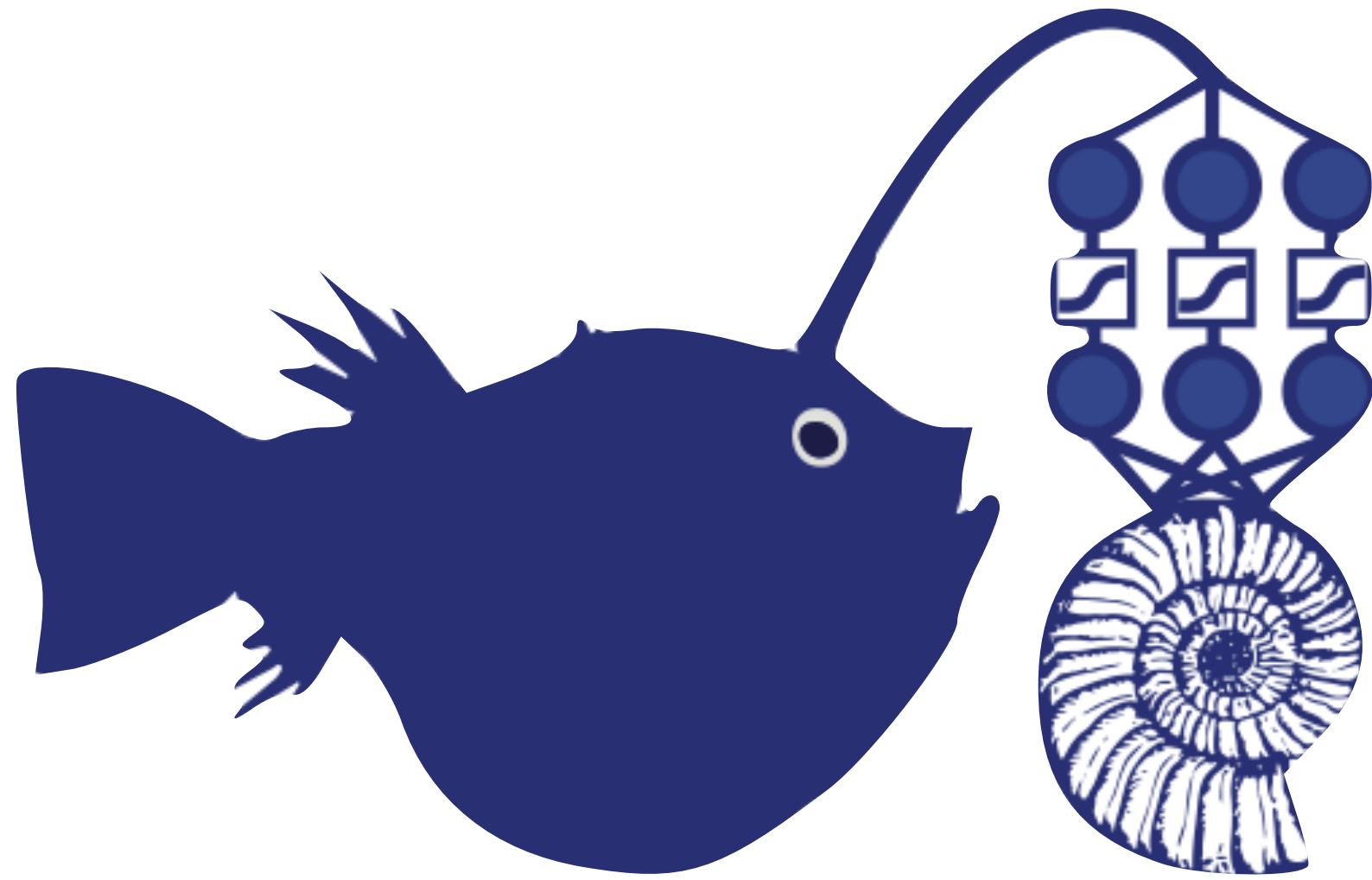
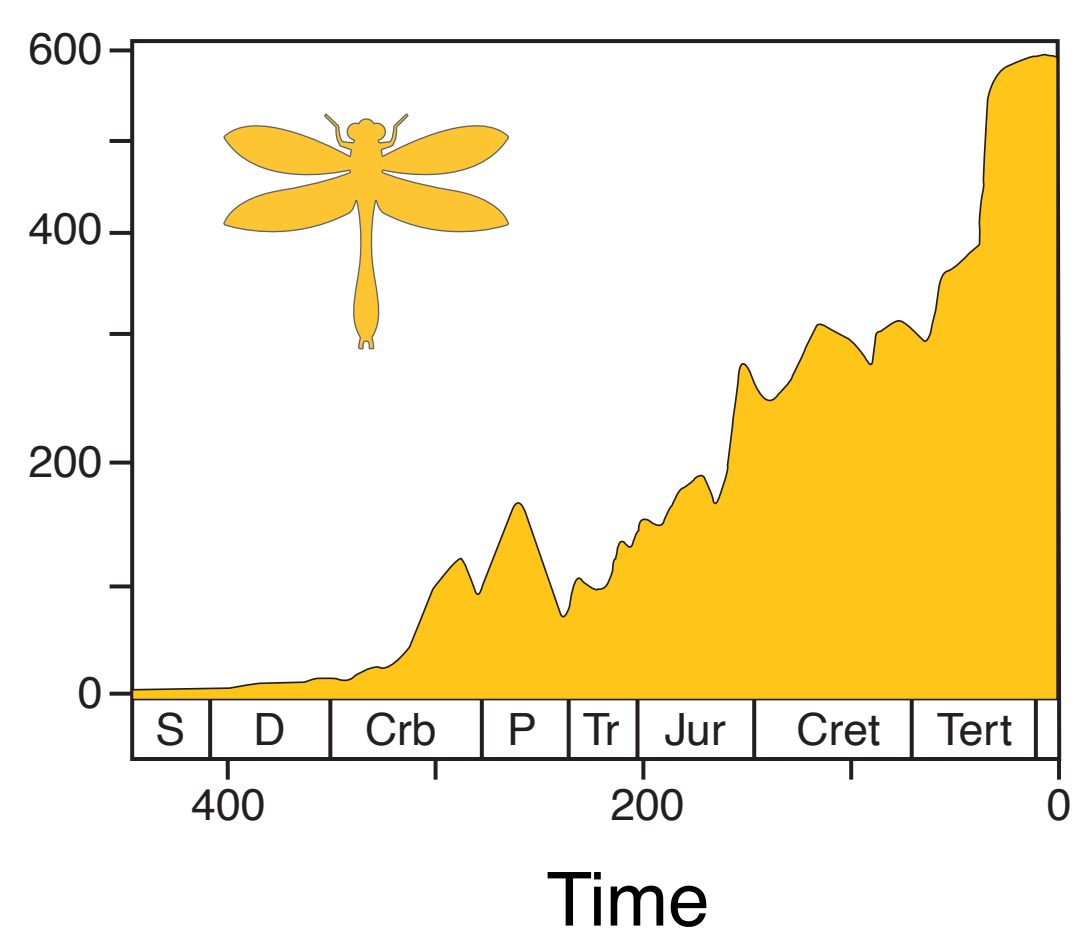
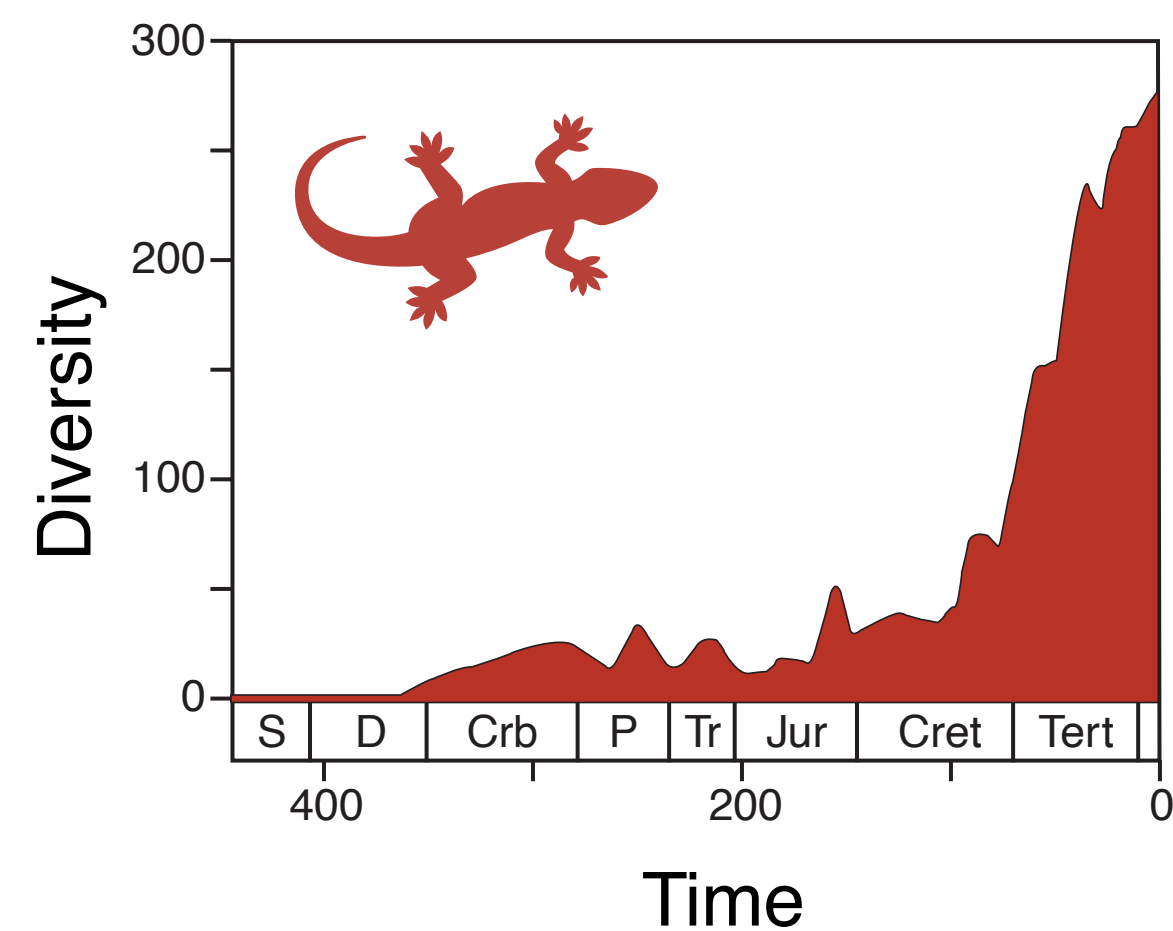
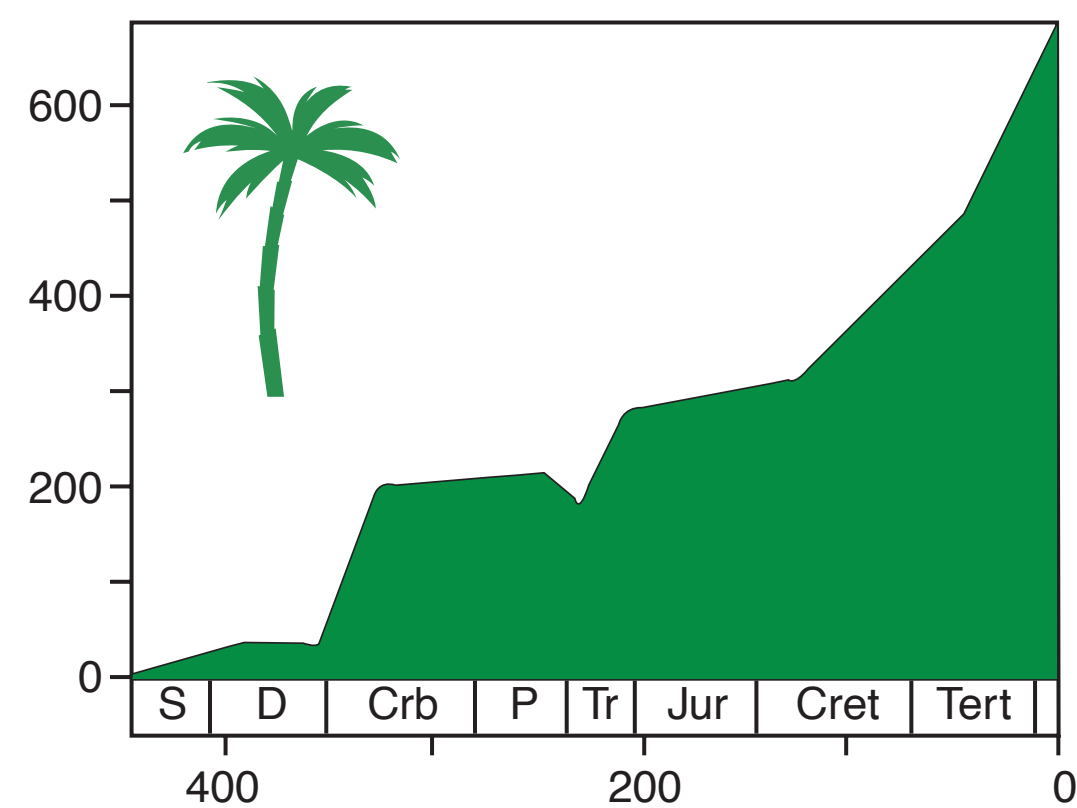
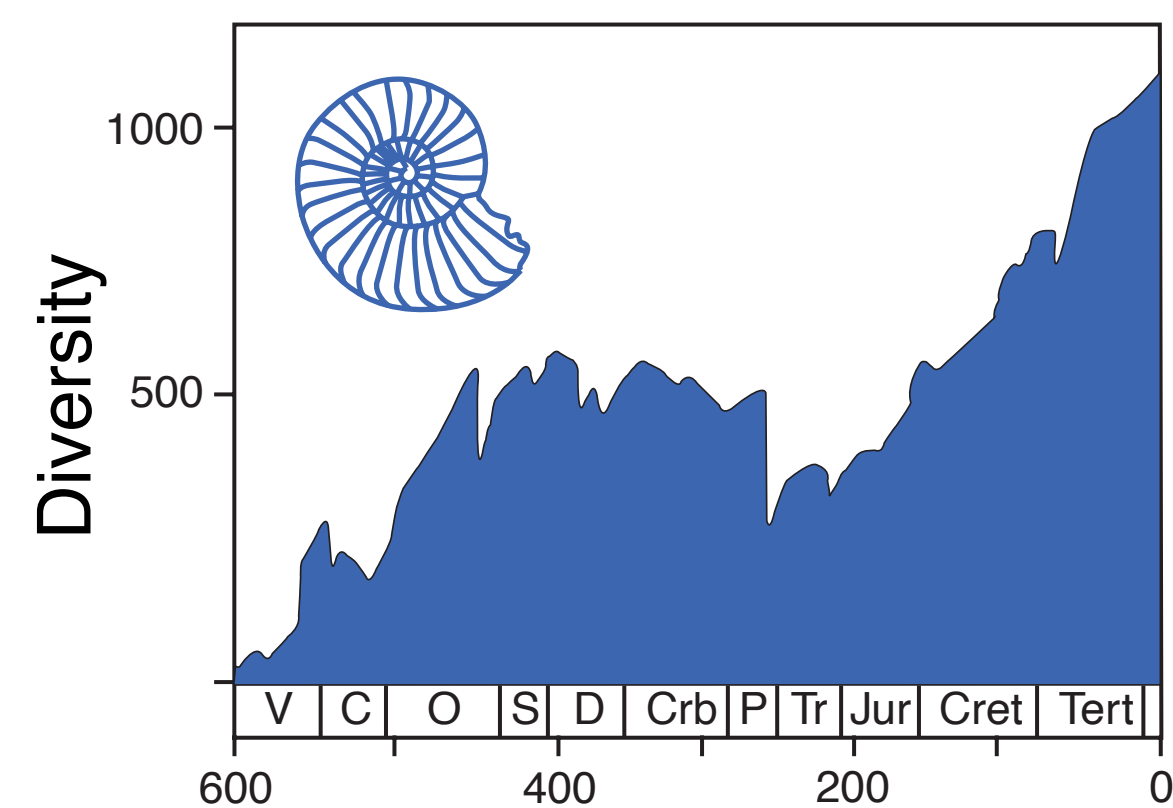




DeepDive – Estimation of diversity trajectories using deep learning

Estimating biodiversity through time

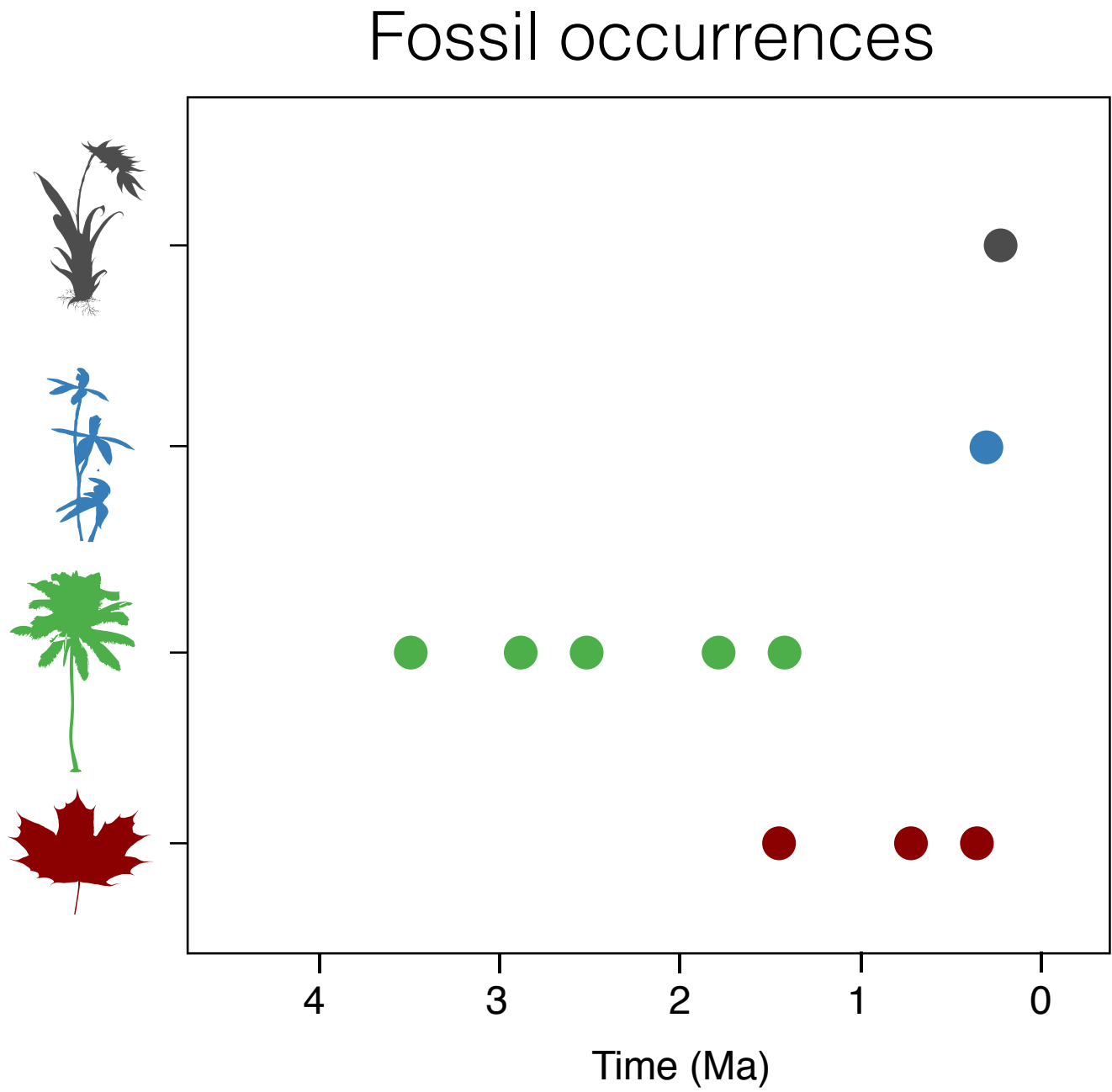


Is there a limit to biodiversity?

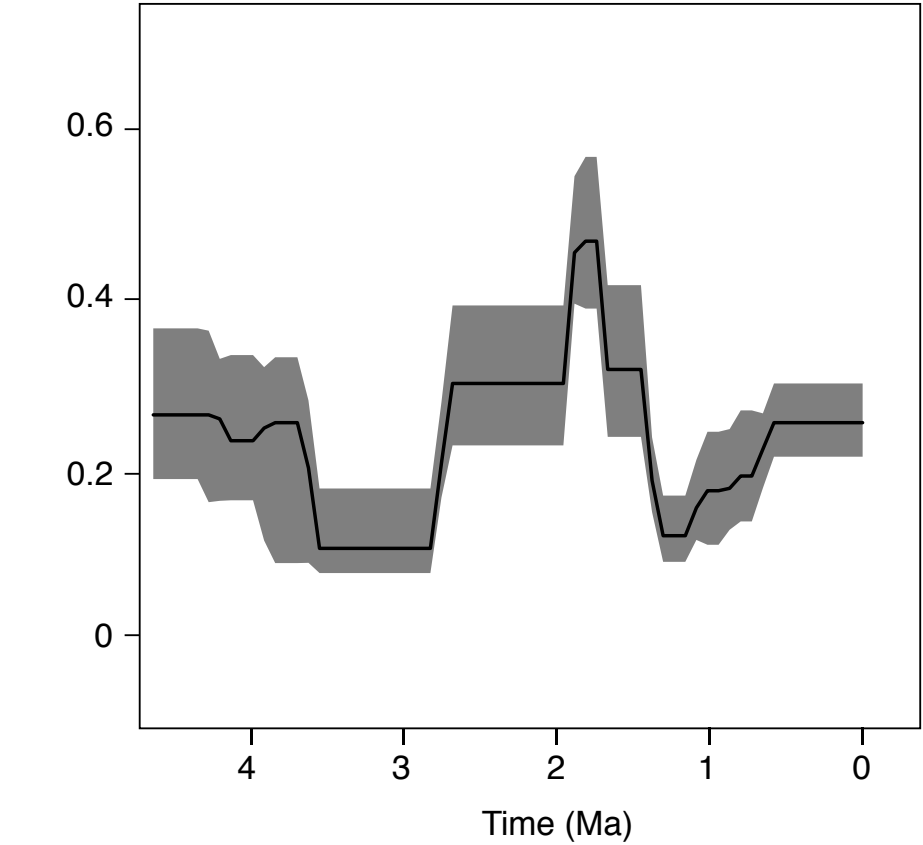
Does biodiversity increase over time?

What are the mechanisms controlling biodiversity?

