
**Obtaining rates of glacial isostatic adjustment from
Unequally spaced data**

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University of Waterloo Earth Sciences Honours Thesis

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1 Abstract

The ground surface underlying the Laurentian Great Lakes is currently undergoing vertical adjustment after being depressed by the weight of an ice sheet formed in the most recent glacial period during the Wisconsinan. The rate of glacial isostatic adjustment (GIA) varies by location, and strongly influences the flow of water in the Laurentian Great Lakes (LGL) as the inclination of the ground surface changes. Previous attempts to estimate the rate of GIA between sites used geologically recent water gauge data from the past 200 years in order to measure the rate of GIA. In contrast, by inferring GIA from measurements of the water level in the geological record over the past 5000 years, a more accurate estimate of the long term process of GIA can be obtained. These measurements are made by measuring the elevation of a subsurface sedimentary contact relating to past lake levels, which are then age dated with optically stimulated luminescence (OSL) to provide an age for sediments. Elevation and age data are then compiled to create site paleohydrographs for each location around the lake basin.

The focus of this paper is to analyze the data compiled by Johnston et al, 2014 which measured past elevation of shorelines by interpreting water levels recorded in the sediment record. Each site paleohydrograph is now extended between points where the data was measured directly, then subtracted from one data point to a modelled elevation in order to create a plot of relative elevation

over time. Once this is done, the rate of change per unit time is obtained from a linear regression, representing an estimate of the value of GIA between each pair of sites. This process is repeated for all possible combinations of the four sites used, Grand Traverse Bay (GTB), Au Train Bay (ATB), Batchawana Bay (BATB), and Tahquamenon Bay (TAHB).

The results of this process were in strong agreement at the 95 % confidence level for GIA rates obtained from forward and reverse regressions for the combination of ATB-BATB (23.5 to 31 cm/century) and BATB-TAHB (11 to 17 cm/century). Agreement was also seen at the 95 % confidence level for GTB-TAHB (anywhere from -3 to 8.5 cm/century), ATB-GTB (9 to 13 cm/century) and ATB-TAHB (19.5 to 29 cm/century).

2 Introduction

The Earth's crust rests on top of the mantle, its elevation rising and falling with the amount of mass weighing on it. During glacial periods, a significant portion of the water on earth is transferred in form from water in the oceans to glacial ice sheets, weighing down the continental crust and causing the mantle to dynamically adjust with it. This causes the crust to ride relatively lower in elevation, a change which reverses when the weight is removed as the ice sheets melt. This vertical motion of the crust while attempting to return to its previous position is known as glacial isostatic adjustment (GIA) (Scott et al, 2010).

The process of GIA has implications for the routes that the flow of water on the Earth's surface takes; the "tilting" of the surface caused by uneven rates of GIA in different locations (the thickness and residence time of the ice sheet impacts how much the crust subsides and how rapidly it rebounds) may open or close drainage outlets from basins, causing some rivers and lake outlets to go dry, while opening new outlets for water to flow through as the 'tilt' of the lake basin changes and some locations rise and fall in elevation. Additionally, the change in "tilt" has potential to change shorelines of existing basins, which has implications for land usage and long term engineering projections for structures such as locks and dams, where the lifetime of the structure may extend over decades or centuries. In the case of locks providing access to a canal, a local change in water level on the order of centimeters to meters over a century

could cause the structure to become submerged below the rising water level, making the structure obsolete well within its intended usage lifetime.

3 Previous Work

Mainville & Craymer (2005) used water gauge data collected around the LGL over the past 150 years to create monthly means of water level. Differences in these values between sites were then plotted against time to calculate a rate of elevation change between sites over time (This value being interpreted as the current impact of the GIA process on the crust underlying the LGL, although the actual process extends over a much longer timescale). Combinations of sites were shown to produce inconsistent results, so a second method using a least squares adjustment process was used, repeatedly removing some monthly mean outliers which fell some arbitrary residual distance away from the linear regression line until none remained "too far away" from the final regression line. A third, and ultimately optimal method for calculating GIA was developed by Mainville & Craymer in their 2005 paper, using the original method of directly comparing monthly water level means, but this time with adjustments for the epoch, site, and month of the year. Their findings with this method showed a general agreement with the post glacial ICE-3G global model of GIA at that time, while (Mainville & Craymer, 2005).

Johnston et al. (2012) attempted to provide a value for GIA in the LGL with better accuracy than previous estimates made using water gauge data. In order to accomplish this, the data used to measure the process of GIA needed to extend over a much longer timescale. In this method, water levels were inferred

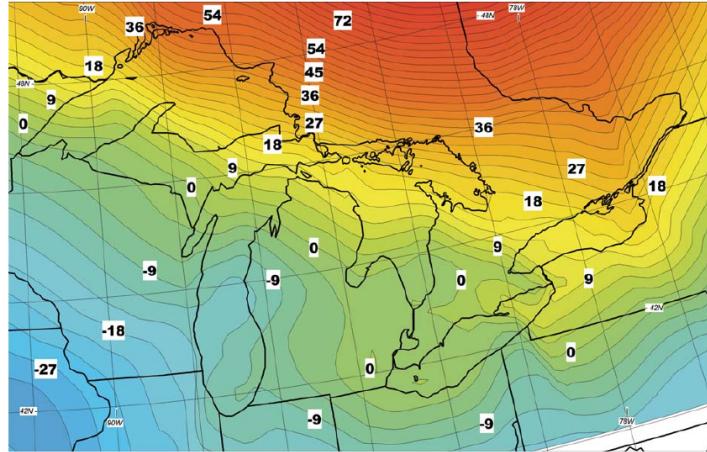


Figure 7. Contour map of vertical velocities derived from water level gauges over the Great Lakes surrounded with ICE-3G-derived velocities. Contour interval—3 cm/century.

Figure 1: GIA values reported by Johnston et al 2012. All values are in cm/century.

from the elevation of relict shorelines in beach ridge strandplains from the late Holocene sediment record surrounding Lake Superior, the ages for each data point being inferred from dating samples from these beach deposits (known as strandplain sequences) with Optically Stimulated Luminescence (OSL) age dating. This data differed from that used by Mainville & Craymer in their 2005 paper, in that data collected did not have elevations sampled at the same points in time for calculation of relative rates. As a result, the elevation vs time data was modelled with a linear regression for each site, the difference in slopes representing the GIA rate between sites. Individual regressions were created per site for a series of four ranges of time related to lake level phases (Johnston et al, 2012). The results reported from this process are summarized in Figure 1.

In order to project the future impact of this process on the Great Lakes Basin, an estimate of the historical rate of GIA is needed. This estimate is obtained by comparing the elevation of the water mark at two different locations around a basin, and observing how this difference changes over time. The elevation of the water can be inferred by a variety of indicators in the sediment record, in this case, beach deposits known as strandplain sequences are used, their ages determined by optically stimulated luminescence (OSL) dating. This raw data is presented in Figure 2.

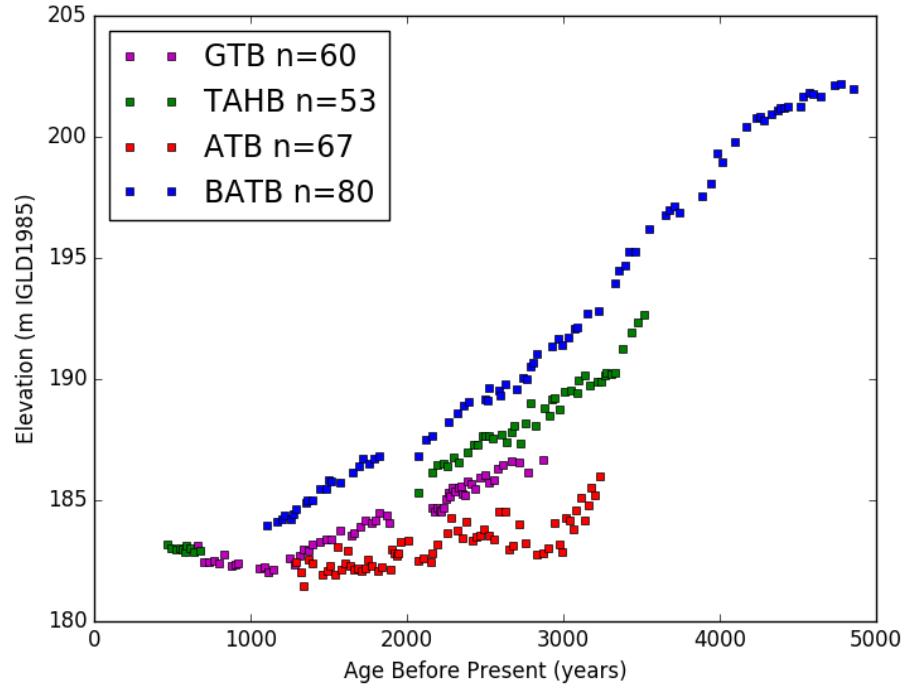


Figure 2: Current day elevation of relict shorelines with respect to time before present over the last 5000 years. Sites ATB, BATB, TAHB, and GTB surrounding Lake Superior are plotted individually.

The data used for this paper was previously published in Johnston et al, 2012, being sampled from four separate locations around Lake Superior: Au Train Bay, Michigan (known in this paper as ATB), Grand Traverse Bay, Michigan (GTB), Batchawana Bay, Ontario (BATB), and Tahquamenon Bay, Michigan (TAHB).

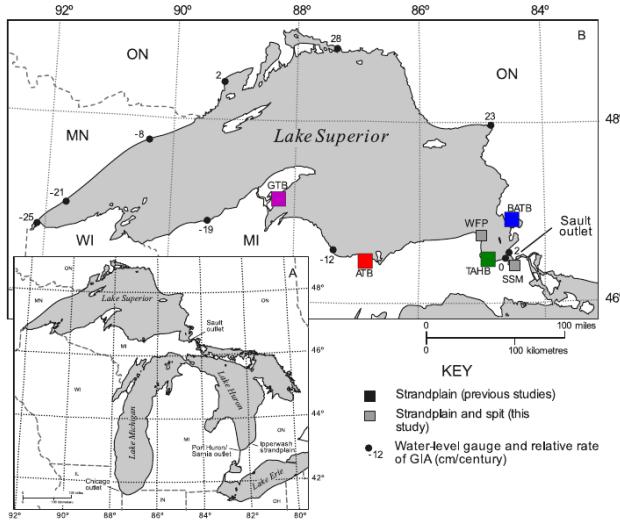


Figure 3: Map of the Upper LGL, showing locations of the modern Sault and Port Huron outlets as well as the ancient Chicago outlet since closed. Strandplains used in this study are marked with the colours blue, green, purple, and red (Note that the colour of each site marker will remain constant throughout the rest of this paper for convenience). Reproduced with modifications from Johnston et al, 2012

Observing Figure 3, it can be seen that all four datasets follow some-

what linear trends, decreasing in elevation as the time before present approaches the current day. This is because the crust underlying the LGL has been rebounding upwards over the past 5000 years, which can be interpreted to imply that areas that were at the elevation of the water surface in the past have been shifted upwards in elevation above the current water surface elevation. The rate of this upward trend varies by site, generally increasing for sites closest to the north and east extremes of the range studied. (this is due to the rate of rebound increasing with closeness to the center of the Laurentide Ice Sheet, roughly near current day James Bay in Northern Ontario).

Between the four study sites, the most common feature is good data coverage between approximately 1000 and 3500 years before present, and a common gap in coverage around 2000 years before present. This was due to conditions which worked against the formation of strandplain sequences during the Algoma lake level fluctuation (Johnston et al, 2014), thus causing most of our datasets to have interrupted records of lake level elevation at this time. While BATB has data coverage extending far beyond this range up to 5000 years before present, the others do not, which makes relative GIA comparisons impossible over that time range. The only dataset which does not follow this pattern of data coverage is TAHB, which will be discussed later in this section.

In order to measure a relative rate of GIA between sites, the rate at which these trends diverge must be measured. In the previous work done on this dataset, this was accomplished by representing the trend of increasing elevation with age as a straight line using a linear regression over the age range of each lake phase

(Johnston et al, 2012). This was an effective first approximation, but failed to take into account that the behaviour of each dataset may not necessarily have been truly linear over each lake phase, being affected by global water level changes caused by climate and other factors.

In order to produce an estimate of GIA which better reflects the rate of GIA from comparisons made between elevations measured at similar times, a better strategy would be to simply subtract the differences in elevation between sites and plot these differences with respect to time, similar to the method used by Mainville & Craymer (2005), using water gauge data. Unfortunately however, none of the datasets have elevations sampled at the same times, an estimate of elevation being needed for times where one dataset has a data point present, but the other does not. The goal of the research presented in this paper was to develop a new method of calculating direct differences in elevation between sites for the strandplain paleohydrographs published in Johnston et al (2012), and compare the results with rates for the process of GIA as calculated in prior studies concerning the LGL.

