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# Using the past to restore the future: Quantifying historical vegetation to assist in tidal freshwater wetland restoration Former Lake Charles at the VCU Rice Rivers Center

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# Using the past to restore the future: Quantifying historical vegetation to assist in tidal freshwater wetland restoration

## Former Lake Charles at the VCU Rice Rivers Center



Christopher D. Gatens and Edward R. Crawford, Ph.D.



### Background

Wetlands have been providing humans with critical natural ecosystem services throughout our time on Earth. Nevertheless, these invaluable ecosystems have been habitually altered as a cost of human progression. Two of the most common alterations to wetlands are hydrologic, in the form of damming, and filling. Both occurred along Kimages Creek in Charles City County, VA during the 19<sup>th</sup> and 20<sup>th</sup> centuries. In 2010 the Lake Charles dam was partially removed, restoring the creek's tidal communication with the James River and beginning tidal forested freshwater wetland restoration. Upon the recession of the body of water, numerous woody stumps were revealed.

#### Objectives:

We studied these stump remnants in an attempt to assess the spatial structure and vegetative community of this forested freshwater tidal wetland before perturbation.



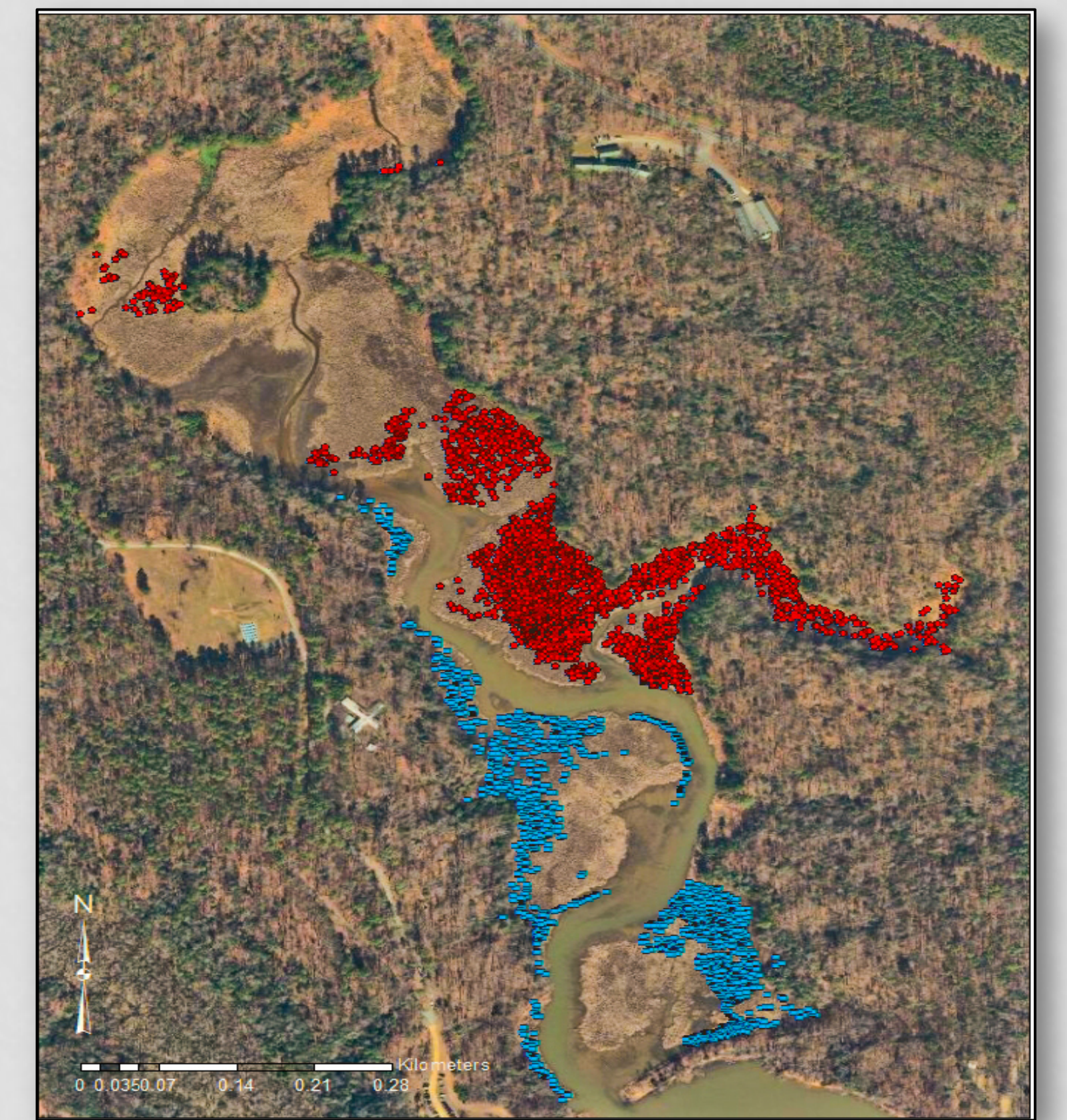
Picture of the Kimages Creek wetland under Union occupation during the Civil War (Egghart 2009)



