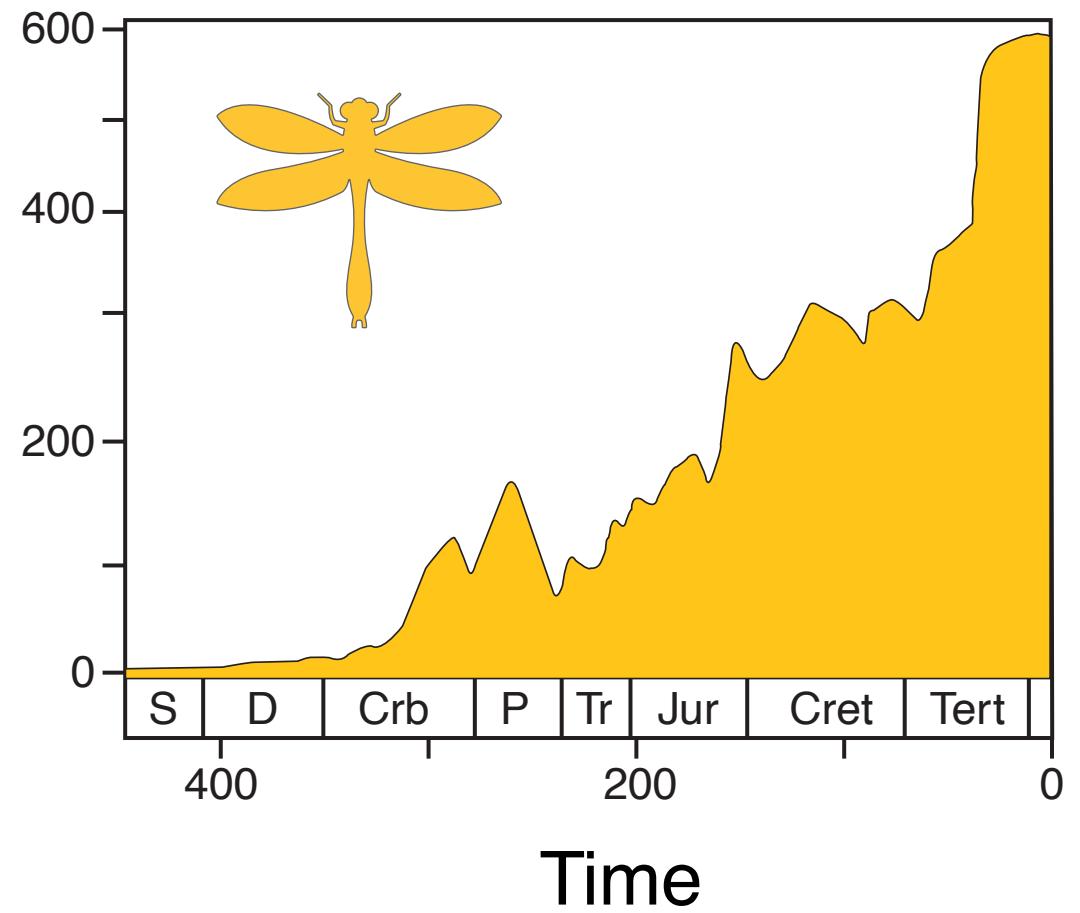
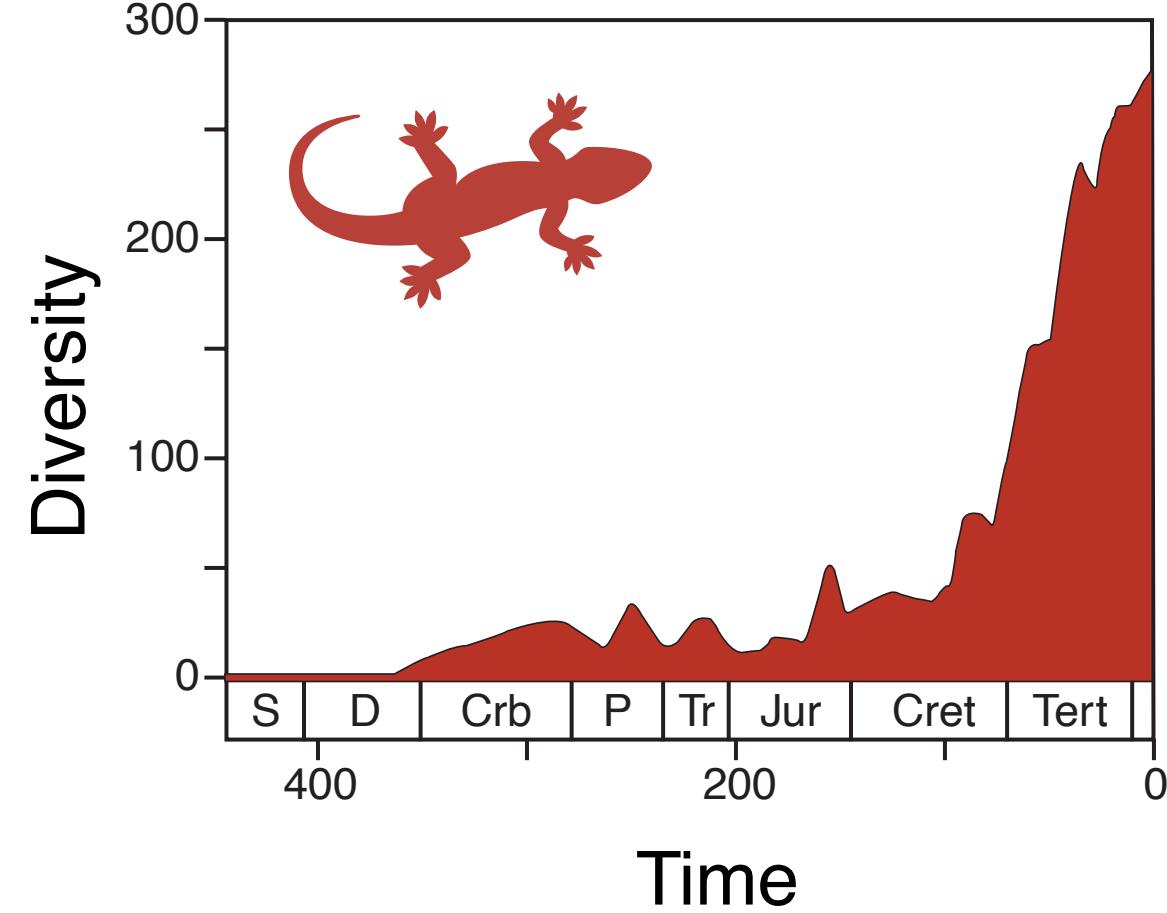
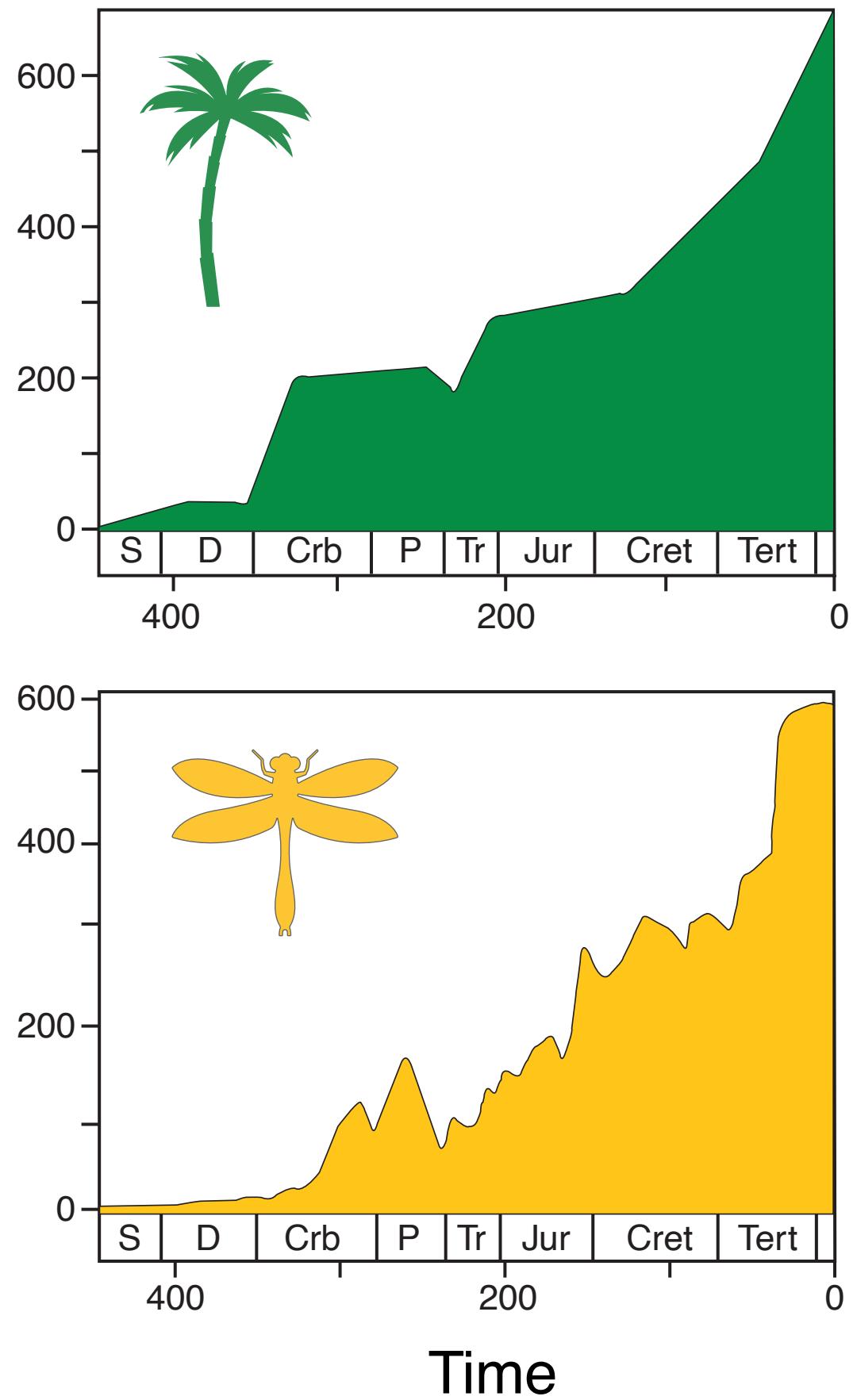
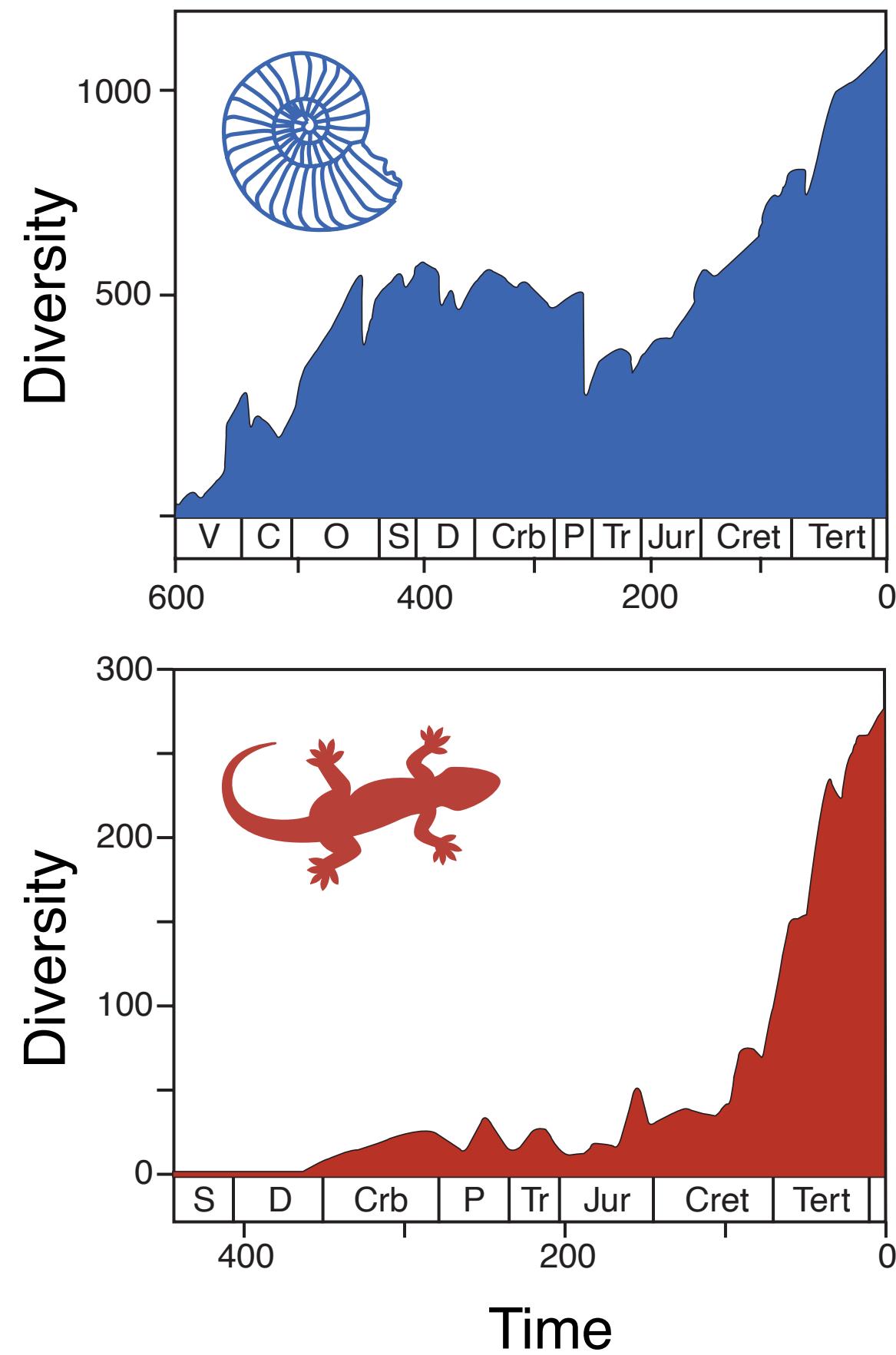


