# Molecular Resources and Protocols: From PLS4178 to PLS4172

Alina Avanesyan

Lab Meeting 10/12/2020





## DNA Barcoding Work: Location

- Room 4172, 4th floor
- Dr. David Hawthorne's lab



David Hawthorne, Associate Professor & Director of Education at SESYNC

Office: 4132 Plant Sciences Building

Phone: 301-405-2401 E-mail: djh@umd.edu

**Research**: The Hawthorne Lab uses population genetics to understand how insects become pests, how they evolve to counter control efforts, and how to use evolutionary thinking to manage them. Additionally, research in the Hawthorne Lab dissects the genetic basis of host-plant associated divides among pest populations and uses phylogeographic analyses to investigate issues in conservation genetics.







- Host plant DNA detection from potato leafhopper, *Empoasca fabae*
- Host plant DNA detection from the spotted lanternfly,
   Lycorma delicatula
- Wetland-stream connectivity: DNA barcoding of isopods
- Prey-predator interactions: DNA barcoding of prey items in Odonata gut contents and feces









## DNA Barcoding Team (at different times over the past 2.5 years )

- 1. Brock Couch\*\*
- 2. Kevin Clements\*
- 3. Nina McGranahan ♦
- 4. Bryan Stancliff♦
- 5. Omar Abdelwahab\*
- 6. Jessica Ho\*
- 7. Darsy Smith\*\*
- 8. Margaret Hartman\*\*
- 9. Nurani Illahi\*
- 10. Hannah Sutton\*









## DNA Barcoding Team: Research results

#### Journal articles

- Avanesyan, A., Illahi, N. and W.O. Lamp. (2020) Detecting ingested host plant DNA in potato leafhopper, Empoasca fabae: potential use of molecular markers for gut content analysis. Journal of Economic Entomology (In Press).
- Avanesyan, A., and W.O. Lamp. (2020) Use of molecular gut content analysis to decipher the range of food plants of the invasive spotted lanternfly, *Lycorma delicatula*. Insects: Special Issue "Molecular Gut Content Analysis: Deciphering Trophic Interactions of Insects", 11(4), 215, doi.org/10.3390/insects11040215.

#### Manuscript in revision

Avanesyan, A., Sutton, H., and W.O. Lamp. (2020) Choosing an effective molecular approach to diet analysis of insect herbivores: a systematic review.

#### Manuscript in preparation

Avanesyan, A., Sutton, H., Lamp, W.O., and D. Hawthorne. Identification of host plant use by the invasive spotted lanternfly (*Lycorma delicatula*) using next-gen DNA sequencing technology.





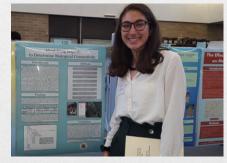




## Conference and Research Symposium Presentations

- Avanesyan, A., and W. Lamp (2018) Use of molecular markers for plant DNA to determine host plant usage for potato leafhopper, Empoasca fabae. Annual Meeting of the Entomological Society of America: 2018 ESA, ESC, and ESBC Joint Annual Meeting, Vancouver, BC, Canada. Oral presentation
- McGranahan, N. (2019) Identifying isopods to determine biological connectivity. The ERHS Research Symposium. Poster Presentation.

Stancliff, B. (2019). The John Carroll School Research Symposium. Oral presentation











