

Empirical seed transfer zones require conventions for data sharing to increase their utilization by practitioners

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Abstract

Empirical seed transfer zones (eSTZs) are being developed more frequently to help guide both the agricultural development of native plant materials and the selection of these materials for restoration projects. Despite the widespread use of eSTZs, standards for distributing these data are lacking, leading to inconsistent data products. To maximize the utilization of eSTZs, we propose standards to guide their distribution in the United States, thereby increasing the focus on seed collection efforts and fostering the utilization and accessibility of the most appropriate commercially available seed sources. Further, we propose that sharing metrics of model uncertainty for these data, which can help practitioners identify the best alternatives for a seed transfer zone, should become common practice. Finally, we briefly introduce the R package eSTZwritR ('easy rider'), which implements our core suggestions for data dissemination.

IMPLICATIONS FOR PRACTICE:

- Developing a restoration plan in a short time period, as required after a natural disturbance, can be difficult. To decrease the chances of simple mistakes being introduced into plans, we developed standards to increase the consistency between eSTZ data products, making their usage in GIS software more consistent.
- We implement these suggestions in an R package 'eSTZwritR' which should facilitate adherence to the guidelines for scientists developing eSTZ products, allowing for a rapid uptake of these conventions.

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- We also suggest incorporating estimates of uncertainty for spatial eSTZ data products so that practitioners have sufficient support for selecting materials from non-target seed zones, as is often required.

INTRODUCTION

[Figure 1 about here.]

Empirical seed transfer zones (eSTZs) are gaining popularity among restoration practitioners as tools to help identify the most appropriate seed source for a species at a restoration site (McKay et al. 2005). eSTZs are becoming more widely used for two primary reasons: 1) they are based on empirical data, such as phenotypes in a common garden, population genetics, or the correlation between occurrences of the species and environmental variables; and 2) they are generally fewer zones than provisional seed transfer zones, thereby reducing the number of lineages requiring cultivation in agricultural settings. Although popular, the development of eSTZs for a species is a costly and time-consuming process, most often involving common gardens or genetic studies, with many populations from across the species range incorporated as samples (Kramer et al. 2015).

In western North America, the majority of eSTZs have been developed by just a couple of lab groups, whereas the remainder have been developed by other assorted groups. While practices for developing eSTZs are becoming more defined, to our knowledge, no standards exist for *sharing* eSTZ results (Supplemental 1). Although eSTZs are produced by a relatively small pool of lab groups and individuals, inconsistencies vary across the spatial data products used to report eSTZs.

The success of a restoration project relies on the timely application of techniques suitable for the site at hand. The implementation of relevant techniques requires not only intrapersonal communication between a practitioner and themselves in time, for example avid note-taking, but also interpersonal communication between practitioners. Hence, the dissemination of ideas during and after a restoration project is the best opportunity to improve restoration outcomes (Figure 1). However, ideas have varying levels of complexity that may hinder their transmission.

For example, seeding rates may be verbally communicated, while seed mixes are likely to require written documentation, spatial data require both written and geographic data (e.g., coordinates and relations between them) in the form of spatial data products (e.g., rasters and shapefiles) to accurately convey their meaning. Given the relative complexity of communicating precise spatial information, standards should exist to ensure not only its accuracy and precision, but also its ease of interpretation and use.

Using 23 sets of eSTZs produced for 22 taxa, we showed that most of the spatial data developed and disseminated to share the results of an eSTZ are inconsistent (Supplemental 1). We have already observed significant hindrances to the uptake of these data at the practitioner level and searched for consensus within these data. Subsequently, using any consensus (wisdom of the masses) from these data combined with standard conventions of data sharing, we present a set of guiding standards for researchers in the United States to make the results more consistent.

Current Condition

[Figure 2 about here.]

We conducted a review of all eSTZs on the Western Wildland Environmental Threat Assessment Center (WWETAC) website as of May 1, 2024 (<https://research.fs.usda.gov/pnw/products/dataandtools/datasets/seed-zone-gis-data>). Each data product (file name structure, field naming conventions, and directory structure) was manually scored, and all analyses were performed in R version 4.2.1.

