Optimizing plant size:biomass allometric relationships at the Sevilleta LTER

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Background: Accurately estimating primary production is a core research theme of all Long-Term Ecological Research (LTER) sites. Non-destructive methods of measuring productivity are preferable in arid regions where plants recover slowly from disturbance. Allometric equations allow researchers to relate non-destructive measurements such as plant height or cover to plant biomass. Here, we present the updated workflow used to generate allometric equations for plant species at the Sevilleta LTER in central New Mexico.

LTER Data: Over the past 20 years, Sevilleta researchers have revisited thousands of sampling points to record well over a million observations of plant identity, abundance and size during the peak of each spring and fall/monsoon growing season. These locations span desert grasslands, shrublands, savannahs, and piñon-juniper woodlands. We use allometric equations to convert these <u>non-destructive</u> <u>observations</u> of plant size into a common unit of <u>biomass</u>.

In order to create species-specific allometries, we used a <u>destructive harvest</u> dataset of >22,000 individual plants representing nearly 200 species. Specimens growing in a range of biomes and climatic conditions are measured, cut at ground level, dried, and weighed.

All observations and harvested samples are paired with weather data from a nearby meteorological tower.

In order to make Sevilleta LTER data as understandable and transparent as possible, all data manipulation and analyses were conducted in a free, open-source software, R, and all code was committed to GitHub.

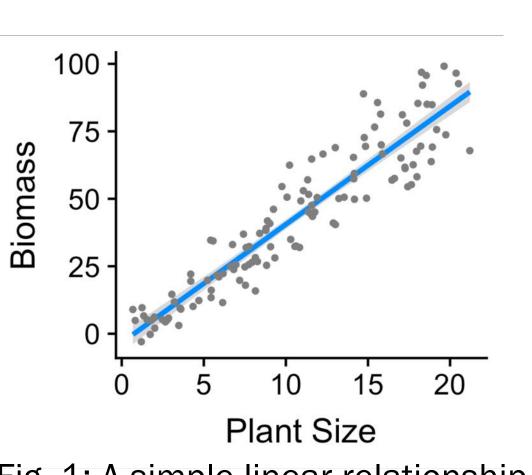


Fig. 1: A simple linear relationship between plant size and mass

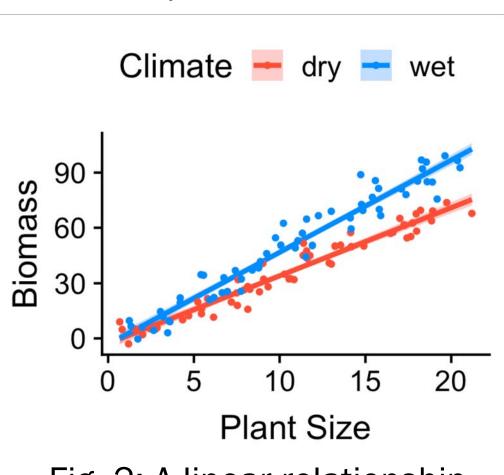


Fig. 2: A linear relationship between plant size and mass that varies with climate

Allometric Models: For each species we first considered the simple linear relationship between plant size (either ground cover or plant volume) and biomass. We also tested whether the size:mass relationship varied between years, biomes, or in response to climate. Climate variables considered include seasonal precipitation, growing degree days (GDD, a measure of accumulated temperature), or 6month SPEI_{comp}, (a measure of how hot and dry the preceding 6 months were at a given site, compared to all other 6-month periods from all sites.

