

Using genetics to inform invasive weed management

Tales from flowering rush, common mullein, and a new thistle species



John Gaskin
Botanist
Sidney, MT

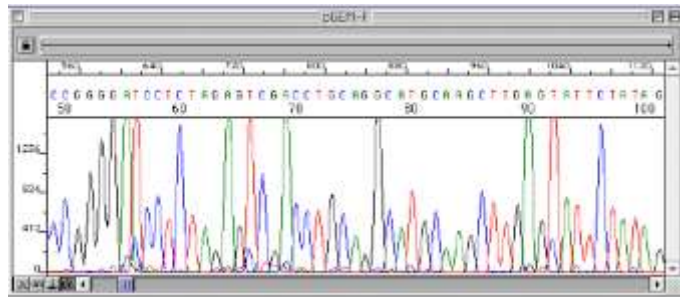


Questions that DNA can help answer

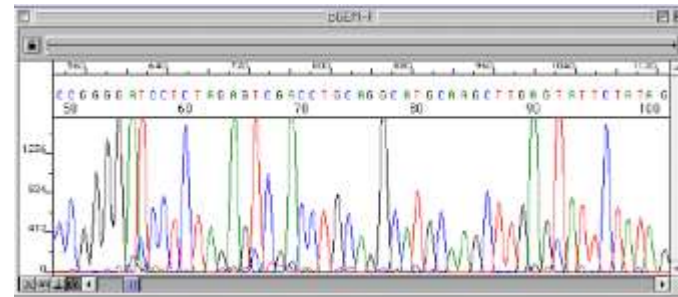
- What species is this weed?
 - Know your enemy
- How does it reproduce?
 - Helps us design control options
- Are there different genotypes?
 - Differences in resistance or tolerance to control
- How are genotypes distributed?
 - What control to use where
- Where did the invasion come from?
 - Finding host-specific enemies

Which DNA method/marker?

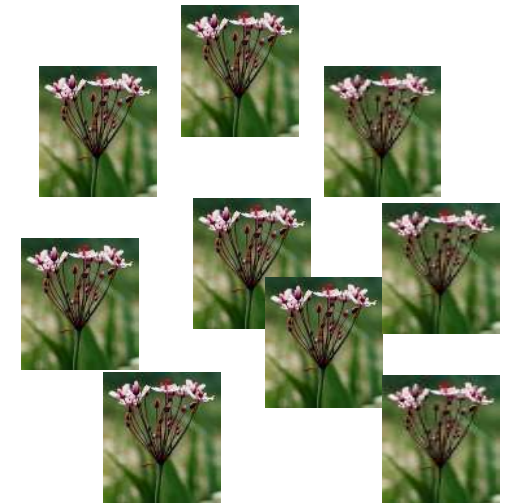
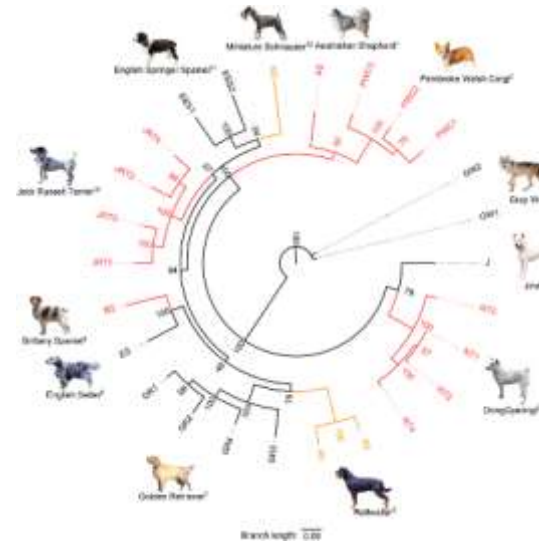
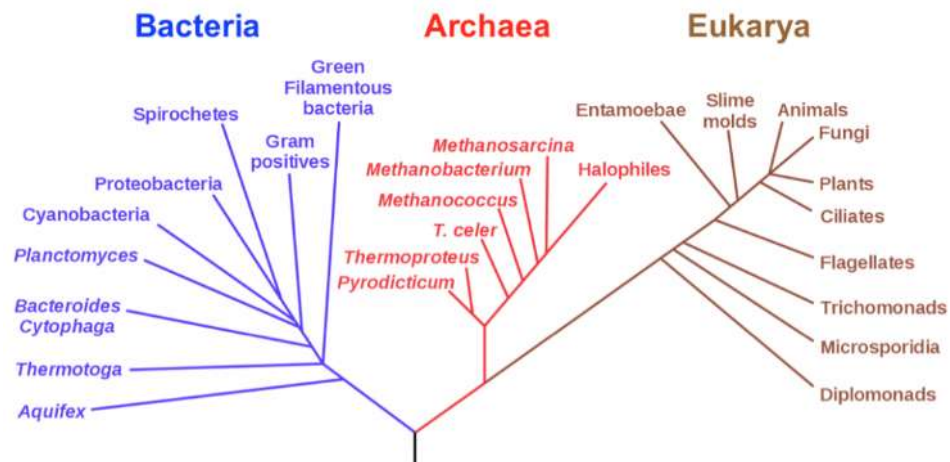
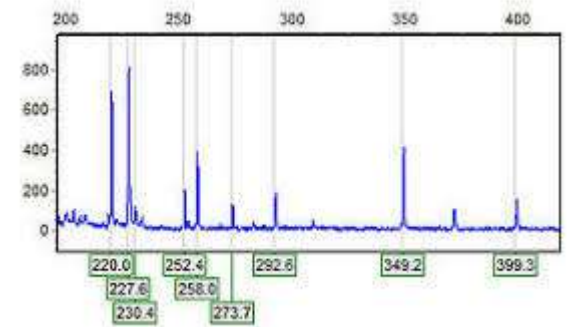
Slowly evolving DNA



