

Investigating the Correlation between Drosophila Genus Biodiversity and Geographic Regions Using Latitude and BIN Richness

Storyboard by Iroayo Toki

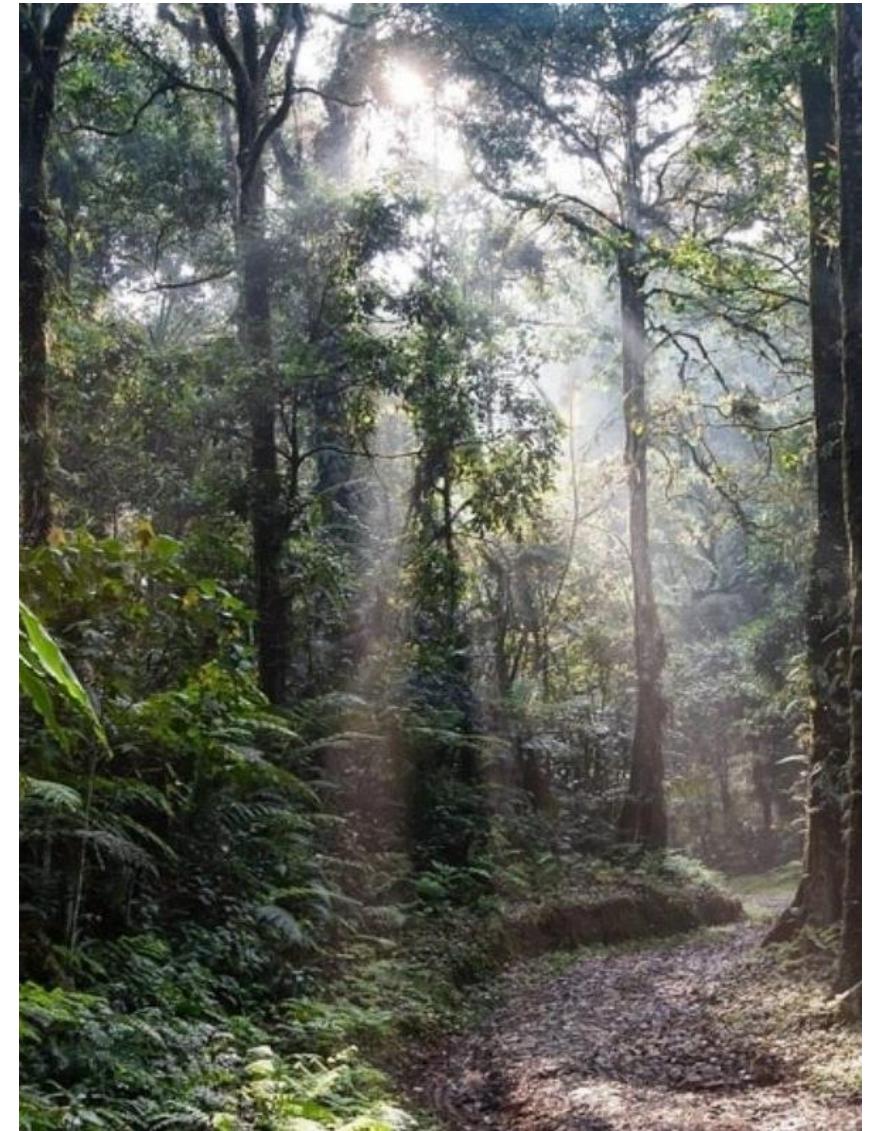
Introduction



- Fruitflies are widely used as research organisms in genetics due to the similarity in their genes with human beings. (Crowley, 2024)
- This makes them great for researching diseases like cancer, diabetes and Alzheimer's Disease. (Crowley, 2024)
- Factors like Temperature, presence of fruit bearing trees and climate affect the survival and reproduction of fruitflies. (Yu et al., 2022 , Roche et al., 2024)
- They have a genus name Drosophila with over 1000 species the most common being *Drosophila melanogaster*. (Tolwinski, 2017)

Hypothesis

- Since environmental factors can affect their survival and reproduction it is plausible to infer that certain ecosystems and by extension certain climatic or geographic regions would improve the chances of survival of fruitflies.
- Tropical regions are described to be warm all year round, with various kind of fruit bearing plants which are just the right conditions for fruitfly survival. (Smith, 2025)
- Since long term survival is important in creating biodiversity as useful mutations only occur once in an extended period of time, we can then generate the following hypothesis. (Kardos et al., 2021)

