

# Plant invasion genetics; what DNA can tell us about some of Washington's worst plants

John Gaskin

Botanist

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# Snake River





# How do plants spread??



*Tamarix ramosissima*  
saltcedar  
Kazakhstan

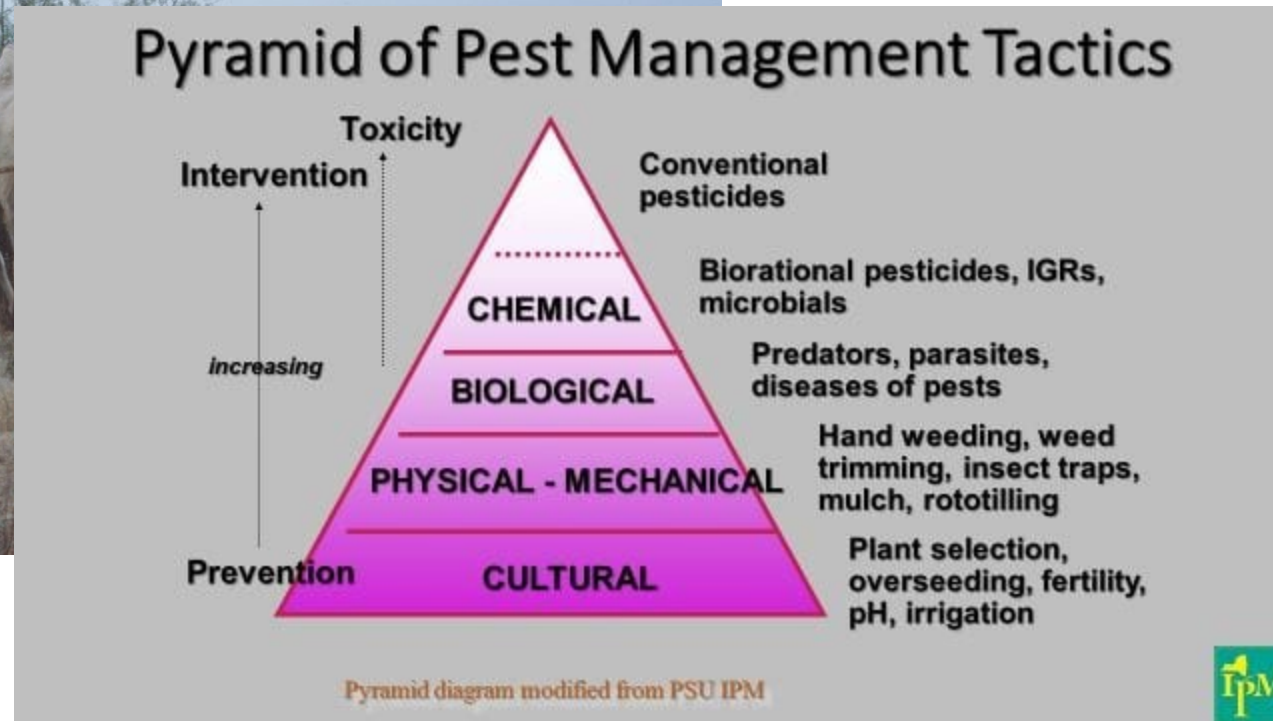
*Tamarix ramosissima*  
saltcedar  
USA







*Tamarix ramosissima*  
saltcedar  
Kazakhstan



*Tamarix ramosissima*  
saltcedar  
USA



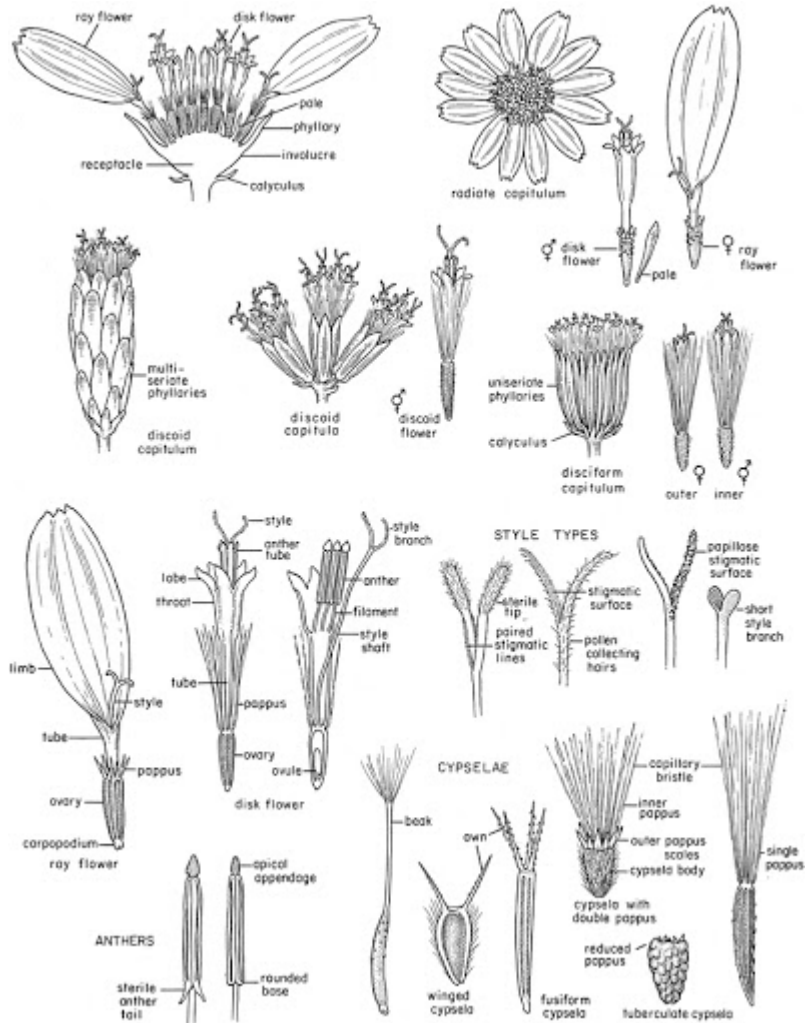


# Snake River

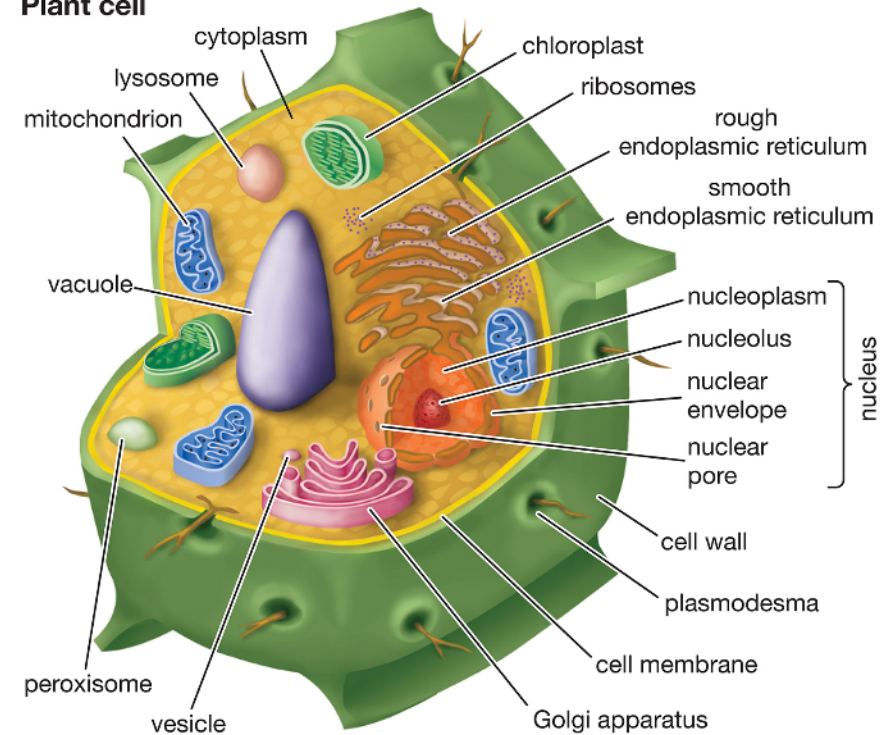




# Morphology can tell us a lot about weeds; DNA can tell us some hidden secrets



**Plant cell**



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# Questions that DNA can help answer

- What species is this weed?
  - Know your enemy



*Euphorbia spp.*  
leafy spurge

# Questions that DNA can help answer

- What species is this weed?
  - Know your enemy
- How does it reproduce?
  - Helps us design control options



*Rhaponticum repens*  
Russian knapweed



# Questions that DNA can help answer

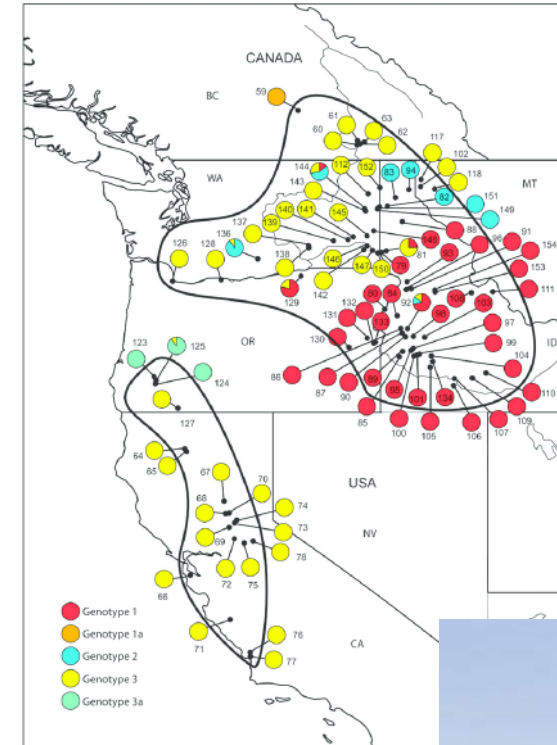
- What species is this weed?
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- How does it reproduce?
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- Are there different genotypes? Hybrids?
  - Differences in resistance or tolerance to control