Present day photo of the Kimages Creek wetland. Photo credit: Rick Ward

### Results/Discussion

During this ongoing study, over 4,500 stumps have been geo-located and 413 samples have been processed. There were 11 unique genera identified, among which 15 were identified to the species. The most abundant genus of trees was *Fraxinus* spp. with a relative density of 73.24%, and the next most abundant was *Carya* spp. with a relative density of 11.79%. The remaining samples were comprised of small densities of various species. The majority of the samples were of obligate or facultative wetland species (63.1%).



Geospatial Points:  
n~4,546

Stump "cookies":  
413 total  
258 identified  
156 age-identified

Historical Forest Analysis (to date):  
11 genera; 15 species  
Most abundant: *Fraxinus* spp. (RD=73%)  
Age range: 10-102 yr

We will soon compile the geospatial coordinates onto a GIS map and use the species data to better understand the native community. Recreating the natural historical vegetative community could help guide current restoration efforts in other locations in other mid-Atlantic formally impounded wetlands. Ultimately our goal is to be able to build a functioning virtual wetland model.

### Methods

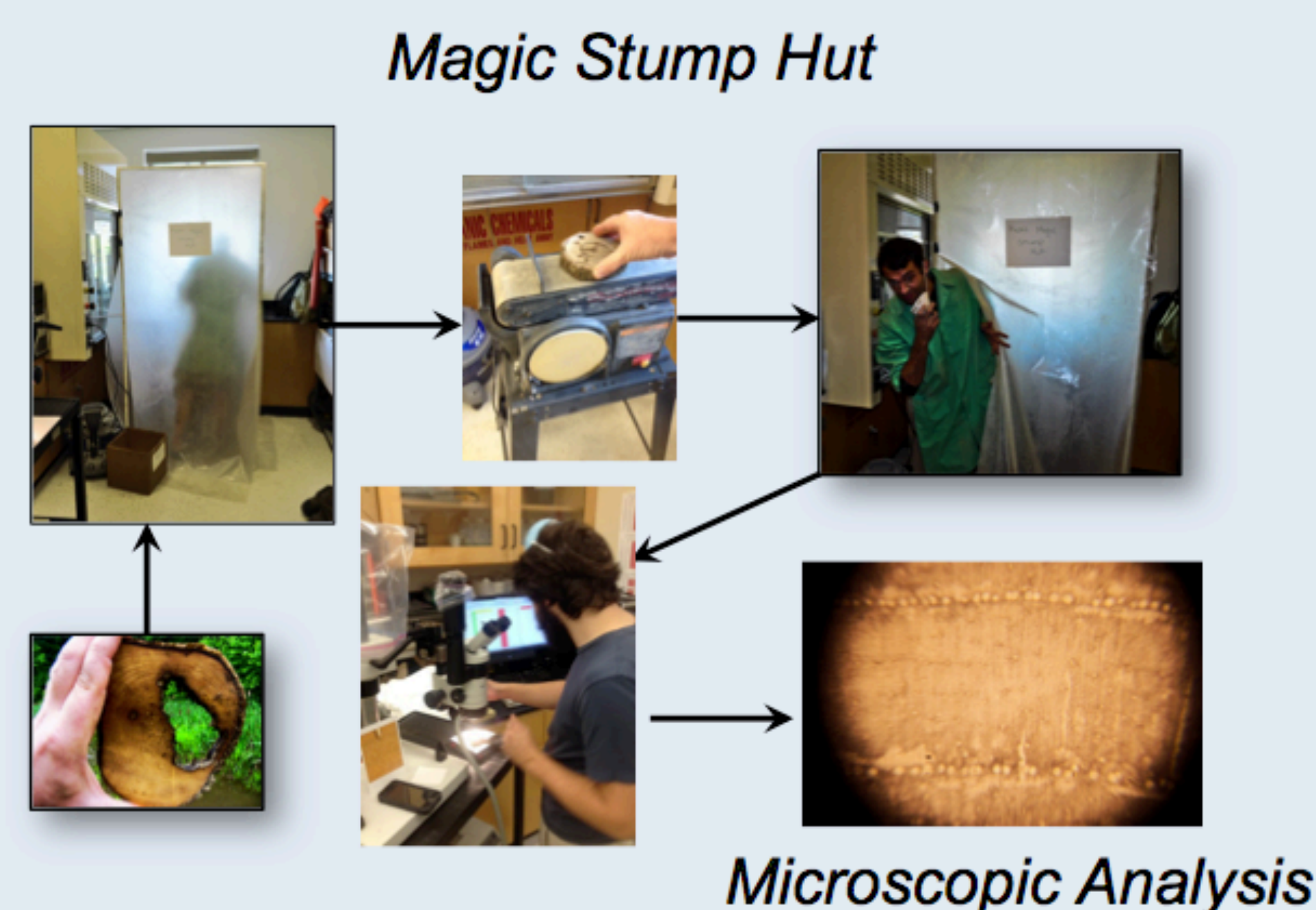
#### Obtain geospatial points



Obtain stump "Cookie"

1) We began by obtaining the geospatial location of each discovered stump with a handheld GPS unit. Every 10 stumps we took a cross-sectional wood sample using a handsaw (a stump "cookie").

