

Final-Project-Paper

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Following Alligatoridae Family Species Diversity Through Time



Figure 1: Alligator mississippiensis

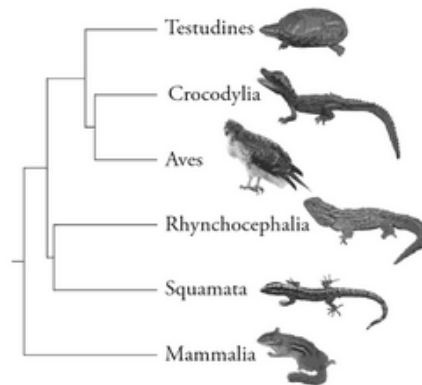
Abstract

Introduction

The *Alligatoridae* family is made up of species classified as alligators and caimans. Alligatoridae belongs to the order Crocodilia which is a sister taxa to birds(aves) and turtles(Testudines) due to the presence of the amniotic sac (Dodd 2016) (Figure) (Fig2). All members of the Alligatoridae are carnivorous. Today the *Alligatoridae* family is distributed in Central and South America as well as the Southeastern United States (*The natural history collections* n.d.). Their distinguished features such as massive skull and short, broad snout have not change significantly since the late Triassic. Members of *Alligatoridea* do not tolerate salt water, therefore their main way of dispersal was via land bridges (Benton 2015). There is evidence that within *Alligatoridea* the caiman subclass evolved from the alligator subclass after its dispersal southward from North America to South America.



labelfig:pop2



Methods

In order to obtain data to analyze, I used the Paleobio Database that was shown in class. From here I searched for occurrence data for the *Alligatoridae* family and found that my original data file from the pbdb website contained 444 specimens using the following shell command:

```
tail -n +19 alligatoridae_pbdb_data.csv | cut -d "," -f7 | sort | wc -l
```

Specimens were identified to the family, genus or species rank within my dataset. Therefore, before I began to analyze the data, I sorted through each occurrence and selected only data which was identified to the species rank. This allowed the creation of a species ranges dictionary using the following python code. Something that was important to include in the function was an if statement that served to only collect data which was identified to the species rank.

```
species_ranges=defaultdict(list)
for line in alligator:
    items = line.split(',')
    min_ma = round(float(items[15]),3)
    max_ma = round(float(items[14]),3)
    species_name = items[9]
    if re.search(r"species", line):
        species_ranges[species_name].append(str(min_ma))
        species_ranges[species_name].append(str(max_ma))
```

From this dictionary, I wrote a .csv file that contained only the genus, species, min_ma and max_ma for specimens identified to the species rank.

```
output=open("alligatoridae_ranges.csv", "w") #i am making the output file
for key, values in species_ranges.items():
    values.sort()
    #the largest value appears last in list and smallest appears first in list
    max_age = values[-1]
    min_age = values[0]
    genus=key.split(" ")[0]
    outline= "{},{},{},{}\n".format(genus, key, min_age, max_age)
    print(outline)
    output.write(outline)
```

The "alligatoridae_ranges.csv" file was read into in Rstudio to first label the data with the appropriate labels then to create an occurrence through time plot using ggplot and forcats:

```
library(ggplot2)
alligatoridae <- read.csv("/home/eeb177-student/Desktop/eeb177-final-project/alligatoridae_ranges.csv",
names(alligatoridae) <- c("genus", "species", "minage", "maxage")

library(forcats)
alligatoridae_occ <- ggplot(alligatoridae, aes(x = fct_reorder(species, minage, .desc = T), maxage, col=
alligatoridae_occ + geom_linerange(aes(ymin = minage, ymax = maxage)) + coord_flip() + theme(axis.text.x =
```

Furthermore, using Pyrate a series of plots was constructed showing speciation, extinction rates for the extant species belonging to *Alligatoridae*. First the *Alligatoridae data was processed in R so that pyrate could be run. A total of 1,000,000 simulations were run to create the plots.