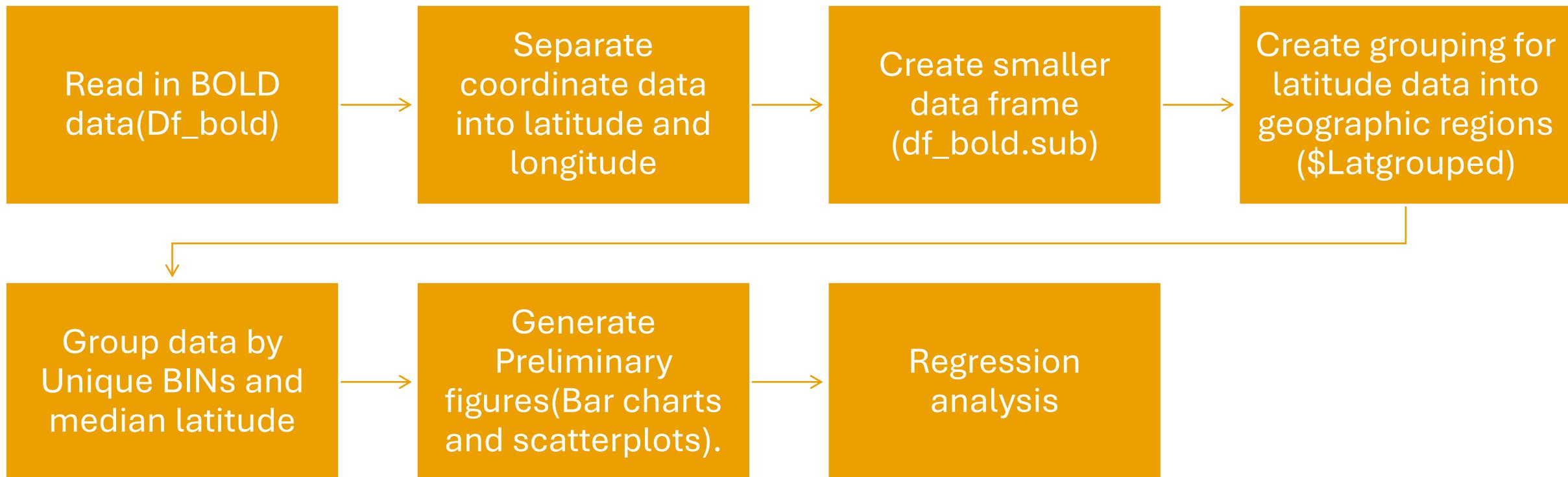




Hypothesis

- Lower environmental variability at lower latitudes as we get closer to the equator leads to increased biodiversity of fruitflies .
- This will be tested by comparing the number of BINs to the average(median) latitude they were found in data from BOLD.
- Studying biodiversity by geographic region can help us understand and improve species conservation by learning the environmental factors best for fruitflies.

