Plant Cell Lineage
Specification based on the
Interactive Visualization
of Hybrid 3D and 2D Data
and with the Support
of Machine Learning

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NAVISCOPE

















Collaboration









Alain Trubuil









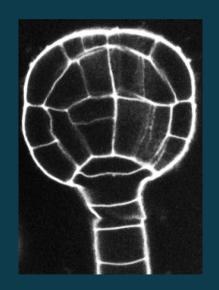


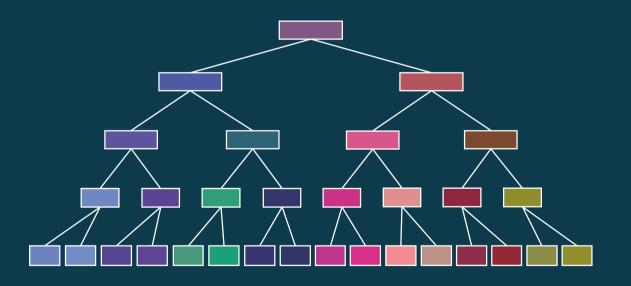






Plant Cell Division











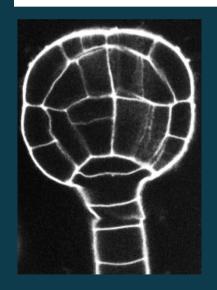


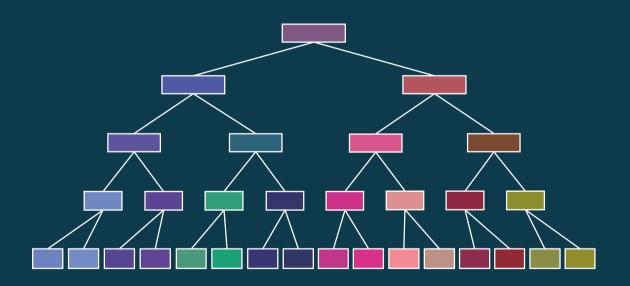




Plant Cell Lineage

For a specific embryo, what do biologists do to build the hierarchy?







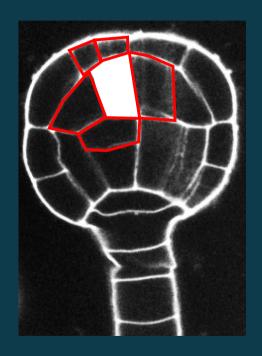












Biologists need to find the right sister cell for every cell in an embryo.















Once decided, they would merge cells and continue assigning the remaining cells.