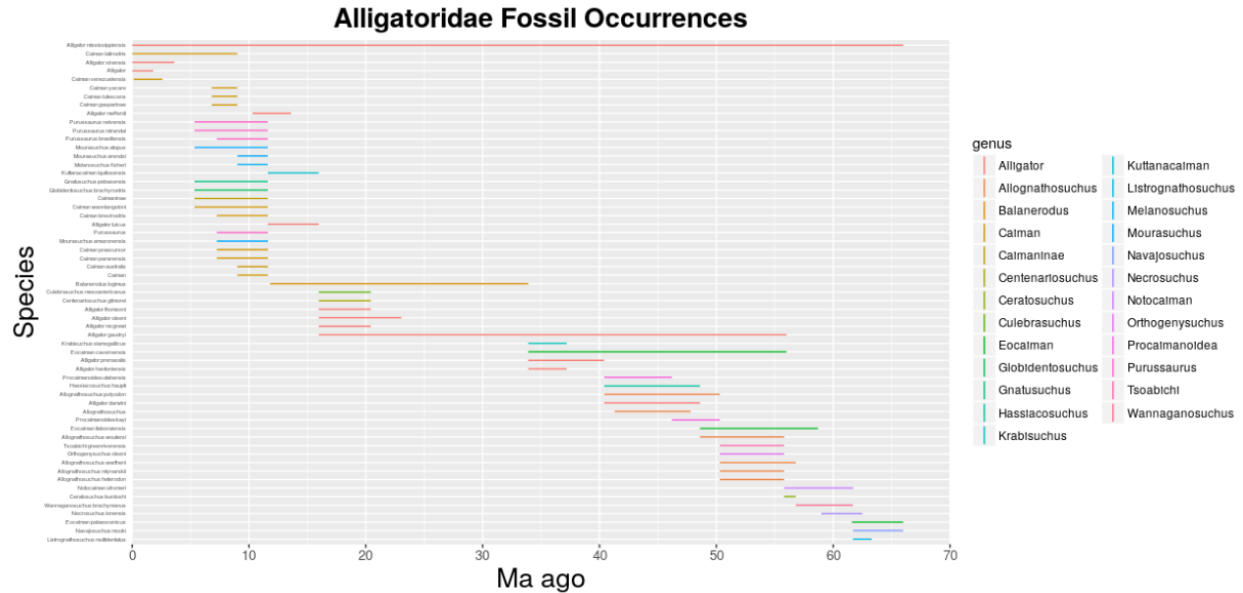
```
source("~/PyRate/pyrate_utilities.r")
extant_alligatoridae = c("Alligator mississippiensis", "Alligator sinensis", "Caiman latirostris", "Caiman
extract.ages.pbdb(file = "alligatoridae_occ.csv", extant_species = extant_alligatoridae)
```

Orthogenysuchus, Orthogenysuchus olseni, 50.3, 55.8
 Caiman, Caiman australis, 11.62, 9.0
 Alligator, Alligator mcgrewi, 15.97, 20.43
 Caiman, Caiman wannlangstoni, 11.608, 5.333
 Alligator, Alligator hantoniensis, 33.9, 37.2
 Tsoabichi, Tsoabichi greenriverensis, 50.3, 55.8
 Caiman, Caiman gasparinae, 6.8, 9.0
 Caiman, Caiman venezuelensis, 0.126, 2.588
 Notocaiman, Notocaiman stromeri, 55.8, 61.7
 Allognathosuchus, Allognathosuchus woutersi, 48.6, 55.8
 Caiman, Caiman praecursor, 11.62, 7.246
 Alligator, Alligator thomsoni, 15.97, 20.43
 Wannaganosuchus, Wannaganosuchus brachymanus, 56.8, 61.7
 Listrognathosuchus, Listrognathosuchus multidentatus, 61.7, 63.3
 Melanosuchus, Melanosuchus fisheri, 11.608, 9.0
 Allognathosuchus, Allognathosuchus wartheni, 50.3, 56.8
 Alligator, Alligator mississippiensis, 0.0, 66.0
 Procaimanoidea, Procaimanoidea kayi, 46.2, 50.3
 Allognathosuchus, Allognathosuchus mlynarskii, 50.3, 55.8
 Allognathosuchus, Allognathosuchus heterodon, 50.3, 55.8
 Alligator, Alligator sinensis, 0.012, 3.6
 Eocaiman, Eocaiman itaboraiensis, 48.6, 58.7
 Mourasuchus, Mourasuchus amazonensis, 11.62, 7.246
 Purussaurus, Purussaurus mirandai, 11.608, 5.333
 Hassiacosuchus, Hassiacosuchus haupti, 40.4, 48.6
 Ceratosuchus, Ceratosuchus burdoshi, 55.8, 56.8
 Alligator, Alligator, 0.012, 1.8
 Caiman, Caiman, 11.62, 9.0
 Allognathosuchus, Allognathosuchus polyodon, 40.4, 50.3
 Purussaurus, Purussaurus neivensis, 11.608, 5.333
 Caiman, Caiman paranensis, 11.62, 7.246
 Caiman, Caiman latirostris, 0.012, 9.0