#### GenBank Submissions



- 1. Avanesyan, A. and W. O. Lamp. (2020) *Betula pendula* isolate 1E4a ribulose-1,5-bisphosphate carboxylase/oxygenase large subunit (rbcL) gene, partial cds; chloroplast. Direct Submission, GenBank Accession no. MT119453
- 2. Avanesyan, A. and W. O. Lamp. (2020) *Acer pseudoplatanus* isolate 1F4b ribulose-1,5-bisphosphate carboxylase/oxygenase large subunit (rbcL) gene, partial cds; chloroplast. Direct Submission, GenBank Accession no. MT108179
- 3. Avanesyan, A. and W. O. Lamp. (2020) Vitis vinifera isolate 1B3 ribulose-1,5-bisphosphate carboxylase/oxygenase large subunit (rbcL) gene, partial cds; chloroplast. Direct Submission, GenBank Accession no. MN862495
- 4. Avanesyan, A. and W. O. Lamp. (2020) *Ailanthus altissima* ribulose-1,5-bisphosphate carboxylase/oxygenase large subunit (rbcL) gene, partial cds; chloroplast. Direct Submission, GenBank Accession no. MN853649
- 5. Avanesyan, A. and W. O. Lamp. (2020) *Celastrus orbiculatus* isolate TT4a ribulose-1,5-bisphosphate carboxylase/oxygenase large subunit (rbcL) gene, partial cds; chloroplast. Direct Submission, GenBank Accession no. MN862496
- 6. Illahi, N.\*, Avanesyan, A. and W. O. Lamp. (2020) *Lonicera maackii* ribulose-1,5-bisphosphate carboxylase/oxygenase large subunit (rbcL) gene, partial cds; chloroplast. Direct Submission, GenBank Accession no. MN631052
- 7. Smith, D.K.\*\*, Avanesyan, A. and W. O. Lamp. (2020) Eupatorium serotinum tRNA-Leu (trnL) gene, partial sequence; chloroplast. Direct Submission, GenBank Accession no. MN395725
- 8. Smith, D.K.\*\*, Avanesyan, A. and W. O. Lamp. (2020) Lonicera maackii tRNA-Leu (trnL) gene, partial sequence; chloroplast. Direct Submission, GenBank Accession no. MN365276
- 9. Smith, D.K.\*\*, Avanesyan, A. and W. O. Lamp. (2020) *Pisum sativum* isolate slf-2 tRNA-Leu (*trn*L) gene, partial sequence; chloroplast. Direct Submission, *GenBank* Accession no. MN335637
- 10. Smith, D.K. \*\*, Avanesyan, A. and W. O. Lamp. (2020) *Acer platanoides* tRNA-Leu (trnL) gene, partial sequence; chloroplast. Direct Submission, GenBank Accession no. MN450067
- 11. Smith, D.K. \*\*, Avanesyan, A. and W. O. Lamp. (2020) *Acer rubrum* tRNA-Leu (trnL) gene, intron; chloroplast. Direct Submission, GenBank Accession no. MN450068
- 12. Illahi, N.\*, Avanesyan, A. and W. O. Lamp. (2019) *Ailanthus altissima* isolate BC4b ribulose-1,5-bisphosphate carboxylase/oxygenase large subunit (rbcL) gene, partial cds; chloroplast. Direct Submission, GenBank Accession no. MN856629
- 13. Illahi, N.\*, Avanesyan, A. and W. O. Lamp. (2019) *Ailanthus altissima* ribulose-1,5-bisphosphate carboxylase/oxygenase large subunit (rbcL) gene, partial cds; chloroplast. Direct Submission, GenBank Accession no. MN853649
- 14. Stancliff, B.♦, Avanesyan, A. and W. Lamp. (2019) *Vicia faba* tRNA-Leu (trnL) gene, partial sequence; chloroplast. Direct Submission, *GenBank* Accession no. MK934667
- 15. Stancliff, B.♦, Smith, D.\*\*, Avanesyan, A. and W. Lamp. (2019) *Pisum sativum* tRNA-Leu (*trnL*) gene, partial sequence; chloroplast. Direct Submission, *GenBank* Accession no. MK919208
- 16. Stancliff, B.♦, Abdelwahab, O.\*, Avanesyan, A. and W. Lamp. (2019) Vigna unguiculata tRNA-Leu (trnL) gene, partial sequence; chloroplast. Direct Submission, GenBank Accession no. MK883492
- 17. Stancliff, B.♦, Ho, J.\*, Avanesyan, A. and W. Lamp. (2019) *Helianthus annuus* tRNA-Leu (*trn*L) gene, partial sequence; chloroplast. Direct Submission, *GenBank* Accession no. MK875279
- 18. Avanesyan, A., and W. Lamp. (2019) *Vicia faba var. major* isolate PLH\_fb tRNA-Leu (*trn*L) gene, partial sequence; chloroplast. Direct Submission, *GenBank* Accession no. MK837073

