

Mast Cell Tumors in Guiding Eyes for the Blind Dogs

Heather McDonough, McKenzie Jones, and Stephanie Wang

University of Richmond

May 27, 2016

Introduction

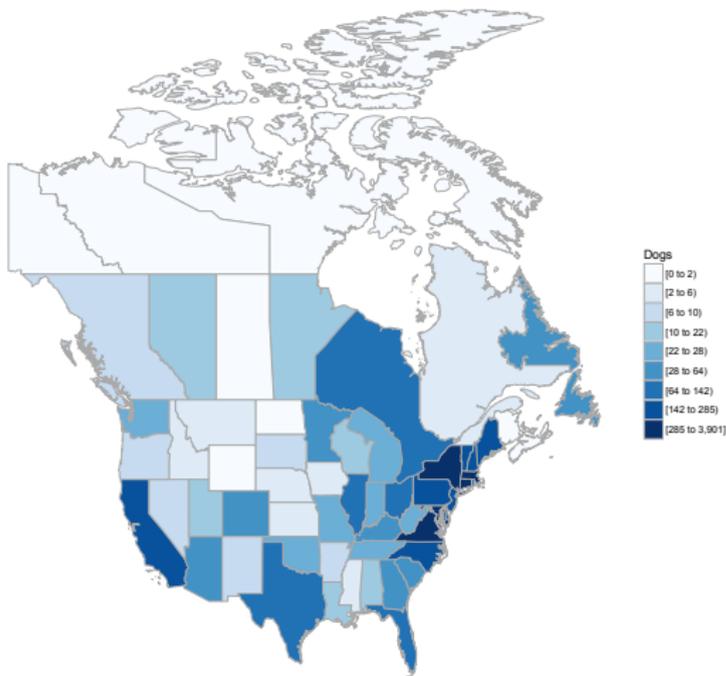
Our lab has been working on analyzing data on mast cell cancer in Guiding Eyes for the Blind dogs using R programming. This presentation will illustrate the work we have done so far and what we hope to accomplish in the upcoming weeks.

Data

We have information on over 10,000 dogs used in the Guiding Eyes for the Blind program that have been diagnosed with Mast Cell Cancer. Within this data set, we have focused on date of birth, sex, death date (if applicable), city, state, zip code, age at diagnosis, and MCT Score Value. Using this data, we have been able to create choropleth maps to illustrate where the dogs are from and ggplots to look at the correlation between date of birth and age of diagnosis in the dogs.

Where The Dogs Are From

Choropleth of Dog Data in Canada and Continental United States by State



Choropleth of Dog Data in Northeast

Choropleth of Dog Data in Northeastern Region per State



Choropleth by Zip Code NY

Choropleth of Dog Data from New York



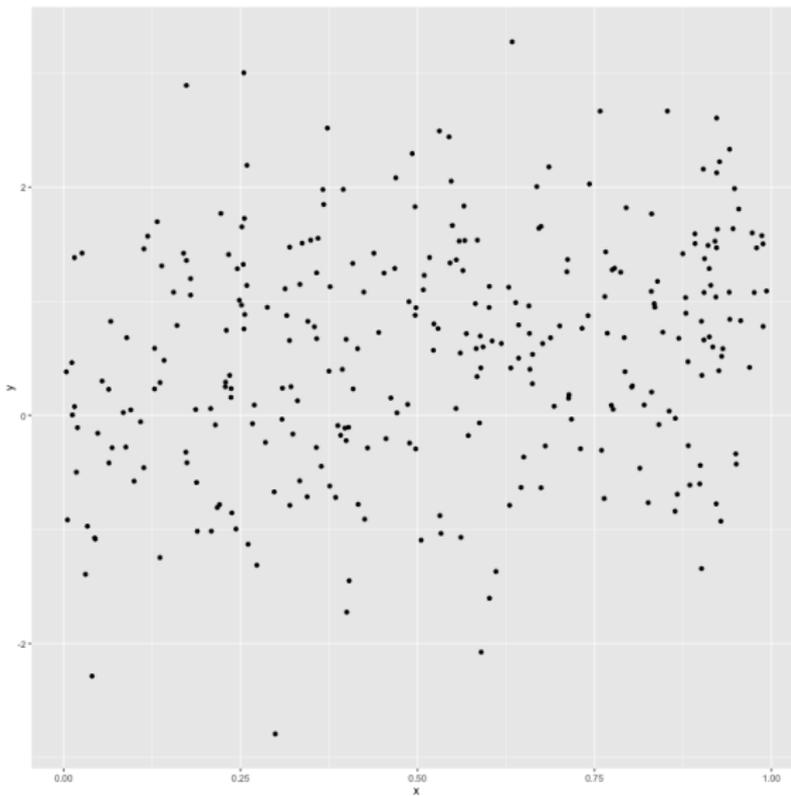
Choropleths in R

To create this map, we:

