



Workflows and Personnel

Gil Nelson

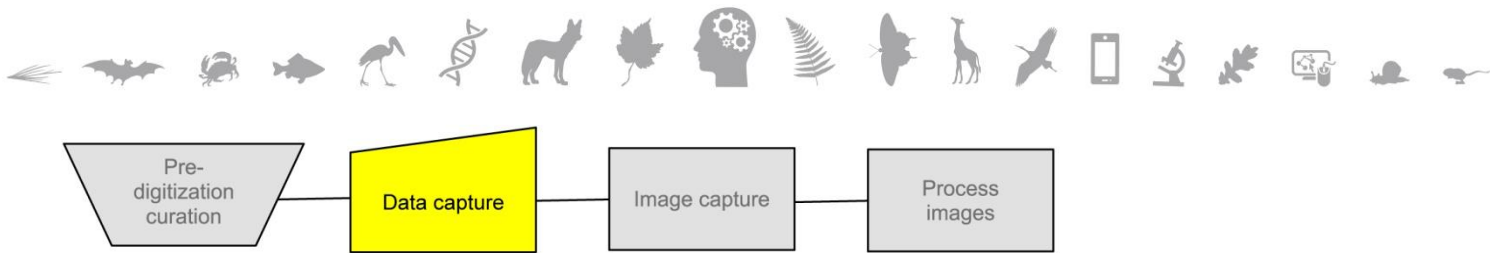
Institute for Digital Information and Scientific Communication
Integrated Digitized Biocollections
Florida State University

Bristol, England

9 March 2018



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Guiding Principles

Follow a modular approach

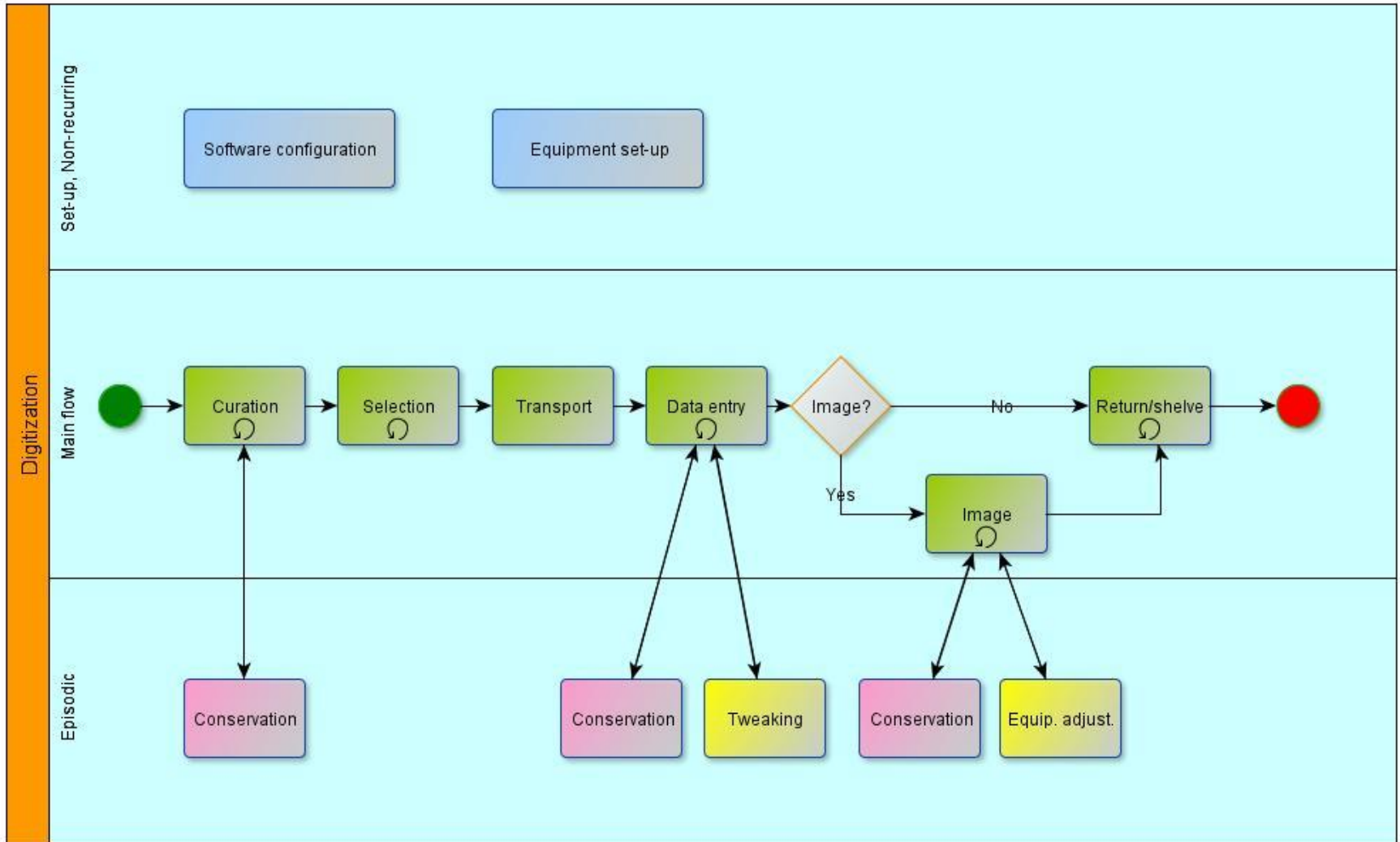
- “Plug and play” modules are preferred.
- Simple modules involving a limited number of tasks are easier to troubleshoot and maintain.
- Divide large modules into sub-modules.
- Modules are generally self-contained but tangential.
- There is no consensus workflow, virtually all workflows are customized.