[Figure 3 about here.]

In Figures 2 through 4, we present inconsistencies that we believe or have observed to be the most likely to interfere with practitioners' workflows. We encountered considerable inconsistencies within file names (Figure 2), in directory structure and naming (Figure 3), and cartographic elements of the 20 maps available (Figure 4). While some consensus existed around the use of USDA NRCS-Plants codes for denoting the taxon contained in the file (Figure 2), the lack of file names mentioning what attribute about the taxon they contained (e.g. 'zones,' 'seed_zone,' 'sz'), and the lack of specified geographic extents can make determining the specifics of the file difficult unless it is explicitly opened in a Geographic Information System (GIS) software.

The naming of the fields (columns) within shapefiles likely presented the most problematic of all results (Figure 3), although many inconsistencies exist; here, we focus on three. Different uses of polygon geometry were implemented to represent the individual seed transfer zones, that is, sometimes all portions of a seed transfer zone, when at least some components are disconnected, were stored within the same object or row (a multipolygon). At other times, each discontinuous portion of the range was stored as its own polygon. For most infrequent Geographic Information System (GIS) users, we observed that multipolygons can be

confusing and require them to use several moderately advanced spatial techniques to interact with. Surprisingly, within each shapefile, the field denoting the seed zones was often ambiguously labelled or entirely lacking any indication (Figure 3). In several instances, it took several minutes to determine which field was the seed zone by toggling through and visualizing many fields, despite having already been interfaced with all of these products multiple times.

[Figure 4 about here.]

Recommendations

Some consensus exists among eSTZ developers for a range of attributes related to the distribution of data products. Combining these opinions with best practices for data sharing and experience as users of each of the existing empirical products results in the following recommendations.

Directory Structure

[Figure 5 about here.]

eSTZs should be distributed using a predictable directory structure that allows users to be immediately familiar with where to find the content (Figure 5).

We recommend that all directories (folders) have two main subdirectories (Figure 5), one containing the essential data products, preferably in both raster and vector Data Formats (*see ‘Data Formats’*).

The second directory contains information related to the product, including a formatted citation for data use, a map for quick reference, and any materials describing the development of the product, both as a paper and a text file of quick metadata attributes.

File Naming

[Figure 6 about here.]

The files within the directory should follow a naming convention that is easy for users to interpret and import to various software while also describing the essential attributes of the data product.

We recommend (Figure 6) that each file name has three main components in addition to the file extension.

The first component is the USDA PLANTS code, and the second is the method used to develop the STZ

- currently one of ‘g’, ‘cg’, ‘cm’ (for genetic, common garden, and climate matched, respectively), and the final is up to the two main regions which the product overlaps.

In the United States, we recommend the use of the 12 Department of Interior regions as they cover contiguous geographic expanses, are few enough to be easily remembered, and balance L2 Omernik ecoregions with easier to remember state lines. However, in other nations, the use of ecoregions may be more desirable.

Maps

Maps of the data product should be included in the ‘Information’ directory. Many questions about eSTZs can be answered quickly and simply by a practitioner consulting a map saved as a PDF with the essential cartographic components; fortunately most developers already supply these. We recommend that each map contains the following elements: north arrow, scale bar, state borders, geographically relevant cities, coordinate reference system information, sensible categorical color schemes for the seed zones (e.g. from ColorBrewer <https://colorbrewer2.org>), a legend, the taxons name as a title, and the maps theme (‘Seed Transfer Zones’) as a subtitle.

Data Formats

We recommend that the spatial data associated with an eSTZ be distributed using both popular spatial data models, vector and raster. For vector data, we advocate the continued usage of the shapefile format, whereas for raster data, we propose the use of geoTIFFs (‘tifs,’ the. tif extension). In our experience, tifs seem to be the most widely used raster data models in ecology for non-time series data, and are supported by virtually all GIS software.