#### "Cookie" Processing & Analyses

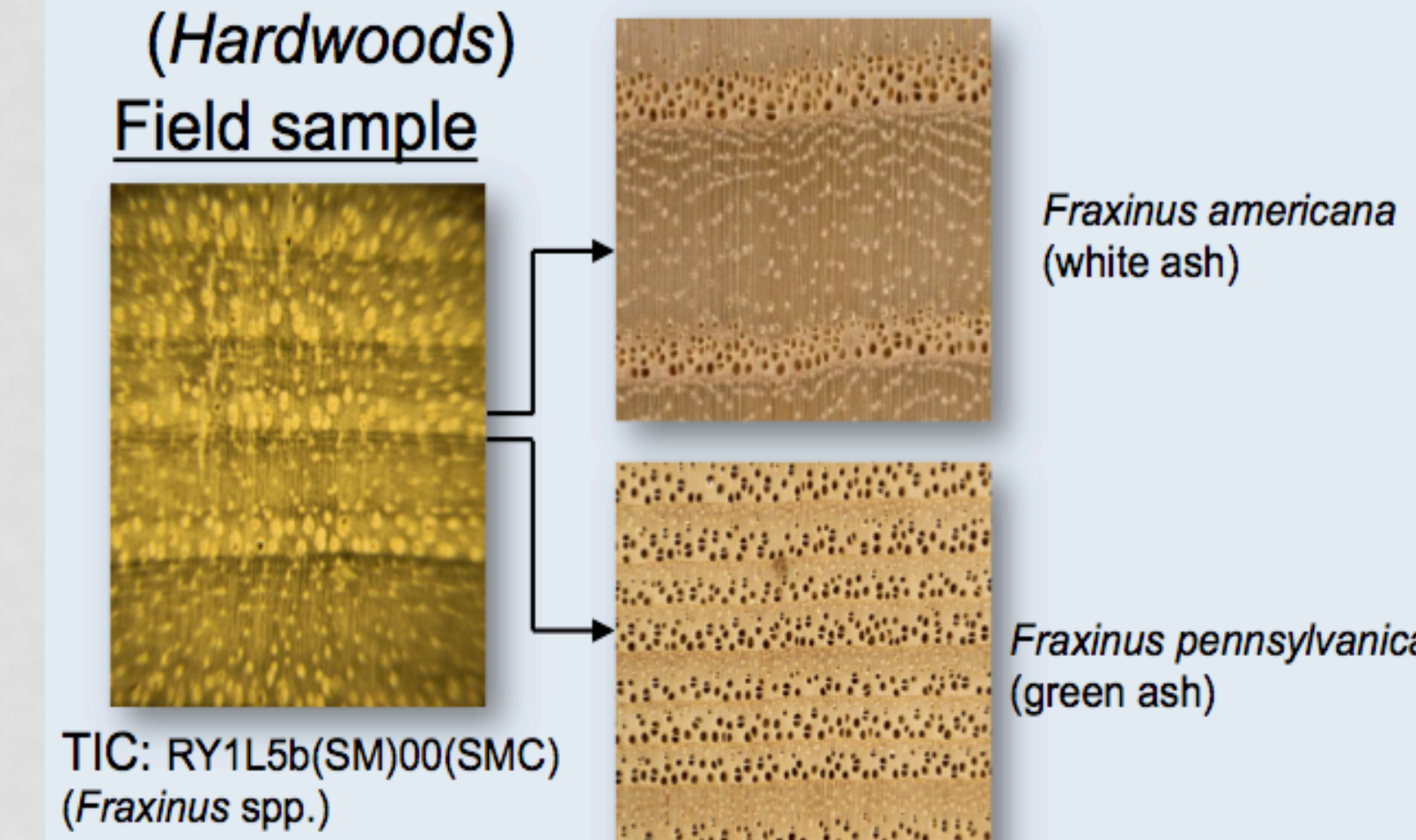


2) In the lab we first sanded the stumps for easier microscopic analysis. We then used microscopy and a Taxonomic Identification Code process. This process yields a unique alphanumeric code (a TIC), each of which corresponds to a certain species or genus of tree.