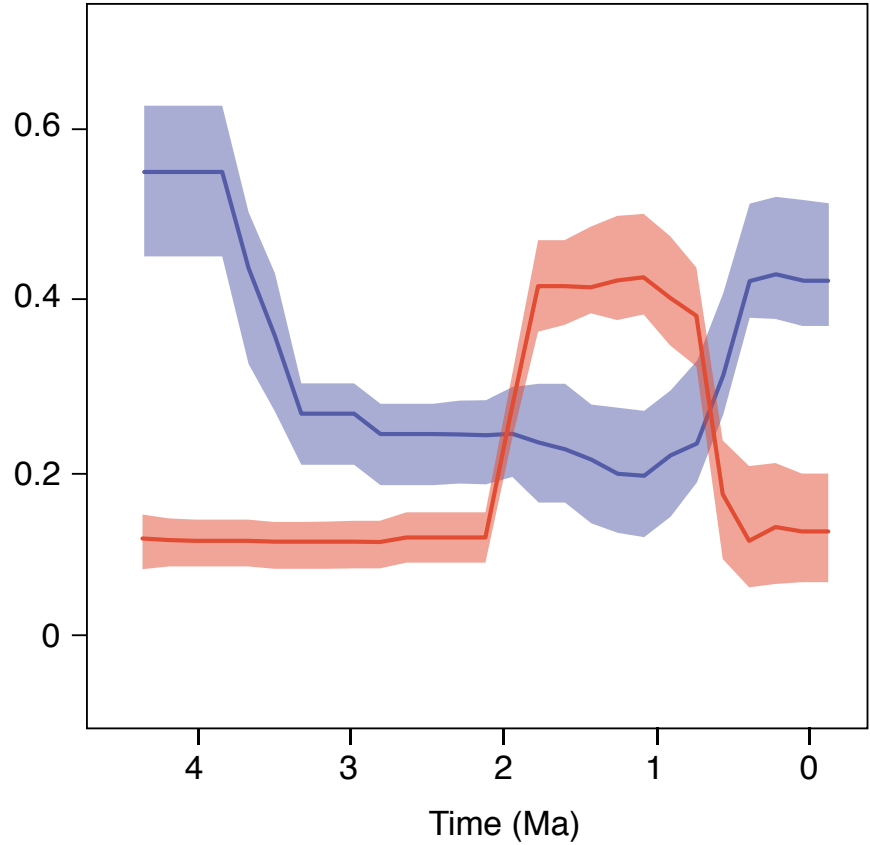
Bayesian estimation of diversity trajectories – mcmcDivE



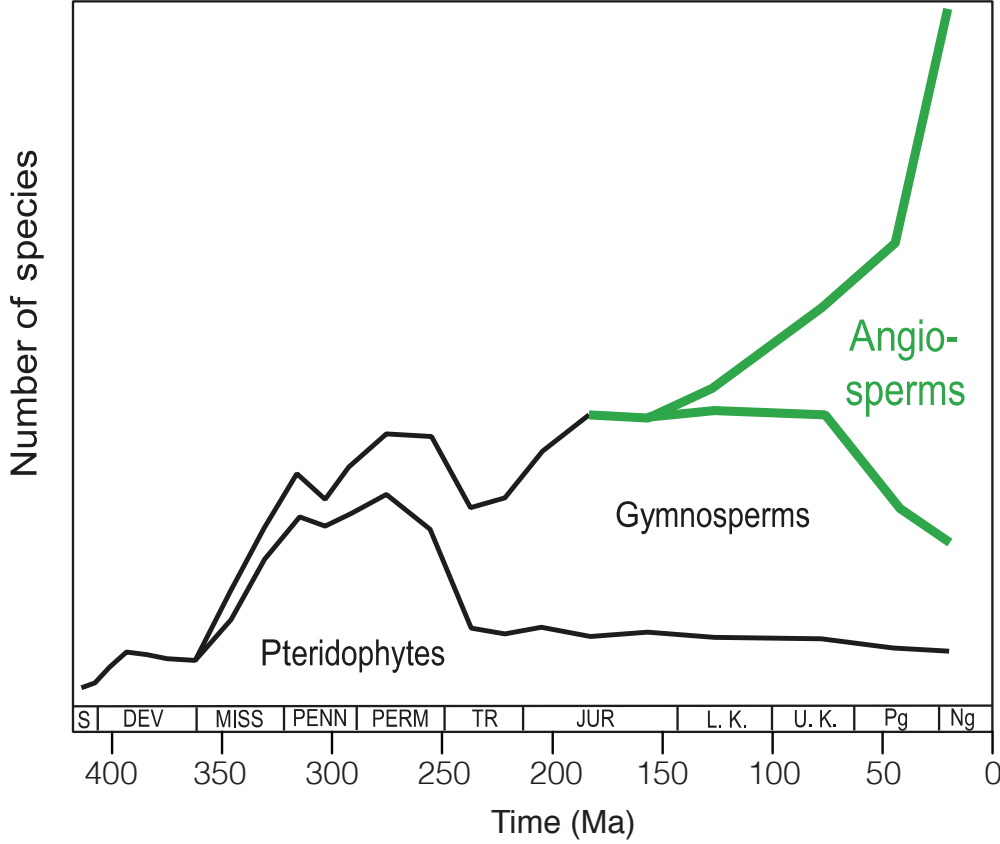
Preservation



Speciation & extinction

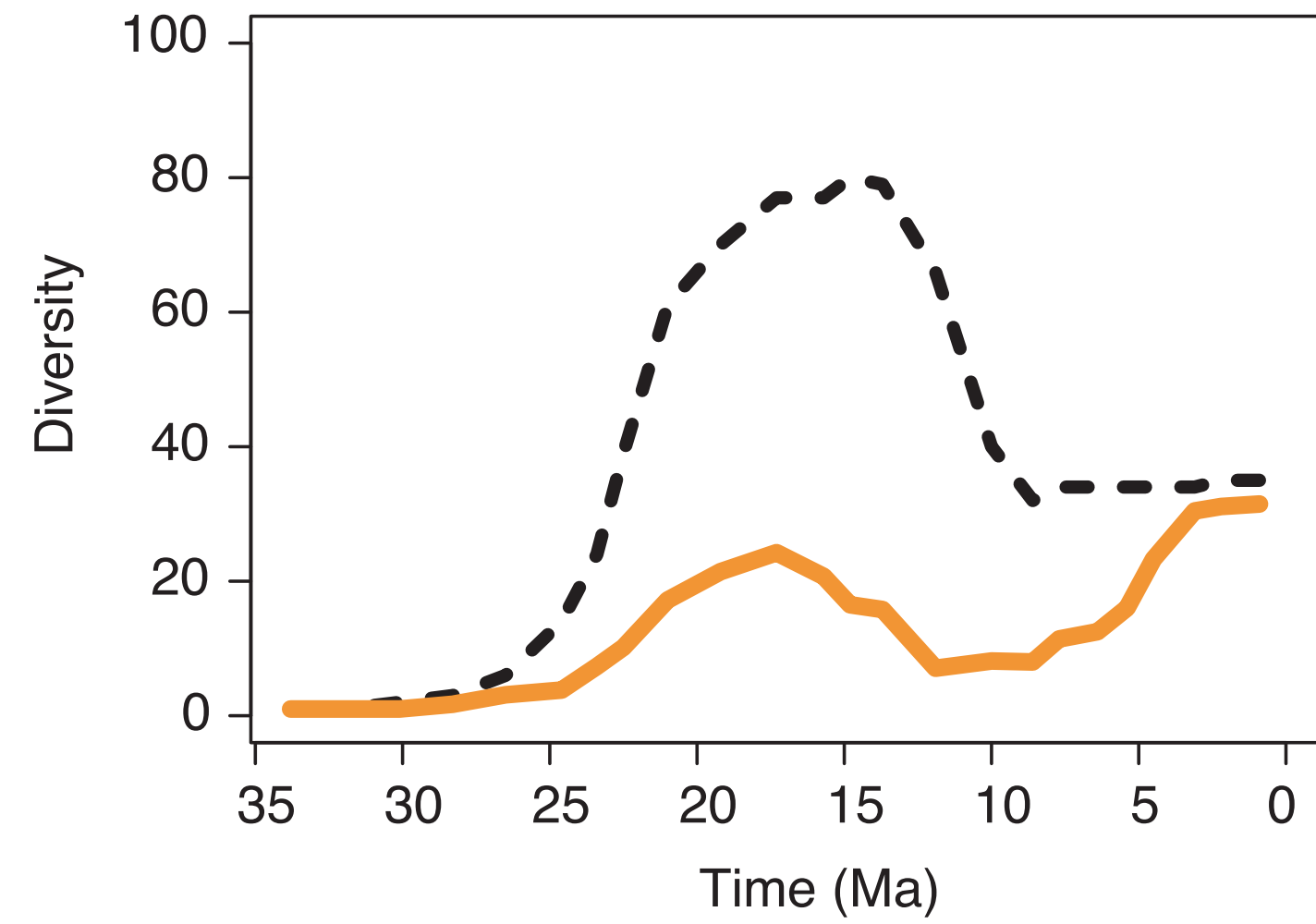


Diversity

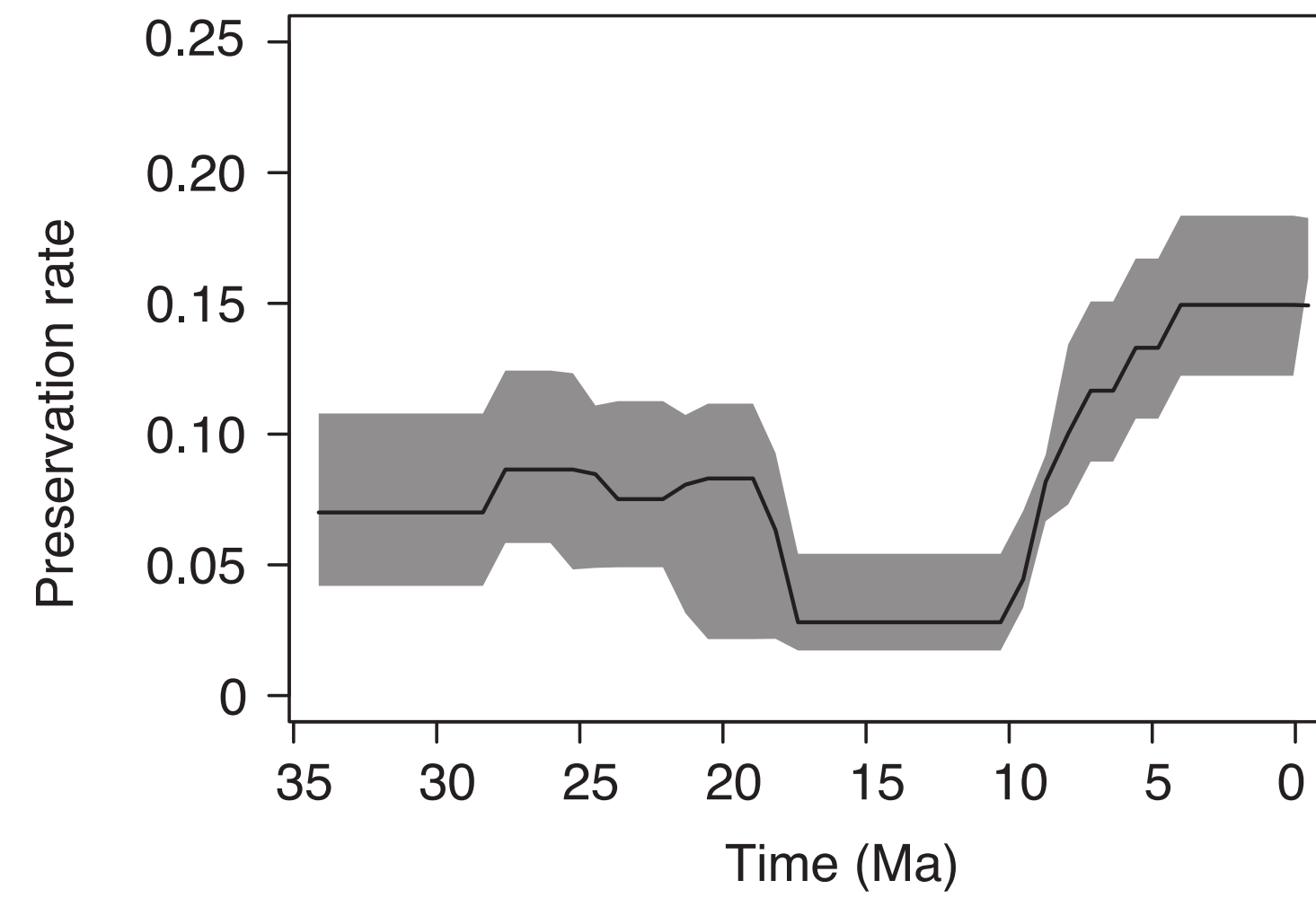


What true diversity could have generated the observed fossil record?

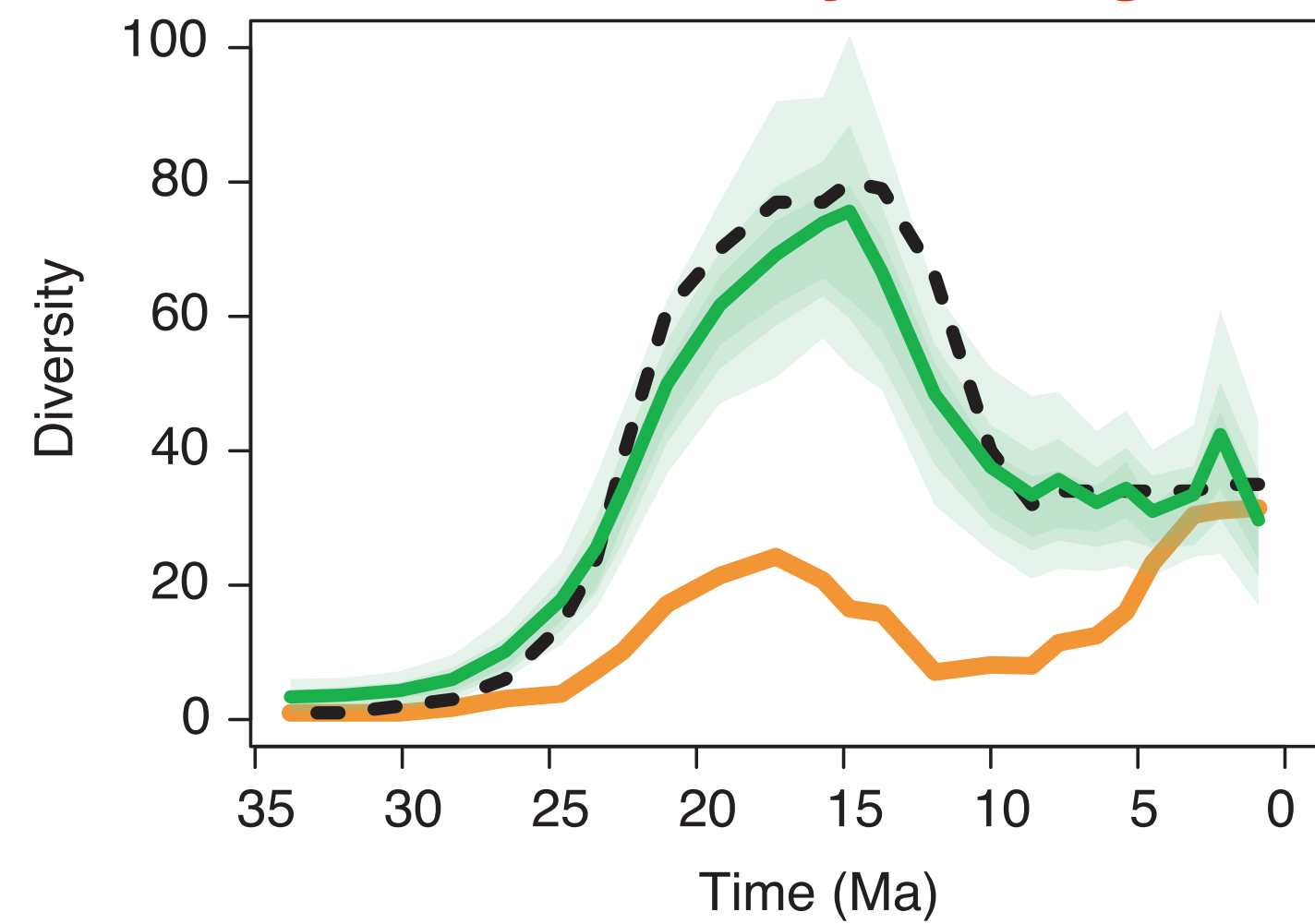
Observed diversity through time



Preservation rates



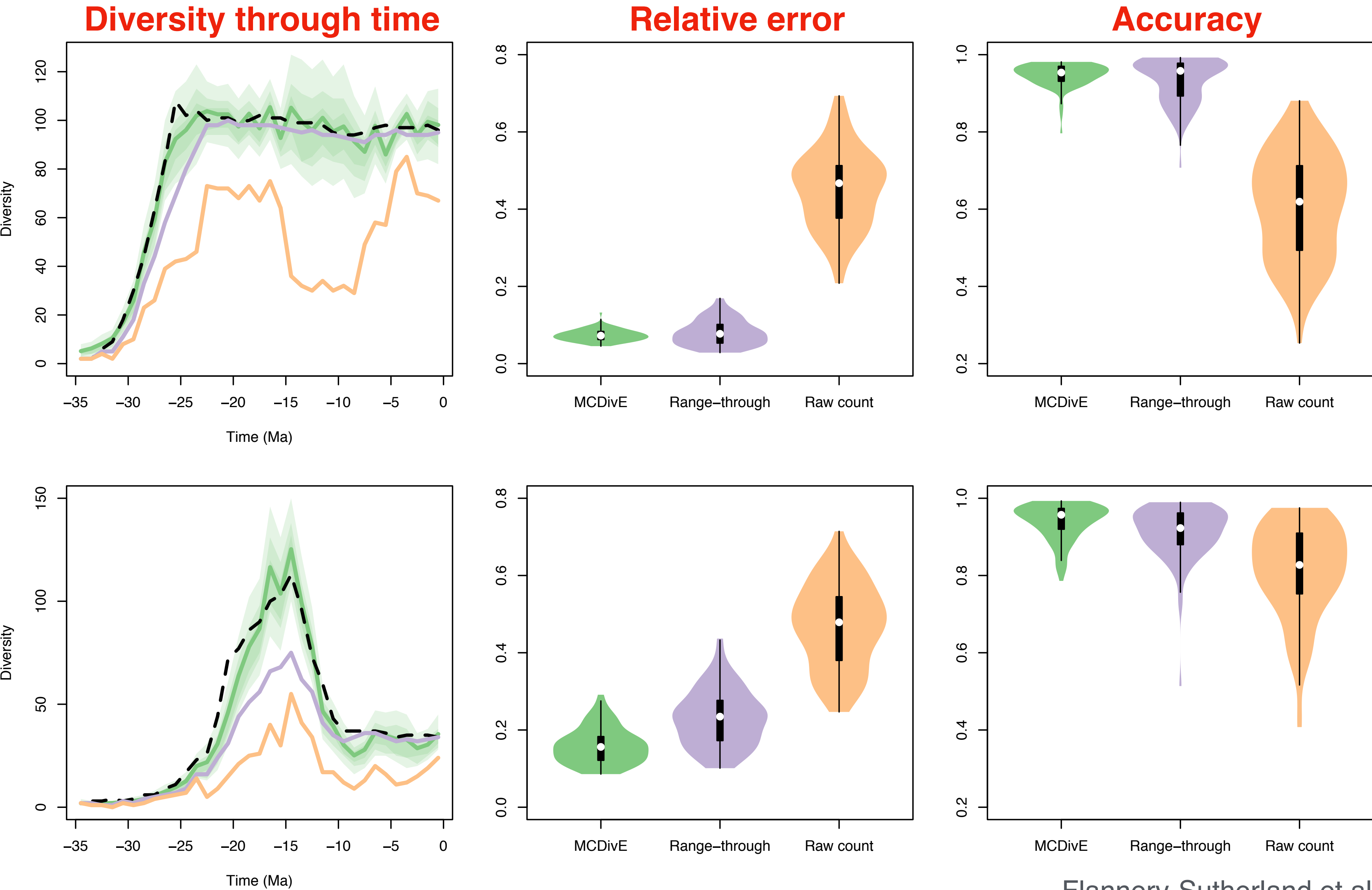
Inferred diversity through time



Likelihood function: Binomial distribution

Prior (smoothing) function: Brownian process

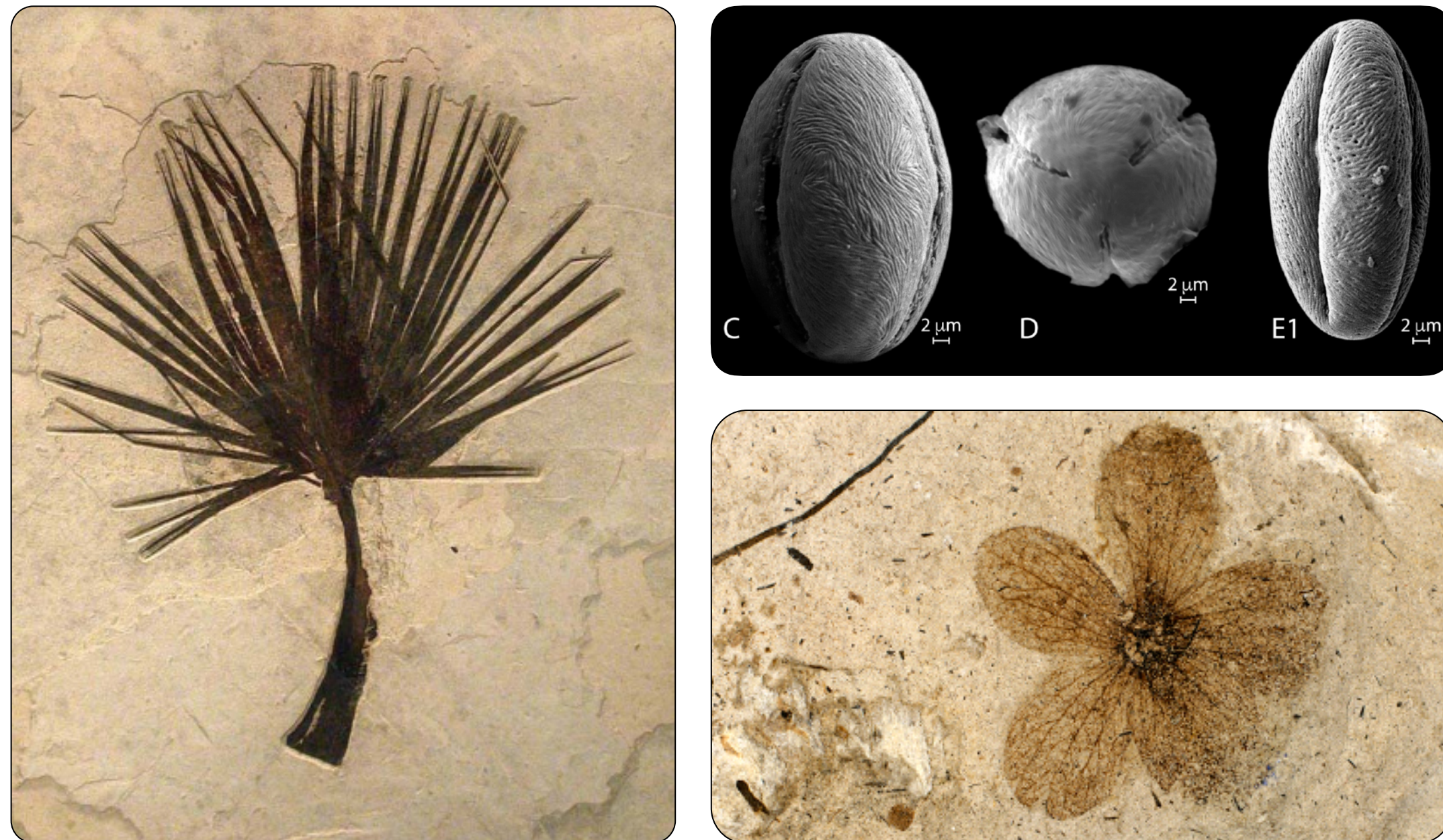
Bayesian estimation of diversity trajectories – mcmcDivE



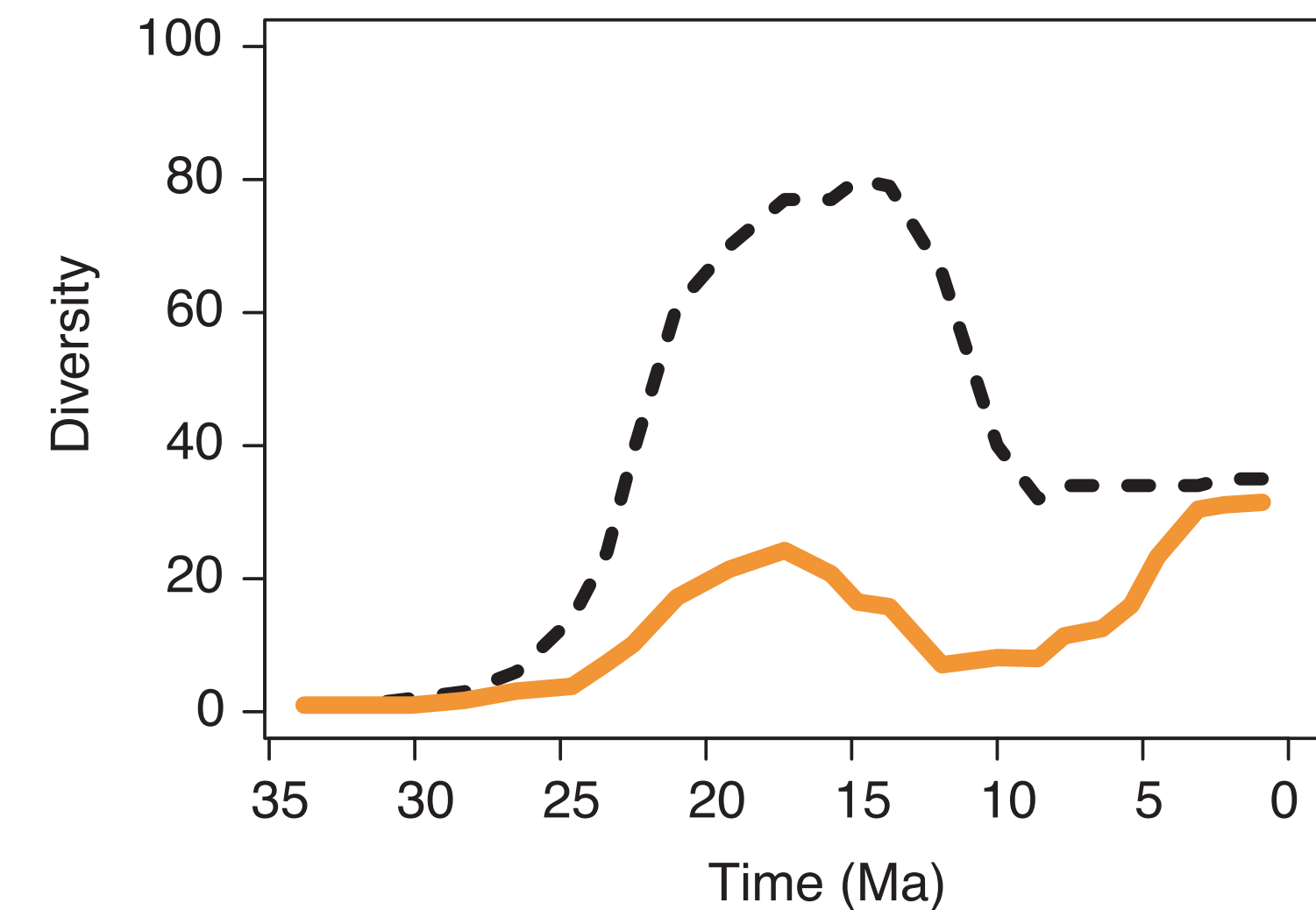
Flannery-Sutherland et al. 2022 Nature Comm

