



# Ecoregions and Endemism to define Gene Conservation Guidelines for Longleaf Pine Ground-layer Restoration

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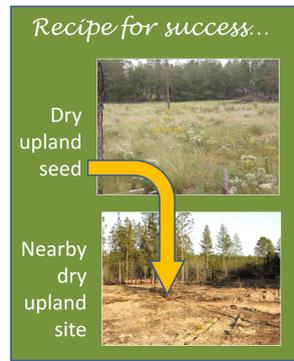


The lack of ecologically suitable plant materials for restoration represents a formidable barrier to achieving range-wide restoration goals for longleaf pine. There is a clear need for developing sources for plant materials that will survive and thrive in the restored setting. In addition to selecting the “right” species for a project, it is also important to consider the source of materials. Botanists, foresters, and geneticists recognize variations within a species. Variation can be associated with the source location and related to adaptations for success in different conditions. A key to selecting suitable plant material lies in understanding the species’ genetics through common garden studies. However, managers often must collect seed or purchase whatever seed is available from commercial sources, and deploy seed of species for which there are no research based guideline.

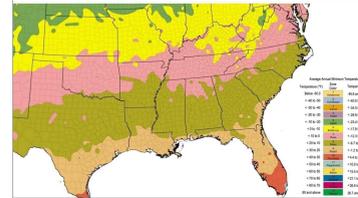
What do you do if no seed transfer guidelines have been established by genetics research?

## THE SAFE SOLUTION

Select seed sources from the closest possible location and collect from a site that physically matches the target site. This may minimize risks of getting individuals maladapted to the restoration site, but provides no guidance for seed sources for wider distribution.



USDA Plant Hardiness Zone Map for the Southeastern United States



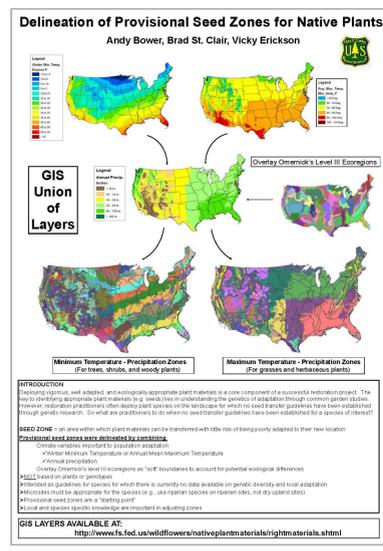
Hardiness Zones, which are based on this principle. Seed transfer recommendations for loblolly pine have been based on a climate model (Schmidting 2003) and are available for all southern pines.

## ANOTHER CLIMATE-BASED MODEL

Bower, St. Clair, and Erickson used climate data coupled with EPA Level III ecoregions to develop 2 models for native plants. The model for trees, shrubs, and woody vegetation was based on minimum winter temperature and annual precipitation. The herbaceous species model was based on average maximum temperature and annual precipitation, climatic factors that affect herbaceous species viability. They suggested that Level III ecoregions could be used to guide transfer decisions within climate defined zones. Note the large zone for southeastern grasses and forbs.

## CLIMATE BASED MODELS

Plant distributions are generally limited by climate, so it is not surprising that seed transfer guidelines based on climate patterns have proved useful, especially for trees. Gardeners recognize the USDA Plant



## ECOREGIONS FOR GUIDANCE

Level III ecoregions define areas with repeating patterns of vegetation and landforms. They are defined by geology, soils, physiography, and to some degree major vegetation types. Ecoregions DO NOT consider within-species genetic variability. However, some studies have shown genetic patterns correlate with ecoregions in the Pacific Northwest.

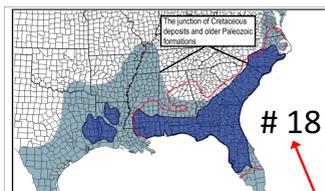
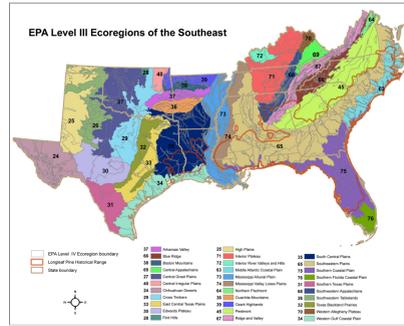


Figure 5. Phylogenetic pattern #18. Dark blue stippled area: core region of longleaf pine (Pinus palustris). Red dotted line: area of discontinuous or distinct occurrences of longleaf pine. After Ware, Frost, and Dow (1993). Light blue: Atlantic Coastal Plain Gulf Coast Plain, including the entire Mississippi Embayment. Adapted from Somers & Weakley (2001).

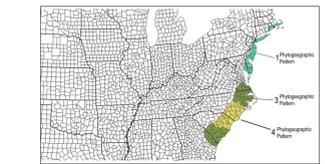


Figure 2. Phylogenetic patterns #1, #3, and #4. #1 (light blue): southeastern Mississippi to southern New Jersey and southern Delaware. #3 (green): southeastern Virginia to southeastern Georgia. #4 (dark green): southeastern South Carolina. Adapted from Somers & Weakley (2001).

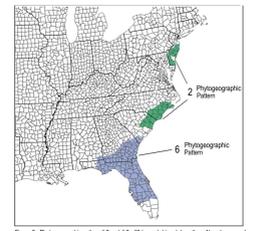


Figure 3. Phylogenetic pattern #2 and #6. #2 (green): broad southern New Jersey and Delaware to North and South Carolina. #6 (blue): southern South Carolina to peninsula Florida westward to the eastern panhandle of Florida and southeastern Alabama. Adapted from Somers & Weakley (2001).