Vector Data Field Attributes

[Figure 7 about here.]

The order of the fields (or columns) of the vector data should follow a predictable pattern (Figure 7), allowing humans to interact with the data in a graphical user interface (GUI) to quickly detect their field of interest.

We recommend that each shapefile have at least four fields in the following order and of the following data types. 1) The numeric integer (ID) is a unique number associated with each individual polygon in the file. 2) Seed Zone (numeric integer) is a unique identifier for each of the eSTZs delineated by the product

128 developers, which allows for quick filtering of the data based on a simple numeric value that is difficult to
129 misspecify. 3) SZName (character): a human-developed name for the zone that may refer to an axis of a
130 principal component analysis, for example, ‘LOW MEDIUM LOW’, or be defined by the product developers.
131 We propose that semi-informative names should be developed before data distribution to help practitioners
132 more easily convey important attributes without relying on numeric values, which may be more difficult to
133 remember because of their nondescriptive nature.
134 4) AreaAcres (numeric - integer) of each polygon.

135 In addition to these standard field naming and placement conventions, we recommend a series of standards
136 for the contents within these essential fields and how to format any additional fields relevant to the project
137 (see package website).

138 **Adapting our Proposals to other Regions**

139 We believe that our core set of proposals is readily adaptable to a variety of regions globally. However, some
140 modifications are required to ensure a seamless implementation. Notably, our suggestions for file names
141 contain three parts, two of which are specific to data sources limited in scope to the United States. The
142 first is the use of a set of abbreviated codes that reliably identifies a plant taxon (‘NRCS PLANTS code’).
143 The second is the use of administrative (‘DOI’) regions for denoting a general geographic area which the
144 specific product covers. An alternative solution that we considered for the US States is the use of ecoregions.
145 However, this was rejected because of the number of ecoregions (e.g., 10 conterminous US DOI regions to
146 20 level 2 Omernik ecoregions) and, most importantly, because many ecoregions are not contiguous, which
147 may lead to confusion. Possible alternatives should be evaluated based on the above points before they are
148 selected for use.

149 Additionally, we believe that many other countries may be able to immediately start distributing their vector
150 data as an OGC GeoPackage (‘gpkg’) files rather than shapefiles. The GeoPackage format is a geodatabase
151 (SQLite Database file) and an open-source standard that entails all data being stored in a single file (relative
152 to the several files a shapefile requires), contains projection data, and contains spatially indexed data. In fact,
153 the data associated with ‘eSTZwritR’ are all shared as GeoPackage files, and anecdotal evidence suggests
154 that its utilization by US Government agencies is increasing.

155 For the US, we suggest that seed zones have numeric representations following a generally decreasing pattern
156 in aridity because the success of restorations often follows this gradient (Shriver et al. 2018). However, in
157 other regions, other climatic conditions may be more important drivers of restoration outcomes, and they

should be used instead of this variable.

Estimating Uncertainty

[Figure 8 about here.]

We have witnessed confusion from seed collection crews, curators, and restoration practitioners alike over the appropriate classification for a new seed source and the selection of a seed source for a restoration. Generally, these hesitations relate to a source located on the border of multiple seed zones. We predict that with the increasing availability of fine-resolution spatial data that more accurately reflect local ecological heterogeneity-seed zones will become finer, increasing the perimeter-to-surface area ratio and the prevalence of this already common confusion (Gibson et al. 2019).

Currently, eSTZs are distributed exclusively as polygon vector data (e.g., shapefiles). Vector data convey a sense of separation between the entities they represent, that is, they are composed of mutually exclusive categories with discrete borders between them. Common examples of polygon vector data usage include administrative units (e.g., zip codes, states, and countries), watersheds, and the geographic range of a species. Raster data, or gridded surfaces, are used to represent continuous phenomena, that is, gradients. Common examples of raster data usage include climate variables, land cover classes, and predictions of species-modeled habitat suitability.

