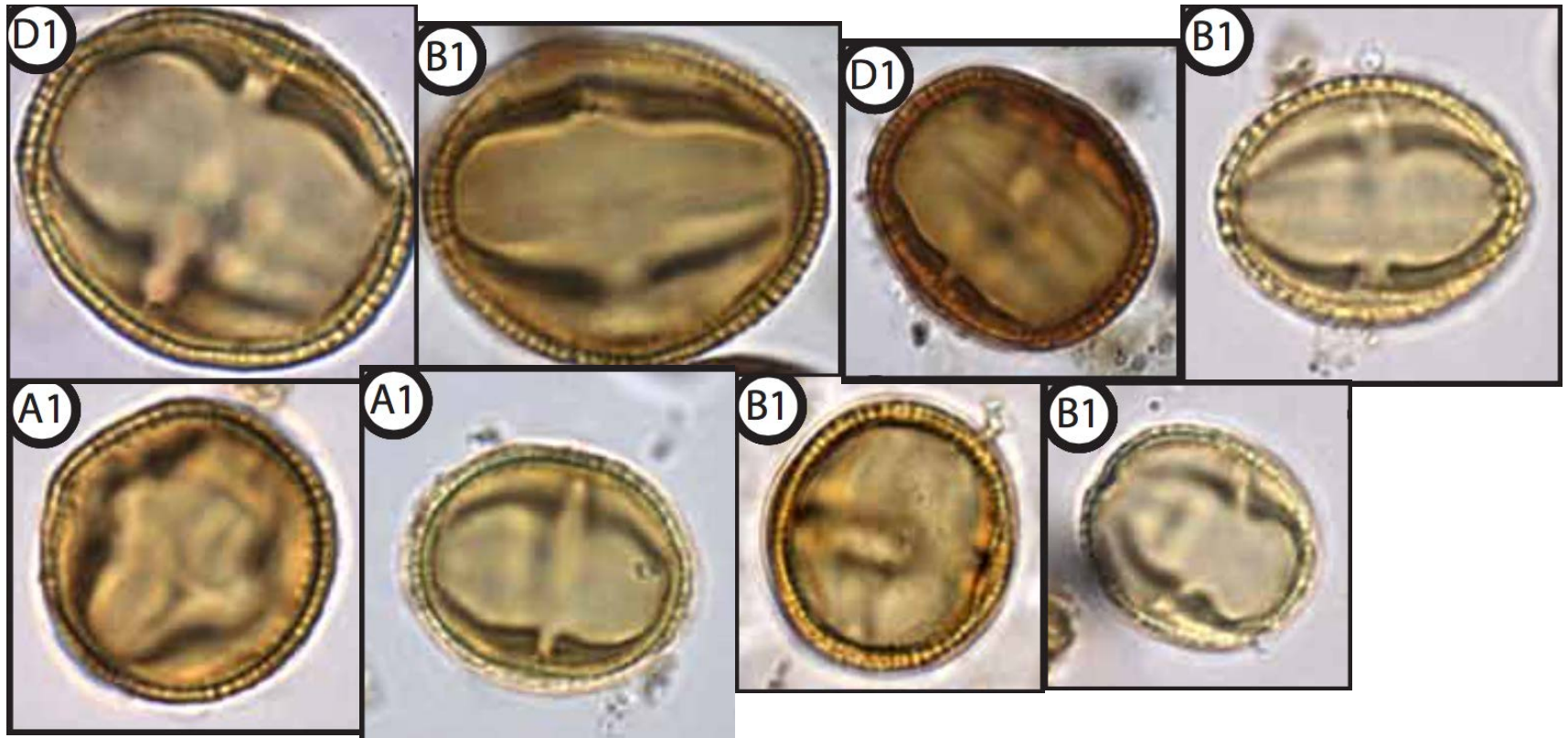
confusion matrix on test set (acc=81.16%)