## ENDEMISM IN THE SOUTHEASTERN COASTAL PLAIN

About 27% of the about 6100 taxa native to the Coastal Plain, are endemic to the region. Somers and Weakley (2001, 2006) examined distributions of >1000 coastal plain endemic species (> 90% of known locations in the coastal plain) and described common distribution patterns (with ≥ 8 taxa) as centers of endemism.

Pattern #18 nearly coincides with the range of longleaf pine in the coastal plains. It includes at least 440 taxa (not counting regional subsets), 15 endemic genera. Within this core area, subsets were recognized (see Figures 2-5; not shown is a West Gulf Coast center). Although some of the endemics are considered rare, many are distributed broadly within pattern #18.

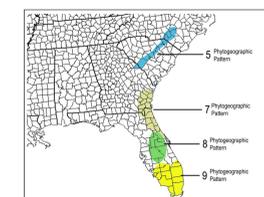


Figure 4. Phylogenetic patterns #5, #7, #8, and #9. #5 (blue): Sabine and sandhills of the California and Colorado. #7 (blue): southern Georgia to northeastern Florida. #8 (green): Lake Wales Ridge. #9 (yellow): subtropical Florida. Adapted from Somers & Weakley (2001).

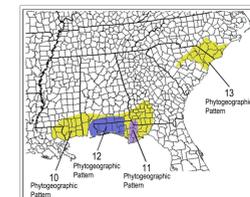


Figure 5. Phylogenetic patterns #10, #11, #12, and #13. #10 (yellow): East Gulf Coastal Plain. #11 (orange): central panhandle of Florida. #12 (dark blue): western panhandle of Florida. #13 (yellow): wide spread on East Gulf Coastal Plain-disrupt to the Carolines. Adapted from Somers & Weakley (2001).

## ENDEMISM--INDICATOR OF GENETIC STRUCTURE?

Endemic taxa arise through speciation via the development of altered gene combinations leading to isolation and further genetic change. Speciation can be associated with adaptations to local conditions, result from long-distance dispersal events, follow from isolation during periods of major climate changes. Processes resulting in speciation, may also lead to intraspecific genetic shifts that are not sufficient for reproductive isolation, but nevertheless lead to differentiation among populations. Genetic structure within a widespread species may lead to differences in traits associated with successful establishment and growth when introduced into a restoration setting. Intraspecific variation may be continuous & gradual (clinal) or local & abrupt, associated with habitat variability (ecotypic variation). Both kinds of variation may affect planting success.

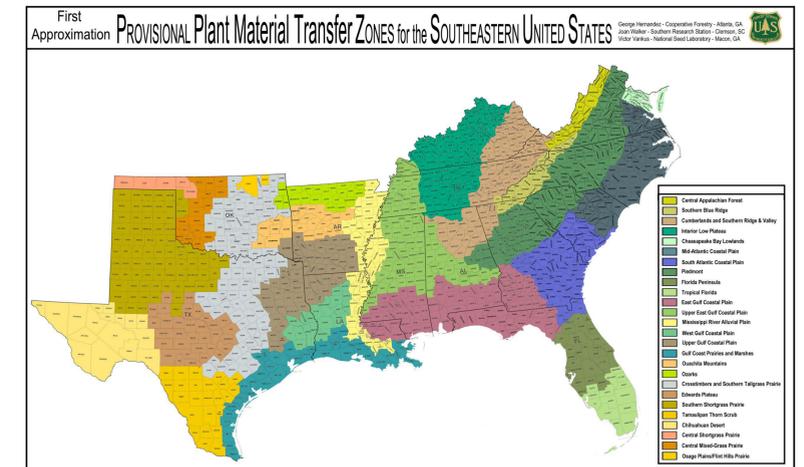
**Hypothesis:** Centers of endemism are indicators for potentially significant genetic structure within widespread species  
**Proposal:** Incorporate the distribution of endemics in the delineation of provisional plant material transfer zones. This approach may conserve genetic diversity within widespread species, and help ensure the establishment of new populations.

## FIRST APPROXIMATION



We examined available ecoregion models, USFS, EPA, NRCS, and The Nature Conservancy models. The TNC model was designed to (1) reflect the distribution of conservation targets at large scales and (2) to be implementable for conservation planning. TNC developers modified EPA

Level III Ecoregions and to meet their dual objectives. Modifications accommodated recognized phylogeographic patterns, e.g. in the decision to split South Carolina. The TNC foci of **conservation and implementation** paralleled the needs for delimiting plant material transfer guidelines, specifically zones within which plant materials (seed or others) might be conservatively moved. We modified the TNC ecoregion model to follow county lines, in order to support implementation of county administered land owner support programs.



## PROVISIONAL ZONES WITH CAVEATS

- They are not species specific. Individual species must be evaluated to determine suitability.
- They are large, and therefore are applicable for species that are naturally distributed across several zones.
- The zones are ecologically heterogeneous, with a range of habit conditions embedded. Within a zone, plant materials should be placed in habitat conditions that match their origins. For example, sites from well-drained sands will be most suitable for well-drained sandy sites within the same zone.
- The zones are provisional. Zones are expected to change with research. And as noted, transfer rules for individual species are likely to be developed for important restoration species.