*Phragmites australis*  
common reed

# Questions that DNA can help answer

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- How does it reproduce?
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- Are there different genotypes? Hybrids?
  - Differences in resistance or tolerance to control
- How are genotypes distributed?
  - What control to use where



*Chondrilla juncea*  
rush skeletonweed





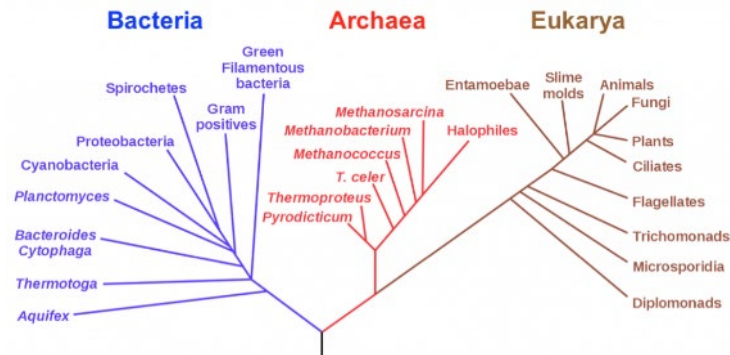
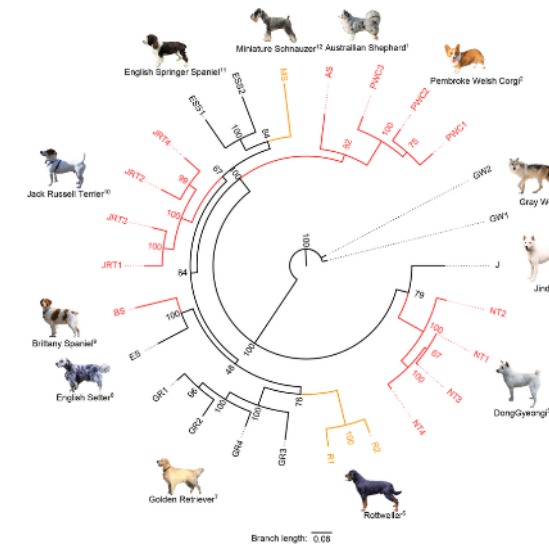
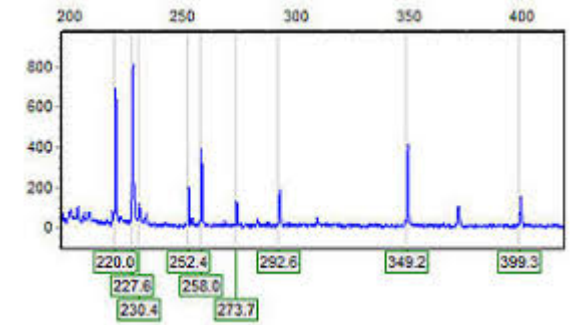
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- Where did the invasion come from?
  - Finding host-specific enemies



# Which DNA method/marker?

- DNA sequencing
- AFLPs
- RAPDs
- microsatellites
- RAD-seq
- ...

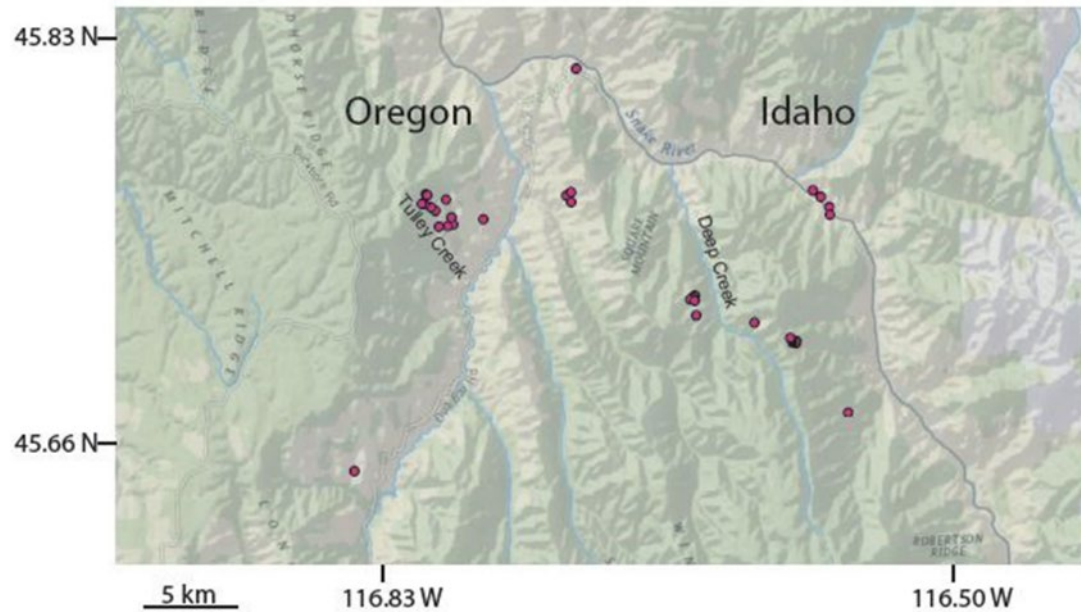




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# What thistle is this?



Hells Canyon  
40km upstream from WA





# Refresher course:

## *Carduus* in N. America

All non-native, state noxious

- *Carduus nutans*
- *Carduus acanthoides*
- *Carduus crispus*
- *Carduus tenuiflorus*
- *Carduus pycnocephalus*





- Carduus nutans
- Carduus acanthoides
- Carduus crispus
- *Carduus tenuiflorus*
- *Carduus pycnocephalus?*
- *Carduus? sp.???*



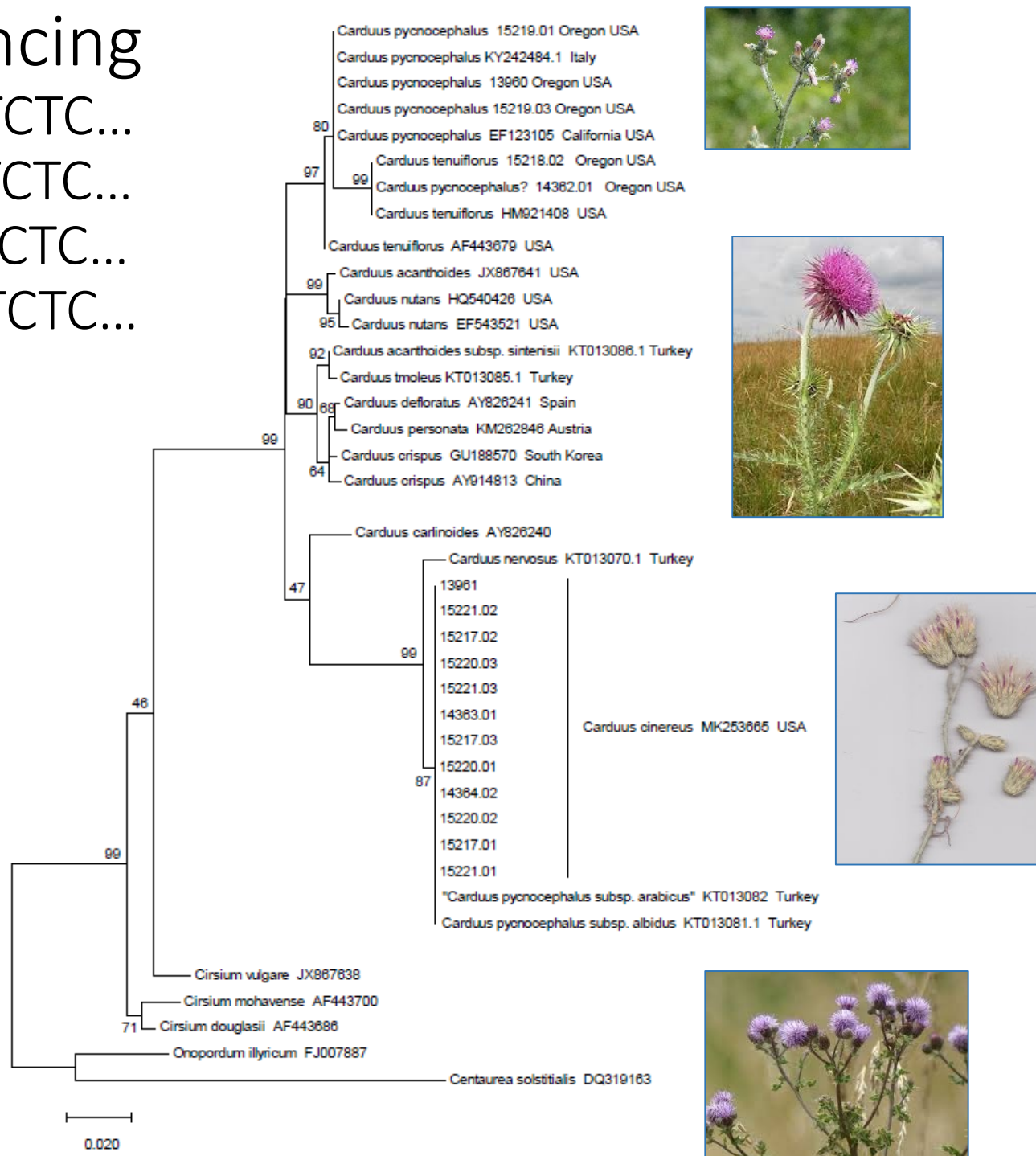
# DNA sequencing

...GATC**A**GACGTCTC...

...GATC**C**GACGTCTC...

...GATC**T**GACGTCTC...

...GATC**G**GACGTCTC...