#### 3) The Taxonomic Identification Process

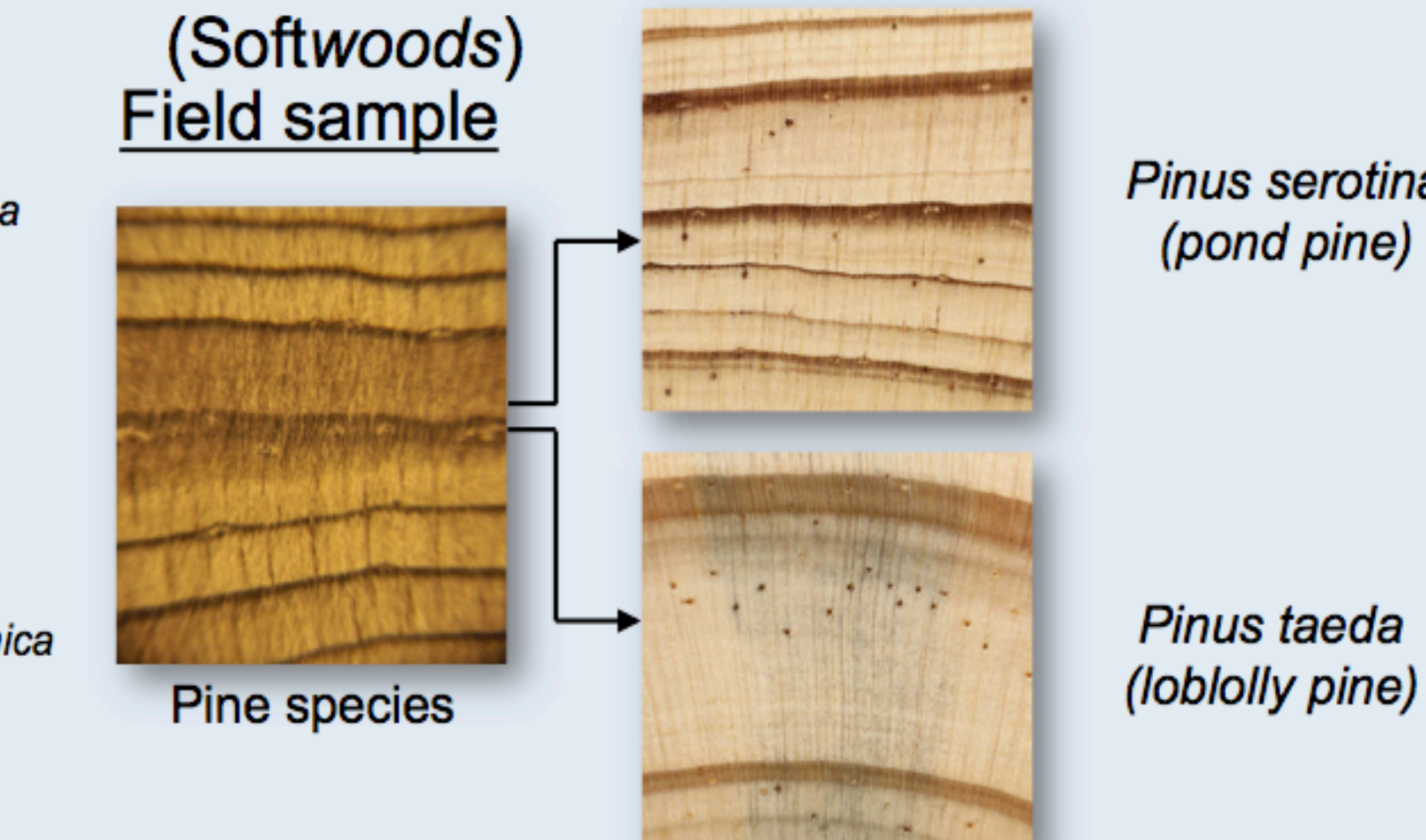
Diagnostic characteristics		TIC	Latewood Diagnostics		TIC
Step 1	Ring-porous	R	Step 7	Width between uniseriate rays > average latewood pore width	G
	Semi-ring-porous	S		Width between uniseriate rays = average latewood pore width	E
	Diffuse-porous	D		Width between uniseriate rays < average latewood pore width	L
Step 2	Visible (>10X) growth rings	Y	Step 8	Even distribution	1
	Microscopic (>10X) growth rings	N		Uneven distribution	2
Step 3	Uniseriate rays	1	Step 9	Solitary pores	S
	Multiseriate rays	2		Multiple pores	M
Step 4	Distinct uniseriate rays (<10X)	E		Chain pores	C
	Indistinct uniseriate rays (>10X)	L		Nested pores	N
	Microscopic uniseriate rays (>10X)	M		Wavy pores (irregular)	W
<b>Earlywood Diagnostics</b>					
Step 5	Minuscule pores, single row, intermittently spaced	1	Step 10	Continuous "lines" run parallel to growth rings connecting pores and uniseriate rays	3
	Small pores, 2 to 4 pores wide, adjacently arranged	2		Intermittent horizontal/wavy lines connecting latewood pores	4
	Large pores, 1 - 2 pores wide, intermittently spaced	3		Two previous characteristics are absent	5
	Large pores, 1 - 2 pores wide, adjacently arranged	4	Step 11	Growth rings are defined by a continuous "line" of pores	P
	Large pores, 2 - 4 pores wide, adjacently arranged	5a		Growth rings are definable bands of parenchyma cells	L
	Large pores, 2 - 4 pores wide, intermittently spaced	5b		Growth rings are neither composed of pores or definable bands of parenchyma cells	O
	Large pores, 3 - 8 pores in width	6			
Step 6	Solitary pores	5			
	Multiple pores	M			

#### Cookie Analyses (Hardwoods)



TIC: RY1L5b(SM)00(SMC) (*Fraxinus* spp.)

#### Cookie Analyses (Softwoods)



### Works Cited

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