DeepDive – Estimation of diversity
trajectories using deep learning

Estimating biodiversity through time

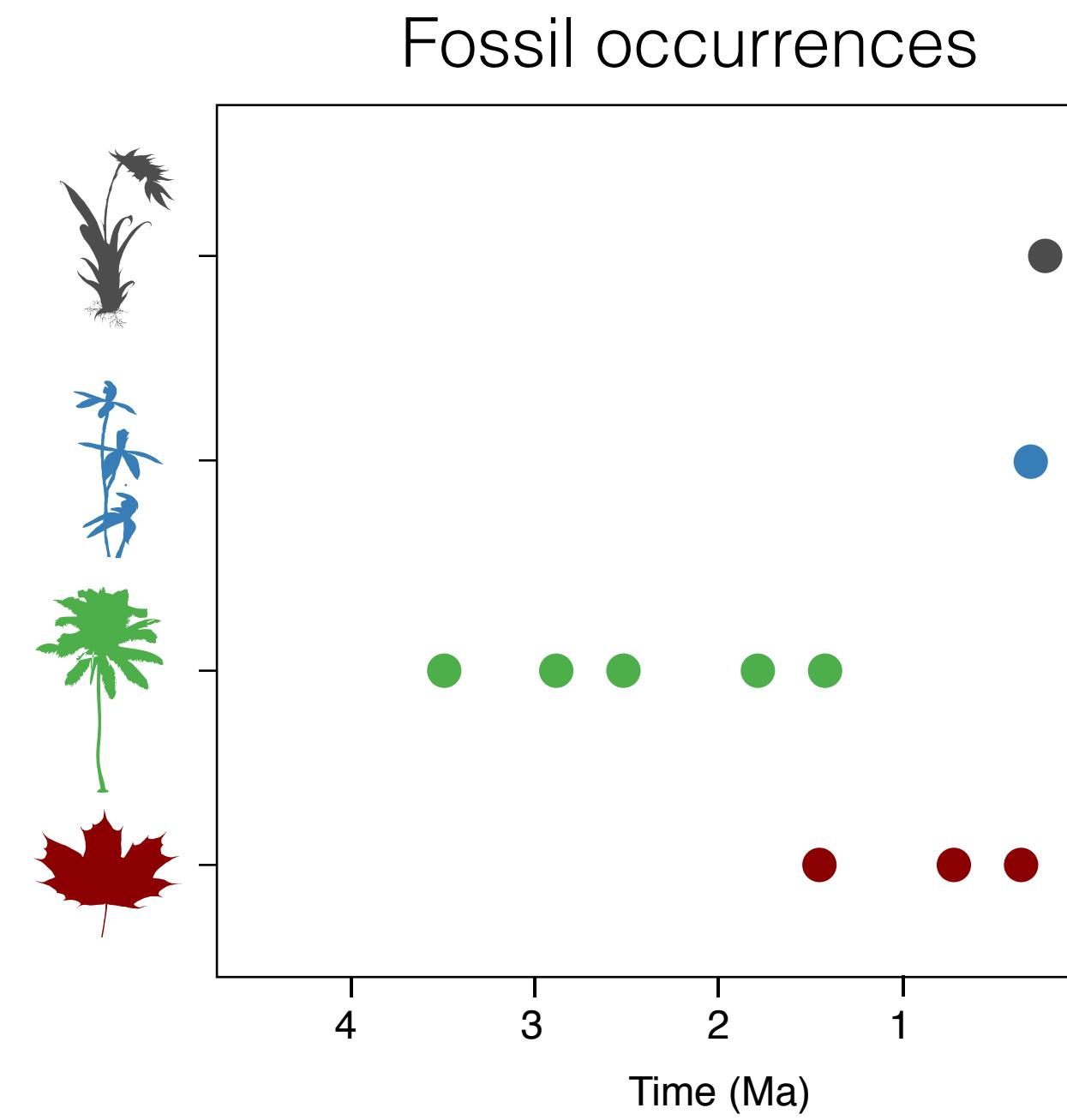


Is there a limit to biodiversity?

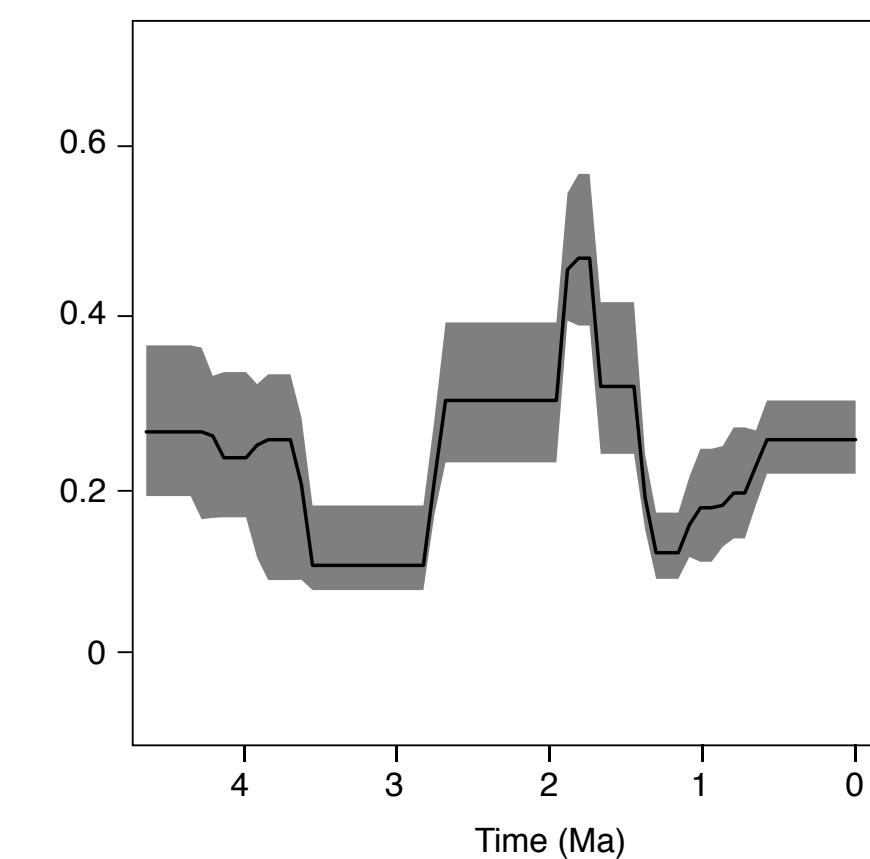
Does biodiversity increase over time?

What are the mechanisms controlling biodiversity?

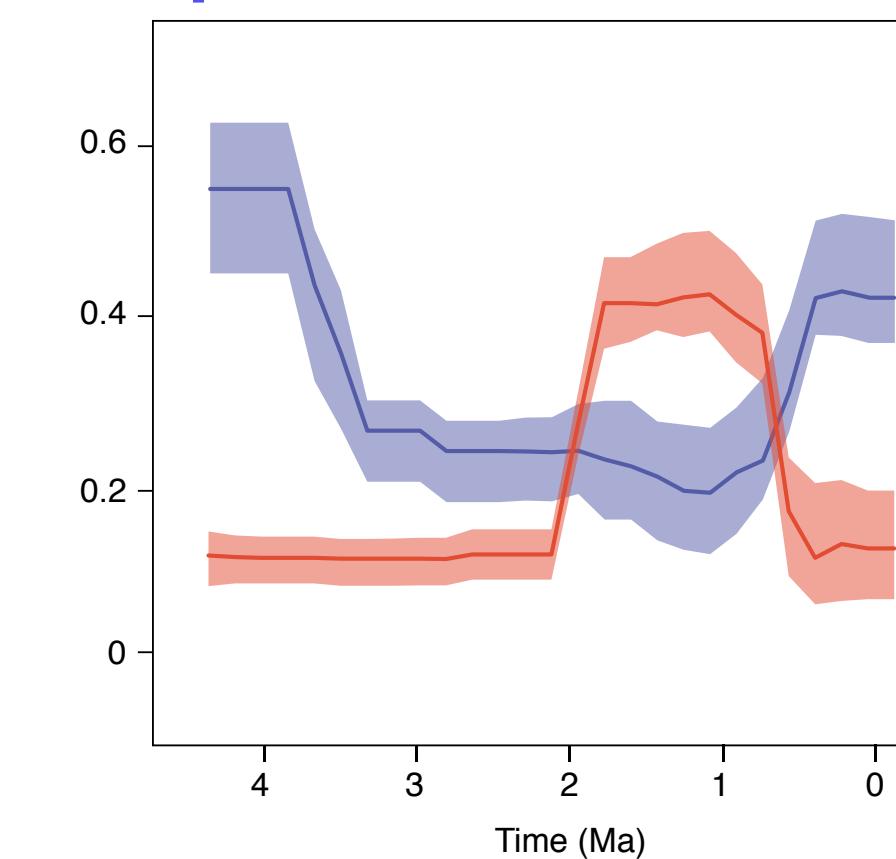
Bayesian estimation of diversity trajectories – mcmcDivE



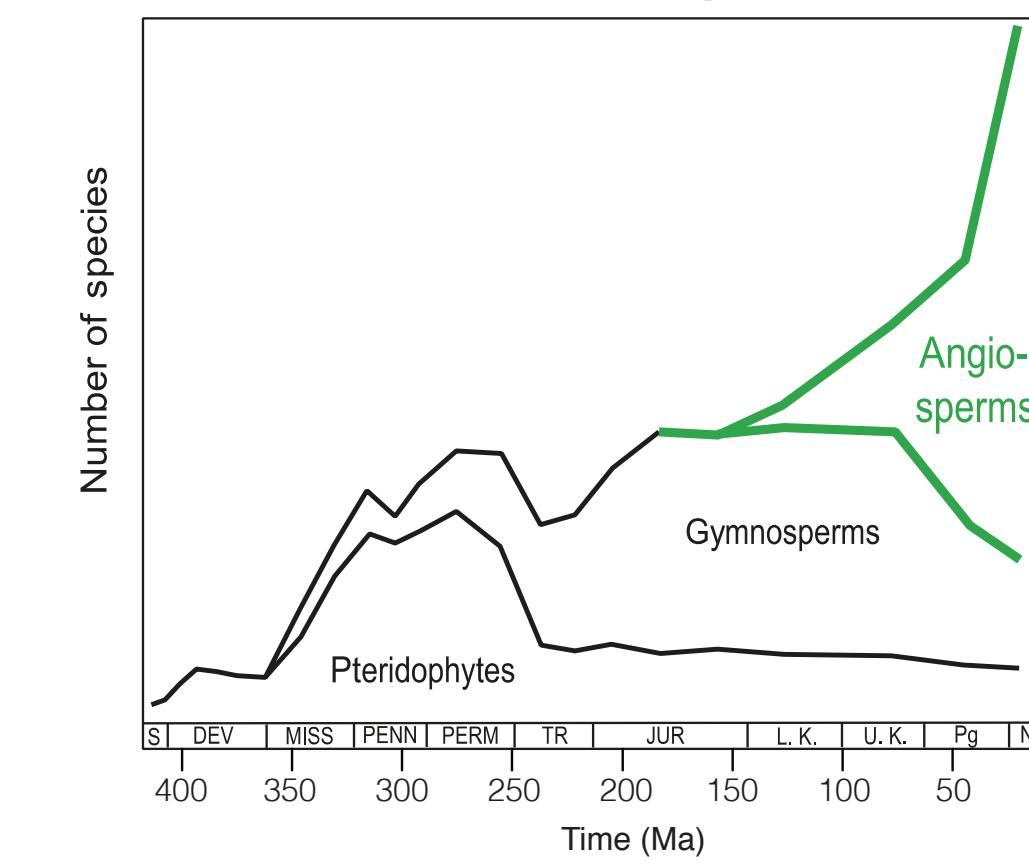
Preservation



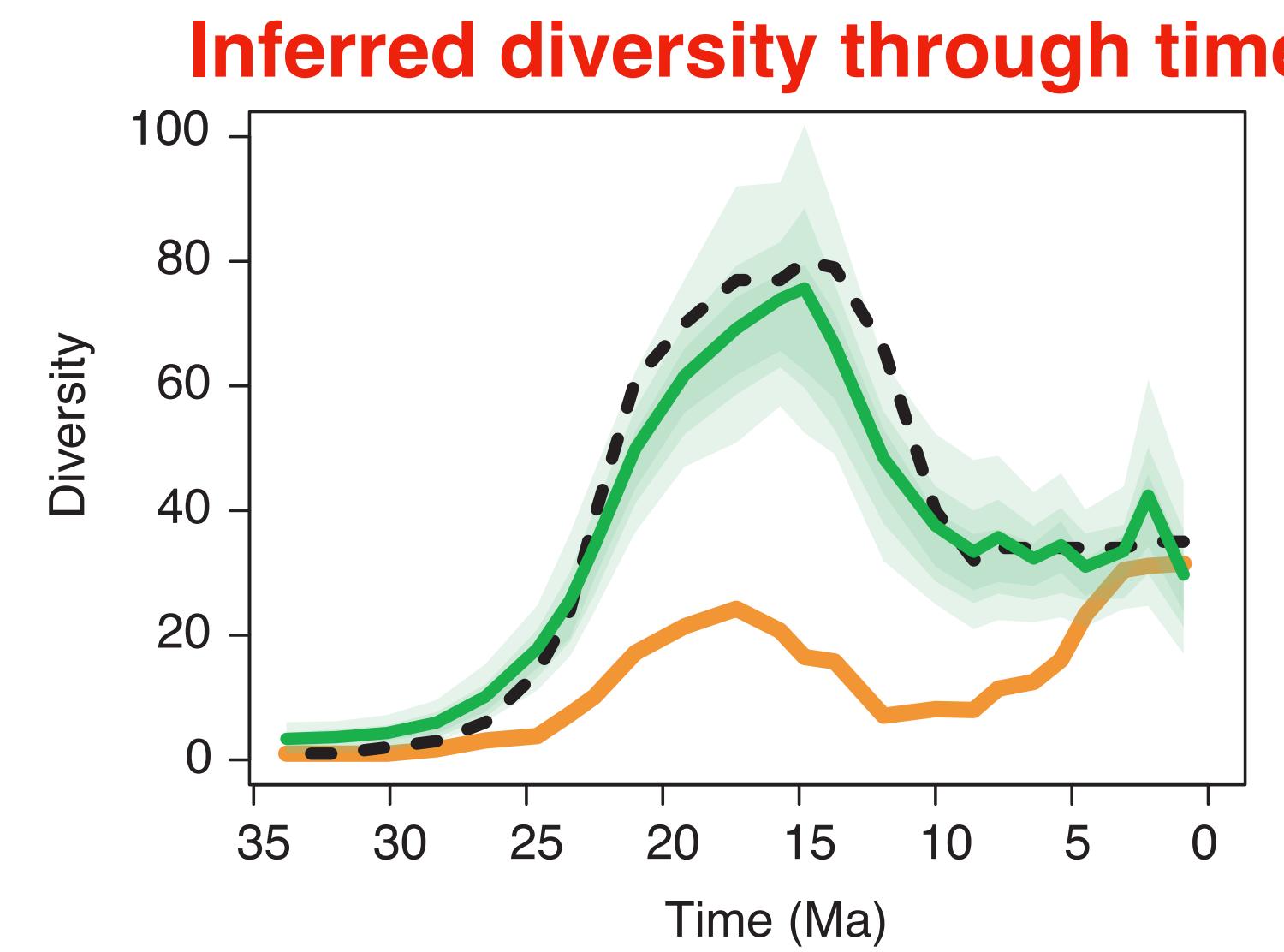
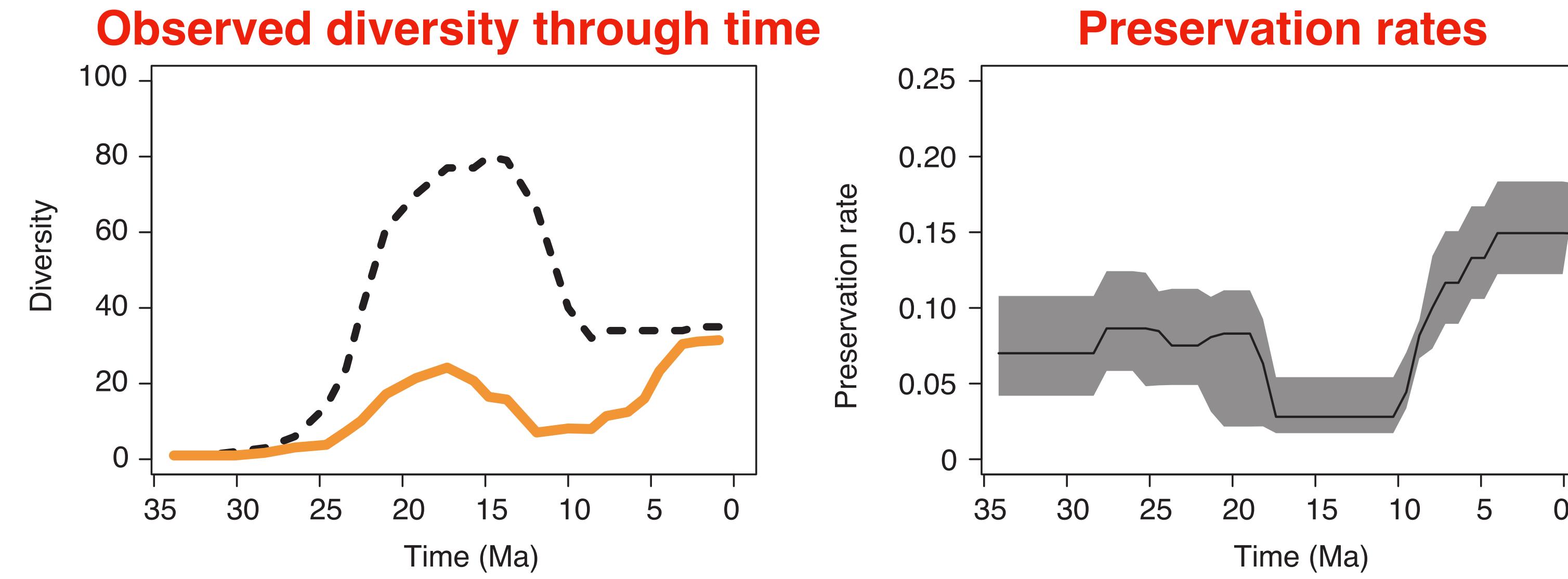
Speciation & extinction