Assign roles deliberately

- Adjust to strengths of each technician--using students and volunteers requires flexibility in role assigned to personnel rather personnel assigned to role.

Create task lists

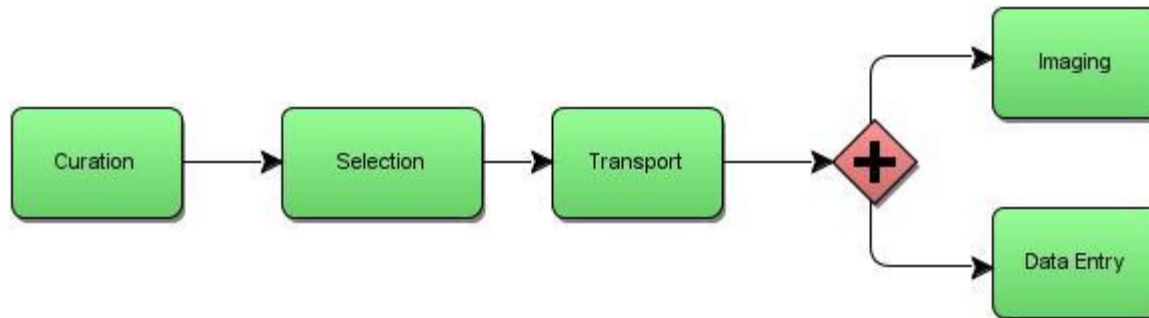
- Complete.
- Clear.
- Succinct.
- Ordered.
- Reusable.