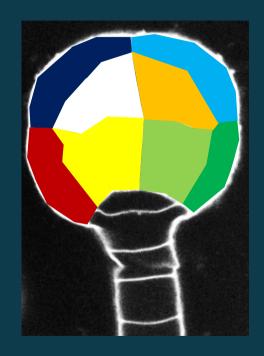




























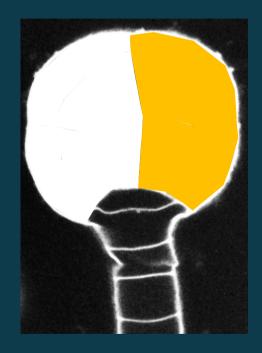














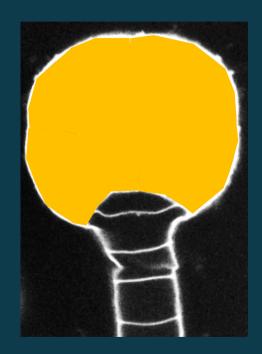
















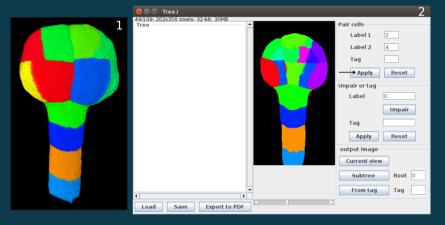


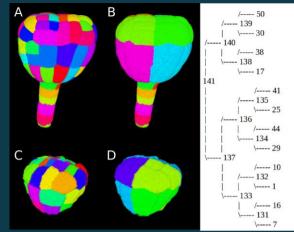


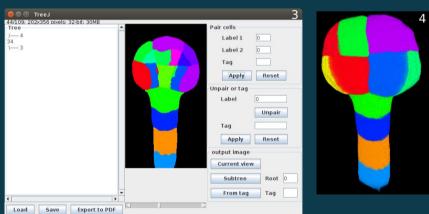




Traditional Tools











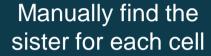






Traditional Workflow

Get segmented 2D slices











Check every cell

Write down the hierarchy on paper













Traditional Workflow

Manually find the Get segmented Interaction Visualization sister for each cell 2D slices Visualization Visualization Check every cell Write down the hierarchy on paper







ML

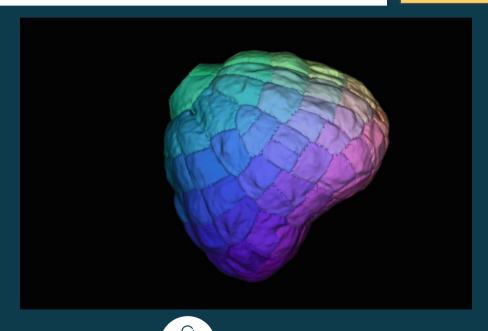




Problem with Plant Cell Embryos

Cells are densely packed together.

Interaction







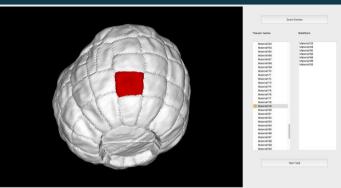


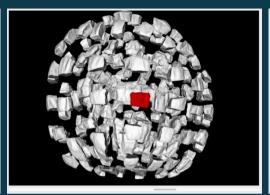


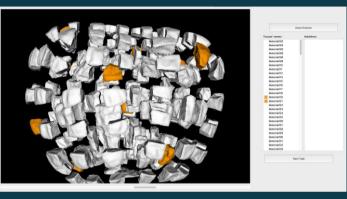




Exploration







1D

Select from the list

3D

Select from the explosion view

1D + 3D

Select from both views











Study Results

- The explosion technique makes it possible to access objects in such a tightly packed 3D environment.
- Combination (List + Explosion) seems to combine these advantages of the single technique.













Traditional Workflow

Manually find the Get segmented Interaction Visualization sister for each cell 2D slices Visualization Visualization Check every cell Write down the hierarchy on paper