Methods



Analytical pipeline for regression analysis



GROUP LATITUDE DATA
INTO BANDS OF 10°



RUN PEARSONS
CORRELATION TEST
`COR()`

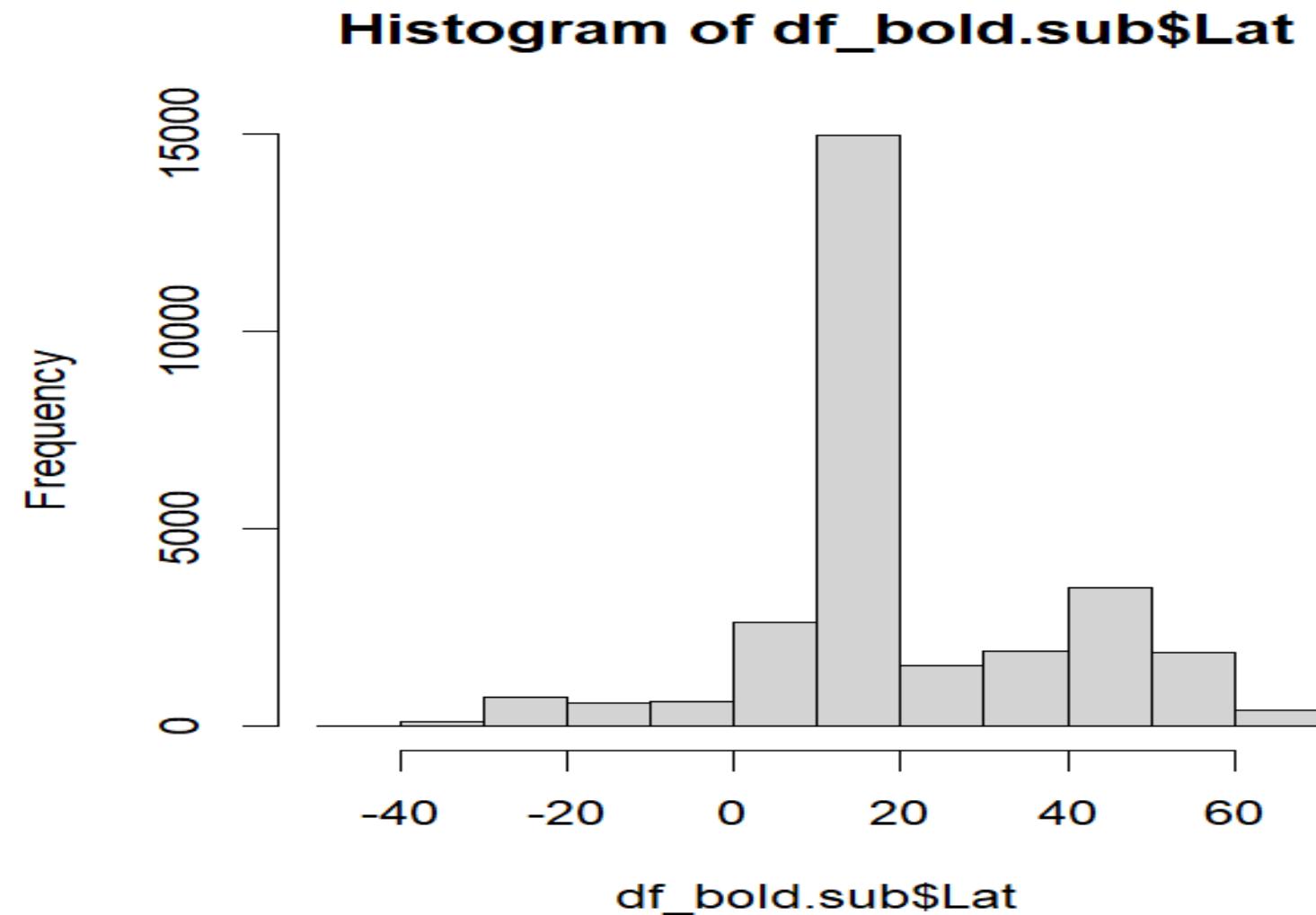


CREATE SIMPLE
REGRESSION MODEL.
`MODEL <- LM()`