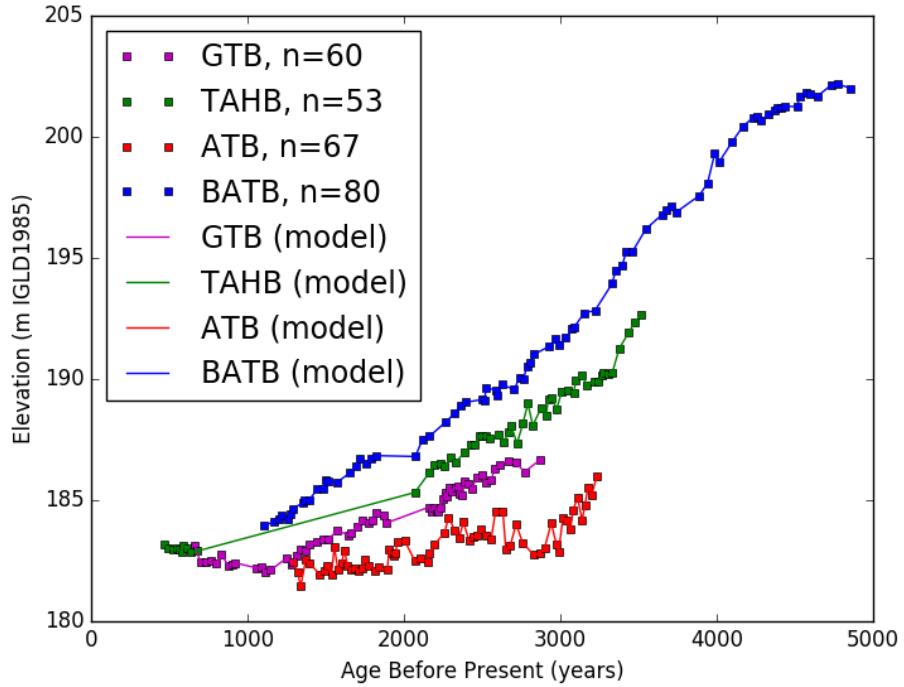


Figure 4: Strandplain paleohydrographs showing current day elevation of relict shorelines with respect to time before present over the last 5000 years. Linear interpolation used between data points, known in this paper as modelled data

4 Methods

In this paper, the modelled elevation for points where elevation was not directly measured is created by using linear interpolation between datapoints, represented as a solid line between points as seen in Figure 4. Once this estimate of elevation for times between sampled datapoints at each site was created, the difference in GIA between sites were calculated by subtracting the elevation of

a measured data point from the modelled elevation of another dataset at that point in time. This difference is shown as a dashed line in Figure 5: To avoid

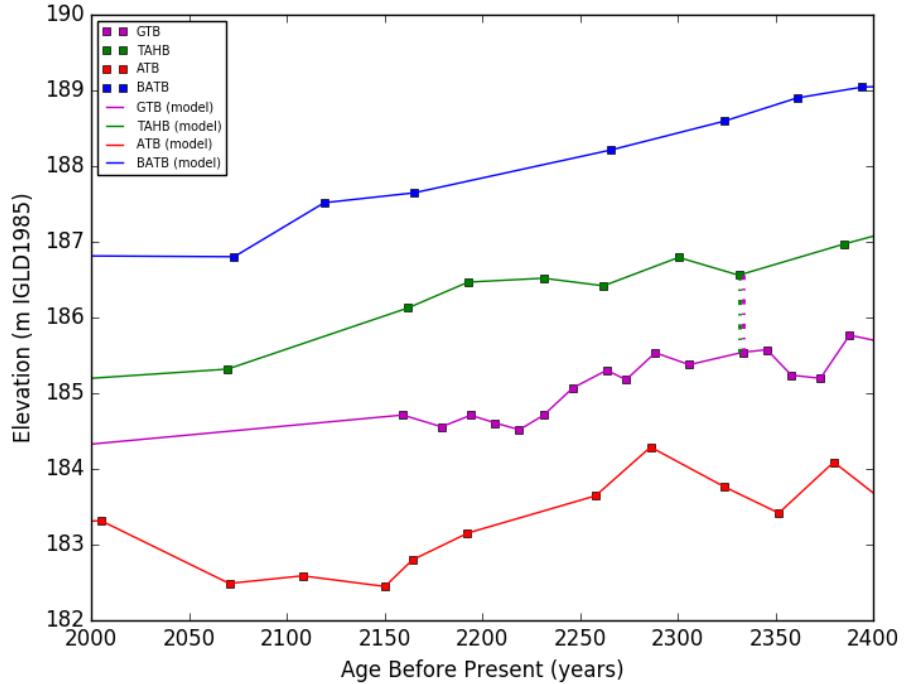


Figure 5: Example GIA comparison between strandplain datasets. Comparison made from a data point to the linear interpolation model, represented by a vertical alternating dashed line

making comparisons between data in one dataset and a model for another in areas where the model stretched over long distances between measured data (such as the 1500 year gap in TAHB), the data for all four sites was grouped into a series of bins starting at 450 years before present with a width of 200 years for each bin. The start point was chosen by taking the youngest age value

recorded in any of the datasets and rounding down to the nearest ten, in this case 450 years before present. The 200 year bin width was chosen as the error on the age models for shorelines published in Johnston et al (2012) had error bounds of approximately 200 years. for the values reported in the age models published by Johnston et al (2012)). If any bin had no data available for one site or the other in a comparison, none of the datapoints in that bin range were used to make comparisons, thus ensuring that areas like the gap in TAHB were not included when making comparisons. In addition, a second rule was created, stipulating that the counts for the bins needed to be within 75% of one another to be considered in calculating relative elevation between sites. This avoided a few areas where the datasets for both bins compared poorly, but produced valid comparison windows (On Figure 4 see the small zone of GTB-TAHB overlap at $\tilde{600}$ years before present).

In order to implement the methods described above, a python script was used to process the data provided by Professor Johnston from his 2012 paper. The source code used can be seen in the appendix of this paper.

Using comparisons from measured data points to linear interpolation model for pairs of study sites, two graphs of the relative difference over time was created for each pair of sites. For example, for the site combination of ATB & BATB, ATB was first compared to BATB, followed by a comparison from BATB to ATB. The rates of GIA produced by these comparisons should be of opposite signs, but similar magnitudes.

The GIA rates are determined by applying a linear regression to each comparison, the slope of each regression representing the relative rate of GIA between sites.

5 Results

5.1 GIA Calculation Results

Listed in this section are the results of each of the possible combinations of sites.

Since 4 sites were used, a total of 6 distinct combinations of sites were studied.

5.1.1 ATB-BATB

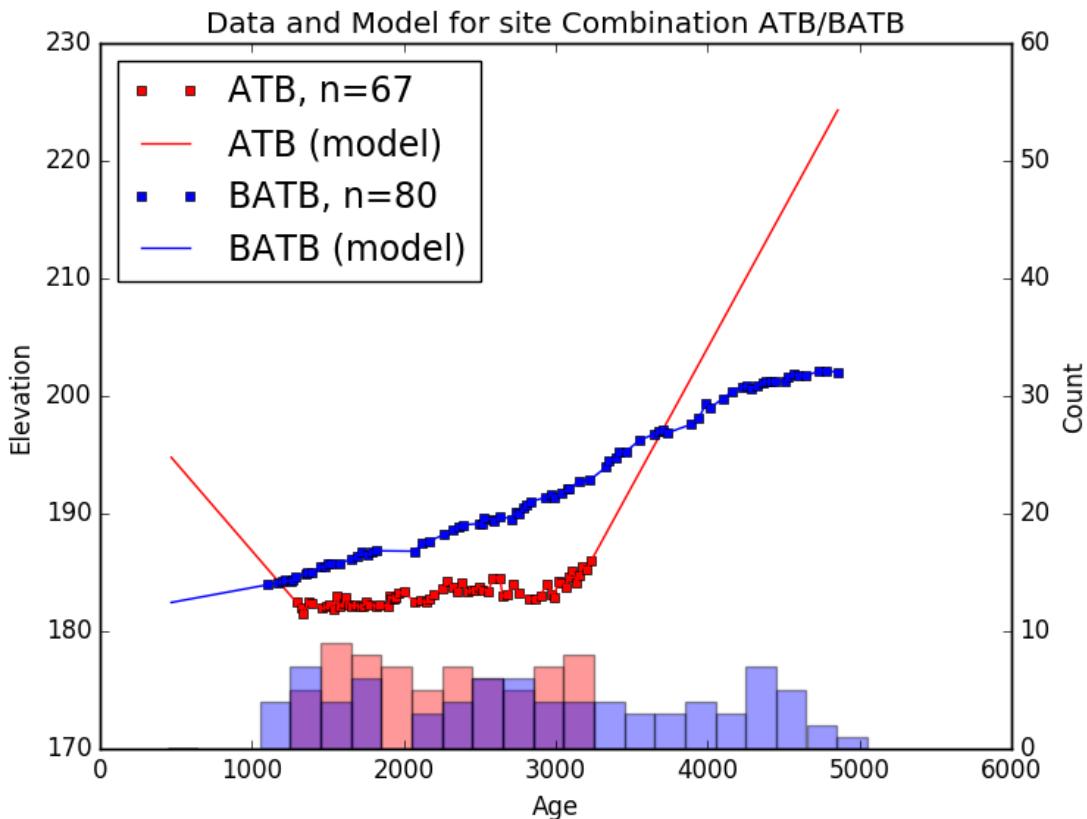


Figure 6: Measured and modelled elevation data plotted against age for sites ATB & BATB. Data grouped into bins with widths of 200 years starting at 450 years before present. Bin Counts shown as a histogram at bottom of graph.

The data available for sites ATB and BATB shows two of the most common trends in the data used in this paper; Data is available for both sites from approximately 1000 to 3300 years before present, with a gap in the record at

around 2000 years before present (caused by a low water level period preferentially not forming beach deposits during this time) (Johnston et al, 2014). With the data divided up into bins of 200 years width starting at 1050 years before present, the data from every bin between 1250 and 3250 was used in calculating a rate of GIA, save for the gap from 1850-2050 years before present caused by the Algoma low water level (where comparisons between data in the ATB dataset would be subtracting against a modelled value for BATB that is unreliable given the distance to the nearest datapoint in BATB). The regressions derived from this pair of data sets, seen in Figures 8 & 9, are well constrained, and produce a moderately well constrained value on relative GIA between ATB and BATB of 23.5-31 cm/century. In order to produce this range, a regression was done on both comparisons, the results of which can be seen in Figure 7. The absolute rate of GIA was determined from the absolute value of the slopes of each regression, reported here as a 95% confidence interval in cm/century. Once these 95% intervals were created, a final value was created combining both sites A complete plot of the confidence intervals for the slopes obtained from each linear regression can be seen in Figure 30.

name	Slope Estimator	Slope Error	r Squared	Slope C.I. (95p)
ATB relative to BATB model	-27.46700	1.32601	0.881	-24.86803 to -30.06597
BATB relative to ATB model	26.83591	1.74490	0.858	30.25591 to 23.41592