Diversity



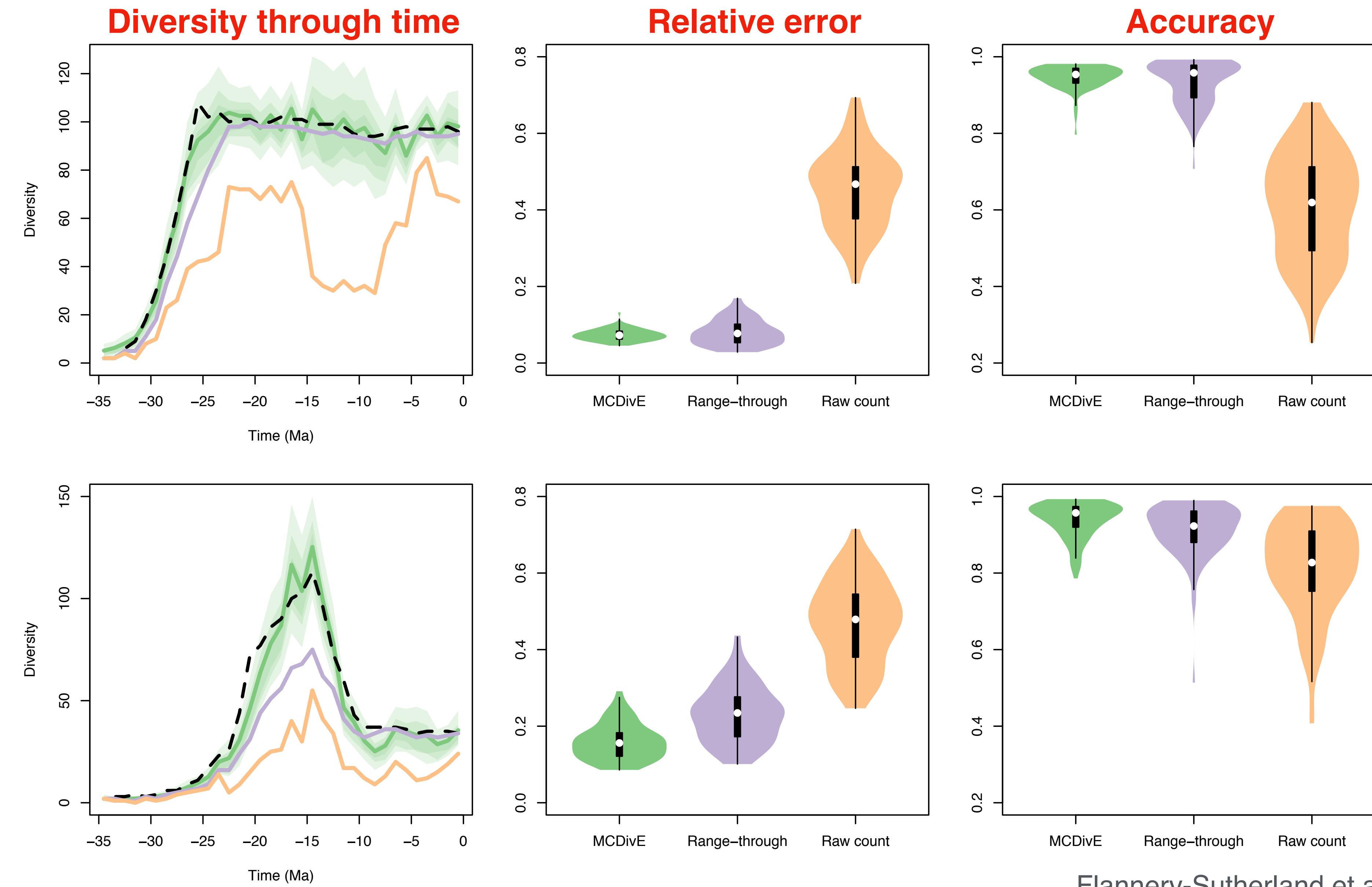
What true diversity could have generated the observed fossil record?



Likelihood function: Binomial distribution

Prior (smoothing) function: Brownian process

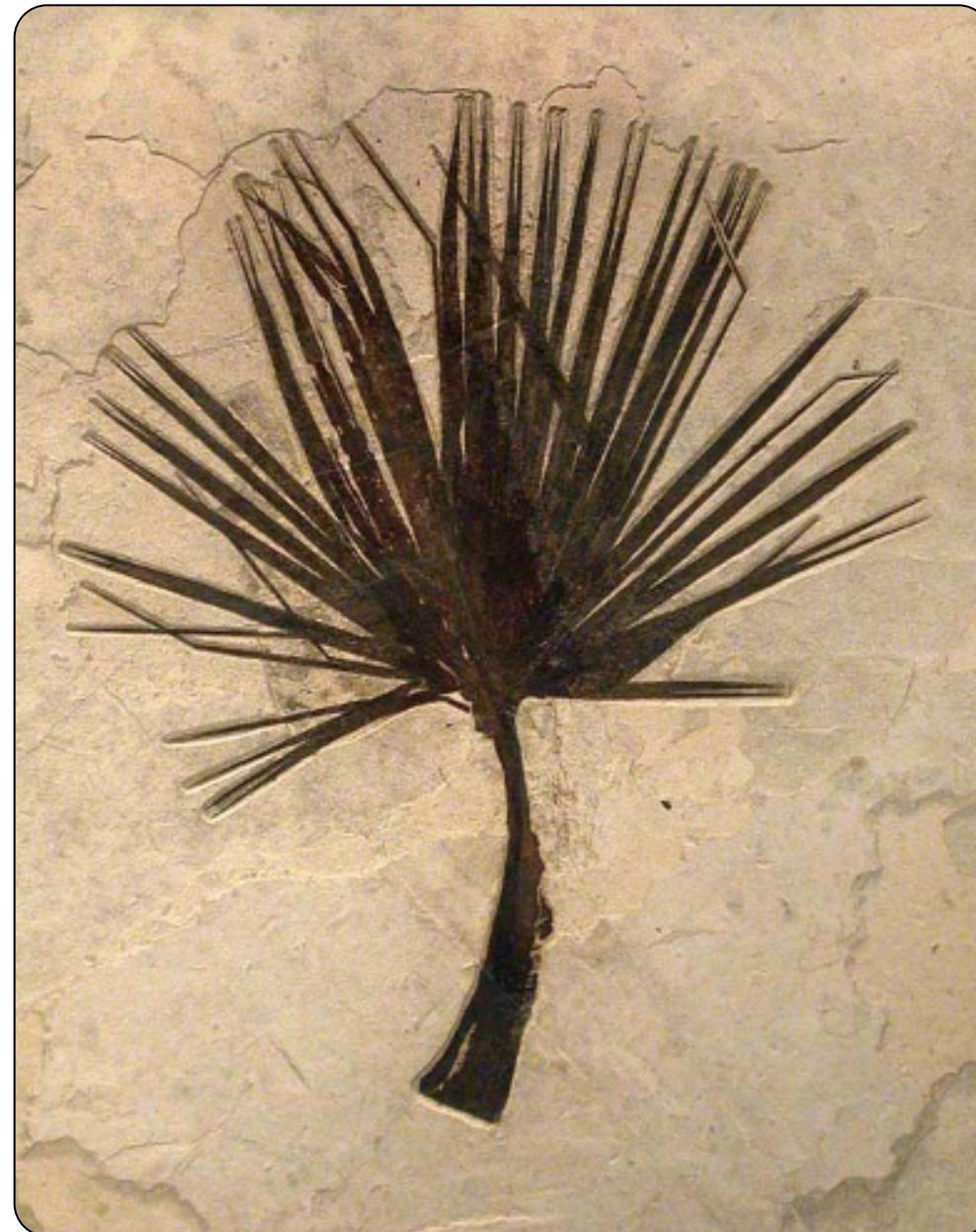
Bayesian estimation of diversity trajectories – mcmcDivE



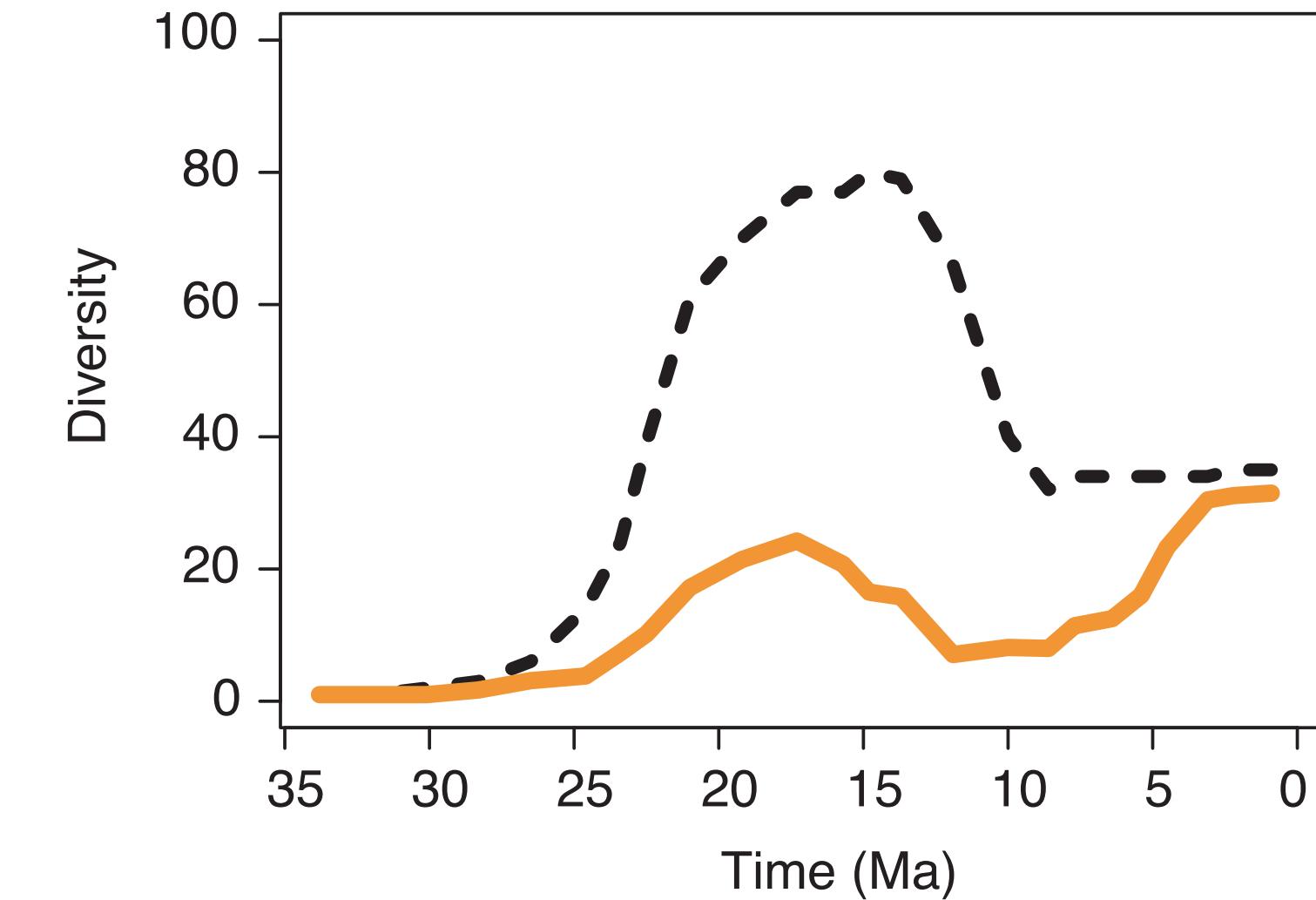
Flannery-Sutherland et al. 2022 Nature Comm

