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Abstract

Extinction within the Paleozoic era has been studied in the past, but there still lacks a comprehensive understanding of how extinction risk changed throughout it. Our research project aims to bridge this gap by exploring extinction risk in relation to major Paleozoic phyla and ecological characteristics.

Using R, we analyzed the Stanford Earth Body Size dataset, which includes extensive data (n=8816) on Paleozoic marine animals. In Step 1, regression coefficients were formed, indicating whether being in one of the 6 phyla in each period of the Paleozoic era conferred greater or less extinction risk. In Step 2, the examined ecological characteristics included ocean acidification resilience, feeding patterns, body volume, length, surface area, motility, tiering, circulatory systems, and respiratory organ type. In Step 3, 6 binomial machine learning models were created using the traits from Step 2 to determine whether an individual genus went extinct in a particular period. Our Step 1 results confirm that within these timeframes, while certain phyla have greater extinction risk, extinction risk was not uniform across these groups. Our Step 2 results show certain traits provided advantages and disadvantages for an organism's extinction risk. One interesting pattern was that the only consistently non-significant traits were body length, area, and volume. Likewise with Step 1, extinction risk for each ecological characteristic varied across the Paleozoic. Finally, in Step 3, the results were largely successful. Most of the six models had an accuracy above 80% with the highest being 92% in the Silurian. The areas under the Precision-Recall and the Receiver Operating Characteristic Curves were all in the acceptable (>0.6) range, demonstrating that the model has low false positive/negative rates and is able to distinguish between what trait indicates extinction or survival for each period.

Our research project identified phyla at risk of extinction in each period of the Paleozoic, determined which natural traits incited greater extinction risk, and demonstrated machine learning models trained on fossil descriptors can predict when an individual genus became extinct. Our results confirmed that extinction risk is not consistently dependent on a singular factor nor is it constant across every period of the Paleozoic era.

Methodology

The dataset we used included nine biological and ecological traits, and it also includes taxonomic groupings and phyla. After making logistic regression models for each trait, we then made regularized regression models predicting extinction in each period based on these characteristics. Below are the Paleozoic periods that we are analyzing.

Cambrian
Ordovician
Silurian
Devonian
Carboniferous
Permian

All analyses and plots were made using the programming language R. During stages 1 and 2, the following were our categories of analysis.

1. Phyla - Echinodermata, Mollusca, Chordata, Arthropoda, Brachiopoda, Foraminifera (Taxonomic Group)
2. Descriptors - buffering, feeding patterns, motility, oceanic tiering, respiratory organ type, circulatory system type, length, surface area, volume

For Stage 3, we built regularized binomial regression models.