ground-truth label	alc	als	ant	cas	cec	cor	fic	fra	gus	hir	hvr	lue	lyc	mic	mor	oen	pip	ply	qua	sim	tch	trm	vir	vra	zna
alc	0.78	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.03	0.00	0.03	0.00	0.00	0.04	0.01	0.00	0.00	0.01	0.00	0.00	0.01	0.00	0.00	0.00	0.07
als	0.00	0.82	0.01	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.01	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.16
ant	0.00	0.04	0.54	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.04	0.00	0.00	0.00	0.17	0.00	0.00	0.00	0.00	0.04	0.00	0.00	0.00	0.00	0.17
cas	0.00	0.00	0.00	0.61	0.00	0.00	0.00	0.00	0.00	0.00	0.04	0.00	0.00	0.14	0.00	0.00	0.00	0.00	0.00	0.00	0.14	0.00	0.00	0.00	0.07
cec	0.00	0.00	0.00	0.00	0.86	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.13	0.00	0.01	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
cor	0.05	0.00	0.00	0.00	0.00	0.81	0.00	0.02	0.00	0.00	0.02	0.08	0.00	0.00	0.00	0.00	0.00	0.02	0.00	0.00	0.00	0.00	0.00	0.00	0.02
fic	0.00	0.00	0.00	0.00	0.02	0.00	0.66	0.00	0.00	0.00	0.00	0.00	0.00	0.02	0.29	0.29%	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
fra	0.03	0.00	0.00	0.00	0.00	0.00	0.00	0.89	0.00	0.03	0.00	0.00	0.00	0.03	0.00	0.00	0.00	0.00	0.00	0.00	0.03	0.00	0.00	0.00	0.00
gus	0.13	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.63	0.00	0.08	0.00	0.00	0.05	0.03	0.00	0.00	0.03	0.00	0.00	0.05	0.00	0.00	0.00	0.00
hir	0.00	0.00	0.00	0.00	0.00	0.03	0.00	0.00	0.03	0.79	0.03	0.00	0.00	0.00	0.00	0.00	0.00	0.05	0.05	0.00	0.03	0.00	0.00	0.00	0.00
hvr	0.01	0.00	0.00	0.00	0.00	0.01	0.00	0.00	0.00	0.00	0.93	0.00	0.00	0.02	0.03	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.01
lue	0.00	0.05	0.00	0.00	0.00	0.05	0.00	0.00	0.00	0.00	0.00	0.91	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
lyc	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.01	0.00	0.94	0.00	0.00	0.00	0.00	0.05	0.00	0.00	0.00	0.00	0.00	0.00	0.01
mic	0.02	0.00	0.00	0.00	0.01	0.00	0.00	0.00	0.01	0.00	0.03	0.00	0.00	0.81	0.12	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.02
mor	0.00	0.00	0.00	0.00	0.11	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.01	0.83	0.00	0.00	0.00	0.00	0.00	0.00	0.04	0.00	0.00	0.00
oen	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.01	0.01	0.84	0.00	0.13	0.00	0.00	0.00	0.00	0.00	0.00	0.00
pip	0.00	0.00	0.00	0.00	0.57	0.57%	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.16	0.00	0.27	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
ply	0.00	0.00	0.00	0.00	0.00	0.01	0.00	0.00	0.00	0.01	0.00	0.00	0.03	0.00	0.00	0.04	0.00	0.89	0.01	0.00	0.00	0.00	0.00	0.00	0.00
qua	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.01	0.99	0.00	0.00	0.00	0.00	0.00	0.00	0.00
sim	0.02	0.11	0.00	0.00	0.02	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.02	0.00	0.00	0.00	0.00	0.00	0.70	0.00	0.00	0.00	0.00	0.14
tch	0.09	0.00	0.00	0.11	0.00	0.02	0.00	0.00	0.00	0.02	0.02	0.00	0.00	0.00	0.00	0.00	0.04	0.00	0.00	0.70	0.00	0.00	0.00	0.00	0.00
trm	0.01	0.01	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.50	0.50%	0.00	0.00	0.00	0.00	0.00	0.47	0.00	0.00	0.01
vir	0.00	0.07	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.89	0.04	0.00
vra	0.00	0.04	0.04	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.12	0.04	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.04	0.64	0.08
zna	0.03	0.10	0.01	0.00	0.00	0.00	0.00	0.01	0.00	0.00	0.01	0.00	0.00	0.00	0.03	0.00	0.00	0.00	0.00	0.00	0.00	0.01	0.00	0.00	0.81

