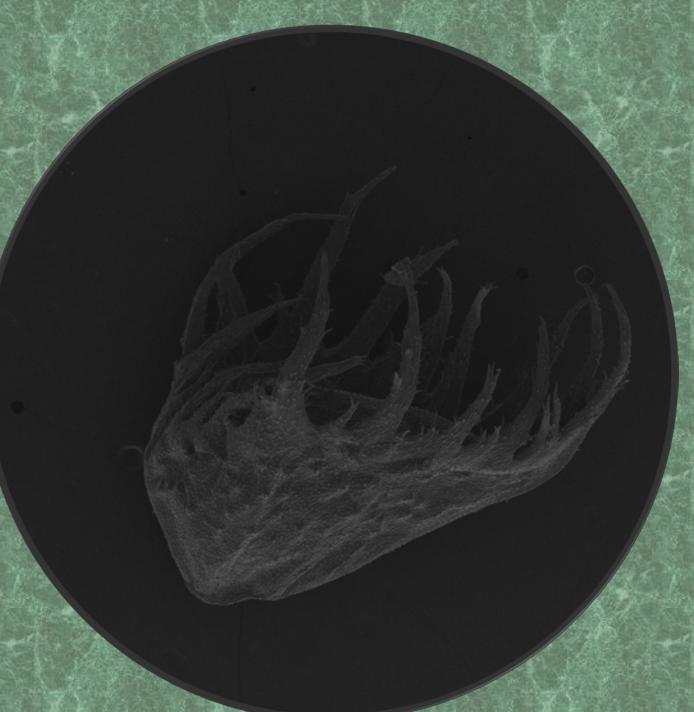
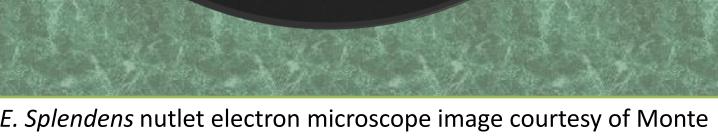
Alaskan Eritrichium Systematics

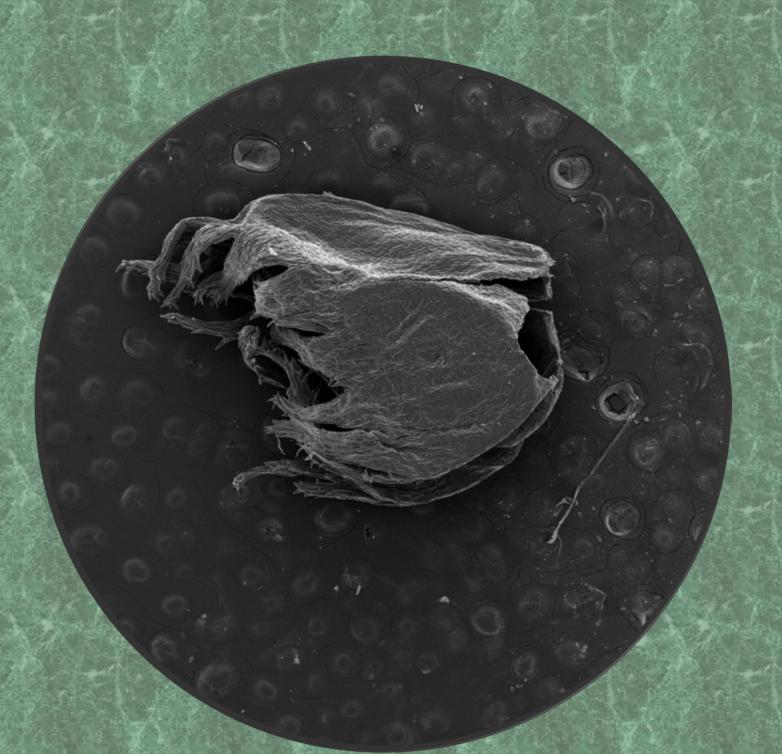
By Samantha Wilson, Monte Garroutte, and Kristen O' Brien

Introduction

The *Eritrichium* project is a small part of the Flora of North America project. There are three types of Eritrichium (Alpine forget-me-not) in Alaska; *E. splendens*, *E. aretioides*, and *E. chamissonis*. Our project is dealing with a division within *E. splendens* relating to size differences between regions. We want to genetically compare the two to determine if they are different species. We are also doing a morphological study see if we can identify the different species by nutlet size and hair structure and ultimately create an evolution tree using morphology and genetics.