Figure 7: ATB-BATB Regression output parameters

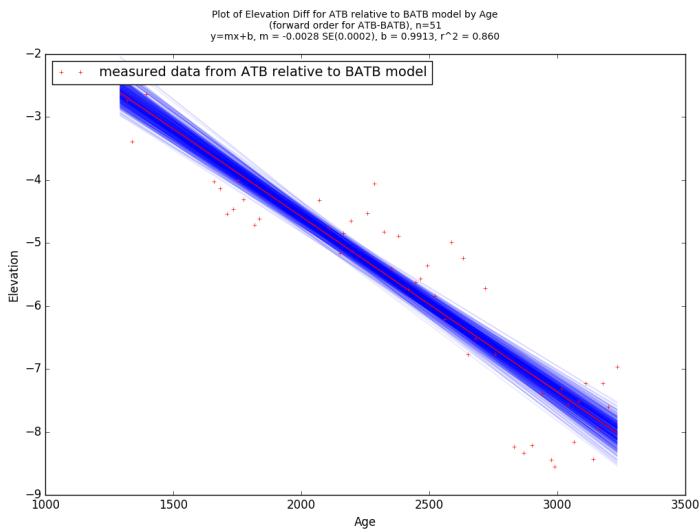


Figure 8: Differences in elevation measured from the ATB data to the ATB model

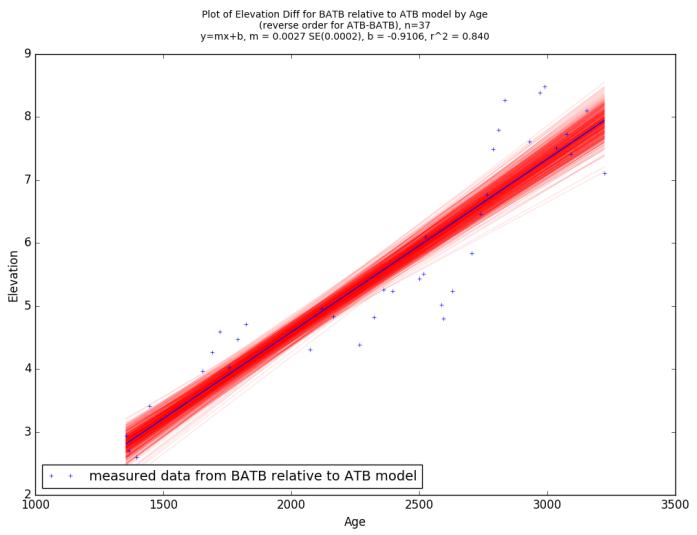


Figure 9: Differences in elevation measured from the BATB data to the ATB model

5.1.2 TAHB-BATB

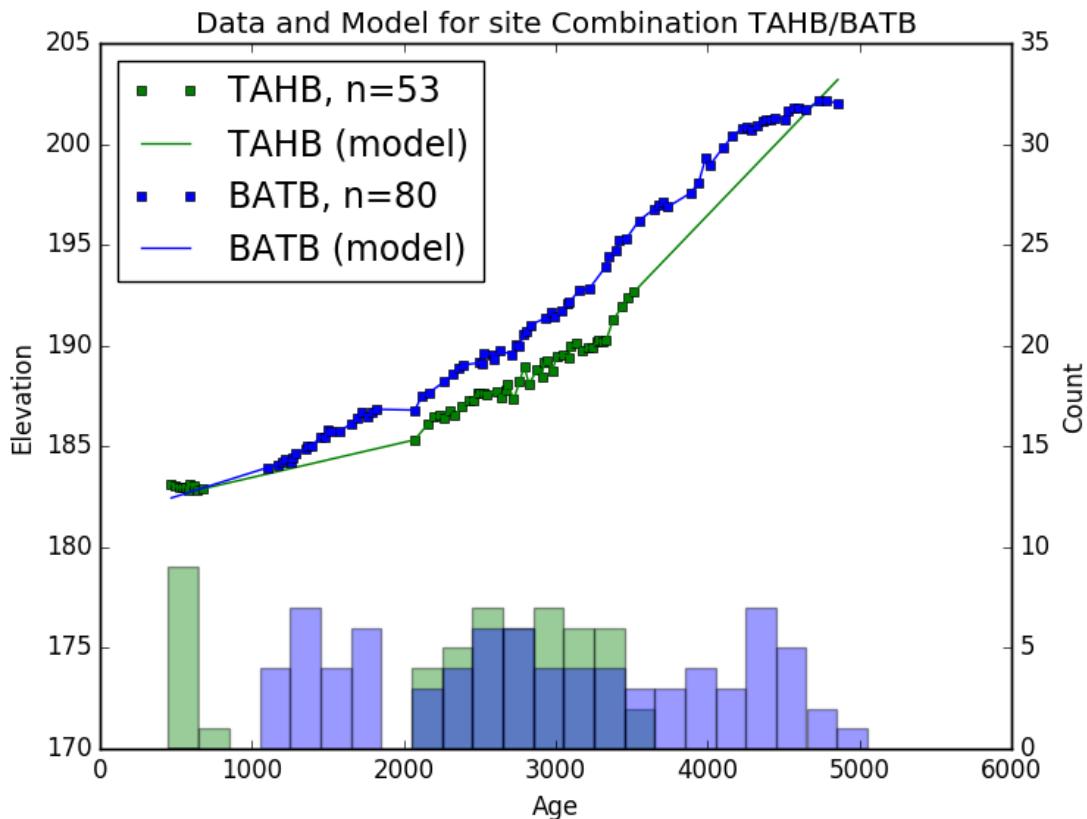


Figure 10: Measured and modelled elevation data plotted against age for sites TAHB & BATB. Data grouped into bins with widths of 200 years starting at 450 years before present. Bin Counts shown as a histogram at bottom of graph.

The data plot for the site combination of TAHB and BATB shows a common issue with comparing datasets, as the data with ages more recent than the 2000 year before present gap is unusable. This is because the regions where data is

available for one dataset are empty of datapoints for the other, making the modelled prediction of the other dataset highly unreliable. As a result, a filter is applied to the data to prevent this, grouping data points into bins 200 years wide, and ignoring the data points from bins in which either data set had no datapoints, as well as any which had bin counts differing by more than 75% for that bin. As a result, only the data from 2050 to 3650 years before present were used in creating the GIA comparisons.

The linear regressions produced from this dataset seen in Figures 12 & 13, are well constrained, and report a value for relative GIA of between 11-16.7 cm/- century.

name	Slope Estimator	Slope Error	r Squared	Slope C.I. (95p)
TAHB relative to BATB model	-14.32814	1.24125	0.765	-11.89530 to -16.76099
BATB relative to TAHB model	14.00018	1.54265	0.733	17.02377 to 10.97660

Figure 11: TAHB-BATB Regression output parameters

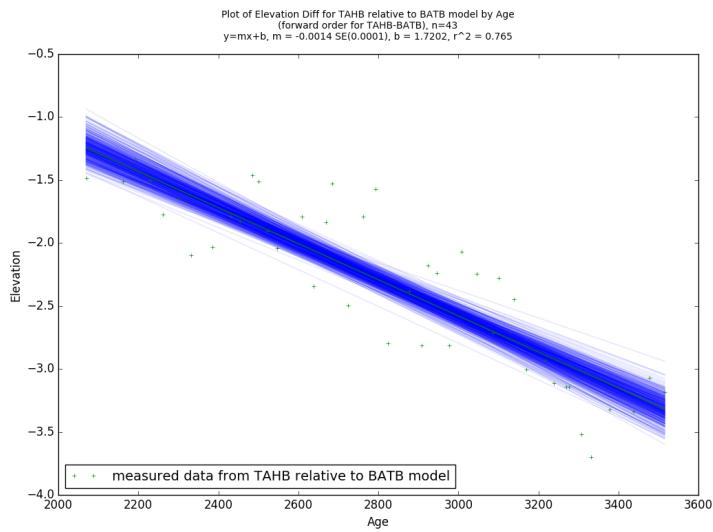


Figure 12: Differences in elevation measured from the TAHB data to the BATB model

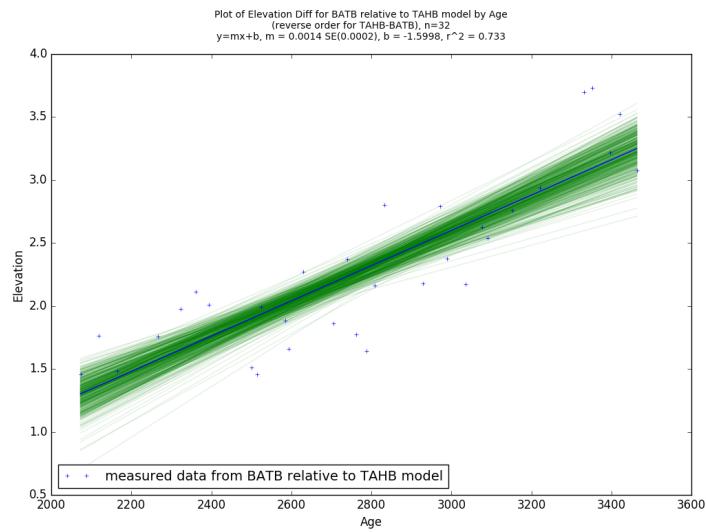


Figure 13: Differences in elevation measured from the BATB data to the TAHB model

5.1.3 TAHB-ATB

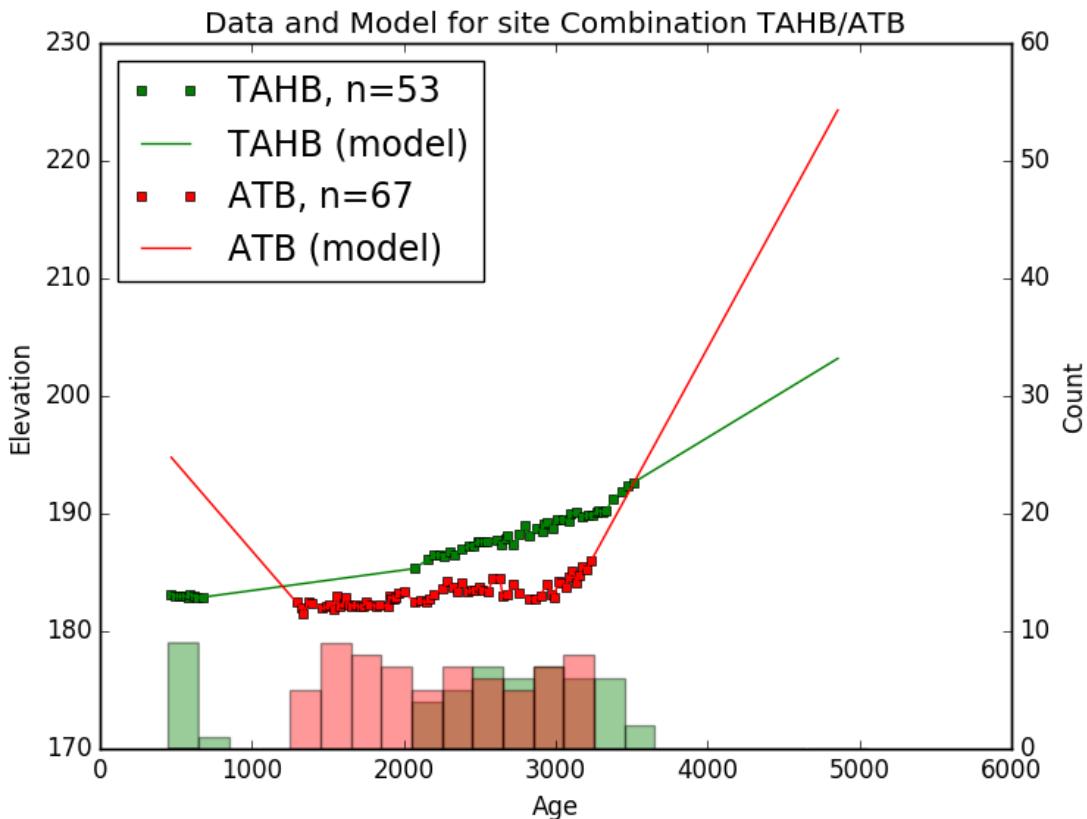


Figure 14: Measured and modelled elevation data plotted against age for sites TAHB & ATB. Data grouped into bins with widths of 200 years starting at 450 years before present. Bin Counts shown as a histogram at bottom of graph.

Similar to previous datasets, the combination of TAHB and ATB are constrained to ages older than 2050 years before present by the Algoma gap, but also have a much shorter range of age values that can be considered for GIA calculation,

ending at around 3100 years before present. This is due to TAHB having no data available between 1250-2050 ybp, while ATB has a great deal of data in this range that can not be considered for this comparison. Using only datapoints between 2050 and 3250 years before present results in relatively poor regressions that are not as well constrained, giving a wide range for relative GIA of between 19.4-29 cm/century.

name	Slope Estimator	Slope Error	r Squared	Slope C.I. (95p)
TAHB relative to ATB model	26.20553	3.45025	0.643	32.96802 to 19.44304
ATB relative to TAHB model	-23.06696	3.14849	0.599	-16.89592 to -29.23801

Figure 15: TAHB-ATB Regression output parameters

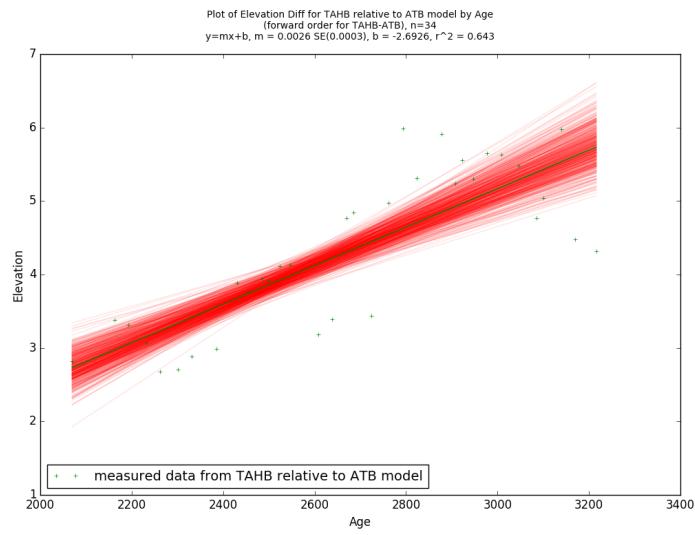


Figure 16: Differences in elevation measured from the TAHB data to the ATB model

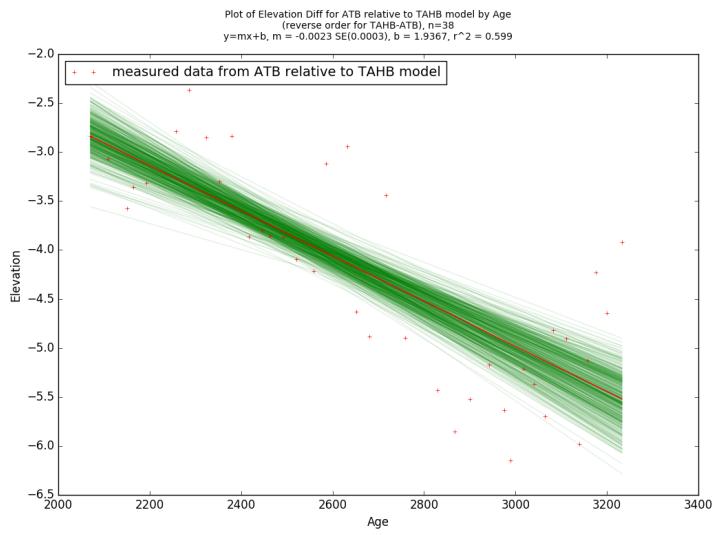


Figure 17: Differences in elevation measured from the ATB data to the TAHB model

5.1.4 GTB-BATB

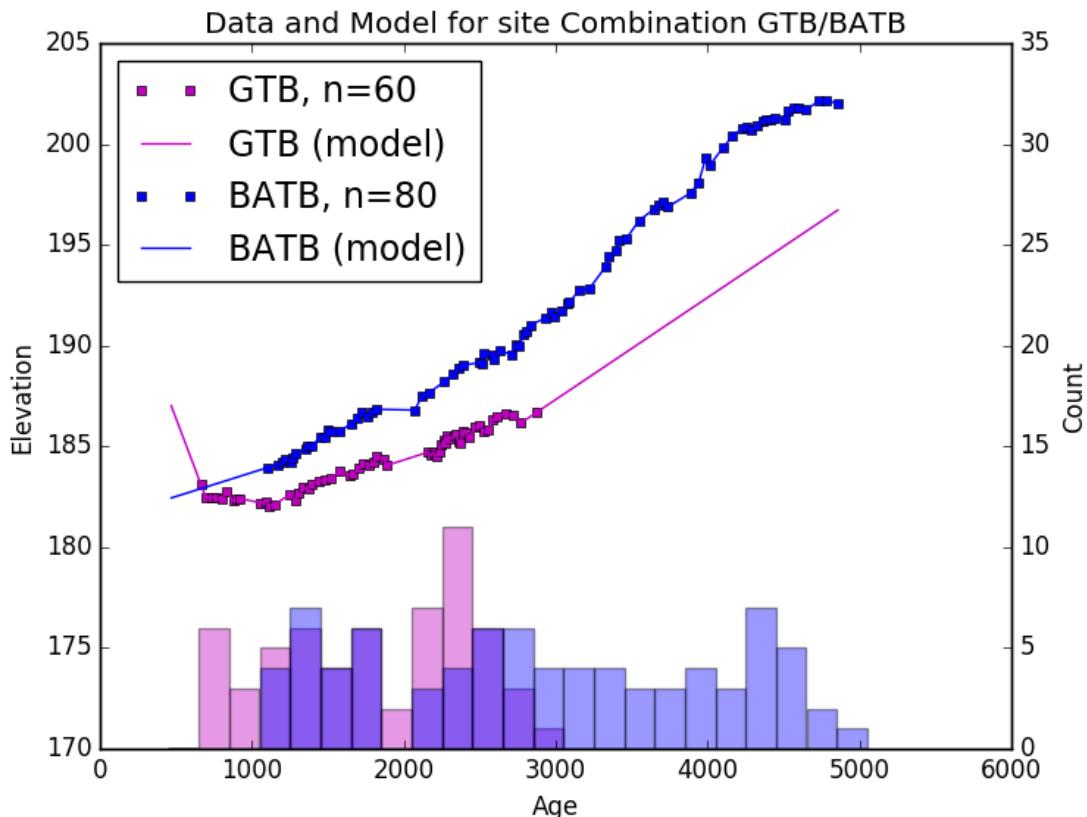


Figure 18: Measured and modelled elevation data plotted against age for sites GTB & BATB. Data grouped into bins with widths of 200 years starting at 450 years before present. Bin Counts shown as a histogram at bottom of graph.