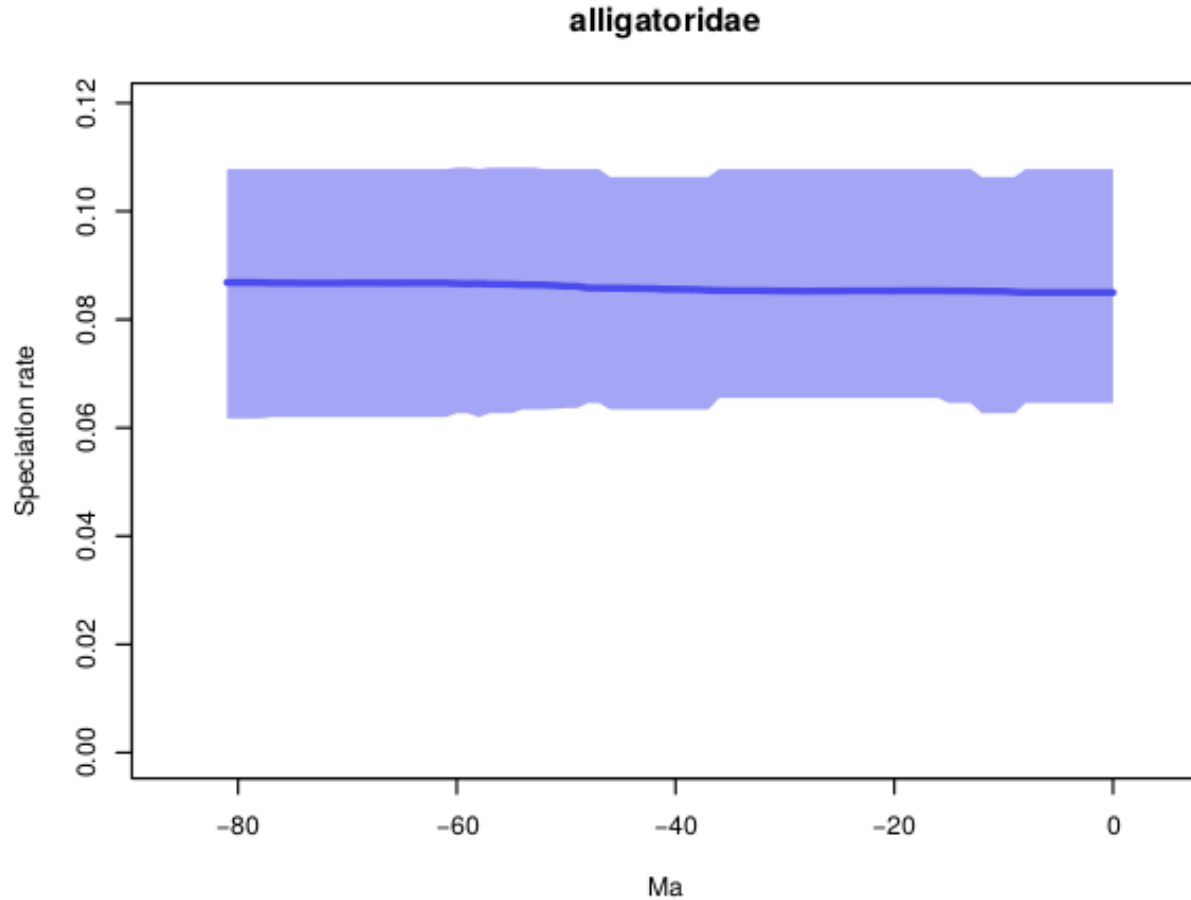
Figure 2: Alligatoridae_ranges.csv

Results

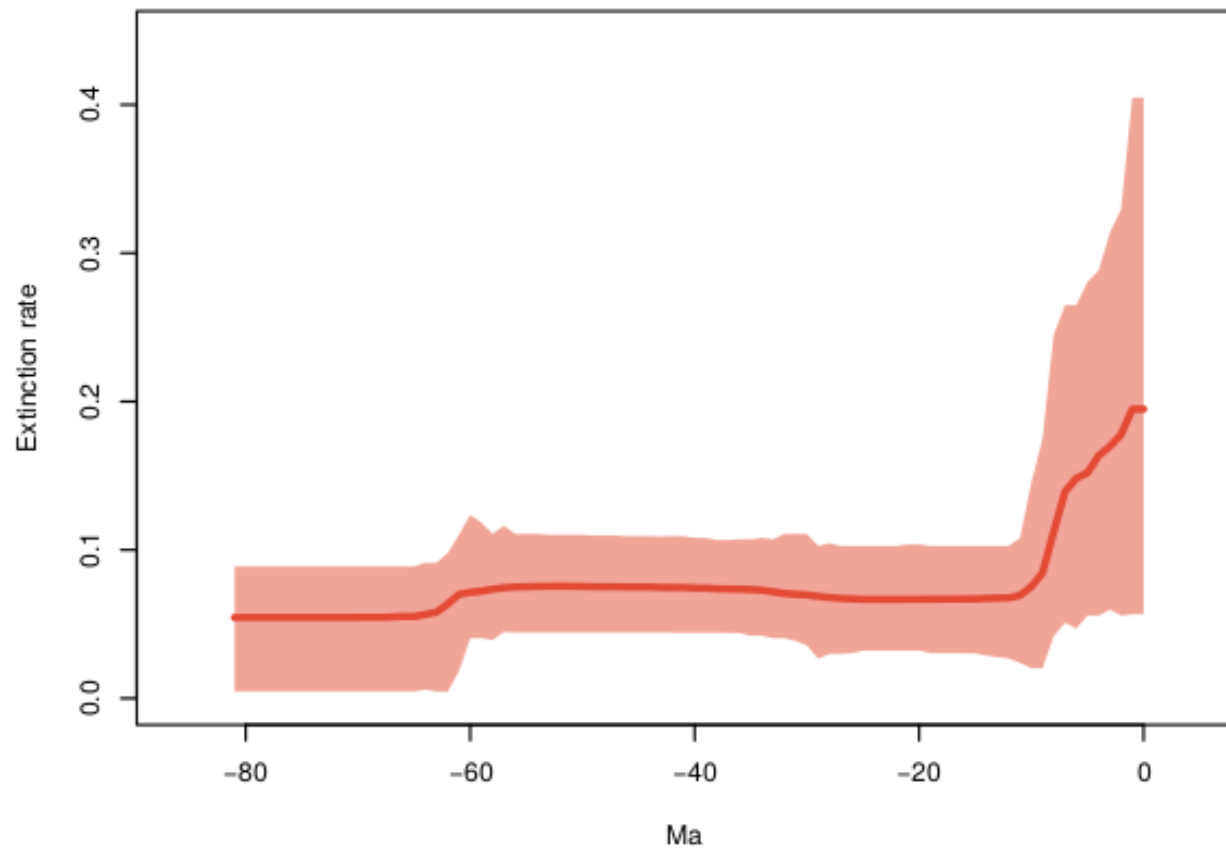
The Alligatoridae Fossil Occurrence plot shows that we have fossil evidence showing that the *Alligatoridae* family has species that date almost to 70 million years into the past around the time of the Late Cretaceous. Many of the species belonging to family *Alligatoridae* have since gone extinct except for the following species: Alligator mississippiensis, Alligator sinensis, Caiman latirostris, and Caiman venezuelensis. Of these, Alligator mississippiensis has been the longest surviving species as, according to this data, it has been in the fossil record since the oldest known *Alligatoridae* fossils nearly 70 million years old.



