

## Introduction



- The plant family Onagraceae is divided into two subfamilies, Ludwigioideae, with only genus Ludwigia of 82 species and 22 sections, and Onagroideae, which consists of six tribes and 21 genera.
- This research is taking a closer look at fruit and seed morphology in 35 species representing all sections chosen to fill gaps in the previous studies (Edye, 1978).

# Introduction

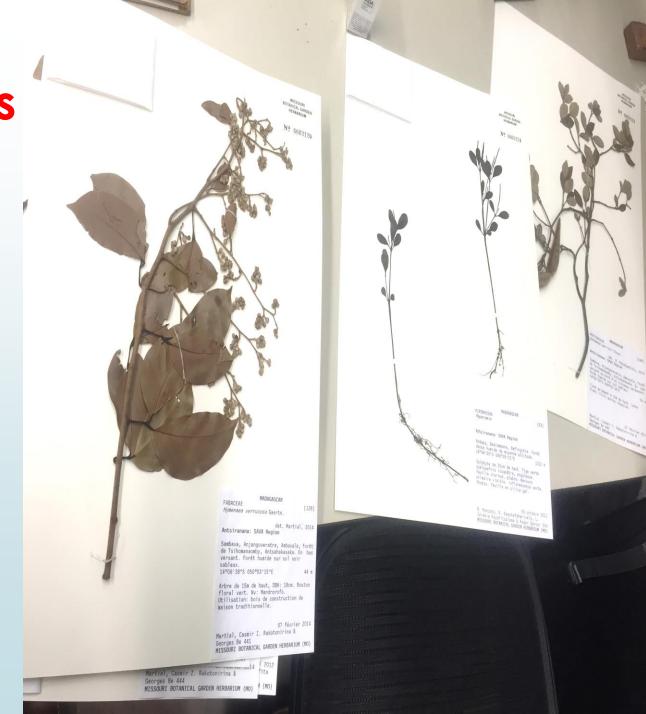
The objective is to discover how fruit and seed morphology can advance knowledge of the relationships among and within the sections of Ludwigia.



■ Why: Our understanding of the phylogenetics of Ludwigia is advancing and within this context there is a need for more morphological work to be done, to complement and focus molecular phylogenetic studies.

# **Materials & Methods**

- Samples
- **■** Literature
- R Program
- Light Microscope
- Sputter Coater
- Scanning Electron Microscope (SEM)