predicted label

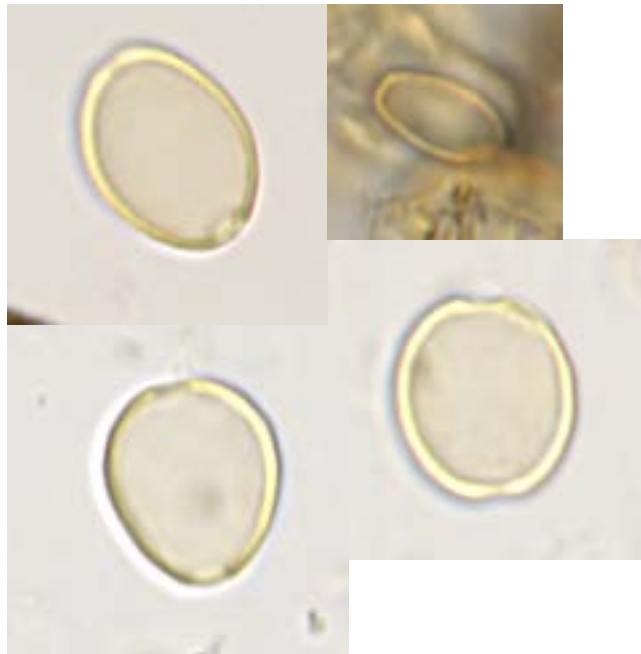
COMPLETE AUTOMATION: CONFUSION MATRICES

<50% accuracy

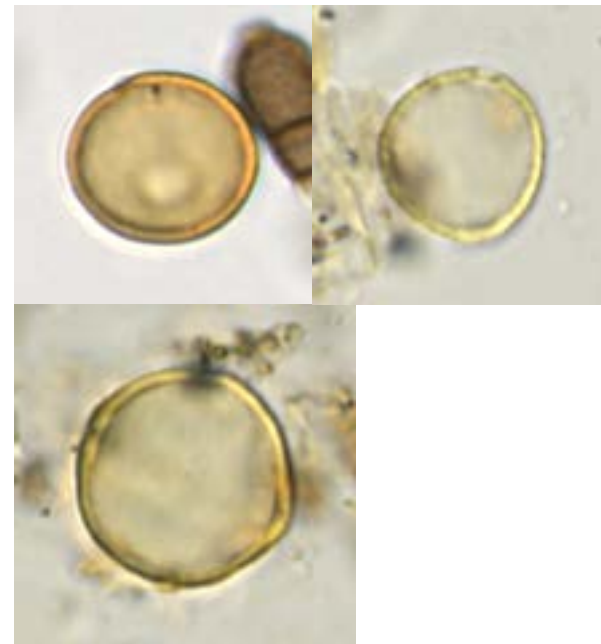


THE PAIRWISE COMPARISONS IN THE CONFUSION MATRIX SHOWING THE MOST DISAGREEMENT ARE MORPHOLOGICALLY VERY SIMILAR

Ficus (66% accuracy) was misclassified 29% of the time as *Brosinum*-type



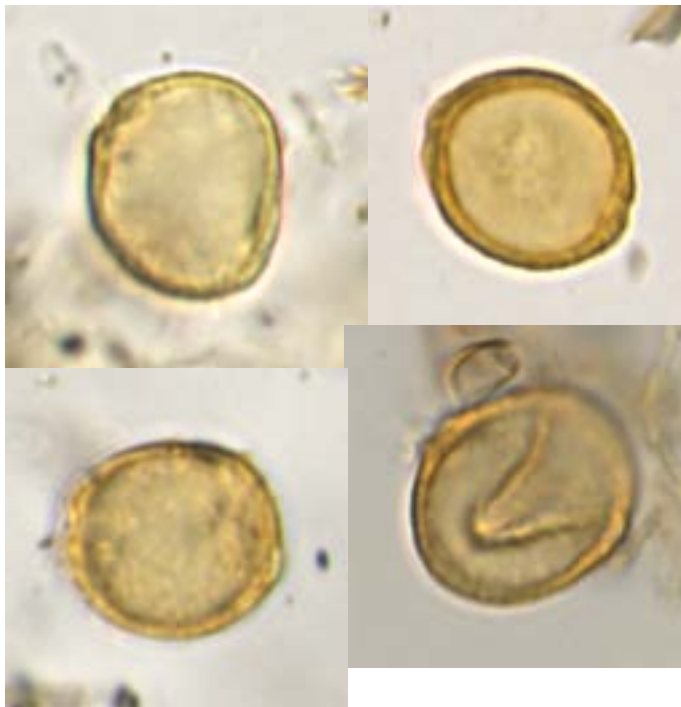
Ficus (Moraceae)