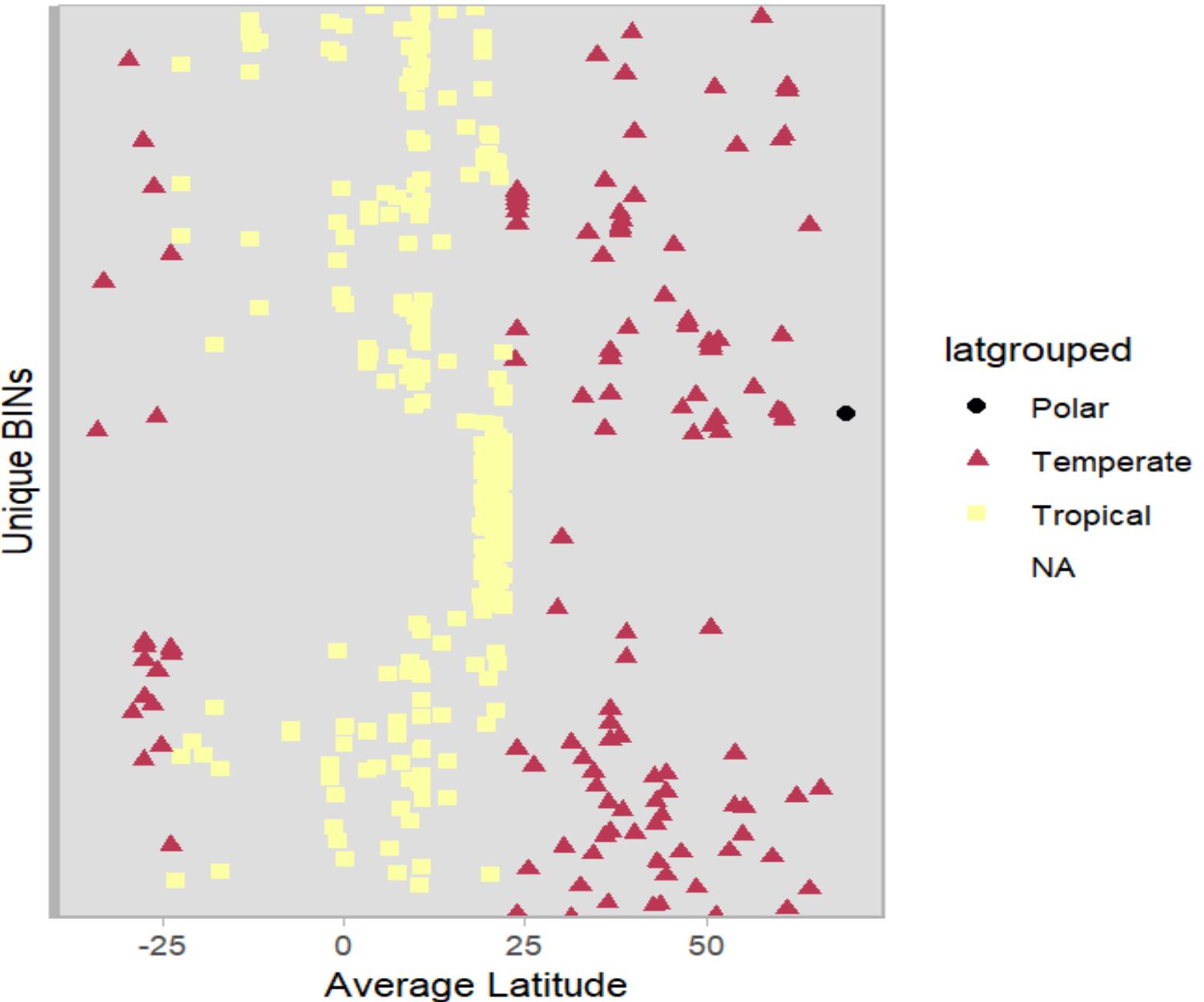
GENERATE SCATTERPLOT
SHOWING REGRESSION
LINE

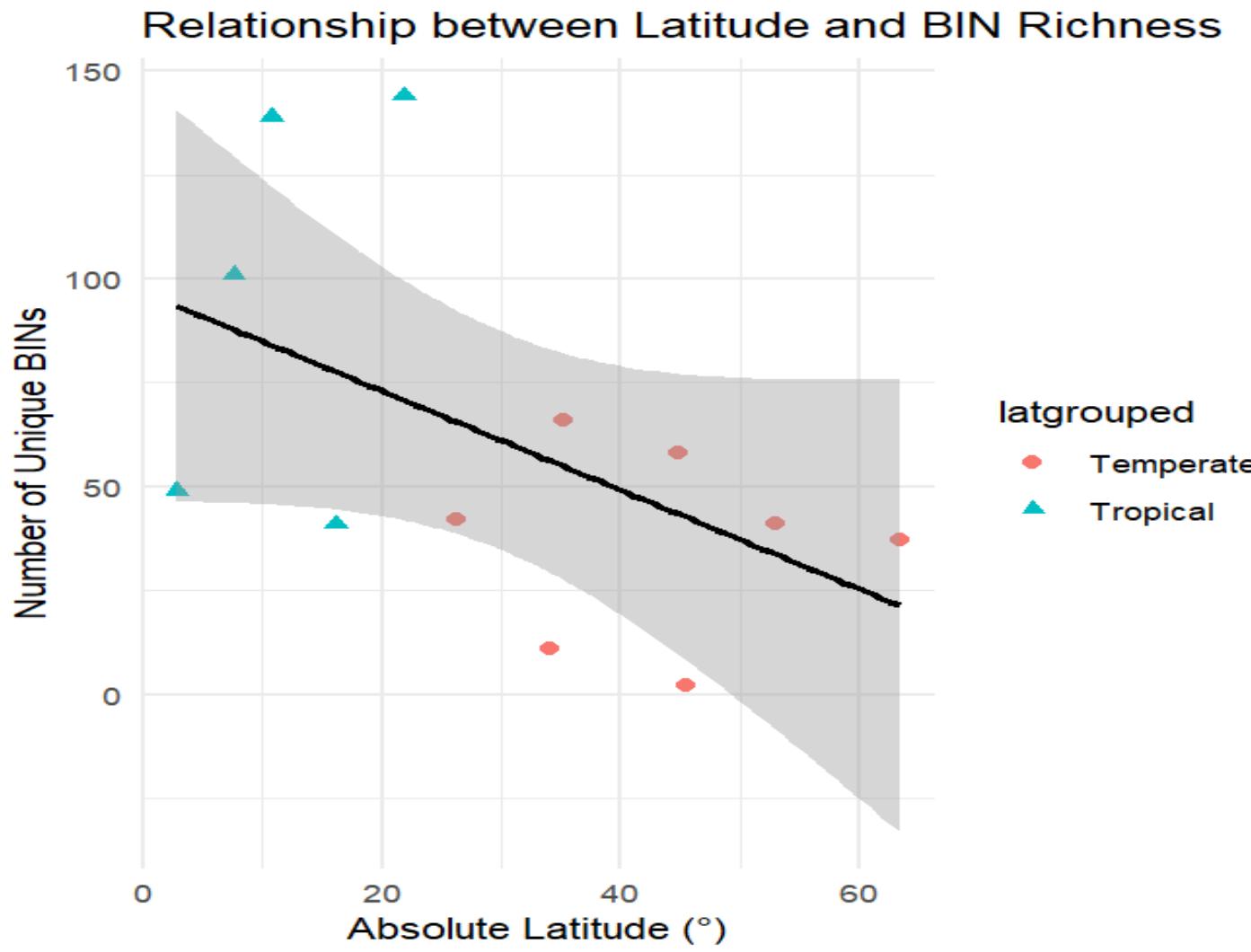
**Sampling
disparity (Over
15000 samples)
in the dataset
between
latitude 10-20°N**