The GTB BATB combination has a data spread similar to that of ATB & BATB, with data available for both datasets from 1050 to 3050 years before present with an Algoma related gap from 1850 to 2050 years before present.

The first case of the 75% difference cutoff has its first appearance here as the oldest shoreline available from GTB just falls inside of the 2850-3000 ybp window, causing the entire window to be used if the only criteria was both dataset counts within that window being non-zero. The 75% cutoff prevents this window from being used in this case, as the counts for the 2850-3000 ybp window differ by 120%. This rule is useful in identifying areas of the dataset where both sites have data available, but the density of one of the datasets in that region is low enough to cause issues with the models ability to make accurate predictions in between measured datapoints. The regressions in Figures 20 & 21 bear this out, producing one of the better constrained values at 10.4-12 cm/century.

name	Slope Estimator	Slope Error	r Squared	Slope C.I. (95p)
GTB relative to BATB model	-11.94343	0.77163	0.842	-10.43103 to -13.45582
BATB relative to GTB model	12.15145	0.82899	0.850	13.77627 to 10.52663

Figure 19: GTB-BATB Regression output parameters

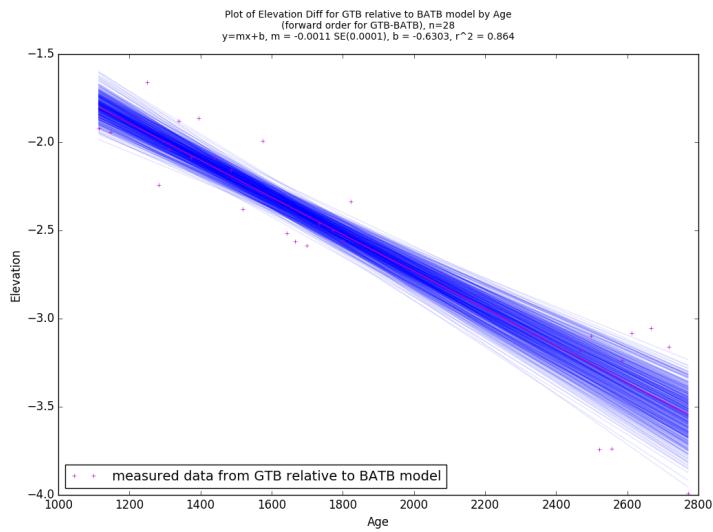


Figure 20: Differences in elevation measured from the GTB data to the BATB model

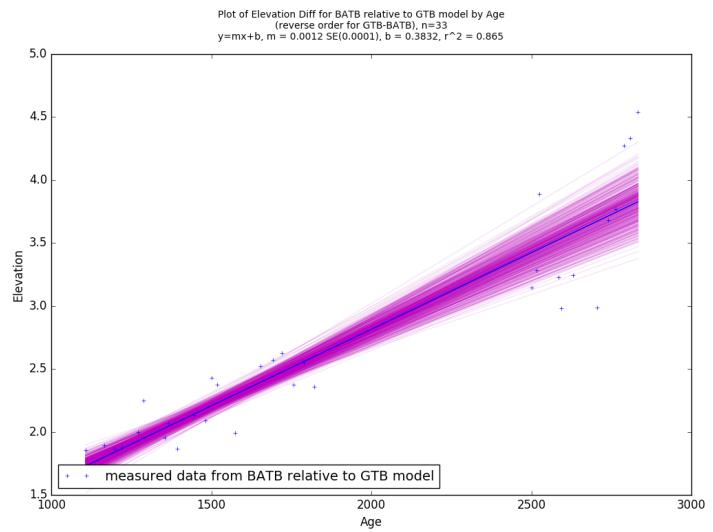


Figure 21: Differences in elevation measured from the BATB data to the GTB model

5.1.5 GTB-ATB

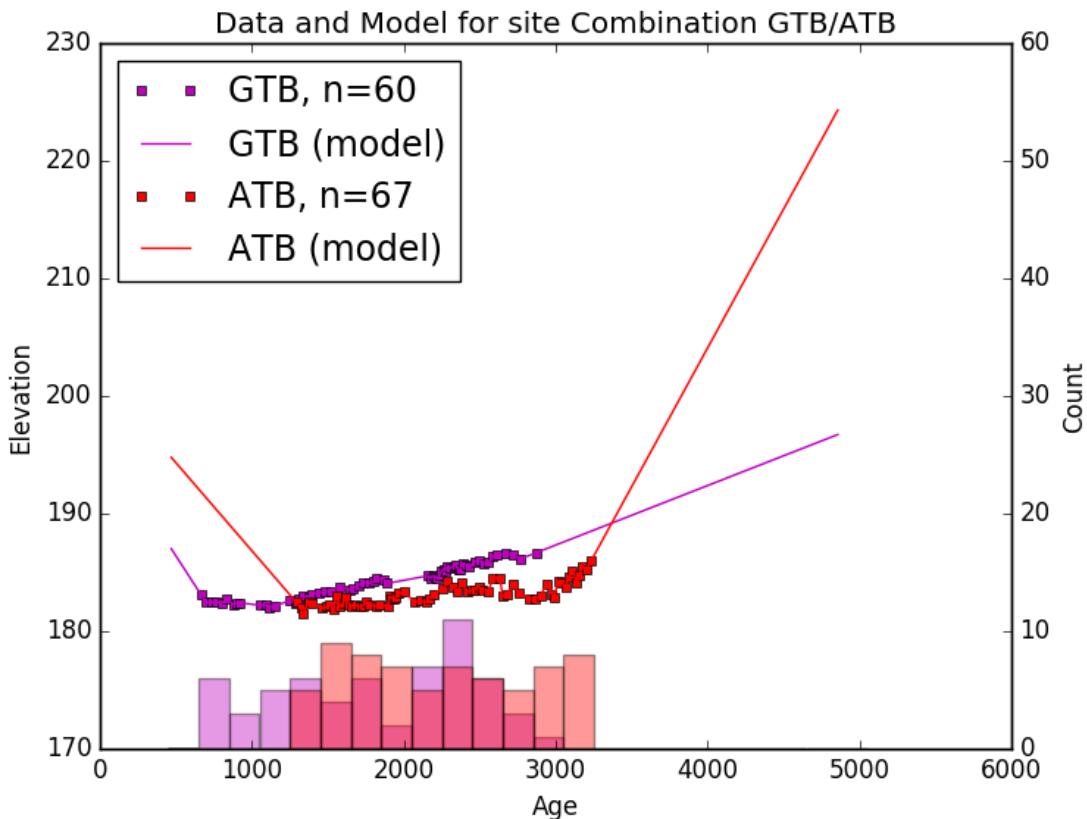


Figure 22: Measured and modelled elevation data plotted against age for sites GTB & ATB. Data grouped into bins with widths of 200 years starting at 450 years before present. Bin Counts shown as a histogram at bottom of graph.

The GTB ATB combination is similar to most of the combinations looked at so far, windows from 1250 to 3050 ybp containing data for both sites. Two of these windows fail to qualify for use under the filter due to the site counts differing by

more than 75%, one from 1850-2050 ybp, and a second one from 2850 to 3050 ybp. Looking at the graph, it can be seen that both of these windows coincide with ranges of time where GTB has sparse data, making the GTB models predictions unreliable. Possibly due to this, the regressions in Figures 24 & 25 are not the best constrained constrained, giving a rate of GIA of 9-13.4 cm/century.

name	Slope Estimator	Slope Error	r Squared	Slope C.I. (95p)
GTB relative to ATB model	10.07182	1.77919	0.427	13.55902 to 6.58461
ATB relative to GTB model	-13.58889	1.56930	0.595	-10.51306 to -16.66471

Figure 23: GTB-ATB Regression output parameters

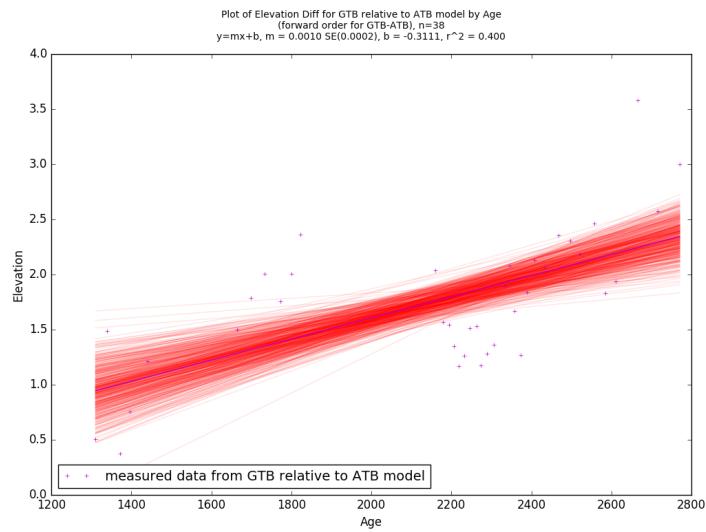


Figure 24: Differences in elevation measured from the GTB data to the ATB model

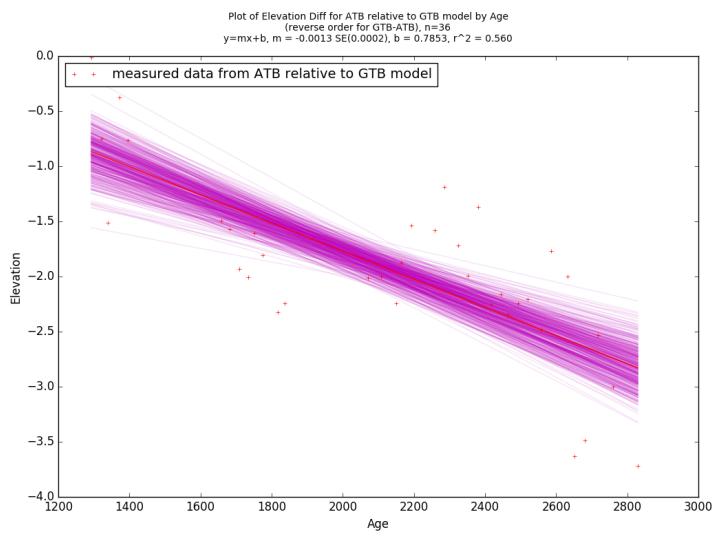


Figure 25: Differences in elevation measured from the ATB data to the GTB model

5.1.6 GTB-TAHB

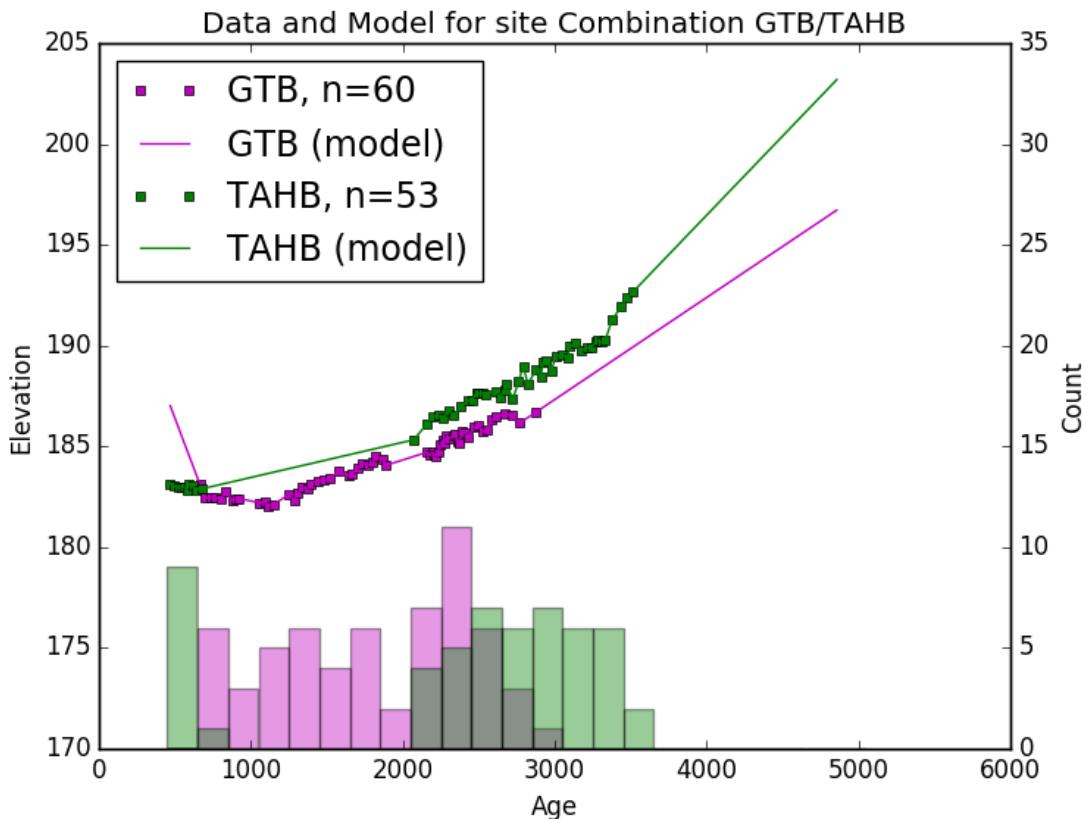


Figure 26: Measured and modelled elevation data plotted against age for sites GTB & TAHB. Data grouped into bins with widths of 200 years starting at 450 years before present. Bin Counts shown as a histogram at bottom of graph.