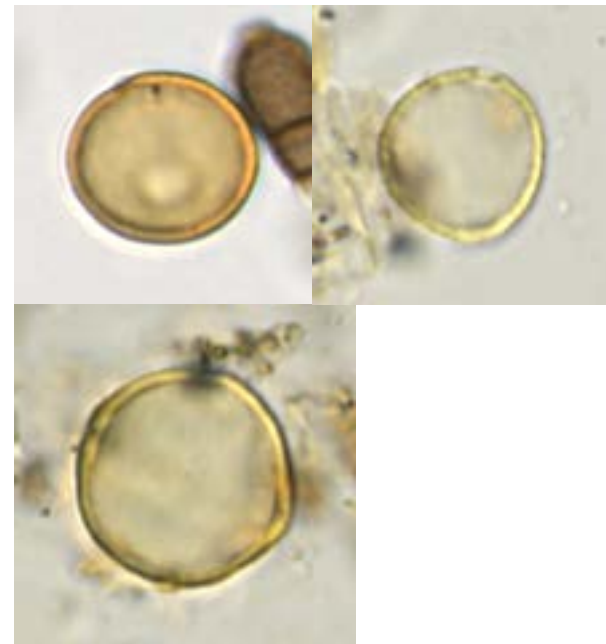
Brosinum-type(Moraceae)

THE PAIRWISE COMPARISONS IN THE CONFUSION MATRIX SHOWING THE MOST DISAGREEMENT ARE MORPHOLOGICALLY VERY SIMILAR

Trema (47% accuracy) was misclassified 50% of the times as *Brosinum*-type



Trema (Ulmaceae)



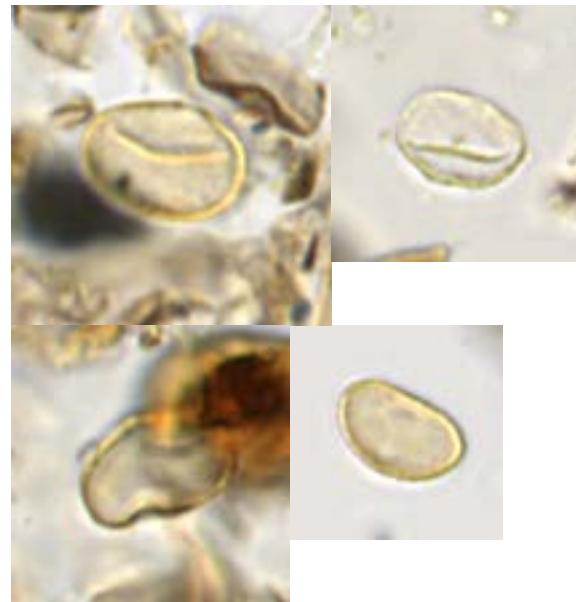
Brosinum-type (Moraceae)

THE PAIRWISE COMPARISONS IN THE CONFUSION MATRIX SHOWING THE MOST DISAGREEMENT ARE MORPHOLOGICALLY VERY SIMILAR

Piperaceae (**27% accuracy**) was misclassified **57%** of the times as *Cecropia*



Piperaceae



Cecropia (Urticaceae)



CONCLUSIONS

- Overall, our results are very promising given this difficult classification problem
- Our preliminary results show that automated segmentation and classification models can distinguish pollen types from hyper-diverse samples
- Model performance is poorest on pollen types that are morphologically very similar
- Predicted outputs should improve as the neural nets are trained on more tagged examples



FUTURE DIRECTIONS

- The same system can be implemented using training data from herbarium and reference material collections
- Apply the methodology to fossil records
- Create a collaborative pollen identification database that harnesses the expertise of multiple palynologists
- Expand to other proxies
 - Diatoms
 - Phytoliths
 - Cuticles



ACKNOWLEDGEMENTS

NSF Macrosystems Ecology
NSF Advances in Biological Informatics

