Conserving Genetic Diversity ex situ: Updates on Practical Advice

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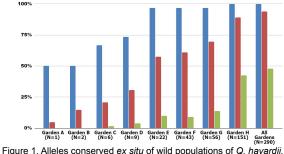
Overview: Conserving rare or threatened plants with limited conservation resources requires data-driven, effective, and efficient strategies. At the Morton Arboretum, we assess genetic diversity safeguarded in *ex situ* collections (left column), test sampling approaches using simulations (middle), and scale up our work across species (right). Our results indicate how multiple avenues of work combine together to contribute *practical advice for seed collectors*!

Assessing the Outcome of an Intensive Collection

Goal: Quantify genetic diversity in a large ex situ collection of a desert oak *Q. havardii* (photo at right, showing its small stature) across multiple botanic gardens (a meta-collection).



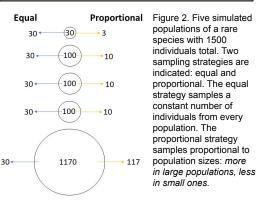
■ Very Common (>10%) ■ Low Frequency (5-1%) ■ Rare (<1%)



Key finding: We conserved *Q. havardii* by sampling >1000 seed, from >20 locations via USFS/APGA funding in 2016. We quantified wild genetic diversity conserved in gardens with different numbers of seedlings. 25 plants conserve common alleles (Fig 1, blue) but >100 are needed for conserving rare but valuable genetic variation (red).

Using Simulations to Test Sampling

Goal: Determine practical sampling guidelines for rare species with unequal sized populations.



Key finding: Sampling proportional to population size captures significantly more genetic diversity than sampling equally across all populations when population sizes vary. Seed samplers should allocate more effort in larger populations to efficiently collect genetic diversity, when feasible.

Suggested reading: BGJournal issue 17.2, pg 23, Hoban, Cavender and Griffith **See also:** BGCI publication "Towards the Metacollection" Griffith et al 2019 (bgci.org/resources/)

Looking Forward - Gene Conservation in 10 Oaks

Goal: Build on recent work showing that we can quantify genetic diversity conserved *ex situ*. Most botanic garden collections do not yet capture enough genetic diversity and we need better seed sourcing guidance for each genus.

Table 1. Allele capture *ex situ* from wild populations of three species of oaks (from Hoban et al. (2020), Proceedings B).

Species	N in situ	N ex situ	Common	Low Frequency	Rare
Quercus boyntonii	244	77	100	66	29
Quercus georgiana	223	36	92	66	39
Quercus oglethorpensis	187	145	100	97	67



Ongoing work, Q. acerifolia Image: Morton Arboretum

Key Plans: We will test seed sampling guidelines that can apply to all threatened oaks. We are examining 7 additional IUCN Red List threatened oaks from the US (pacifica, hinckleyi, tomentella, acerifolia, ajoensis, graciliformis, havardi). We will see how these oaks are safeguarded *ex situ* and design better guidelines for this important genus.