The final pair of sites GTB and TAHB has by far the most poorly constrained set of regressions, likely due to the alignment of most of both datasets only giving sample sizes of $n=22$ (Figure 29) and $n=27$ (Figure 28). This was due to

the valid range of data extending only from 2050 to 2850 ybp. Two potential windows at 650-850 ybp and 2850-3050 ybp were thrown out due to not meeting the 75% rule, as both would have produced comparisons between areas of data in one dataset and a poorly constrained model in the other. This resulted in an estimate of GIA that ranges anywhere from -2.8-8.6 cm/century, possibly implying that there may be no difference in vertical adjustment rates between the TAHB and GTB sites.

name	Slope Estimator	Slope Error	r Squared	Slope C.I. (95p)
GTB relative to TAHB model	-6.33388	0.90547	0.605	-4.55916 to -8.10860
TAHB relative to GTB model	7.56282	2.10691	0.380	11.69236 to 3.43327

Figure 27: GTB-TAHB Regression output parameters

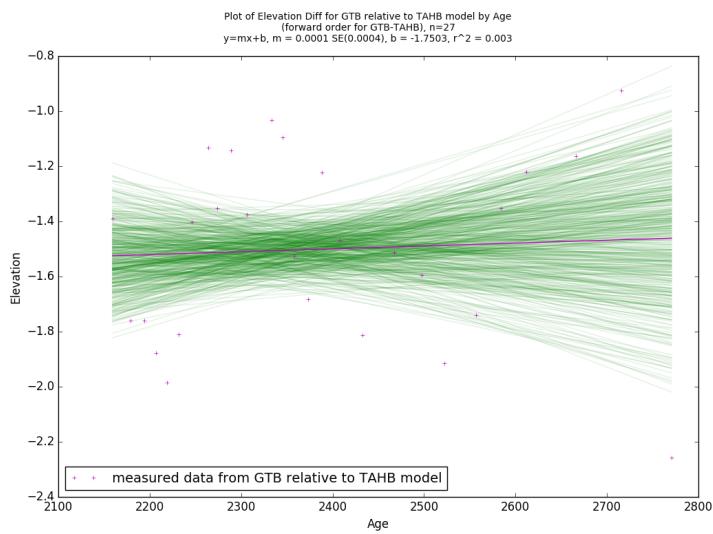


Figure 28: Differences in elevation measured from the GTB data to the TAHB model

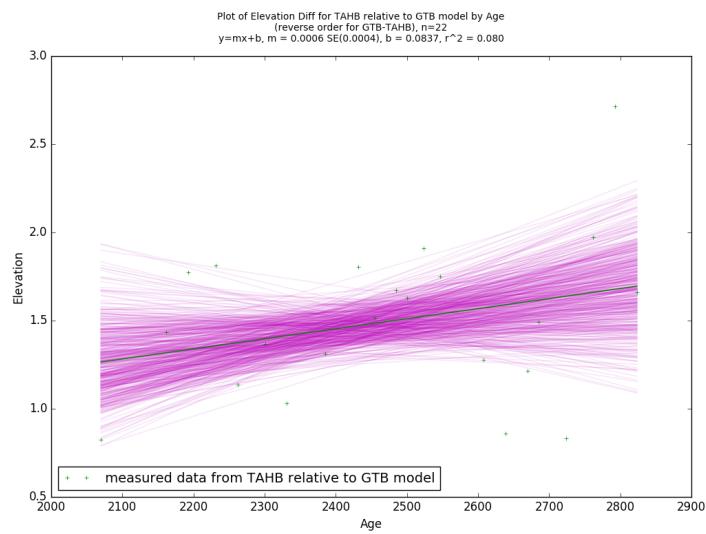


Figure 29: Differences in elevation measured from the TAHB data to the GTB model

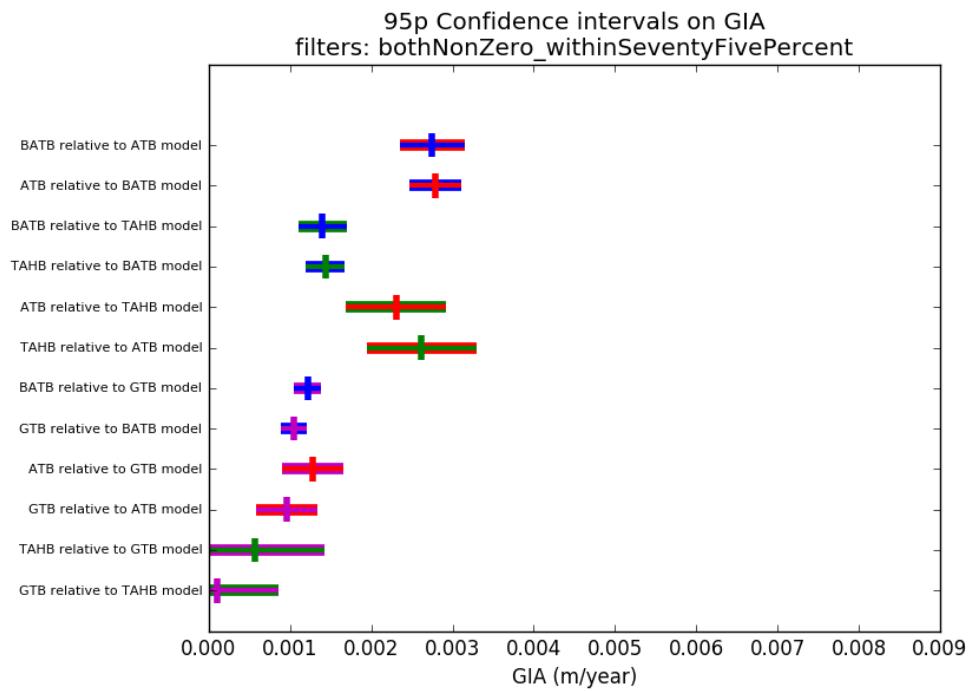


Figure 30: 95p Confidence intervals on GIA rates obtained from site comparisons

In the following section, the values for relative GIA produced by this paper are contrasted with those previously obtained by Mainville & Craymer by plotting the difference between each site as a line between sites with the corresponding value next to it on a map.

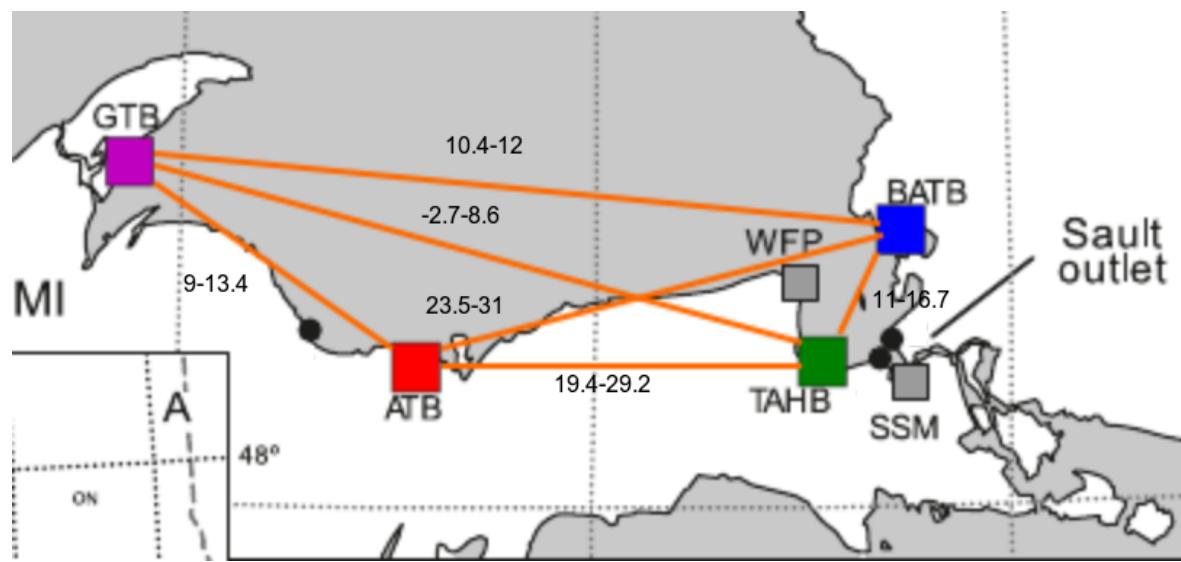


Figure 31: Relative GIA Rates produced by this papers method, all values reported in cm/century

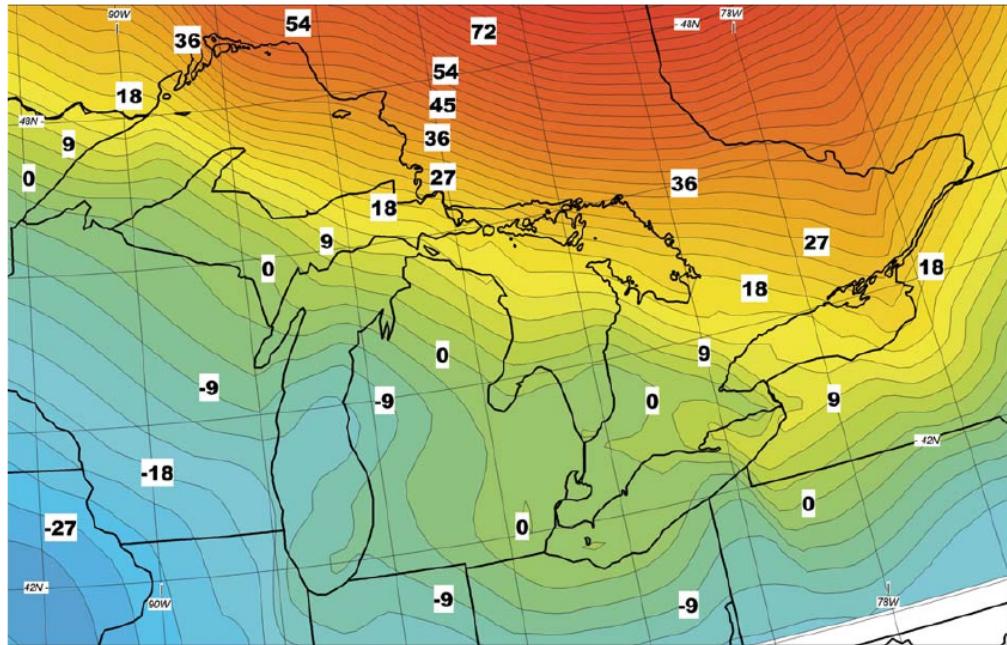


Figure 7. Contour map of vertical velocities derived from water level gauges over the Great Lakes surrounded with ICE-3G-derived velocities. Contour interval—3 cm/century.

Figure 32: Relative GIA Rates produced by Mainville & Craymer, all values reported in cm/century (reproduced from Mainville & Craymer, 2005)

The equivalent values for rates between sites as produced by Mainville & Craymer are inferred from subtracting the difference in contour between sites as shown in Figure 32, and are presented in Figure 33.