Estimating biodiversity through time from fossil data

Evidence of past biodiversity in the fossil record



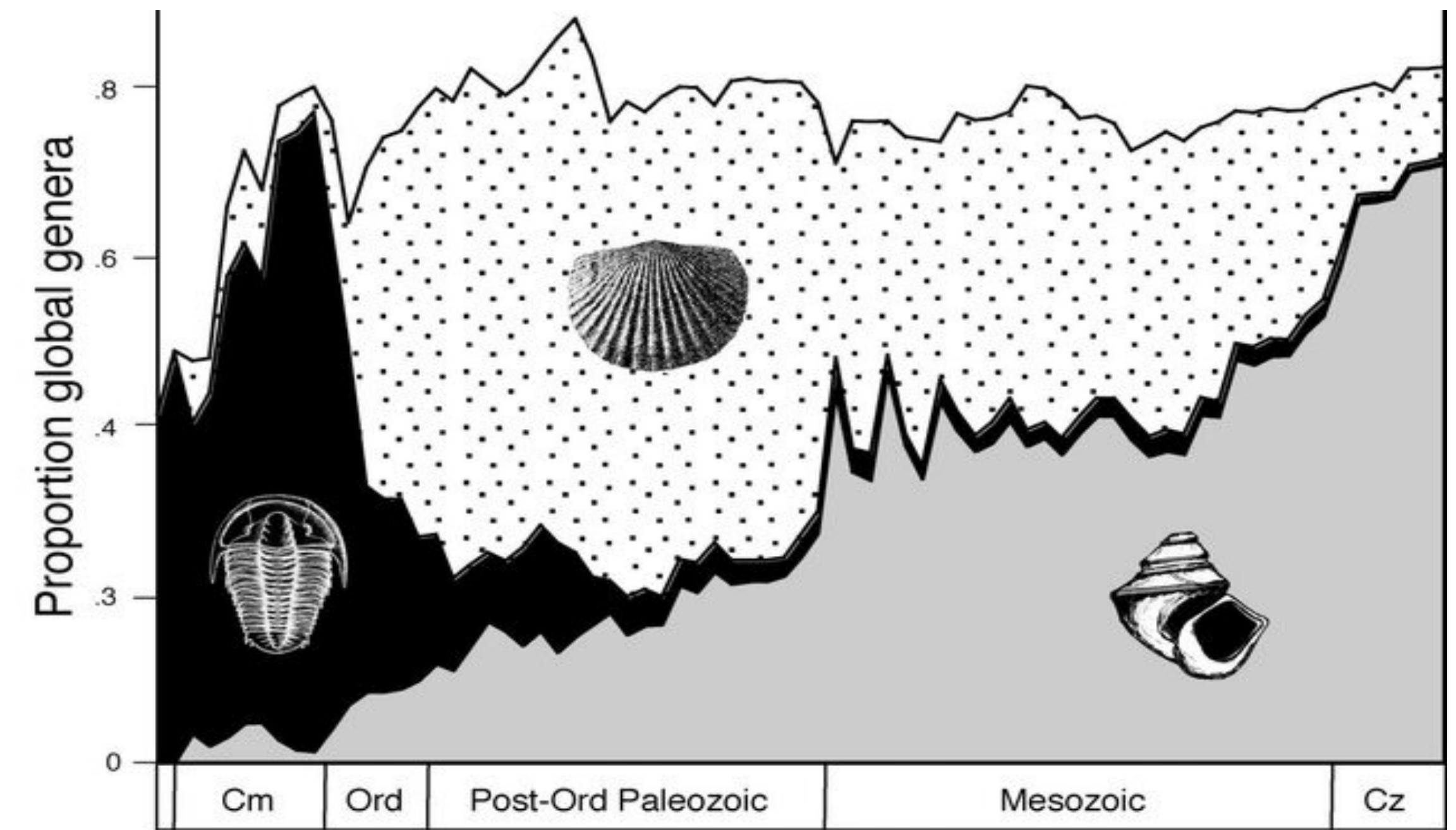
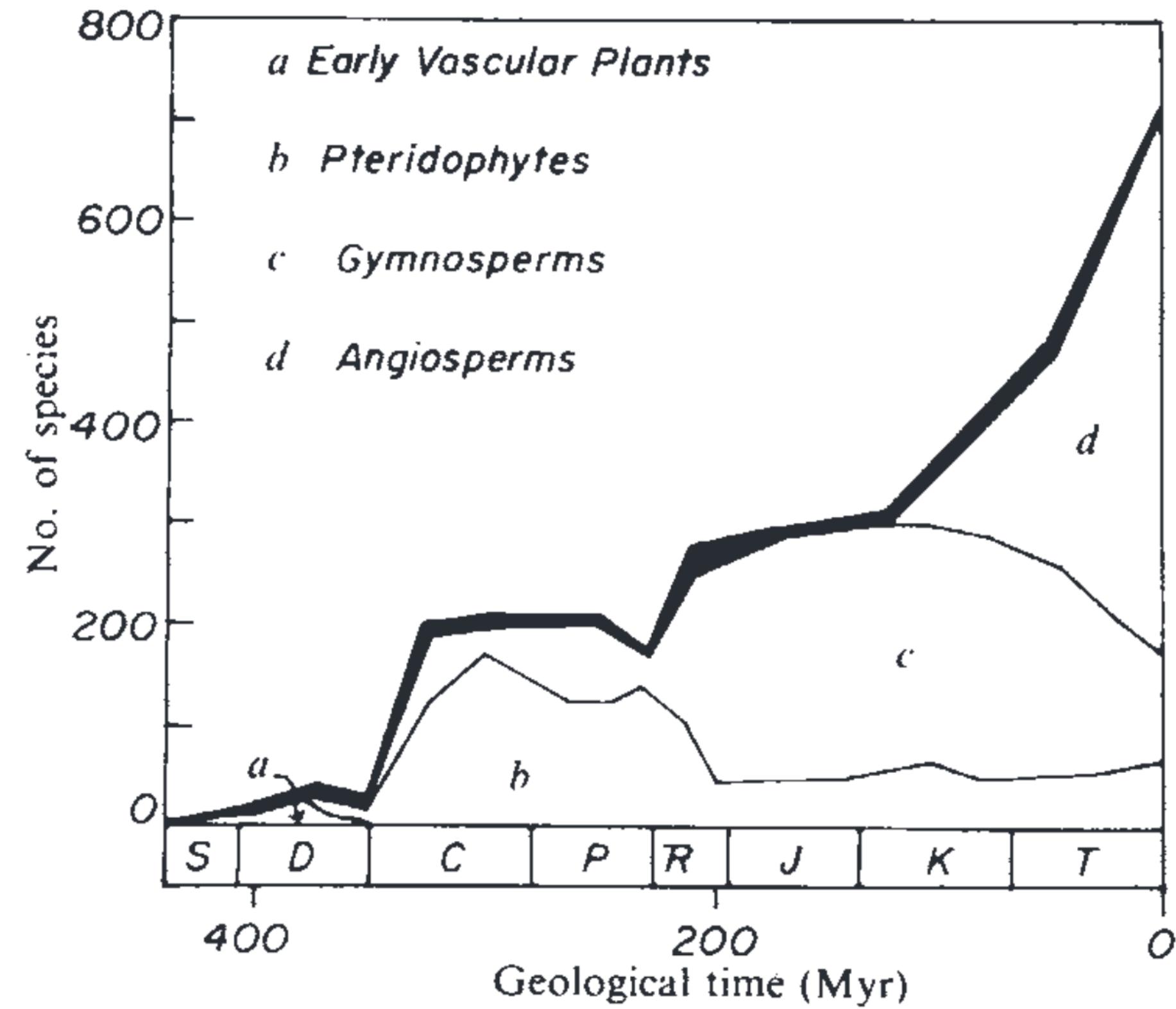
Observed vs true diversity through time



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

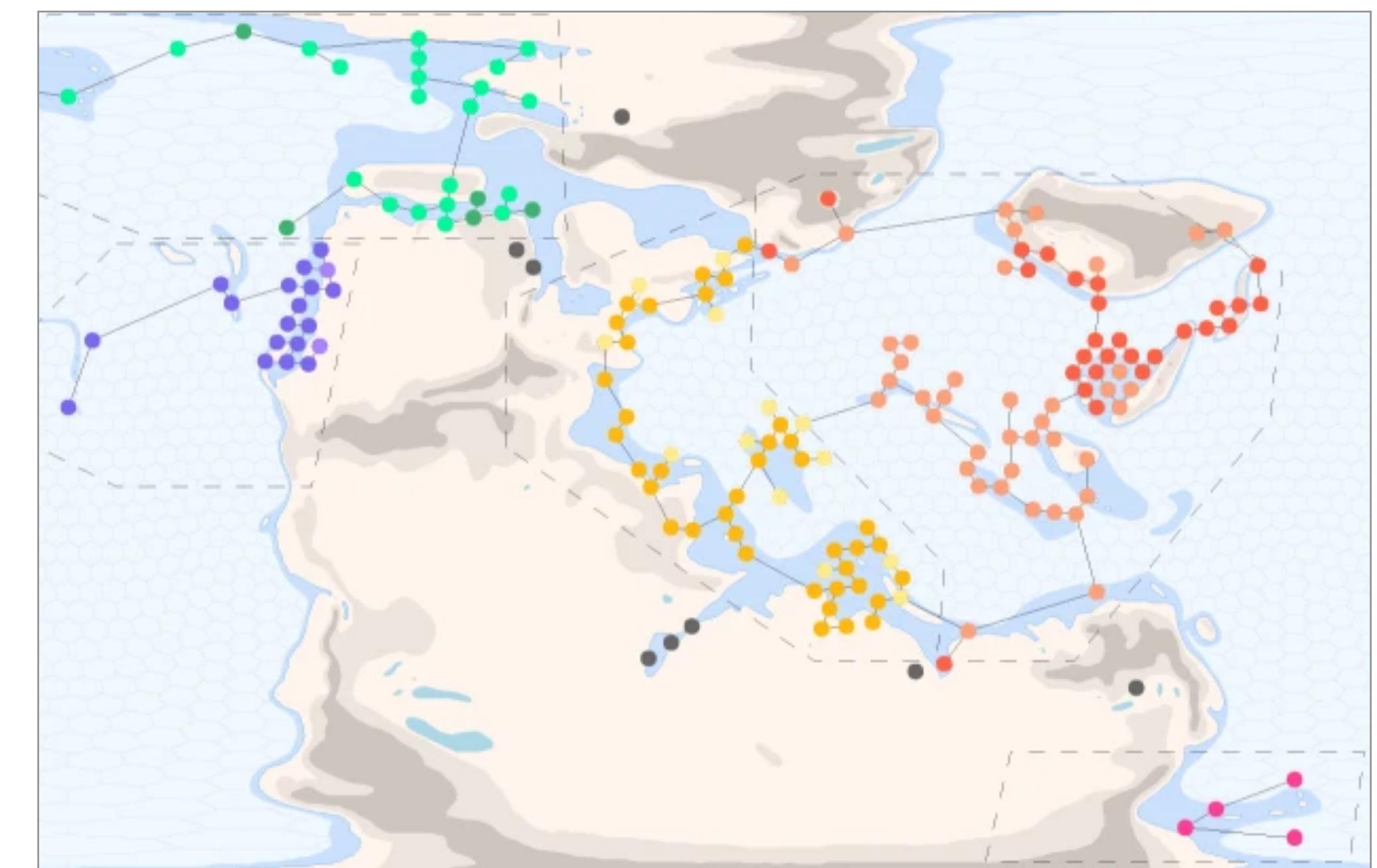
Available methods are based on rarefaction
or simple models assuming random sampling

Inferring biodiversity dynamics from the fossil record



Problems in estimating biodiversity through time from fossil data

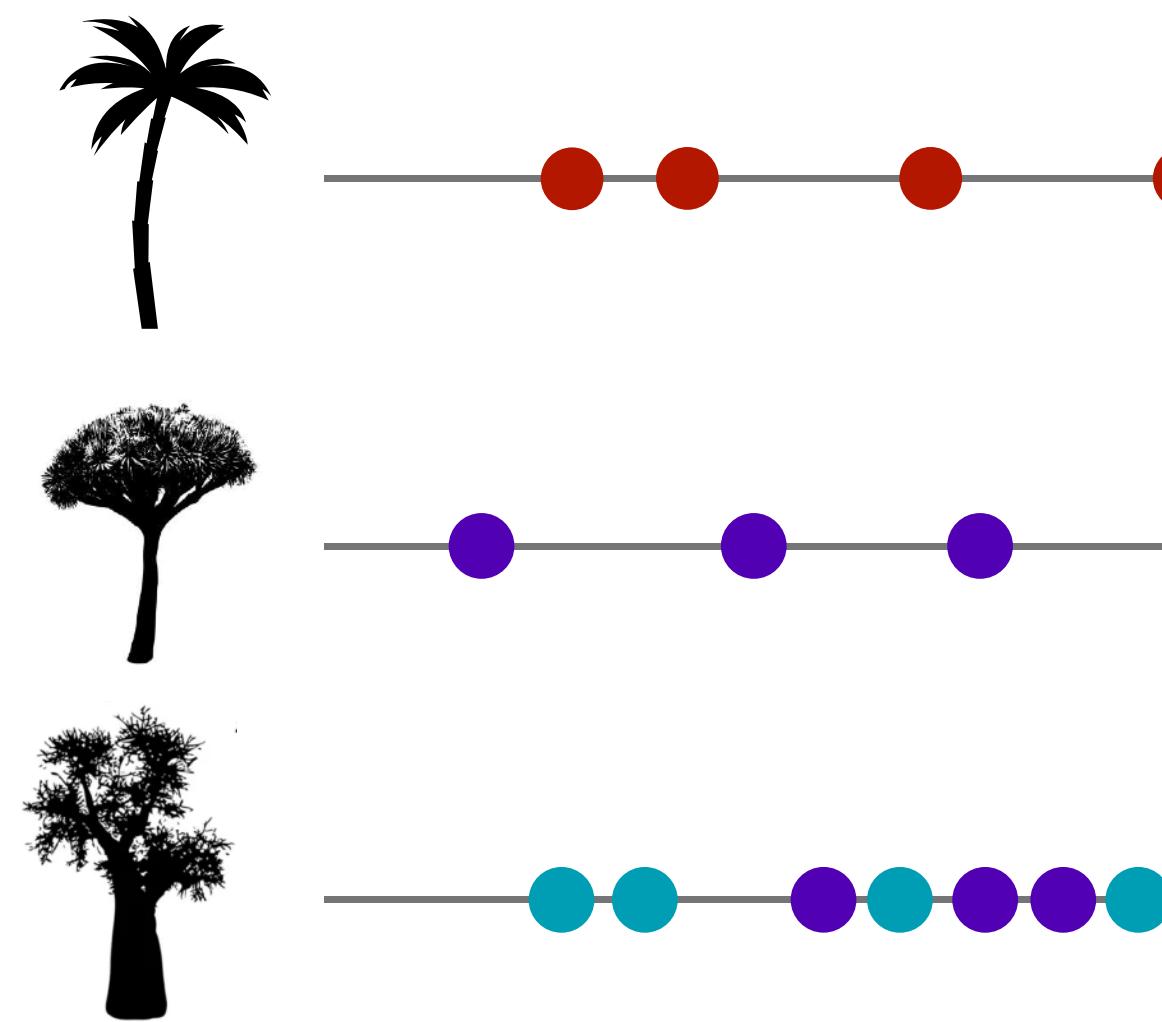
Spatial biases make global biodiversity patterns unidentifiable using the current methods