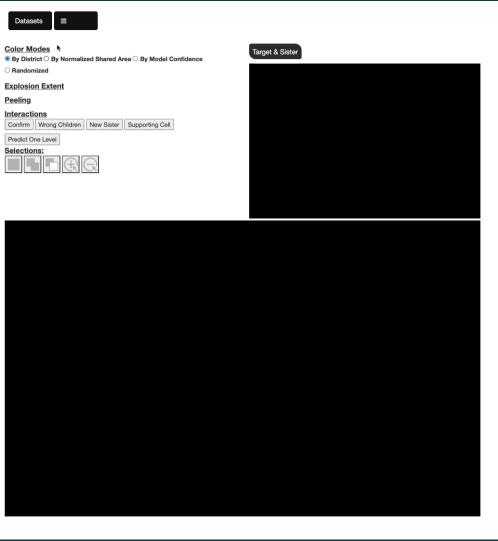




ML

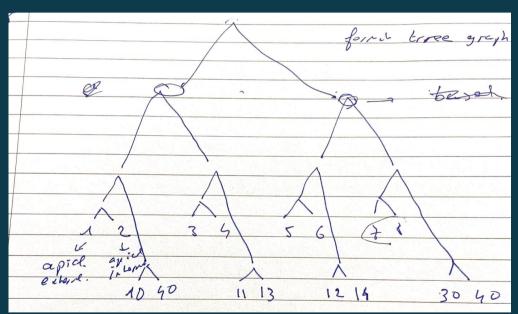


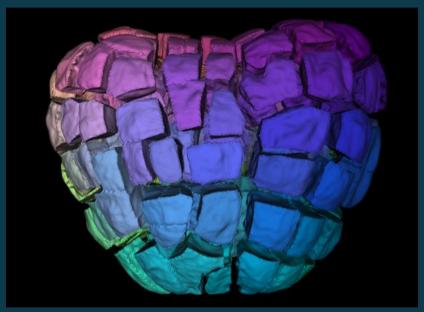




Interactions Confirm Wrong Children New Sister Supporting Cell Predict One Level

Traditional Hand-written Tree









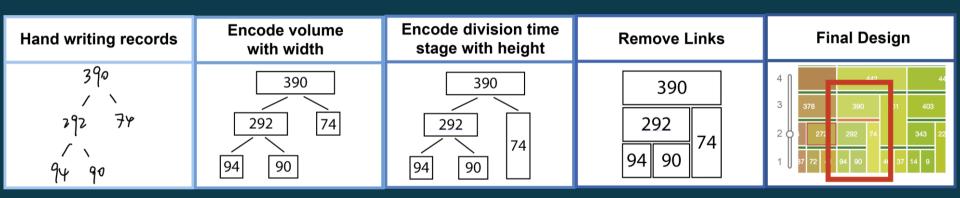








2D Abstract Hierarchy







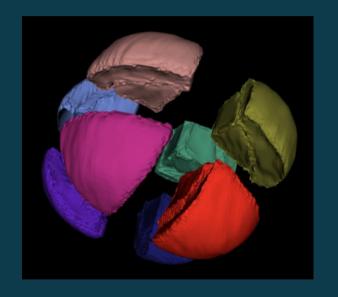






Visually and Interactively Connected









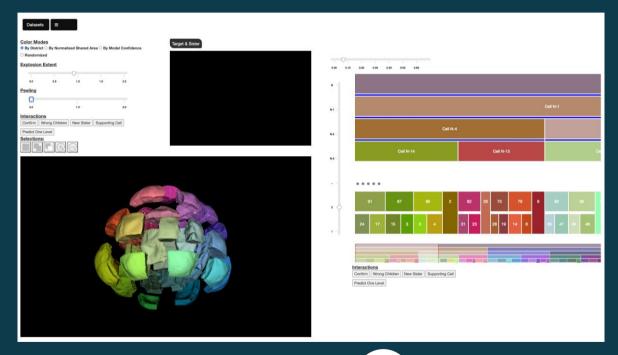








Visually and Interactively Connected







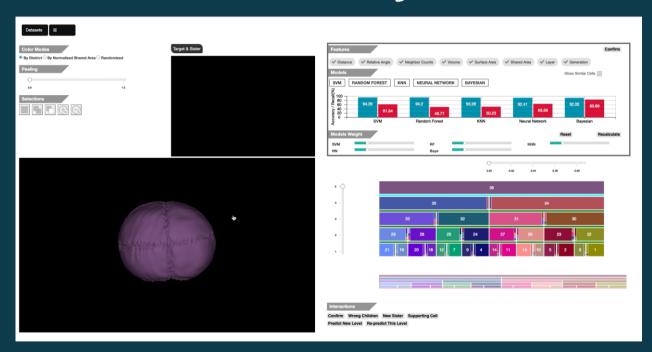








Visually and Interactively Connected















Advanced Workflow



Two-direction of Hierarchy Building

Enable biologists to build the hierarchy tree from both top-down and bottom-up approaches.

Machine Learning Predictions

Use ML to predict a single level as a basis for biologists to check and correct.