Example Processes (Modules), their Cycles and Dependencies

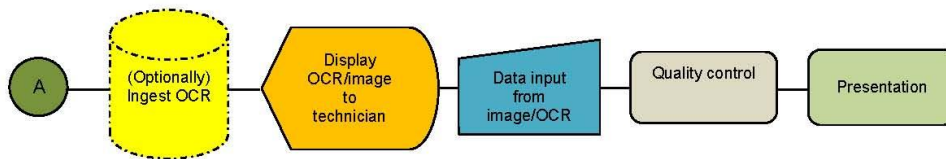
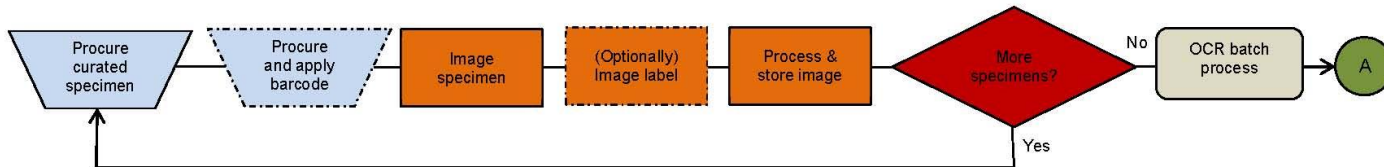
Process	Cycle	Dependency
Software configuration	Once/non-recurring	
Equipment set-up	Once/non-recurring	
Specimen curation	Recurring	
Specimen selection	Recurring	Pre-digitization curation
Specimen transport	Recurring	Specimen selection, imaging, data entry
Conservation	Episodic	Curatorial processes, imaging, data entry
Data entry	Recurring/tasks iterative	Specimen transport
Imaging	Recurring/tasks iterative	Specimen transport
Equipment adjustment	Episodic	Data entry/imaging
Software update/tweaking	Episodic	QC
Specimen return/shelving	Recurring	Imaging or data entry





O2I2D(1)—Existing Specimen Workflow Using Optical Character Recognition: Object to Image to Data

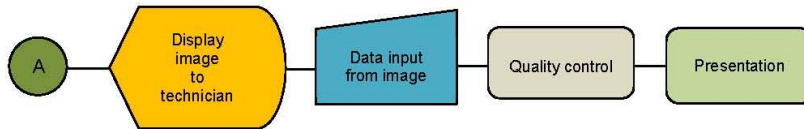
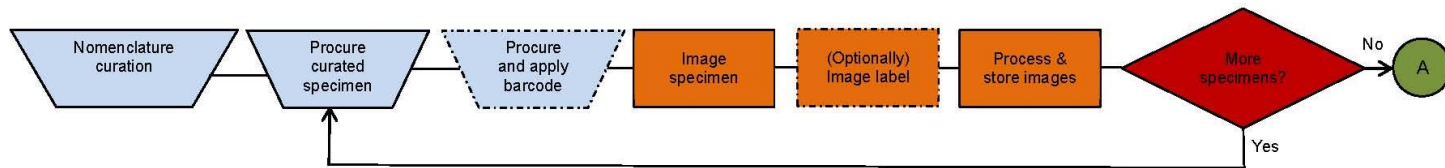
This workflow is designed to capture images of existing specimens, pass the images through optical character recognition (OCR) software, and use the combination of image and OCR output to capture data. There are variations on this workflow. For example, depending on preparation type, barcodes are sometimes applied inline as the step immediately previous to imaging (shown optionally below) and other times en masse within an independent step during which several dozen or several hundred barcodes are applied in preparation for imaging. OCR may also occur in various ways: 1) in batch (as shown below), with numerous images being processed following the close of one or more imaging sessions, 2) "on the fly" as a record and its associated image are loaded for data entry, or 3) one image at a time as a step immediately following the imaging of each specimen. OCR output may be ingested into a field in the database (shown optionally below), stored as individual text files within the computer's file system, or virtually processed at the time the image is presented to the data entry technician. The presentation of images and OCR to data entry technicians occurs in a single interface in which database fields, OCR output, and specimen image are simultaneously visible. Pre-digitization curation and annotation is particularly important in this workflow to ensure that the current nomenclature to be used in data entry is obvious and clearly visible in the image and/or OCR output.