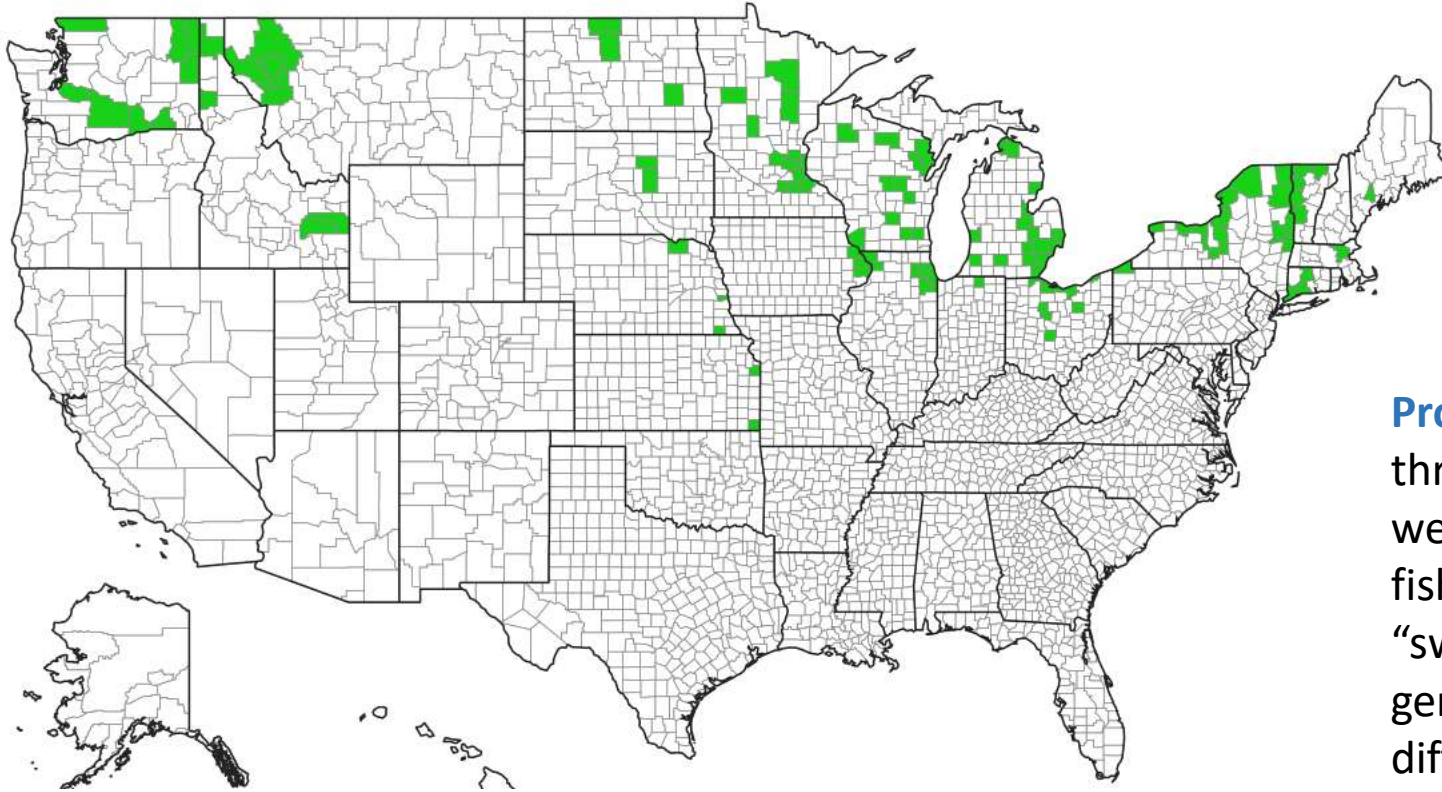
Moderately fast evolving DNA



Really fast evolving DNA
 Microsatellites
 RadSeq
 AFLP



Flowering rush (Butomus)



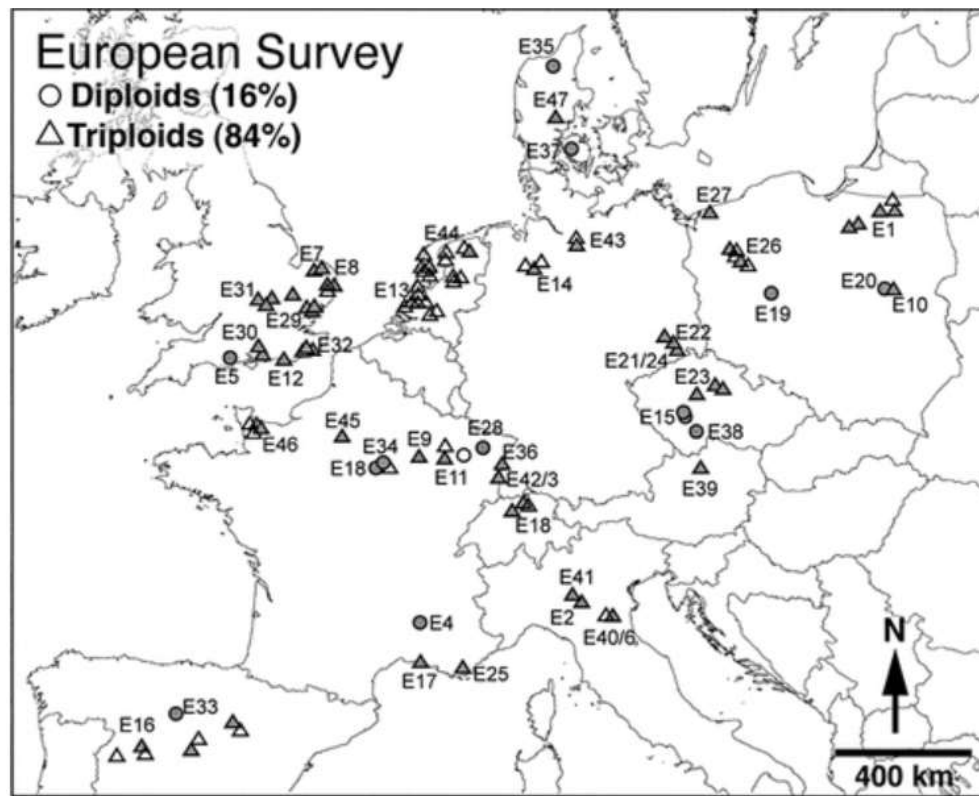
Problem: Flowering rush is now spreading throughout river and lake systems in western North America, eliminating native fish habitat and enhancing diseases such as “swimmers itch”. Different flowering rush genotypes are invading that behave differently in terms of reproduction and management needs, such as potential biological control.

DNA analysis questions

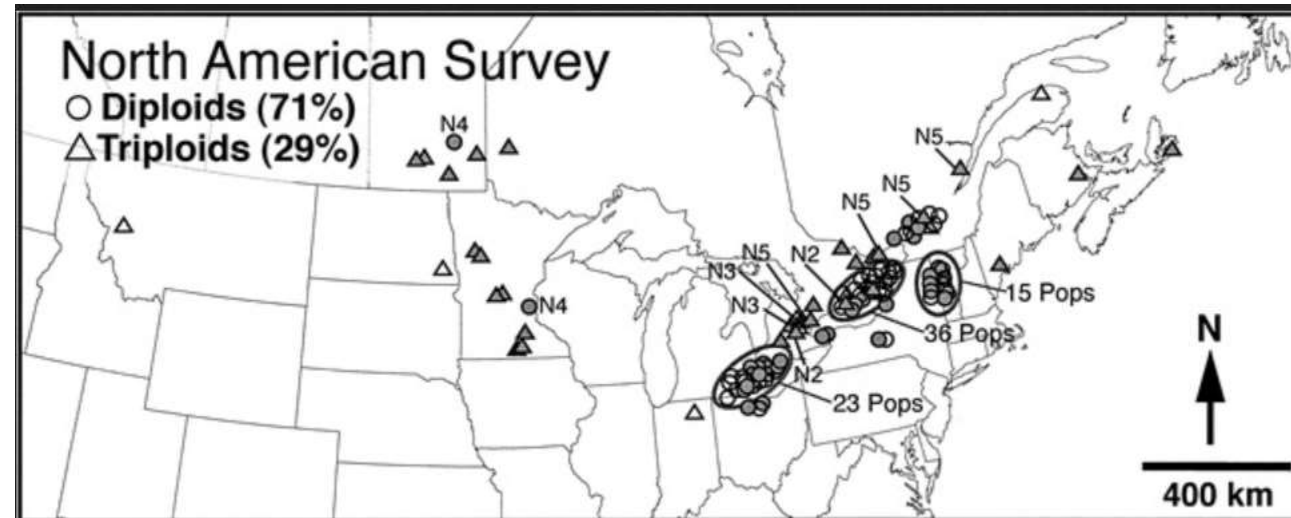
- What ploidy is the invasion (2x-diploid/3x-triploid)?
- What genotypes exist in N. America?
- Where are genotypes distributed?
- Where did genotypes come from in Eurasia?
 - Native to: Afghanistan, China, India, Kashmir, Kazakhstan, Kyrgyzstan, Mongolia, Pakistan, Russia, Tajikistan, Uzbekistan; SW Asia, Europe. More?

We already know some stuff:

- Most studies focused on eastern North America
- Changes from diploid to triploid may be historical and recent and recurring in Europe (some diploid and triploid plants genetically similar). Enzyme study. Kirschner et al. 2002.
- Most natives triploid, most invasives diploid. Kliber and Eckert 2005.
- Both diploid and triploid NA pops usually clonal, even though diploid seed viable. Eckert et al. 2003.

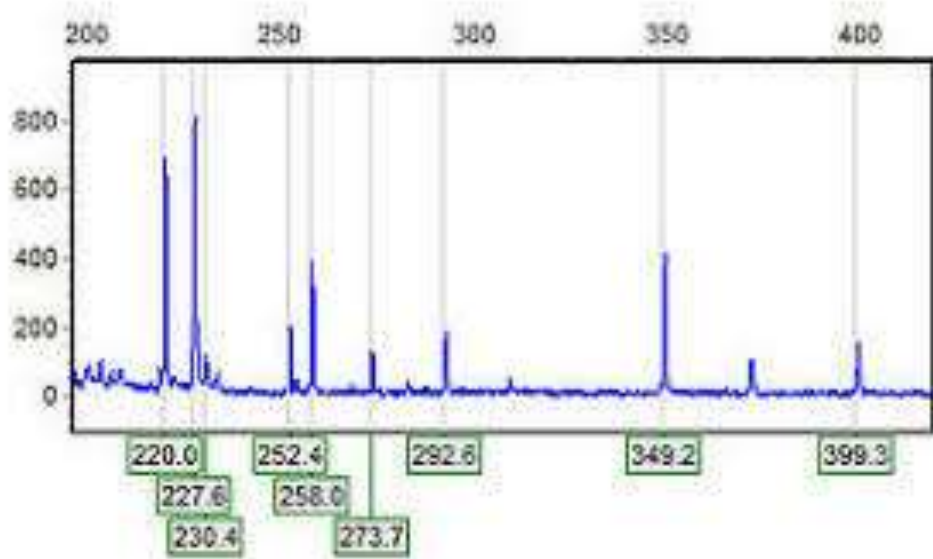


Kliber and Eckert 2005.