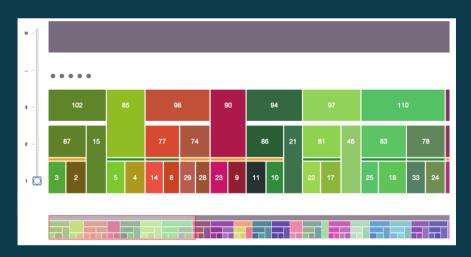


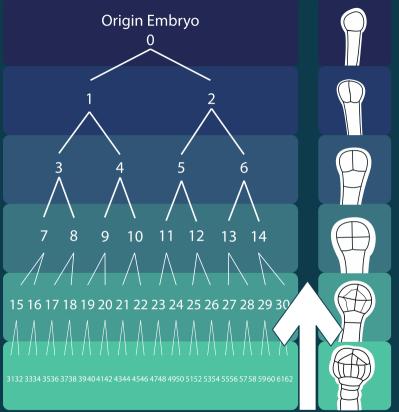




Bottom-up Approach

The traditional way to build the hierarchy.











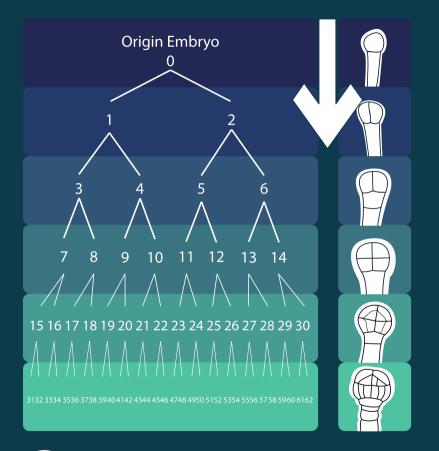




Top-down Approach

Biologists have ideas about how the embryo could be divided in the beginning.







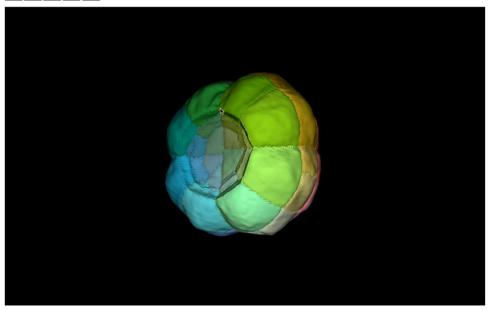


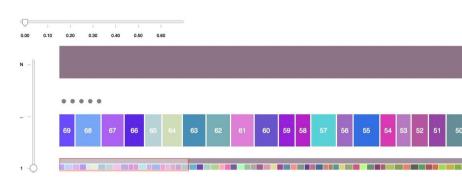












Interactions

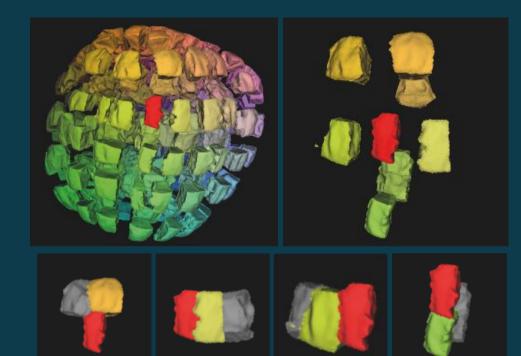
Confirm Wrong Children New Sister Supporting Cell

Predict One Level

Machine Learning Model

Binary Classification Problem: Sisters (1) / Non-sisters (0)

- Sequential Neural Network
- 93 Embryo datasets
- 47132 pairs
- 12 features





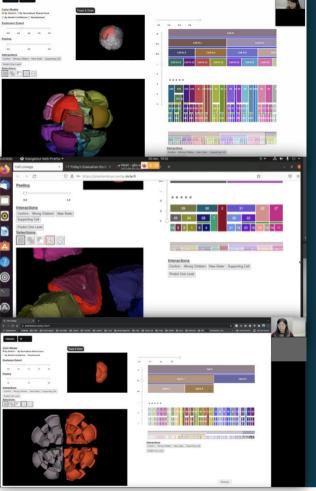






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Evaluation Study

- Biologists took time to learn the functionalities of LineageD.
- They used both the 2D hierarchy and 3D views.
- They valued the visual representations in 2D and 3D.
- They thought LineageD helps to understand the embryos' development better.