Although we agree with the consensus that vector data are generally the best method of distributing data, given the number of times we have observed classification confusion, we believe that the inclusion of raster data is also warranted. Raster data have an additional benefit that they can intuitively incorporate multiple layers (a ‘raster stack’) for each of their pixels (although this functionality is easily achievable with GeoPacks). This allows for the first layer of consensus predictions (the data conveyed in a vector dataset) and other levels of raw predictions. As a simple example, a raster with four layers would have three layers of raw model output, while the final layer is a consensus of these products. In the case of regression-type analyses, two layers could represent predictions at the lower and upper confidence intervals, and the final layer could represent a model prediction, whereas in the case of a classification algorithm, the three classes with the highest predicted probabilities and a consensus class would be present.

We believe that conveying these uncertainties will allow users to better understand and explore caveats using model predictions. This practice is further grounded in best scientific practice because the spatial data used to develop the initial zones are imperfect representations of the modelled phenomenon itself, which

introduces a variety of errors, and the classification (modelling) process itself is a generalization (Ascough et al. 2008; Li et al. 2018; Lechner et al. 2012).

A meta-analysis of genetic data sets, collected over large spatial domains in Western North America, found that the probability of mixing populations of more distantly related evolutionary lineages (i.e. different clusters, or K , identified by STRUCTURE analyses) increases as a function of distance between the prospective populations (Massatti et al. 2020). These analyses suggest that to avoid the possibility of outbreeding depression at a restoration site the most appropriate seed source will likely be the most proximate source, relative to a more distal source from within the same pSTZ. This work provides a precedent which supports restoration ecologists in leaning towards using closer, rather than further away seed sources even if they are located within the same seed zone; combining the above findings with metrics of uncertainty from eSTZ classifications will further bolster decision making.

The use of raster surfaces are already implemented in the Seedlot Selection Tool and Climate-Smart Restoration Tool, which blend attributes of pSTZs and eSTZs, or can be used to select the most appropriate eSTZ based seed source factoring in observed and forecast (e.g. RCP pathways) climatic values and custom functions (St. Clair et al. 2022). These tools offer both an alternative format for guiding native germplasm development (e.g. spatially where should germplasm be developed from?) via gap analysis, and especially for choosing a priority seed source for a restoration when multiple are available. However, one drawback of the existing implementation of these tools is that they calculate climate similarity from a single focal cell, e.g. a restoration site, to the entire surface. In many instances seed lots need to be developed *before* a restoration site is known, for example in the case of reseeded severe disturbances (e.g. wildfires). Combining the development of seed sources within zones, which allows for decision making to develop seed availability, with the Climate-Smart Restoration Tool to select potential seed source options after a natural disaster may lead to the most pragmatic decisions.

IMPLEMENTATION

Software

To make these suggestions easy to implement we have created an R package, eSTZwritR (pronounced ‘easy rider’), which can implement all of them, less the statistical processing, with minimal user inputs. The package is installable from GitHub <https://github.com/sagesteppe/eSTZwritR> and has a GitHub Pages website (<https://sagesteppe.github.io/eSTZwritR/>) for users interested in better understanding its functionality, and

which includes supplemental figures and details not discussed here.

FOR DEVELOPERS

The package requires only 5 functions to produce a directory with the contents discussed above, with minimal data entry. Most importantly, the entries are well outlined and easily entered without requiring close attention to detail.

FOR PRACTITIONERS

These results should allow for simple utilization of existing empirical seed-transfer zone resources. We have re-processed all eSTZ data products that we are aware of following these standards, with the exception of creating uncertainty raster layers. We provided some sample code that showcases loading these data into a non-GUI GIS on the website above.

CONCLUSIONS

Seed-based active restoration will always be a relatively expensive yet necessary option for terrestrial restoration. Here, we present simple standards for scientists developing eSTZs to use in order to standardize the data products they are developing to assist in their uptake. Although these conventions should be easy to implement for a sufficiently motivated individual, we also present an R package that can quickly achieve these results.