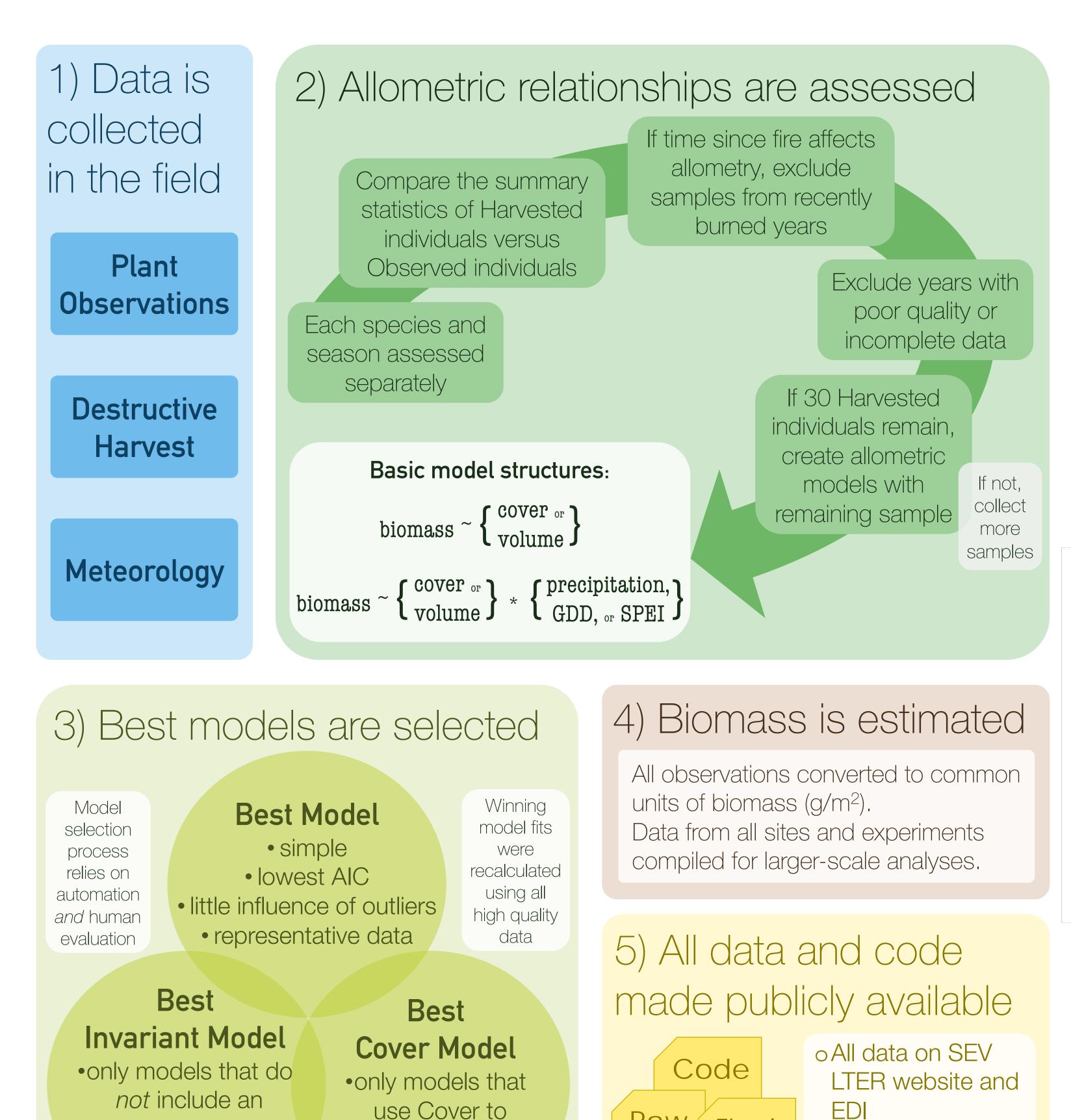


Fig. 3: Diagram of workflow used to calculate biomass at the Sevilleta LTER.