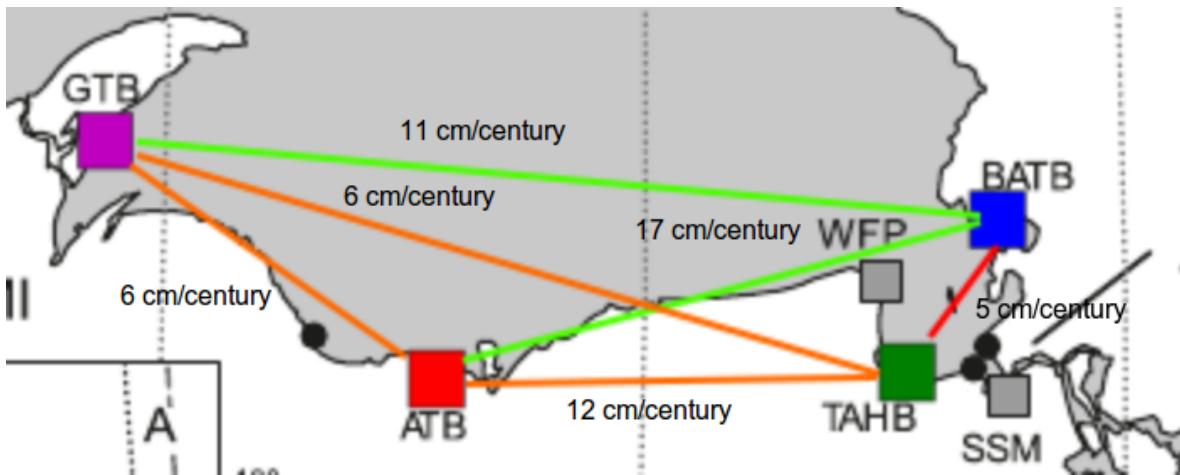


Figure 33: Relative GIA Rates produced by Mainville & Craymer

While most of the site comparisons agree reasonably well between the method employed by Mainville & Craymer and this paper, one area where significant disagreement is seen is between sites ATB, BATB, and TAHB, especially in the much larger values produced by this paper between ATB-BATB and ATB-TAHB. Given that both of these site combinations are separated by an East-West line, this could imply that the location of the center of the Laurentide Ice Sheet during the last glaciation being to the north and west of Lake Superior had a stronger effect on the overall process of rebound than the simple fact that areas to the north were more likely to be depressed by the weight of ice sheets than areas further south.

6 References

- Mainville, A., & Craymer, M. R. (2005). Present-day tilting of the Great Lakes region based on water level gauges. *Geological Society of America Bulletin*, 117(7-8), 1070-1080.
- Scott, T. W., Swift, D. J., Whittecar, G. R., & Brook, G. A. (2010). Glacioisostatic influences on Virginia's late Pleistocene coastal plain deposits. *Geomorphology*, 116(1-2), 175-188.
- Johnston, J. W., Argyilan, E. P., Thompson, T. A., Baedke, S. J., Lepper, K., Wilcox, D. A., & Forman, S. L. (2012). A Sault-outlet-referenced mid-to late-Holocene paleohydrograph for Lake Superior constructed from strandplains of beach ridges. *Canadian Journal of Earth Sciences*, 49(11), 1263-1279.
- Johnston, J. W., Thompson, T. A., & Wilcox, D. A. (2014). Palaeohydrographic reconstructions from strandplains of beach ridges in the Laurentian Great Lakes. Geological Society, London, Special Publications, 388(1), 213-228.

7 Appendix

7.1 Source code for `giaModel.py`

```
## giaModel.py #####  
## attempt to model the gia between sites using the data #####  
## in reformattedData.ods #####  
#####  
#####  
import pyexcel_ods  
  
##import sys  
  
  
import csv  
  
  
import matplotlib.pyplot as plt  
import numpy as np  
from scipy import stats  
  
  
from matplotlib.font_manager import FontProperties  
  
  
import itertools  
## used to generate the number of links between sites  
import random  
from random import sample  
## used in the random choice feature of the zoomed site comparisons  
  
  
from rawhide import bootstrapper  
## get the custom bootstrap plotting function from this projects code
```

```
from linearInterpolationModel import *
from giaUtils import *

fontP = FontProperties()
fontP.set_size('small')

def trendline(x, gradient, intercept):
    ## return a y given an mx+b
    output = gradient*x + intercept
    return output

## getLinearModel: listof(Num) listof(Num) -> listof(Num) Num Num Num listof(Num)
listof(Num)

def getLinearModel(x_values, y_values, k=1.0, l=1.0):
    gradient, intercept, r_value, p_value, std_err = stats.linregress(x_values,
        y_values)

    y_model = []
    yModelHigh = []
    yModelLow = []

    grad = k*gradient
    interc = l*intercept
```

```
for x in x_values:

    y = trendline(x, grad, interc)

    yHigh = trendline(x, grad+(1.96*std_err), interc)

    yLow = trendline(x, grad-(1.96*std_err), interc)

    y_model.append(y)

    yModelHigh.append(yHigh)

    yModelLow.append(yLow)

rSquare = r_value**2

return y_model, grad, interc, std_err, yModelHigh, yModelLow, rSquare

## yModelHigh and yModelLow are the y model built with a slope at the
## extreme of the error bounds on the gradient
```

```
def plotGradientConfidenceIntervals(giaRegressionsByCombo, keys,
                                     giaRegressionDescriptions, outputPathDict):
    def plotInterval(ax, y, xstart, xstop, intervalLabel, colord, colord):
        """Plot interval at y from xstart to xstop with given color."""

        ax.hlines(y, xstart, xstop, colord, lw=7)
        ax.hlines(y, xstart, xstop, colord, lw=3, label=intervalLabel)
        ## plots the interval in the colours of both sites

    outputPath = convertListToRelativePath([outputPathDict[setting] for setting in
                                             getCurrentSettingOptions()])

    y = 0
    ## used in spacing out the intervals for each site vertically through the
    ## graph

    fig,ax = plt.subplots(1)

    for combo in keys:
        y += 1
        combo1 = combo.split('->')[0]
        combo2 = combo.split('->')[1].split(':')[0]
        order = combo.split('->')[1].split(':')[1]
```

```
if(order == 'forward'):

    direct = combo1

    modelled = combo2

else:

    direct = combo2

    modelled = combo1

est = giaRegressionsByCombo[combo]['gradientEstimator']

ciStart = giaRegressionsByCombo[combo]['gradient'][0]

ciEnd = giaRegressionsByCombo[combo]['gradient'][1]

if(est < 0):

    est = -est

    ciStart = -ciStart

    ciEnd = -ciEnd

if(order == 'forward'):

    plotInterval(ax, y, ciStart, ciEnd, "", mapSiteToColour(direct),
                 mapSiteToColour(modelled))

else:

    plotInterval(ax, y, ciStart, ciEnd, "", mapSiteToColour(direct),
                 mapSiteToColour(modelled))

ax.vlines(est, y+0.3, y-0.3, mapSiteToColour(direct), lw=4)

ax.set_xlabel('GIAU(m/year)')

ax.set_xlim([0,0.009])
```

```
plt.yticks(list(np.arange(1, len(keys)+1, 1.0)), [giaRegressionDescriptions[
    key] for key in keys], rotation=0)

fileNameIdentifier = "_" .join([outputPathDict[setting] for setting in
    getCurrentSettingOptions()])

plt.title("95p Confidence intervals on GIA\nfilters: %s" % fileNameIdentifier)

for item in ax.get_yticklabels():
    item.set_fontsize(8)

outputFilePath = filePathOnRelativePath(outputPath+"gias/", fileName='
    intervals', ext="png")
print "Saving_gia_intervals_plot_at_%s" % outputPath
verifyPath(outputPath+"gias/")

plt.savefig(outputFilePath,bbox_inches='tight')

outputFilePath = filePathOnRelativePath(outputPath+"gias/", fileName='%
    s_intervals' % fileNameIdentifier, ext="png")
print "Saving_gia_intervals_plot_at_%s" % outputPath
```

```
plt.savefig(outputFilePath,bbox_inches='tight')

plt.close()

def getDatasetsModelsAndObjects(filenameToLoad):
    lookupTable = pyexcel_ods.get_data(filenameToLoad)
    ## open up the excel file to get the data as a dict of 2-lists
    locations = ['BATB', 'TAHB', 'GTB', 'ATB']
    ## the first key for the lookupTable is the site location

    datasets = {}

    for loc in locations:
        datasets[loc] = [row for row in lookupTable[loc]]
        ## under each key is a rectangular list with two columns to each row,
        ## the first one is elevation, the second one is age

    for d in datasets:
        print d, datasets[d], "\n\n\n"

datasetObjects = {}

datasetModels = {}
```

```
for d in datasets:

    datasetObjects[d] = siteData(d, datasets[d])

    ## build the dataset containers using the data retrieved for each site


    ## note that the siteData object automatically filters the data received
    ## to get rid of the first few non data lines and any empty spaces


return datasets, datasetModels, datasetObjects


if(__name__ == "__main__"):

    datasets, datasetModels, datasetObjects = getDatasetsModelsAndObjects("./"
        reformattedData.ods")

    allAgesSampled = [datasetObjects[d].getAgeValues() for d in datasets]
    allAgesSampled = [item for sublist in allAgesSampled for item in sublist]
    ## flatten out the 2-list with some list comprehension
    print min(allAgesSampled), max(allAgesSampled)

    ## create the raw plot of data points #####
    for site in datasetObjects:
        print site, datasetObjects[site].data
        x = datasetObjects[site].getAgeValues()
        y = datasetObjects[site].getElevationValues()
```

```
n = len(datasetObjects[site].getAgeValues())

plt.plot(x, y, mapSiteToColour(site) + 's', label=site+"n=%i" % n,
          markersize=4.0)

datasetModels[site] = siteModelConnectTheDots(datasetObjects[site])

##plt.title("Plot of Elevation by Age\nRaw Data only")
plt.ylabel('Elevation_(m_IGLD1985)')
plt.xlabel('Age_Before_Present_(years)')
plt.legend(loc=2, prop={'size': 17})
plt.savefig('./theDataRaw.png')
plt.close()

#####
##### create the raw plot with the model included #####
#####

for ds in datasetObjects:
    print ds, datasetObjects[ds].data

    x = datasetObjects[ds].getAgeValues()
    y = datasetObjects[ds].getElevationValues()
    plt.plot(x, y, mapSiteToColour(ds) + 's', label="%s,n=%i" % (ds, len(x)),
              markersize=4.0)

    for d in datasets:
```

```

plt.plot([age for age in sorted(allAgesSampled) if datasetModels[d] .
          ageValueIsInRangeCoveredByModel(age)], [datasetModels[d] .
          getModelledElevation(age) for age in sorted(allAgesSampled) if
          datasetModels[d].ageValueIsInRangeCoveredByModel(age)], mapSiteToColour
          (d), label=d+"_(model)")

##plt.title("Plot of Elevation by Age\nRaw Data with Model")
plt.ylabel('Elevation_(m_IGLD1985)')
plt.xlabel('Age_Before_Present_(years)')
plt.legend(loc=2, prop={'size': 17})
plt.savefig('./theData.png')
plt.close()

#####
#####

zoomXRange = (2000, 2400)
zoomYRange = (182, 190)

for site in datasetObjects:
    print site, datasetObjects[site].data
    x = datasetObjects[site].getAgeValues()
    y = datasetObjects[site].getElevationValues()
    plt.plot(x, y, mapSiteToColour(site) + 's', label="%s" % (site), markersize
              =4.0)

```

```
for site in datasets:

    plt.plot(sorted(allAgesSampled), [datasetModels[site].getModelledElevation(
        age) for age in sorted(allAgesSampled)], mapSiteToColour(site), label=
        site+"_(model)")

    ## plot the dataset models as straight lines

siteCodeOptions = [site for site in datasetObjects]

exampleSites = sample(siteCodeOptions, 2)

print exampleSites

for order in ['forward', 'reverse']:

    if(order == 'forward'):

        direct = exampleSites[0]

        modelled = exampleSites[1]

    elif(order == 'reverse'):

        direct = exampleSites[1]

        modelled = exampleSites[0]

    agesToConsider = [age for age in sorted(allAgesSampled) if ( ((age >= min(
        zoomXRange))and(age <= max(zoomXRange))) and (datasetModels[direct] .
        ageValueInRawData(age) and datasetModels[modelled] .
        ageValueIsInRangeCoveredByModel(age)) )

    for age in agesToConsider:

        print age

demoComparisonPoint = random.choice(agesToConsider)
```

```

print "-> ", demoComparisonPoint

directElevation = datasetObjects[direct].getElevationByGivenAge(
    demoComparisonPoint)

modelledElevation = datasetModels[modelled].getModelledElevation(
    demoComparisonPoint)

print "Direct[%s]:" % direct, directElevation
print "Modelled[%s]: " % modelled, modelledElevation

##plt.plot([demoComparisonPoint, demoComparisonPoint], [directElevation,
    modelledElevation], "%s" % mapSiteToColour(direct), linewidth=3.0)
plt.plot([demoComparisonPoint, demoComparisonPoint], [directElevation,
    modelledElevation], "%s-." % mapSiteToColour(direct), linewidth=2.0)

##'--'
##plt.title("Plot of Elevation by Age\nRaw Data with Model")
plt.ylabel('Elevation_(m_IGLD1985)')
plt.xlabel('Age_Before_Present(years)')
plt.axis((zoomXRange[0], zoomXRange[1],zoomYRange[0],zoomYRange[1]))
plt.legend(loc=2, prop={'size': 7})
plt.savefig('./theDataZoomed.png')
plt.close()

#####
#####
```