Estimating biodiversity through time from fossil data

Evidence of past biodiversity in the fossil record



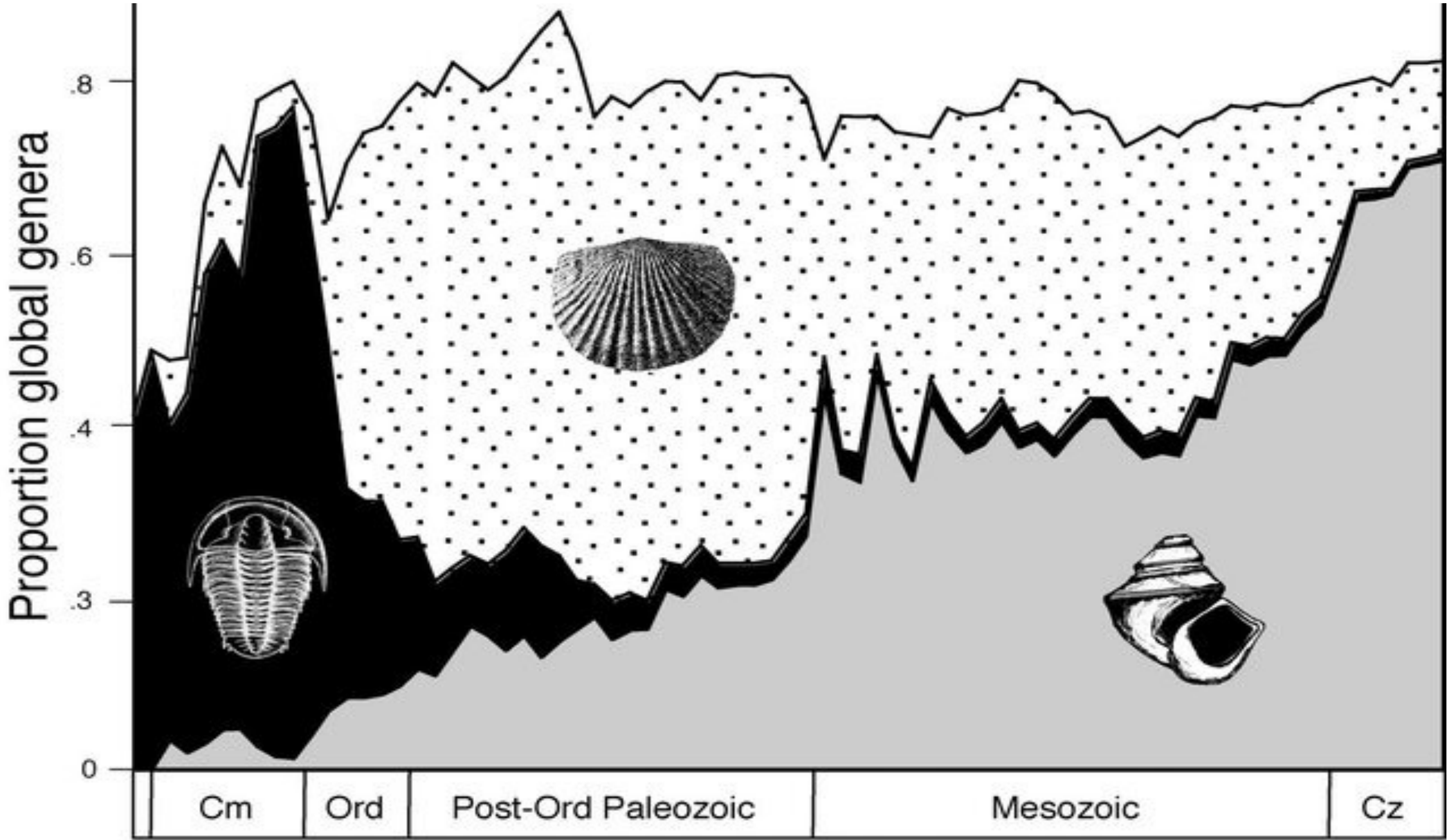
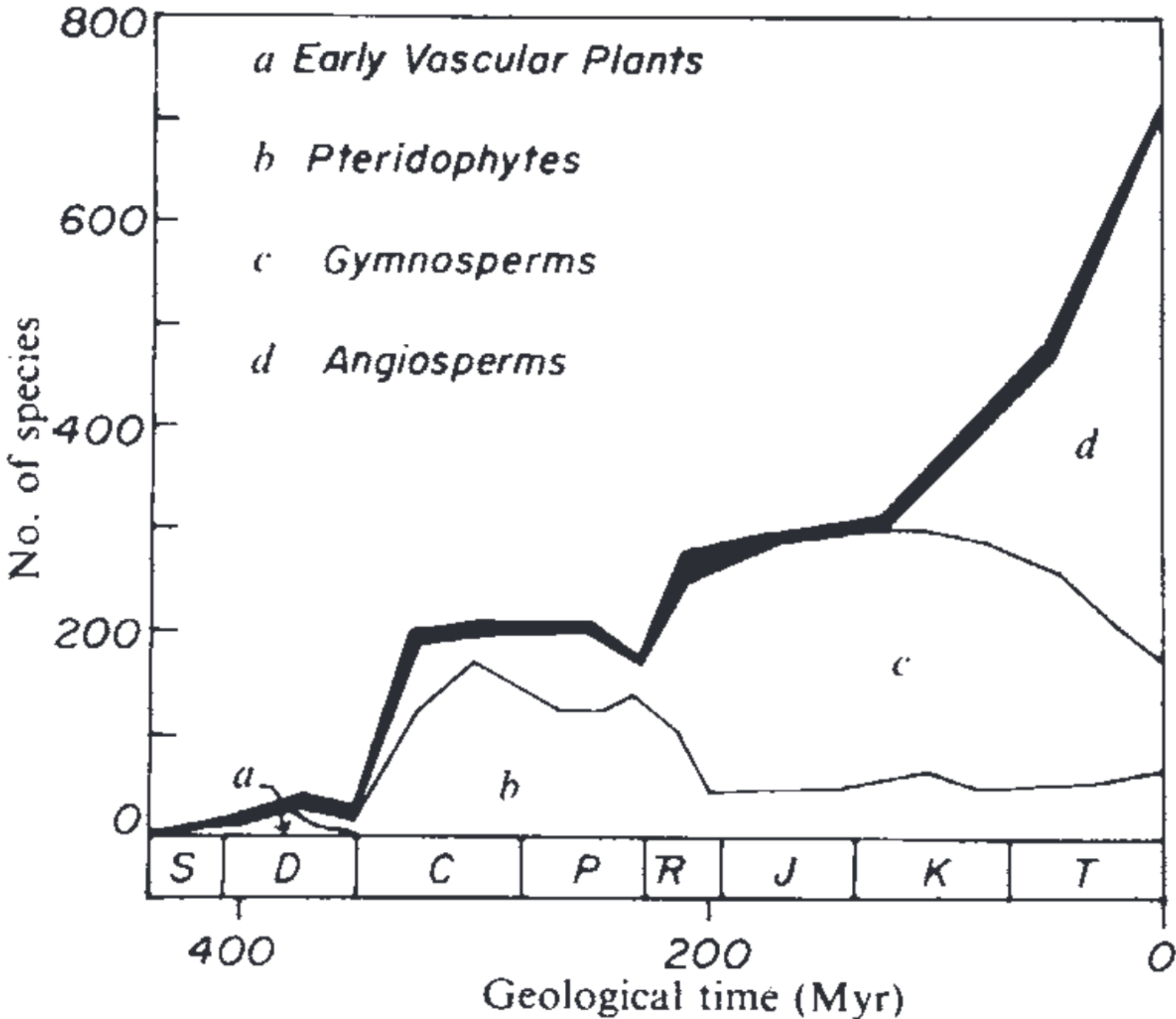
Observed vs true diversity through time



Non-random biases at taxonomic, temporal, and spatial scales 🤖

Available methods are based on rarefaction or simple models assuming random sampling

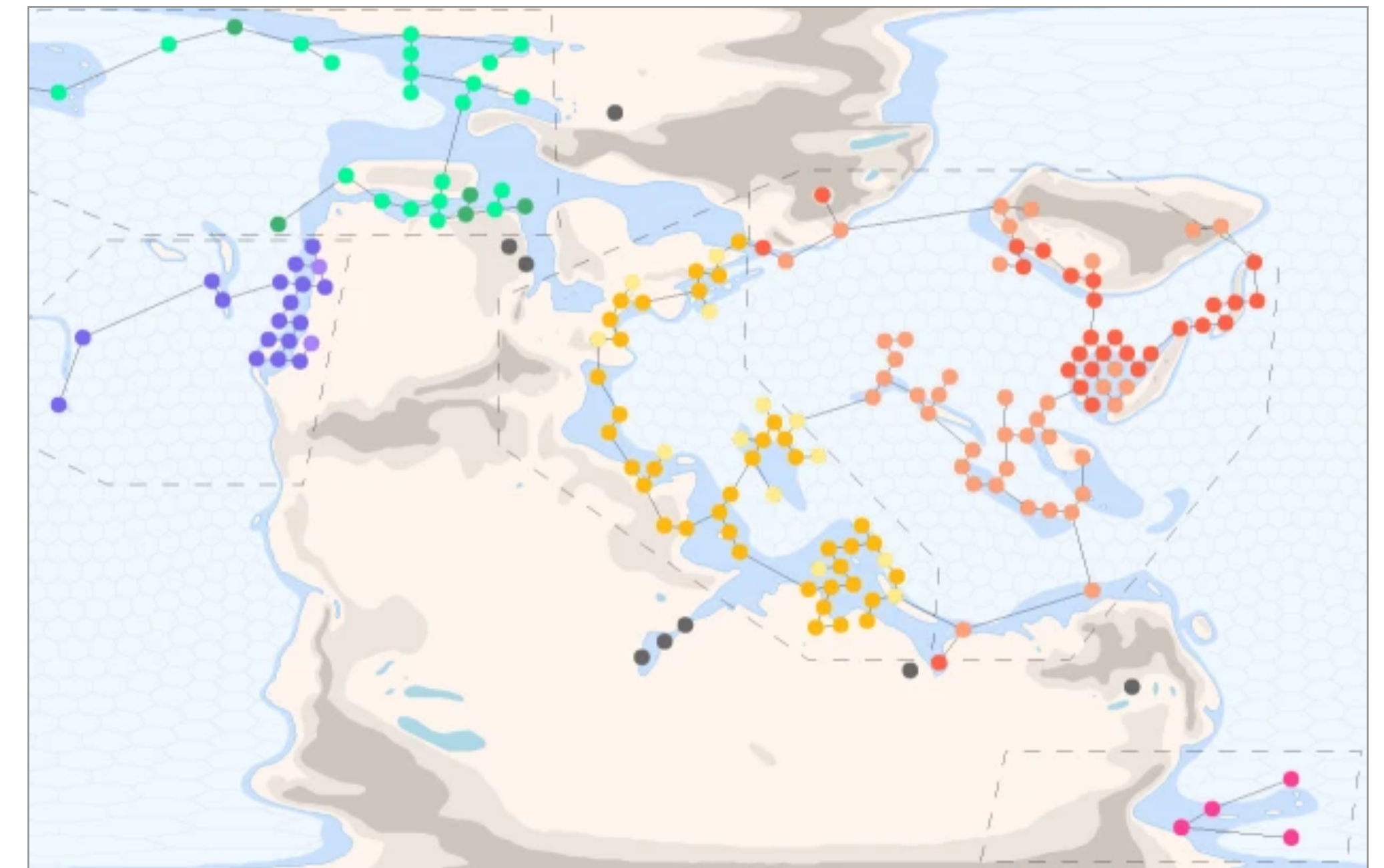
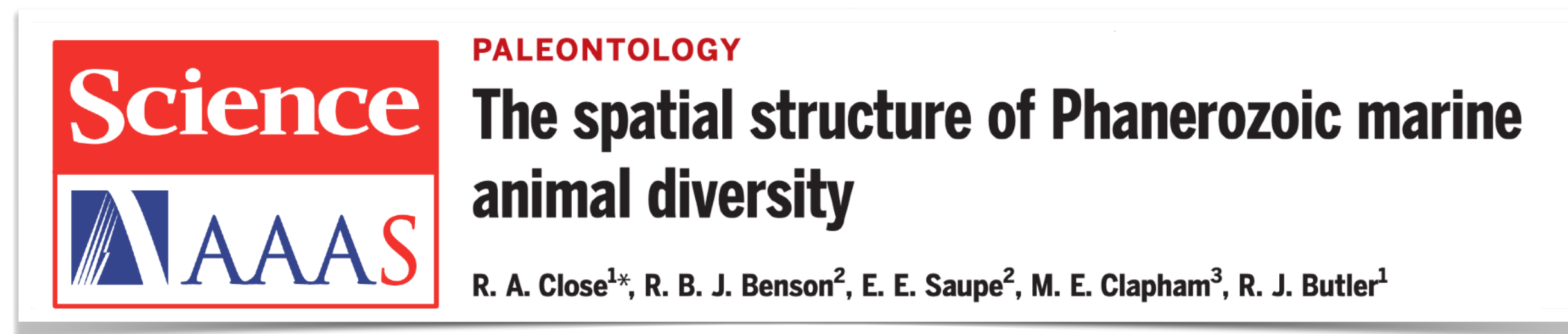
Inferring biodiversity dynamics from the fossil record



Niklas et al. 1983 Nature; Sepkoski 1981 Paleobiology

Problems in estimating biodiversity through time from fossil data

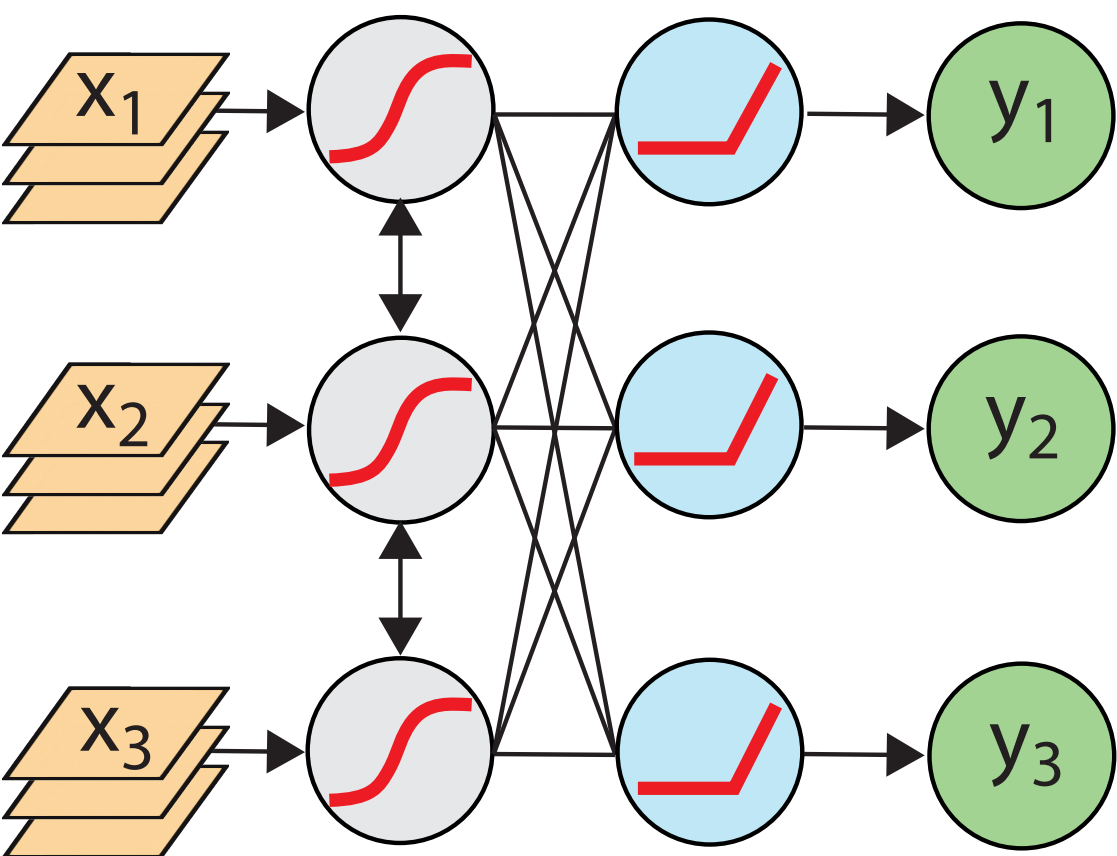
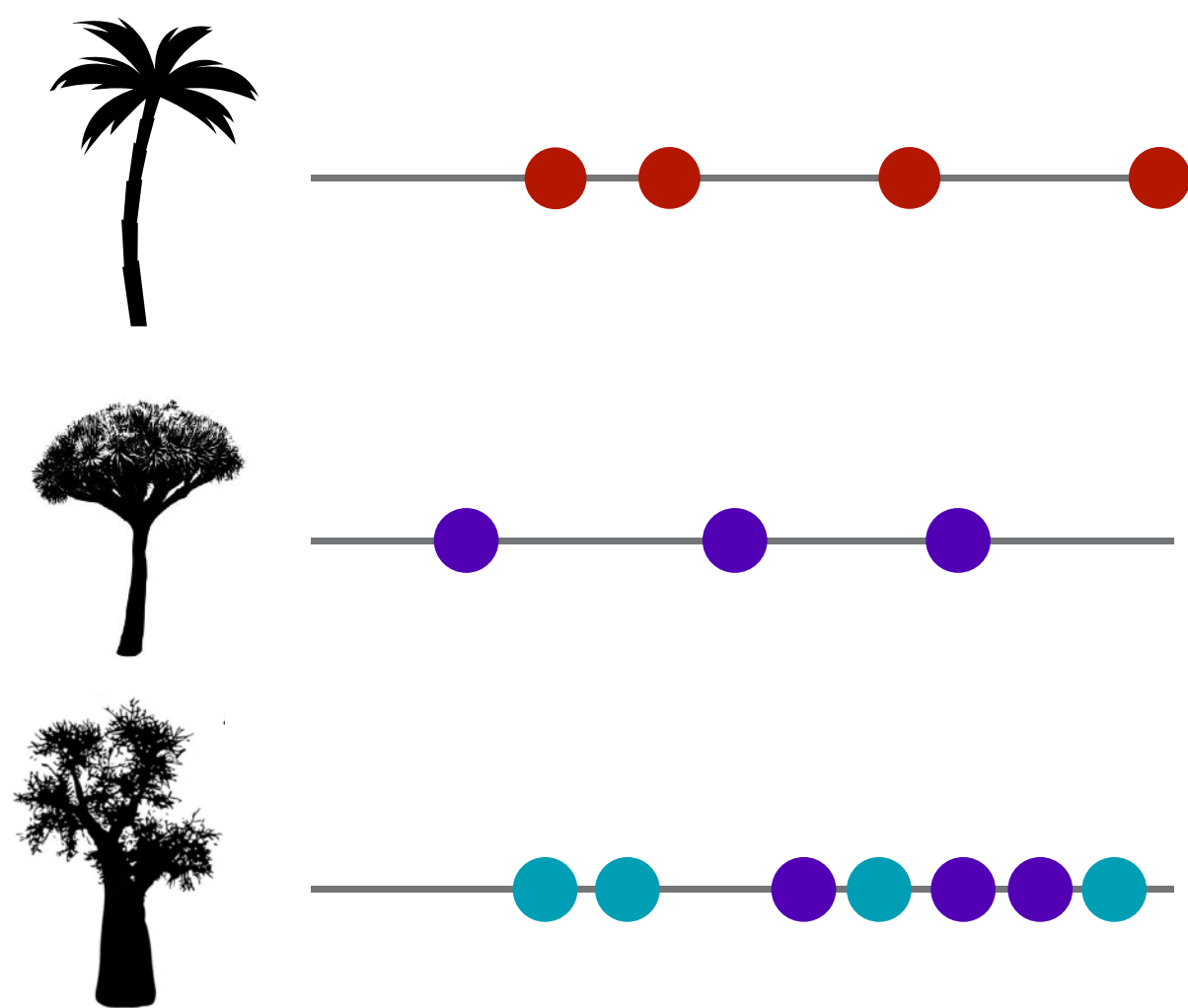
Spatial biases make global biodiversity patterns unidentifiable using the current methods



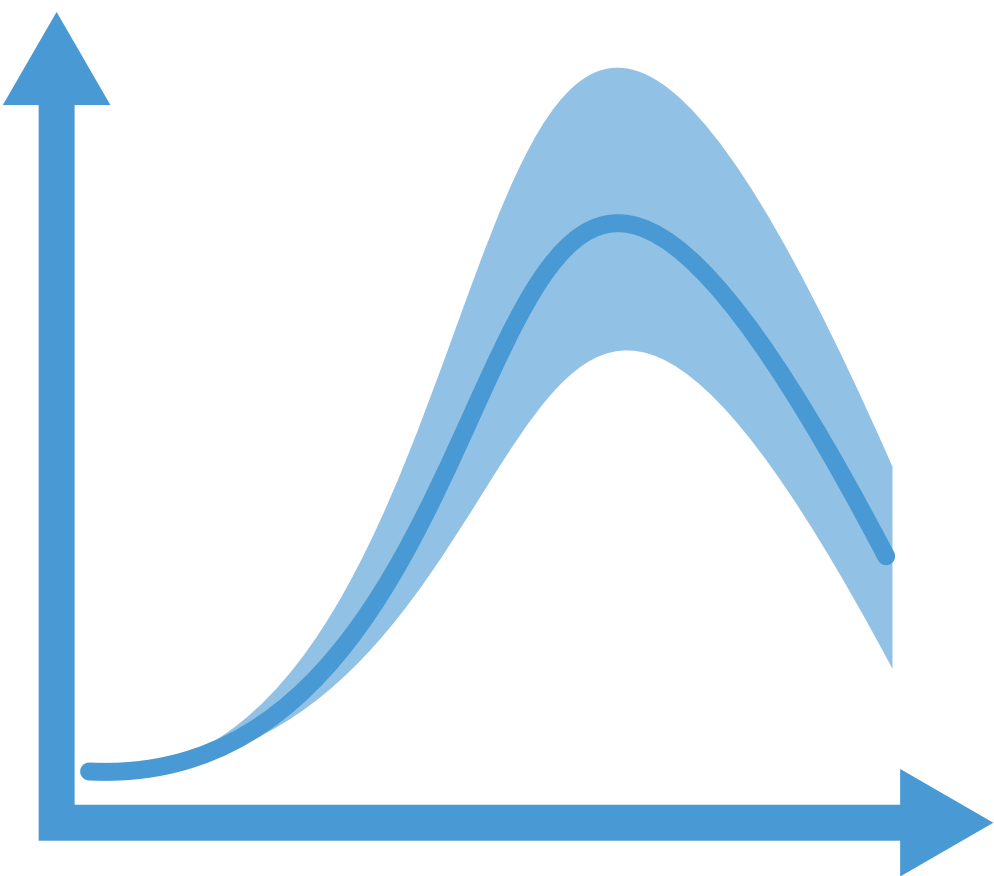
Alroy 2010 Science; Starrfelt & Liow 2016 Phil Trans B; Flannery-Sutherland et al. 2022 Nature Comm

DeepDive: deep learning estimation of biodiversity through time from fossil data

Input: fossil distribution in space and time



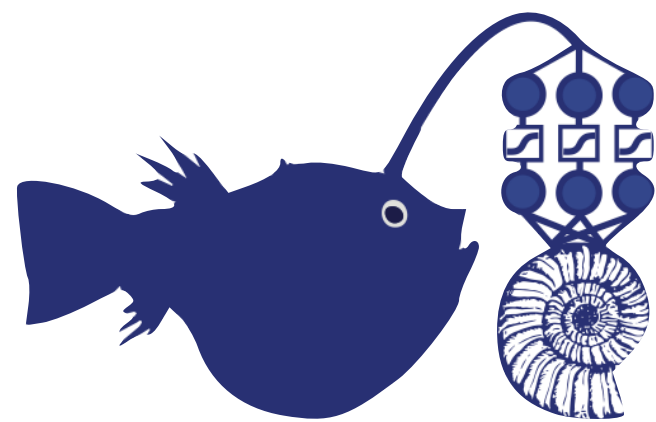
Output: diversity estimates through time