predict biomass

o Easy-to-read R

GitHub

Markdown files on

data

interacting climate

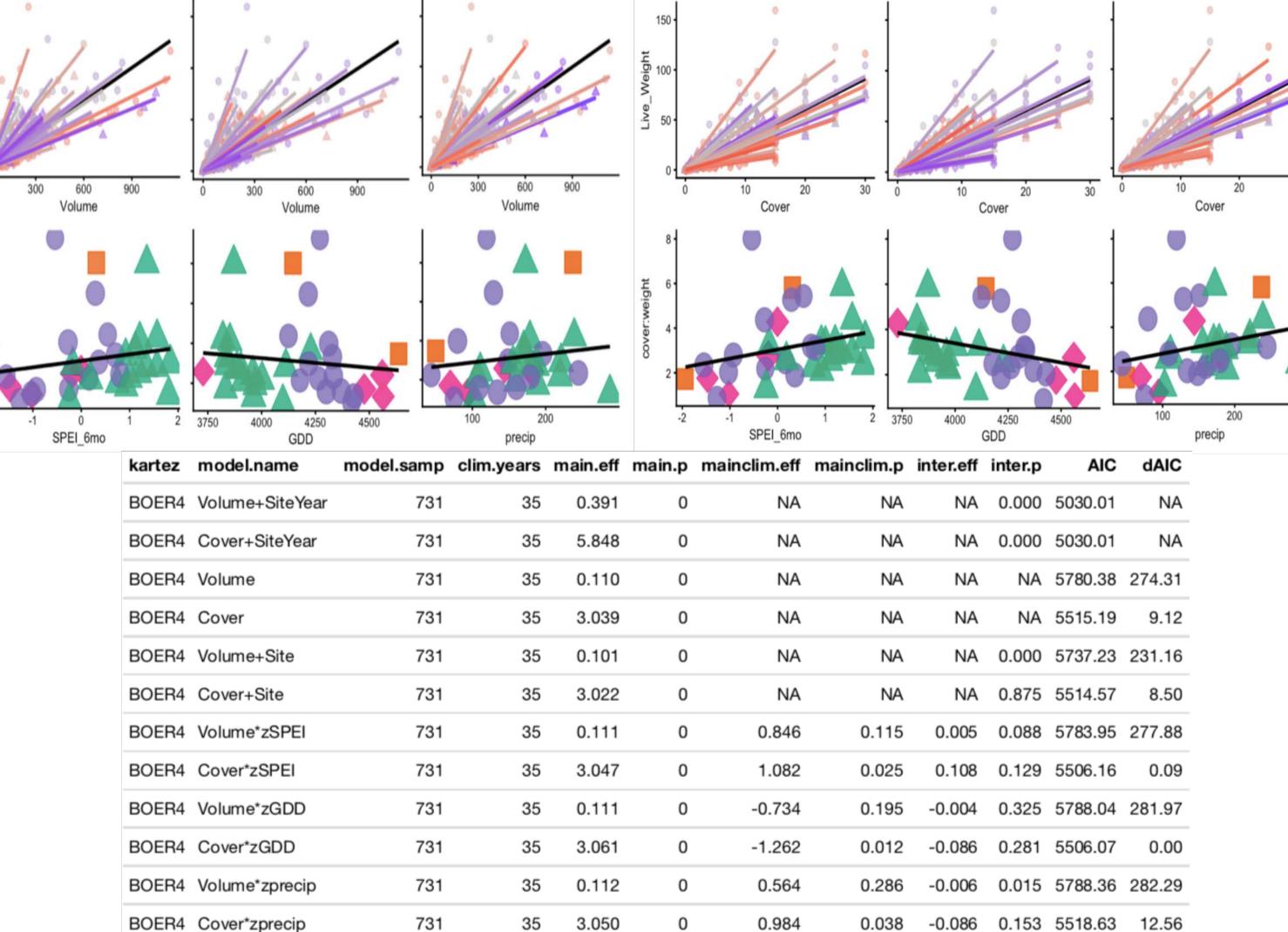


Fig. 4: Diagnostic plots and model summaries used to select Best Model for black grama (Bouteloua eriopoda)

"Winning" Models: We were able to create strong species- or genus-specific allometric models for 174 species-seasons, representing over 90% of all Sevilleta core site observations. Aggregated plant functional type allometric equations were used for rarer species.

Cover predicted biomass better than volume in 67% of cases. Climate-invariant models performed best in 77% of cases. Of those models that did show strong climate interactions, 47.5% varied more with seasonal precipitation, 37.5% with temperature (GDD), and 15% with aridity (SPEI).