create a list of the shortform name of all the sites, then use

```
## itertools to make a list of the sites!, all possible combinations of #####
## sites ##### ##### ##### ##### ##### ##### ##### ##### ##### ##### ##### #####
## ie [A,B,C] -> [[A,B], [B,C], [C, A]]  
sites = [ds for ds in datasetObjects]  
siteCombinations = list(itertools.combinations(sites, 2))  
##### ##### ##### ##### ##### ##### ##### ##### ##### ##### ##### ##### ##### #####  
  
for combo in siteCombinations:  
    print combo  
    for i in range(len(combo)):  
        print i, combo[i]  
  
globalHistogramFloor = None  
## ayy lmao  
  
histogramFloorsList = []  
  
histogramFloorsByCombo = {}  
  
totalAges = []
```

```
#####
## loop through the site combinations and use the data to decide on a #####
## floor for the age bins and the bounds on the plot axes #####
for combo in siteCombinations:

    histogramFloor = None

    ageFloor = None

    for site in combo:

        x = datasetObjects[site].getAgeValues()

        totalAges += x

        y = datasetObjects[site].getElevationValues()

        if(histogramFloor == None):

            histogramFloor = min(y)

        else:

            histogramFloor = min([min(y), histogramFloor])



        if(ageFloor == None):

            ageFloor = min(x)

        else:

            ageFloor = min([min(x), ageFloor])


def roundFloatDownToNearestTen(someFloat):

    someFloat /= 10

    someFloat = int(someFloat)

    someFloat *= 10

    someFloat -= 10
```

```
        return someFloat

    histogramFloor = roundFloatDownToNearestTen(histogramFloor)
    ageFloor = roundFloatDownToNearestTen(ageFloor)
    histogramFloorsByCombo[combo] = histogramFloor

    histogramFloorsList.append(ageFloor)
    print "histogramFloor\u00d7for\u00d7site\u00d7combo", combo, ":\u00d7", histogramFloor
#####
#globalHistogramFloor = min(histogramFloorsList)

print "global\u00d7bin\u00d7floor\u00d7set\u00d7at\u00d7", globalHistogramFloor

globalBins=range(globalHistogramFloor, int(max(totalAges))+200, 200)
## build a list of bin endpoints starting at the floor value and ending at
## one bin width above the last age value of any of the dataset

## example of how this works if you run
## range(450, 4857+200, 200)
print "global\u00d7bins:\u00d7", globalBins

#####
## for debug output print out bin counts for each dataset #####
for i in globalBins:
    print "bin\u00d7",i, ", ", i+200, ":"
    print "-"*80
    for combo in siteCombinations:
```

```
for site in combo:

    thisSiteDataset = datasetObjects[site]

    siteName = '{:4s}'.format(site)

    print "site: %s count: %i" % (siteName, thisSiteDataset.

        getCount(i, 200))

    print "-"*80

    print "\n\n"

#####
#####

giaRegressions = {}

giaRegressionComboMappingsByConditions = {}


giaRegressionKeys = []
giaRegressionDescriptions = {}
giaKeysByDescriptions = {}


for combo in siteCombinations:

    for order in ['forward', 'reverse']:

        if(order == 'forward'):

            direct = combo[0]

            ## d is the site we are using as our direct comparison

            ## ie MUST have a measured data point at this age

            modelled = combo[1]

            ## ds is what we are comparing against, so it can just be a
```

```
## modelled point

else:
    direct = combo[1]
    modelled = combo[0]

thisRegressionKey = "%s-%s:%s" % (combo[0], combo[1], order)

giaRegressionKeys.append(thisRegressionKey)
thisComparisonGiaDescription = "%s\u2014relative\u2014to\u2014%s\u2014model" % (direct,
    modelled)
giaRegressionDescriptions[thisRegressionKey] =
    thisComparisonGiaDescription
giaKeysByDescriptions[thisComparisonGiaDescription] = thisRegressionKey

sortedKeys = sorted(giaRegressionKeys)
print "sortedKeys:", sortedKeys

#####
## plot the raw data plots with counts for each bin #####
for combo in siteCombinations:
    print "\nPlotting raw data for site combo: "
    for site in combo:
        print site

    for site in combo:
        x = datasetObjects[site].getAgeValues()
        y = datasetObjects[site].getElevationValues()
```

```

plt.plot(x, y, mapSiteToColour(site) + 's', label="%s,\n n=%i" % (site, len
(x)), markersize=4.0)

## plot the raw data for each site


plt.plot(sorted(allAgesSampled), [datasetModels[site].getModelledElevation(age) for age in sorted(allAgesSampled)],
mapSiteToColour(site), label=site+"_(model)")

## plot the linear interpolation model for each site


plt.hist(x, bottom = histogramFloor, normed=False, bins=globalBins, alpha
=0.4, color=mapSiteToColour(site))

## plot the histogram of data set counts on the plot alongside the
## data itself

## histogram floor was chosen here as a nice looking spot to put the
## count histogram so it doesnt overlap the main data


plt.title("Data_and_Model_for_site_Combination_%s/%s" % (combo[0], combo
[1]))

plt.ylabel('Elevation')
plt.xlabel('Age')
plt.legend(loc=2, prop={'size': 17})

axes1 = plt.gca()
yScaleRange = max(axes1.get_ylim()) - min(axes1.get_ylim())

```

```
axes2 = plt.twinx()
axes2.set_ylabel('Count')
axes2.axis([None,None,0,yScaleRange])
## set the left axis to be elevation relative to datum

## and the right axis to be count of each dataset in each bin

outputFilePath = filePathOnRelativePath("./", fileName='%s-%s_DataAndModel'
                                         % (combo[0], combo[1]), ext="png")
print "Saving rawData plot at '%s'" % outputFilePath
verifyPath("./")
## umm ok
plt.savefig(outputFilePath)
plt.close()
## save the raw data combo graph
#####
#####
#####
## plot the gia graphs and store the raw regression numbers used to create #
## them #####
for conditions in [{"valueCounts": "bothNonZero"}, \
                    ##{"valueDifference": "withinThirtyPercent", "valueCounts": "bothNonZero"}, \
                    ##{"valueDifference": "withinFiftyPercent", "valueCounts": "bothNonZero"}, \
                    ##{"valueDifference": "withinTwentyPercent", "valueCounts": "bothNonZero"}, \
                    ]:
```

```
        {"valueDifference": "withinSeventyFivePercent", "valueCounts": "bothNonZero"}]:  
  
outputPathDict = populateConditionsDict(conditions)  
  
outputPath = convertListToRelativePath([outputPathDict[setting] for setting  
in getCurrentSettingOptions()])  
  
conditionIdString = "_" .join([outputPathDict[setting] for setting in  
getCurrentSettingOptions()])  
  
giaRegressionsByCombo = {}  
  
for combo in siteCombinations:  
  
    print "\nPlotting_gia_for_site_combo:"  
    for site in combo:  
        print site  
  
    ## now the gia calculations  
    for order in ['forward', 'reverse']:  
        ## each comparison has a forward A to B, and reverse B to A,  
        ## comparison, the CIs on the absolute value of slope for this must  
        ## be statistically similar for the comparison to work  
  
        if(order == 'forward'):  
            direct = combo[0]  
            ## d is the site we are using as our direct comparison
```

```
## ie MUST have a measured data point at this age
modelled = combo[1]

## ds is what we are comparing against, so it can just be a
## modelled point

else:
    direct = combo[1]
    modelled = combo[0]

allowableAgeValues = []

## for each comparison, there are only a small number of data values
## from the initial dataset that can be used for valid comparison

## each datapoint used for a gia comparison must be:
## -from the direct dataset
## -in the range covered by the modelled dataset (meaning that if
## the direct comparison dataset has a datapoint available, but the
## modelled one has just been hanging off the end in a straight line
## from the last known datapoint, it cant be considered valid
## -given that theres a bin from startAge to startAge+binWidth that
## the datapoints age is in, that bin needs to hit some criteria for
## the number of datapoints in the bin from both

for age in sorted(allAgesSampled):
    if(datasetModels[direct].ageValueInRawData(age) and datasetModels[
        modelled].ageValueInRangeCoveredByModel(age) and
        datasetModels[modelled].ageComparisonValidForThisBin(
```

```
datasetModels[direct], globalBins, age, conditions) ):

allowableAgeValues.append(age)

else:

    continue

## the case where we have an overlap of the models, but

## either A: no datapoint is actually present for either

## dataset at this age, so comparisons are not honouring

## the raw data, or

## B: we have a datapoint on the set to compare against

## but not the one we are comparing

##else:

##continue

## if the datapoint in question is outside the bounds

## covered by these two datasets, they cant be considered

elevationDiffs = [(datasetModels[direct].getModelledElevation(age) -

datasetModels[modelled].getModelledElevation(age)) for age in

allowableAgeValues]

bootstrapper.plotBootstrapsOnDataPlot(plt, allowableAgeValues,

elevationDiffs, mapSiteToColour(modelled), mapSiteToColour(direct

));

thisComparisonGiaDescription = "%s measured data from %s relative to %s %s"

model" % (direct, modelled)

plt.plot(allowableAgeValues, elevationDiffs, mapSiteToColour(direct)+''

+'', label=thisComparisonGiaDescription, markersize=4.0)
```

```

linRegressYValues, gradient, intercept, gradientError, yModelHigh,
yModelLow, rSquare = getLinearModel(allowableAgeValues,
elevationDiffs)

if(direct != modelled):
    giaRegressionKey = "%s-%s:%s" % (combo[0], combo[1], order)

giaRegressionsByCombo[giaRegressionKey] = {"N": len(
    allowableAgeValues), "gradientEstimator": gradient, "gradientError": gradientError, "gradient": [gradient+(1.96*
    gradientError), gradient-(1.96*gradientError)], "intercept": intercept, "rSquare": rSquare}

plt.suptitle("Plot of Elevation Diff for %s relative to %s model by %s\norder for %s - %s, n=%i\ny=mx+b, m=% .4f SE(%.4f), b=% .4f, r^2=% .3f" % (direct, modelled, order, combo[0], combo[1],
len(allowableAgeValues), gradient, gradientError, intercept, rSquare), fontsize=10)
plt.ylabel('Elevation')
plt.xlabel('Age')

if(direct == "ATB"):

    plt.legend(loc=2, prop={'size': 14})

else:

    plt.legend( loc=3, prop={'size': 14})

##plt.savefig('./theGIA_%s_relative_to_%s.png' % (d, ds))
## ^ this was creating a ton of clutter

```

```
outputFilePath = filePathOnRelativePath(outputPath+"gias/", fileName='

    theGIA_%s_relative_to_%s' % (direct, modelled), ext="png")

print "Saving_gia_plot_at_%s" % outputPath
verifyPath(outputPath+"gias/")

plt.savefig(outputFilePath)

plt.close()

giaRegressionMappingsByConditions[conditionIdString] =
    giaRegressionsByCombo

plotGradientConfidenceIntervals(giaRegressionsByCombo, giaRegressionKeys,
    giaRegressionDescriptions, outputPathDict)

#####
print "Finished_gia_plots"

#####
## Check for any exact age matches in the datasets provided #####
## Spoiler: there aren't any #####
ageMatches = []

for d in datasets:

    for dv in datasetObjects[d].getAgeValues():

        for od in datasets:

            if(od != d):

                if((dv in datasetObjects[od].getAgeValues())and dv not in

                    ageMatches):

                    ageMatches.append(dv)

print "Exact_age_matches_between_datasets:", ageMatches

#####
```

```
#####
## now that values have been generated for GIA for each site comparison, ##
## convert them to intervals for each site combination and save the result #
## to file #####
for idString in giaRegressionComboMappingsByConditions:
    print "\n\n%s:" % idString

giaRegressionsByCombo = giaRegressionComboMappingsByConditions[idString]
siteCombos = ["ATB-BATB", "GTB-ATB", "GTB-BATB", "GTB-TAHB", "TAHB-ATB", "TAHB-
BATB"]

with open("%s_intervals.csv" % idString, "wb") as csv_file:
    writer = csv.writer(csv_file, delimiter=',')

    writer.writerow([ "name", "Slope_UncEstimator", "Slope_Error", "r_Squared",
                     "Slope_C.I.(95p)"])
    for regress in sortedKeys:
        description = giaRegressionDescriptions[regress]
        ciStart = 100*100*giaRegressionsByCombo[regress]['gradient'][0]
        ciEnd = 100*100*giaRegressionsByCombo[regress]['gradient'][1]
        est = 100*100*giaRegressionsByCombo[regress]['gradientEstimator']
        error = 100*100*giaRegressionsByCombo[regress]['gradientError']
        rSquare = giaRegressionsByCombo[regress]['rSquare']

        print regress
```

```
for param in giaRegressionsByCombo[regress]:
    print param

writer.writerow([ description, "%.5f" % est, "%.5f" % error, "%.3f" %
    rSquare, "%.5f_to_.5f" % (ciStart, ciEnd)])

for combo in siteCombos:
    comboSites = combo.split('-')

print combo, comboSites

with open("%s_regressionTable.csv" % combo, "wb") as csv_file:
    writer = csv.writer(csv_file, delimiter=',')
    writer.writerow([ "name", "Slope_Estimator", "Slope_Error", "r_Squared"
        , "Slope_C.I.(95p)"])

    for order in ["forward", "reverse"]:
        regress = "%s:%s" % (combo, order)
        description = giaRegressionDescriptions[regress]
        ciStart = 100*100*giaRegressionsByCombo[regress]['gradient'][0]
        ciEnd = 100*100*giaRegressionsByCombo[regress]['gradient'][1]
        est = 100*100*giaRegressionsByCombo[regress]['gradientEstimator']
        error = 100*100*giaRegressionsByCombo[regress]['gradientError']
        rSquare = giaRegressionsByCombo[regress]['rSquare']
        writer.writerow([ description, "%.5f" % est, "%.5f" % error, "%.3f"
            % rSquare, "%.5f_to_.5f" % (ciStart, ciEnd)])

with open("%s_mergedIntervals.csv" % idString, "wb") as csv_file:
```

```
writer = csv.writer(csv_file, delimiter=',')

writer.writerow(["siteCombination", "startValue", "endValue"])

for combo in siteCombos:

    for order in ["forward", "reverse"]:

        regress = "%s:%s" % (combo, order)

        print giaRegressionDescriptions[regress], ",",
        print giaRegressionsByCombo[regress]['gradient'][0], ",",
        print giaRegressionsByCombo[regress]['gradient'][1]

        est = giaRegressionsByCombo[regress]['gradientEstimator']
        ciStart = 100*100*giaRegressionsByCombo[regress]['gradient'][0]
        ciEnd = 100*100*giaRegressionsByCombo[regress]['gradient'][1]

        if(est < 0):
            ciStart = -ciStart
            ciEnd = -ciEnd

        if(order == "forward"):

            forwardInterval = {"start":min(ciStart, ciEnd), "end": max(
                ciStart, ciEnd)}

        elif(order == "reverse"):

            reverseInterval = {"start":min(ciStart, ciEnd), "end": max(
                ciStart, ciEnd)}

        mergedInterval = mergeConfidenceIntervals(forwardInterval,
                                                    reverseInterval)

        print combo, ":", mergedInterval

        if(mergedInterval == "NoOverlap"):
```

```
writer.writerow([combo, "%s_merged" % combo, mergedInterval, ""])

else:

writer.writerow([combo, "%s_merged" % combo, "%.3f" %

mergedInterval[0], "%.3f" % mergedInterval[1]])

#####
## plot a legend showing the colour coding system for the sites #####
## this sounded like a decent idea earlier, but it eventually proved #####
## not to be needed #####
for site in sites:

plt.plot([1], [1], mapSiteToColour(site)+'s', label=site, markersize=20)

plt.plot([1], [1], mapSiteToColour(site), label=site+"_model", markersize

=20)

plt.axis('off')

plt.legend(loc=3, prop={'size': 29})

plt.savefig("legendary.png")

plt.close()

#####
```