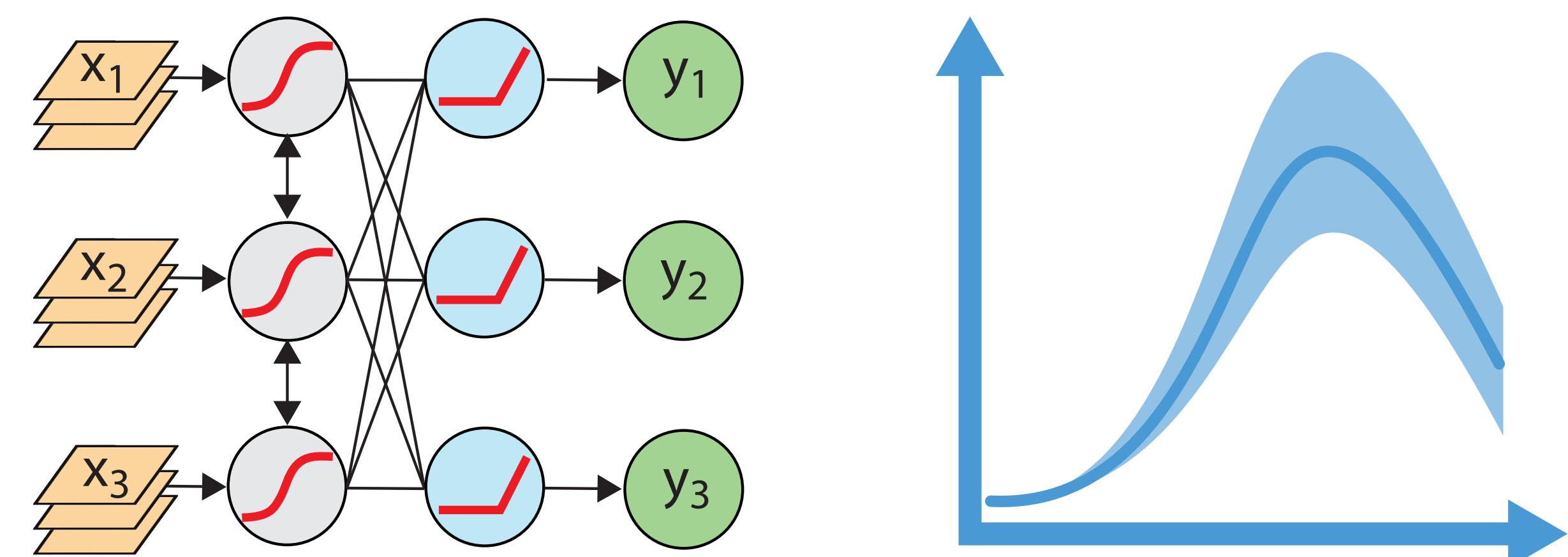
Alroy 2010 Science; Starrfelt & Liow 2016 Phil Trans B; Flannery-Sutherland et al. 2022 Nature Comm

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

Input: fossil distribution in space and time



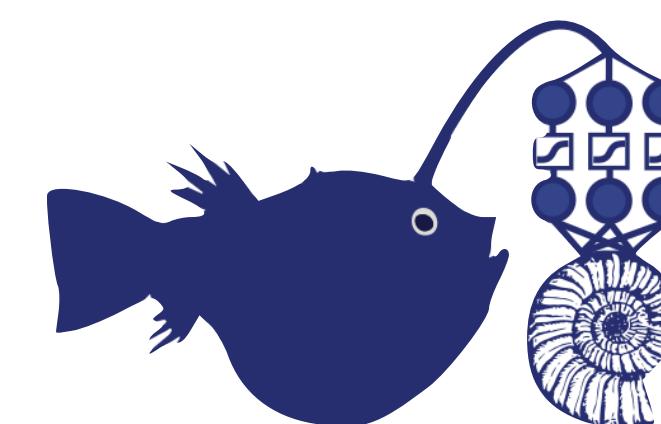
Output: diversity estimates through time



R Cooper



B Allen



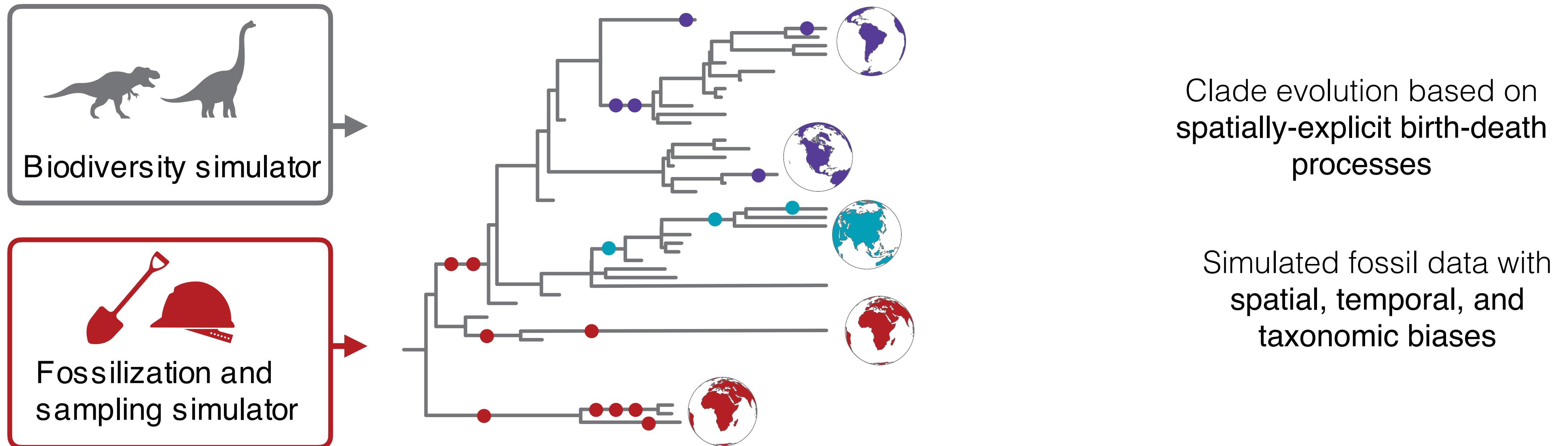
github.com/DeepDive-project

Cooper et al. 2024 Nature Comms

Application note: Cooper et al. 2024 bioRxiv

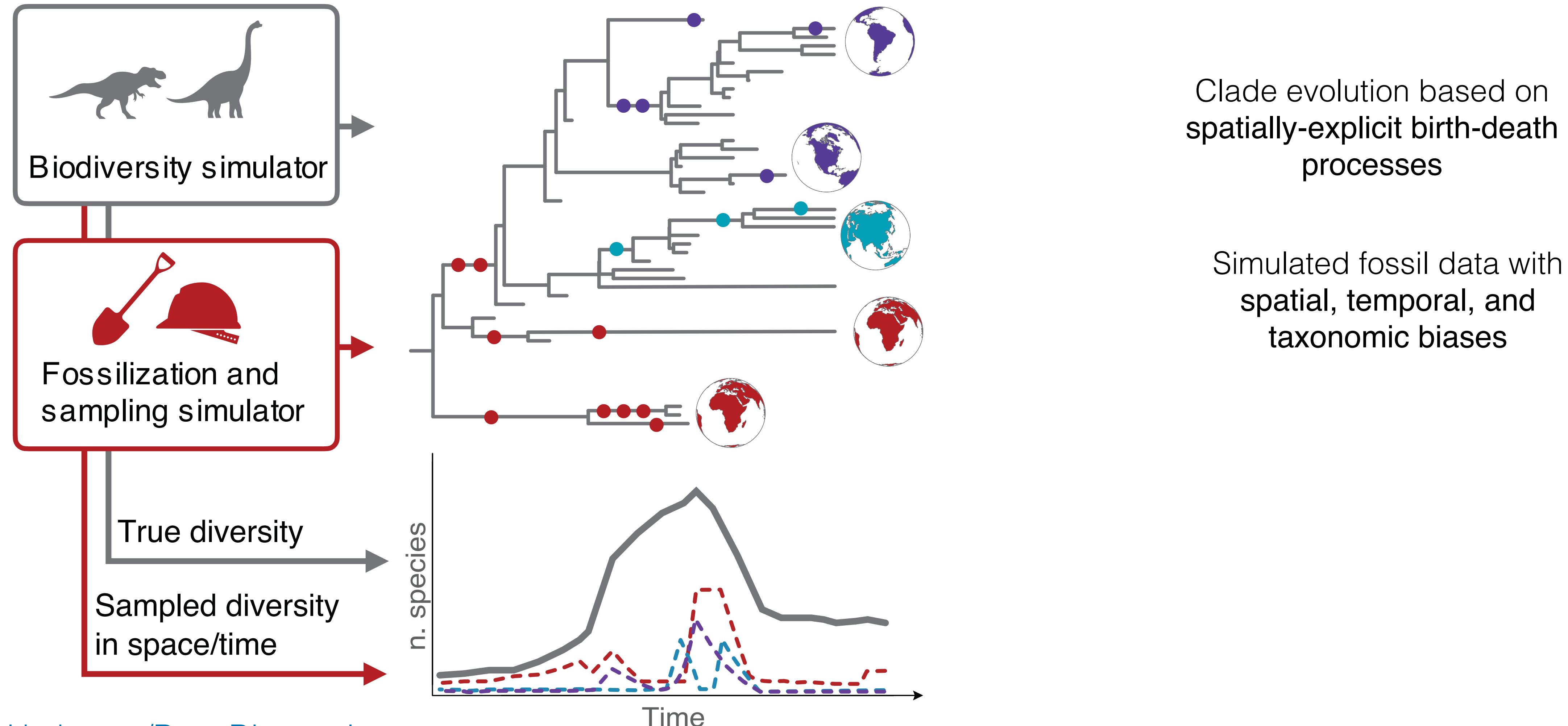
DeepDive – Deep learning models to estimate Diversity trajectories

Mechanistic model of species diversification and extinction in time and space



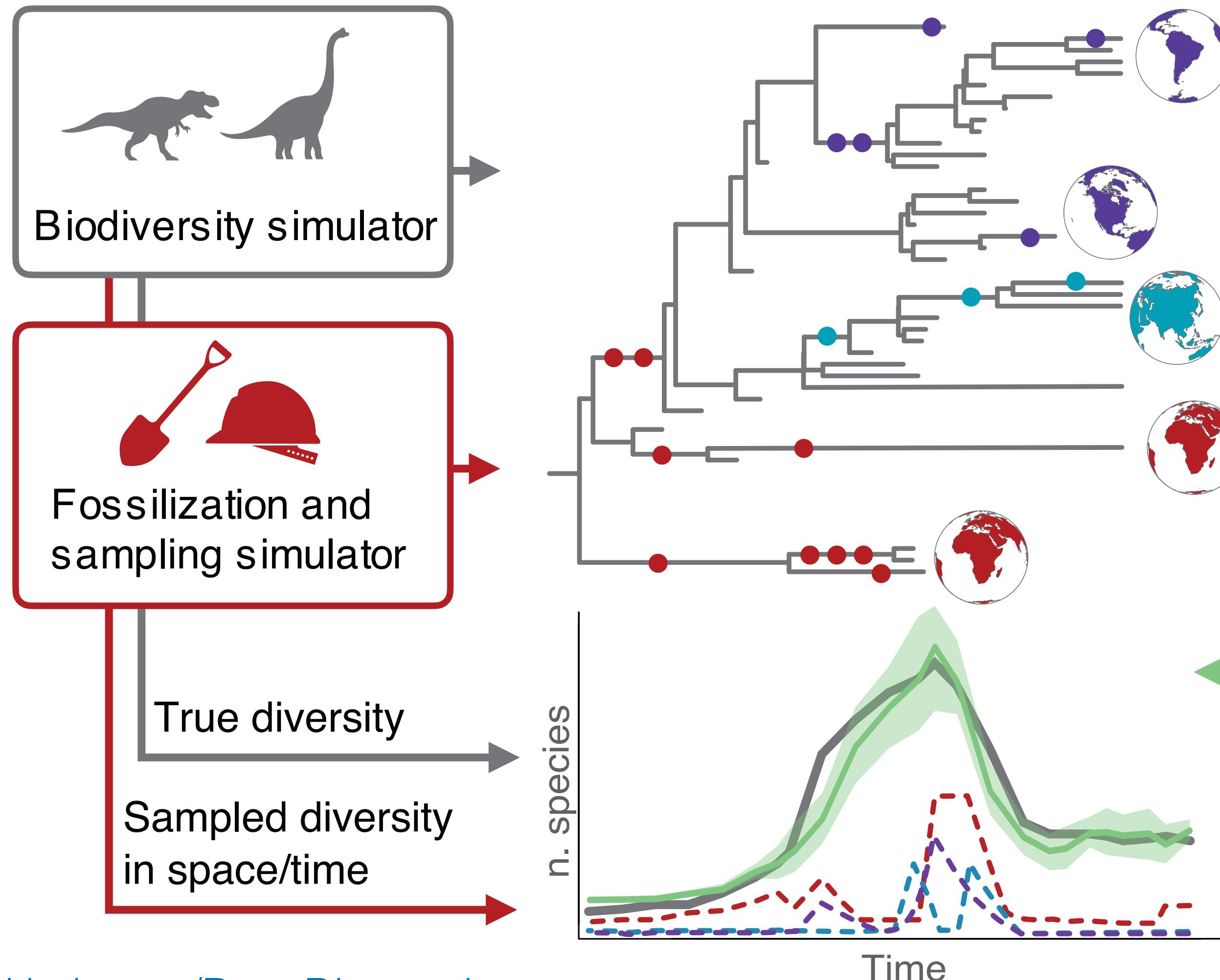
DeepDive – Deep learning models to estimate Diversity trajectories