ACKNOWLEDGEMENTS

Two anonymous reviewers are thanked for their constructive comments on this manuscript.

DATA AVAILABILITY STATEMENT

The scored data for the analysis, and post processed eSTZ data sets are openly available on Dryad at ... (doi)... reference number ...

Term box

Seed transfer zones are intended to improve restoration outcomes by decreasing the chances of using mal-adapted seeds at a restoration site (Kramer & Havens 2009). They seek to characterize (empirically) or estimate (provisionally) local adaption and aim to minimize the variation between prospective seed sources and restoration sites (Leimu & Fischer 2008; Kramer & Havens 2009). Provisional seed transfer zones (pSTZs) can be utilized across all vascular plants and are based on the similarity between several climate variables known to be broadly relevant to plant life in a region (e.g., in the US). Winter Minimum Temperature, and Annual Heat:Moisture index) (Bower et al. 2014). eSTZs have the same goal as pSTZs but are tailored for individual species, allowing for more accurate determination of zones in which seeds can be transferred under existing climate regimes.

The development of seed transfer zones in the US can be traced back to the 1960s when forestry companies, which had to replant timber stands after logging, developed guidance to ensure the success of their replantings (Johnson et al. 2004). However, it was not until the early 21st century that researchers began developing STZs for other groups of plants in the US (McKay et al. 2005), in part because of the increasing size of large wildfires and demand for locally adapted seeds (National Academies of Sciences et al. 2023).”

**Tandem to the development of STZs for non-tree species in the US has been the development of STZs in both Europe (Johnson et al. 2004).*

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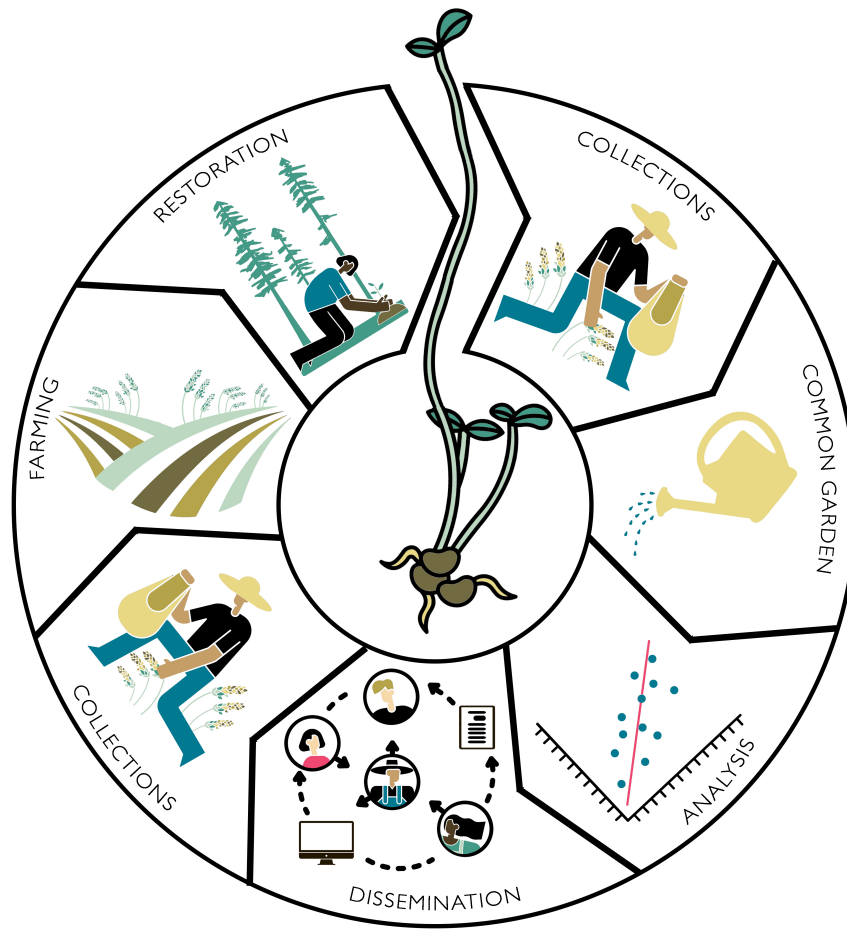