O2I2D(2)—Existing Specimen Workflow: Object to Image to Data

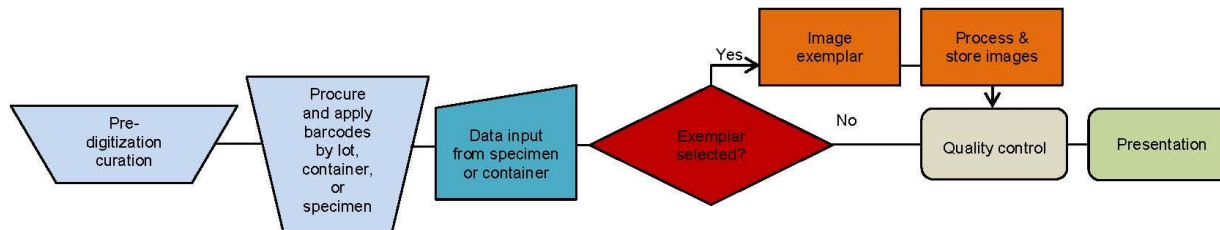
This workflow is designed for capturing images of existing specimens and using these images as the basis for data capture. Depending upon preparation type, barcodes are sometimes applied inline as the step immediately previous to imaging (shown optionally below) and other times en masse within an independent step during which several dozen or several hundred barcodes are applied in preparation for imaging. Pre-digitization curation and annotation is particularly important in this workflow to ensure that the current nomenclature to be used in data entry is obvious and clearly visible in the image.





O2D2EI—Existing Specimen Workflow: Object to Data to Exemplar Images

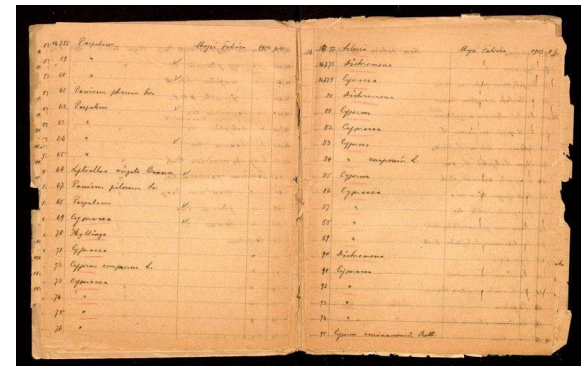
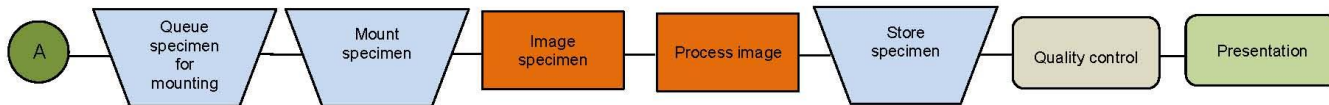
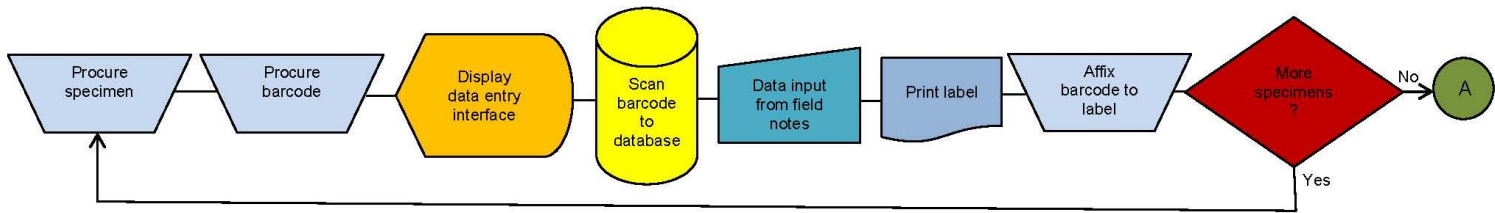
This workflow is in use for collections that capture data in specimen lots, collecting events, taxon container, or other aggregates, but capture images only for exemplar specimens. Data capture is effected from specimen labels. Depending upon preparation type, barcodes are usually applied inline—often to the containing tray or container—as the step immediately preceding data entry. Hence, barcodes may designate a single specimen or an aggregate of specimens, such as a unit tray within an insect drawer or ethanol-filled container in a wet collection. Barcode application is executed prior data entry and image capture usually follows data entry. Pre-digitization curation, including nomenclatural annotations and specimen organization, is usually important in this workflow.