Mechanistic model of species diversification and extinction in time and space

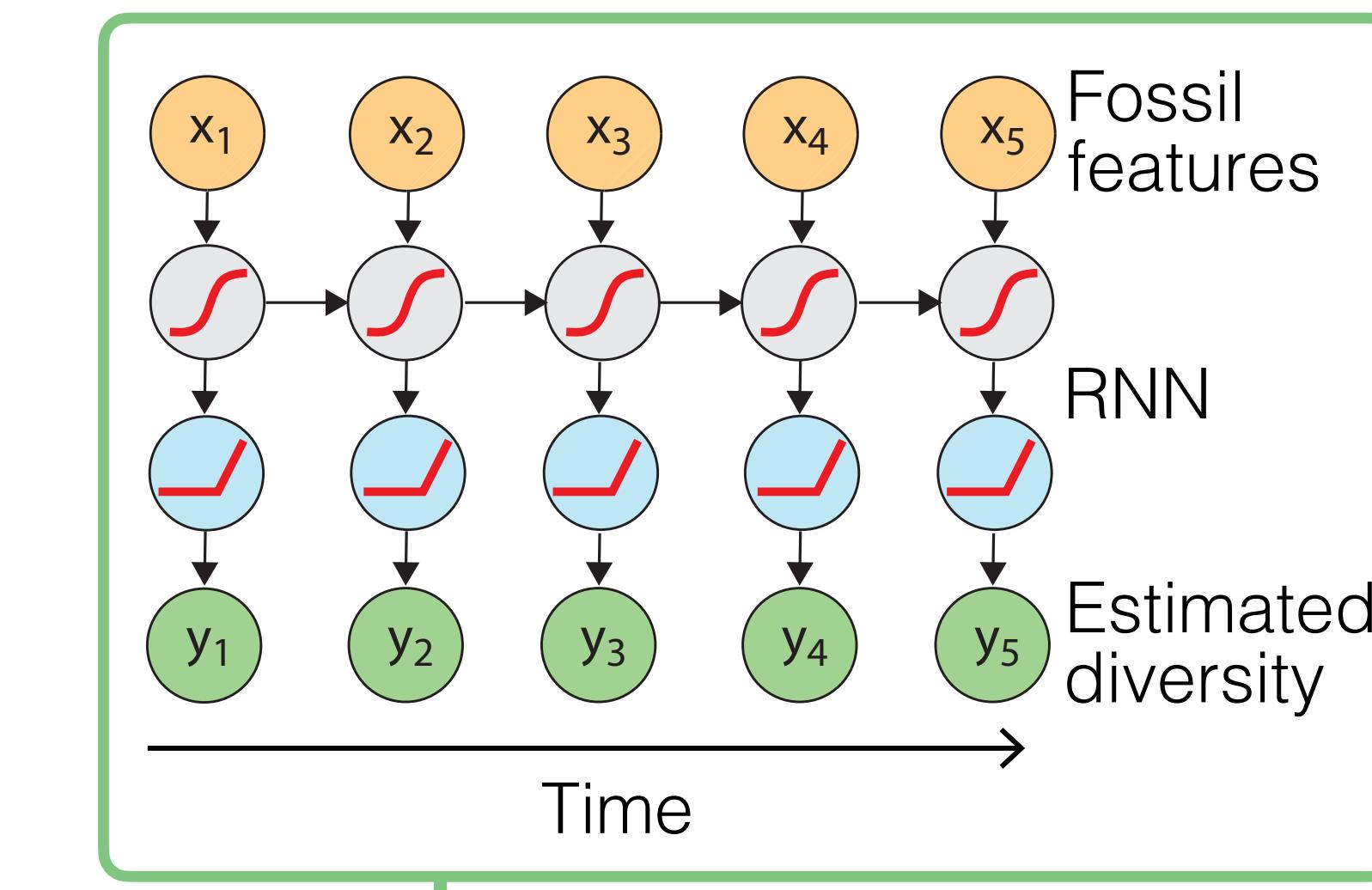


Supervised deep learning models to estimate diversity trajectories

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



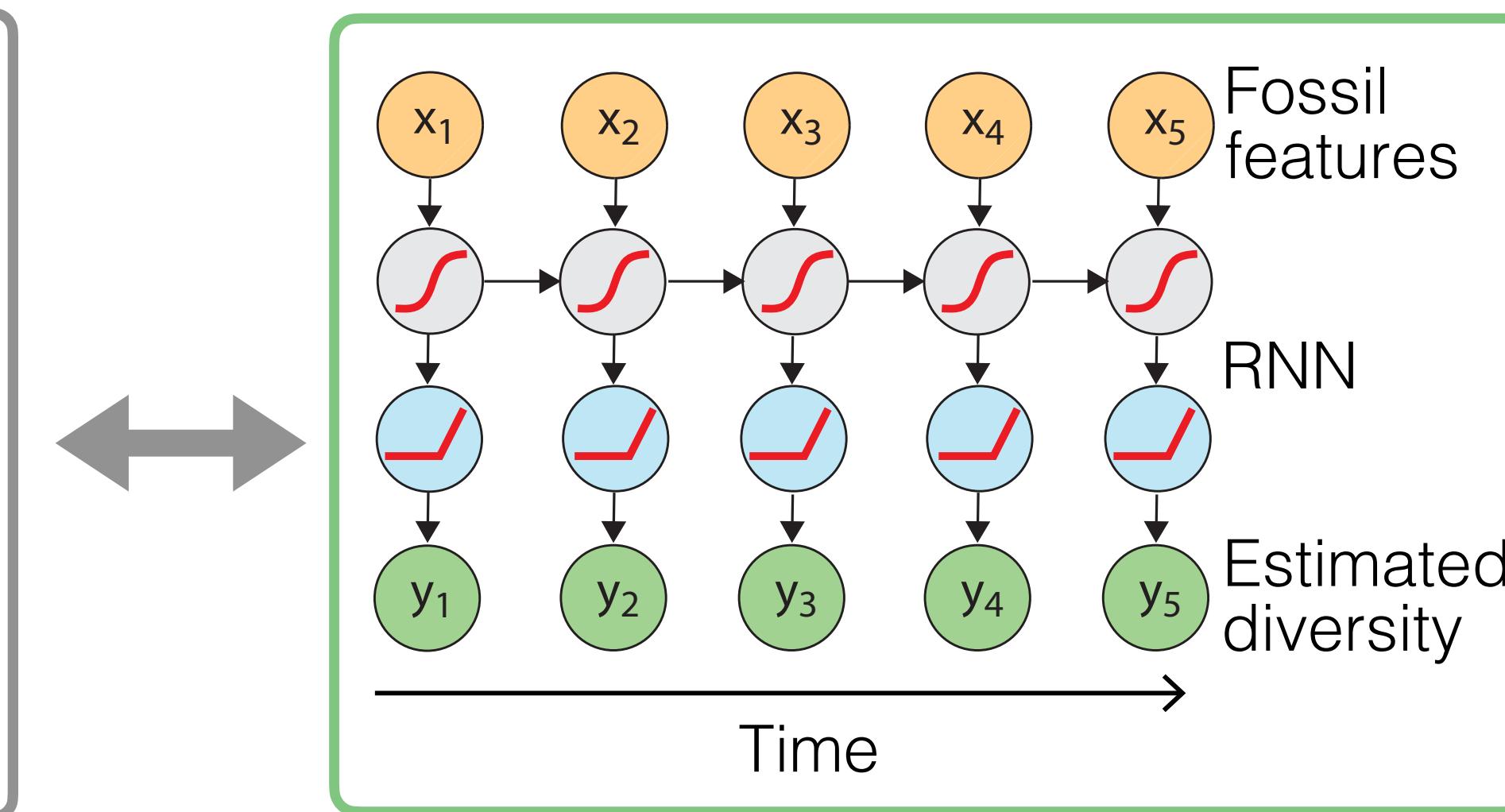
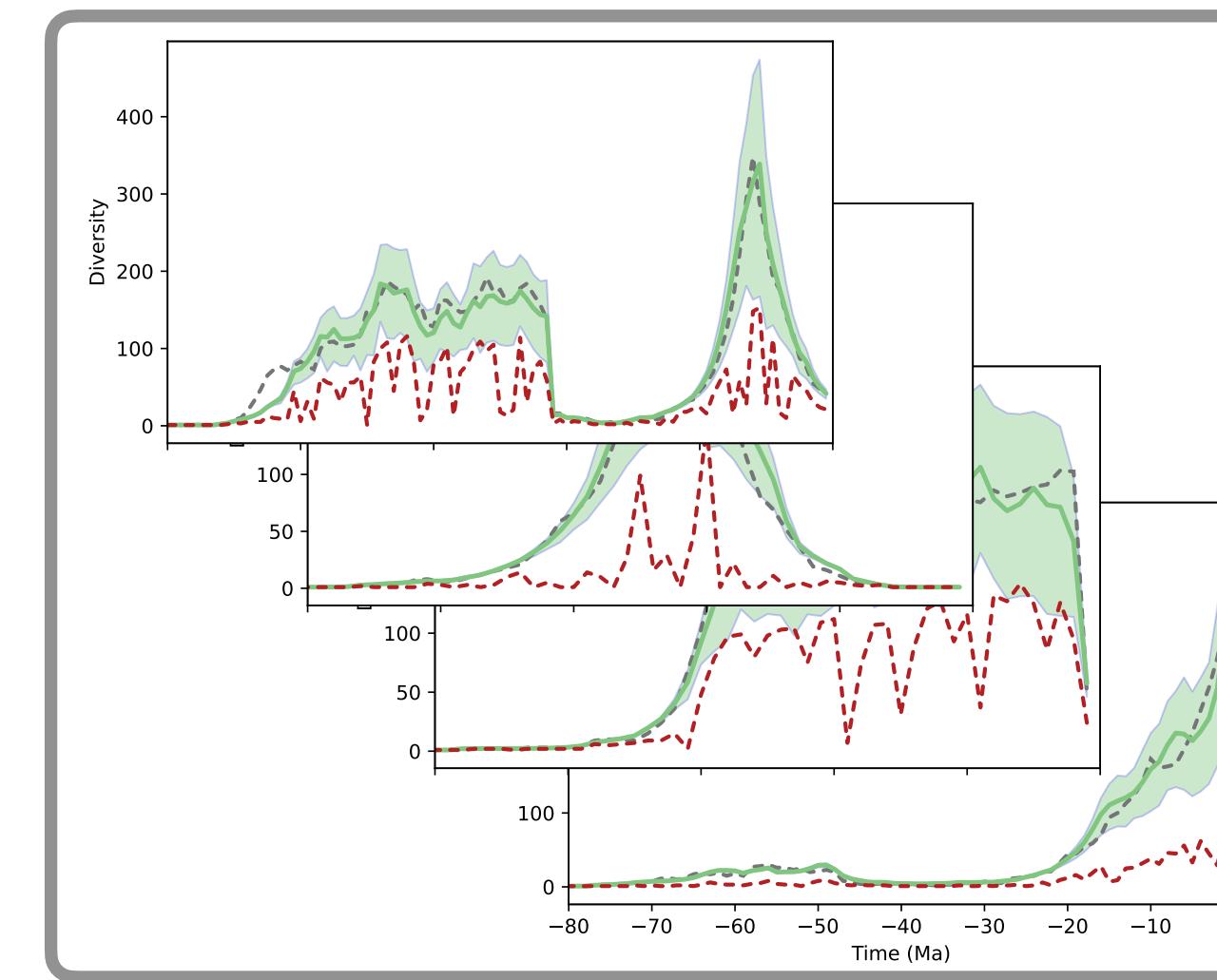
Deep learning model predicting diversity time series from fossil features



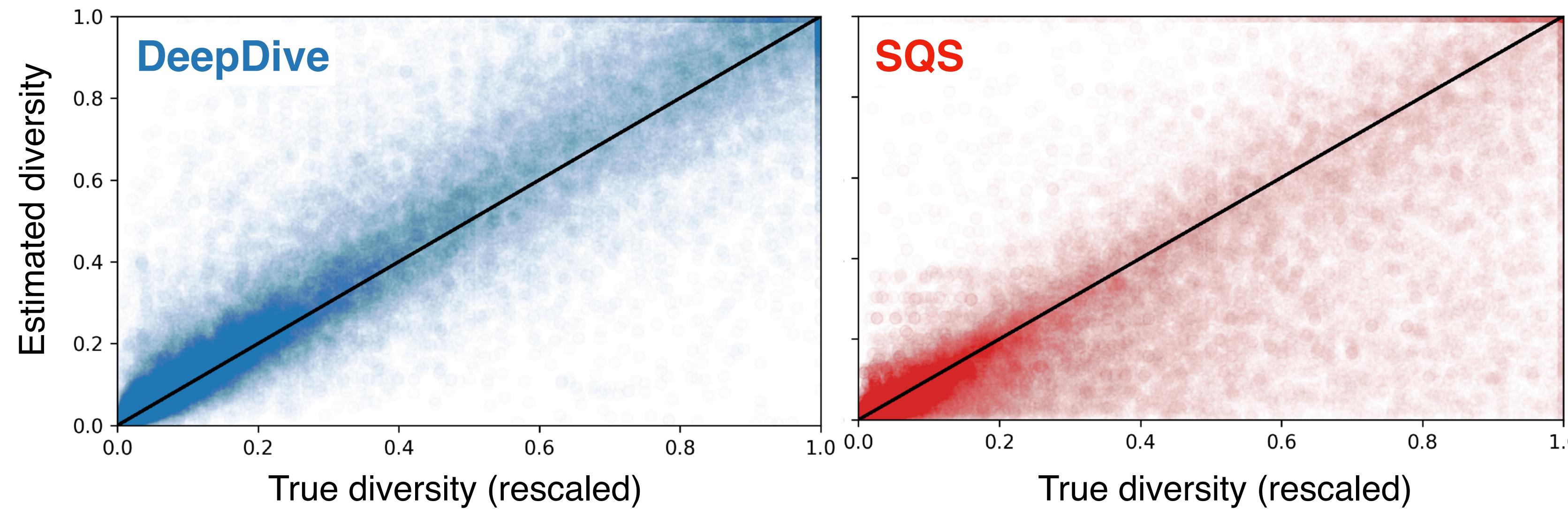
DeepDive – Deep learning models to estimate Diversity trajectories

Model optimization (training) and validation

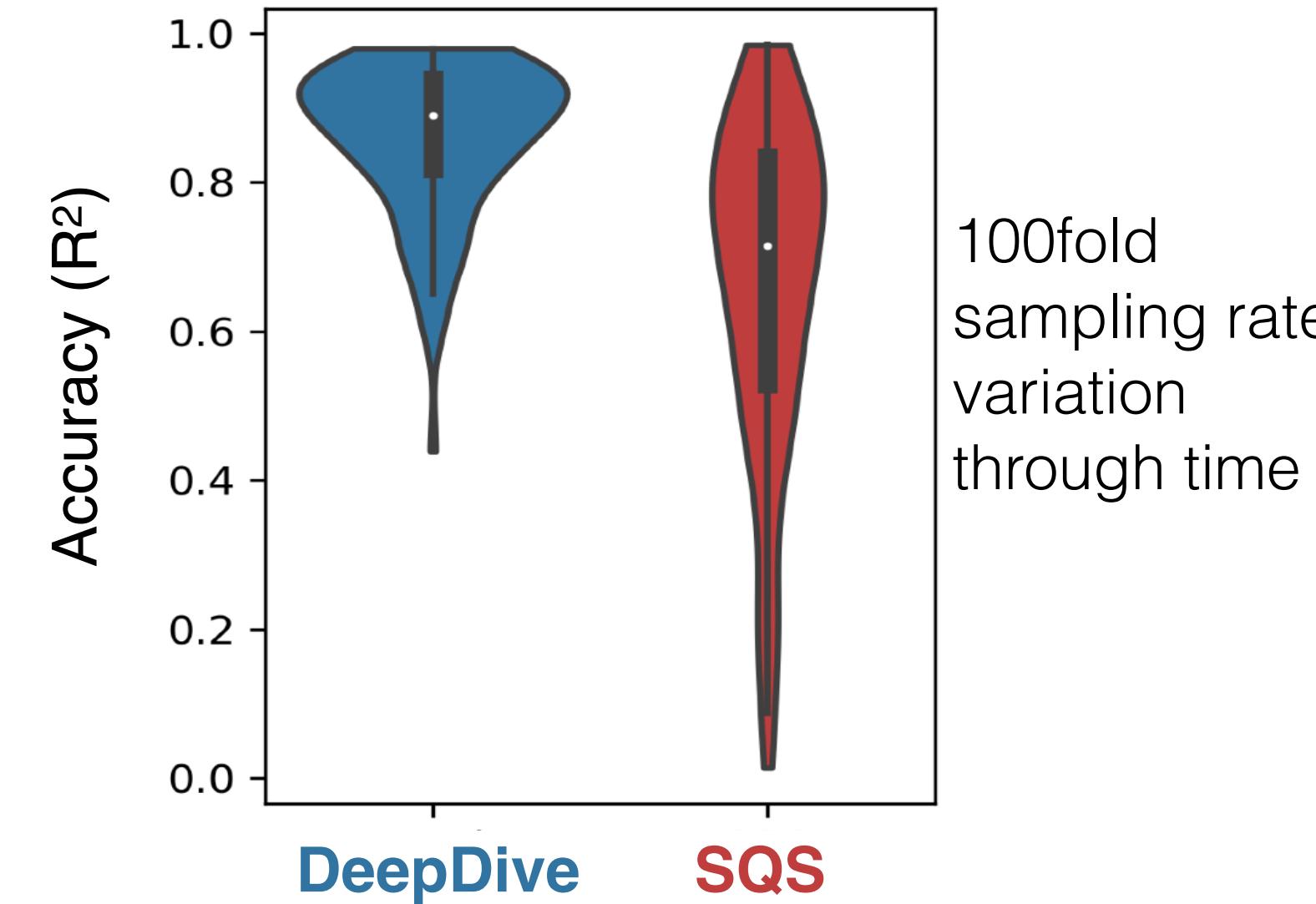
Samples from ‘prior’ distributions of speciation, extinction, migration, preservation rates