Figure 1: Dissemination. The first three panels ('collections', 'common garden', 'analysis') indicate the process of developing an eSTZ, while the 'dissemination' panel showcases the need to share results so they can inform operational seed collections, agricultural increase, and selection of materials for a restoration. Note that 'common garden' represents the act of eSTZ development, but could be replaced with molecular work to develop a genetic product, or modelling for a climate matched product. By Emily Woodworth

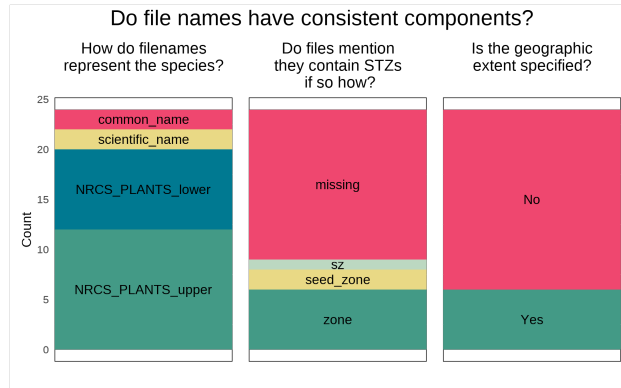


Figure 2: File Naming. Three inconsistencies in file names discussed here, with the advised format for data sharing in green, and the least desirable condition in grey.

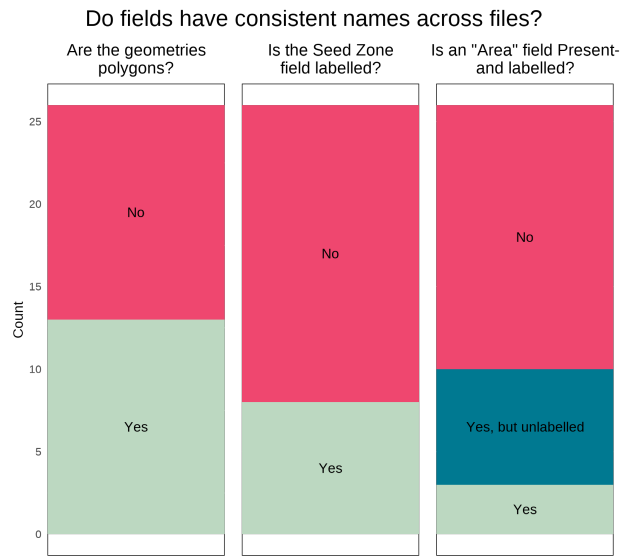


Figure 3: Field Names Shapefile. The three attributes of field names discussed here, with the most desirable condition in green, and the least desirable condition in grey.

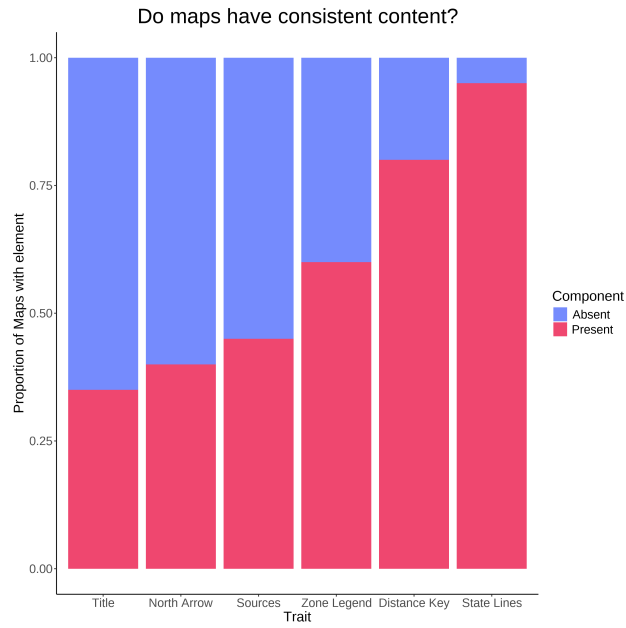


Figure 4: Map Components ($n = 20$). Several essential cartographic elements - most notably a Title, a statement on data sources, and a legend for the seed zones, were missing from at least - or nearly half of the products inspected.