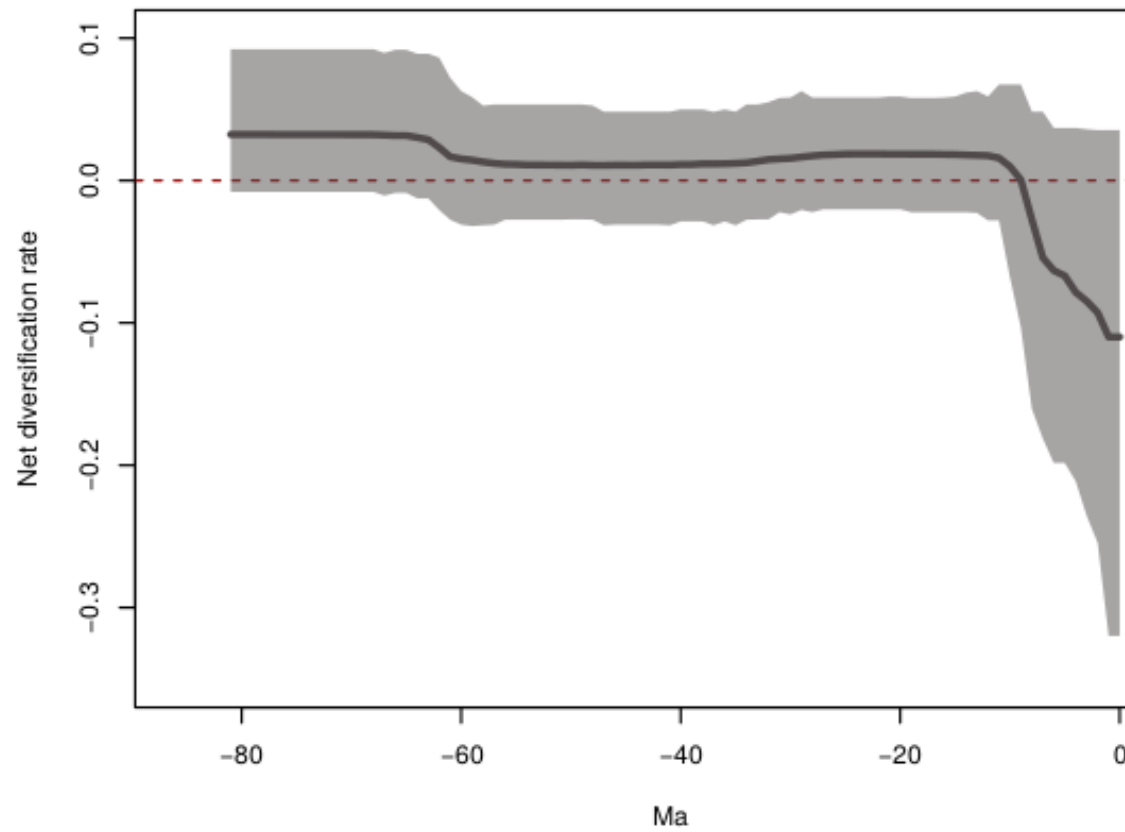
The plots from running PyRate for 1,000,000 simulations show the speciation, extinction, net-diversification and longevity of the *Alligatoridae* species. The first plot shown is the Speciation rate through time for the *Alligatoridae* family. Even with 1 million simulations run, it appears that the speciation rate was most likely to be about 0.085 and constant over the last 80 million years.



The next plot done using PyRate is an Extinction curve for the *Alligatoridae* family. There is a visible trend for increasing extinction rate occurring through time. This could be a result of more species being around in general. Although we do see that at present, only 4 extant species are represented in the fossil occurrence data.