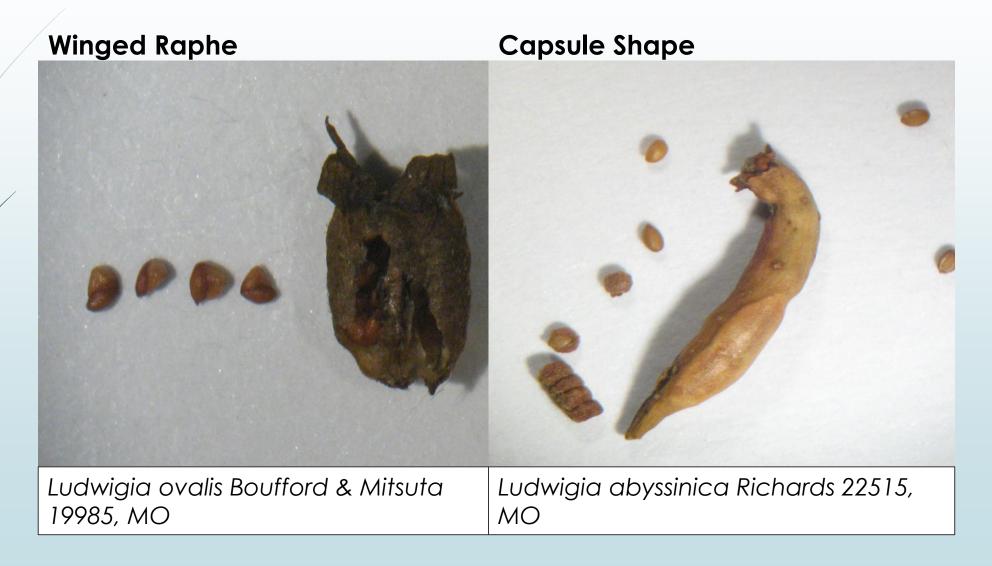


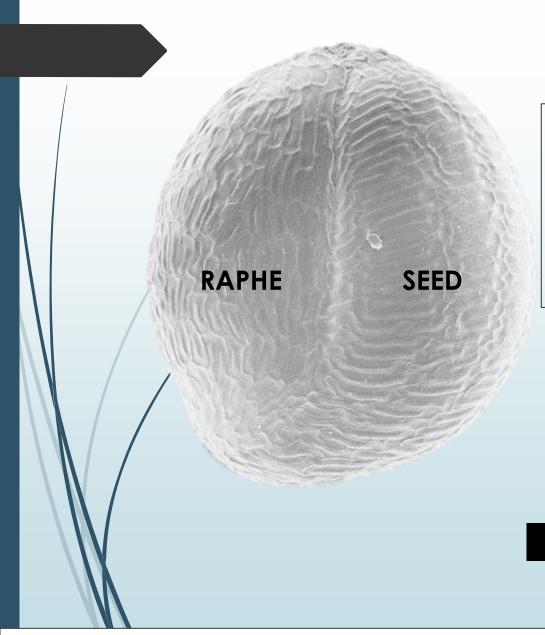
## **Materials & Methods**



- 35 species & 10 specimens per species
- 3 fruits and seeds were measured per specimen
- Measured under Light Microscope and photographed
- SEM images of seeds at several magnifications were taken for species not already imaged

# Examples of morphological characters

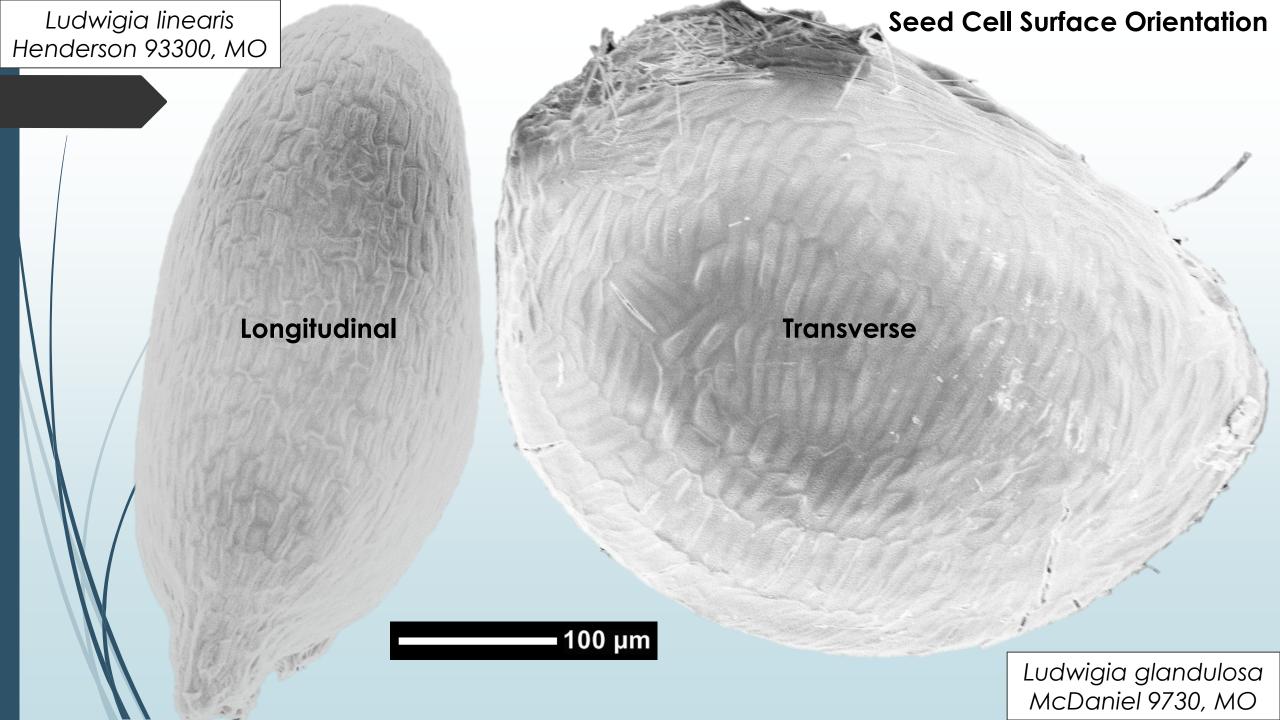




Raphe is a groove attached to the seed, typically marking the line where two halves fused in the embryo.