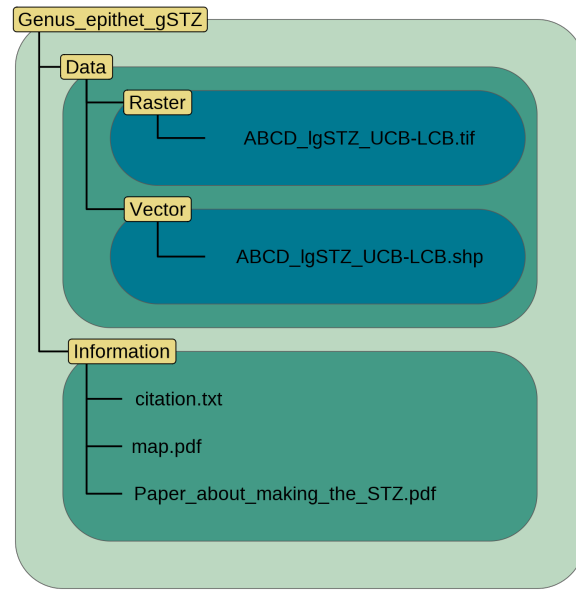


Figure 5: Directory Structure. Each directory is named in yellow, and spans the extent of variously coloured polygons. Individual files (or a set of files in the case of a shapefile) are depicted in black text within these directories.

File naming convention

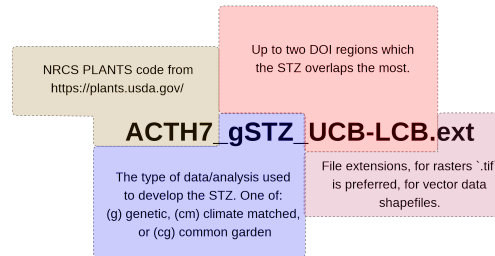


Figure 6: File Naming. The four proposed components of a filename are highlighted in different colours, and with appropriate cases.

Example field names in a shapefile					
ID	SeedZone	SZName	AreaAcres	BIO1_R	BIO2_mean
1	1	Salt Desert	12340	20.2	5.1
2	2	Desert Scrub	14230	19.1	7.1
3	3	Pinyon-Juniper/Oak Brush	30142	15.1	10.1
4	4	Montane	9872	12.3	12.3
The first four (blue) fields should be in every file. More fields are optional.					

Figure 7: Vector Data Field Attributes. The proposed field names for distributing vector data.

Three model predictions
from a classifier
and a consensus layer

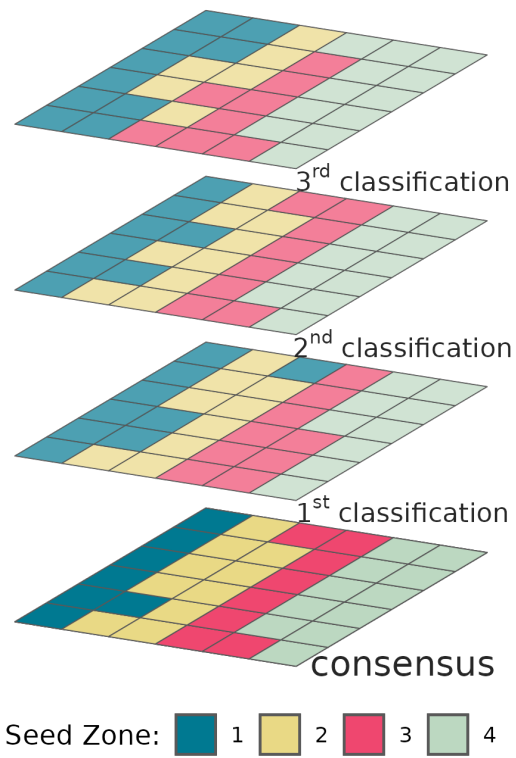


Figure 8: Many common classification algorithms can output probabilities for each class in the model. Reporting a subset of these allows users to interact with prediction probabilities directly.