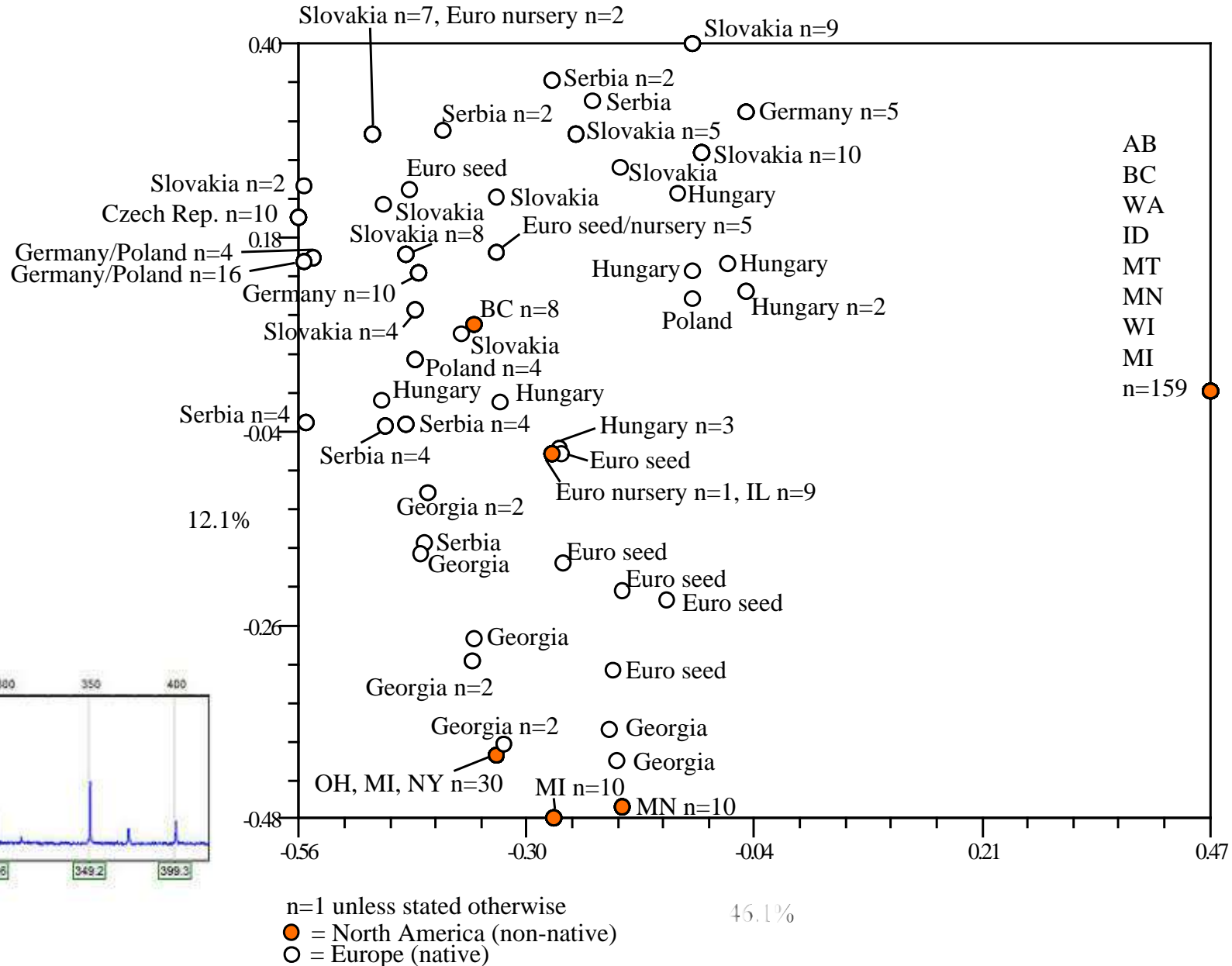


Now, let's use AFLPs to try to answer questions about western North America invasion.

- We have samples from:
 - PNW
 - Europe (looking for matches for biocontrol)
 - Midwest and Eastern USA



Butomus (flowering rush) AFLP genotype similarity - PCOA

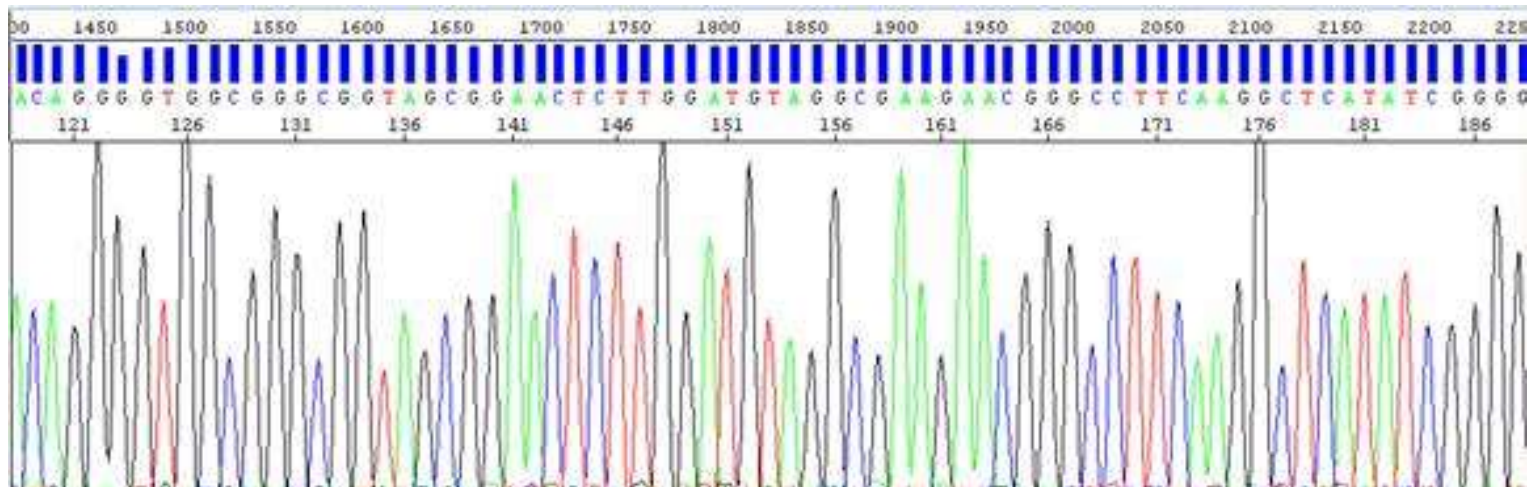


How are they distributed?



There were some unexpected genotypes; are they *Butomus*?

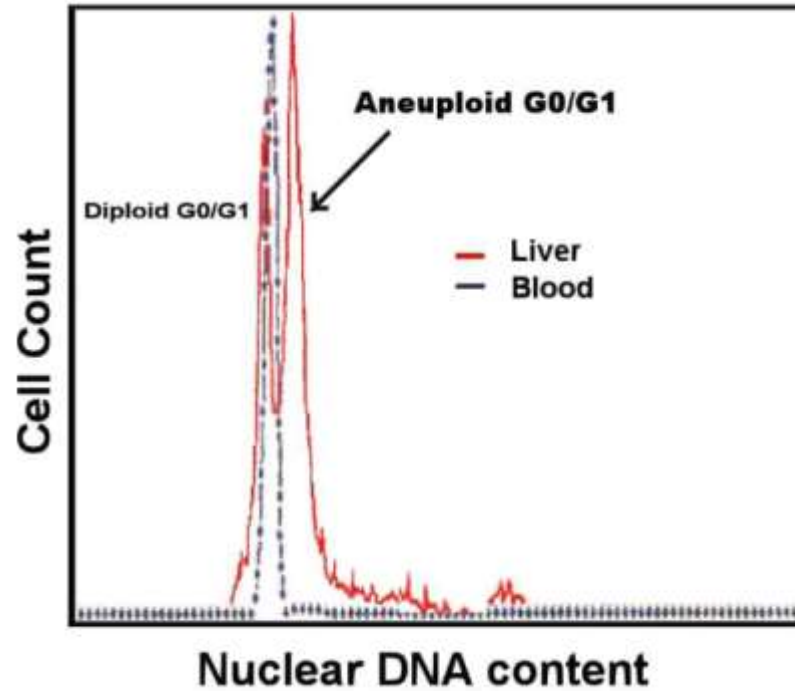
- Some AFLP genotypes are unique and rare (6,7,8,9)
- DNA sequenced (chloroplast marker) them and others (1,2,3,4,5,6,9) identical match to *Butomus* in GenBank
- (7,8) *Sparganium* (aquatic bur-reed)



Diploid or triploid?