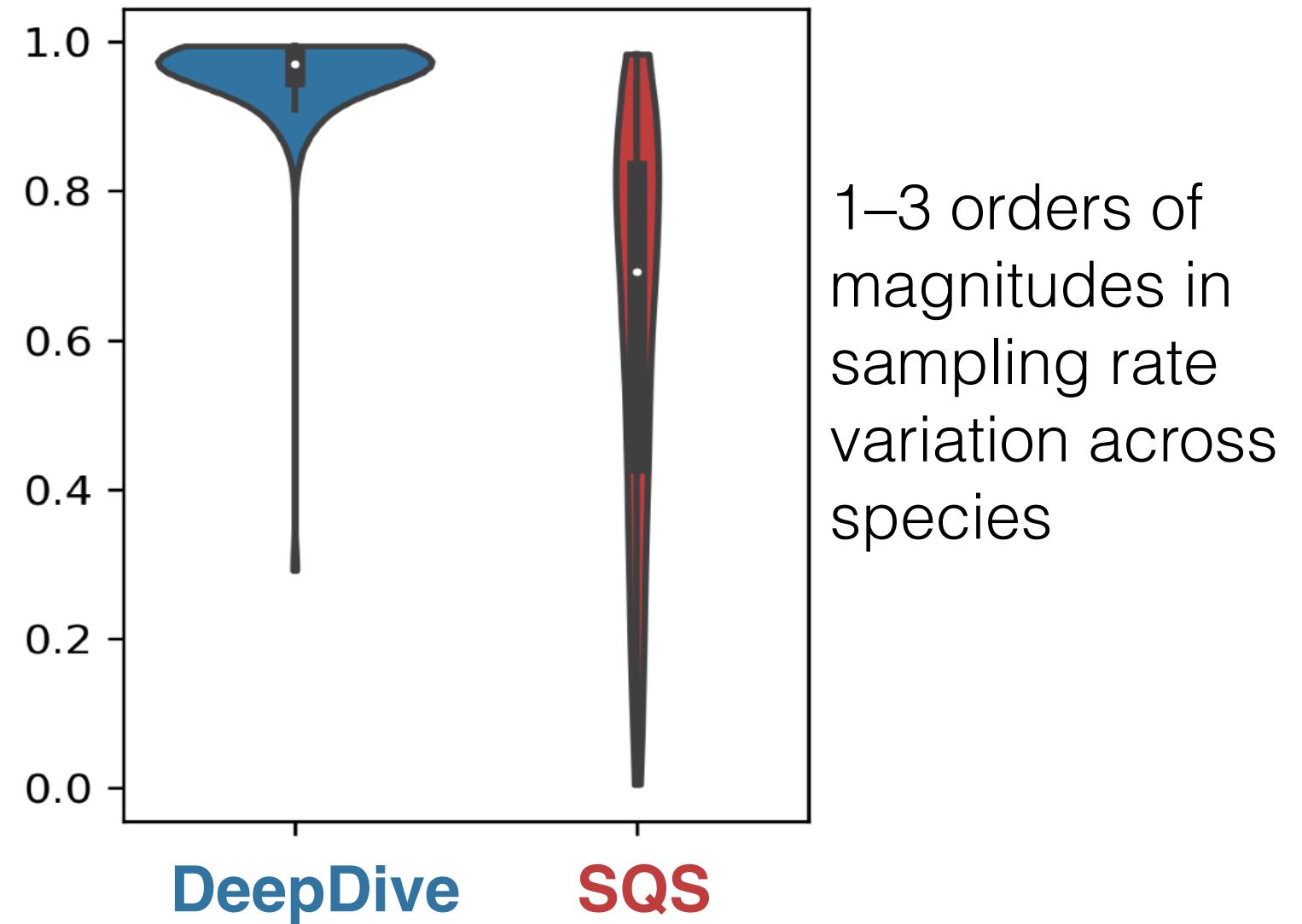
DeepDive performance



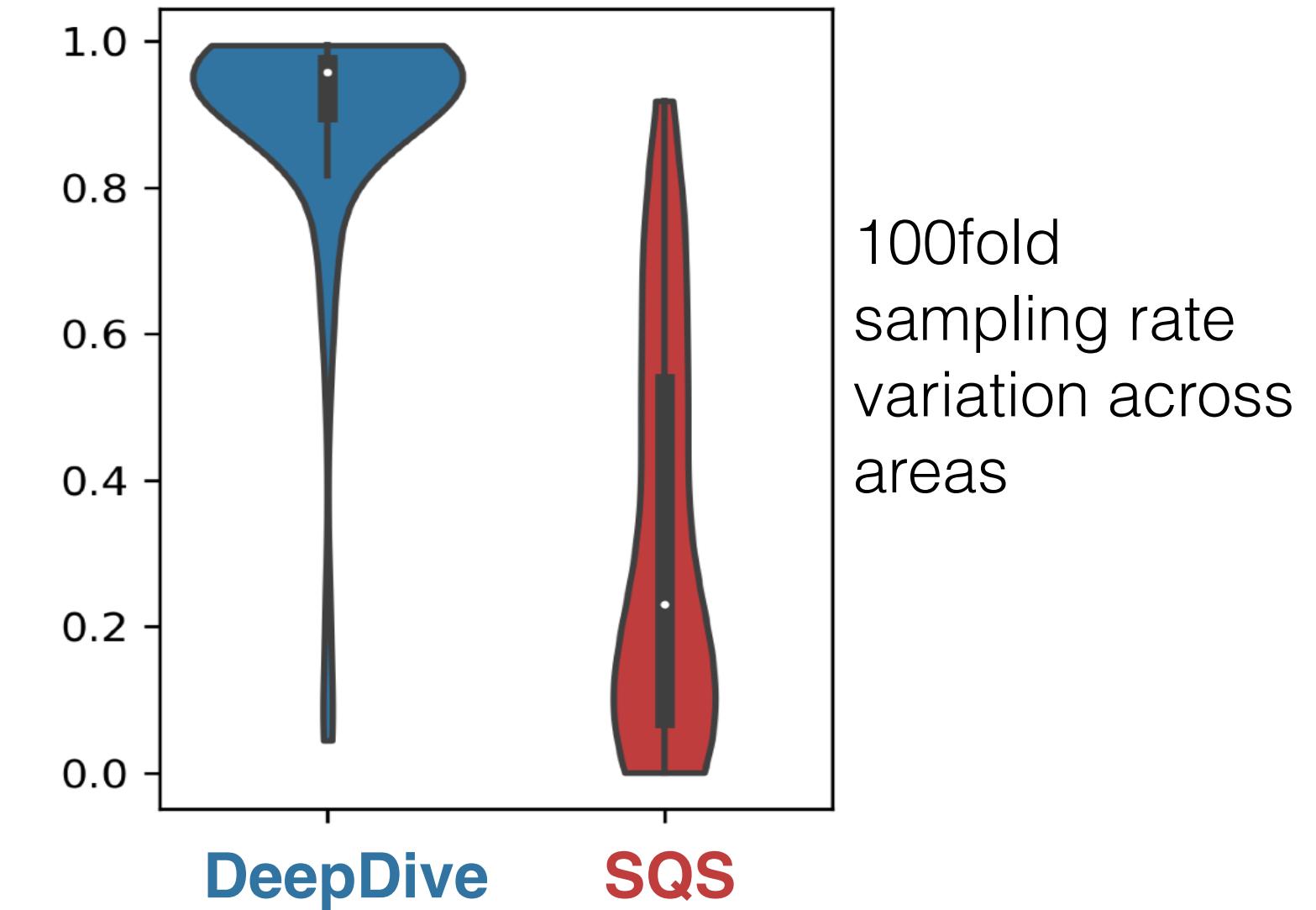
Strong temporal bias



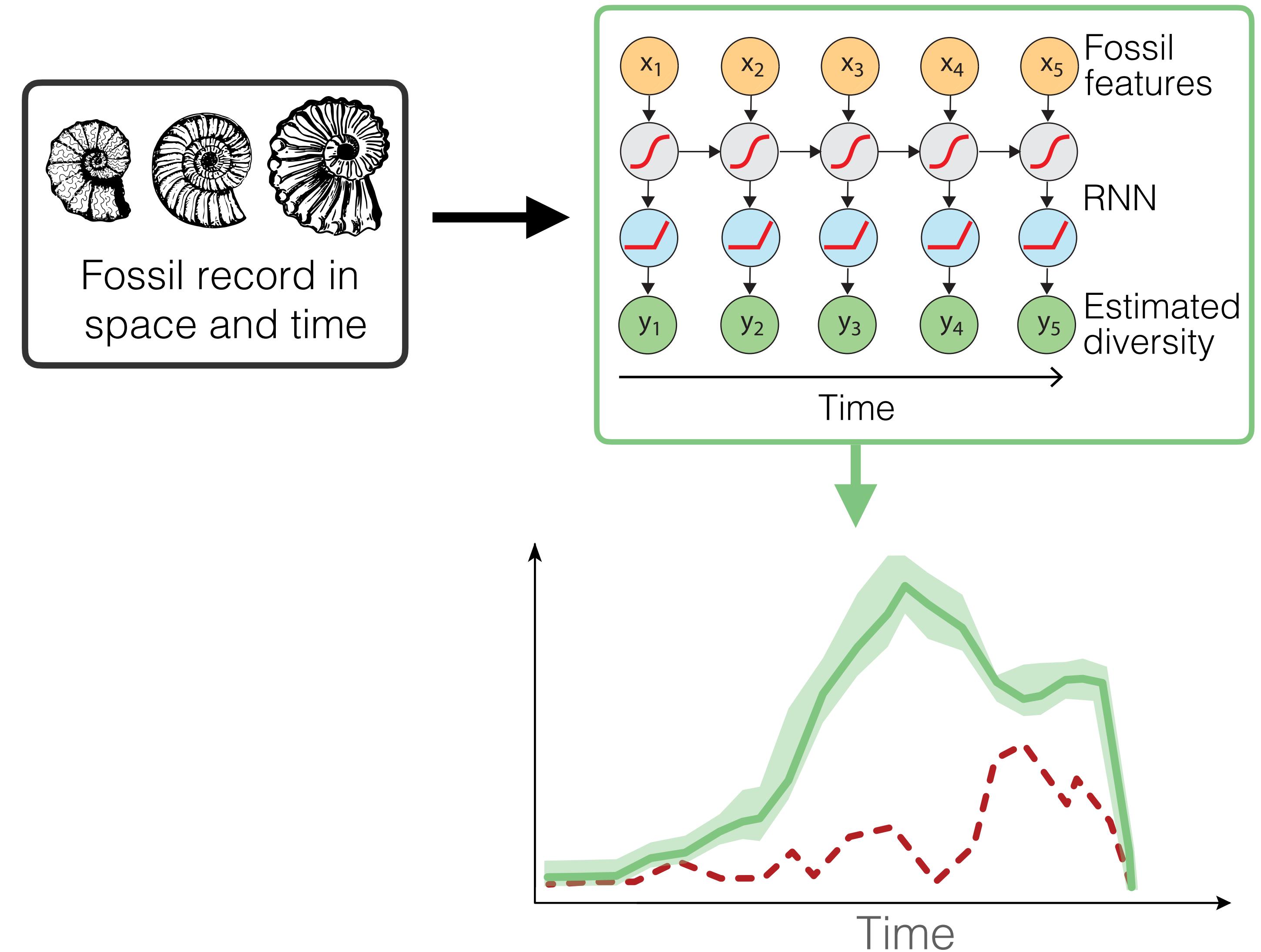
Strong taxonomic bias



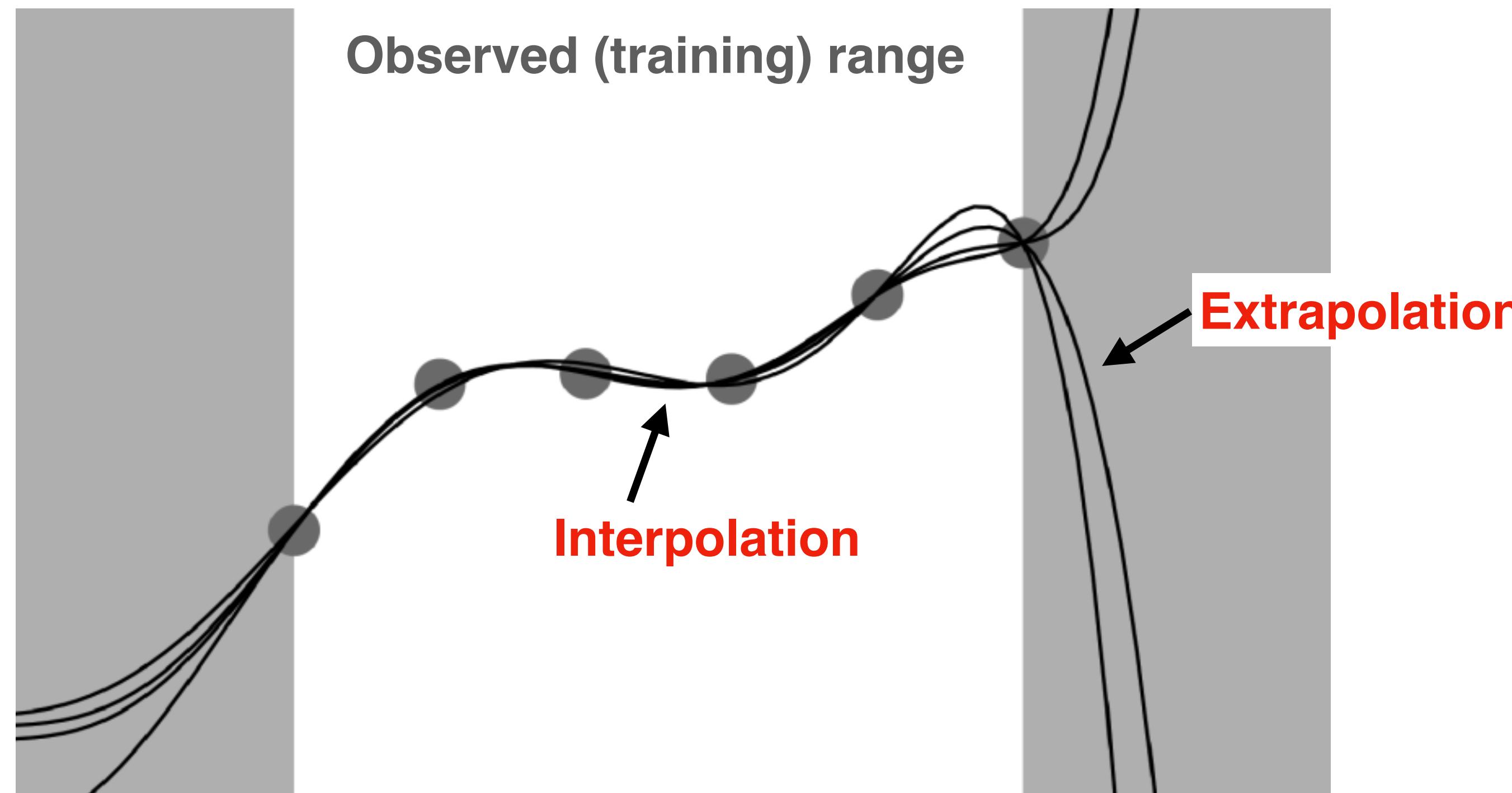
Strong spatial bias