■ 200 µm





## **Results & Discussion**

#### <u>Table Summary of the</u> <u>morphological characters</u>

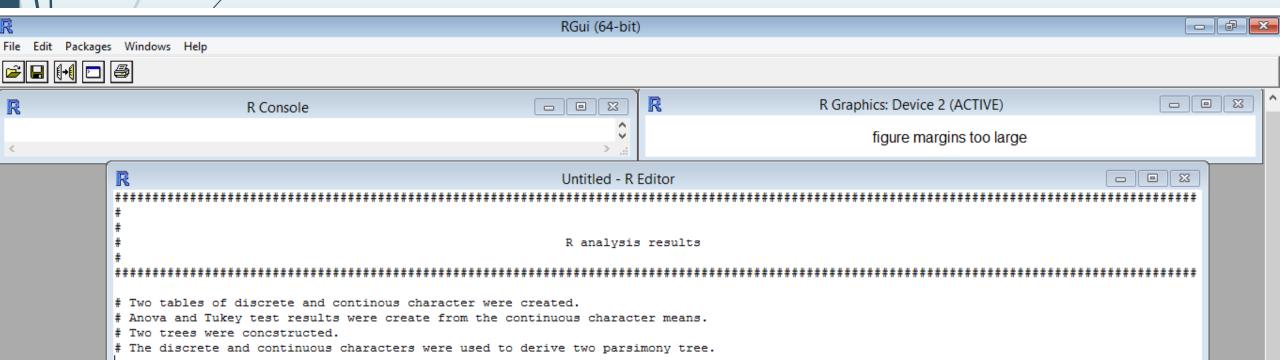
- 3 sections characterized by expanded raphe
- Seed cell surface orientation only groups 4 of the 35 species together
- Five types of capsule shapes and seed arrangements.