**Unique
Drosophila
BINs cluster in
the tropical
regions of the
Northern
hemisphere**

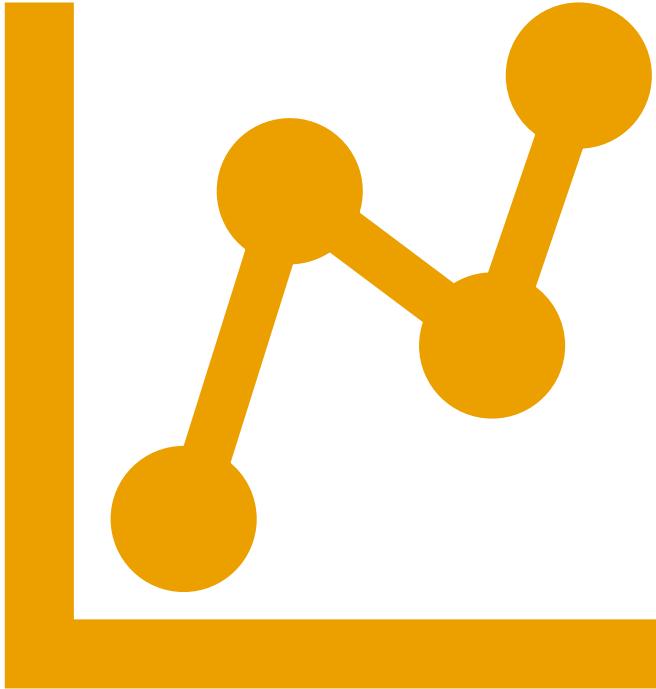
Distribution of Drosophila BINs by latitude





**There is increased
BIN richness in the
Drosophila genus as
we get closer to the
equator(0°)**

Results



- From the Pearson's correlation test we get a value of -0.5043, showing a moderately negative correlation between our variables i.e as our independent variable which was absolute latitude increased there was a decrease in BIN richness.
- For our Linear regression model we get a p-value of 0.09453 which shows that there is an effect however not statistically significant as it does not cross the $p < 0.05$ threshold

Discussion

- Our results show that while there is a moderate negative correlation between latitude and BIN richness but it is not statistically significant. This could be due to a few reasons.
- A. According to a 2023 study there is increased biodiversity for insects in the outer bands of the tropical region and in the subtropical region compared to the inner bands which are close to the equator where they noted precipitation, wettest month and driest month as important factors, it creates this effect where there is first an increase in biodiversity as we move from the equator, then a drop as we move further into the temperate and tropic regions (French et al., 2023).
- B. As seen in the first figure there is a sampling bias in the data with about 15000 samples between 10-20°N while the next highest region has only about 5000 samples which may have affected our analysis. However, we tried to moderate its effect on our analysis by grouping the data into latitude bands before conducting statistical analysis
- The next steps would include
 - i. conducting further sampling on other regions to match the samples in the 10-20°N latitude band
 - ii. Studying environmental factors within the tropical region to understand how they affect biodiversity

Acknowledgements



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