Fresh tissue

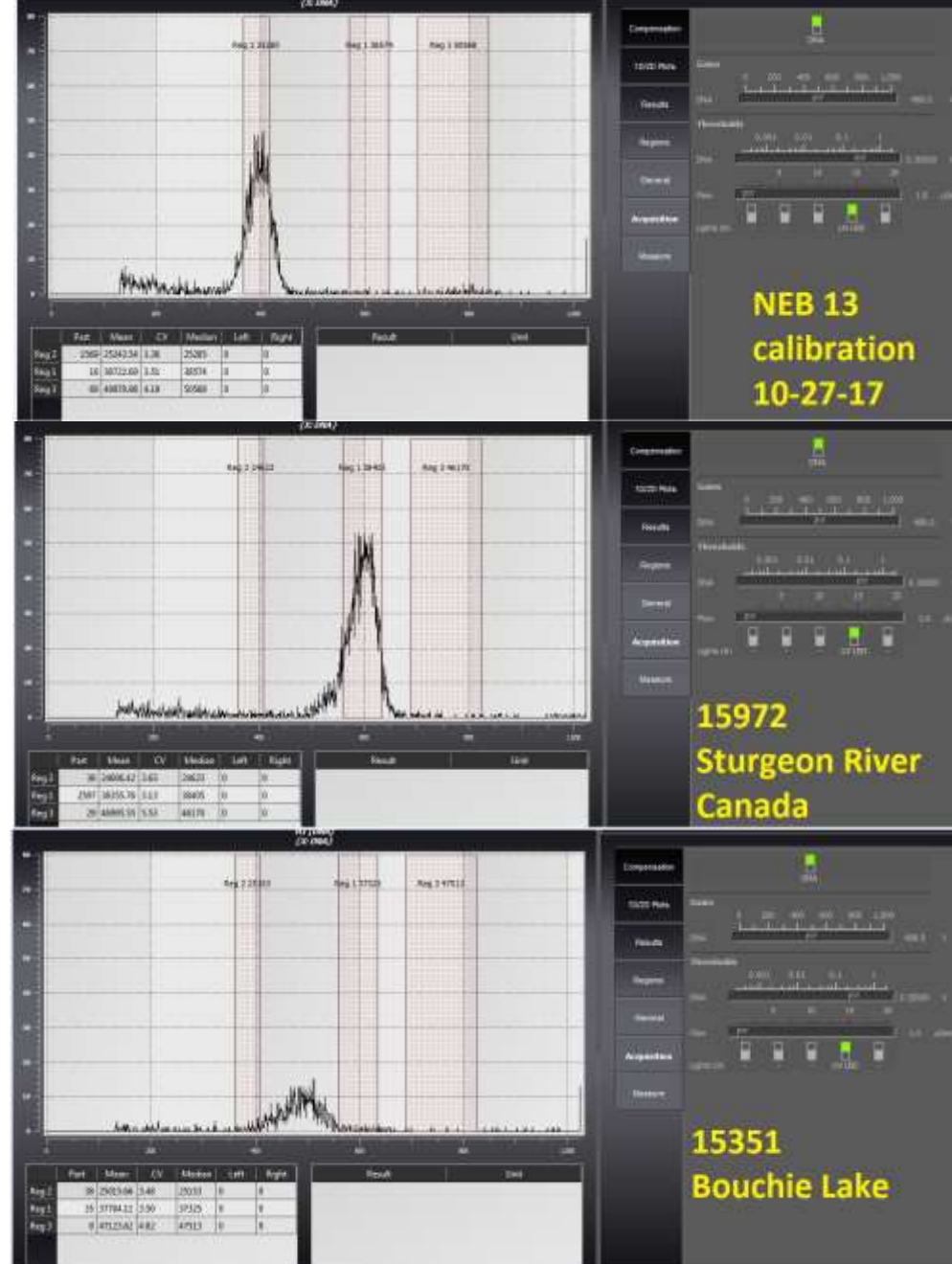
Can also tell by stomata



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

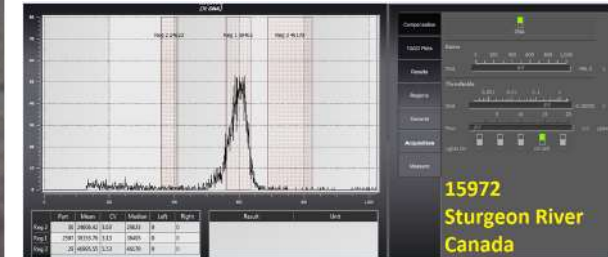
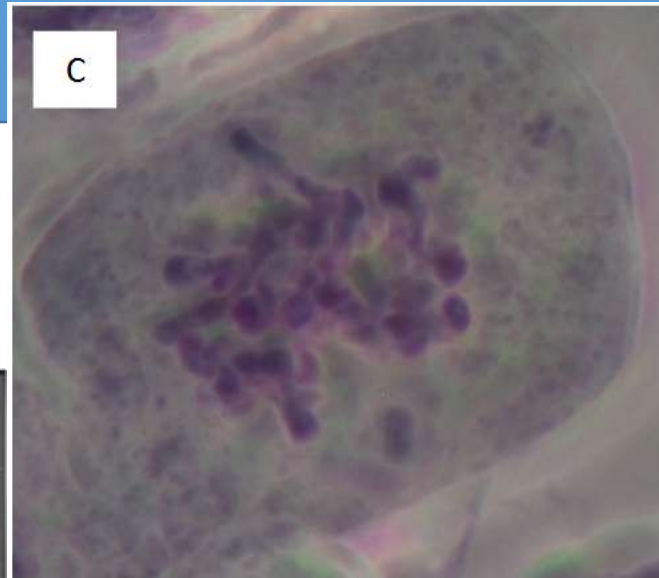









Figure 1. Composite cell images from *Butomus umbellatus* with A-C) 2n=26 chromosomes, D) 2n=39 chromosomes.

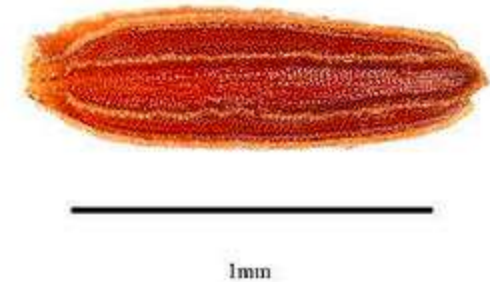


-  1 3x (2x at Prosser Diversion dam) [Jennifer Parsons]
-  2 3x
-  3 2x
-  4 2x
-  5 2x
-  6 ?
-  9 ?

yellow star=favorite restaurants

Who cares about ploidy!?