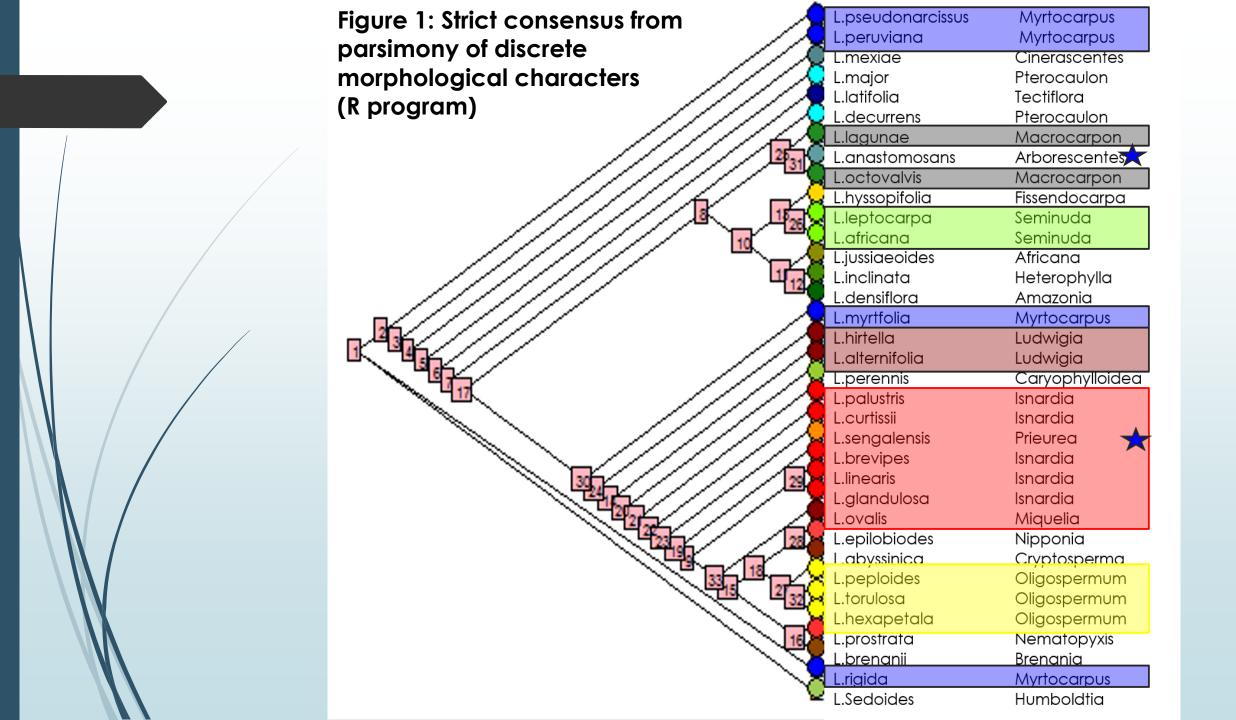
#### **Analysis of variance and means**

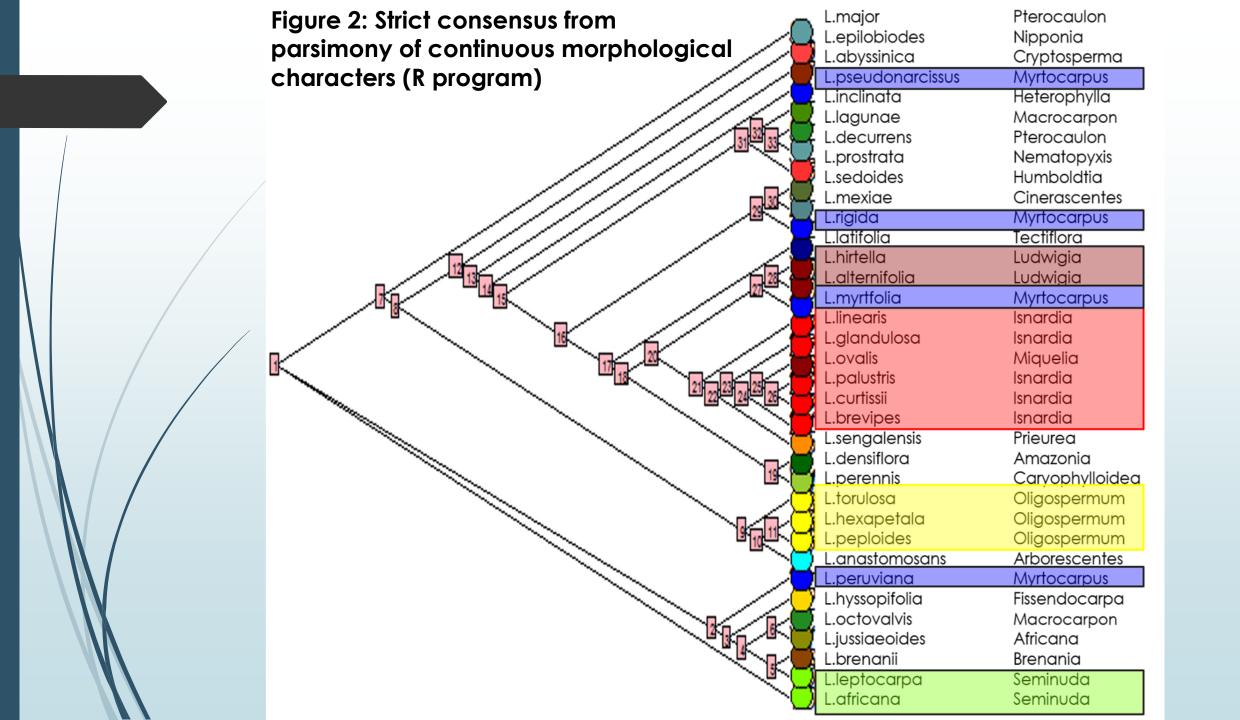
- In searching for differences the trend was that within sections the means were significantly the same
  - **►** Exception: Myrtocarpus
- Within the genus there are fewer sections that are consistently significantly the same
  - ■i.e., Isnardia and Miquelia

# "R" Analysis Results

- Two tables of discrete and continuous character were created. For Anova and Tukey t-test results in the table are created from the continuous character means.
- Two trees were constructed. The discrete and continuous characters were used to derive two parsimony tree.