- *Carduus pycnocephalus*??
  - Garbage can... No.
- Hybrid?
  - Nuclear DNA. No.
- Eastern Euro *C. pycnocephalus*?
- *Revise the genus!*
- ***Carduus cinereus*!!**
- Is it invasive?
  - *Carduus* a notorious genus

MADRONO, Vol. 66, No. 4, pp. 142-147, 2019

## *CARDUUS CINEREUS* (ASTERACEAE) – NEW TO NORTH AMERICA

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# Reproductive mode of Russian knapweed



Invasive Plant Science and Management 2017 10:119–124  
© Weed Science Society of America, 2017



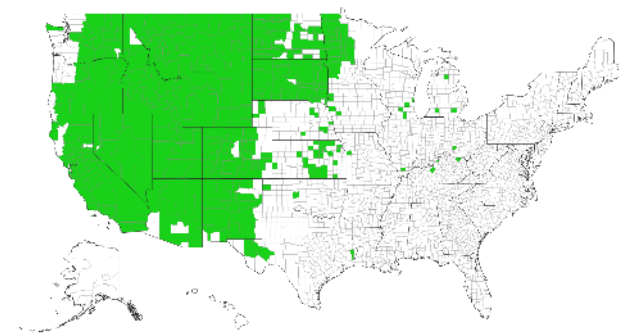
**Invasive Russian Knapweed  
(*Acroptilon repens*) Creates Large  
Patches Almost Entirely by  
Rhizomic Growth**

John F. Gaskin and Jeffrey L. Littlefield\*





Outcrossing  
Selfing  
Apomictic  
Clonal (vegetative)









# How is this plant spreading?

- Locally
- Long distance

# Why do this research?

- Mode of reproduction at times unknowable without genetic analysis
- Effective management can require knowledge of reproductive mode
  - Classical biological control
  - Timing of chemical or manual control
    - E.g. Fall is better spray time maybe but too late for seed control?
- Sometimes results are not what you would predict...



# Goal: Distinguish ramets from genets to test hypotheses:



- If clonal spread, plants genetically identical
- If spread by seed (and self-incompatible), all plants genetically different.
- Some combination of two above.

# Russian knapweed

- Known to reproduce clonally from rhizomes (Watson 1980)
- Also produces seed (50-1200/plant; Beck 2001 Ivanova 1966)
- Self-incompatible (Young & Clements 2003; Harrod & Taylor 1995)
- How much local spread is from vegetative vs. seed propagation? How big can clones get?



*Aulacidea acroptilonica*

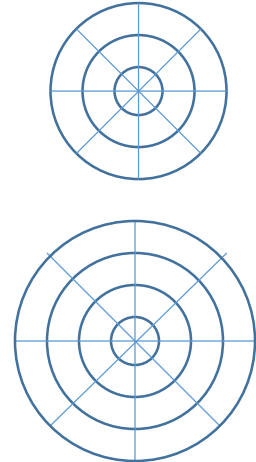


*Jaapiella ivannikovi* gall



# Methods

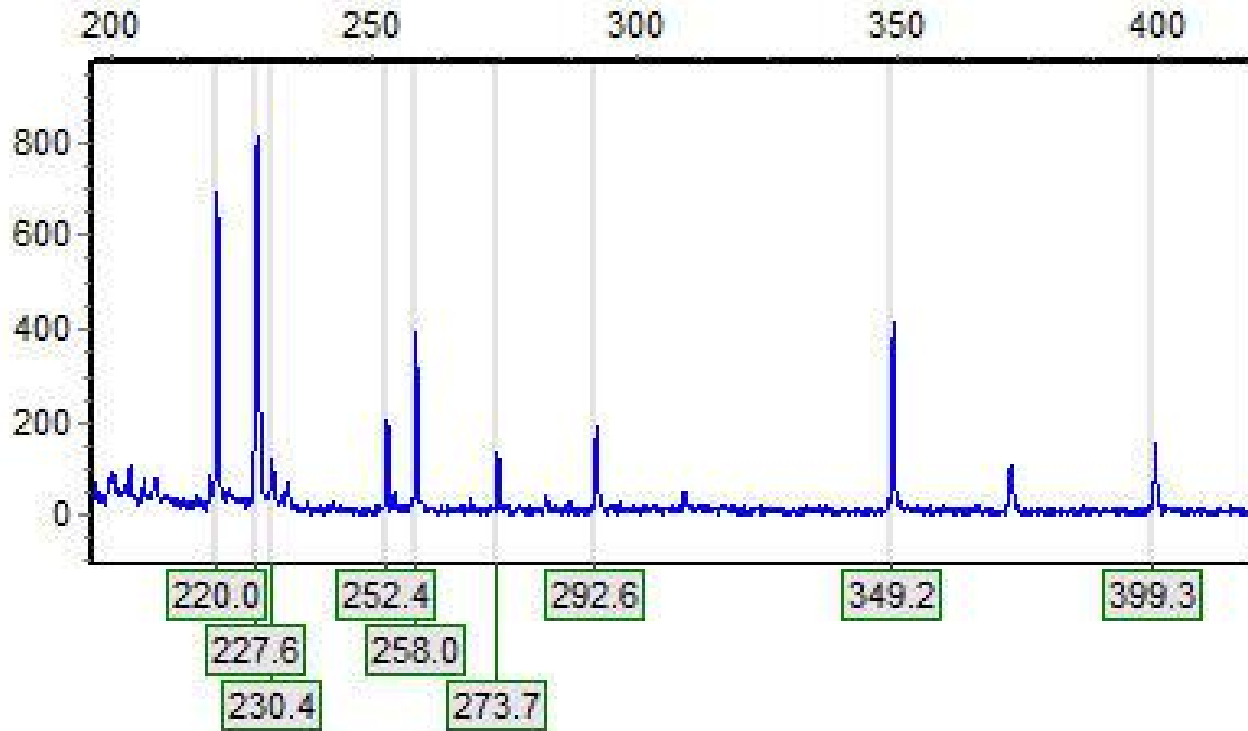
- Collected leaf tissue from 174 shoots from 6 dense patches, 10-56m diameter
- 25-33 samples per patch
- AFLPs (54 loci)



# AFLPs

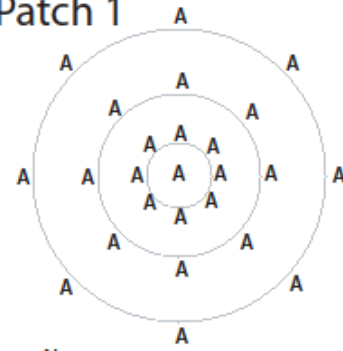
## Amplified Fragment Length Polymorphisms (DNA fingerprint)

- Extract DNA from leaves
- Cut at restriction sites  
TCGT cut TCCGTA
- Make multiple copies of fragments so signal is strong enough
- Run on fragment analyzer (sequencing machine)
- 1, 0 data
- AFLPs are repeatable
- AFLPs are high resolution

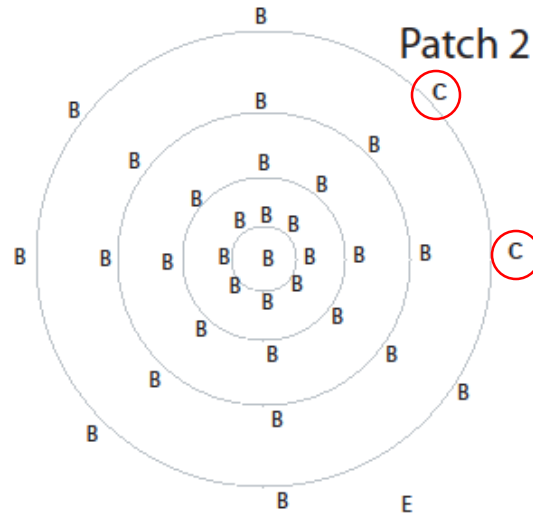




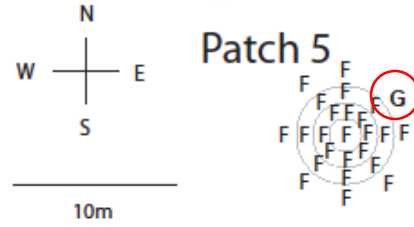
Patch 1



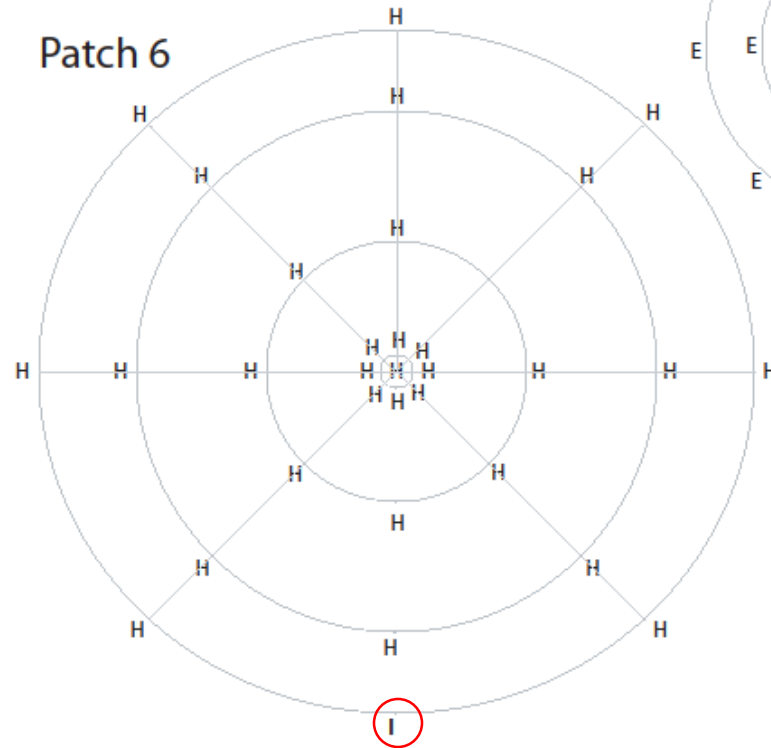
Patch 2



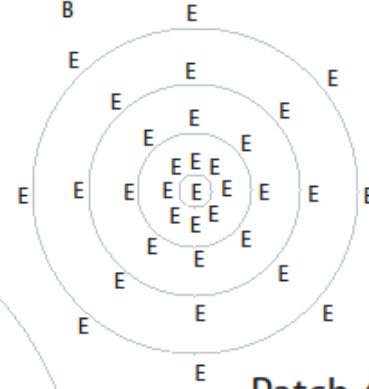
Patch 5



Patch 6



Patch 4



Patch 3

# Conclusions

- Patches are typically one individual plant
- Largest genet size 56m, as big as our biggest patch (unexpected result)
- There are likely even larger individuals