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



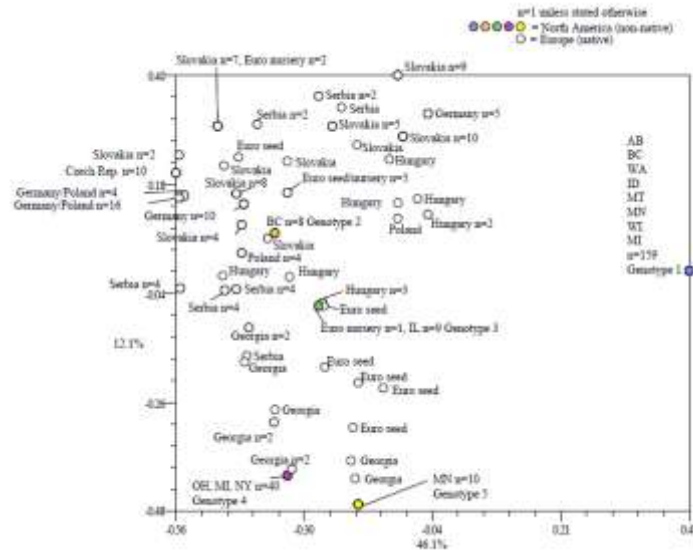
Results so far

- Europe has many genotypes. NA has few.
- Not all Euro pops clonal. Most NA pops clonal. Why??
- We have both diploid and triploid genotype 1 (mostly triploid so far)
- Diploid and triploid of same genotype...do they act the same when it comes to control?
- We have not found a close match for our very common genotype 1
- CABI and USACE will have or do have all genotypes to use in testing of potential agents.



Biocontrol with foreign disease (CABI UK)

Will a disease attack all of our *Butomus*?

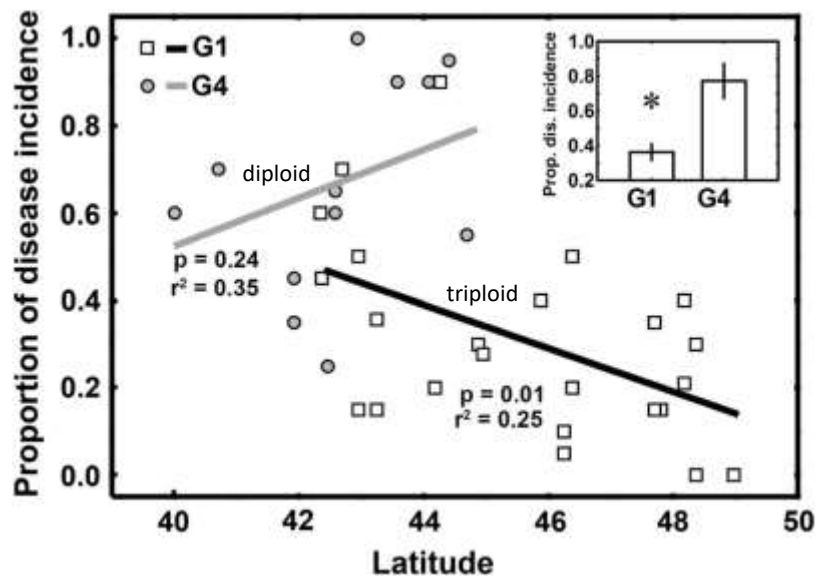


No close genetic match in Eurasia for our genotype 1...yet.

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



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.

Side note!

- Plant invasions are rarely homogenous entities.
- Evolutionary processes such as selection, drift, reproductive mode, and founding events shape the population structure and diversity of an invasion, and that information can help guide weed management.

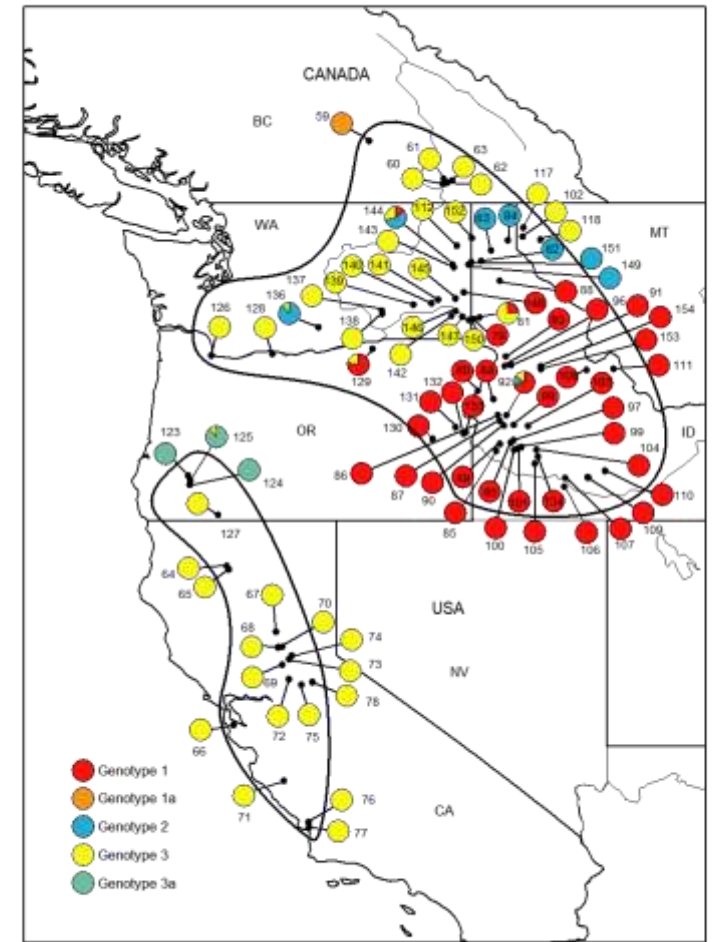


Fig 2a

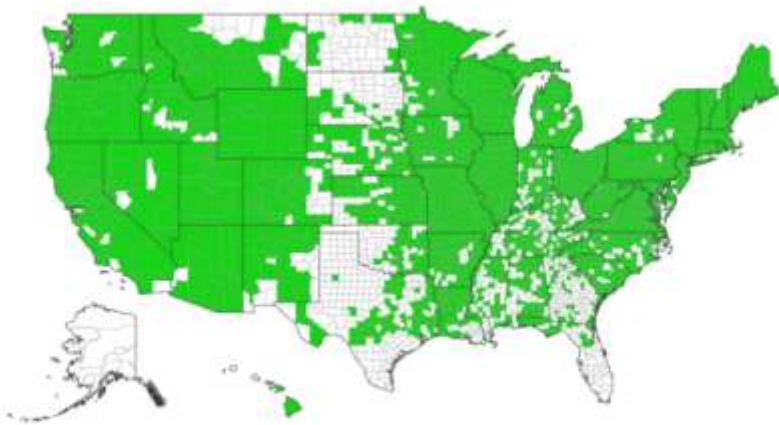
Rush skeletonweed

Verbascum thapsus; common mullein

John F. Gaskin USDA ARS
Stacy B. Endriss CSU
Christa Fettig CSU
Ruth Hufbauer CSU
Andrew Norton CSU
René F.H. Sforza USDA ARS EBCL



Verbascum thapsus
Common mullein



How do weeds reproduce? Self? Outcross? Other? (affects population structure and weed control method)



- Donnelly et al. (1998) demonstrated that this species is fully self-compatible without evidence of inbreeding depression, and that it has a mixed mating system with reproductive strategy (selfing or outcrossing) depending on variables that affect pollinator availability (e.g. plant height or population size).
- Donnelly SE, Lortie CJ, Aarssen LW. Pollination in *Verbascum thapsus* (Scrophulariaceae): the advantage of being tall. *American Journal of Botany*. 1998 Nov;85(11):1618-25.

Why be showy if you don't need or use sex??