ML Results not satisfying

How could we help biologists to have a better collaboration with ML in the case where the training datasets are limited?













Our Solution – Human-Al Teaming



Make use of machine learning

We need to try our best to improve ML performances.



Allow people to control the results

Human beings should have full control over the final decisions.



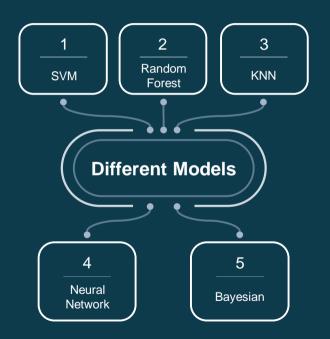








Model Training







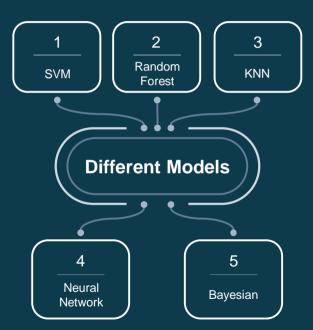


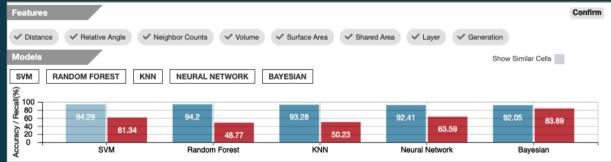






Model Training





Accuracy

Recall











The overall predictions for each cell

Models Weight

The visualization of model prediction confidence



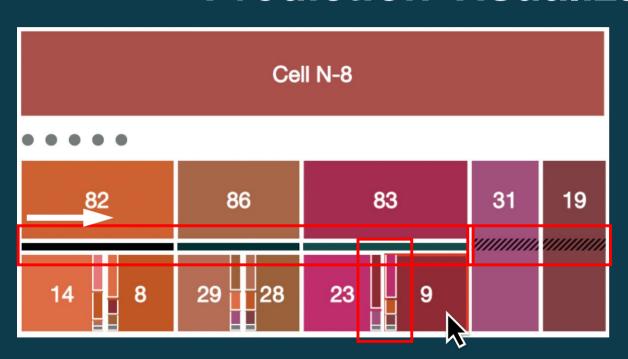












Combining the prediction results from five models, we visualized them with stacked bar charts on each node.

