# Genetic identity and phylogenetic relationships of invasive brooms in California

## Annabelle Kleist and Marie Jasieniuk, UC Davis Department of Plant Sciences

## Background

The horticultural industry introduces and distributes large numbers of non-native plants into new areas and is a pathway for plant invasions. Horticulturallyintroduced invaders are often difficult to identify due to hybridization among ornamental cultivars and naturalized populations. Intra- and inter-specific hybrid identification is critical for invasive species management. This is especially true for biological control programs because incorrect identification of hybrids can lead to ineffective biocontrol and/or searches for biocontrol agents outside of native ranges.

## Study System: Brooms

#### Taxonomy:

- Legumes in the tribe Genisteae (~450 species)
- Invasive brooms belong to genera Cytisus, Genista, and Spartium

## French broom: Genista monspessulana

- Native Range: Mediterranean Region
- Introduction to CA: initial introduction ~1850 in San Francisco
- Current distribution: 23 counties in CA

#### Sweet broom: Genista racemosa

- Ornamental plant currently available in CA
- Putative close relative of French broom • Morphological similarities
- Sold under a variety of scientific names
  - Including: Cytisus spachianus, Genista stenopetala ssp. spachiana, and Genista canariensis

## Predictions

- . The current taxonomic status of ornamental and invasive brooms in CA is confused.
  - Multiple species are being sold as "sweet broom"
  - Invasive French broom populations are comprised of multiple closely related species
- 2. Hybridization occurs between invasive French broom and ornamental sweet broom.
- 3. Sweet broom individuals are present in invasive populations, particularly in urban invasions





## Objectives

**Overall Goal:** Elucidate the identity and origins of invasive brooms in California

## **Specific Objectives:**

- 1. Identify which species are being sold by the horticultural industry and which species comprise invasive populations
- 2. Determine whether hybridization between ornamental cultivars, species, and populations in natural areas has occurred



vtisus

(Scotch broom)

(French broom)

(Spanish broom)

-enista

partium

## Materials and Methods

#### **Overview:**

Phylogenetic analyses of nuclear and chloroplast DNA sequence data were used to determine relationships of invasive and ornamental brooms to each other and to brooms from the native range.

#### Sampling:



- 23 invasive brooms from throughout CA
- 6 landscape plantings
- 6 plants from horticultural industry
- Samples from botanical gardens and arboreta worldwide

Fig 1. French broom and sweet broom collection locations

## **Molecular Tools:**

PCR amplification and DNA sequencing:

- nuclear ETS region
- chloroplast *tRNA-leu* region

## **Phylogenetic Analyses:**

Maximum parsimony tree constructed using PAUP\*









## French broom clade:

Contains majority of invasive French broom from California and G. monspessulana and G. canariensis

#### Sweet broom clade:

Contains all sweet broom, 3 urban invasive broom samples, G. stenopetala, and G. maderensis

#### Scotch broom:

Invasive "Scotch broom" with some similarities to French broom clusters with French broom for *tRNA-leu* and with Scotch broom for ETS





## **Conclusions:**

- 1. Most of the invasive French broom in CA is either G. monspessulana, G. canariensis, or a hybrid between the two
- 2. Ornamental sweet broom most likely contributes directly or via hybridization to invasive broom populations
- 3. Hybridization between French and Scotch broom can occur in natural populations