FN2D2I—New Specimen Workflow: Field notes to data to image

This workflow is designed for actively growing collections in which new specimens are regularly added. Collectors, especially in herbaria, typically keystroke label data from field notes, store the label with the specimen, and queue the specimen for mounting. Following mounting, the specimen is treated as an existing specimen with the data entered into the database by a technician, who re-keys the data previously keyed by the collector. The workflow proposed here eliminates the second keying of label data by capturing label data into the database as the label is prepared, allowing the label to be printed immediately following data entry. The workflow assumes a database management system with functionality for printing labels, as well as a strategy that includes the application of bar codes to the newly printed label rather than to the specimen sheet.

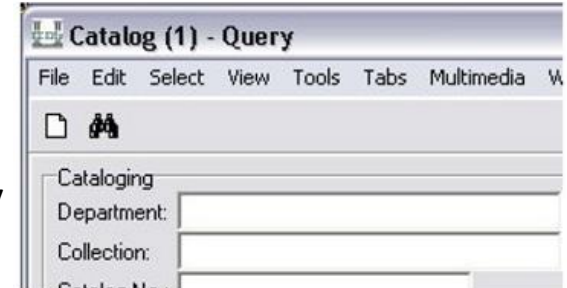
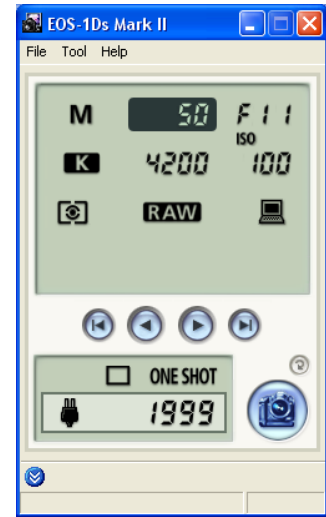




Documentation and Instructions

- **Written Protocols**
 - Essential!
 - Include screen shots and pictures.
 - Attention to detail (leave nothing to the imagination).
 - Express limits on technician authority.

- **Feedback Loops**
 - Technicians: best source of efficiency adaptations, either by show or tell.
 - Easy methods for receiving feedback.
 - Personal copies of the protocol.
 - Master copy available via Google docs or other shared storage for updates and suggestions.





Idigbio.org->Digitization->Documentation->Workflow and Protocols->Workflow Modules and Task Lists

<https://www.idigbio.org/content/workflow-modules-and-task-lists>

Workflow Modules and Task Lists

One outgrowth of the [DROID](#) (Developing Robust Object-to-Image-to-Data) workflow workshop held in May 2012 was the establishment of a series of working groups, each focused on workflow modules and tasks for various preparation types. The first of these groups, informally called the [Flat Sheets and Packets Working Group](#), was charged with fleshing out task lists for digitizing vascular and non-vascular plant collections. The second group, Pinned Specimens in Trays and Drawers, is investing its time developing modules to support effective entomological digitization workflows. Other preservation types will follow, concluding with the development of an overall project management module designed to provide guidance for developing and managing digitization projects across disciplines and preservation types.

read more



Workflow Modules and Task Lists

Researchers

[Browse our specimen portal](#)