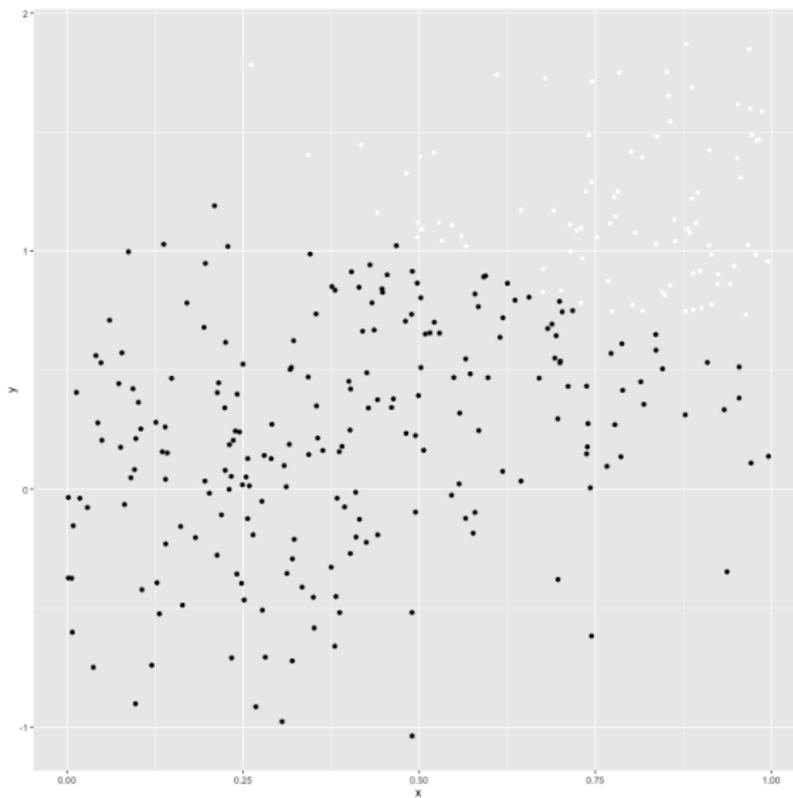
- Installed packages that allow R to work with zip codes, states, and regions, and use GGPlot to create choropleth maps, which are used to display themes such as population density
- Created maps of US states with Canadian provinces, zoomed in on Northeastern US states, and created another map using New York zip codes
- Packages used: `choroplethr`, `choroplethrMaps`, `choroplethrZip`, `admin1_region_choropleth`

ggplot

Truncated Data

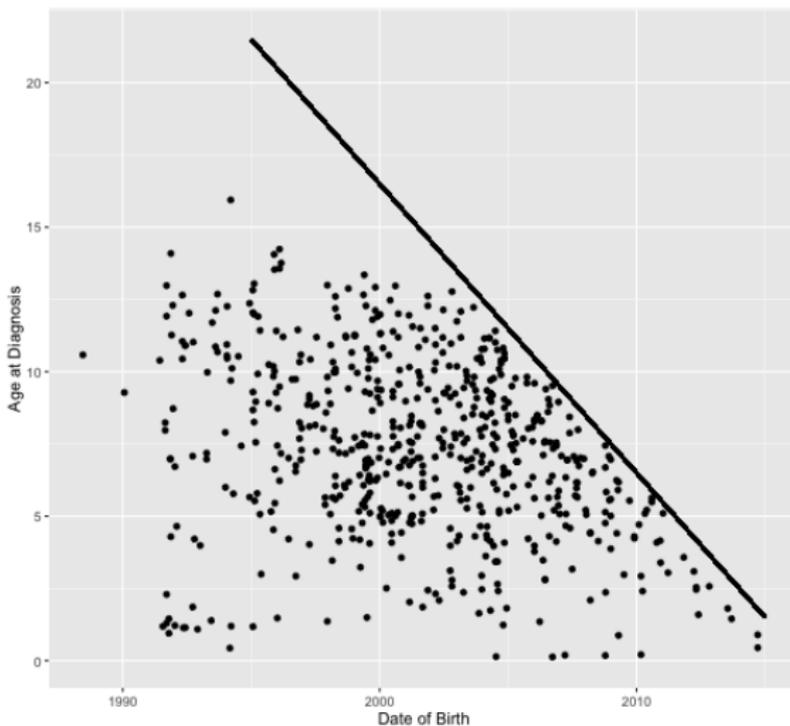


Truncated Data



ggplot

Diagnosis Age by Date of Birth



Truncated Random Variables

A truncated distribution means that at a certain point, the data is cut short, and we do not have complete information about one end of the distribution, as it cannot be observed

In our case, our data is truncated. As we approach the current year, any dogs born in recent years who are diagnosed with cancer are only diagnosed at younger years. For example, given a dog born in 2014 and diagnosed with cancer in its lifetime, it will only be in our data if its diagnosed by the age of two years old. If it will be diagnosed at a later age, we do not have that information yet.

Truncated Random Variables

Our scatterplot of the data makes it seem as though the age of diagnosis has gotten younger over the past decade. This is a false trend because, in reality, dogs that have been born recently cannot be diagnosed with cancer at older ages yet. Although they may develop MCT later in life, we do not have this data, so we cannot make conclusions taking our current data at face value.

Future Work

In general, we want to test the hypothesis that dogs are being diagnosed with cancer at younger ages as years pass. To do this, we have to use statistical tools to account for this truncated data while focus on modeling the data we have from earlier years of the study. We can then focus on analyzing this claim using statistical models to allow us to predict the age of diagnosis for dogs that will develop MCT in the future.