R Cooper



B Allen

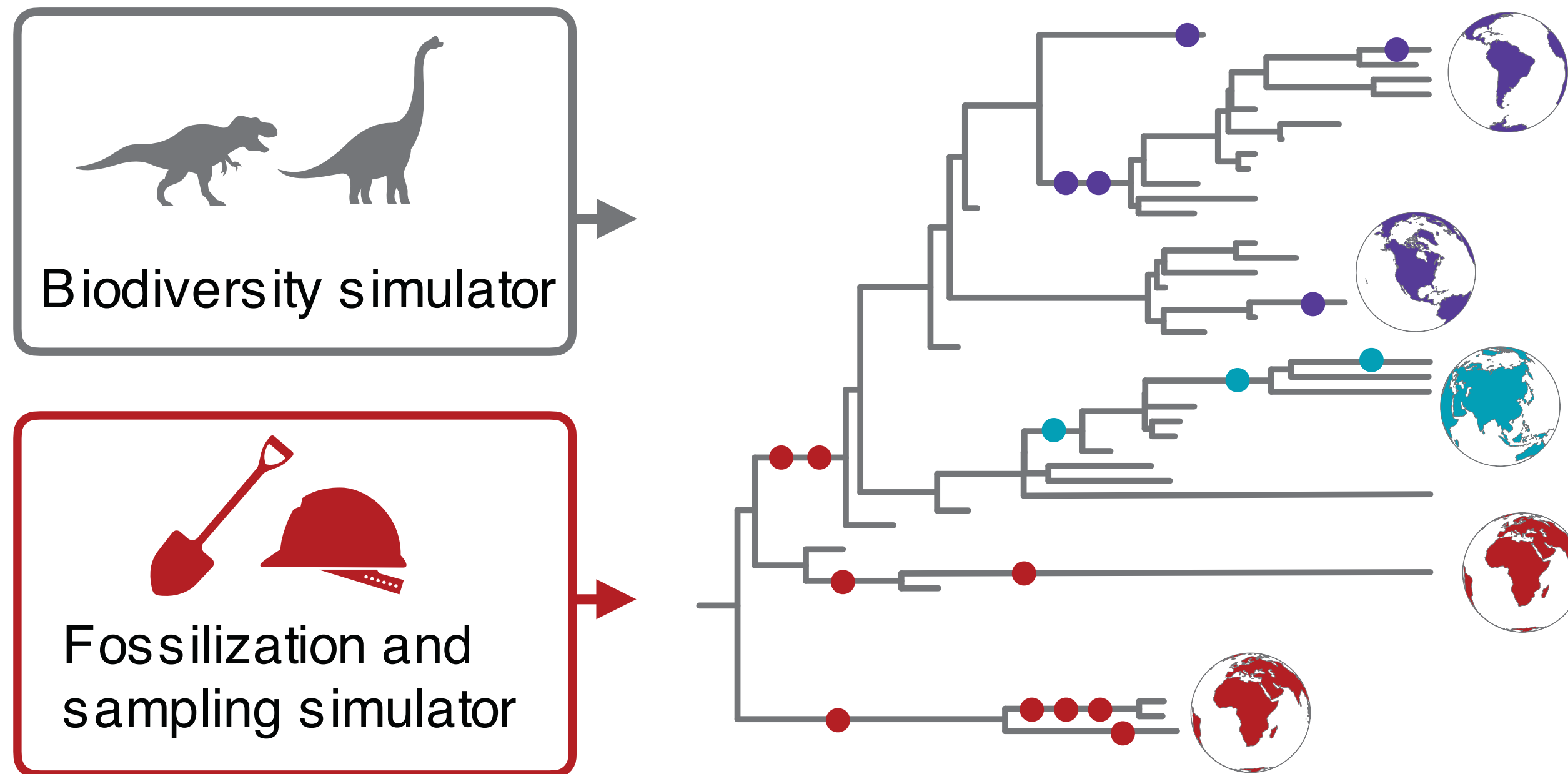


github.com/DeepDive-project

Cooper et al. 2024 Nature Comms
Application note: Cooper et al. 2024 bioRxiv

DeepDive – Deep learning models to estimate Diversity trajectories

Mechanistic model of species diversification and extinction in time and space

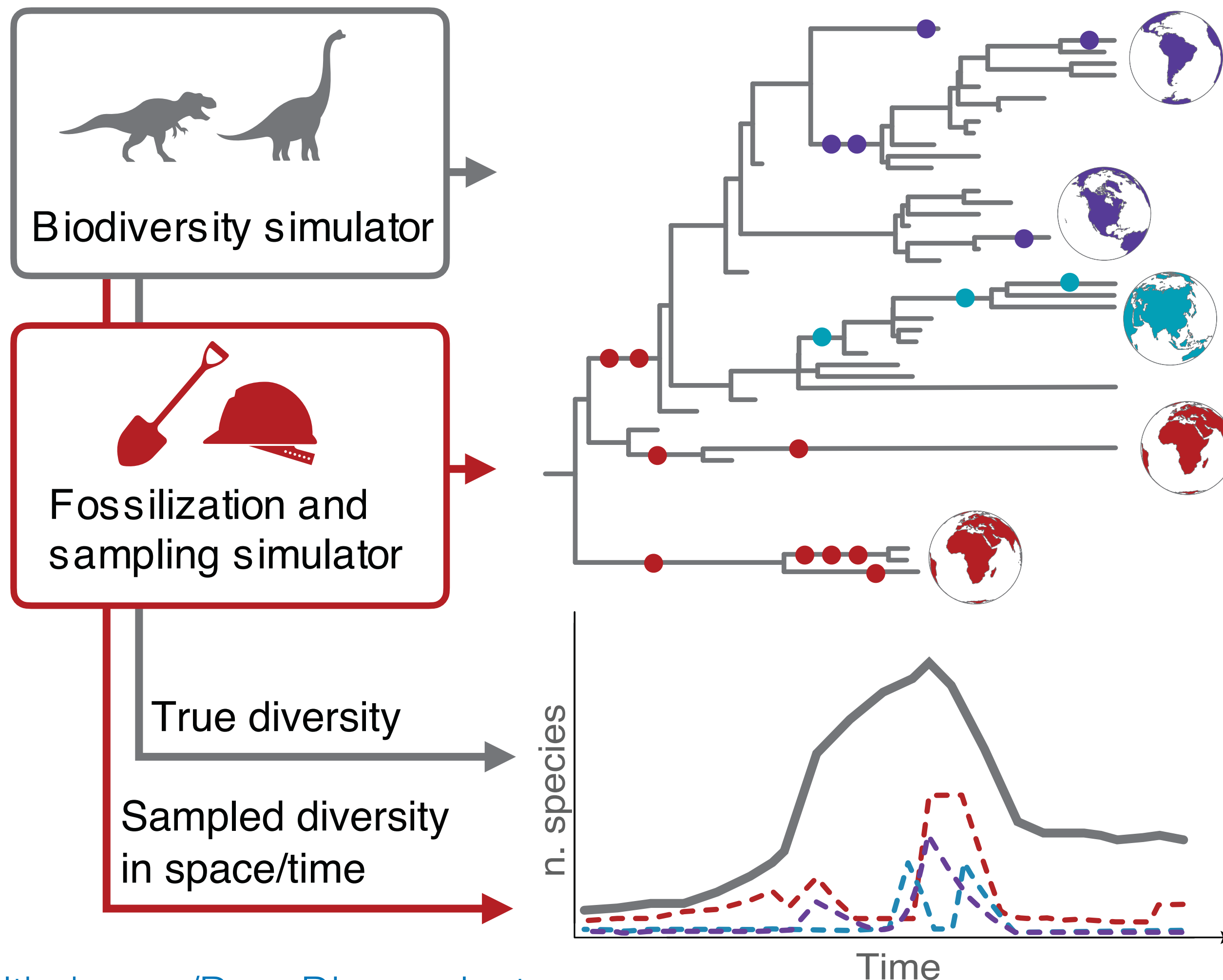


Clade evolution based on spatially-explicit birth-death processes

Simulated fossil data with spatial, temporal, and taxonomic biases

DeepDive – Deep learning models to estimate Diversity trajectories

Mechanistic model of species diversification and extinction in time and space

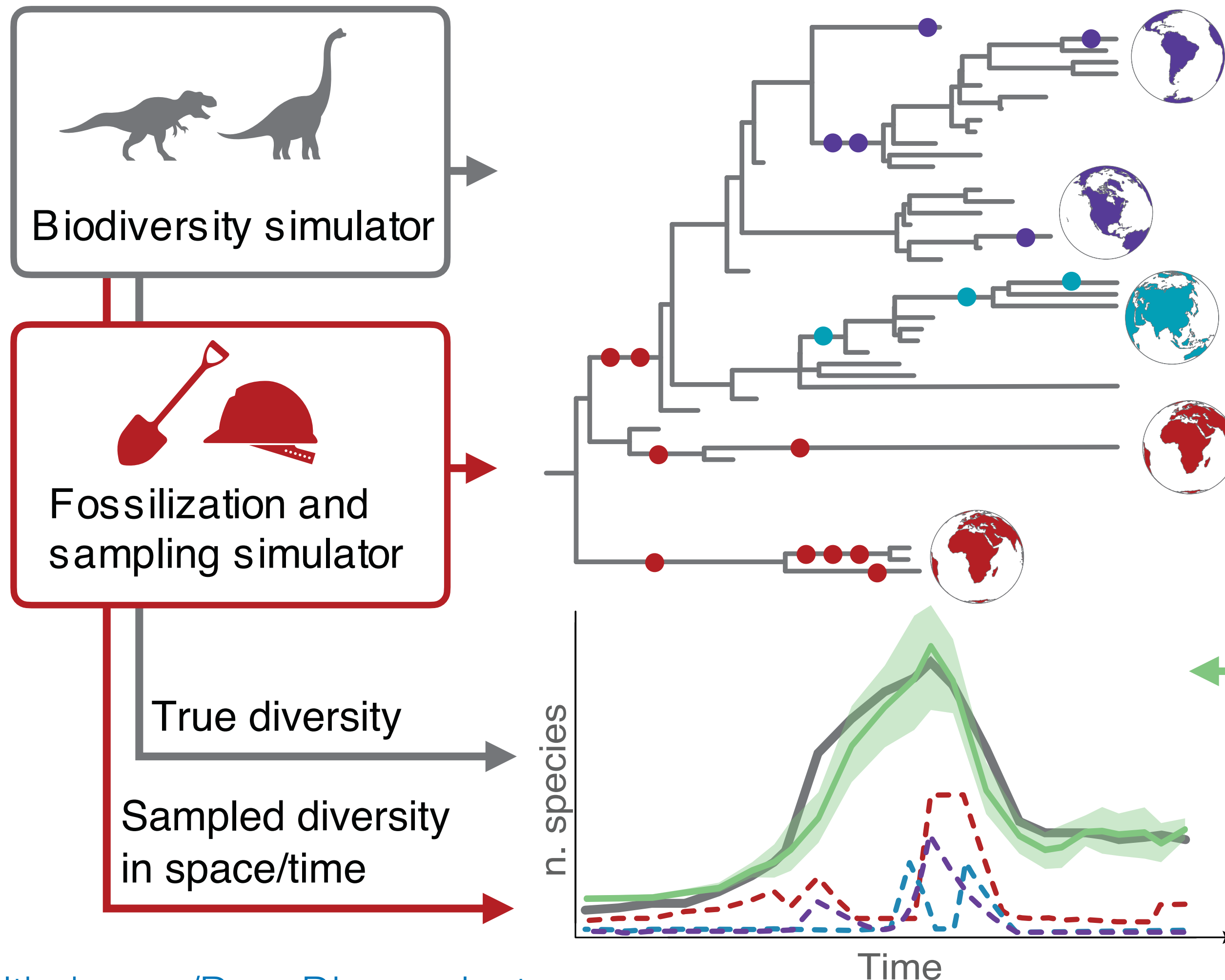


Clade evolution based on spatially-explicit birth-death processes

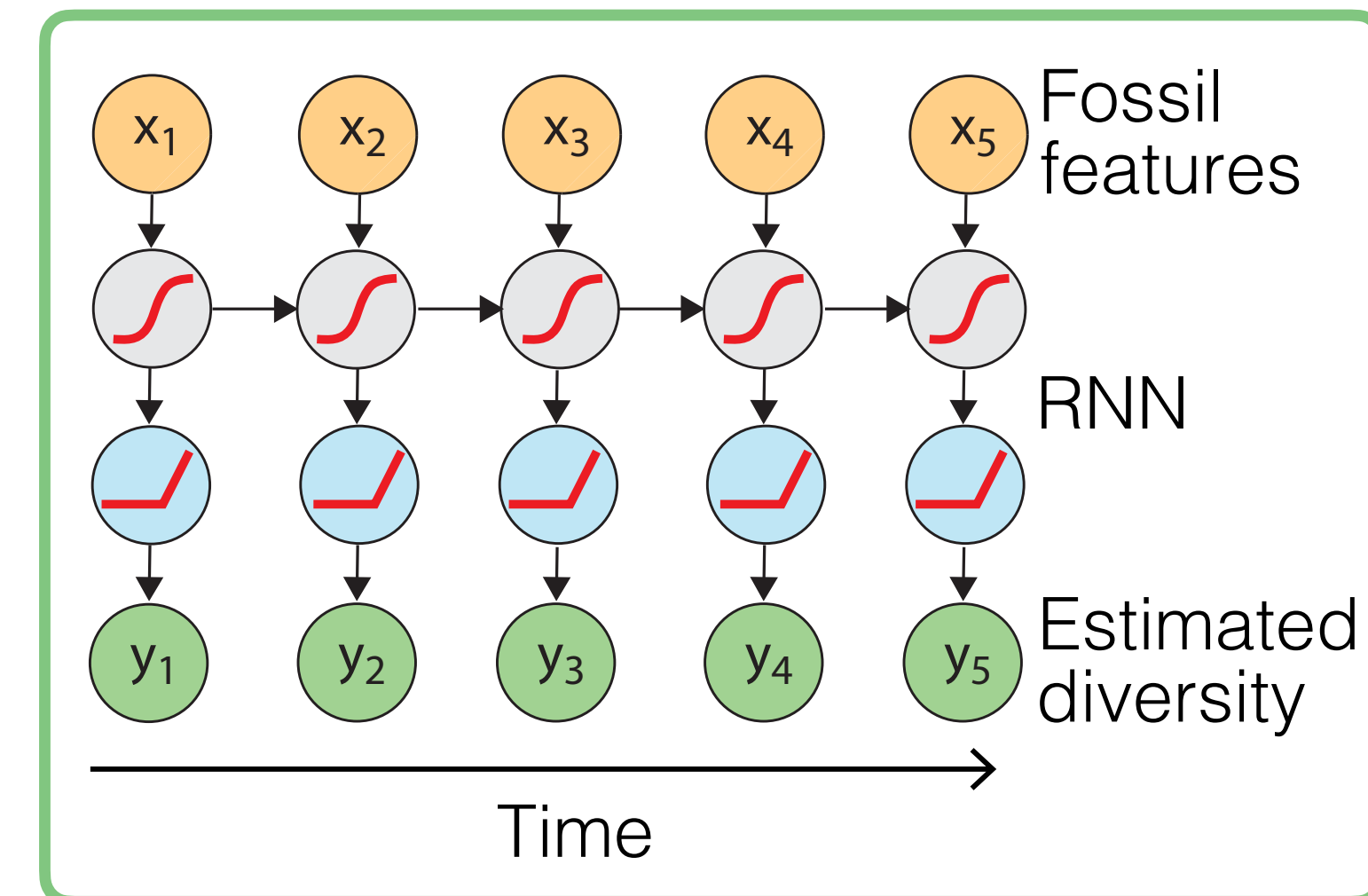
Simulated fossil data with spatial, temporal, and taxonomic biases

Supervised deep learning models to estimate diversity trajectories

Mechanistic models of speciation, extinction, and fossilization in time and space



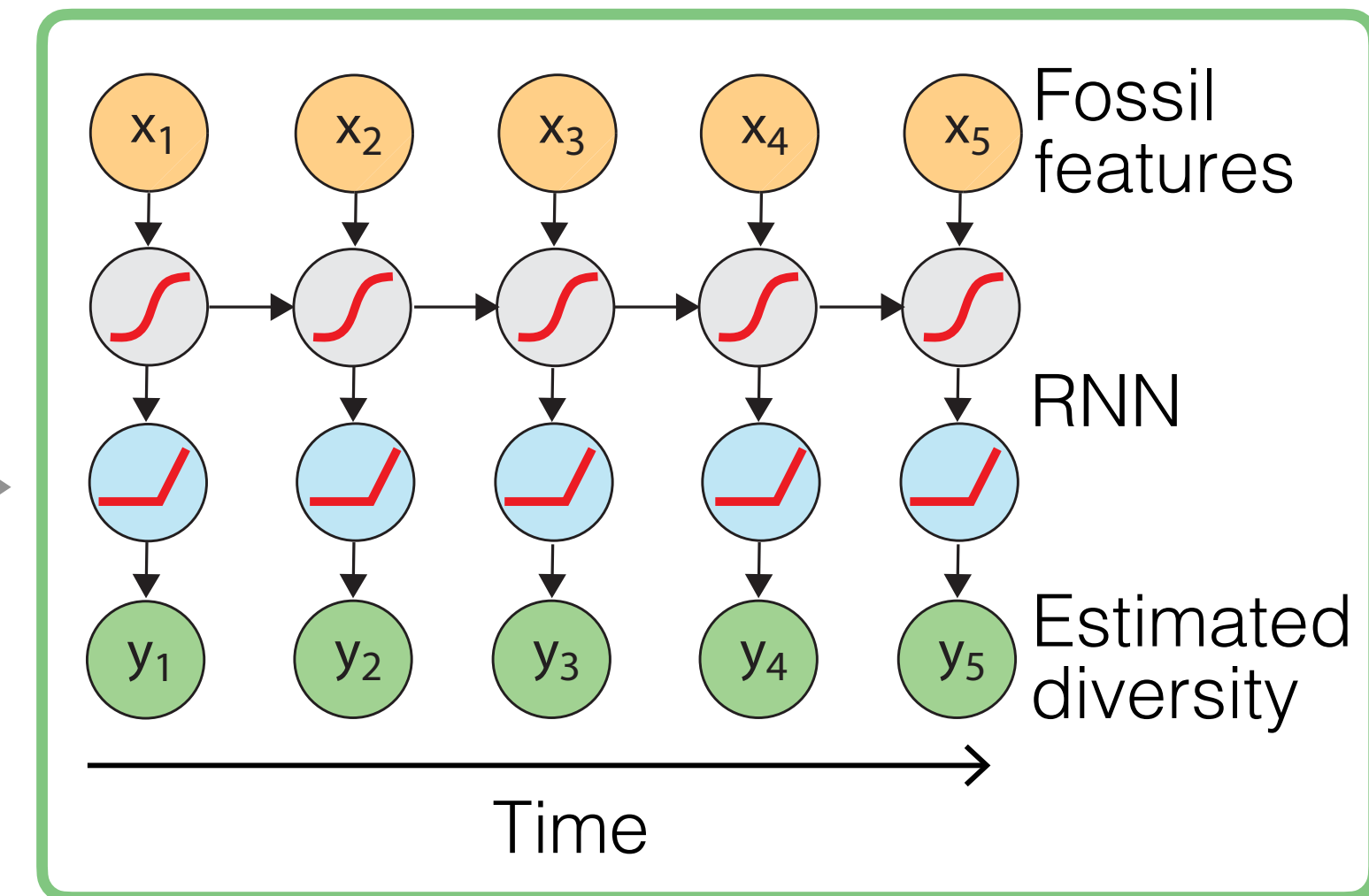
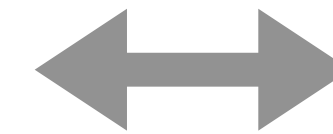
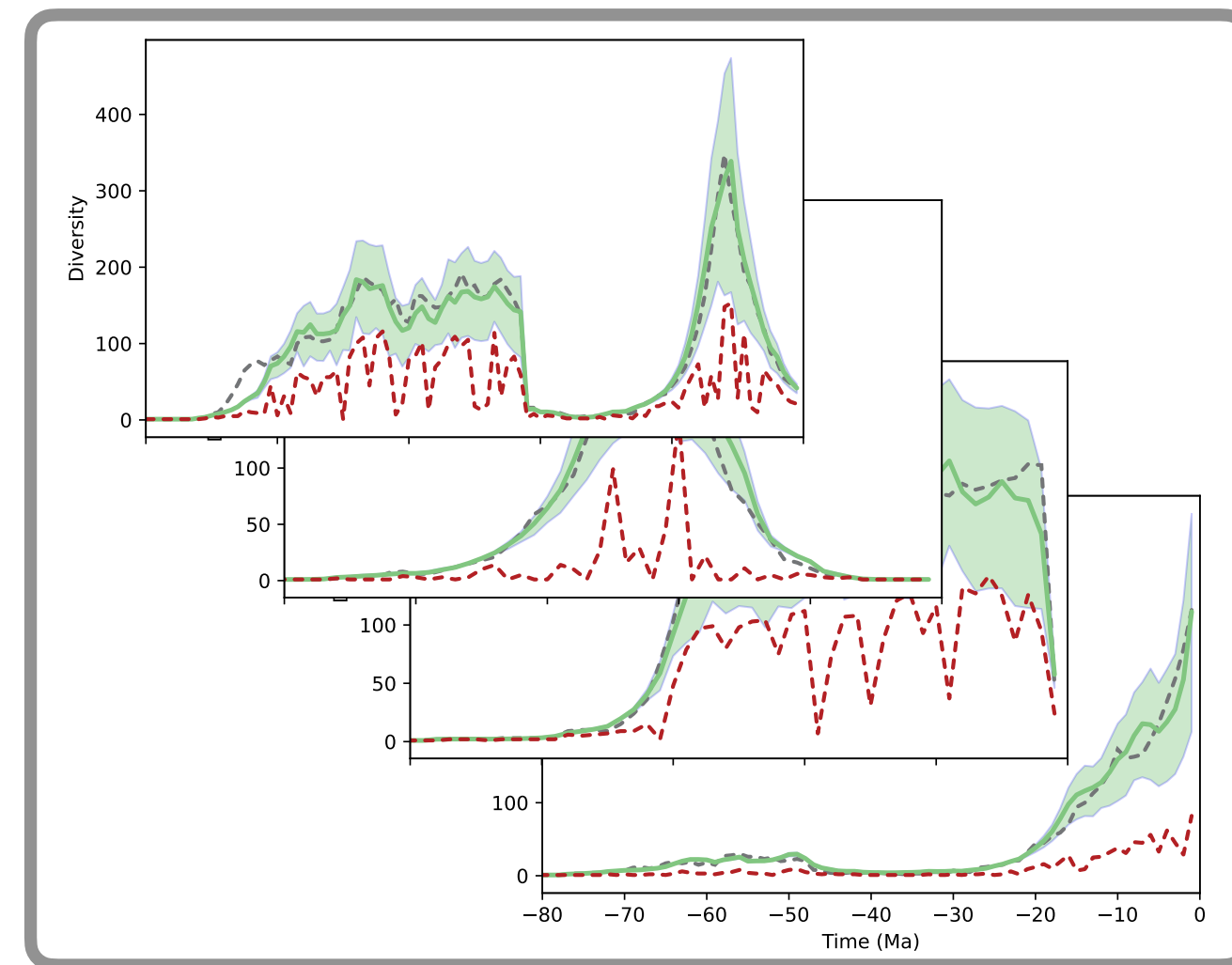
Deep learning model predicting diversity time series from fossil features



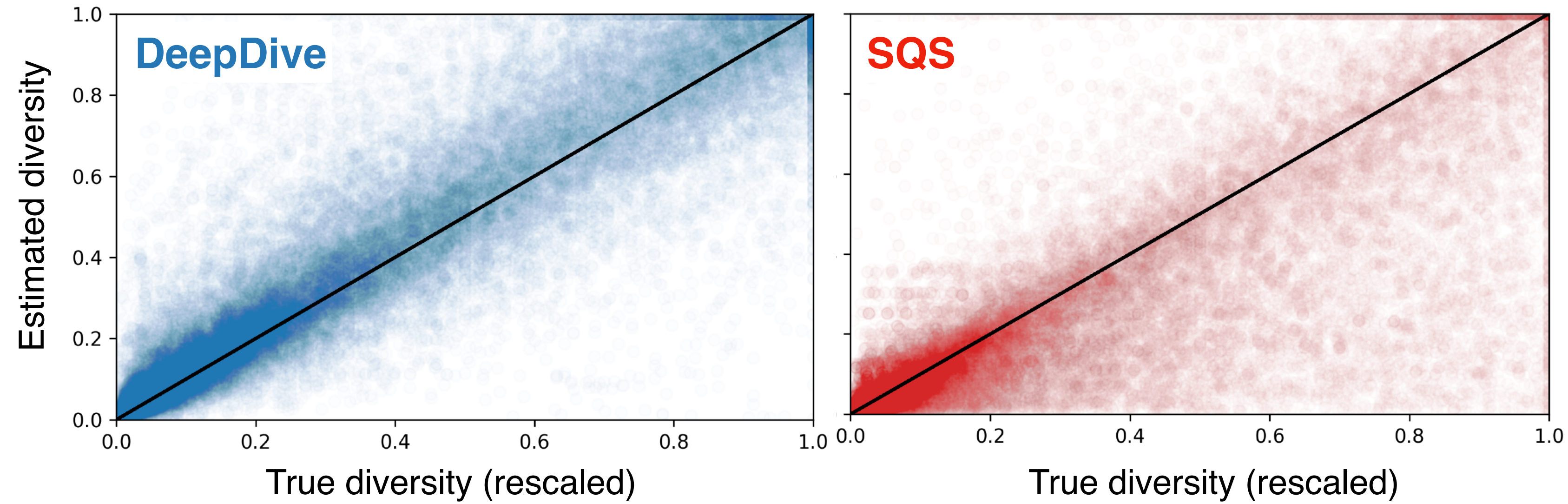
DeepDive – Deep learning models to estimate Diversity trajectories

Model optimization (training) and validation

Samples from 'prior' distributions of speciation, extinction, migration, preservation rates