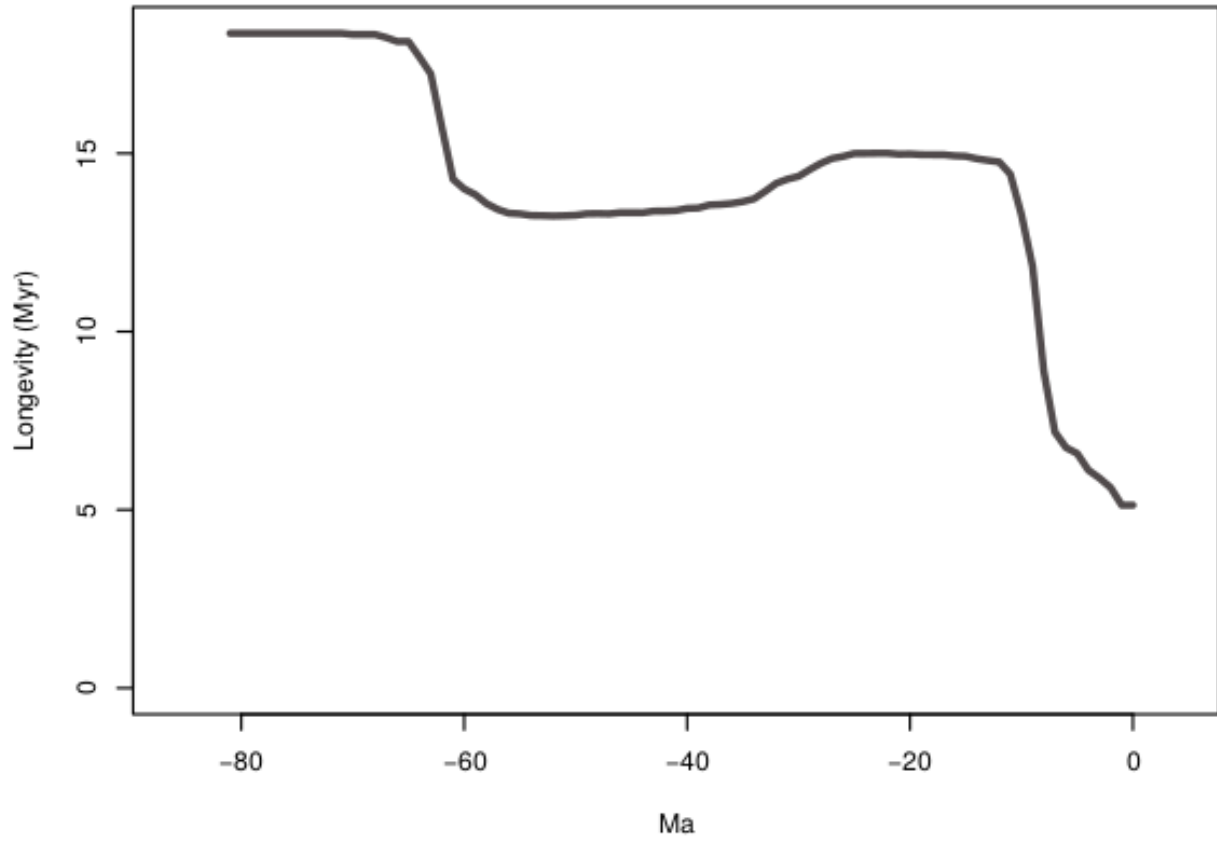


The net diversification through time plot shows that the peak diversification for *Alligatoridae* occurred early in its evolutionary history from about 80-65 million years ago. This is perhaps representative of shifting land