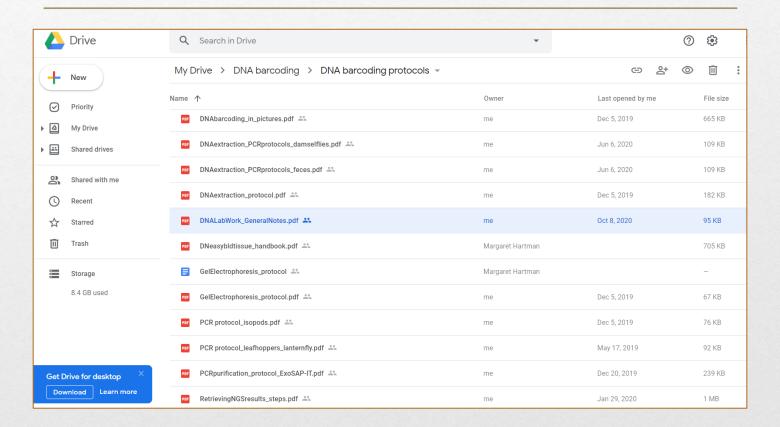
  (\*undergraduate students, \*\*graduate students, ♦ high school students)





## DNA Barcoding Protocols

Shared Google folder





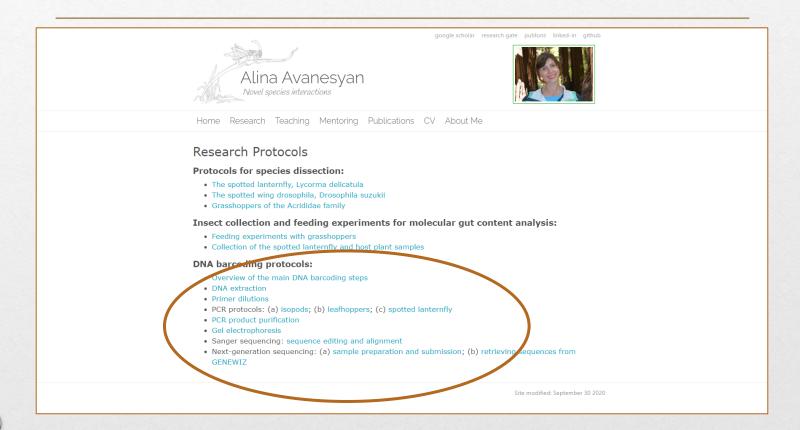






### DNA Barcoding Protocols

http://alinaavanesyan.com/projects/research-protocols/





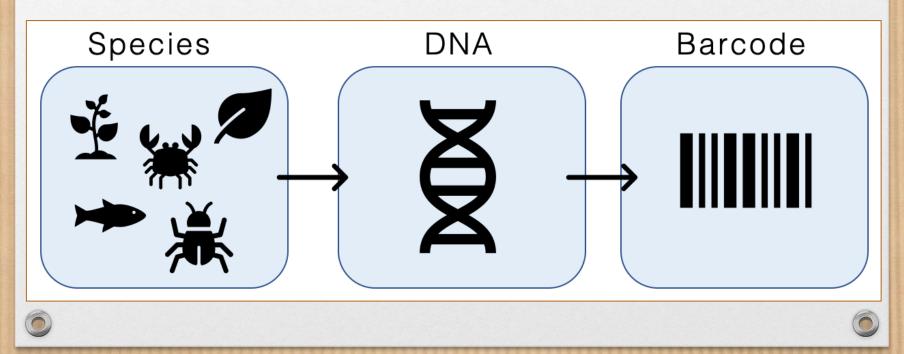






### DNA Barcoding

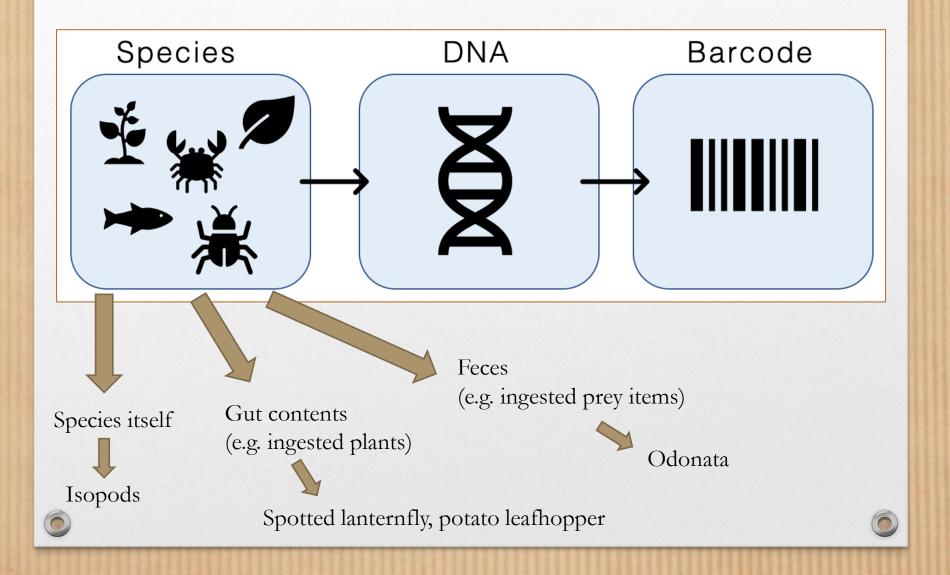
 method of species identification using a short section of DNA from a specific gene or genes







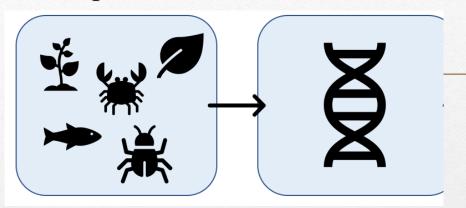