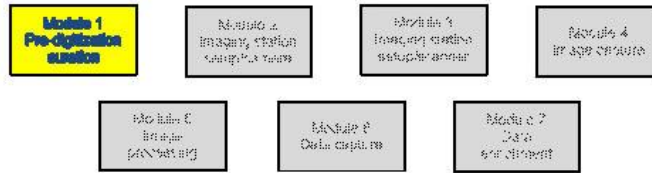
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One outgrowth of the **DROID** (Developing Robust Object-to-Image-to-Data) workflow

workshop held in May 2012 was the establishment of a series of working groups, each focused on workflow modules and tasks for various preparation types. The first of these groups, informally called the **Flat Sheets and Packets Working Group**, was charged with fleshing out task lists for digitizing vascular and non-vascular plant collections. A reconstitution of this working group, convened in January 2015, added 8 modules to this set of workflows and updated the existing ones. The second working group, **Pinned Specimens in Trays and Drawers**, invested its time developing modules to support effective entomological digitization workflows. **Things in Spirits in Jars** devoted time to workflows for fluid-preserved collections. The 3D Objects in Trays and boxes completed its work in spring 2015 and focused mostly on paleontological specimens.

We have chosen a modular approach for presenting our results in order to accommodate the broad range of workflow implementations within the collections community. We recognize that there is no consensus workflow that fits all situations, even within a single preservation type. In light of this, we have attempted to assemble orderly, comprehensive task lists to serve as foundations from which institutionally specific workflows can be created. Not all institutions will use every task, but we hope that the lists we have developed encompass all relevant digitization tasks. We also hope that those in the collections digitization community will provide feedback on these lists, either through forum posts or e-mails to Gil Nelson, alerting us to deficiencies and oversights.

Links to published modules as they are completed are provided below:

Flat Sheets and Packets Working Group - Vascular and Non-vascular Plants

- [Module 1 Pre-digitization Curation Tasks](#)
- [Module 2 Selecting Components for an Imaging Station](#)
- [Module 3 Imaging Station Setup Camera/Copy Stand](#)
- [Module 4 Imaging Station Setup Light box](#)
- [Module 5 Image Station Setup Scanner](#)
- [Module 6 Imaging](#)



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REVIEW ARTICLE

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Gil Nelson, Patrick Sweeney, Lisa E. Wallace, Richard K. Rabeler, Dorothy Allard, Herrick Brown, J. Richard Carter, Michael W. Denslow, Elizabeth R. Ellwood, Charlotte C. Germain-Aubrey, Ed Gilbert, Emily Gillespie, Leslie R. Goertzen, Ben Legler, D. Blaine Marchant, Travis D. Marsico, Ashley B. Morris, Zack Murrell, Mare Nazaire, Chris Neefus, Shanna Oberreiter, Deborah Paul, Brad R. Ruhfel, Thomas Sasek, Joey Shaw, Pamela S. Soltis, Kimberly Watson, Andrea Weeks and Austin R. Mast

1500065

 [Abstract](#)

[Abstract & References](#) : [Full Text](#) : [PDF \(778 KB\)](#) : [Supplementary Materials](#)

APPLICATION ARTICLE

Bioinformatic Identification and Expression Analysis of *Nelumbo nucifera* MicroRNA and Their Targets 


Lei Pan, Xiaolei Wang, Jing Jin, Xiaolu Yu and Jihong Hu

1500046

 [Abstract](#)

[Abstract & References](#) : [Full Text](#) : [PDF \(1342 KB\)](#) : [Supplementary Materials](#)

PRIMER NOTES

Development of 23 Novel Polymorphic EST-SSR Markers for the Endangered Relict Conifer *Metasequoia glyptostroboides* 

Yuqing Jin, Quanxin Bi, Wenbin Guan and Jian-Feng Mao

1500038

 [Abstract](#)

[Abstract & References](#) : [Full Text](#) : [PDF \(492 KB\)](#) : [Supplementary Materials](#)