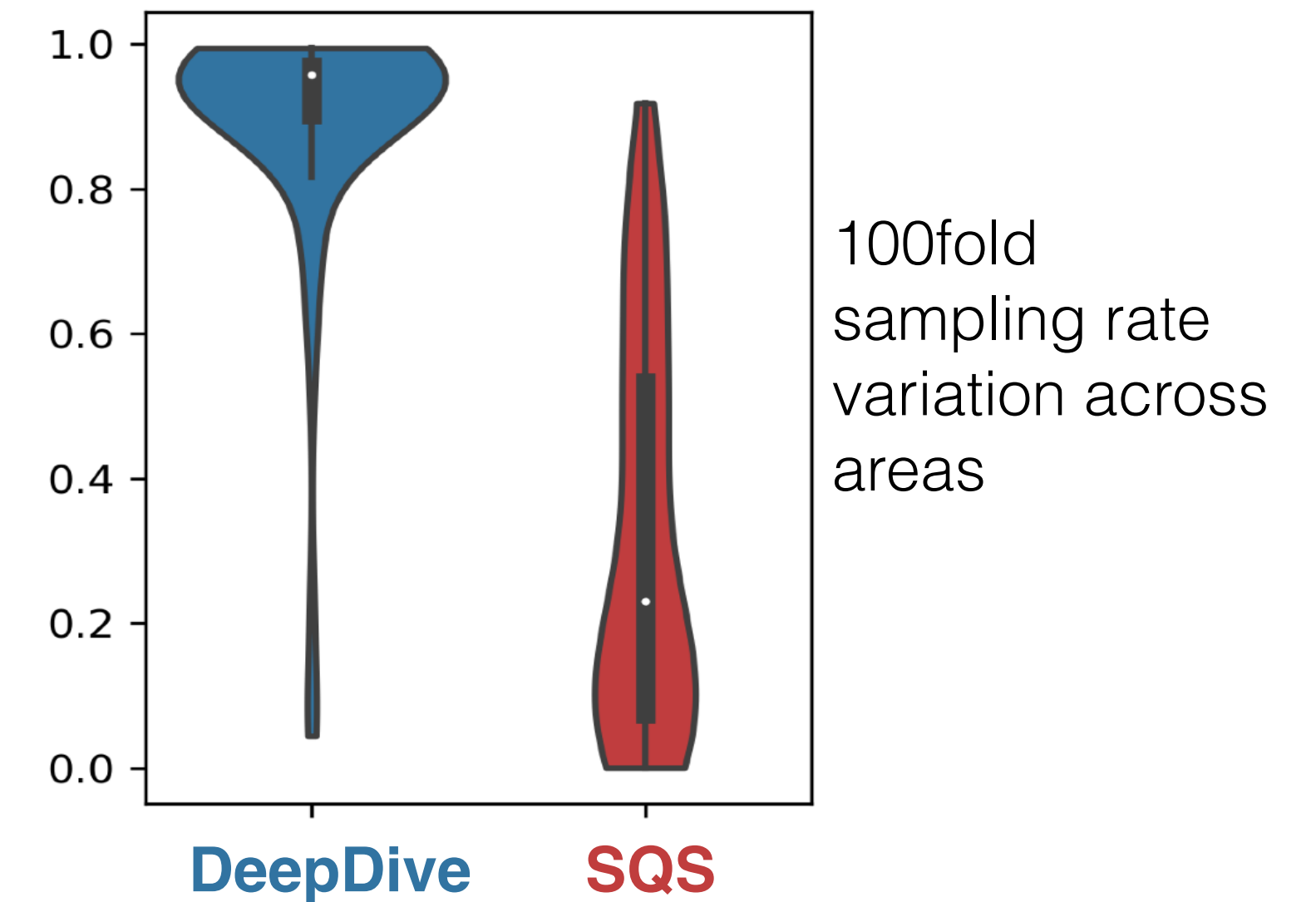
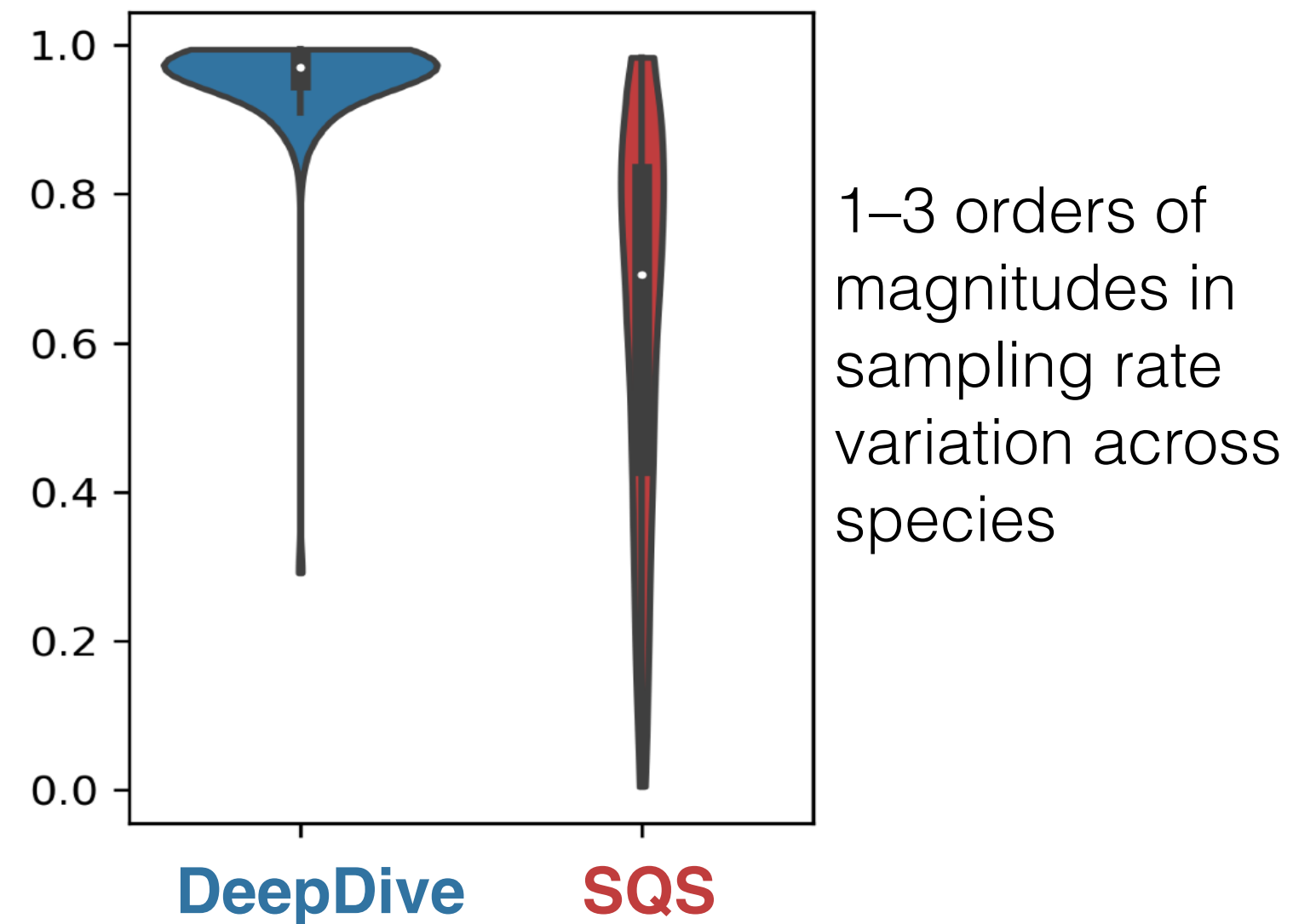
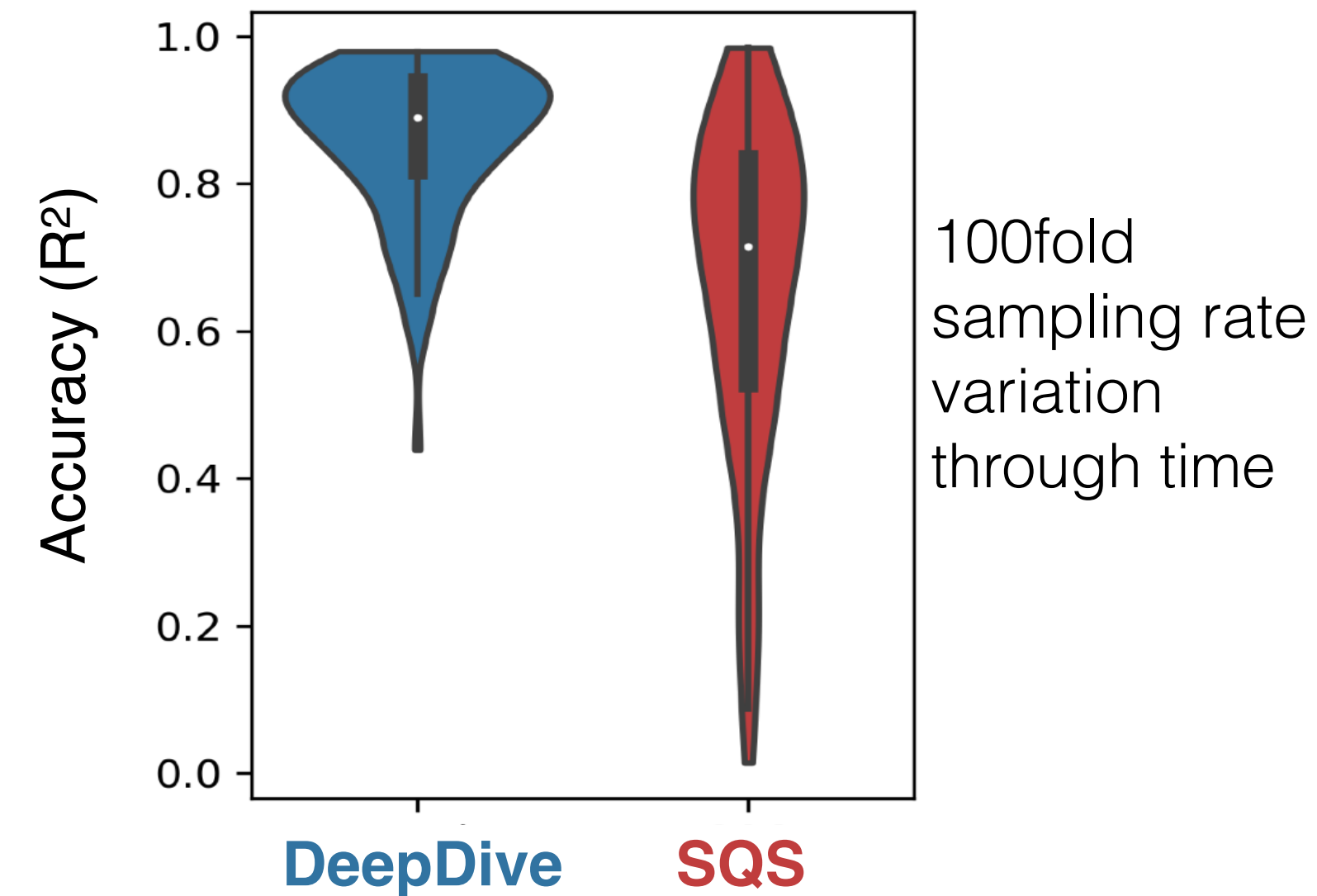
DeepDive performance



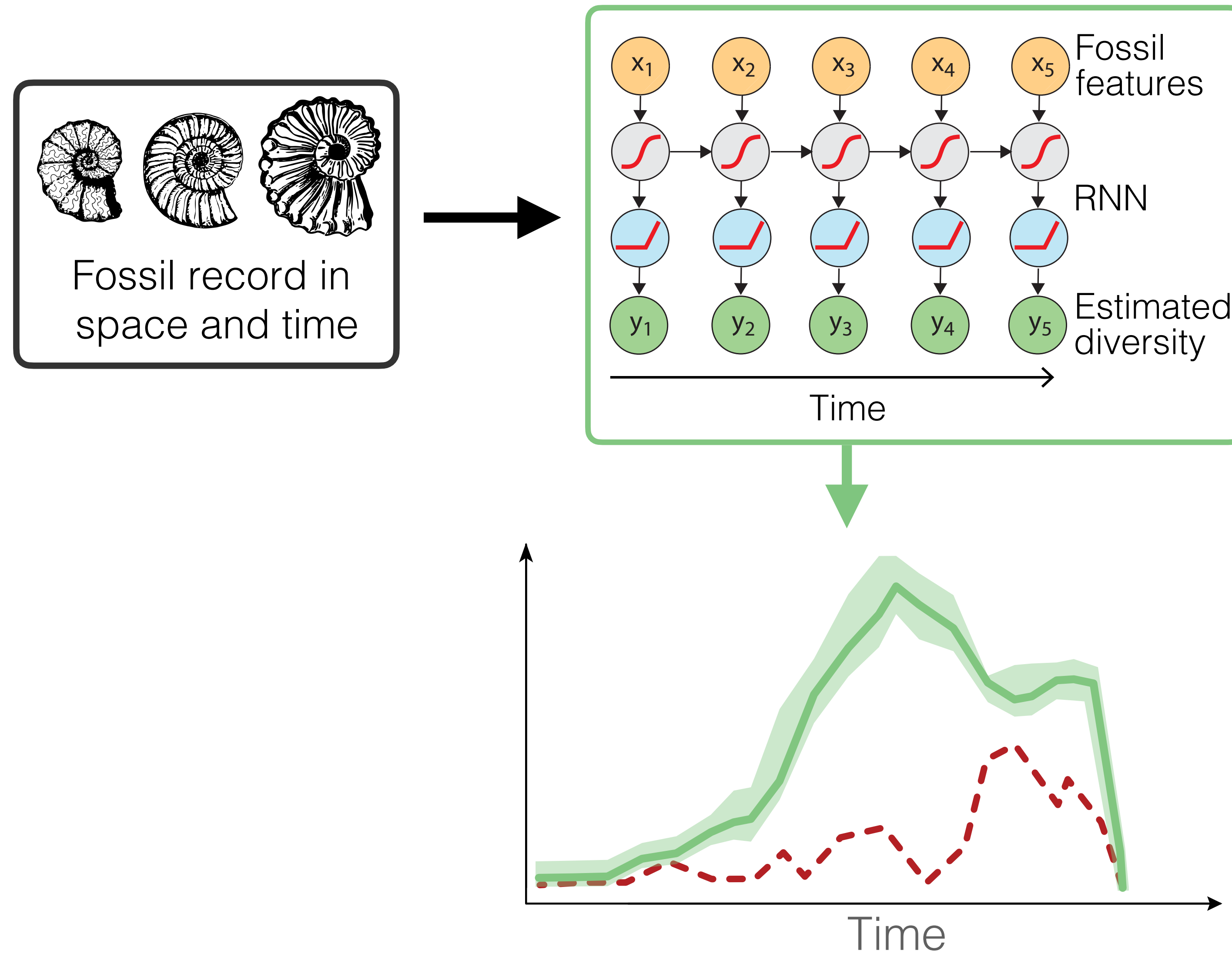
Strong temporal bias

Strong taxonomic bias

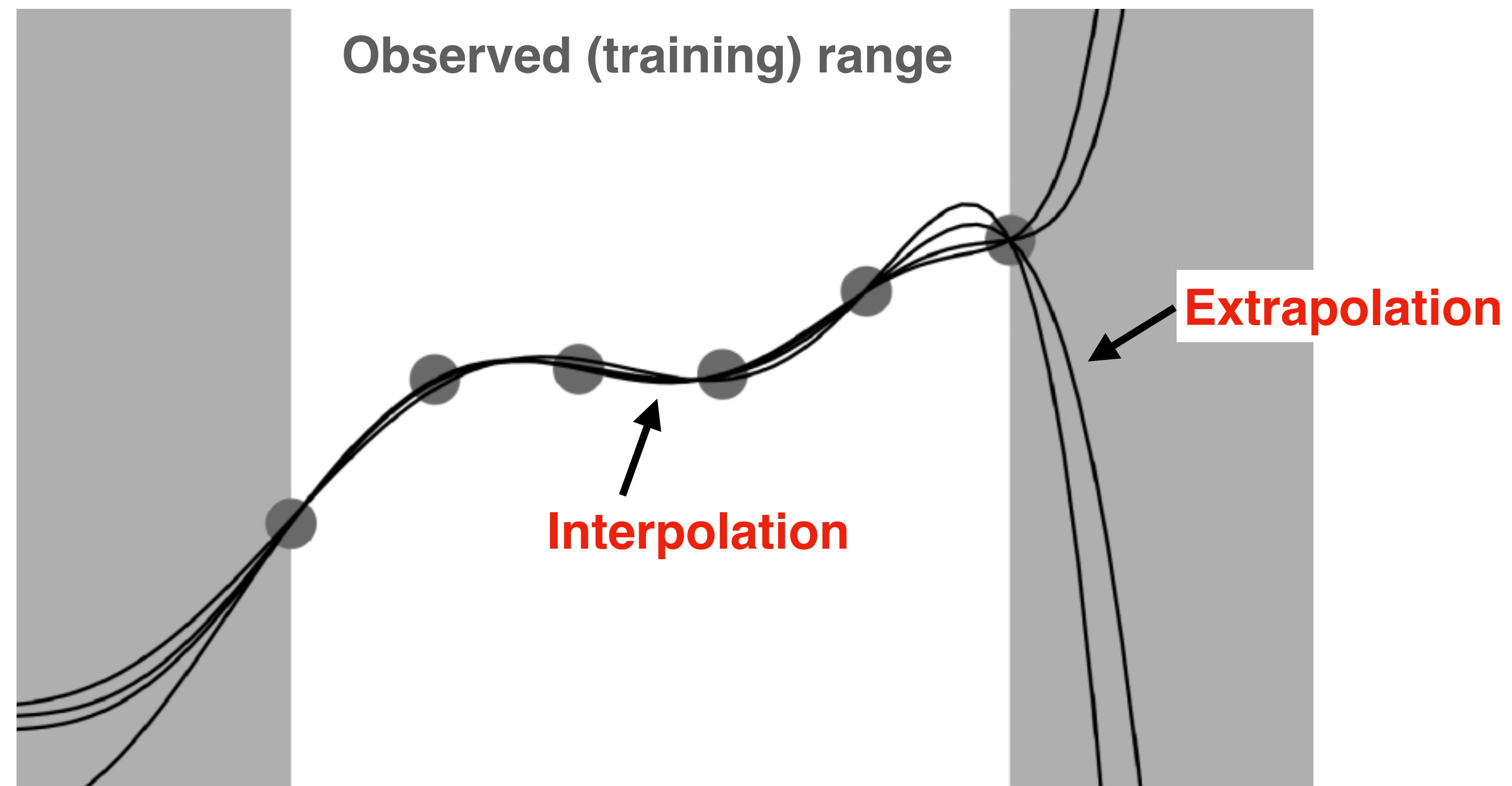
Strong spatial bias



DeepDive – Predictions of empirical data



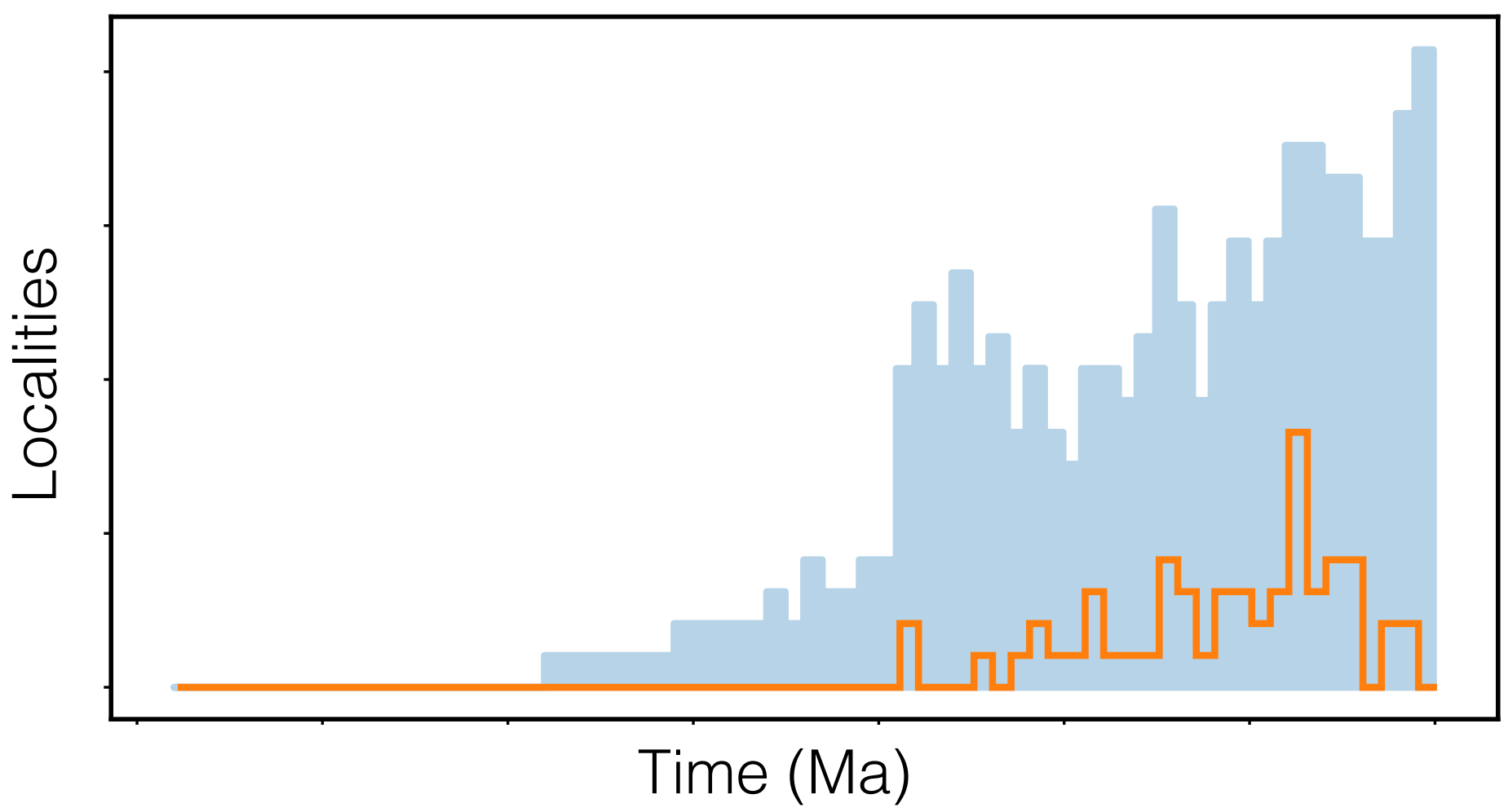
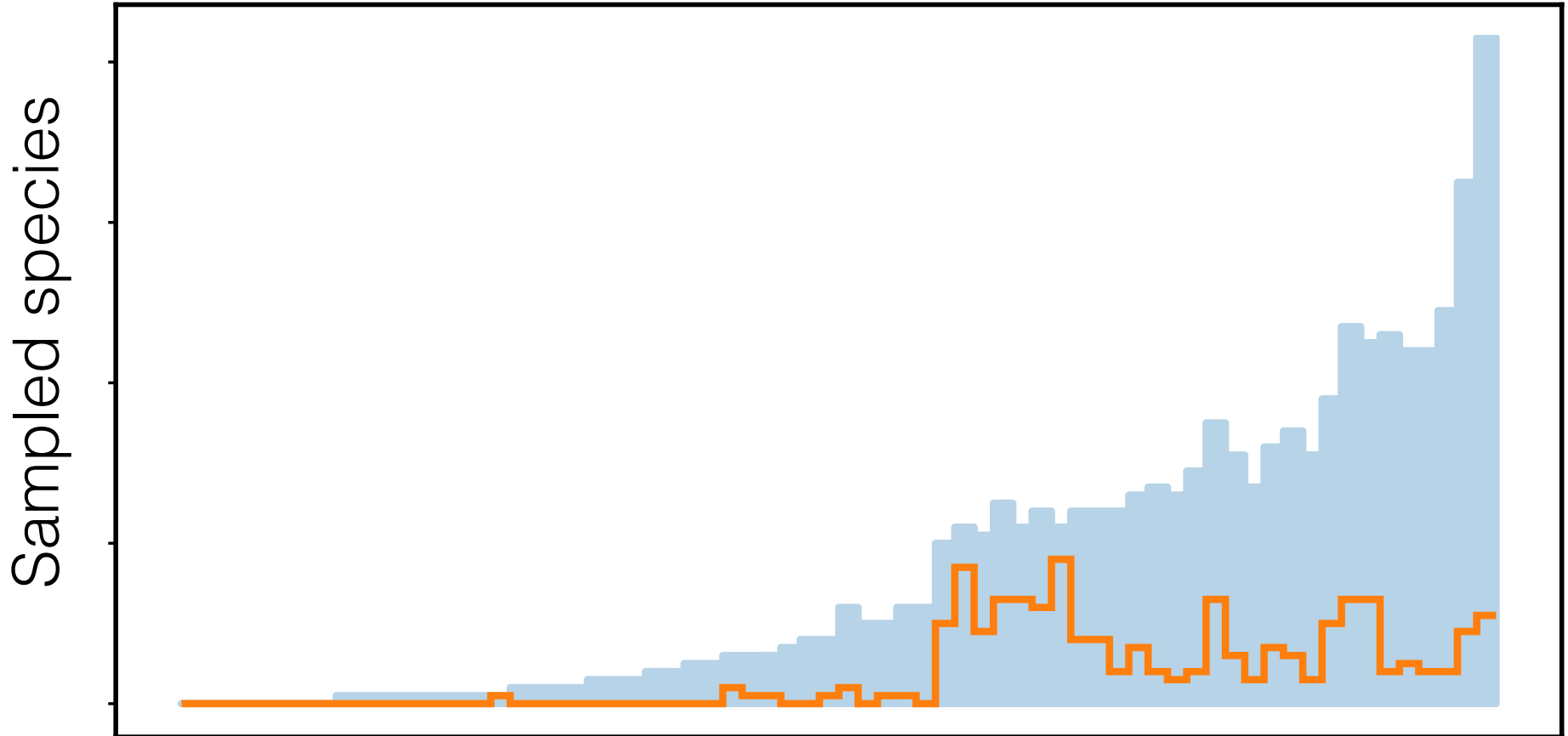
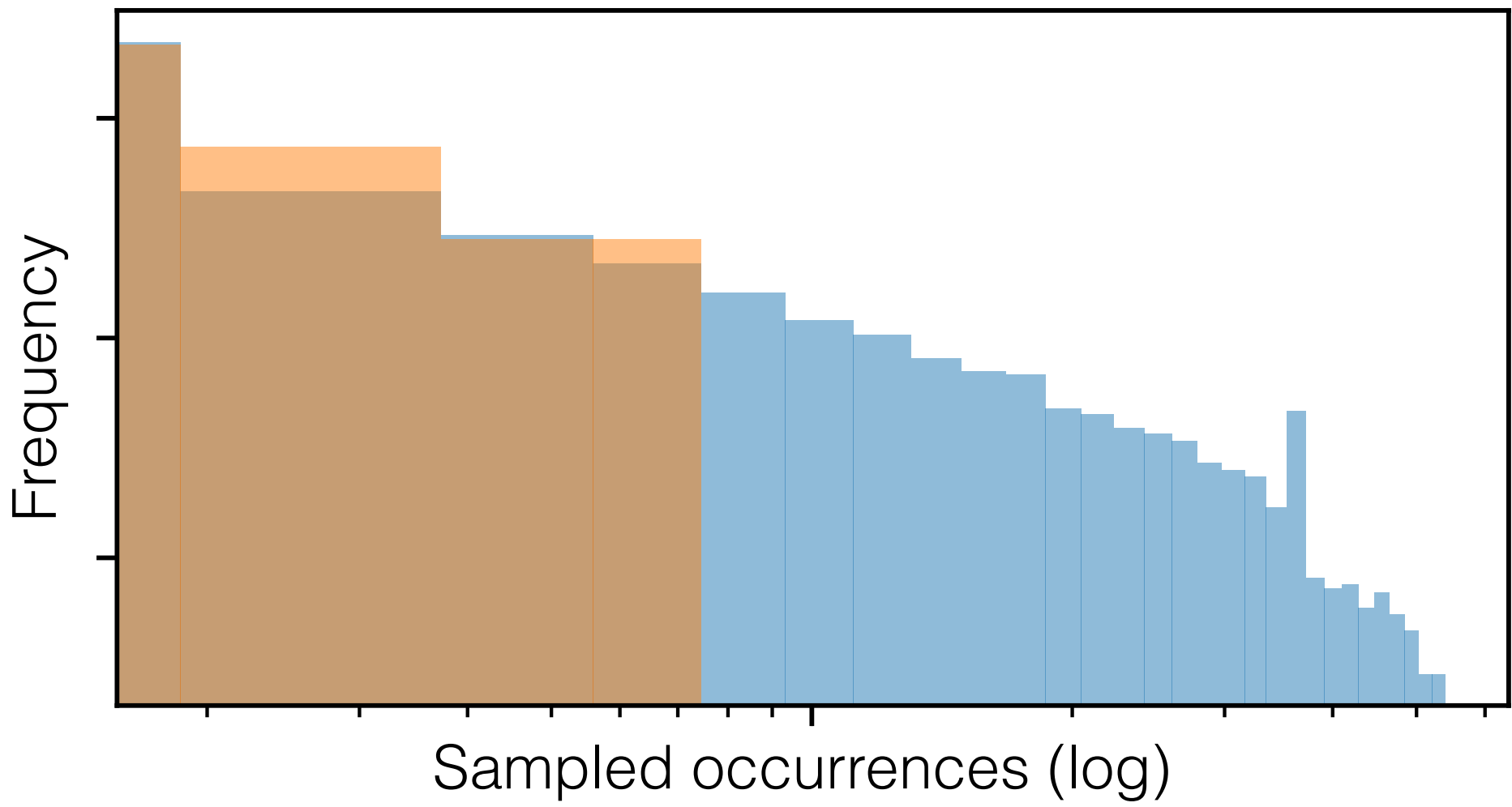
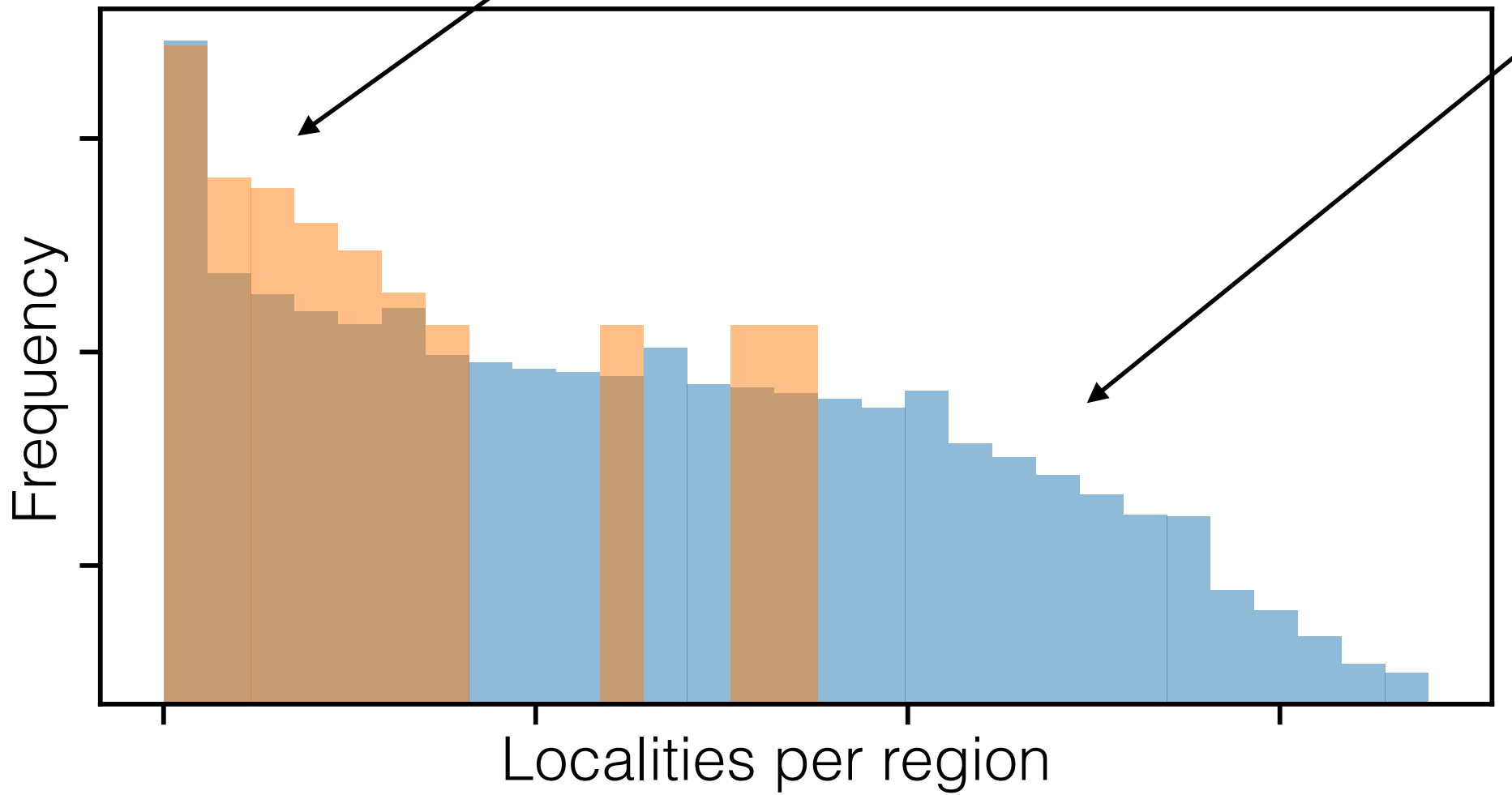
Caveat: the training simulations must resemble the true data