# Conclusions

- New establishment of patches (n=6) likely by seed
- Control of existing patches will have to focus on methods that are effective against root mass
- Control of establishment of patches relies on destruction of seed in non-tilled/harvested environment



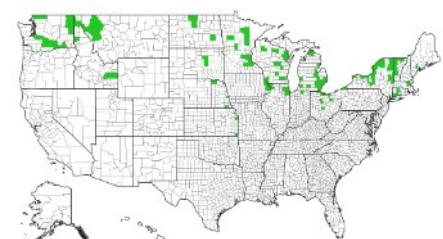


# Questions that DNA can help answer

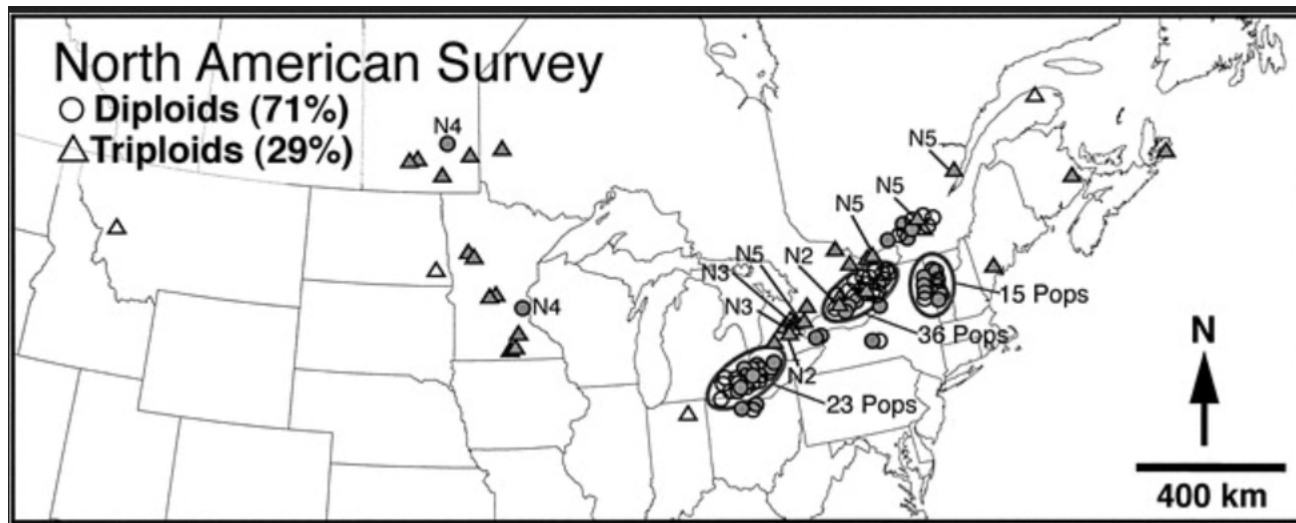
- What species is this weed?
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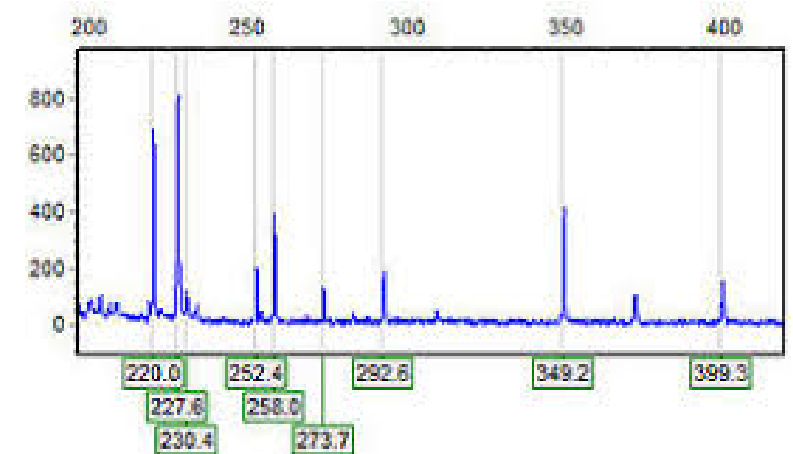
Flowering rush  
*Butomus umbellatus*





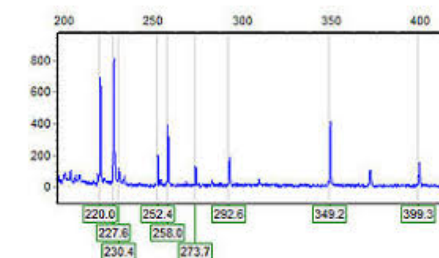
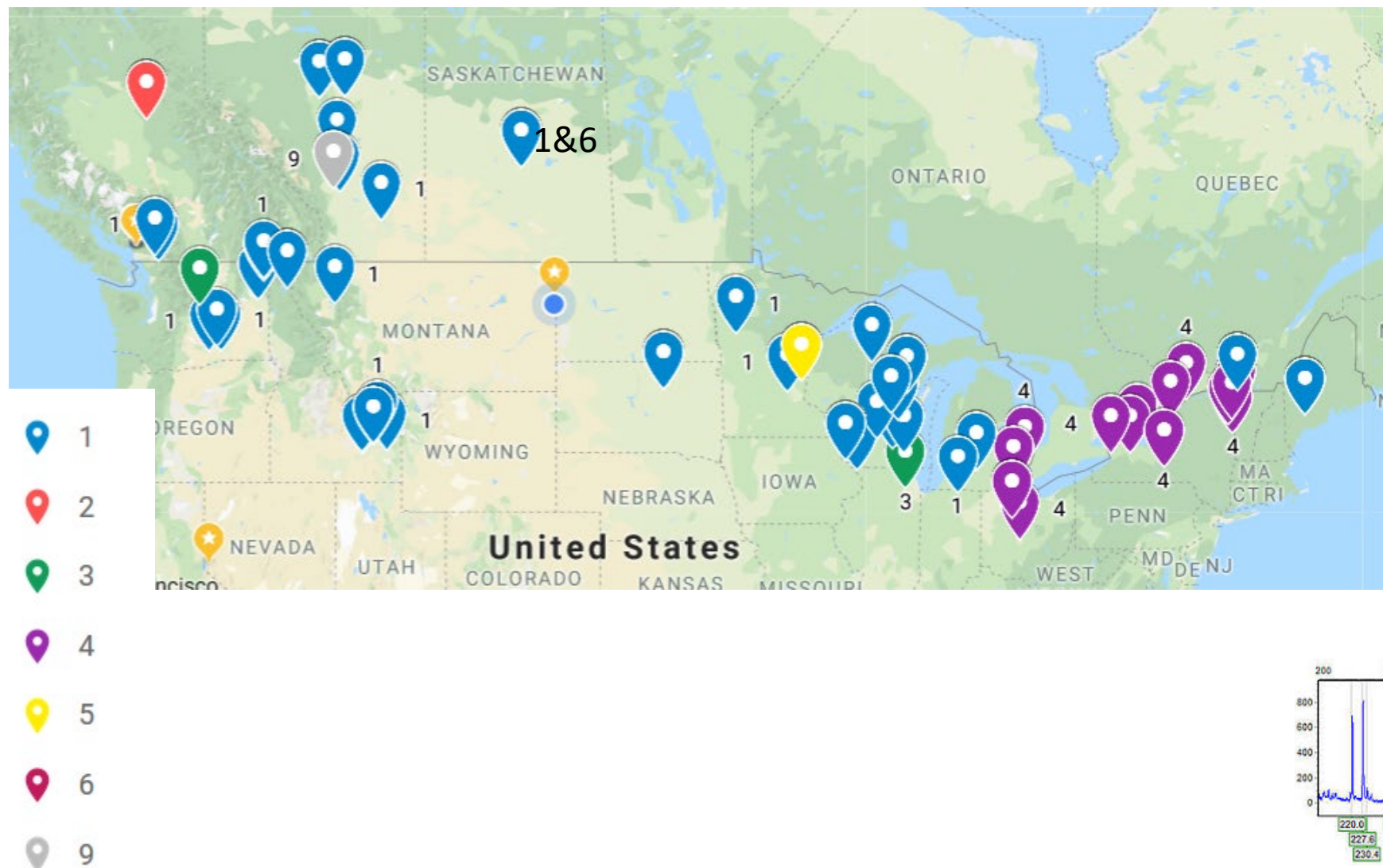


Kliber and Eckert 2005.



AFLPs

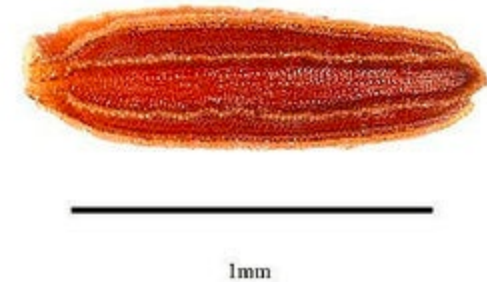
# How are they distributed?