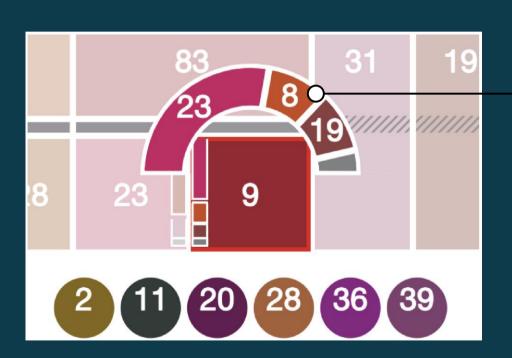












All predicted sisters

Vertical thumbnail of all predictions

All the other neighboring cells

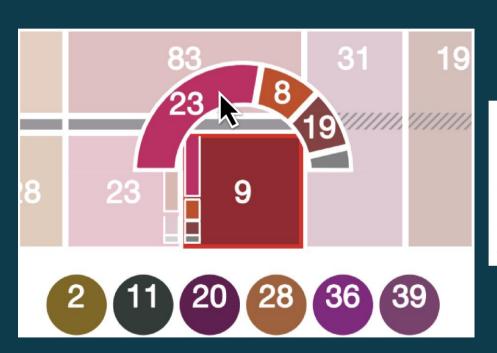












neuralNetwork 23
knn
bayesian 8 19
svm 23
randomForest 23

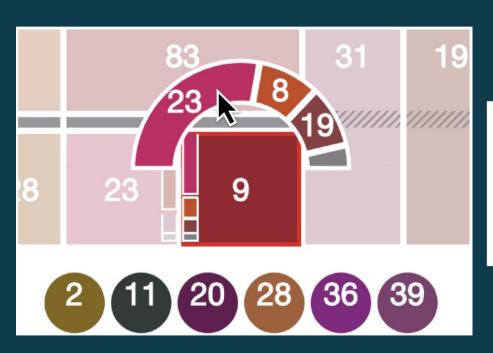












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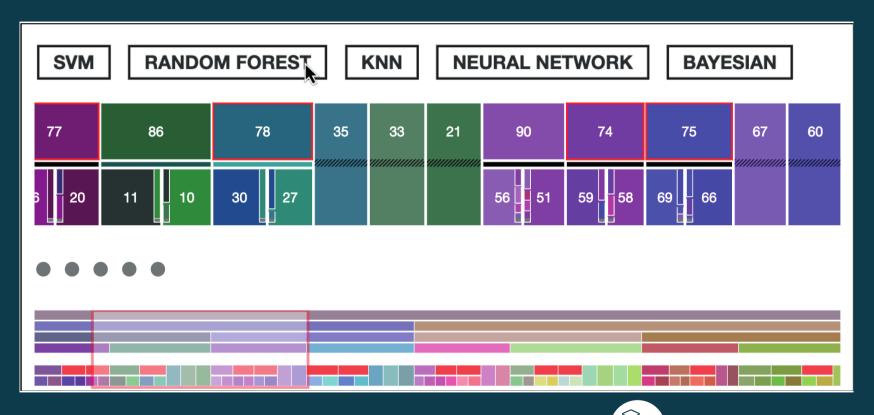








Individual Model Prediction



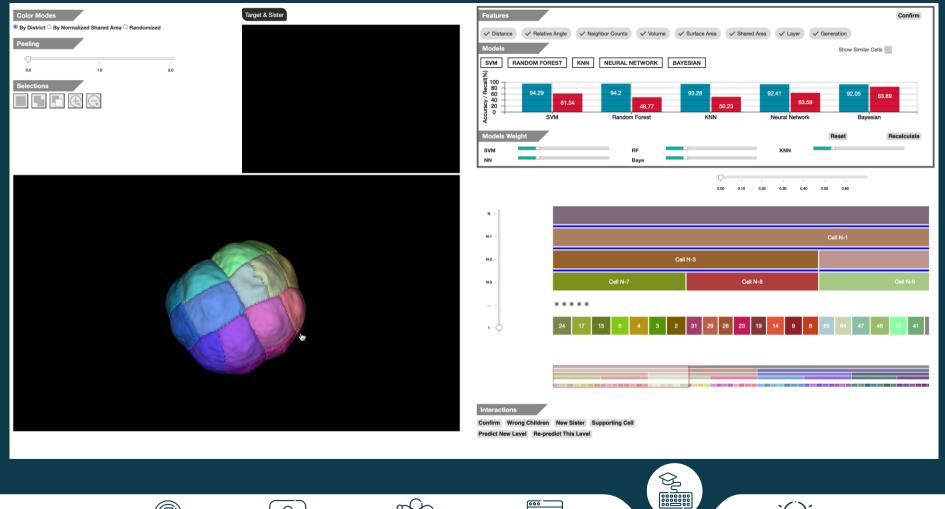
























Evaluation Study

- Biologists appreciated the prediction results and their visualization.
- They thought LineageD+ could help save time and change the traditional approach they used in the assignment process.
- One biologist expressed that interacting with ML made her feel like she was discussing with the computer in making decisions.











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Thèse soutenue à Paris-Saclay, le 14 février 2023, par

Jiayi Hong





