#### Sources of DNA



## DNA Barcoding: How do we do it?

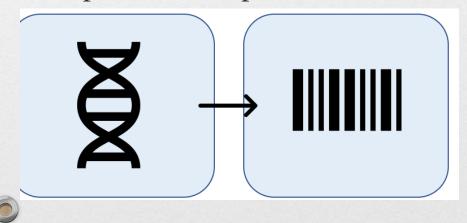


Step 1. DNA extraction



Final product: genomic DNA

Step 2. PCR amplification



Final product: targeted piece of DNA



Piece of plant DNA, piece of insect mitochondrial DNA, etc.



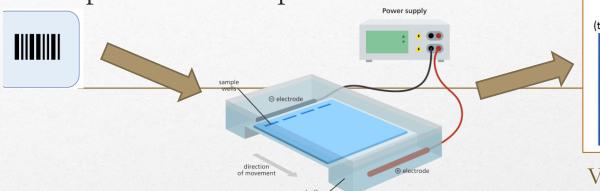


## DNA Barcoding: How do we do it?

Electrophoresis tank



Step 3. Gel electrophoresis

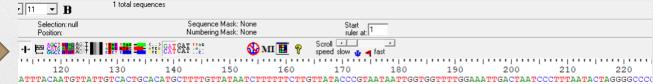


wetland "+"control (terrestrial isopod) specimens 1000 bp 750 bp 500 bp

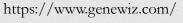
Verification of presence of targeted DNA