# Who cares about ploidy!?

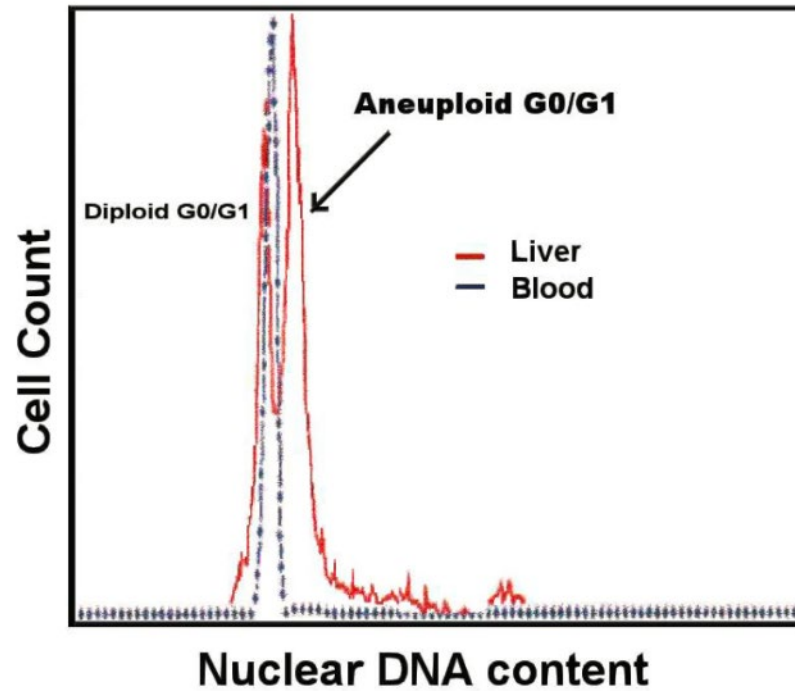
- Reproduction and spread
  - 2x (diploid): seed and bulbil and fragments
  - 3x (triploid): only root fragments?
- Invasiveness?
- Control?
  - Chemical
  - Biological



# Diploid or triploid?

Fresh tissue

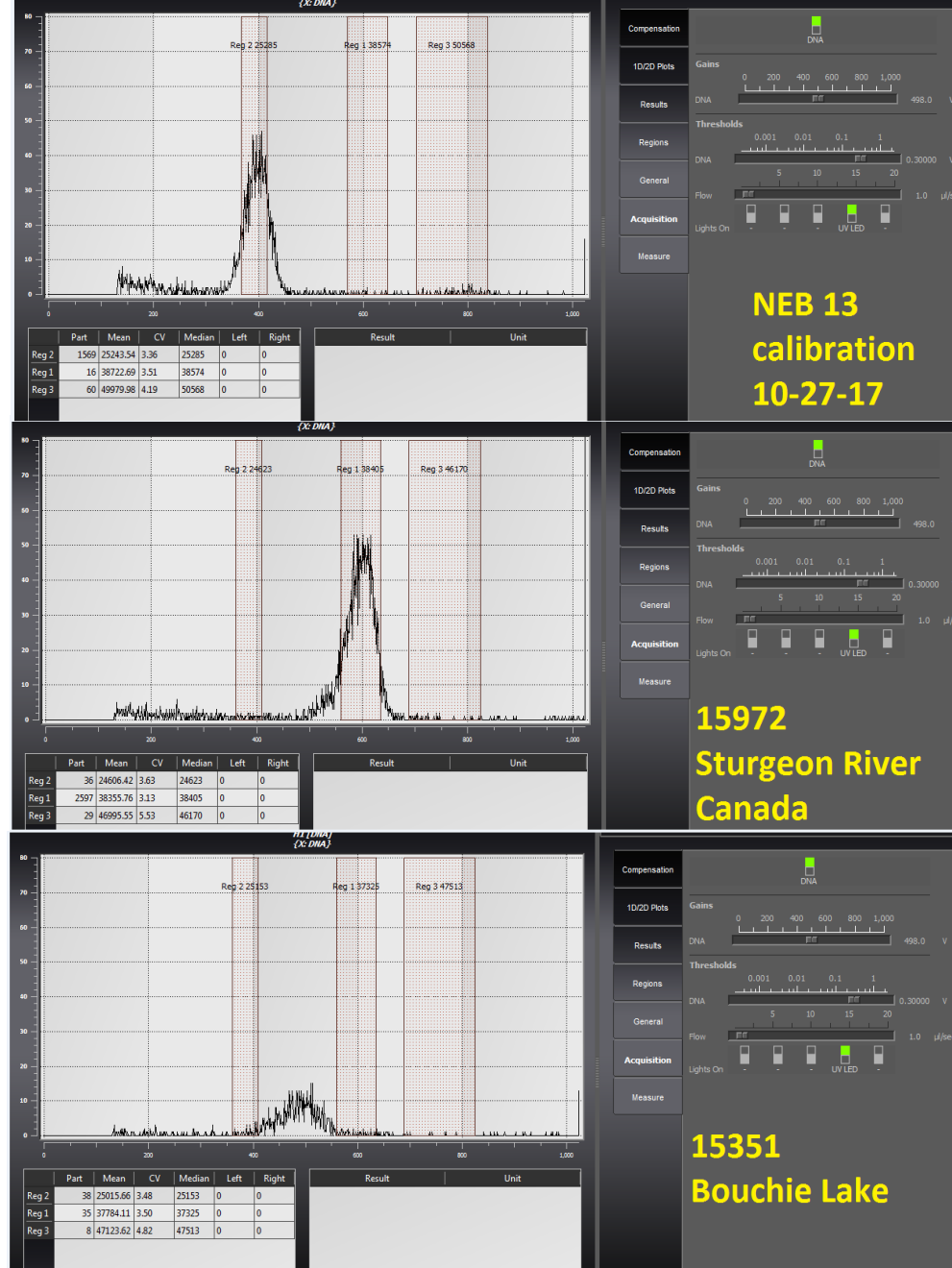
Can also tell by stomata



2x= 400  
3x=600

Fresh leaf  
material

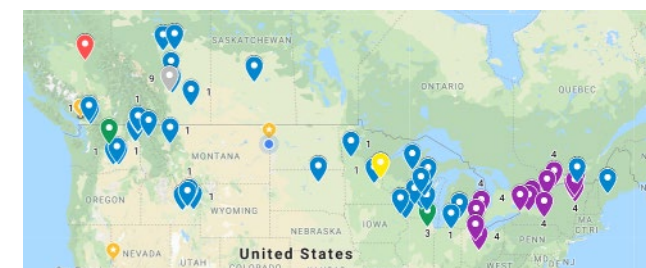
Grrr. Stringy tissue?



Genotype 4 VT

Genotype 1

Genotype 2





Species Code	ARS PMRU DNA Sample #	# Root Tip Evaluations	# Cell Evaluations	Chromosome Count (2n=)
BUUM	15348.00 Yak-Pross	2	15	26
BUUM	15349.00 Yak-Pross	2	15	26
BUUM	15350.00 Yak-Pross	2	15	26
BUUM	15351.00 Bouchie	2	16	39

### 3-D analysis

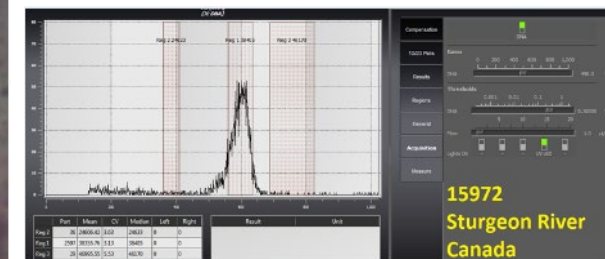
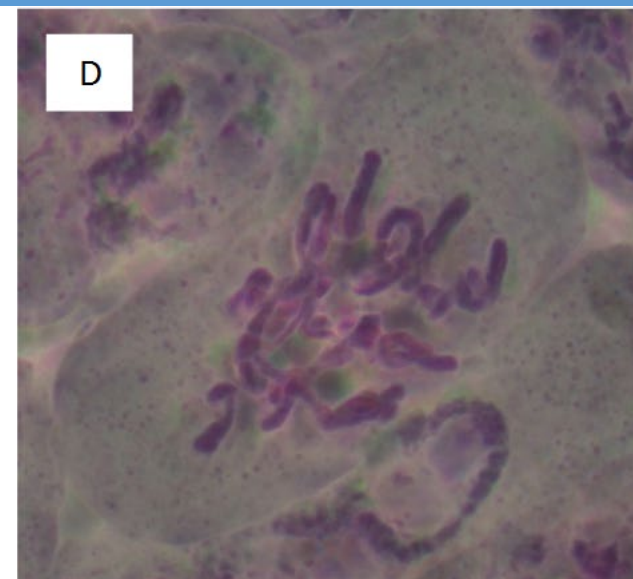
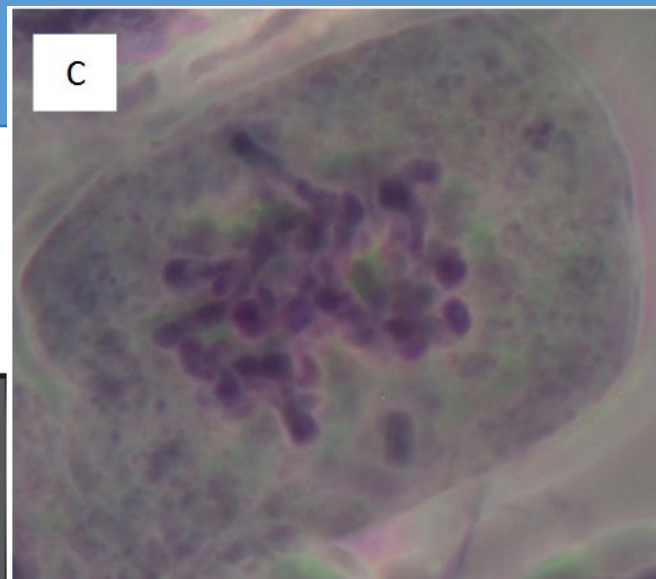
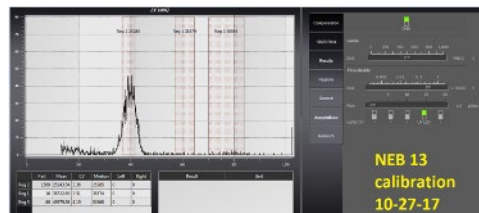
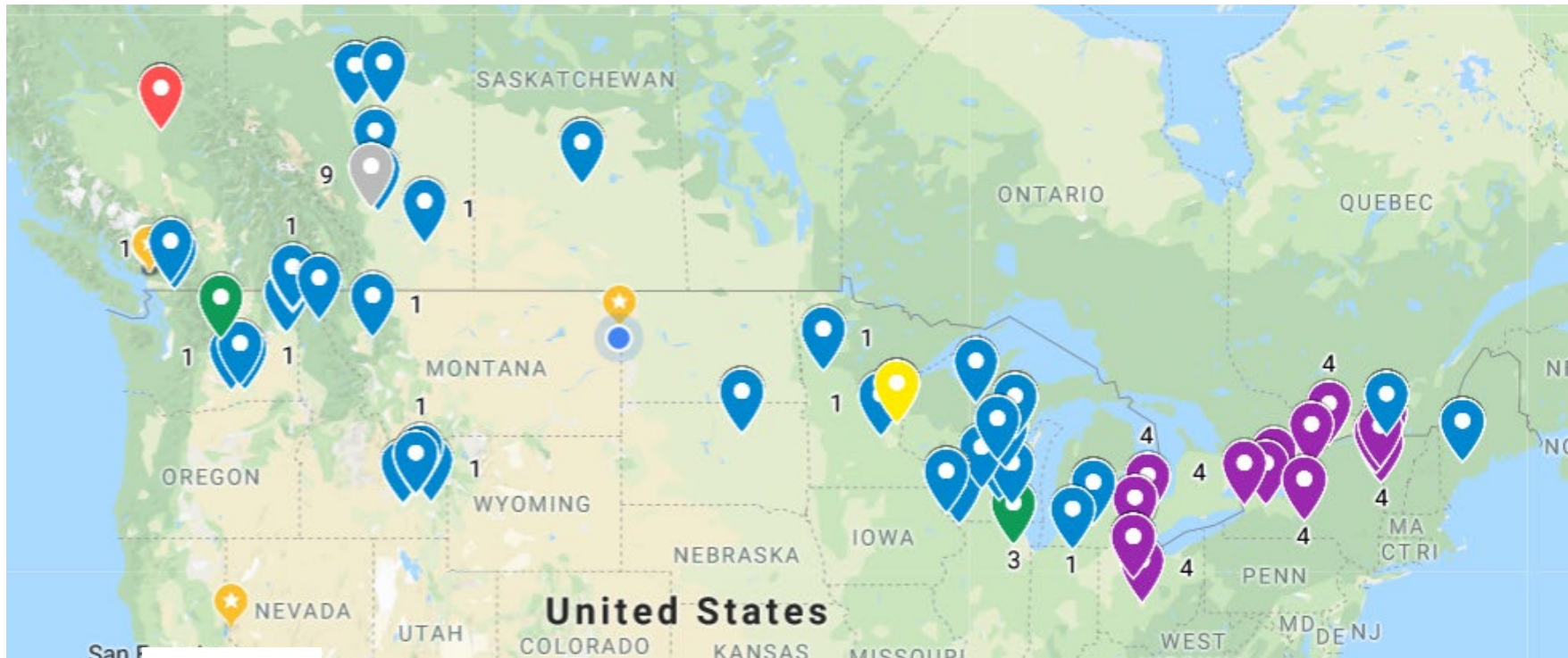