7.2 Source code for `giaUtils.py`

```
## giaUtils.py #####  
## utility functions for gia thesis #####  
#####  
import os  
  
  
def mapSiteToColour(siteLoc):  
  
    siteMappings ={'BATB': 'b', 'TAHB': 'g', 'GTB': 'm', 'ATB': 'r'}  
  
    return siteMappings[siteLoc]  
  
  
def convertListToRelativePath(someListOfStrings):  
  
    output = "./"  
  
    for someStr in someListOfStrings:  
  
        output += "%s/" % someStr  
  
    return output  
  
  
def filePathOnRelativePath(somePath, fileName, ext=None):  
  
    if(ext == None):  
  
        return "%s%s" % (somePath, fileName)  
  
    else:  
  
        return "%s%s.%s" % (somePath, fileName, ext)  
  
  
def verifyPath(somePath):  
  
    if(os.path.exists(somePath)):  
  
        if(os.path.isdir(somePath)):  
  
            return True  
  
        else:
```

```
        print "Path '%s' exists, but is not a directory"
        return False
    else:
        print "Path did not exist, attempting to create it..."
        os.makedirs(somePath)
        return os.path.exists(somePath)

## input dict
## ie {"valueDifference": "withinTwentyPercent", "valueCounts": "bothNonZero"}


def populateConditionsDict(inputDict):
    if("valueDifference" not in inputDict):
        inputDict["valueDifference"] = "any"
    if("valueCounts" not in inputDict):
        inputDict["valueCounts"] = "any"
    return inputDict
    ## technically it mutates the dict, but this is ok too


def getCurrentSettingOptions():
    return [ "valueCounts", "valueDifference"]

def mergeConfidenceIntervals(intervalA, intervalB):
    if((intervalA["start"] >= intervalB["end"]) or (intervalB["start"] >=
        intervalA["end"])):
        return "NoOverlap"
    else:
```

```
## some overlap

if((intervalB["start"] < intervalA["end"]) and (intervalB["end"] >
    intervalA["end"])):
    return (intervalB["start"], intervalA["end"])

elif((intervalA["start"] < intervalB["end"]) and (intervalA["end"] >
    intervalB["end"])):
    return (intervalA["start"], intervalB["end"])

elif((intervalB["start"] > intervalA["start"]) and (intervalB["end"] <
    intervalA["end"])):
    return (intervalB["start"], intervalB["end"])

elif((intervalA["start"] > intervalB["start"]) and (intervalA["end"] <
    intervalB["end"])):
    return (intervalA["start"], intervalA["end"])

if(__name__ == "__main__"):

    print convertListToRelativePath(["withinTwentyPercent", "baseFixedAt450", "
        gias"])


---


```

7.3 Source code for rawhide/bootstrapper.py

```
## bootstrapper.py #####  
## tools to create a bootstrap for a linear regression model, and plot this ##  
## in matplotlib #####  
#####  
  
import numpy as np  
import matplotlib.pyplot as plt  
from sklearn.linear_model import LinearRegression  
  
  
def plotBootstrapsOnDataPlot(pllt, x, y, strapColor='grey', regressColor='red'):  
  
  
    # Extend x data to contain another row vector of 1s  
    x = np.asarray(x)  
    y = np.asarray(y)  
  
    X = np.vstack([x, np.ones(len(x))]).T  
  
  
    plt.figure(figsize=(12,8))  
    for i in range(0, 500):  
        sample_index = np.random.choice(range(0, len(y)), len(y))  
  
        X_samples = X[sample_index]  
        y_samples = y[sample_index]  
  
        lr = LinearRegression()  
        lr.fit(X_samples, y_samples)  
        y_hat = lr.predict(X)
```

```
lr.fit(X_samples, y_samples)

plt.plot(x, lr.predict(X), color=strapColor, alpha=0.1, zorder=1)

lr = LinearRegression()

lr.fit(X, y)

plt.plot(x, lr.predict(X), color=regressColor, zorder=5)

if(__name__ == "__main__"):

    ## Create toy data and bootstrap it to verify everything is working

    ## correctly

    x = np.linspace(0, 10, 20)

    y = x + (np.random.rand(len(x)) * 10)

    plotBootstrapsOnDataPlot(plt, x, y)

    plt.scatter(x, y, marker='+', color='blue', zorder=5)

    plt.savefig('bootstrapDemo.png')
```

7.4 Source code for rawData.py

```
## rawData.py #####\n## object containing the raw data object used to wrap what gets pulled out ####\n## of the spreadsheet file ####\n#####\n\n\nclass siteData(object):\n\n    ## siteData: Str Listof(Any) -> siteData\n\n    def __init__(self, siteName, rawData):\n\n        self.dataHeader = []\n\n        self.data = [row for row in rawData if len(row) == 2]\n\n        ## filter to only those rows that contain exactly two elements in order\n\n        ## to get rid of the unimportant details at the top of each sheet\n\n        self.siteName = siteName\n\n\n\n\ndef getAgeValues(self):\n\n    ## get all available (measured) data points for age\n\n    return [row[1] for row in self.data]\n\n\n\n\ndef getElevationValues(self):\n\n    ## get the entire list of elevation values for this dataset\n\n    return [row[0] for row in self.data]
```

```
def getElevationByGivenAge(self, someAge):

    ## map an age to an elevation if possible
    for row in self.data:

        if(row[1] == someAge):

            return row[0]

    def getThisSiteBinCount(self, binStart, binWidth):

        def withinside(someValue, binStart, binWidth):

            ## determine if someValue is between binStart and (binStart+binWidth)

            delta = someValue - binStart

            if((delta >= 0)and(delta <= binWidth)):

                return True

            else:

                return False

        return len([row[1] for row in self.data if withinside(row[1], binStart,
            binWidth)])

    def getSiteName(self):

        return self.siteName
```

7.5 Source code for dataModel.py

```
## dataModel.py #####
## Base class for building data models (ie interpretations of what the age #####
## is for the entire age range that the data spans) #####
#####
from rawData import *

## ideas for future models:
## -same connect the dots idea, but with binned means every so many years
## -use mean water level of other sites in areas where a site has a gap to
## adjust for global water lows caused by climate, etc.

class siteModel(object):
    ## parent to all models that take a set of siteData and attempt to build a
    ## model of elevations for all of the possible age values in between points
    ## where the elevation is directly sampled for that exact time.

    def __init__(self):
        pass

    ## getModelledElevation: Num -> Num
```

```
def inRange(value, high, low):
    if((value <= high) and (value >= low)):
        return True
    return False

## getAgeBinByAgeValue: float, listof(float) -> float, float

## ageValue is a float
## ageBins is some list of bin endpoints

def getAgeBinByAgeValue(ageValue, ageBins):
    baseAge = ageBins[0]
    ageBinsDelta = ageBins[1] - baseAge
    ## should be consistent throughout the ageBins list

    ## now find the nearest start point for a bin to the ageValue

    binStartValue = baseAge

    if(ageValue < baseAge):
        while(True):
            if(inRange(ageValue, binStartValue+ageBinsDelta, binStartValue)):
                return binStartValue, ageBinsDelta
            binStartValue -= ageBinsDelta
    else:
        while(True):
            if(inRange(ageValue, binStartValue+ageBinsDelta, binStartValue)):
```

```
    return binStartValue, ageBinsDelta  
    binStartValue += ageBinsDelta
```

7.6 Source code for linearInterpolationModel.py

```
## linearInterpolationModel.py #####
## model for elevation v. time values made by connecting point to point, ie #####
## "connect the dots" formally known as linear interpolation #####
#####
from dataModel import *

import numpy as np

def percentageDifference(someValue, anotherValue):
    average = float(someValue + anotherValue)/2.0
    diff = abs(someValue - anotherValue)
    return float(diff/average)

def conditionMet(thisBinCount, otherBinCount, condition):
    if(condition == "any"):
        return True
    elif(condition == "bothNonZero"):
        if((thisBinCount > 0) and (otherBinCount > 0)):
            return True
        return False
    elif(condition == "withinTwentyPercent"):
        if(percentageDifference(thisBinCount, otherBinCount) <= 0.20):
            return True
        return False
    elif(condition == "withinThirtyPercent"):
```

```
        if(percentageDifference(thisBinCount, otherBinCount) <= 0.30):

            return True

        return False

    elif(condition == "withinFiftyPercent"):

        if(percentageDifference(thisBinCount, otherBinCount) <= 0.50):

            return True

        return False

    elif(condition == "withinSixtyPercent"):

        if(percentageDifference(thisBinCount, otherBinCount) <= 0.60):

            return True

        return False

    elif(condition == "withinSeventyFivePercent"):

        if(percentageDifference(thisBinCount, otherBinCount) <= 0.75):

            return True

        return False

class siteModelConnectTheDots(siteModel):


## siteModelConnectTheDots: siteData -> siteModelConnectTheDots


def __init__(self, availableData):

    self.siteName = availableData.getSiteName()

    self.rawDataObject = availableData


def ageValueIsInRangeCoveredByModel(self, someAge):

    if(someAge in self.rawDataObject.getAgeValues()):

        return True

    else:
```

```
    maxAgeCovered = max(self.rawDataObject.getAgeValues())

    minAgeCovered = min(self.rawDataObject.getAgeValues())

    if((someAge <= maxAgeCovered)and(someAge >= minAgeCovered)):

        return True

    else:

        return False


def ageValueInRawData(self, someAge):

    if(someAge in self.rawDataObject.getAgeValues()):

        return True

    return False


def ageComparisonValidForThisBin(self, otherModelToCompareAgainst, globalBins,
                                 ageValue, conditions):

    binStart, binWidth = getAgeBinByAgeValue(ageValue, globalBins)

    thisModelsBinCount = self.rawDataObject.getThisSiteBinCount(binStart,
                                                               binWidth)

    otherModelsBinCount = otherModelToCompareAgainst.rawDataObject.

        getThisSiteBinCount(binStart, binWidth)

    for condition in conditions:

        if(not conditionMet(thisModelsBinCount, otherModelsBinCount, conditions[
            condition])):

            return False

    return True

## the loop successfully met every condition, so we're good to go
```

```
def getModelledElevation(self, someAge):

    if(someAge in self.rawDataObject.getAgeValues()):

        return self.rawDataObject.getElevationByGivenAge(someAge)

    else:

        ## dont have a datapoint available at that age value, so we need to

        ## interpolate linearly between them to get it

        ageValues = np.array(self.rawDataObject.getAgeValues())



        if(ageValues[ageValues < someAge].size == 0):

            ## case where our value to interpolate is off the bottom end

            ## of the dataset, so we extrapolate from the last two values

            ## min, and 2dmin

            minValue = min(ageValues)

            restOfValues = np.array([val for val in ageValues if val != minValue])

            ## maybe npifying the array will make the min/max calls faster

            ## idk

            secondMinValue = min(restOfValues)

            ageDelta = someAge - secondMinValue

            ## distance from second smallest to the point we want to

            ## interpolate

            secondMinAgeElevation = self.rawDataObject.getElevationByGivenAge(
                secondMinValue)

            minAgeElevation = self.rawDataObject.getElevationByGivenAge(minValue)
```

```
        outputElevationGuess = secondMinAgeElevation + ( (ageDelta/(abs(
            secondMinValue-minValue)))*(secondMinAgeElevation -
            minAgeElevation) )

    elif(ageValues[ageValues > someAge].size == 0):
        ## case where our value to interpolate is off the top end
        maxValue = max(ageValues)
        restOfValues = np.array([val for val in ageValues if val != maxValue])
        ## npifying this array could make the min/max calls faster

        secondMaxValue = max(restOfValues)

        ageDelta = someAge - secondMaxValue

        secondMaxAgeElevation = self.rawDataObject.getElevationByGivenAge(
            secondMaxValue)
        maxAgeElevation = self.rawDataObject.getElevationByGivenAge(maxValue)

        outputElevationGuess = secondMaxAgeElevation + ( (ageDelta/(abs(
            maxValue - secondMaxValue)))*(maxAgeElevation -
            secondMaxAgeElevation) )

    else:
        closestAgeBelow = ageValues[ageValues < someAge].max()
        closestAgeAbove = ageValues[ageValues > someAge].min()
```

```
ageDelta = closestAgeAbove - closestAgeBelow

elevBelow = self.rawDataObject.getElevationByGivenAge(closestAgeBelow)
elevAbove = self.rawDataObject.getElevationByGivenAge(closestAgeAbove)

elevDelta = elevAbove - elevBelow

outputElevationGuess = elevBelow + elevDelta*((someAge -
closestAgeBelow)/(ageDelta))

return outputElevationGuess
```