Select time series:

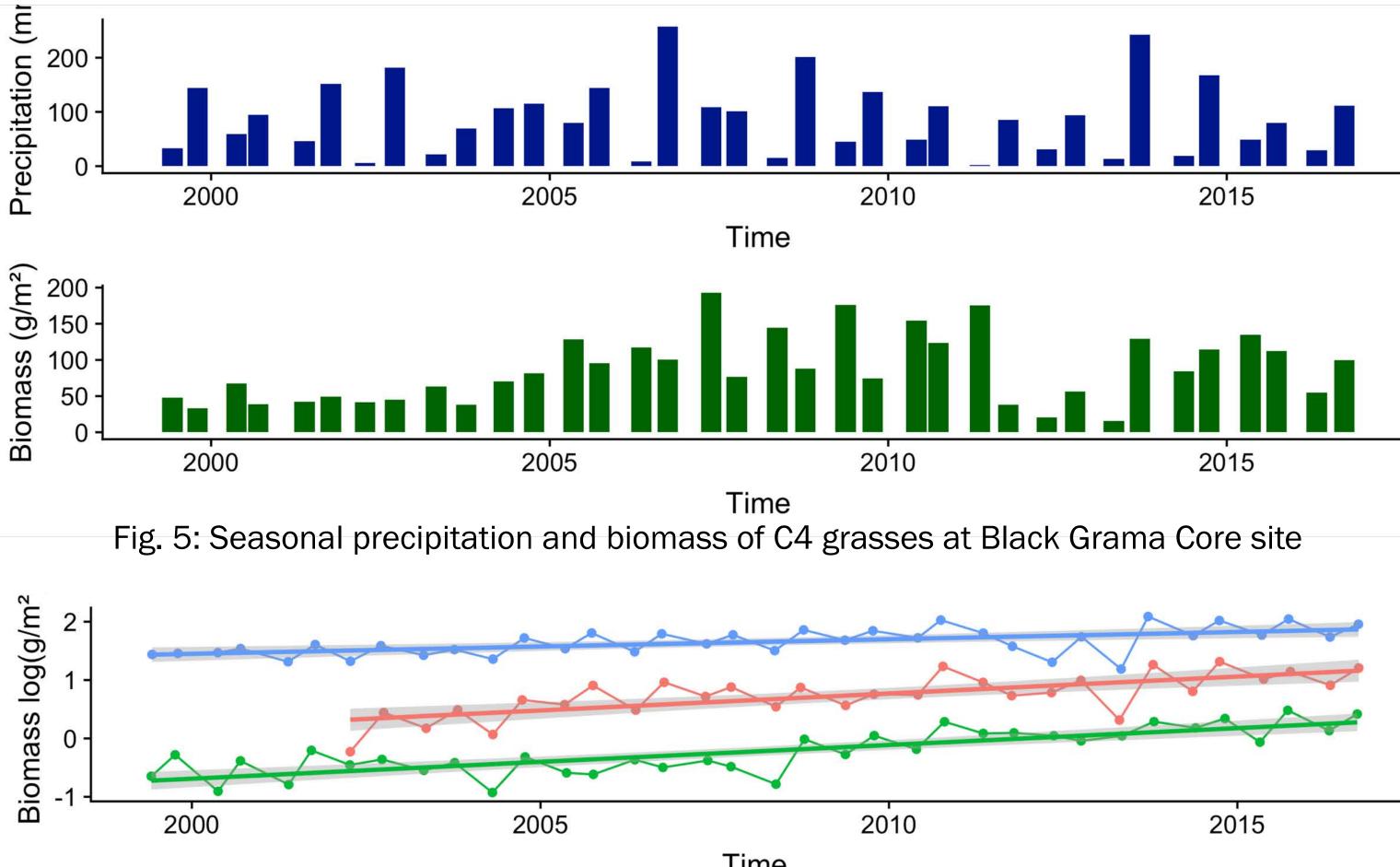


Fig. 6: Biomass of black grama (Bouteloua eriopoda) at core sites increases over time.

Core Site - Blue Grama - Creosote - Black Grama

Outcomes: We were able to find robust allometric relationships for most species (percent of observations) at the Sevilleta LTER. The majority of models were climate-invariant and many only required measurements of cover to predict biomass. This supports the use of these models in novel biomes and climate scenarios. It may also reduce field efforts and aid biomass estimation using aerial drone photography in the future. However, it should be noted that invariant models were selected in multiple cases due to their simplicity and to exclude confounding outliers.

Future work includes the inclusion of more species in our destructive harvest dataset, analysis of the phylogenetic basis of allometric relationships, and exploration of fruiting allometries.









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