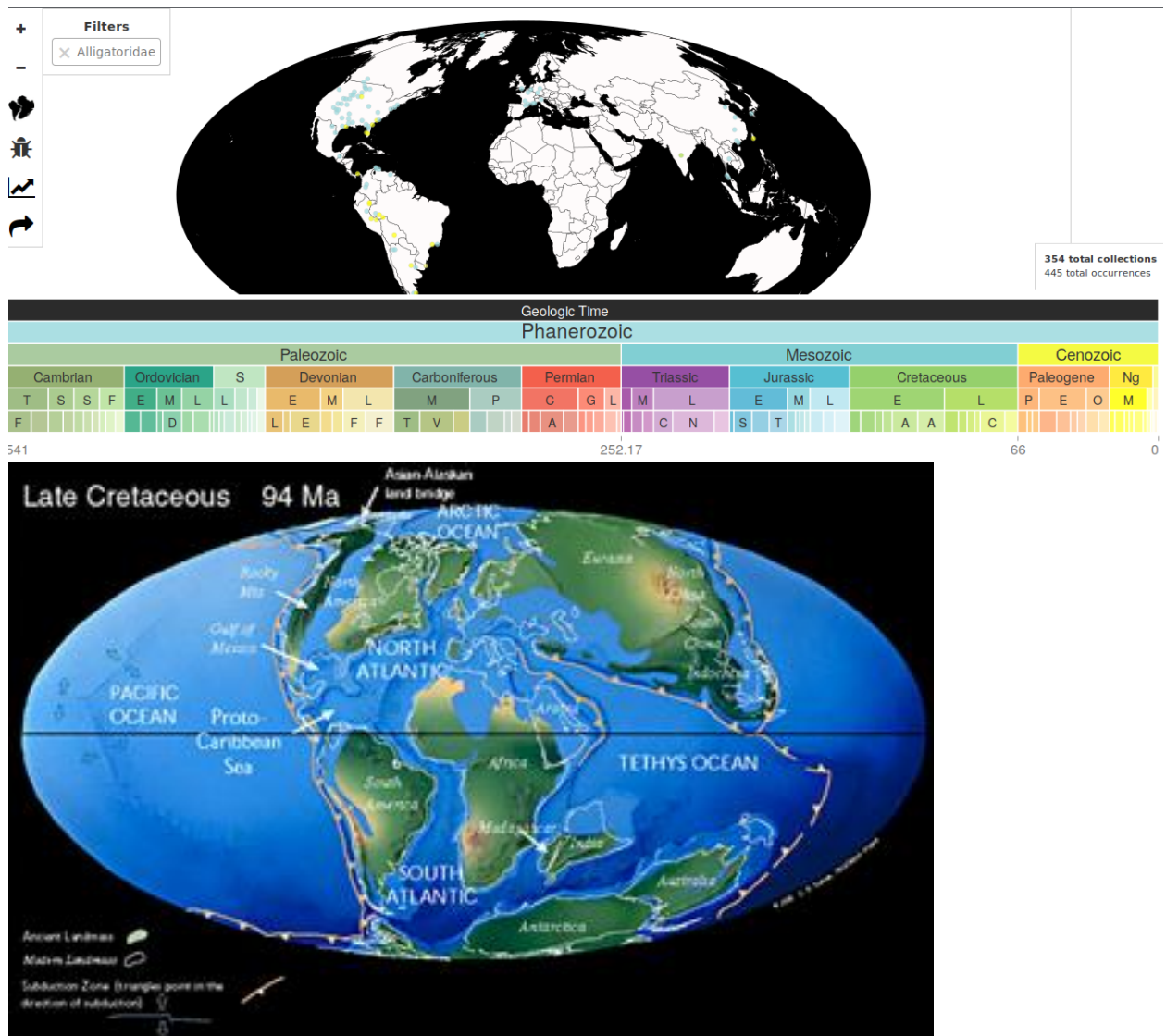
masses and then the ice age?

The final plot created using PyRate was the Longevity curve through time. This curve however does not completely agree with the *Alligatoridae* Fossil Occurences plot as it shows that the more ancient a species, the greater its longevity and this is not fully supported when viewing the occurence through time plot made using ggplot. The longevity curve shows that the more younger species had shorter longevity, this is expected as they have not been around long enough to rank high in longevity.



Discussion

It would be very interesting if the pldb website provided data on the exact locations that each specimen was found in order to recreate species occurrences map. When on the pldb website however, I found a map which showed locations of the occurrences already (it is linked below). We can see that some *Alligatoridae* specimens occurred in regions as far north as Canada. Clearly it is much too cold today in Canada at those high latitudes, therefore finding remains of members of this family either suggests that Canada once had a much warmer climate. This could be evidence supporting that tectonic plates and thus continents have shifted throughout earth's history.



Github link

link

References

1. Benton, E., Terry Jr. (2015). *The white river badlands: Geology and paleontology*. Indiana University Press.
2. Dodd. (2016). *Reptile ecology and conservation: A handbook of techniques*. Oxford University Press.
3. *The natural history collections: Family alligatoridae*. (n.d.). University of Edinburgh.