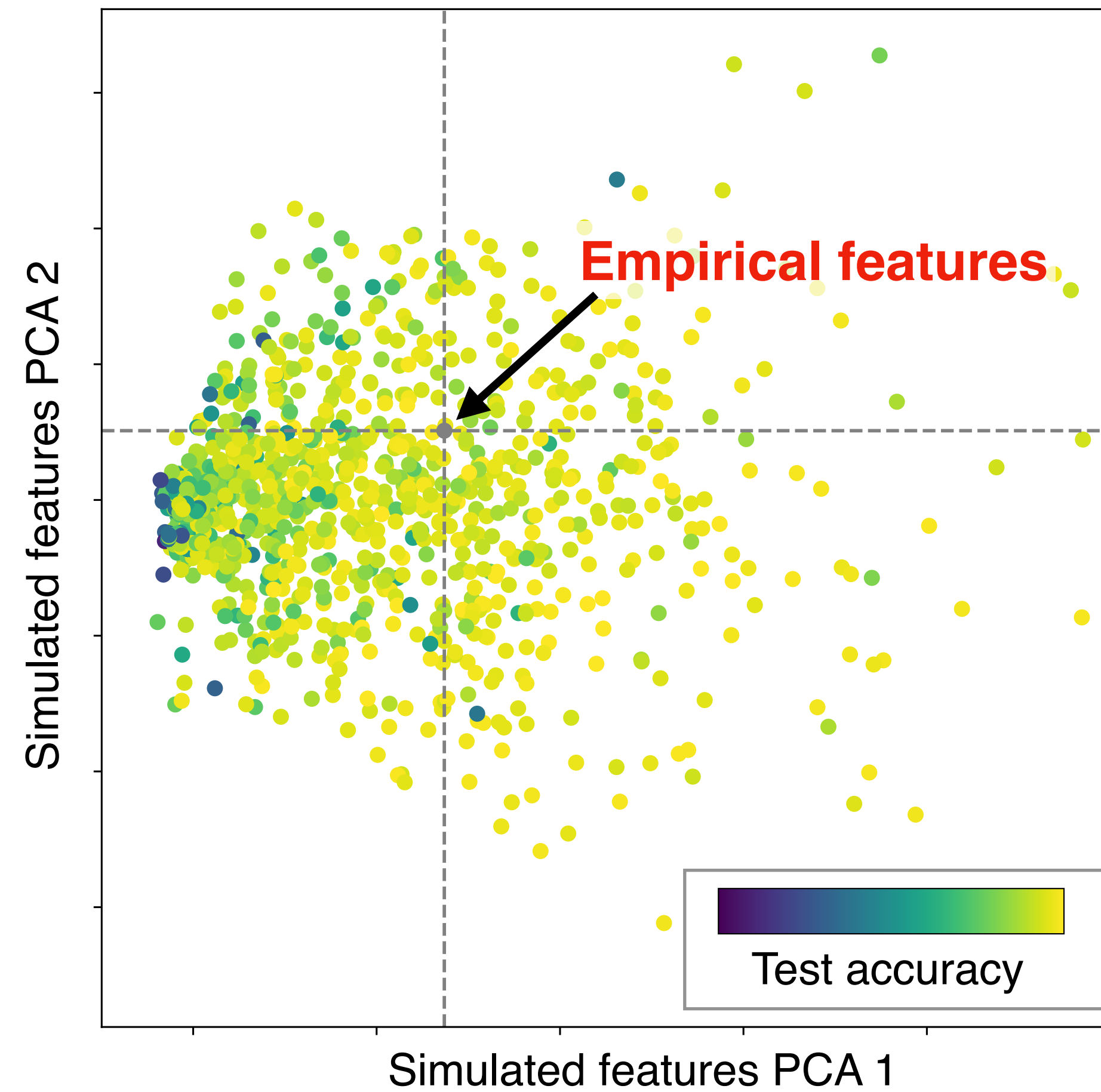
Checking that the simulations are in the right ballpark

Features of the empirical dataset

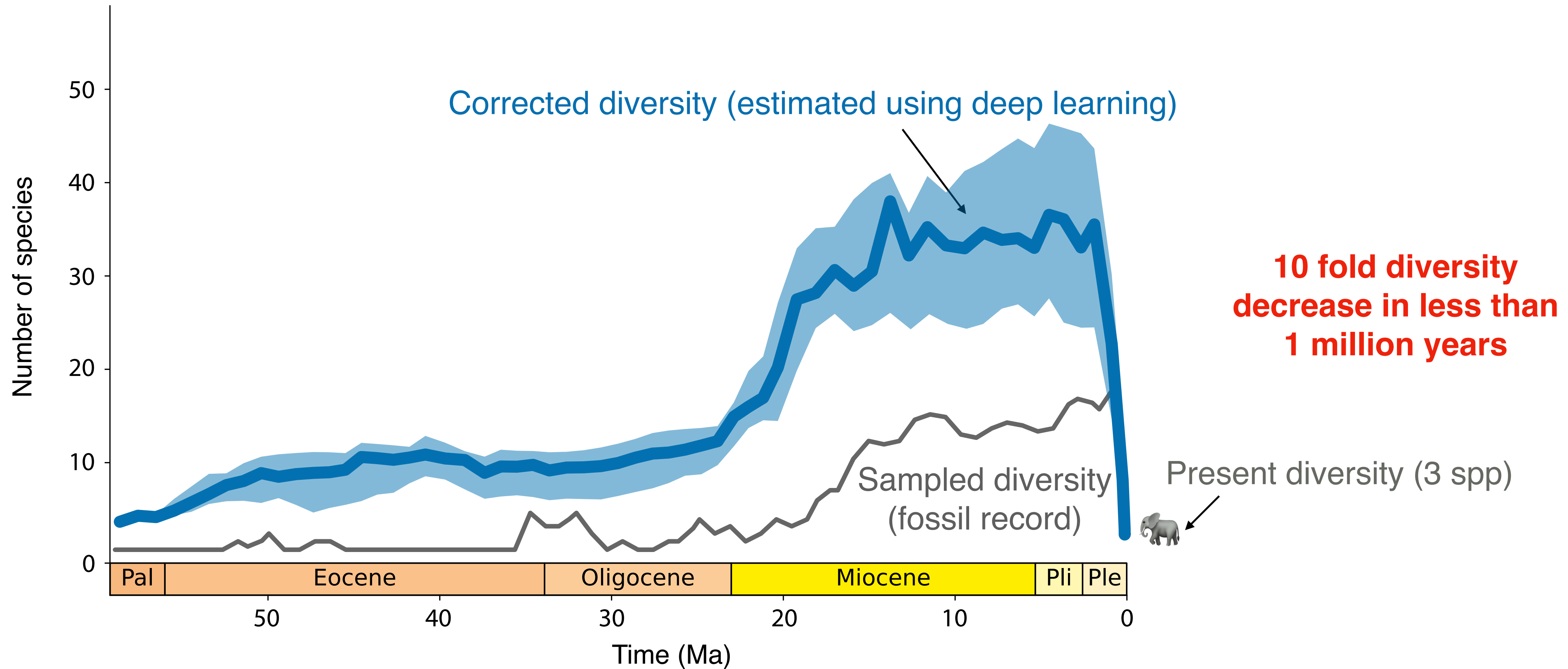
Features of the simulated datasets



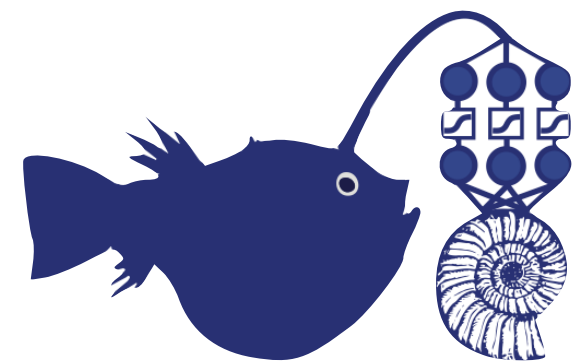
Checking that the simulations are in the right ballpark



The rise and fall of elephants



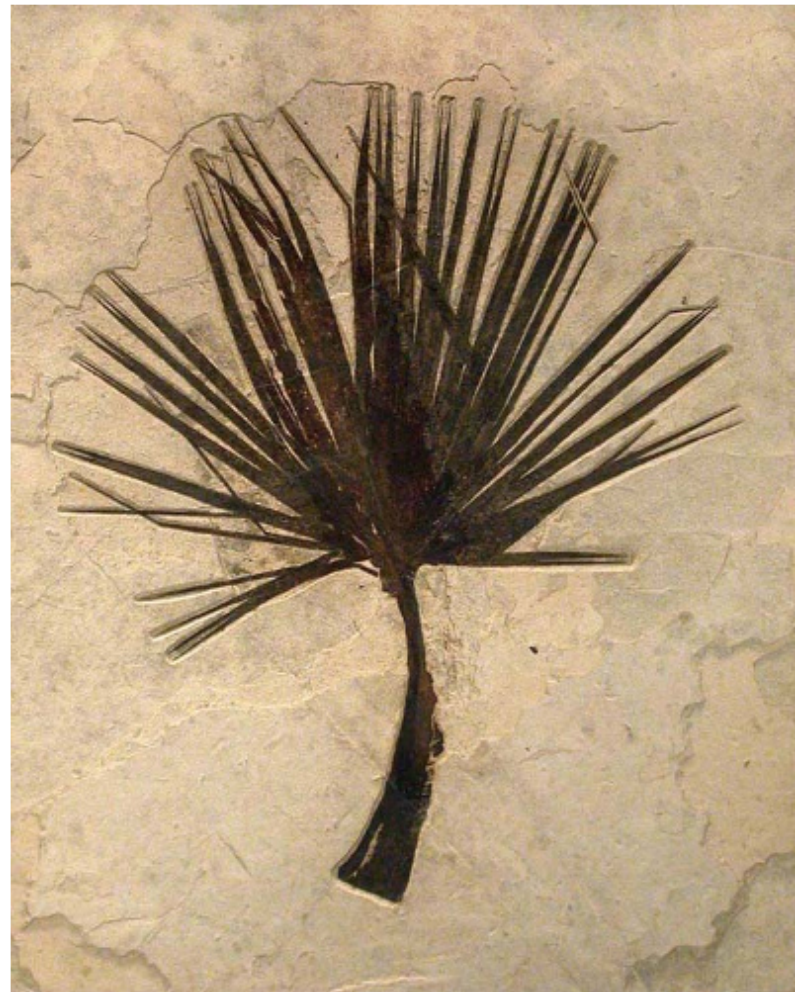
R Cooper



github.com/DeepDive-project

Cooper et al. 2024 Nature Comms

How did palms diversify through time?



A new palm fossil database

>2000 records from 1300 references

>400 verified micro- and macro-fossil occurrences

Cretaceous and Cenozoic localities from all continents



C. D. Bacon

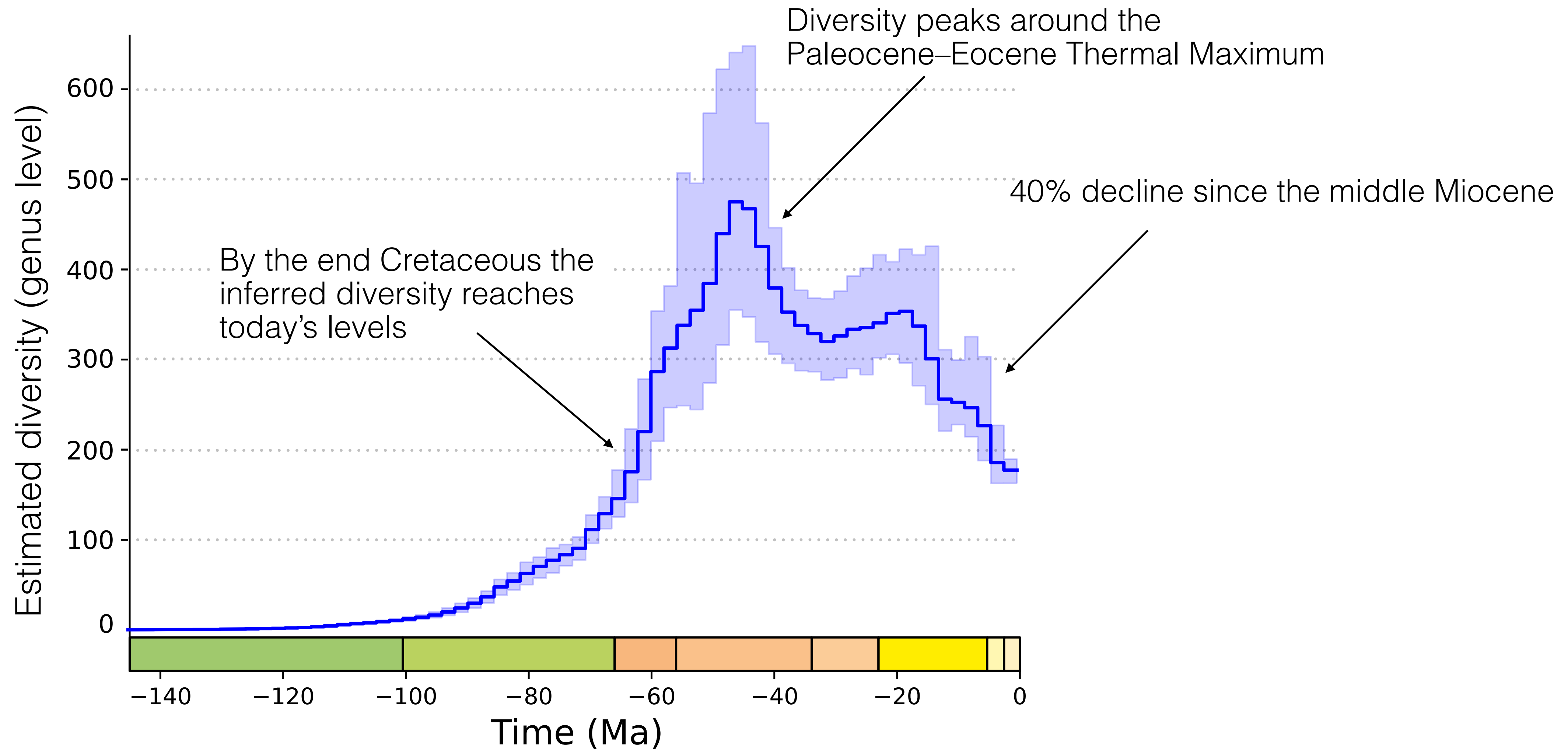
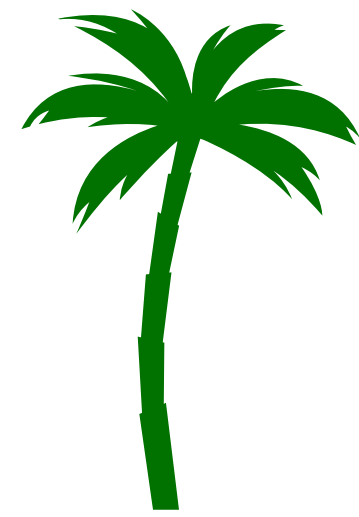
A team (ongoing) effort

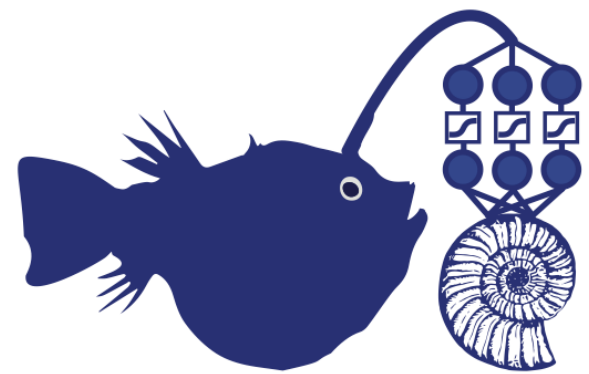
Rosane Collevatti
Carina Hoorn
Huasheng Huang
Viktoria Keller

Luis Palazzesi
Shalani Parmar
Vandana Prasad
David Sunderlin

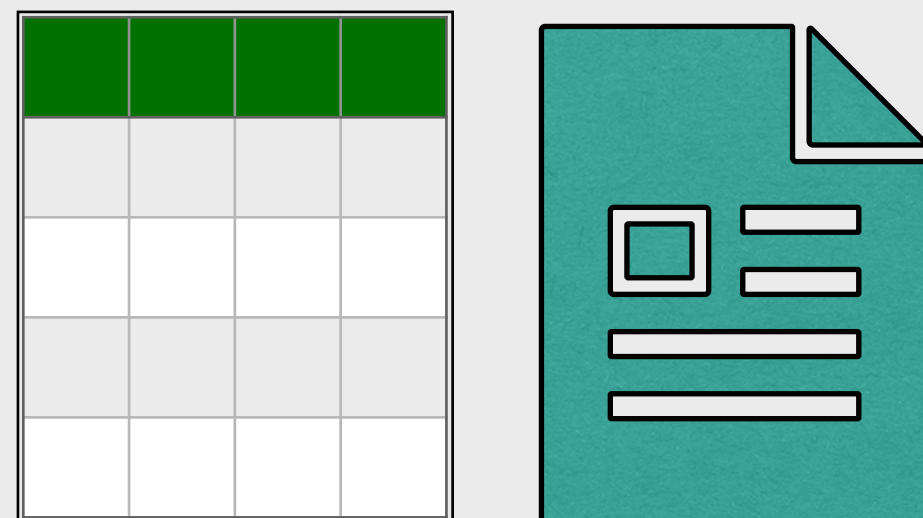
Kelly Matsunaga
Robert Morley
Yaowu Xing