311 **APPENDIX**

312

[Figure 9 about here.]

313 List of Figures

Appendix 1

All eSTZ data products reviewed and scored.

Authors	Year	Publication	Journal
Bradley St. Clair, J., Kilkenny, F. F., Johnson, R. C., Shaw, N. L., & Weaver, G.	2013	Genetic variation in adaptive traits and seed transfer zones for <i>Pseudoroegneria spicata</i> (bluebunch wheatgrass) in the northwestern United States	<i>Evolutionary Applications</i> , 6(6), 933-948
Doherty, K. D., Butterfield, B. J., & Wood, T. E.	2017	Matching seed to site by climate similarity: techniques to prioritize plant materials development and use in restoration	<i>Ecological Applications</i> , 27(3), 1010-1023
Erickson, V. J., Mandel, N. L., & Sorensen, F. C.	2004	Landscape patterns of phenotypic variation and population structuring in a selfing grass, <i>Elymus glaucus</i> (blue wildrye)	<i>Canadian Journal of Botany</i> , 82(12), 1776-1789
Johnson, R. C., Erickson, V. J., Mandel, N. L., St Clair, J. B., & Vance-Borland, K. W.	2010	Mapping genetic variation and seed zones for <i>Bromus carinatus</i> in the Blue Mountains of eastern Oregon, USA	<i>Botany</i> , 88(8), 725-736
Johnson, R. C., Cashman, M. J., & Vance-Borland, K.	2012	Genecology and seed zones for Indian ricegrass collected in the southwestern United States	<i>Rangeland Ecology & Management</i> , 65(5), 523-532
Johnson, R. C., Hellier, B. C., & Vance-Borland, K. W.	2013	Genecology and seed zones for tapertip onion in the US Great Basin	<i>Botany</i> , 91(10), 686-694
Johnson, R. C., Horning, M. E., Espeland, E. K., & Vance-Borland, K.	2015	Relating adaptive genetic traits to climate for Sandberg bluegrass from the intermountain western United States	<i>Evolutionary Applications</i> , 8(2), 172-184
Johnson, R. C., & Vance-Borland, K.	2016	Linking genetic variation in adaptive plant traits to climate in tetraploid and octoploid basin wildrye [<i>Leymus cinereus</i> (Scribn. & Merr.) A. Love] in the Western US	<i>PLoS One</i> , 11(2), e0148982
Johnson, R. C., Leger, E. A., & Vance-Borland, K.	2017	Genecology of Thurber's Needlegrass (<i>Achnatherum thurberianum</i> [Piper] Barkworth) in the Western United States	<i>Rangeland Ecology & Management</i> , 70(4), 509-517
Massatti, R.	2019	Genetically-informed seed transfer zones for <i>Pleuraphis jamesii</i> , <i>Sphaeralcea parvifolia</i> , and <i>Sporobolus cryptandrus</i> across the Colorado Plateau and adjacent regions.	<i>USGS Report</i> , 37
Massatti, R., & Winkler, D. E.	2022	Spatially explicit management of genetic diversity using ancestry probability surfaces	<i>Methods in Ecology and Evolution</i> , 13(12), 2668-2681
Shryock, D. F., Havrilla, C. A., DeFalco, L. A., Esque, T. C., Custer, N. A., & Wood, T. E.	2017	Landscape genetic approaches to guide native plant restoration in the Mojave Desert	<i>Ecological Applications</i> , 27(2), 429-445

Figure 9: All data sets scored