Figure 1. Composite cell images from *Butomus umbellatus* with A-C) 2n=26 chromosomes, D) 2n=39 chromosomes.

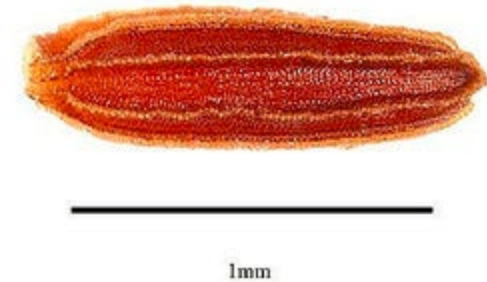


- 📍 1 3x (2x at Prosser Diversion dam) [Jennifer Parsons]
- 📍 2 3x
- 📍 3 2x
- 📍 4 2x
- 📍 5 2x
- 📍 6 ?
- 📍 9 ?

yellow star=favorite restaurants

# Who cares about ploidy!?

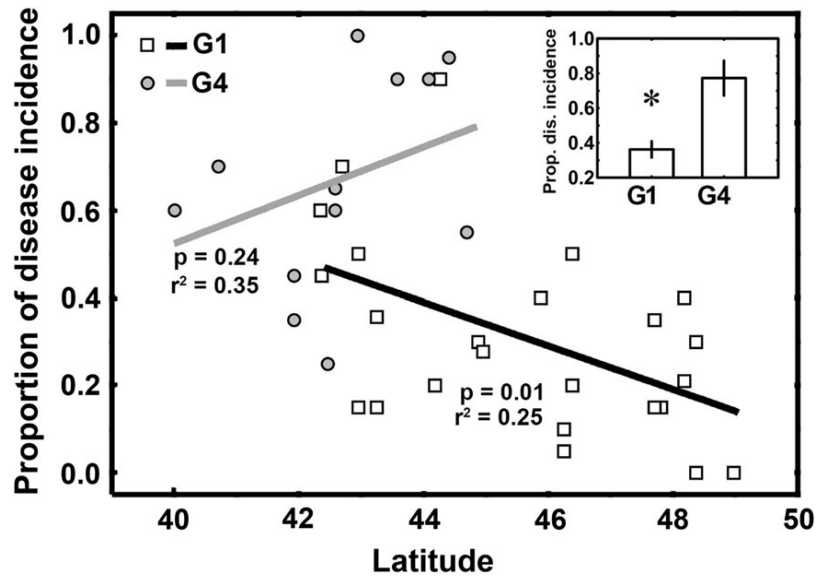
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- Control?
  - Chemical
  - Biological







Flowering rush NA  
diseases. Does genotype  
matter?



Harms, N., Shearer, J., Cronin, J.T. and Gaskin, J.F., 2019. Geographic and genetic variation in susceptibility of *Butomus umbellatus* to foliar fungal pathogens. *Biological Invasions*, pp.1-14.

# Questions that DNA can help answer

- What species is this weed?
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- Are there different genotypes? **Hybrids?**
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# Genetics of Knotweeds

Biol Invasions  
DOI 10.1007/s10530-014-0652-y

ORIGINAL PAPER

## **Extreme differences in population structure and genetic diversity for three invasive congeners: knotweeds in western North America**

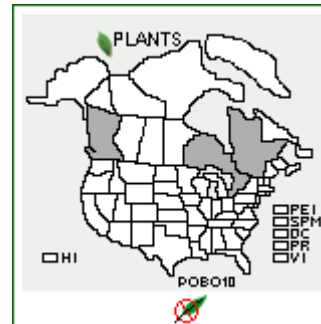
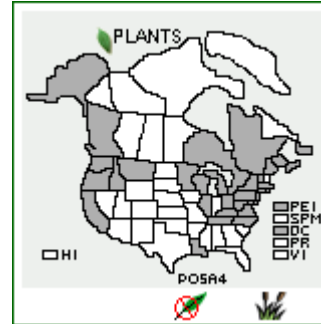
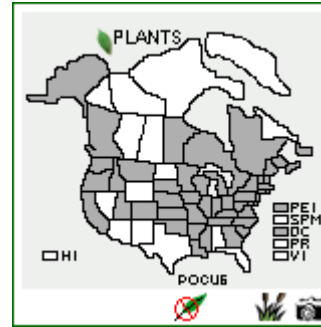
John F. Gaskin • Mark Schwarzländer •  
Fritzi S. Grevstad • Marijka A. Haverhals •  
Robert S. Bouchier • Timothy W. Miller





# Background info: Three knotweed taxa

- Japanese
  - *Fallopia japonica*
- Giant
  - *Fallopia sachalinensis*
- Bohemian (hybrid of the two above)
  - *Fallopia x bohemica*





# Knotweeds in western North America

1. How are species distributed in western North America?
2. Can managers distinguish the different species in the field?



# Biocontrol update



*Aphalara itadori*

- A psyllid biological control agent for Japanese knotweed has been released in the U.K. (Shaw et al. 2009) and Canada (and USA)
- Two strains of the psyllid that are being considered for use in Canada/USA have contrasting preferences for Japanese, giant and Bohemian knotweed (Grevstad et al. 2013).
- The psyllid also shows differential development on different Bohemian knotweed collections (Bourchier and Grevstad pers. obs.)