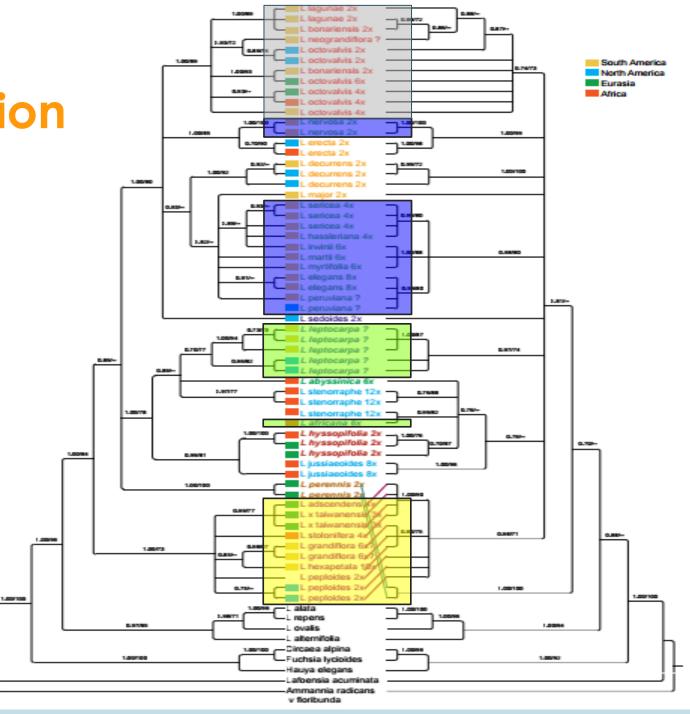




**Results & Discussion** 

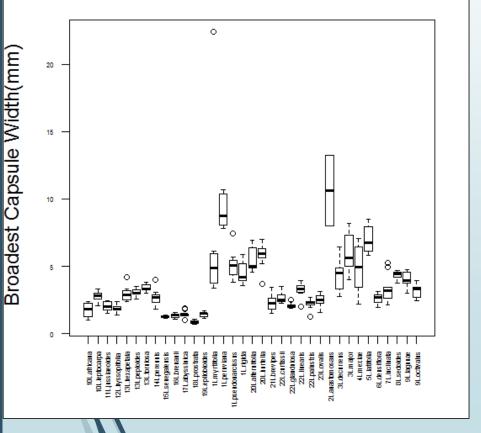
## **Molecular Phylogeny**

**■** The morphological data provide some support for the new molecular phylogeny (S. Liiu et al., unpubl.), but it is still unclear the degree to which either or both datasets support the traditional sectional classification



# **Future Research**

#### Mean Broadest Capsule Width for the genus Ludwigia



- Ludwigia is very diverse morphologically, and a thorough analysis will require morphological data from all or most species, and from additional characters.
- Combining morphological and molecular data is essential for a complete understanding of the phylogeny and evolutionary history of Ludwigia.
- Continued analysis of fruit and seed characters in Ludwigia will provide insight into ecological adaptations that have facilitated the evolutionary success of the genus.

## Conclusion

- Fruit and Seed morphology provides limited support for some sections and the North Temperate haplostemonous group (sections Ludwigia, Isnardia, and Miquelia)
- Recent molecular phylogenetic analysis also provides only limited support for some parts of the traditional sectional classification.
- Several fruit and seed morphological character appear to be associated with the evolutionary radiation of some taxa or groups of Ludwigia

# Thank You!

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- Missouri Botanical Garden
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- Columbia College



# **Questions?**

Thank you for listening!