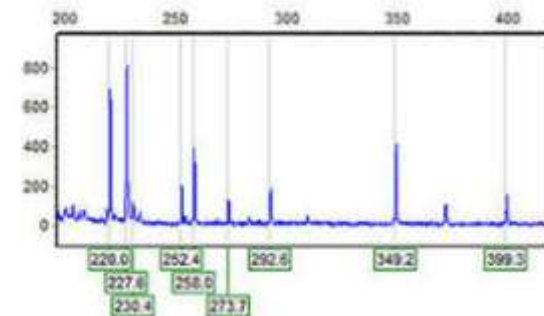
How is it reproducing in the invasion? AFLP study



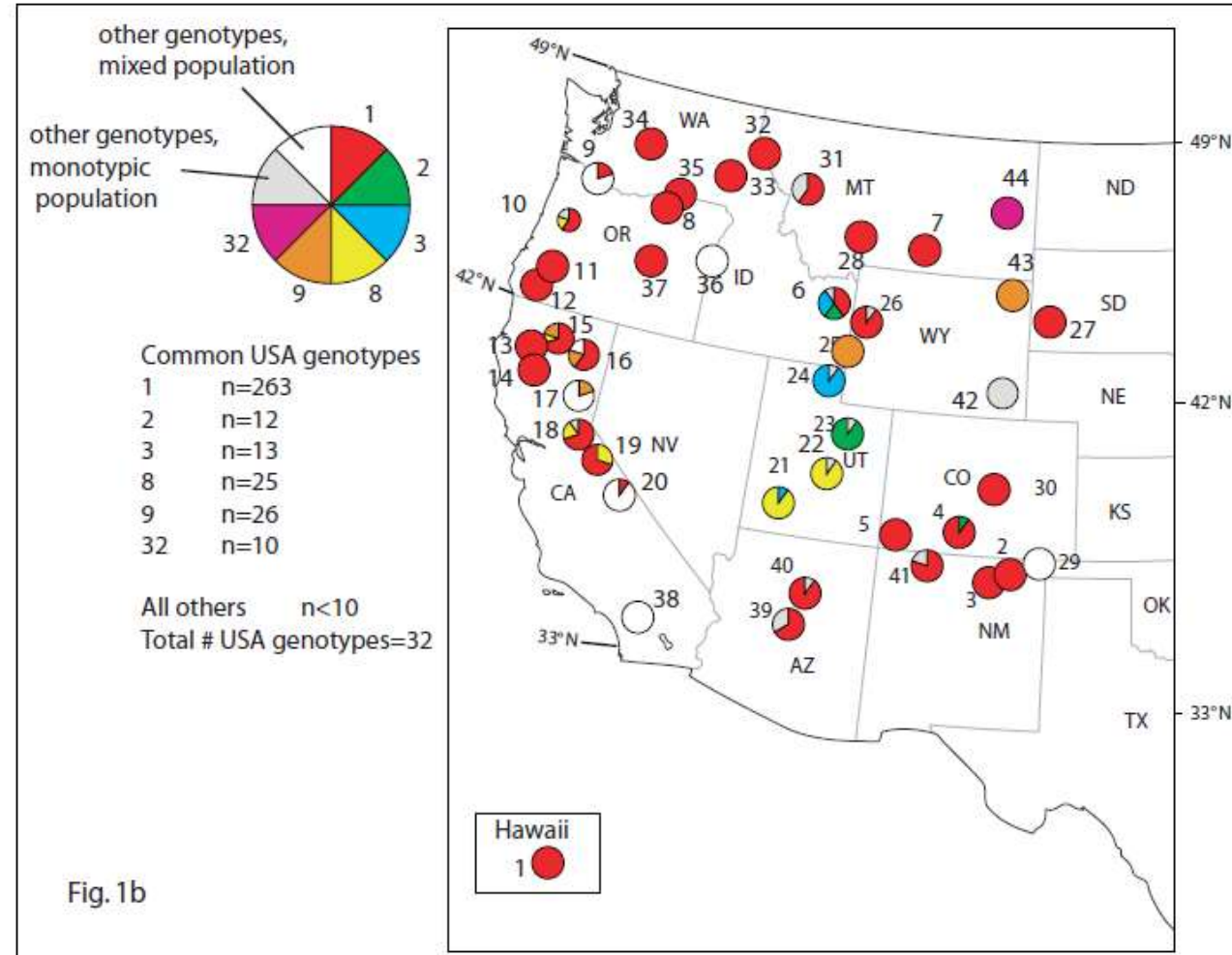
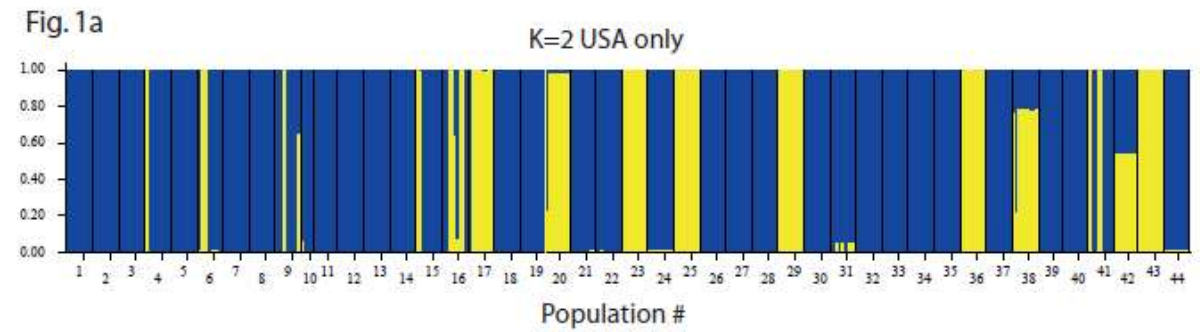
Table 2. Comparison of USA and Eurasian genetic data for *Verbascum thapsus*.

Location	N	G	G/N	PLP	Hj	Fst
USA	431	32	0.0700	12.5%	0.05	0.30
Eurasia	479	394	0.8200	61.5%	0.20	0.72

We found much higher diversity in the native Eurasian range, with 394 genotypes found in 479 plants.



- 50% of invasive populations were monotypic, and we found only 32 genotypes in 431 plants.
- Same genotype can be found across range of invasion.
- Why aren't there more genotypes??
 - founding effect
 - selfing



Origins

- We found highest genetic similarity between the USA invasion and plants from Belgium, Germany and France.

Fig. 2a

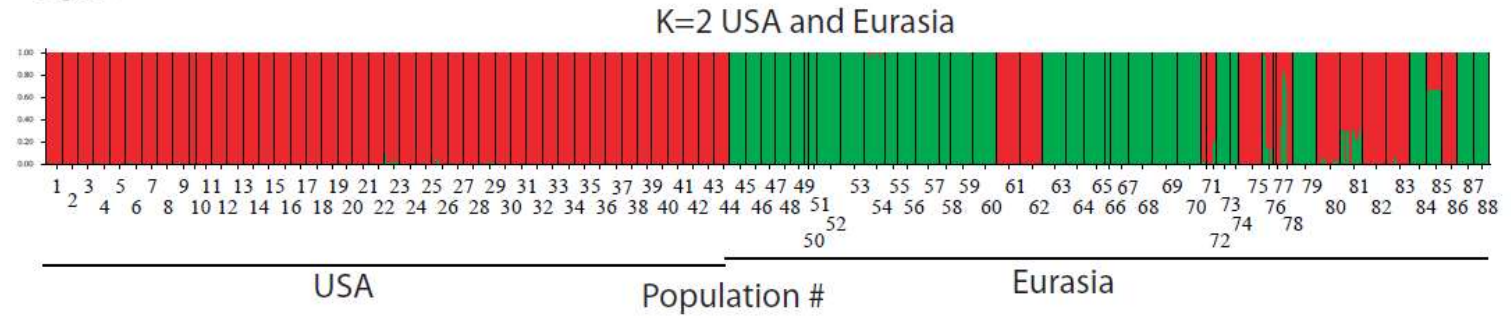
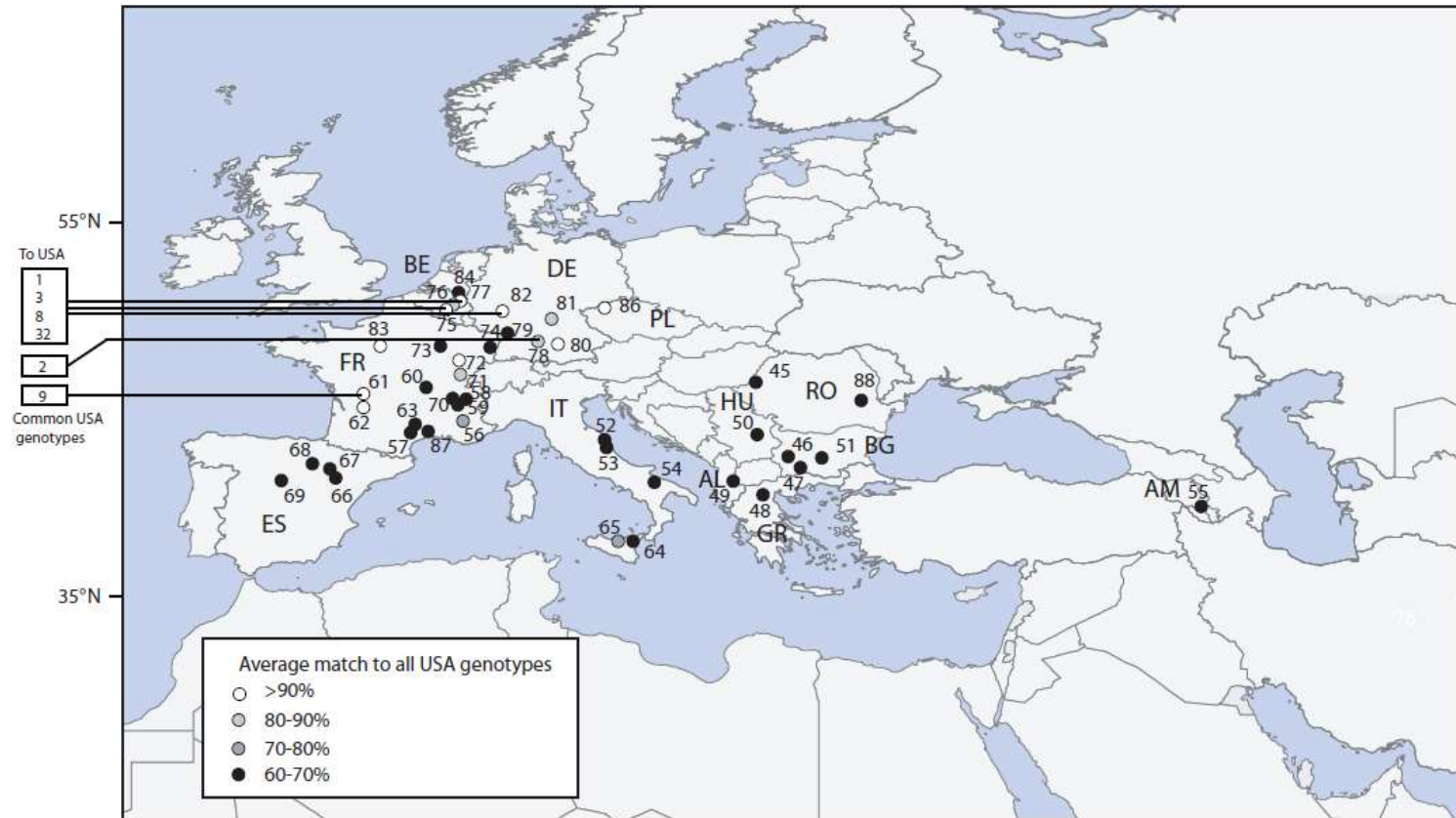