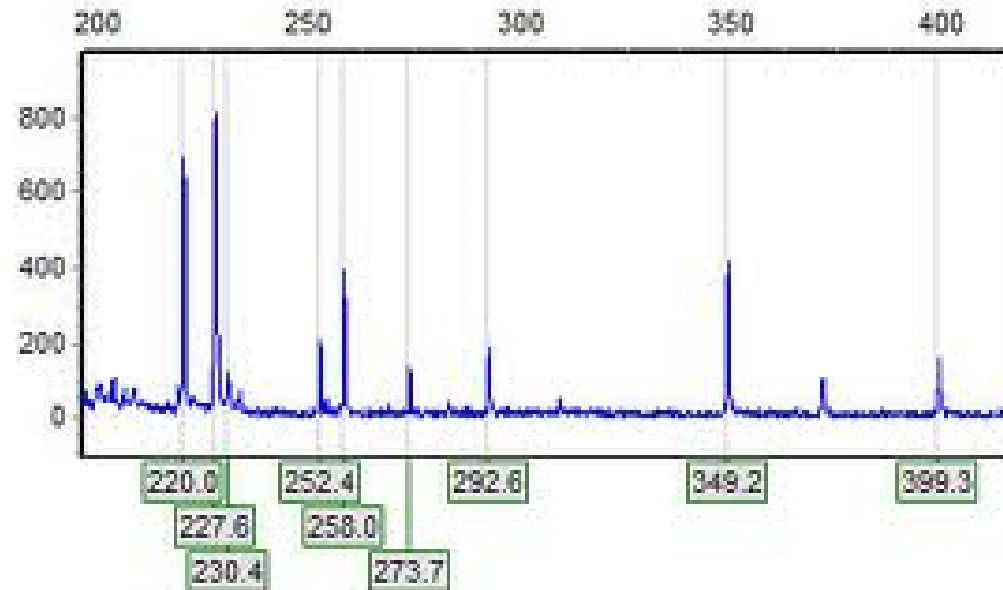


# Key to Identification of Invasive Knotweeds in British Columbia

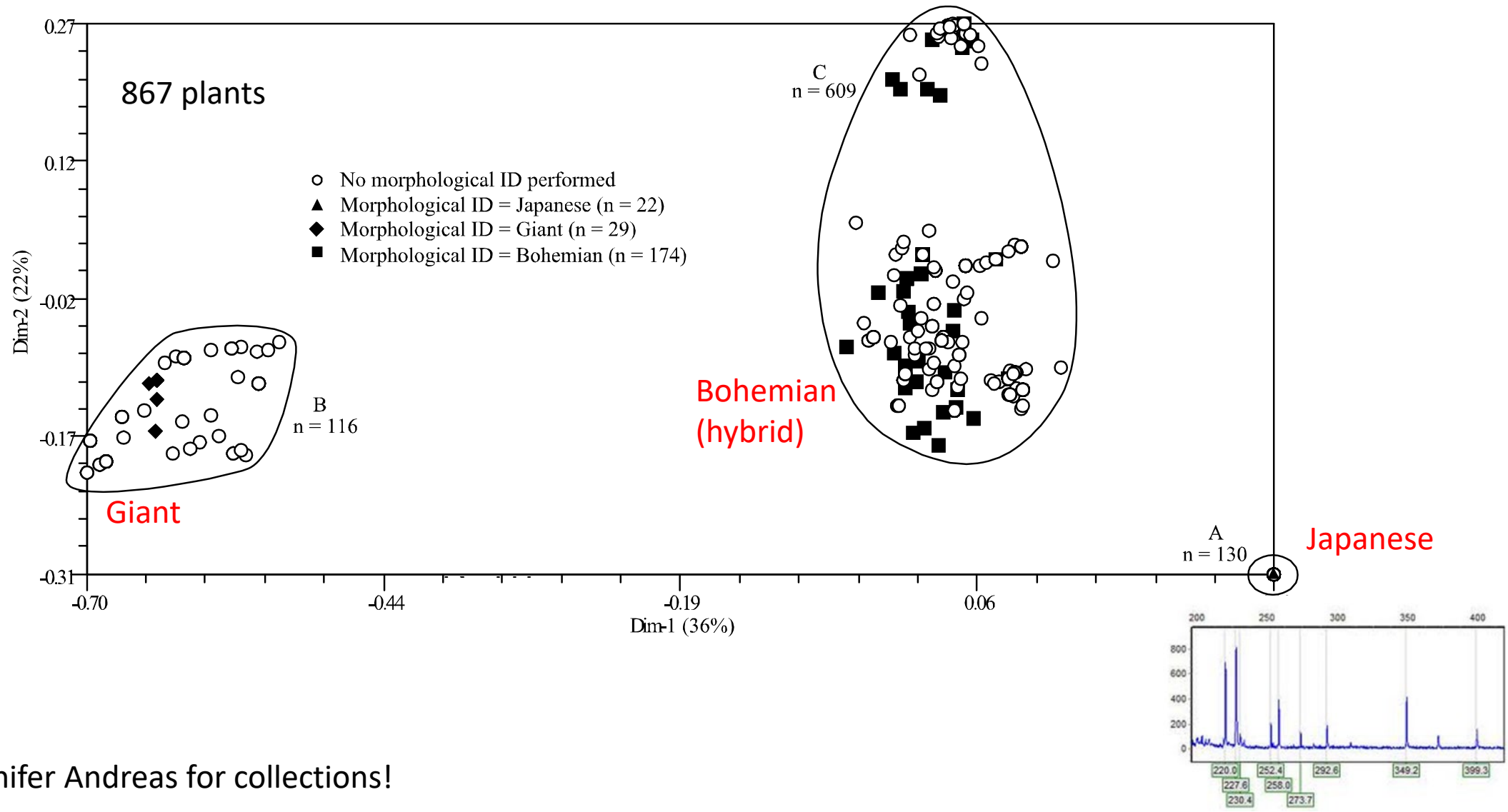


# DNA Method

- Amplified Fragment Length Polymorphisms (AFLPs)
  - 135 repeatable, variable loci, presence/absence (1/0)



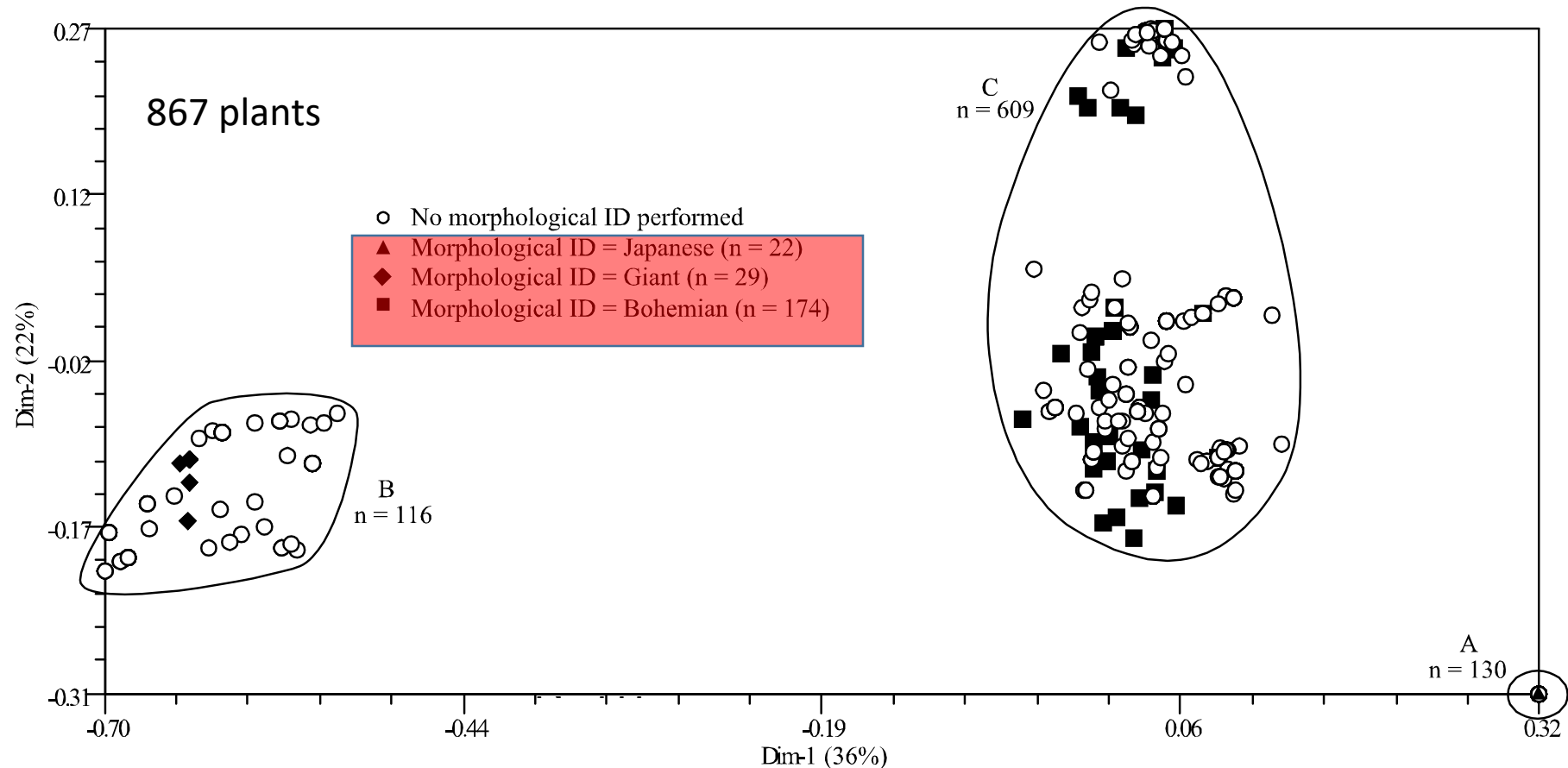
# Principle coordinates analysis of AFLPs



Thanks to Jennifer Andreas for collections!

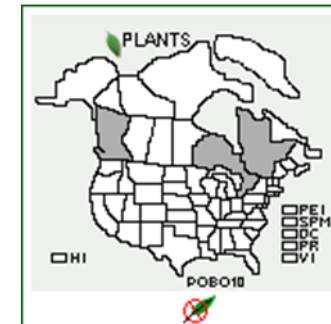
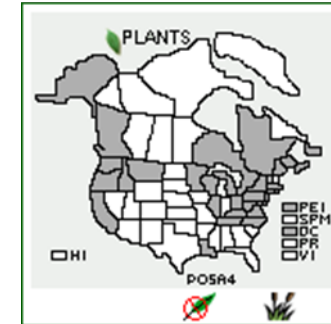
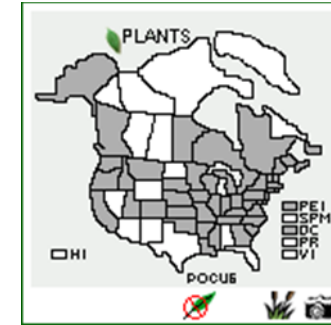