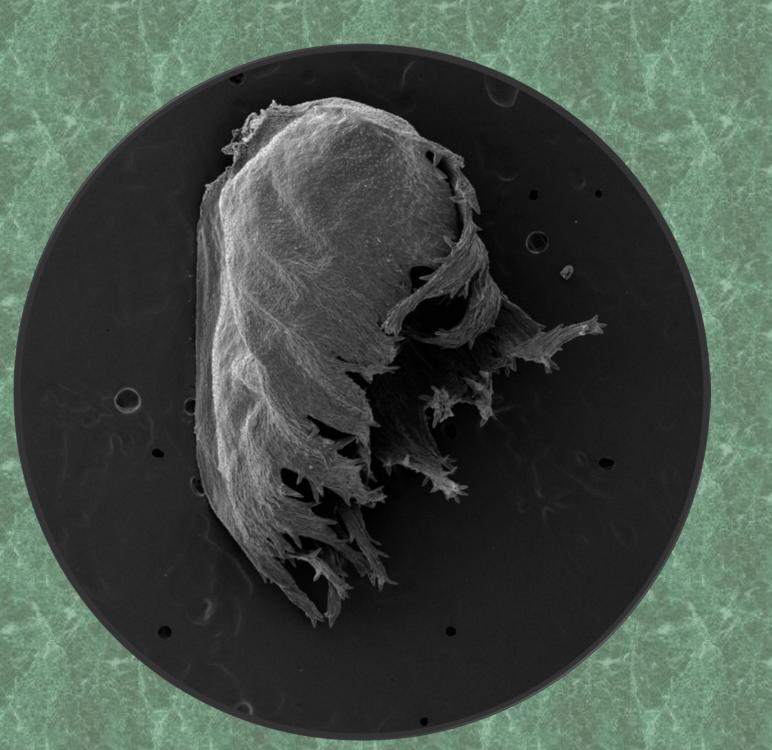






E. chamissones nutlet electron microscope image courtesy of Monte

Garroutte



E. aretiodes nutlet electron microscope image courtesy of Monte Garroutte



E. splendens

Methods

Genetics - We are targeting three genes within *E. splendens* and *E. chamissonis'* plastid DNA. The genes are rps16, TrnRr, and TrnSG. The first step we had to do was extract DNA from *E. splendens*. Next we did PCR, on rps16 and TrnSG in *E. splendens* and *E. chamissonis*. We used an agarose gel to determine if DNA was present.

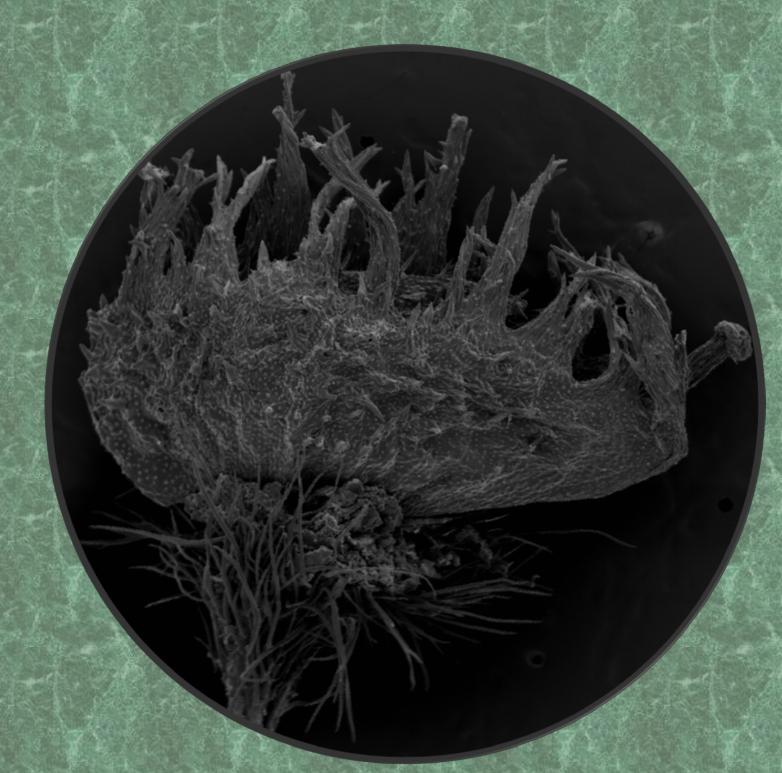
Morphology- We examined scanning electron microscope images to identify the characteristics of nutlets and how they vary between species. We also examined hair thickness variation between species.