Figures

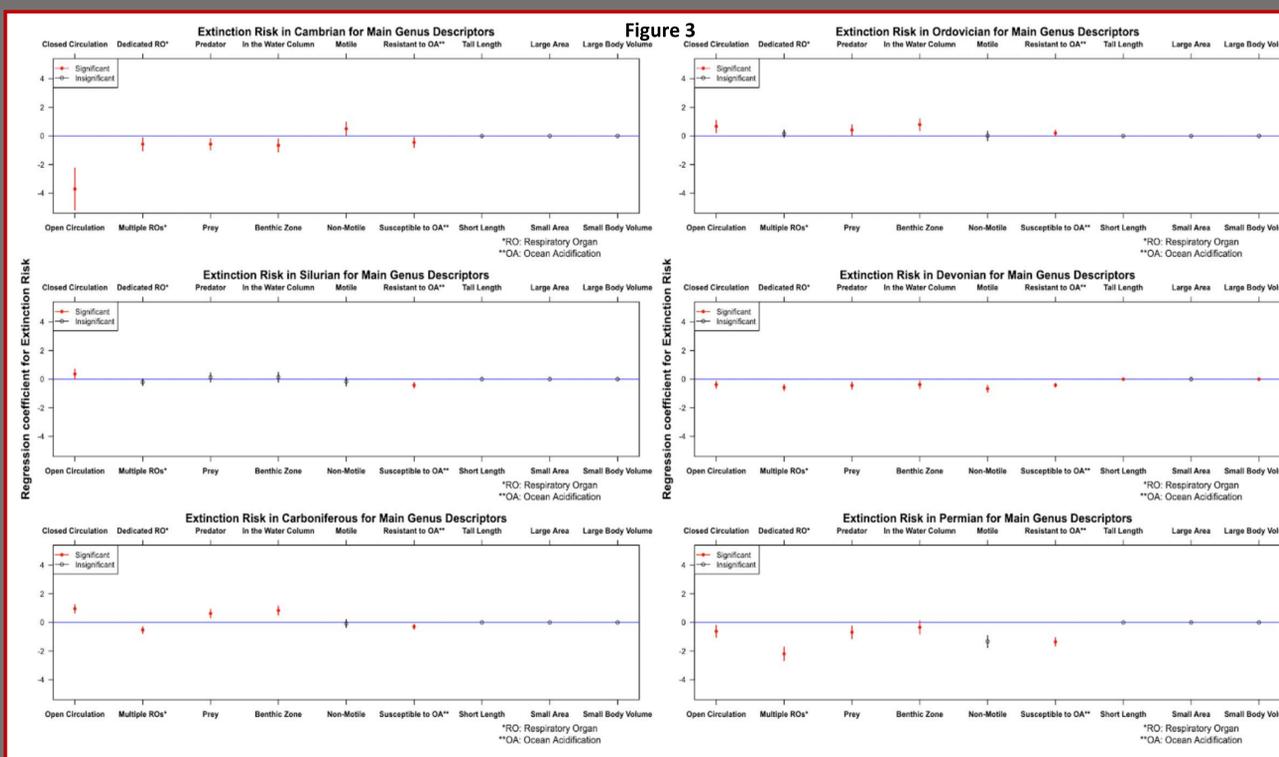
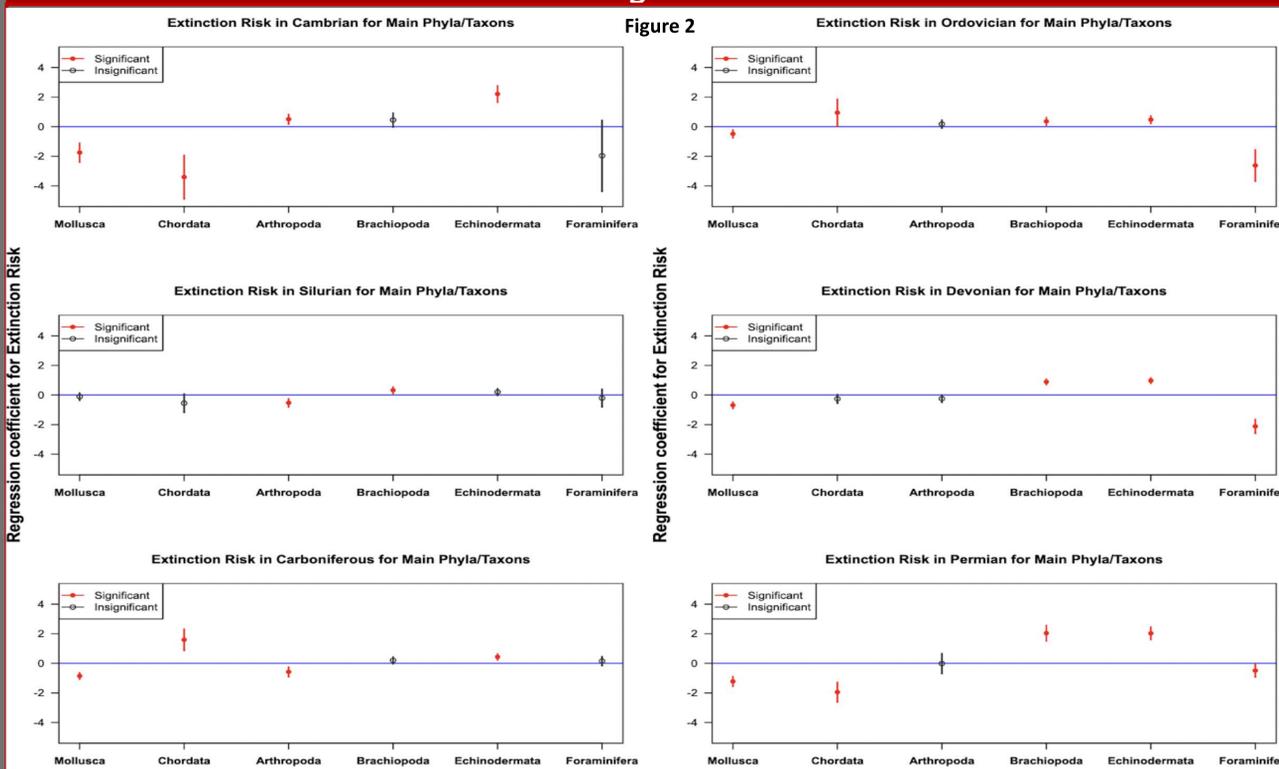