The inferred palm evolutionary trajectory



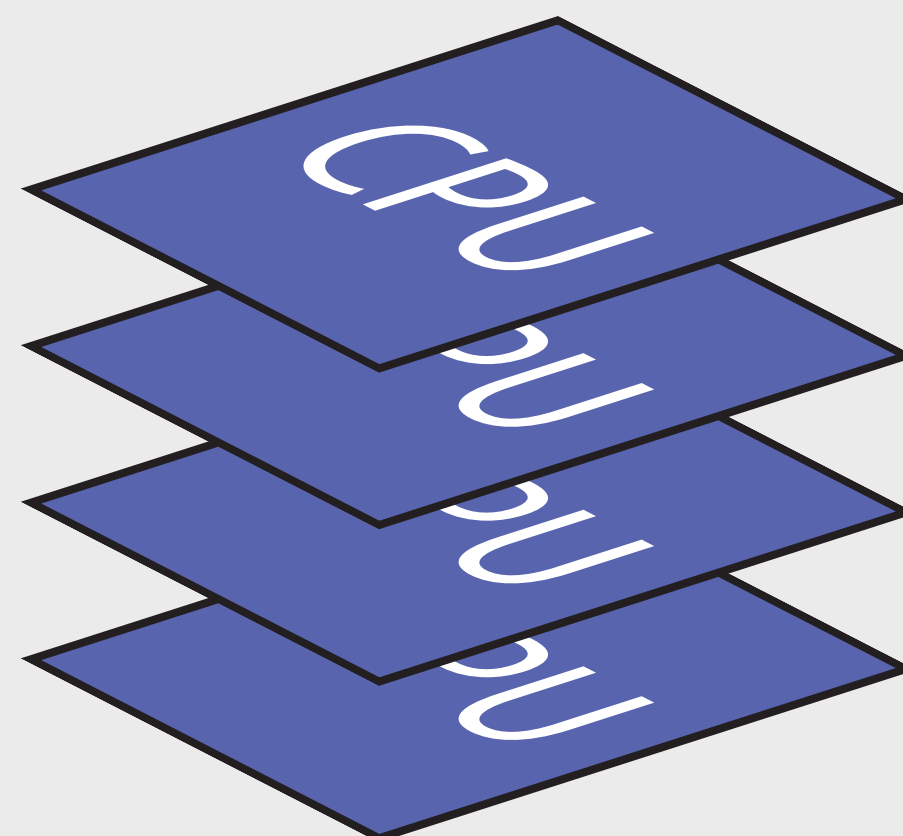


How DeepDive works



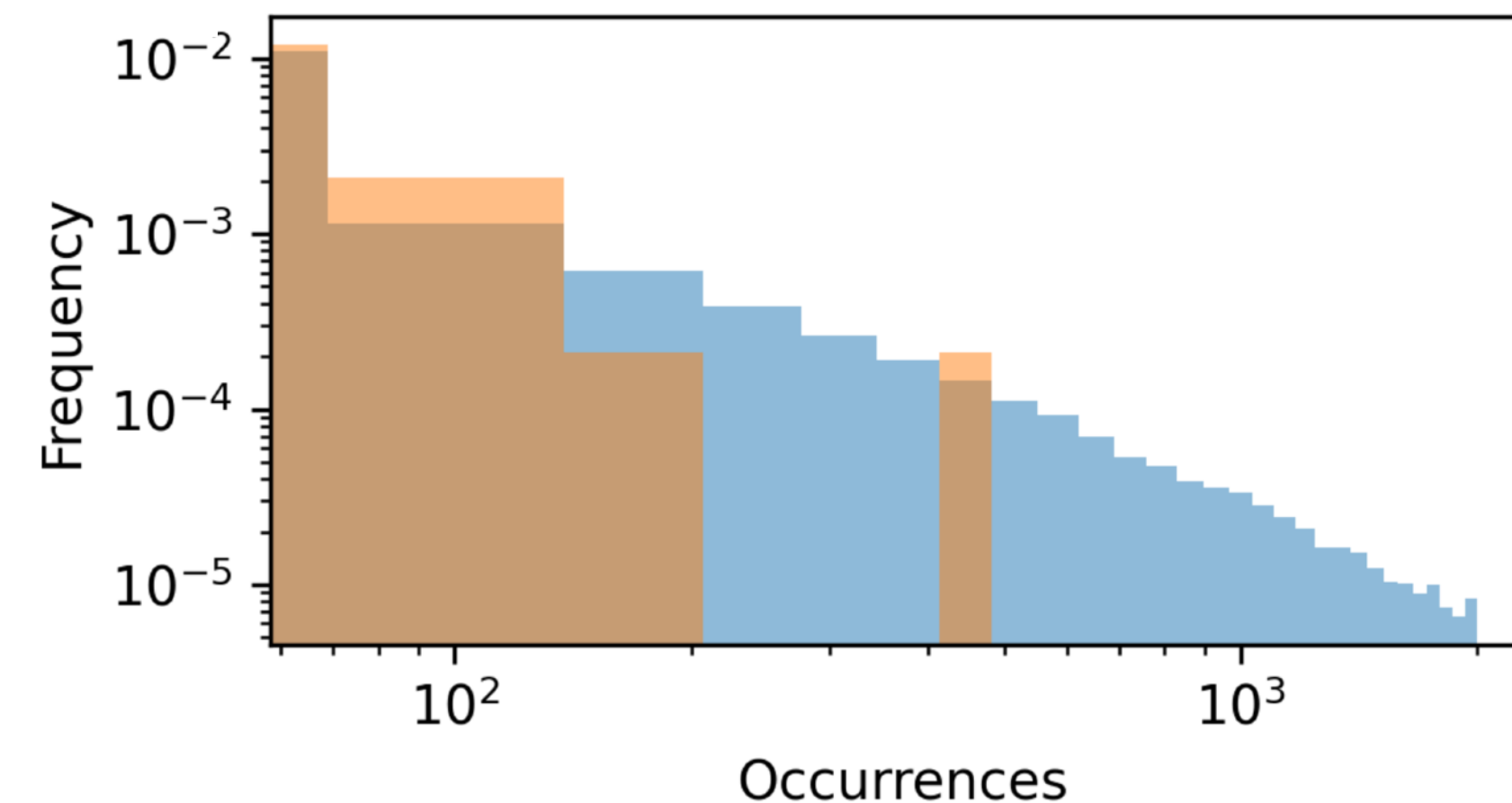
**Fossil table and
config preparation**

DeepDiveR

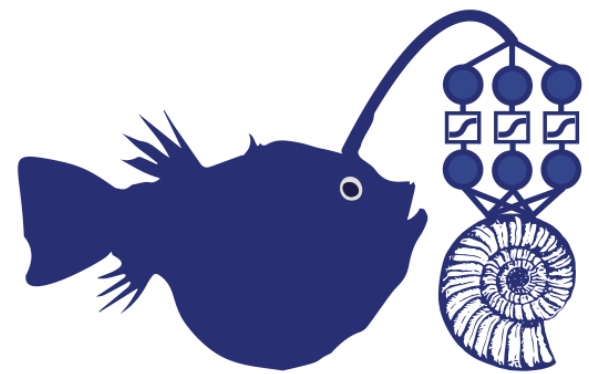


Parallel simulations

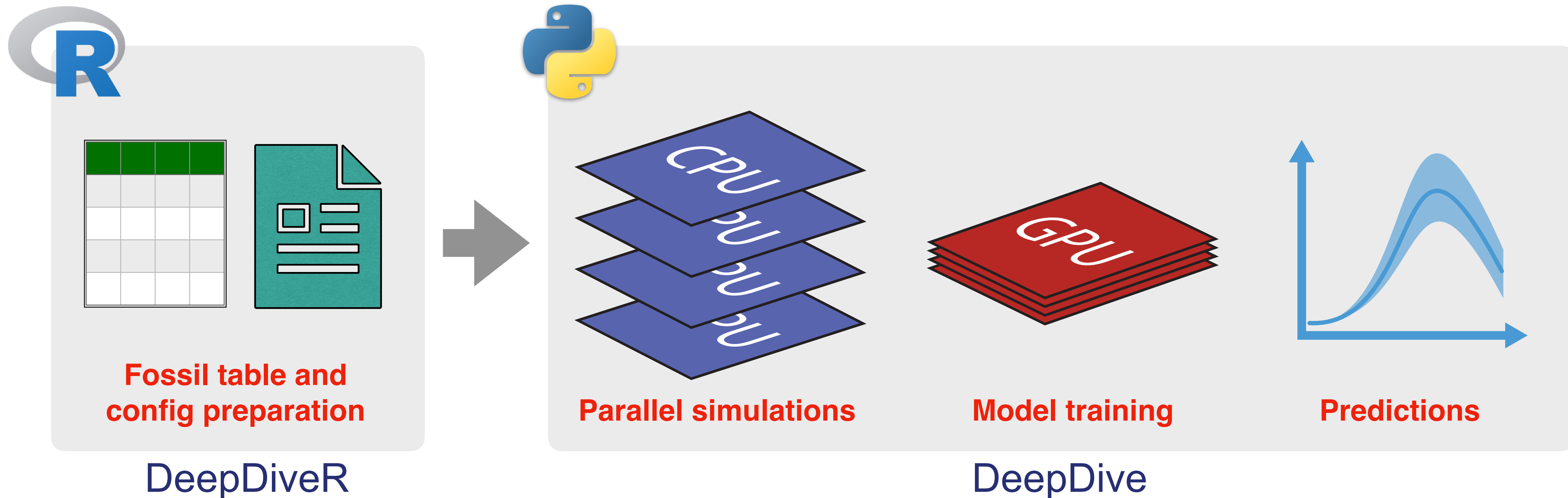
DeepDive

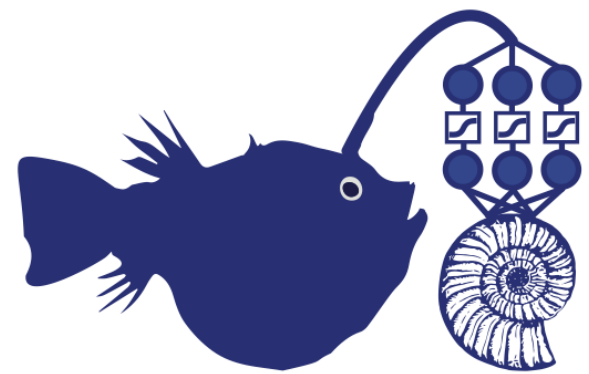


Simulations settings can
be adjusted manually but
are otherwise automatically
calibrated to match the
empirical data



How DeepDive works





Setting up an analysis using DeepDiveR

Taxon	Area	MinAge	MaxAge	Locality
<i>Ailurus</i>	Asia	0.6	1.3	loc_1
<i>Ailurus</i>	Asia	9.5	10.35	loc_2
<i>Alopecocyon</i>	Asia	9.5	10.35	loc_2
<i>Alopecocyon</i>	Europe	9.7	11.11	loc_3
...	...	...	...	...

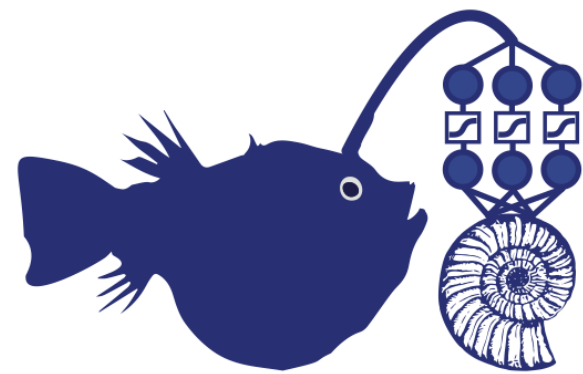
Fossil occurrence data with taxonomic, temporal and spatial info



Parse fossil table and prepare config file

```
1 library(DeepDiveR)
2
3 fossil_tbl <- read.csv("Fossil_table.csv")
4
5 # define time bins (here of 1 myr)
6 bins <- seq(23, 0, by=-1)
7
8 # prep DeepDive-formatted fossil dataset
9 prep_dd_input(fossil_tbl, bins=bins, output_file="fossil_input.csv", r=100)
10
```

Define timelines and number of age randomizations



Setting up an analysis using DeepDiveR

Taxon	Area	MinAge	MaxAge	Locality
<i>Ailurus</i>	Asia	0.6	1.3	loc_1
<i>Ailurus</i>	Asia	9.5	10.35	loc_2
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...	...	...	...	...

Fossil occurrence data with taxonomic, temporal and spatial info



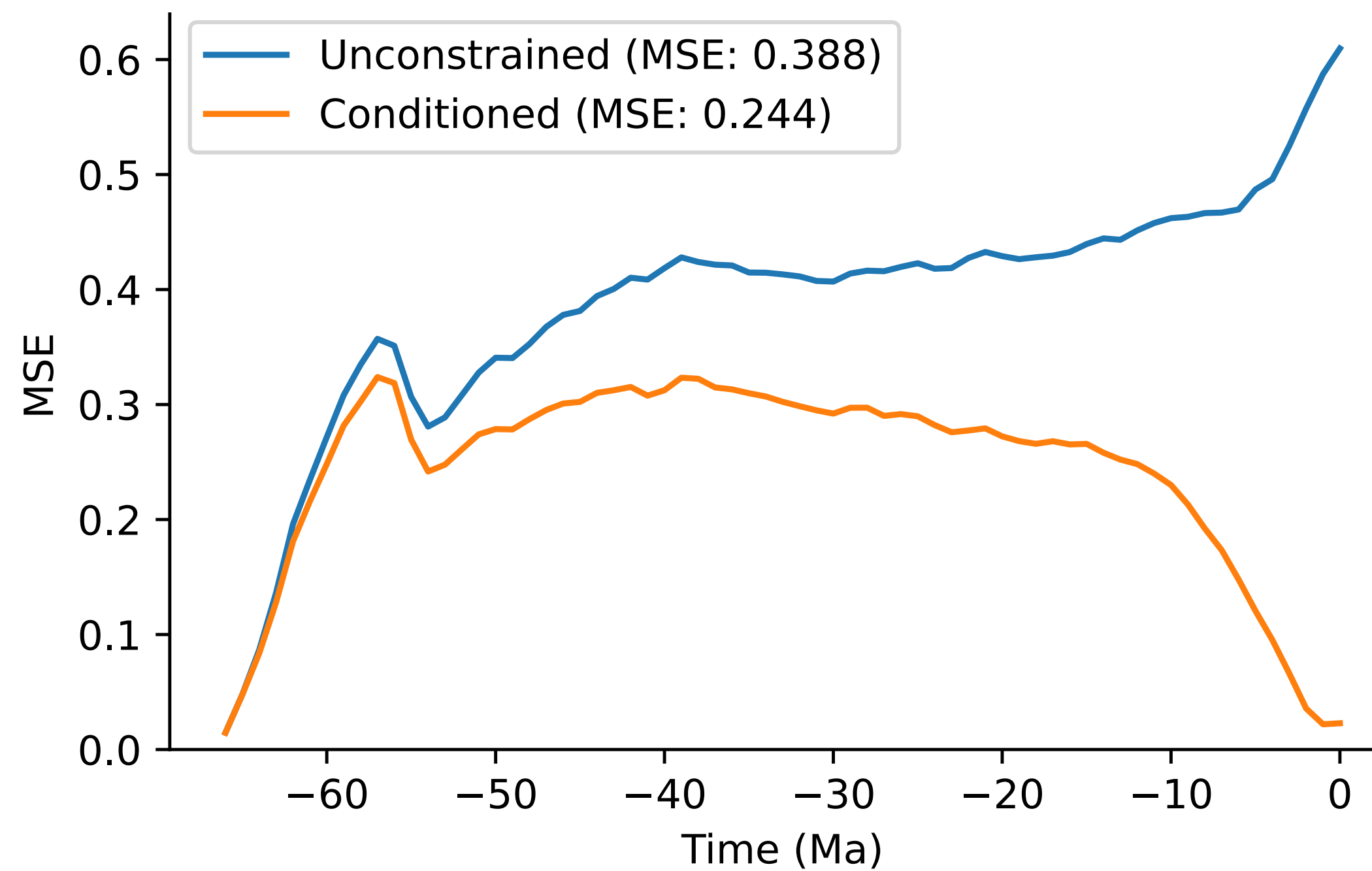
Parse fossil table and prepare config file

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6 bins <- seq(23, 0, by=-1)
7
8 # prep DeepDive-formatted fossil dataset
9 prep_dd_input(fossil_tbl, bins=bins, output_file="fossil_input.csv", r=100)
10
11 # create config file
12 config <- create_config(wd = "my_working_dir",
13                        bins = bins,
14                        present_diversity=137,
15                        include_present_diversity = TRUE)
16
17 config$write("config_file.ini")
```

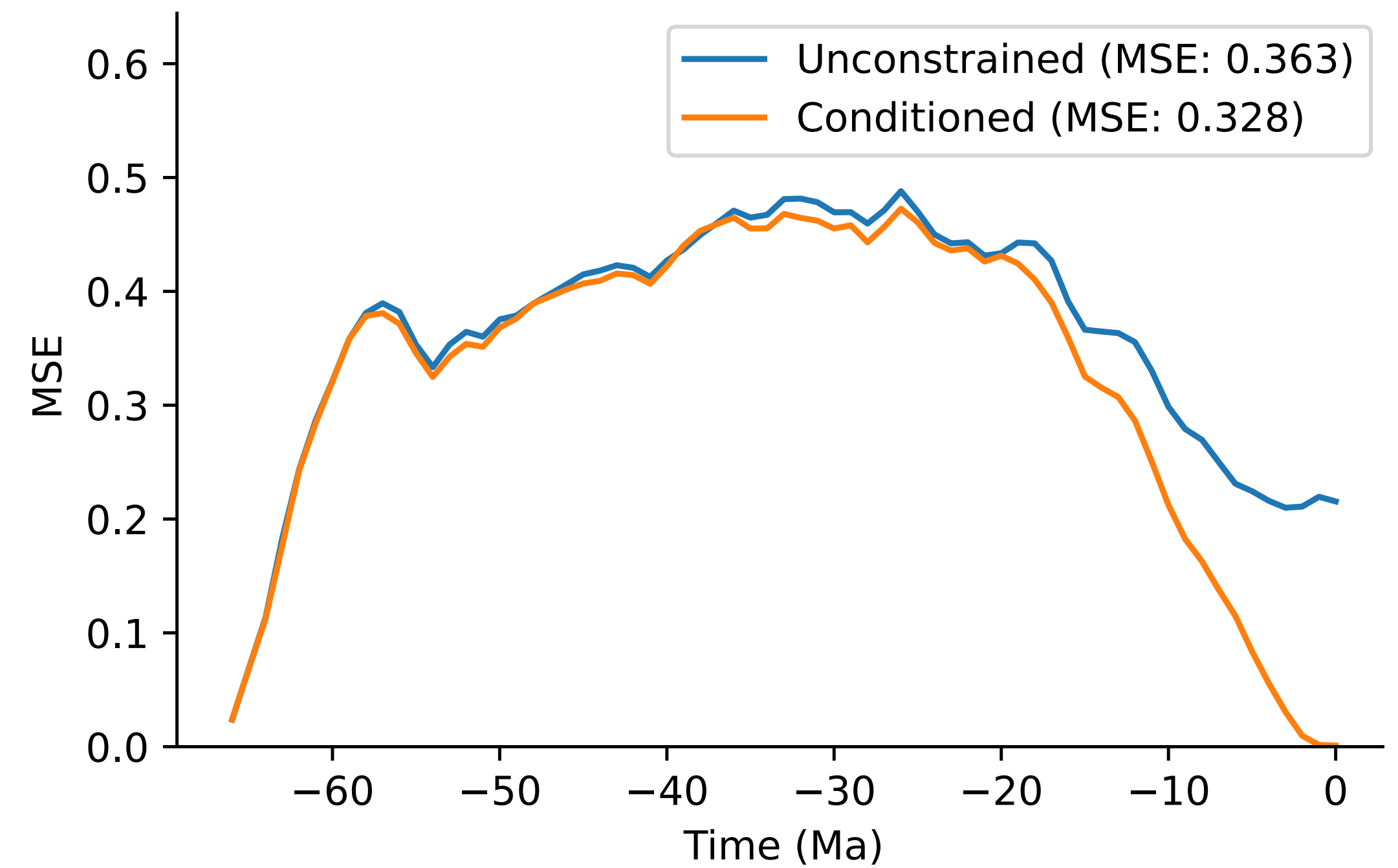
Include present diversity (if applicable)

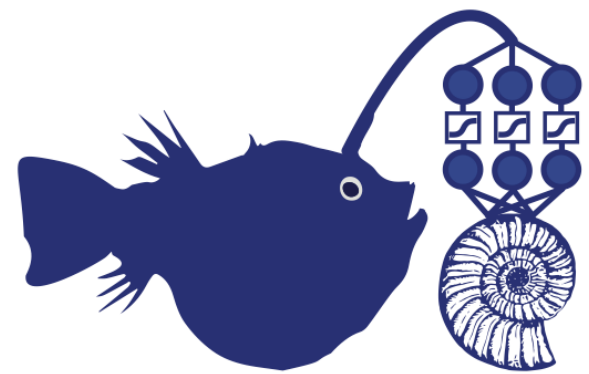
Conditioning on present diversity

Extant clades



Extinct clades





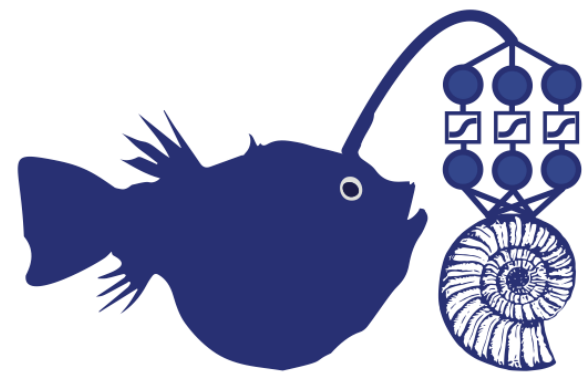
Setting up an analysis using DeepDiveR



**Adding biogeographic
information
(timing and availability
of areas)**

```
1 ▼ area_ages <- rbind(  
2   c(0, 0), # Africa  
3   c(30, 40), # Antartica  
4   c(0, 0), # Asia  
5   c(0, 0), # Europe  
6   c(0, 0), # N America  
7   c(0, 0), # Oceania  
8   c(0, 0) # S America  
9 ▲ )  
10  
11 areas_matrix(area_ages, n_areas = n_areas, label = "end", config)
```

e.g. make Antartica inhabitable
after ice sheet forms

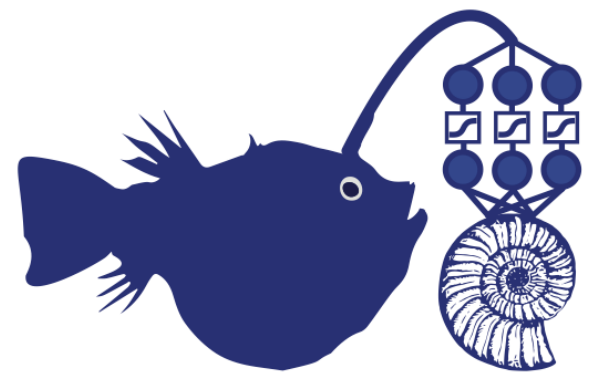


Run in an interactive
Python console

Running a DeepDive analysis from config

```
1 # import the DeepDive python library
2 from deepdive import config_runner
3
4 wd = "my_working_dir"
5
6 # launch the analysis (simulations, training, predictions)
7 config_runner.run_config("config_file.ini", wd=wd, CPU=64)
8
```





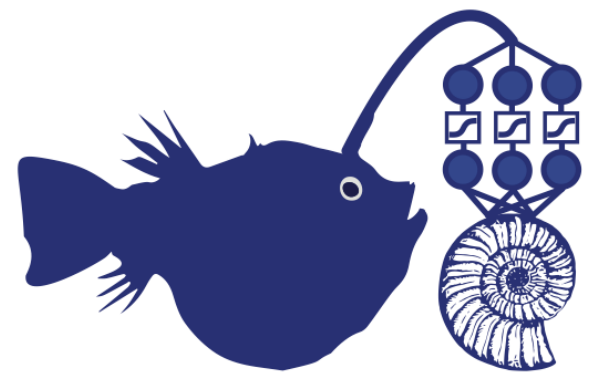
Running a DeepDive analysis from config



Or execute as a
command-line program
from a Terminal window