Fig. 2b



Verbascum

There is an unsupported suggestion that common mullein has some herbicide resistance or tolerance (Burton 1964 in Gross and Werner 1978).

- With this DNA information we can assist common mullein management by
 - identifying and supplying the common, or all, invasive genotypes for herbicide resistance research
 - identification of specific origins for searching for co-evolved biological control agents in the native range.



What thistle is this?

JOHN F. GASKIN

USDA

ERIC COOMBS

OR DEPT. AG

MARK PORTER

OR DEPT. AG

DEAN G. KELCH

UC Berkeley, California Department of Food and Agriculture

DAVID J. KEIL

California Polytechnic State University

ALFONSO SUSANNA

Botanic Institute of Barcelona



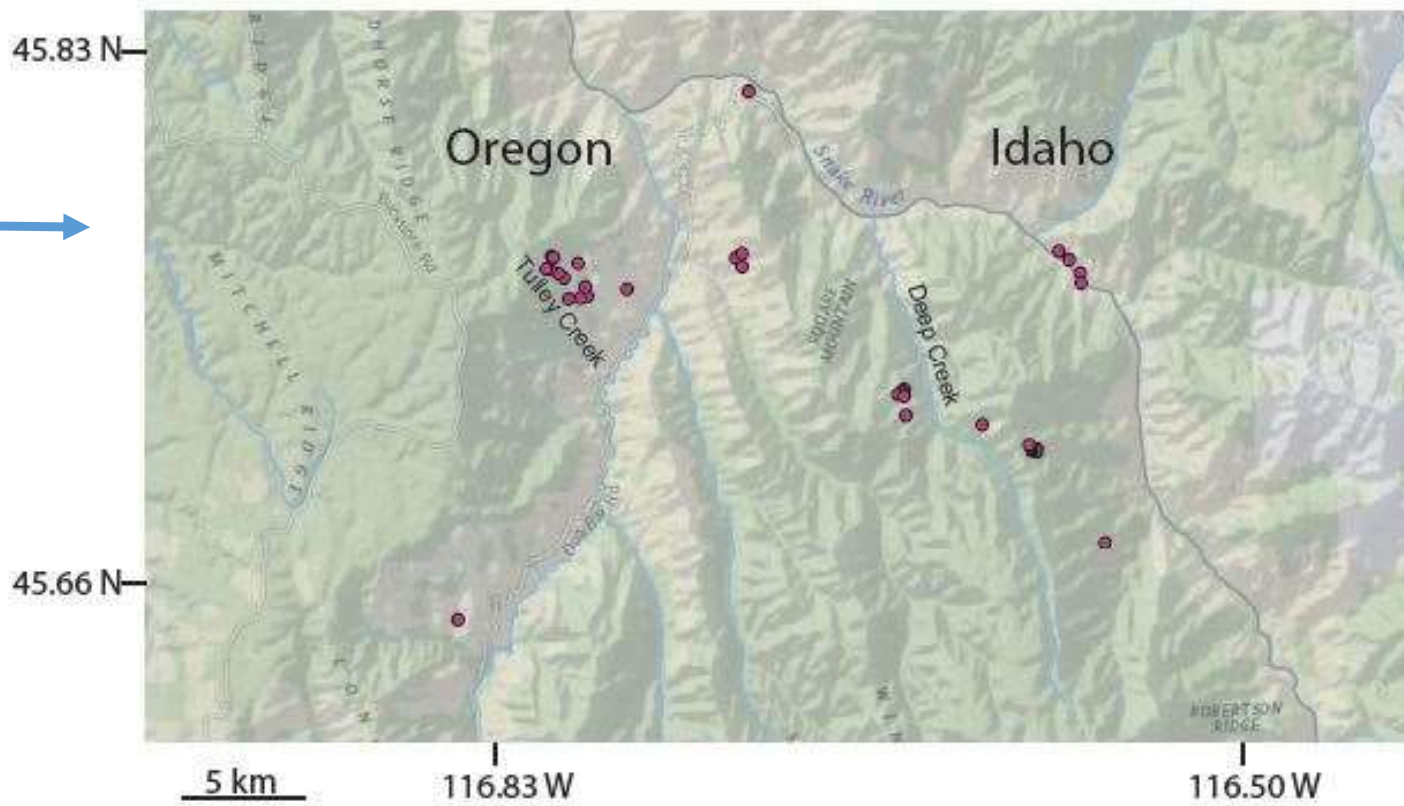
Refresher course: *Carduus* in N. America

