"DOCUMENTING THE OCCURRENCE THROUGH SPACE & TIME OF AQUATIC NON-INDIGENOUS FISH, MOLLUSKS, ALGAE, & PLANTS THREATENING NORTH AMERICA'S GREAT LAKES" [GLIS]

Aim (Plant side): digitize specimens of 43 [49] genera of plants from North America that include species known to be invasive in the Great Lakes region.

19 [28] Great Lakes participants

lead: WIS

"Hubs": F, MICH, MOR, NY

Other participants:

- BHO, BUT, ILL/ILLS, MIL, MIN, MSC, MU, ND, NYS, UWL, UWM, UWSP
- Michigan Small Herbarium Initiative (10 in yellow)
- Canadensys (http://www.canadensys.net/)
- currently has 800K vasc. plant records from 12 herbaria

One advantage: ILL/ILLS, MICH, MIN, MU, WIS worked with NY on imaging for the Tri-Trophic TCN



https://www.idigbio.org/content/digitization-tcn-great-lakes-invasives-collaborator-map

GLIS WORKFLOW — COLLECTION MANAGER/CURATOR

Prepare specimens for imaging [Pre-curation]

- Specimens already barcoded (e.g., in-state specimens) only imaging needed
 - Data may be complete or "minimal"; ideally, assume data is present if barcoded.
 - Nomenclature likely is current if not, see "Check/Update nomenclature" below.
 - "First barcode" tag marked with name of taxon and inserted above 1st specimen of that taxon
 - Mark genus/case as ready for imaging
- Specimens not barcoded (e.g., out-of-state specimens)
 - Will be barcoded (skeletal records created) and then imaged
 - Check/Update nomenclature we use "filed by" names in project
 - Check names against Tropicos/FNA; not a formal annotation (ID NOT checked). Folders tagged so Herbarium Assistant can pencil new name on sheets
 - May result in some collection rearrangement/refoldering (add/remove genera, species)
 - Segregate any needing repair
 - Mark multiple sheet and mixed collections, pull duplicates, discover types
 - "First barcode" tag marked with name of taxon and inserted above 1st specimen of that taxon
 - Mark genus/case as ready for imaging

(Later) Mark genus/case as completed

GLIS WORKFLOW — EIT'S AND PROJECT MANAGER

Electronic Imaging Technicians [EIT]

- Retrieve specimens from collection as directed by CM/Curator; mark pigeonholes
- Barcode specimens + create skeletal record (Excel, Access, App) if needed
 - Skeletal record = barcode + taxon name [+collector,#, state]
- Add barcode # to "First Barcode" tag when starting a new taxon
- Image the specimens [lightbox]; image name = barcode #
- Return specimens to Collection
- Move to next genus

Project Manager

- Process the images Adobe Lightroom; add metadata, color correct, create DNG/JPG
- Rename files, adding "MICH-V-" as a prefix
- Use "First Barcode" tag to help sort/store all images by taxon locally
- Load images + data (complete or skeletal) records to Symbiota portal(s)
- OCR performed at WIS on images with skeletal data loaded to portal
- Perform Label Transcription (via Symbiota GL portal), Georeferencing as necessary