Step 4. Sequencing









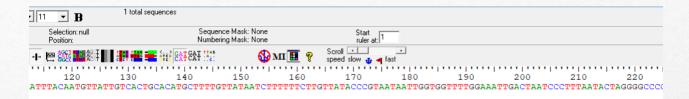






### DNA Barcoding: What's next?







#### **Web BLAST**

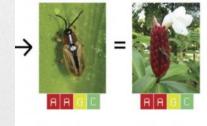


https://blast.ncbi.nlm.nih.gov/

Identification of species

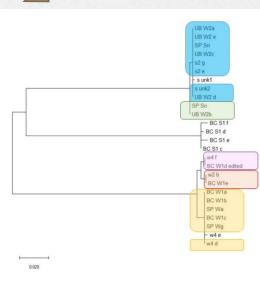


#### Matching DNA sequences and host plant identification



Identification of species interactions





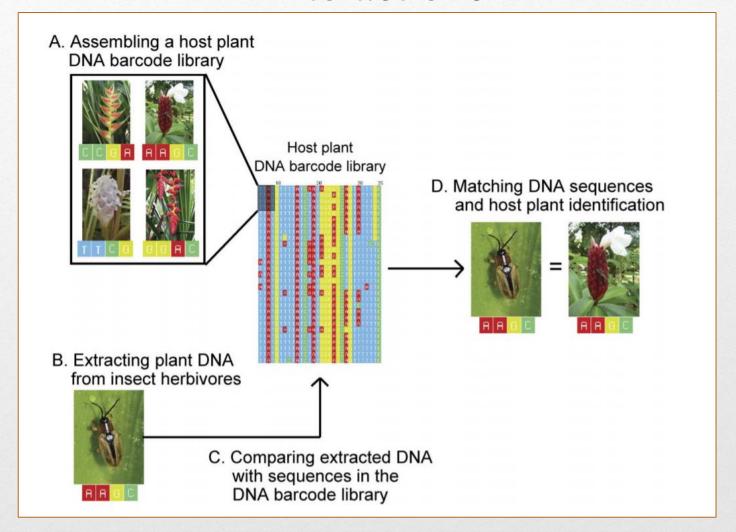
Constructing phylogenetic relationships





## Identification of plant-insect interactions











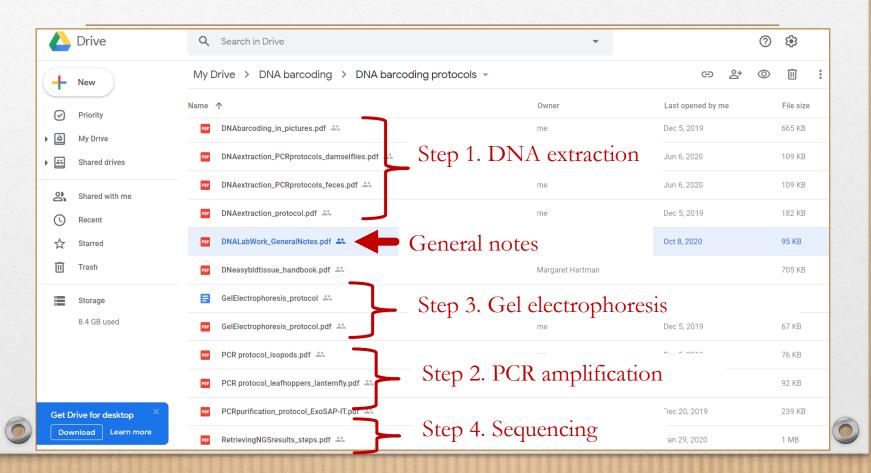


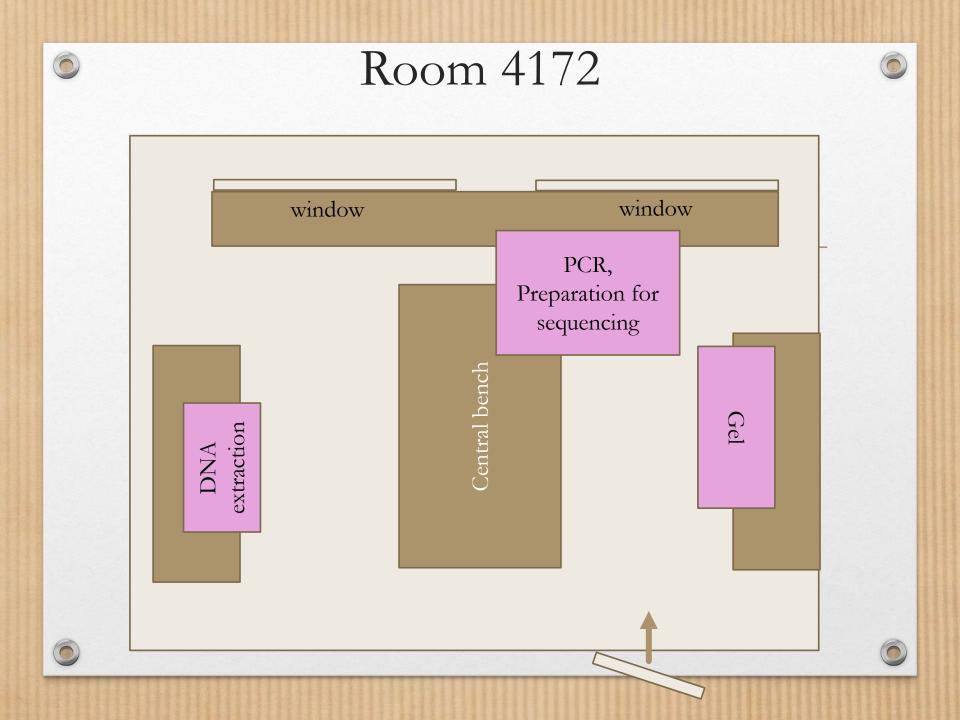


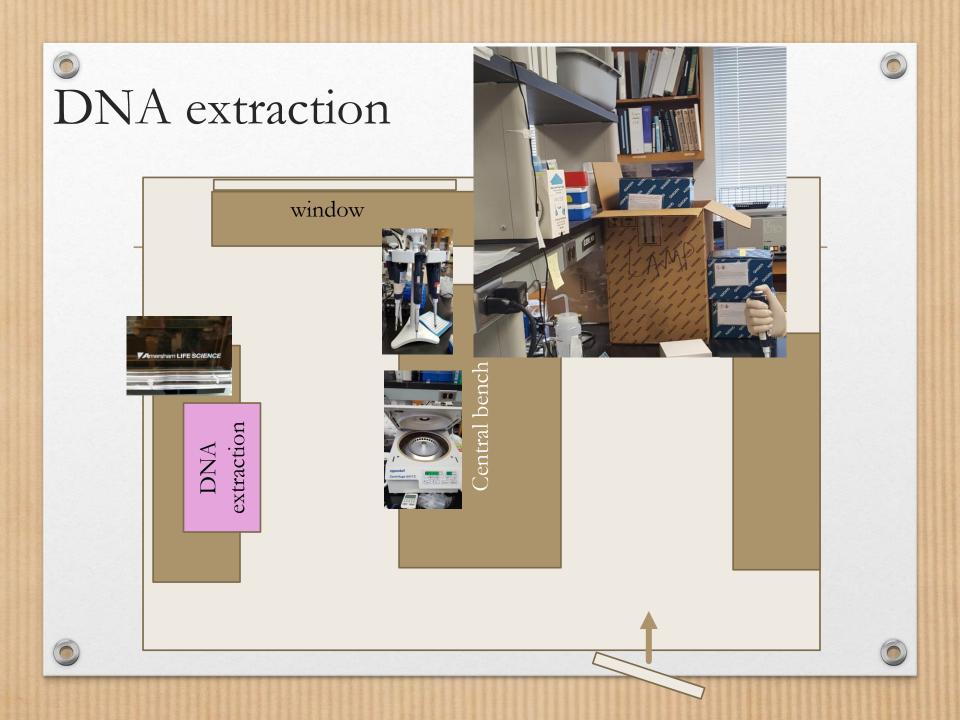