```
dsilvestro — -zsh — 126x38  
mbp:~ $ python run_dd_config.py my_path/config_file.ini -cpu 10 -wd my_path
```



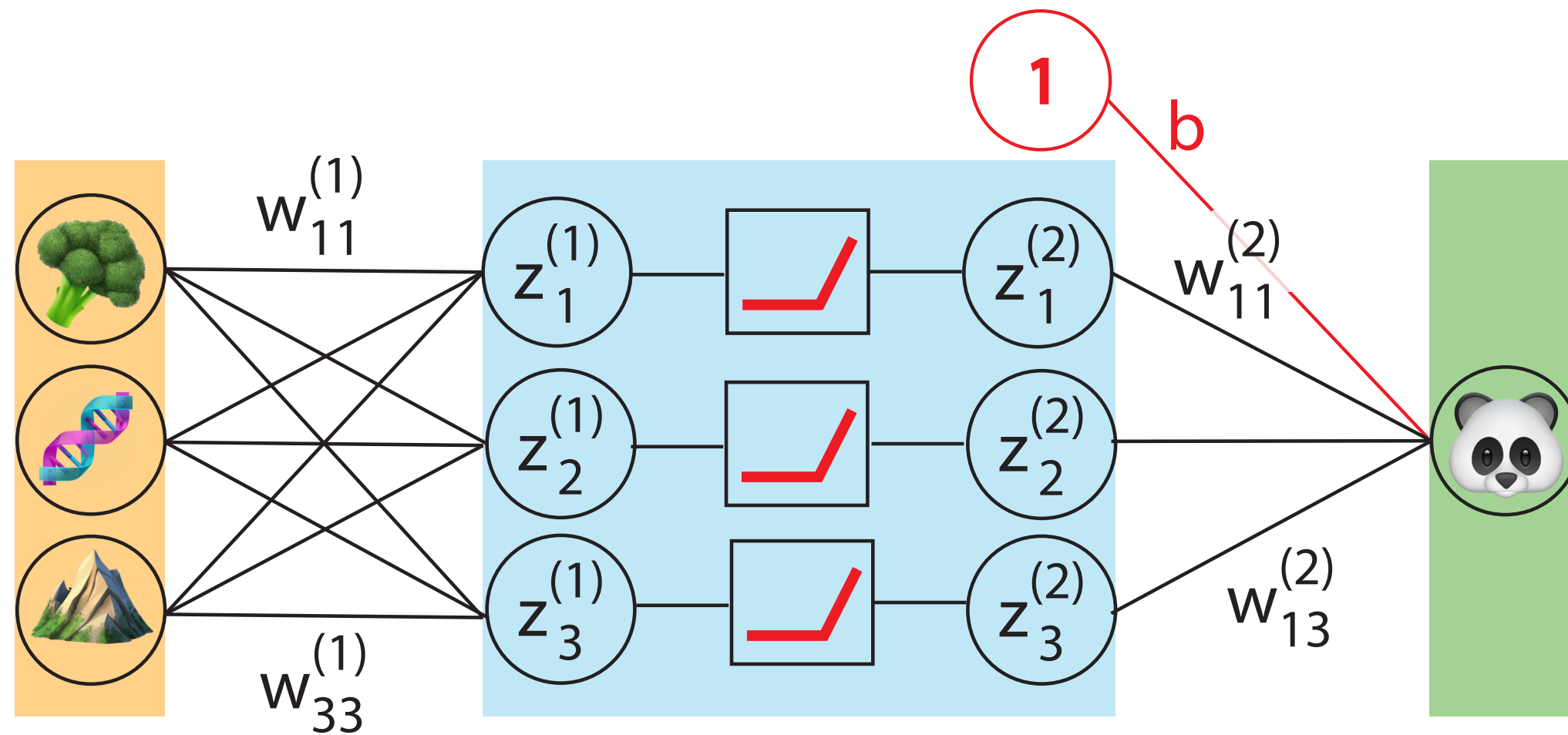


Check out our tutorials on Github

https://github.com/decoding-the-past/decoding_the_past/



Optimization of a NN model

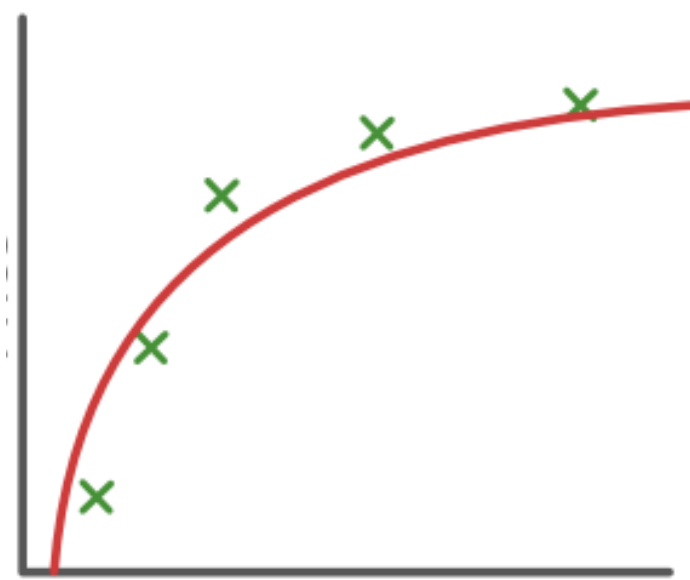


A maximum likelihood optimization of an NN would inevitably lead to over-fitting because NNs are over-parameterized

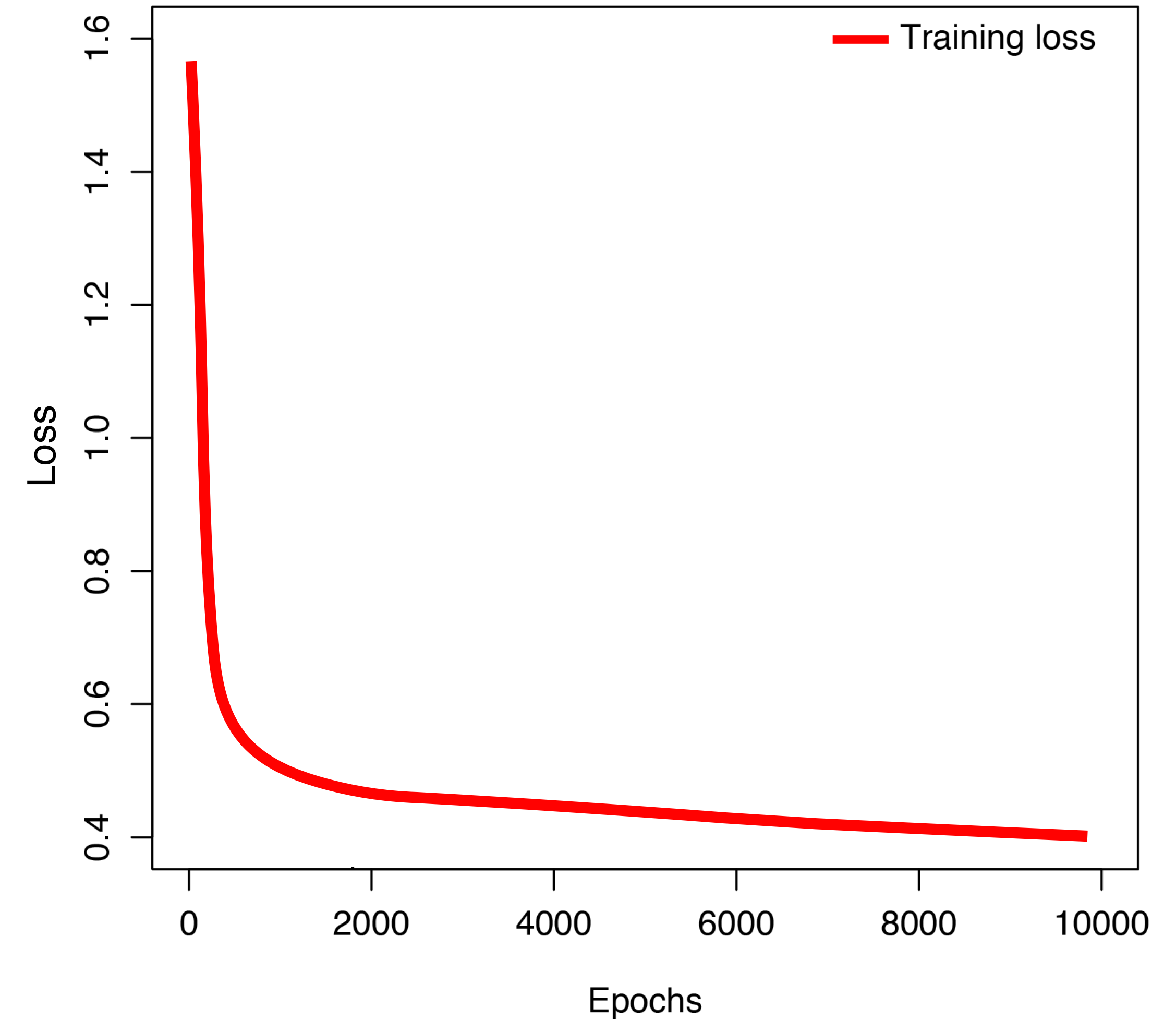
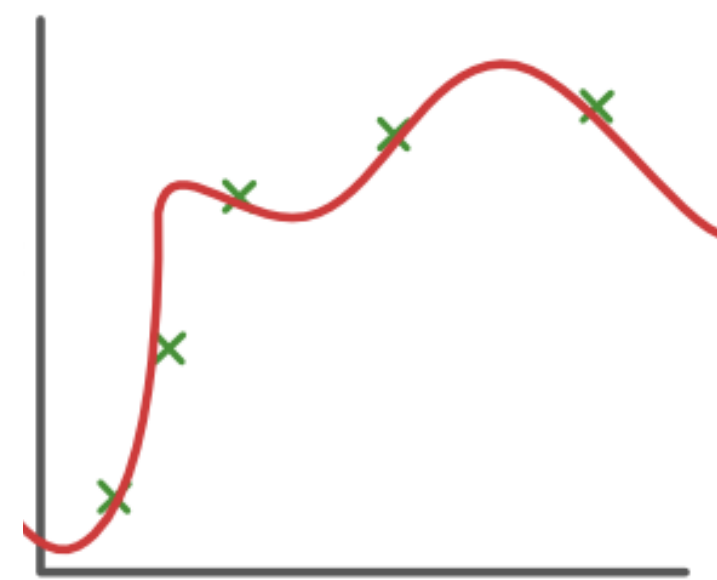
Under-fitting



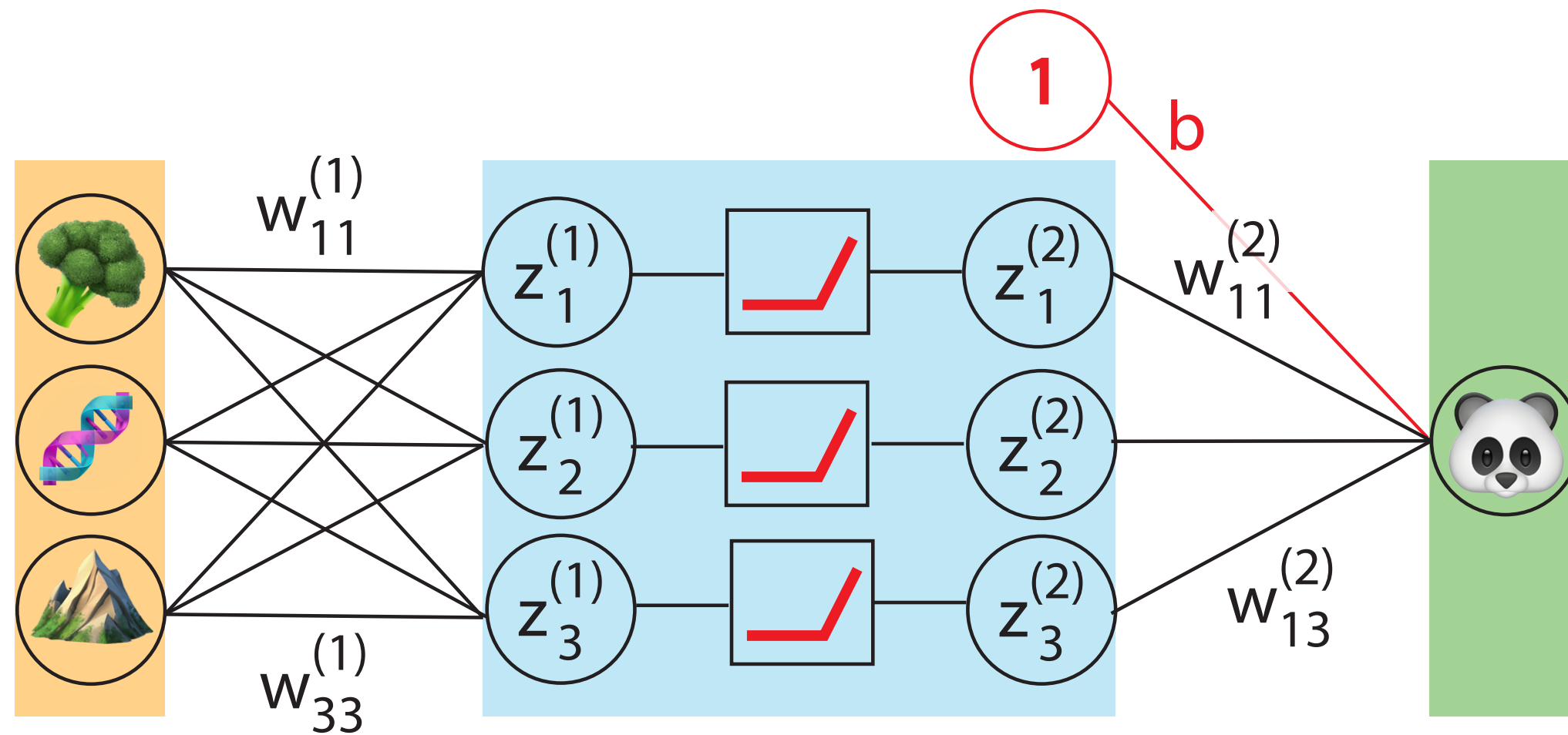
Just right



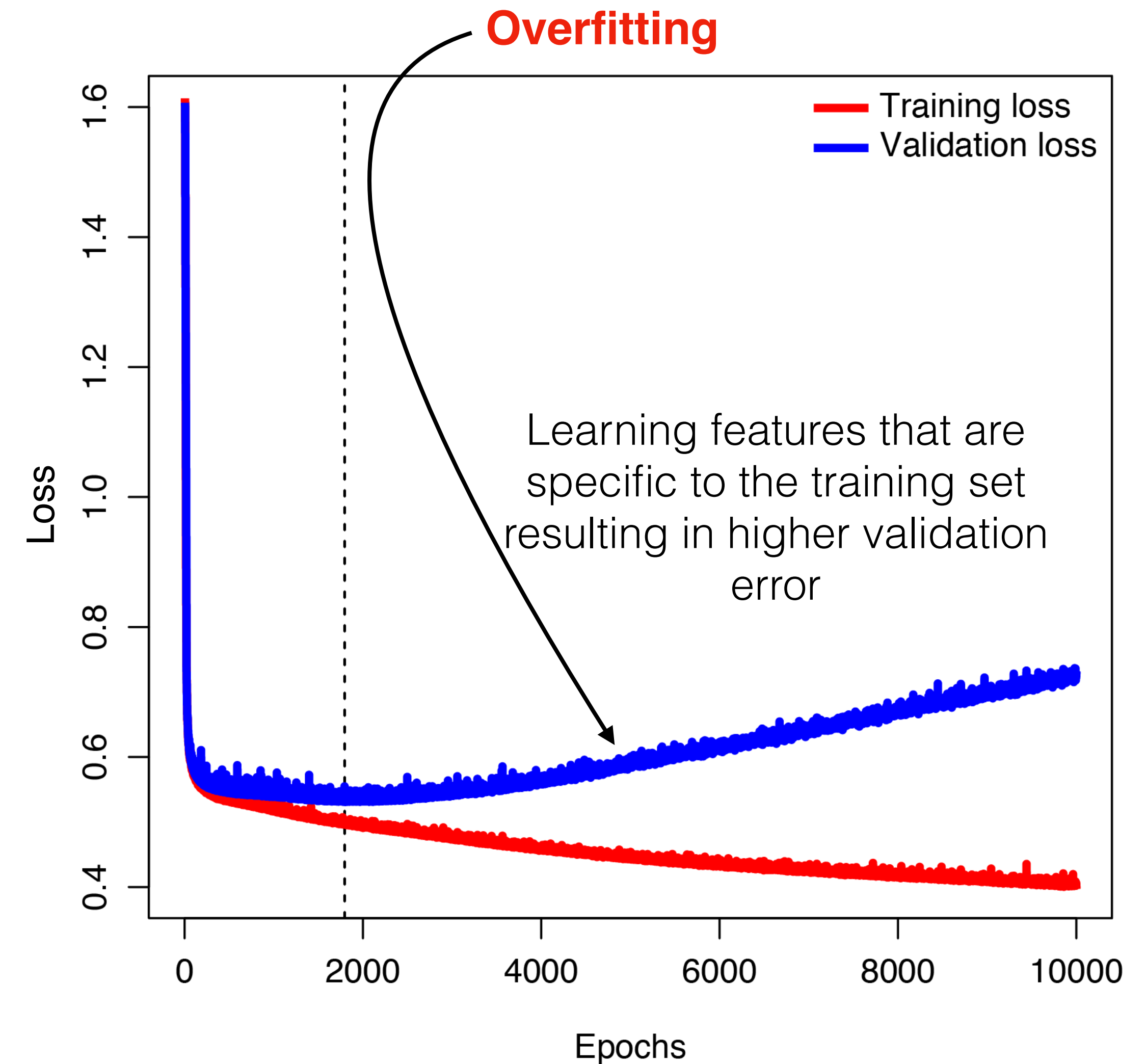
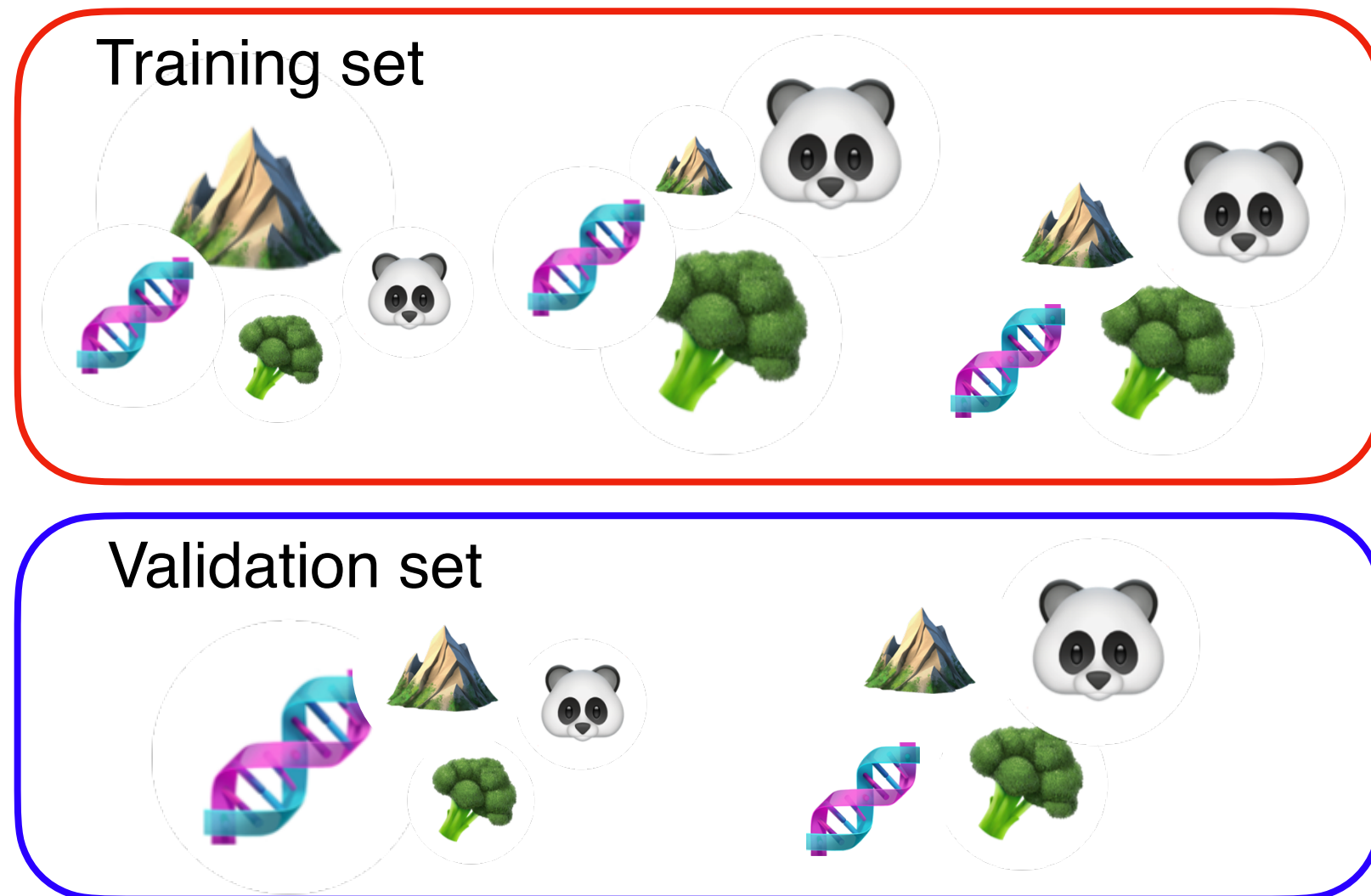
Over-fitting



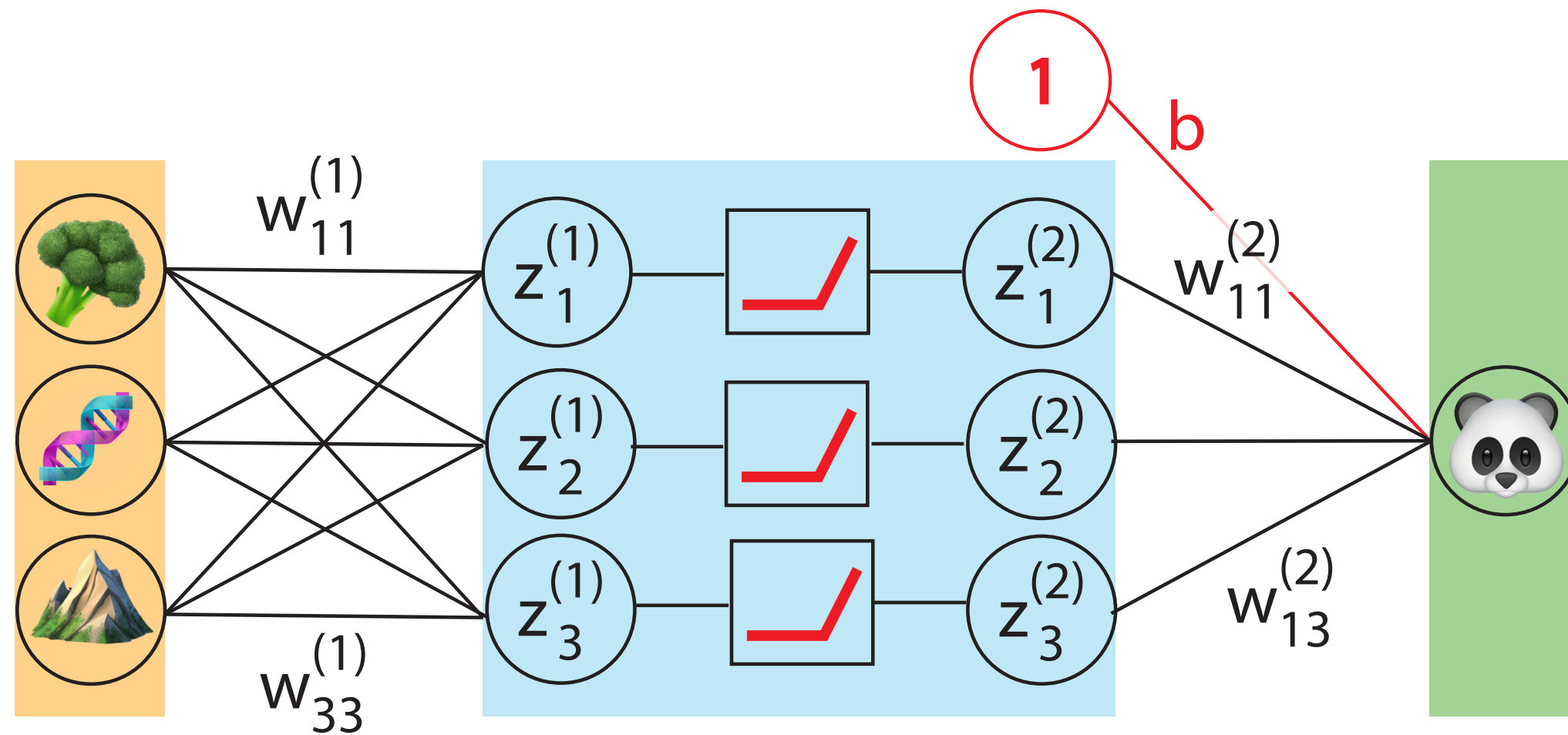
Optimization of a NN model



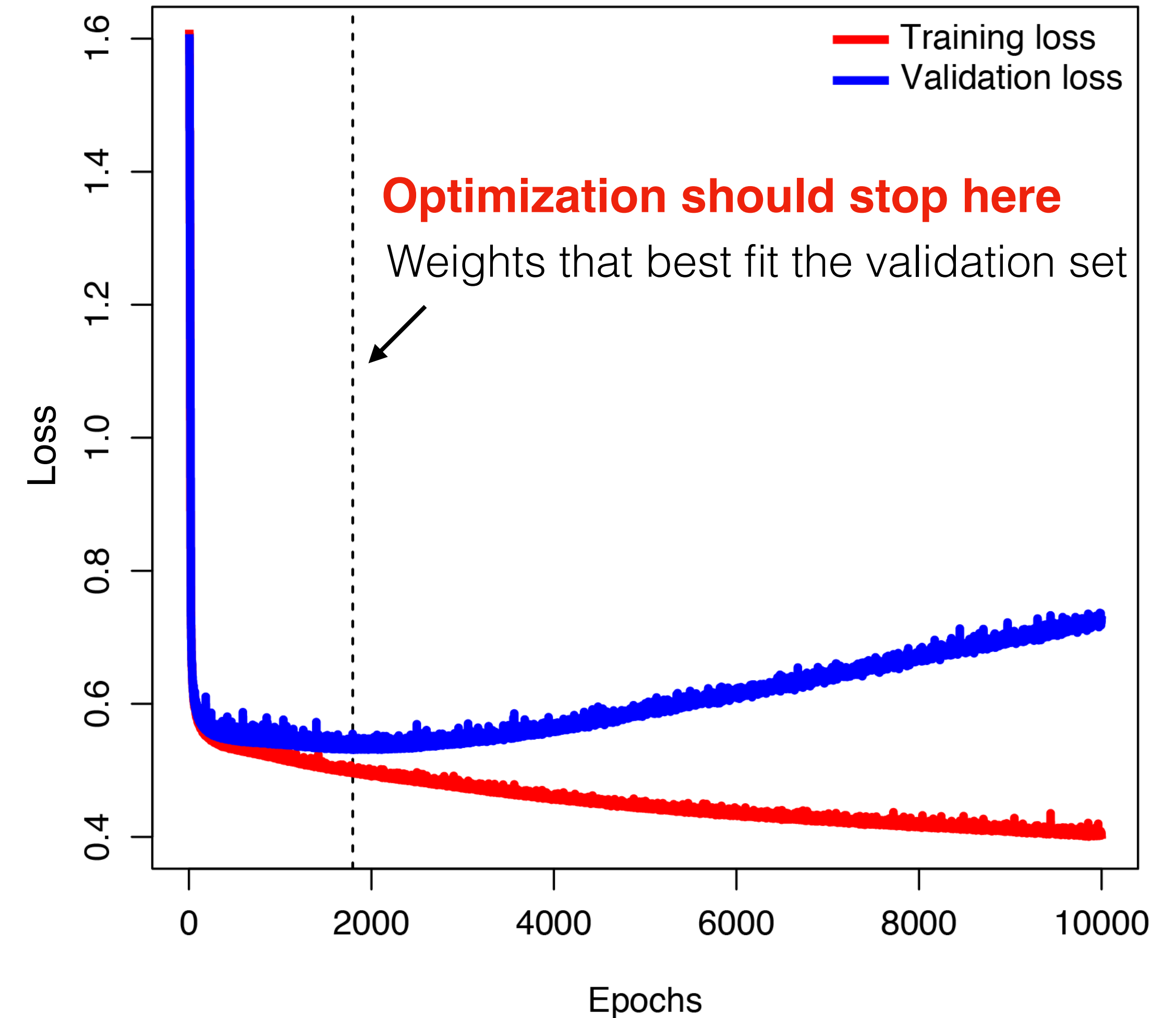
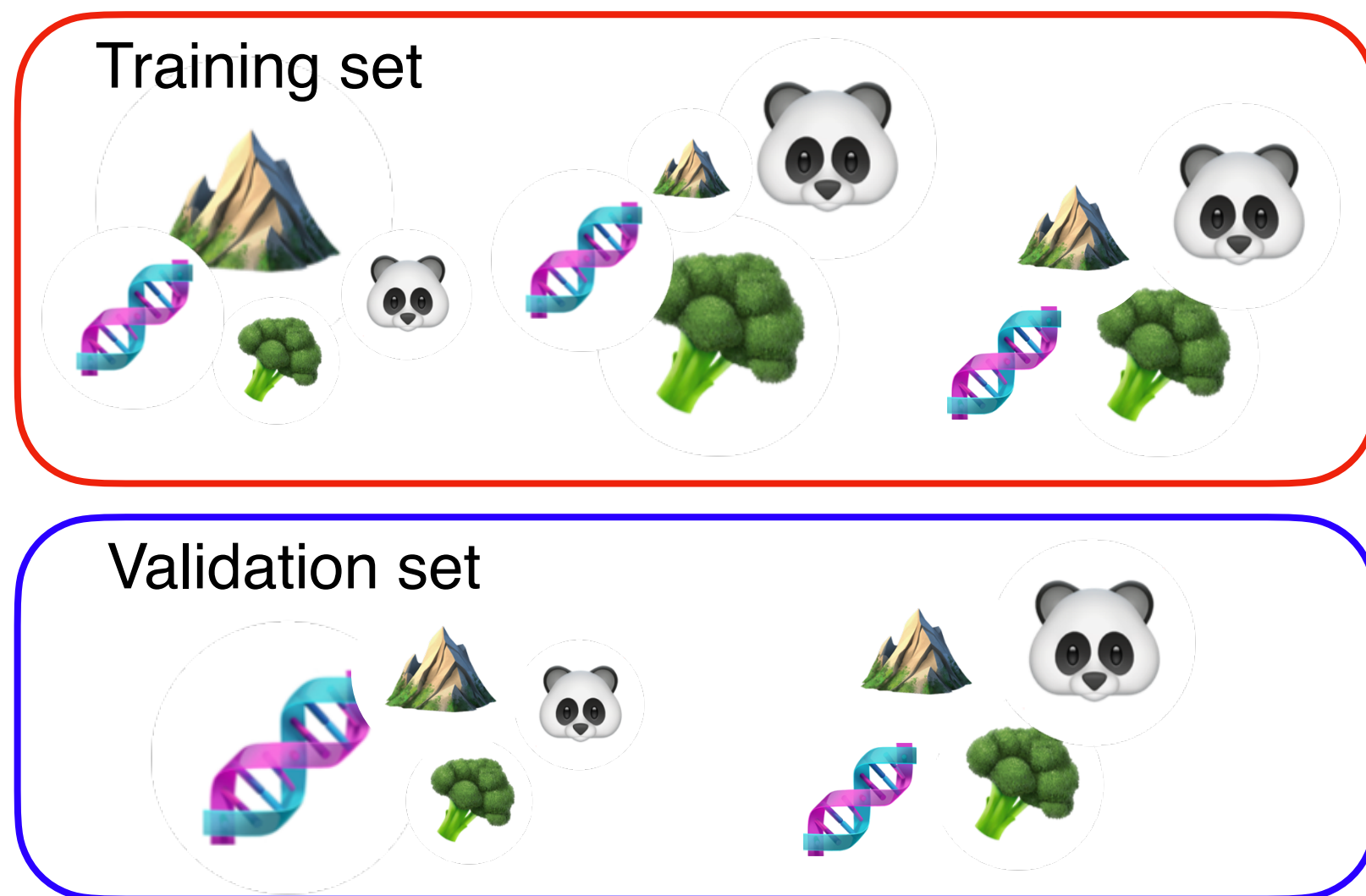
In machine learning a separate validation set is used to stop the optimization process before it starts overfitting

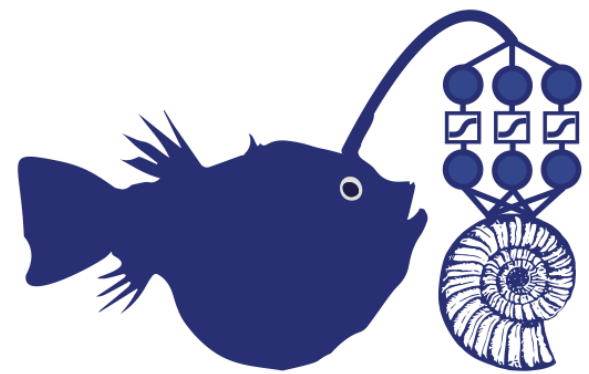


Optimization of a NN model



In machine learning a separate validation set is used to stop the optimization process before it starts overfitting





DeepDive results

Output files:

- A trained model (Keras model)
- Model accuracy (test set)
- Empirical predictions (CSV and PDF)
- Simulated vs empirical features (PDF plots)