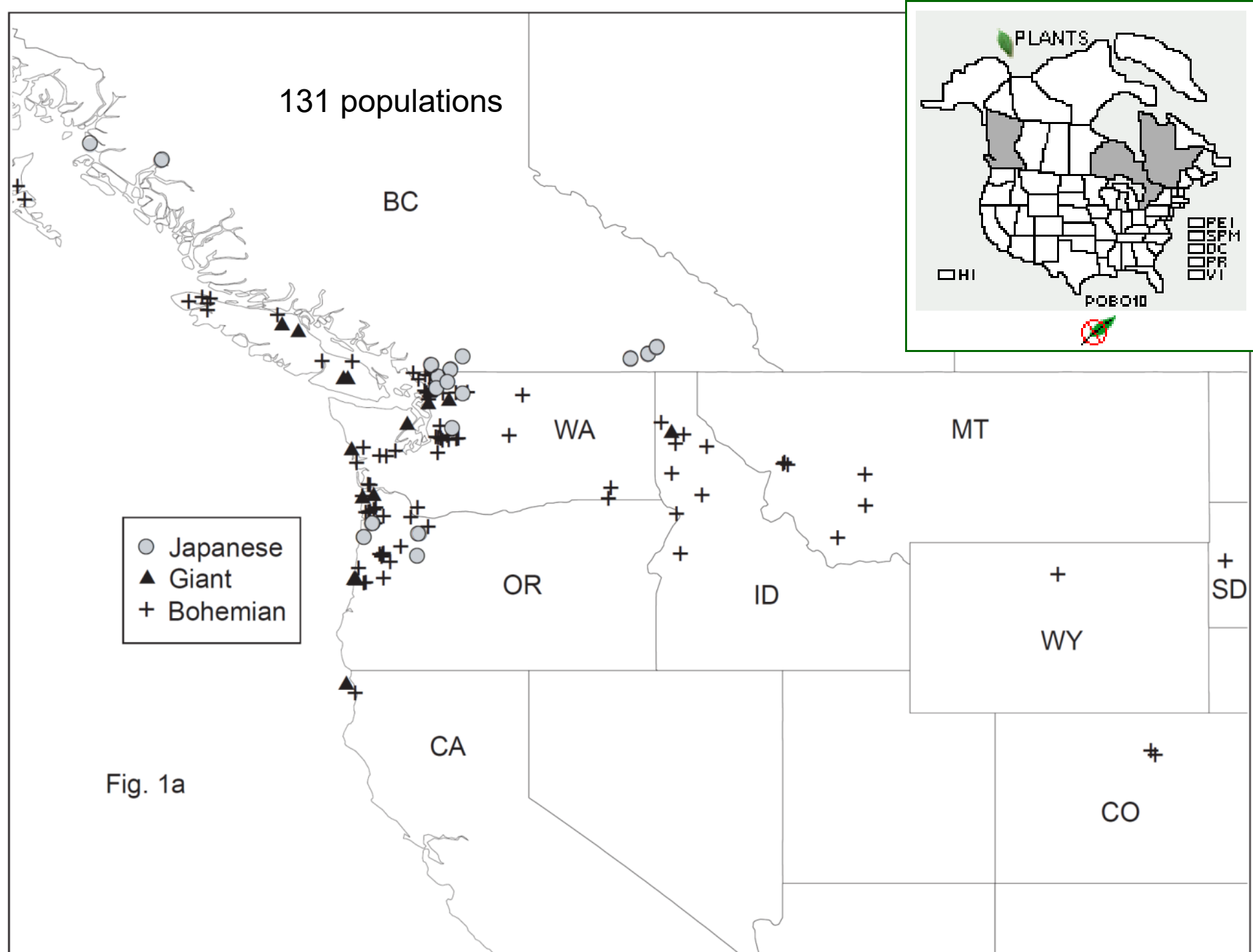
# Principle coordinates analysis of AFLPs



Can managers identify species?  
N=222 identified in field, even in mixed pops.

- How are the taxa distributed?

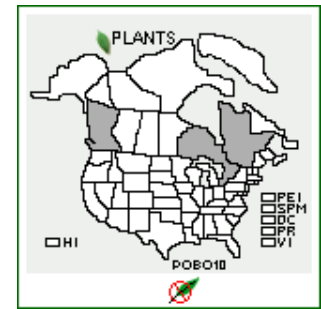






# Implications for biocontrol in the Northwest?

- Plants can be identified with morphology.
- Host-specificity test lineages are genotyped.
- Bohemian most common, most widespread, and most genetically diverse in PNW. Priority target??
- Japanese knotweed is one genotype (Same genotype as in UK).

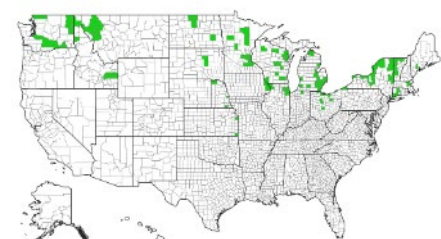


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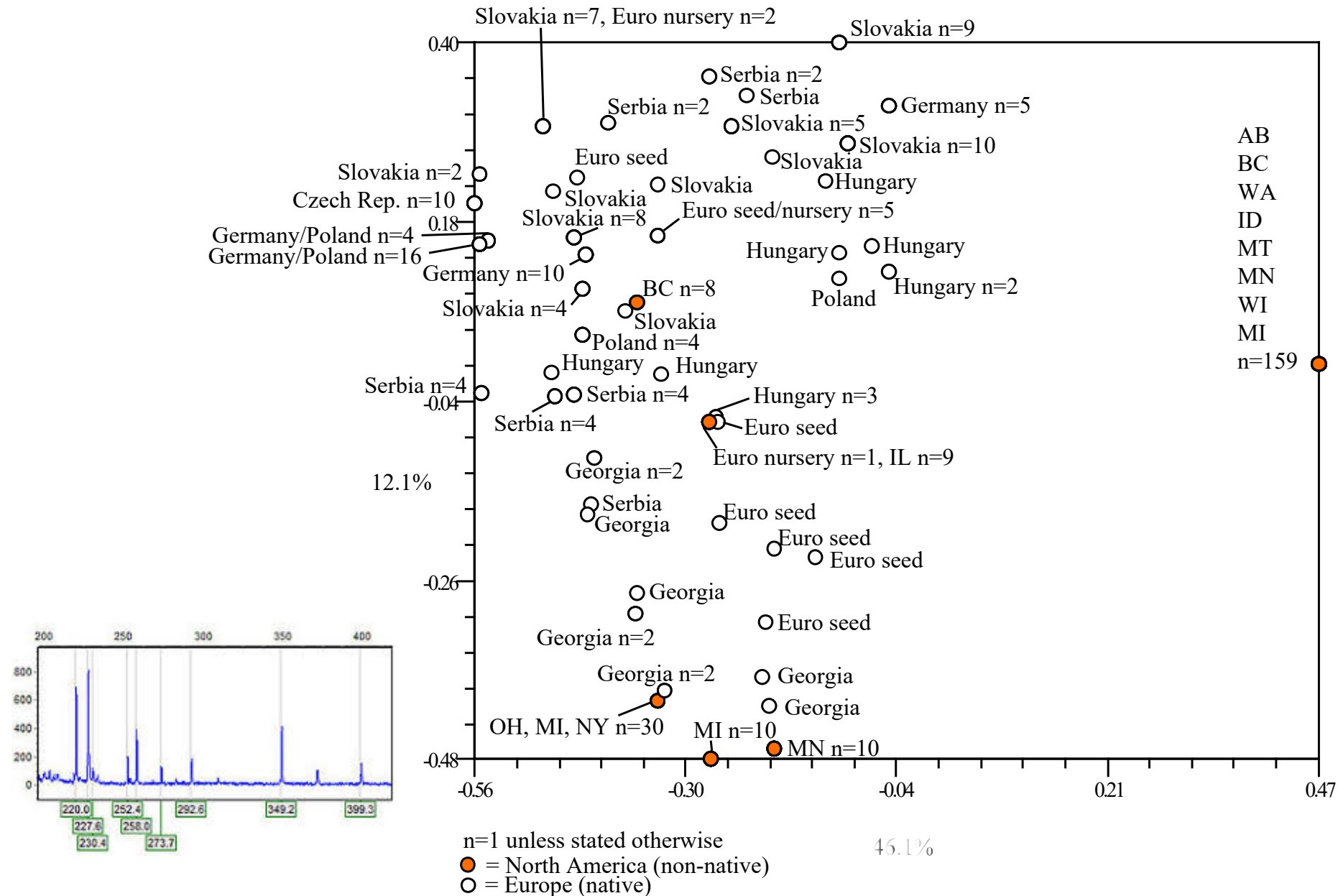


Flowering rush  
*Butomus umbellatus*



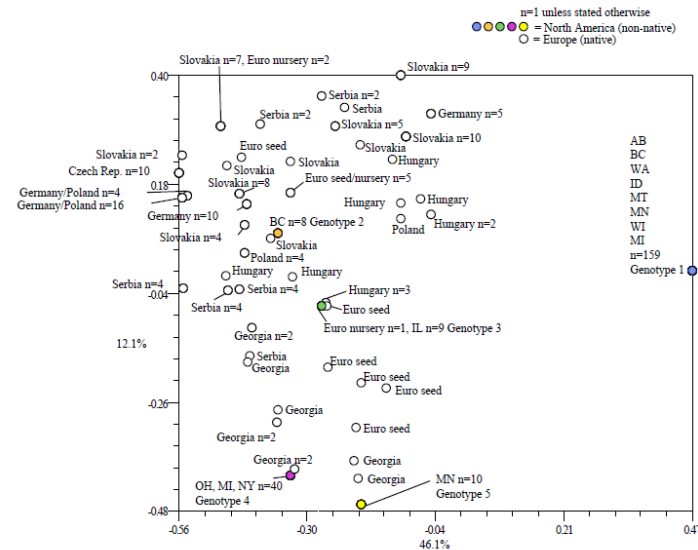


# Butomus (flowering rush) AFLP genotype similarity - PCOA



# Biocontrol with foreign disease (CABI UK)

## Will a disease attack all of our *Butomus*?



No close genetic match in Eurasia for our genotype 1...yet.

- *Doassansia niesslii*
- Smut
- Does not attack our genotype 1...need to find 1 in Eurasia...collect disease from it

# •Questions?

- John Gaskin
- USDA Agricultural Research Service, Sidney, Montana
- Population genetics, origins, and hybridization of invasive plants.



Saltcedar



Perennial  
pepperweed



Rush  
skeletonweed



Casuarina