### DNA Barcoding Protocols

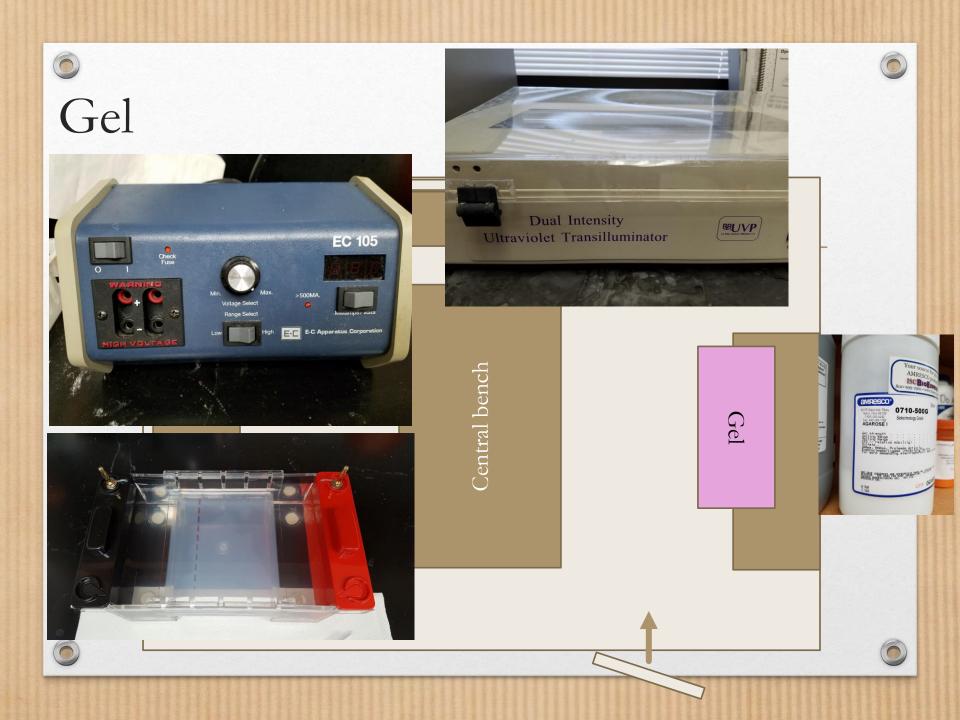
Shared Google folder















## Sequencing

- Open an account at genewiz.com
- Follow the guidelines for sample preparations
- Samples drop-off: dropbox on the 2<sup>nd</sup> floor





- Results are ready on the next day (Sanger sequencing) or in ~3 weeks (NGS)
- Contact Genewiz customer service if you have any questions
- Contact me if you have more questions ©







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- Maggie, mehartma@umd.edu
- Nina, nina.e.mcg@gmail.com
- Hannah, hsutton1@terpmail.umd.edu





#### Helpful Resources

#### Textbook:

• Genetics: Analysis and Principles, Brooker et al, 6th edition

#### Coursera:

- Introduction to Genetics and Evolution: <a href="https://www.coursera.org/learn/genetics-evolution">https://www.coursera.org/learn/genetics-evolution</a>
- DNA decoded: <a href="https://www.coursera.org/learn/dna-decoded">https://www.coursera.org/learn/dna-decoded</a>

#### Review papers on DNA barcoding:

- Taylor, H. R., & Harris, W. E. (2012). An emergent science on the brink of irrelevance: a review of the past 8 years of DNA barcoding. Molecular Ecology Resources, 12(3), 377-388.
- Li, X., Yang, Y., Henry, R. J., Rossetto, M., Wang, Y., & Chen, S. (2015). Plant DNA barcoding: from gene to genome. Biological Reviews, 90(1), 157-166.





#### Thank you!

Happy DNA barcoding!