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Title Tools

Most Read Articles

Digitization Workflows for Flat Sheets and Packets of Plants, Algae, and Fungi 


A Protocol for Targeted Enrichment of Intron-Containing Sequence Markers for Recent Radiations: A Phylogenomic Example from *Heuchera* (Saxifragaceae) 


Biodiversity Comparison among Phylogenetic Diversity Metrics and Between Three North American Prairies 

Plann: A Command-Line Application for Annotating Plastome Sequences 

An Empirical Review: Characteristics of Plant Microsatellite Markers that Confer Higher Levels of Genetic Variation 

Most Cited Articles

A Targeted Enrichment Strategy for Massively Parallel Sequencing of Angiosperm Plastid Genomes 

A Target Enrichment Method for Gathering Phylogenetic Information from Hundreds of Loci: An Example from the Compositae 

Hyb-Seq: Combining Target Enrichment



FluidPreservedWorkingGr x iDigBio Wiki | Small Herbaria x

← → ↻ https://www.idigbio.org/wiki/index.php/Small_Herbarium_Workshop_FSU ☆ ☰

Collaborative Notes

- [Collaborative Notes Doc](#)

Workflow Documents

- [Florida State University Herbarium Imaging Protocol](#)
- [Valdosta State University Herbarium \(VSC\) Vascular Plant Imaging Protocol](#)
- [Valdosta State Herbarium \(VSC\) Bryophyte Packet Imaging Protocol](#)
- [Valdosta Herbarium image processing with Nikon Dust Off process included](#)
- [Increasing the efficiency of digitization workflows for herbarium specimens, Tulig M, Tarnowsky N, Bevans M, et al](#)

iDigBio's Flat Sheets and Packets Working Group Workflows

- [Module 1 Pre-digitization Curation Tasks](#)
- [Module 2: Imaging station setup for camera stations](#)
- [Module 3: Imaging station setup for scanners](#)
- [Module 4: Imaging tasks](#)
- [Module 5: Image processing](#)
- [Module 6: Data capture](#)

Imaging Equipment

- [NYBG Herbarium Imaging and Equipment Specifications](#)
- [Nikon cameras and a related copy stand and lights](#)

Imaging Procedures and Workflows

- [NYBG: Standardized Digital Imaging and Archiving Procedures, Mike Bevans](#)

Image Processing

- [NYBG Image editing guidelines](#)
- [iDigBio's Recommendations for the Acquisition, Processing, Storage, and Distribution of Digital Images](#)

Related Articles and Papers

- [Streamlining Collaborative Digitization, Tulig & Watson, NYBG](#)

Barcodes

- [LBCC barcode document](#)

Sources for Barcodes

- [CompuType 2285 West County Road C St. Paul, MN 55113 \(800\) 328-0852](#)
- [Watson Label Products 3884 Forest Park Blvd St Louis, Mo 63108 \(314\) 652-6715 \(800\) 678-6715](#)
- [University Products blank archival labels](#)

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Word	Update readme.md	14 days ago
README.md	Update README.md	14 days ago

README.md

This folder contains the original set of 14 workflow modules published with the paper Digitization workflows for flat sheets and packets of plants, algae, and fungi, Nelson, G., P. Sweeney, L. E. Wallace, R. K. Rabeler, D. Allard, H. Brown, J. R. Carter, et al., Applications in Plant Sciences 3(9): 1500065. doi:10.3732/apps.1500065 (<http://www.bioone.org/doi/pdf/10.3732/apps.1500065>). Files in this folder are linked to the published paper and will not be edited or revised. Future revisions will be stored in a separate directory. PDF and Word versions are provided in separate folders.

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Continuous Workflow Improvement

Develop written workflows that reflect actual practice

Continuous evaluation of written and actual workflows by:

- Technicians
- Workflow managers
- Collections managers

With particular attention to:

- Bottlenecks
- Redundancy
- Handling time
- Varying rates of productivity



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