**Intro**

~~I linked demographic and species distribution models for the invasive plant~~ *~~Frangula alnus~~* ~~throughout its North American range, and compared simulation results to observed patterns of spatial spread to determine the demographic properties and processes most likely responsible for this species’ success~~.

**Methods**

Across all six sites, 815 individual plants were measured and tagged with a unique identifier, 182 first tagged in 2010, 567 in 2011, and 65 in 2012.

At four of the six sites, I counted the number of fruit on tagged plants during the month of July of 2012 to estimate differences in timing of fruit set within a season and to estimate annual maximum fruit per plant values.

Another advantage to using the IPM framework is that the regression analyses allowed me to estimate parameter uncertainty bounds to use in a GSA (described below in **Exploring parameter uncertainty space via global sensitivity analysis**).

Other records were reported in the Early Detection and Distribution Mapping System database (EDDMapS n.d.), iMap Invasives – New York (© 2013 The Nature Conservancy) database, and the Great Lakes Indian Fish and Wildlife Commission invasive species survey data (Garske and Falck 2007, Garske 2010).

Approximately 75% of occurrences were collected after the year 2000.

The climate/ weather variables matched a set of widely applied bioclimatic variables (Hijmans et al. 2005), and were calculated as 10-year and 50-year aggregated climate variables. All variables were provided as raster layers (WGS84 projection with a decimal degree coordinate reference system) at a spatial resolution of 5 x 5 arc minutes. The variables were continuous measurements with the exceptions of potential vegetation, pH top, and pH sub. While each predictor variable was chosen based on hypothesized importance in plant performance, some variables were not used in the final SDMs because they were highly correlated with other variables. Eliminations were based both on correlations with other variables and results from preliminary SDMs (see appendix for details).

**Discussion (maybe)**

(THIS PASSAGE COMES FROM MY INTRO, BUT SEEMS MORE LIKE DISCUSSION TO ME NOW?) This is similar to patterns found by Larkin (2011) for this species regionally in Michigan and Wisconsin (i.e., a lag of approximately 30 – 40 years).