Figure 4

Cambrian Model Features	Score	Ordovician Model Features	Score	Silurian Model Features	Score
1 Circulation System	3.3422079	1 Motility	1.06918333	1 Motility	1.32779174
2 Motility	2.27727063	2 Ocean Acidification	0.85323525	2 Circulation System	1.06206068
3 Resistance	1.12284708	3 Resistance	0.33049509	3 Respiratory Organ System	0.70851463
4 Feeding Patterns	0.42942449	4 Feeding Patterns	0.32654485	4 Feeding Patterns	0.6192809
5 Respiratory Organ System	0.37019192	5 Resistance	0.15505047	5 Resistance	0.59558913
6 Tiering	0.02890546	6 Circulation System	0.02294367	6 Tiering	0.5959338
7 Maximum Length	0.00479994	7 Maximum Length	0.00124485	7 Maximum Length	0.00258455
8 Maximum Area	4.23E-06	8 Maximum Area	4.10E-06	8 Maximum Area	3.22E-06
9 Calculated Maximum Volume	2.39E-09	9 Calculated Maximum Volume	1.08E-08	9 Calculated Maximum Volume	3.32E-08

Figure 5

Devonian Model Features	Score	Carboniferous Model Features	Score	Permian Model Features	Score
1 Resistance	0.70835892	1 Circulation System	2.32544688	1 Circulation System	2.18390017
2 Respiratory Organ System	0.45876009	2 Respiratory Organ System	0.73146419	2 Respiratory Organ System	1.04649225
3 Motility	0.22309167	3 Respiratory Organ System	0.65025381	3 Resistance	0.93551679
4 Circulation System	0.06226351	4 Circulation System	0.00189793	4 Feeding Patterns	0.86094481
5 Maximum Length	0.00555858	5 Maximum Length	0.00555858	5 Tiering	0.74849748
6 Maximum Area	2.20E-05	6 Motility	0.70593468	6 Motility	0.70593468
7 Calculated Maximum Volume	4.32E-08	7 Motility	0.000241	7 Maximum Length	0.000241
8 Feeding Patterns	0	8 Maximum Area	0	8 Calculated Maximum Volume	2.56E-08
9 Tiering	0	9 Calculated Maximum Volume	0	9 Maximum Area	0

1

Binomial Logistic Regression Analysis on Phyla/Class/Order during each stage of Paleozoic identifying likelihood of extinction on each class

2

Binomial Regression Analysis on Phyla/Class/Order during each stage of Paleozoic identifying likelihood of extinction on each genus descriptor (ie predatory feeding, facultative motility, benthic tiering)

3

Developing simple machine learning model (using regularization) to predict whether a taxonomic group/specimen goes extinct in a specific period

Figure 1: Summary of Analysis Stages

Results / Discussion

In Stage One, we conducted logistic regression analyses for each stage of the Paleozoic era for each of the major phyla. The goal was to identify the phyla that have a predilection for extinction during each stage. As evident in Figure 2, we came across some impressive results as 23 of the 36 data points had a significant regression coefficient. 12 data points had a significantly greater extinction risk while 11 were significantly selected for survival. Among various phyla, coefficient values were high in magnitude, but no groups were consistently significant across all periods. However, specifically, Mollusca was generally selected for survival while Echinodermata was generally selected for extinction. For Brachiopoda, you may notice the relatively low coefficients in background periods but a significant extinction risk during the major extinction events in Devonian and Permian. This is in line with the understanding that these extinction events devastated Brachiopoda populations.

In Stage Two, we conducted a logistic regression analysis and binomial test to determine which natural traits incited greater evolutionary selection. The examined factors included ocean acidification resilience (buffering), predatory nature, body volume, length, surface area, motility, tiering, circulatory systems, and respiratory organ type. The largest coefficient value was around -3.9 which demonstrated a high susceptibility of organisms with open circulatory systems for extinction during the Cambrian period. The majority of the data points were significant, 30 of 54. Among these, the only consistently insignificant characteristics were factors associated with body size. This shows that body size had little impact on the extinction risk of organisms. Surprisingly, two descriptors out of nine were significant across the board, circulatory systems, and buffering. Although the type of circulation and amount of buffering that was selected for extinction varied across the Paleozoic.

Finally, in Stage Three, we built machine learning models for each period of the Paleozoic using the ecological factors that we tested in Stage Two. In the Cambrian, the model with the highest accuracy was 92%. In chronological order, the remaining periods had a model with the highest accuracies of 83%, 91%, 79%, 83%, and 84%. As evident, the Devonian appeared to have the lowest accuracy. We believe that with increased data and testing out alternative models, this accuracy can increase greatly.

Major Take-aways:

- Extinction Risk is not uniform across both geologic history or across taxonomic groups
- Certain traits can act as indicators for higher extinction risk; however, these too vary across geologic history
- These traits can even be used to create relatively accurate predictive models.

Future Research:

For future developments, we believe that completing analysis across the rest of geologic history could allow us to identify patterns of how extinction risk changes for each phylum and each trait across every period in Earth's history. This could allow us to know how anoxic conditions affect extinction risk for each phyla/trait or how mass extinctions affect extinction risk for each phyla/trait.

For the machine learning model, we would like to test out Decision Tree or Random Forest Regression Models to predict the exact first and last appearance in geologic history for each genus. Finally, we will look into Building Neural Nets for this type of prediction

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