All non-native, state noxious

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



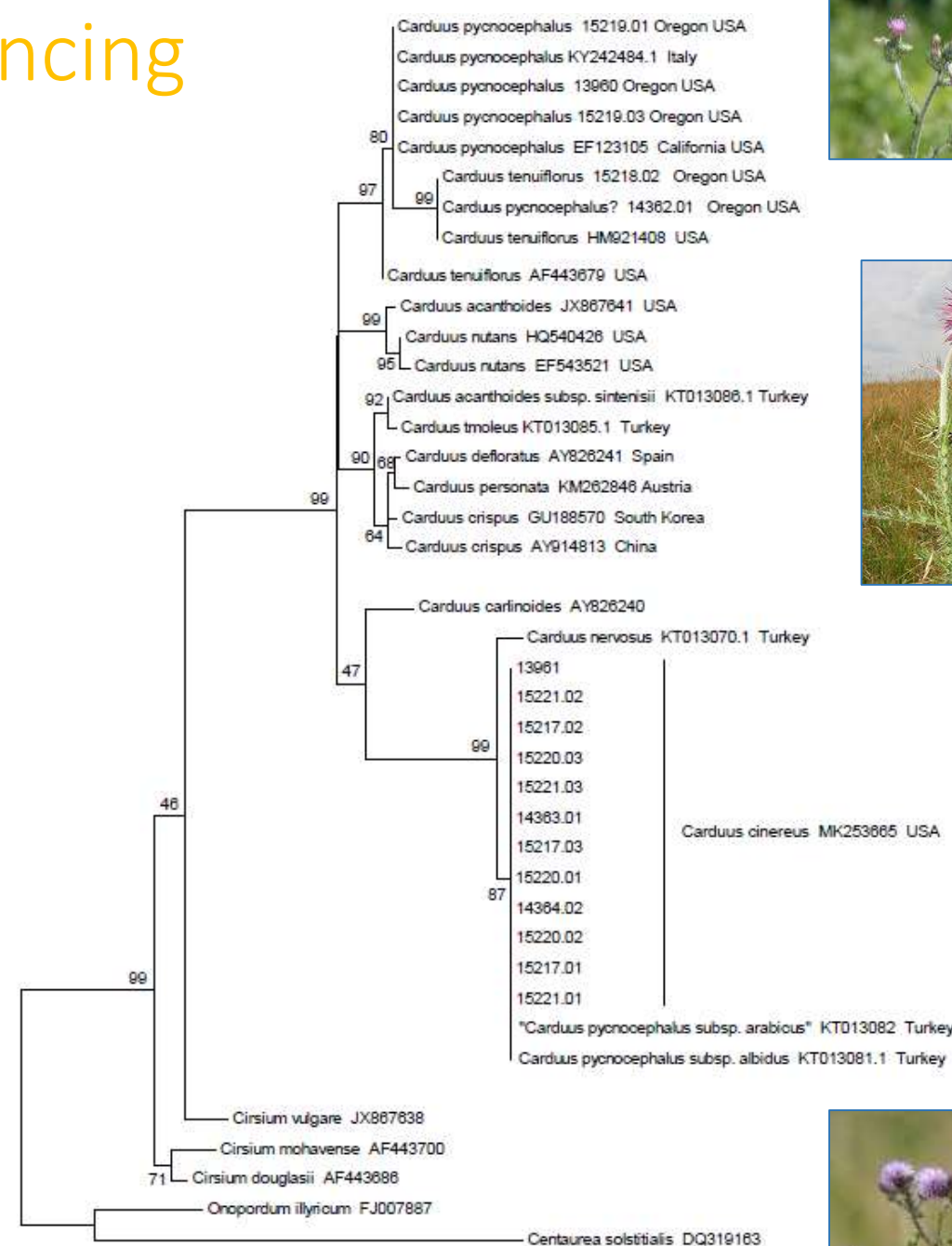




- *Carduus nutans*
- *Carduus acanthoides*
- *Carduus crispus*
- *Carduus tenuiflorus*
- *Carduus pycnocephalus?*
- *Carduus? sp.???*

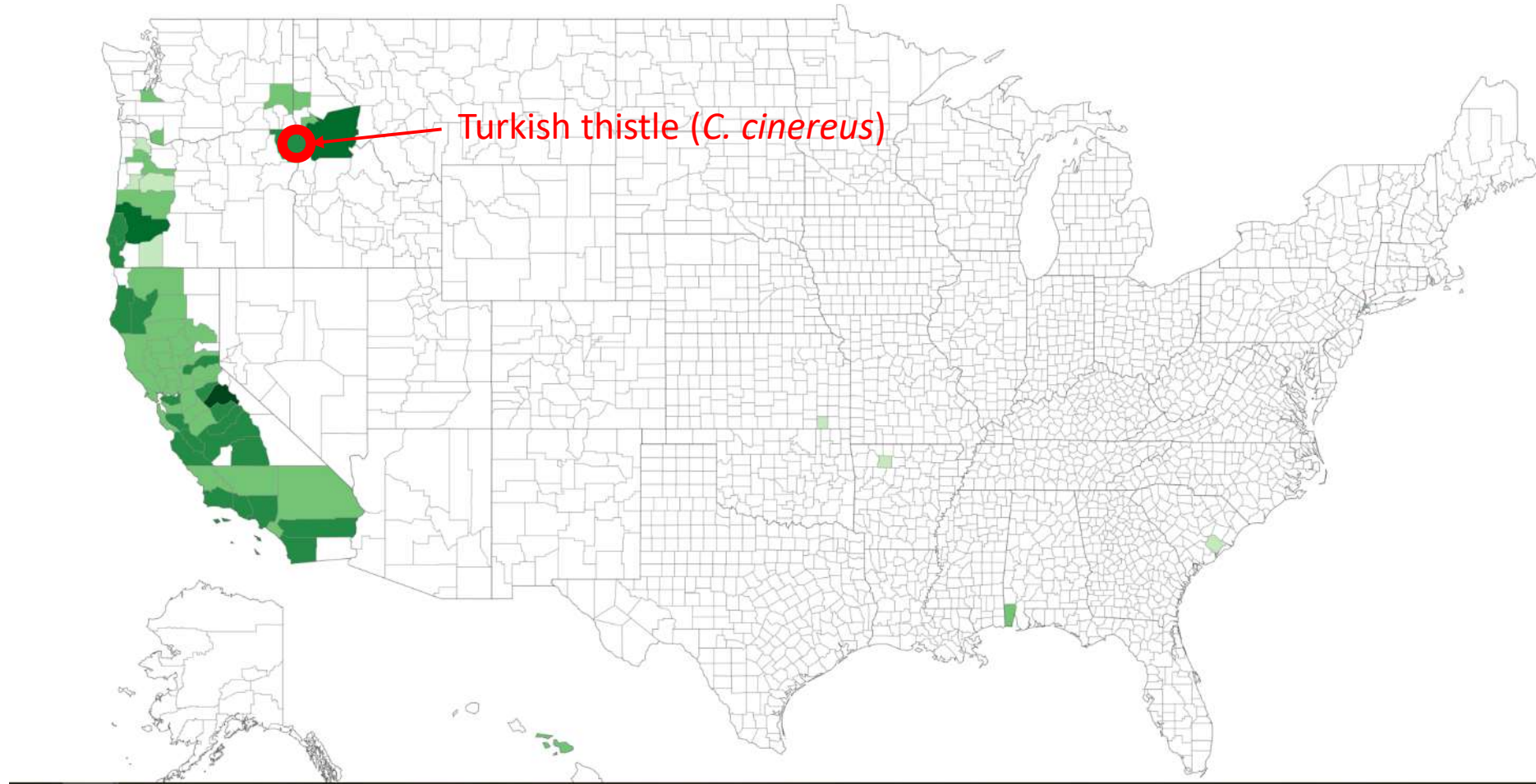


DNA sequencing



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

Italian thistle (*C. pycnocephalus*)



MODIFIED KEY TO *CARDUUS* SPECIES IN NORTH AMERICA

GASKIN ET AL. MADRONO. IN PRESS. *CARDUUS CINEREUS*; A NEW SPECIES TO THE AMERICAS

1. Phyllaries 2–7 mm wide, usually wider than the appressed bases; peduncles often elongate, distally wingless; heads often nodding, usually borne singly or in leafy corymbiform arrays; involucre 20–70 mm diam *Carduus nutans*

1' Phyllaries 0.5–2.0 mm wide, usually narrower than the appressed bases; peduncles short, if present, usually winged throughout, less often sparsely winged or unwinged; heads erect, 1–many, often clustered or loosely clustered at branch tips or upper axils; involucre 7–30 mm diam

2. Involucres spheric or hemispheric

3. Corollas 13–20 mm long; heads 18–25 mm long; involucre 14–20 mm long; abaxial leaf faces glabrate except for long, curled, septate hairs along veins

Carduus acanthoides

3' Corollas 11–16 mm long; heads 15–18 mm long; involucre 12–17 mm long; abaxial leaf faces sparsely to densely tomentose with fine, non-septate hairs and often with curled, septate hairs along veins as well

Carduus crispus

2' Involucres cylindric or narrowly ellipsoid

4. Heads 5–20 at ends of branches; phyllaries glabrous or sparingly tomentose, distally ciliolate or glabrous *Carduus tenuiflorus*

4' Heads 1–5 at ends of branches; phyllaries ± persistently tomentose, distally scabrous on margins and faces

5. Heads usually pedunculate and loosely clustered; phyllaries scarious-margined

Carduus cinereus

5'. Heads usually sessile and tightly clustered; phyllaries not scarious-margined

Carduus pycnocephalus

Using genetics to inform invasive weed management

What are your needs or questions?



John Gaskin
Botanist
Sidney, MT