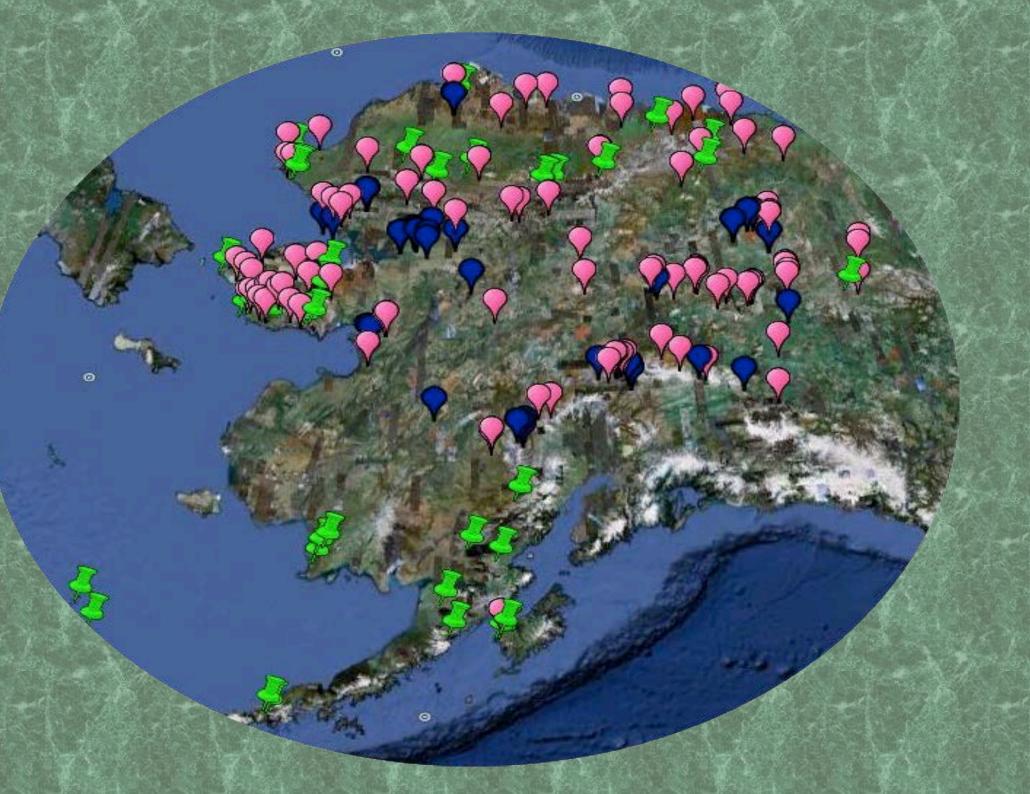
E. aretioides



E. chamissonis



E. Splendens nutlet (large flower variety) electron microscope image courtesy of Monte



Green is *E. chamissonis*, blue is *E. splendens*, pink is *E. aretioides* courtesy of Monte Garroutte and Arctos

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Results

Genetics- We managed to isolate and perform PCR on the rps16 gene for *E. chamissonis* and the TrnSG gene for *E. chamissonis*. We found that a higher annealing temperature during PCR tended to work best. We have not yet managed to successfully perform PCR on *E. splendens*.

Morphology- We have started the process of examining and putting quantifiable information in to a data base on a variety of *Eritrichium*. We hope to not only be able to confidently identify an *Eritrichium* to the species by looking at the nutlets, but to also be able to use these features to create an evolutionary tree to see which *Eritichium* are most related. However, many more data sheets are needed to make this possible.

Conclusion

We do not yet have enough information to conclude that *E. splendens* has two separate species under that name. However, we have narrowed down the ways to get a working PCR for genetic proof of our hypothesis. We have also made significant progress in retrieving genetic information for *E. chamissonis*.

Nutlet examination, if proved effective in identification, can, working together with our genetic research, be an effective way to identify the various types of *Eritrichium*.