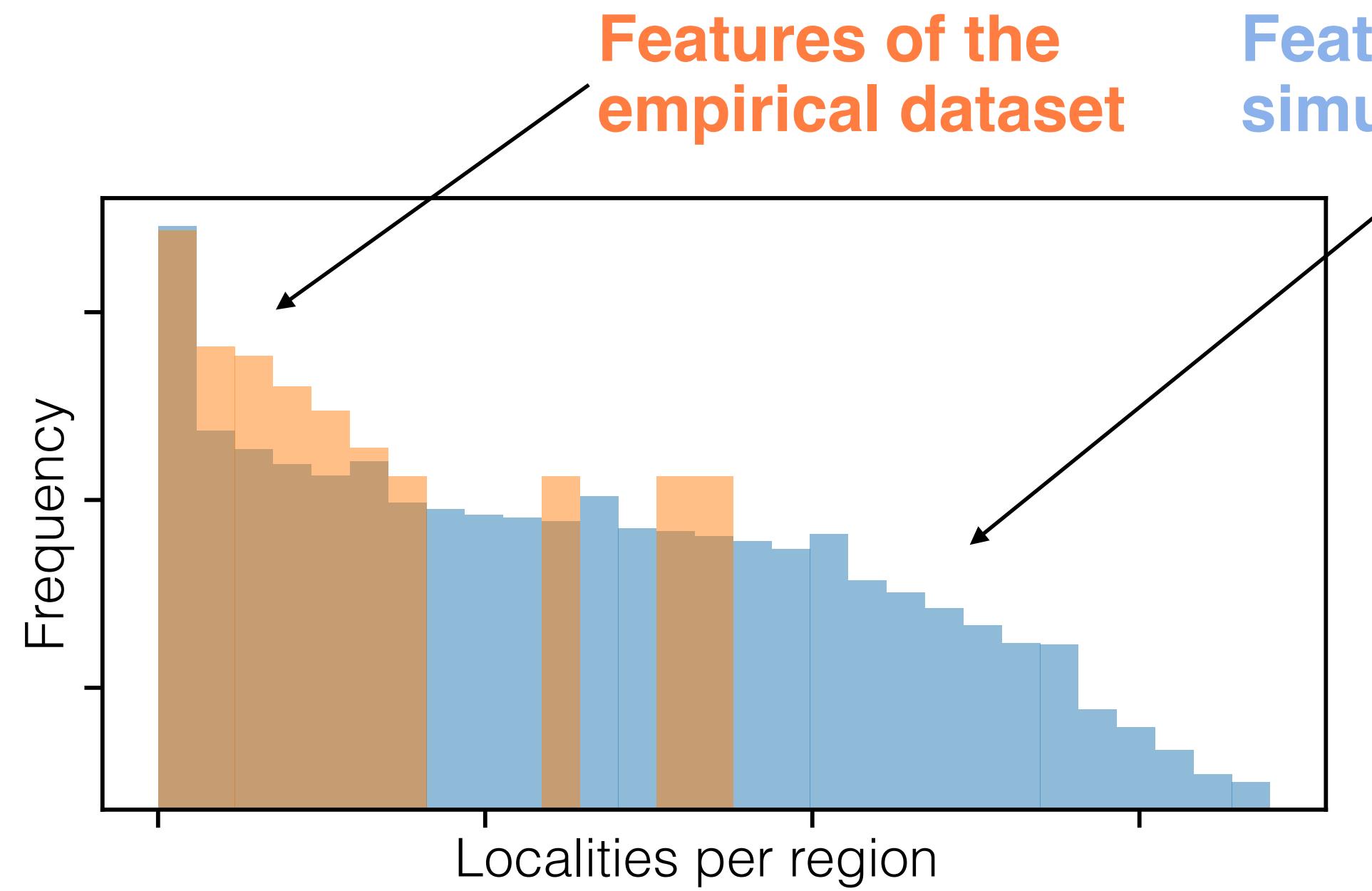
DeepDive – Predictions of empirical data



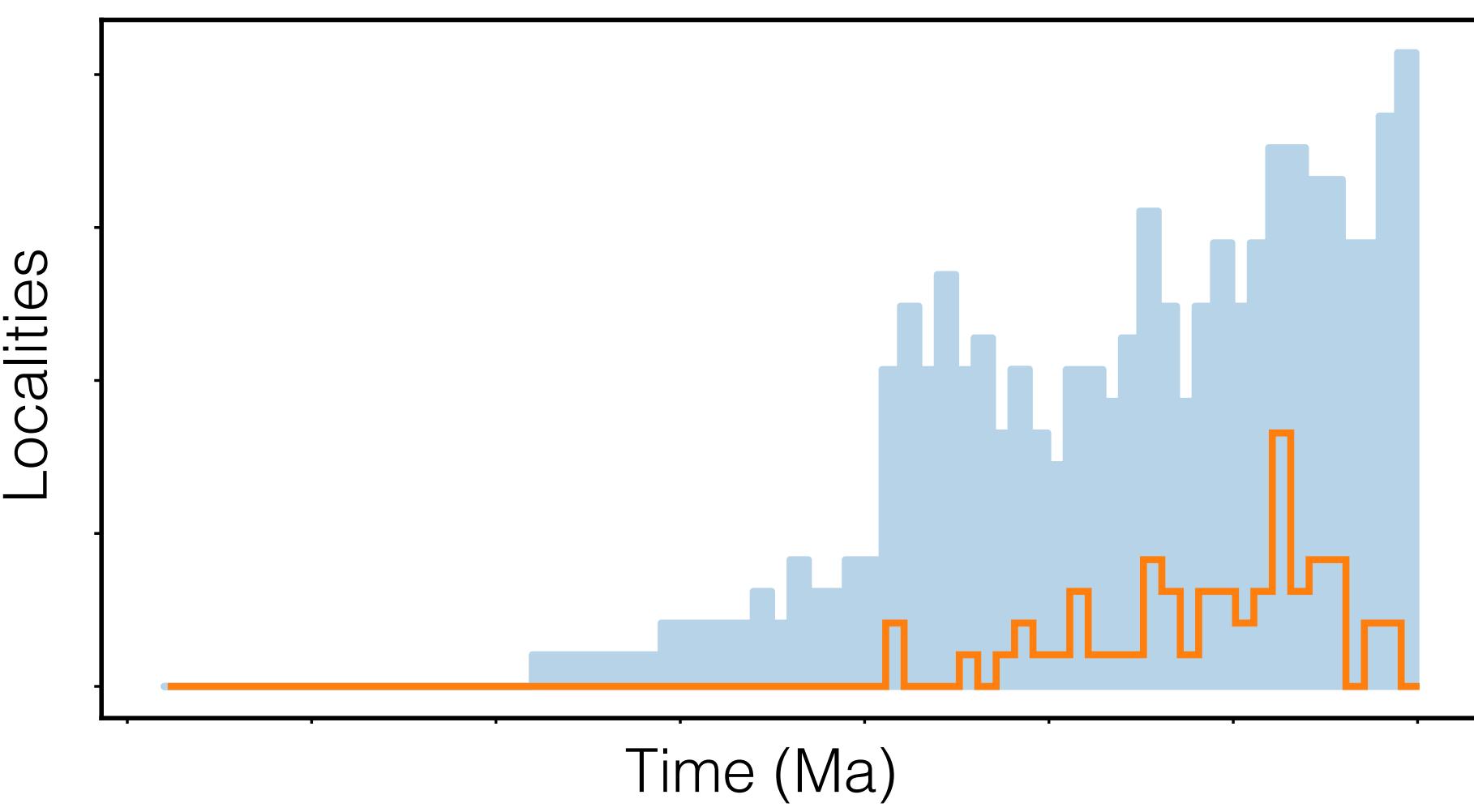
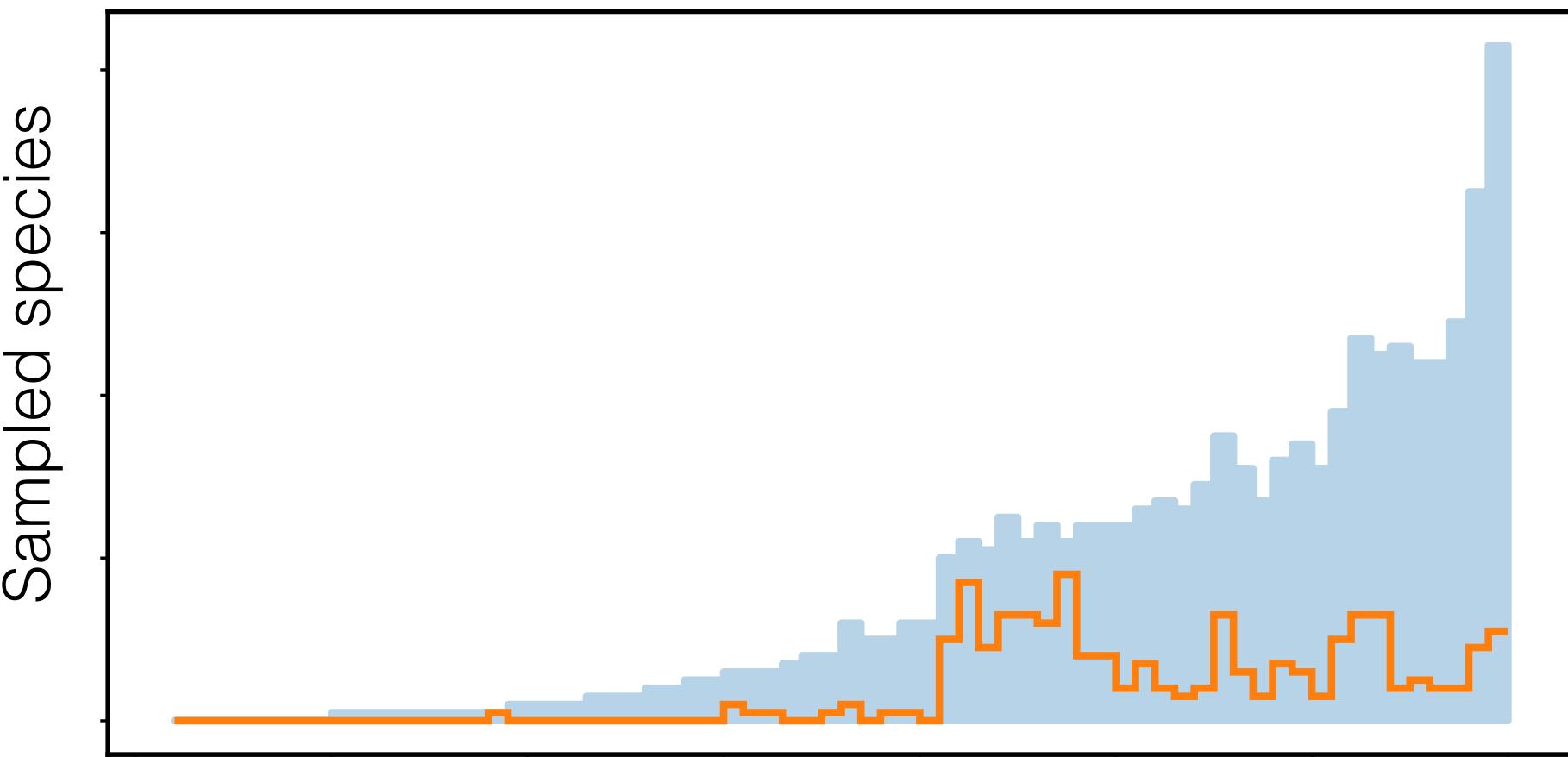
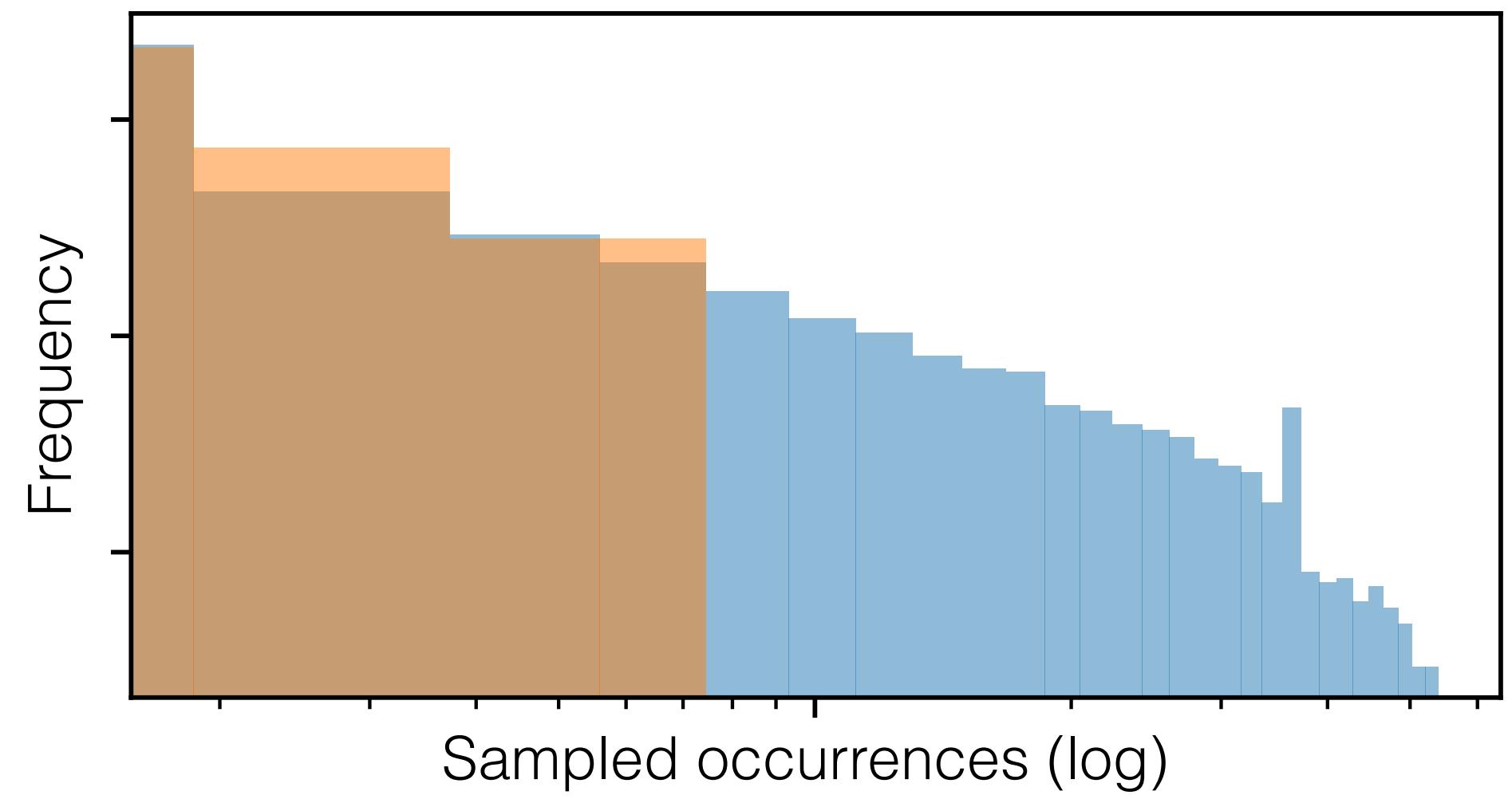
Caveat: the training simulations must resemble the true data



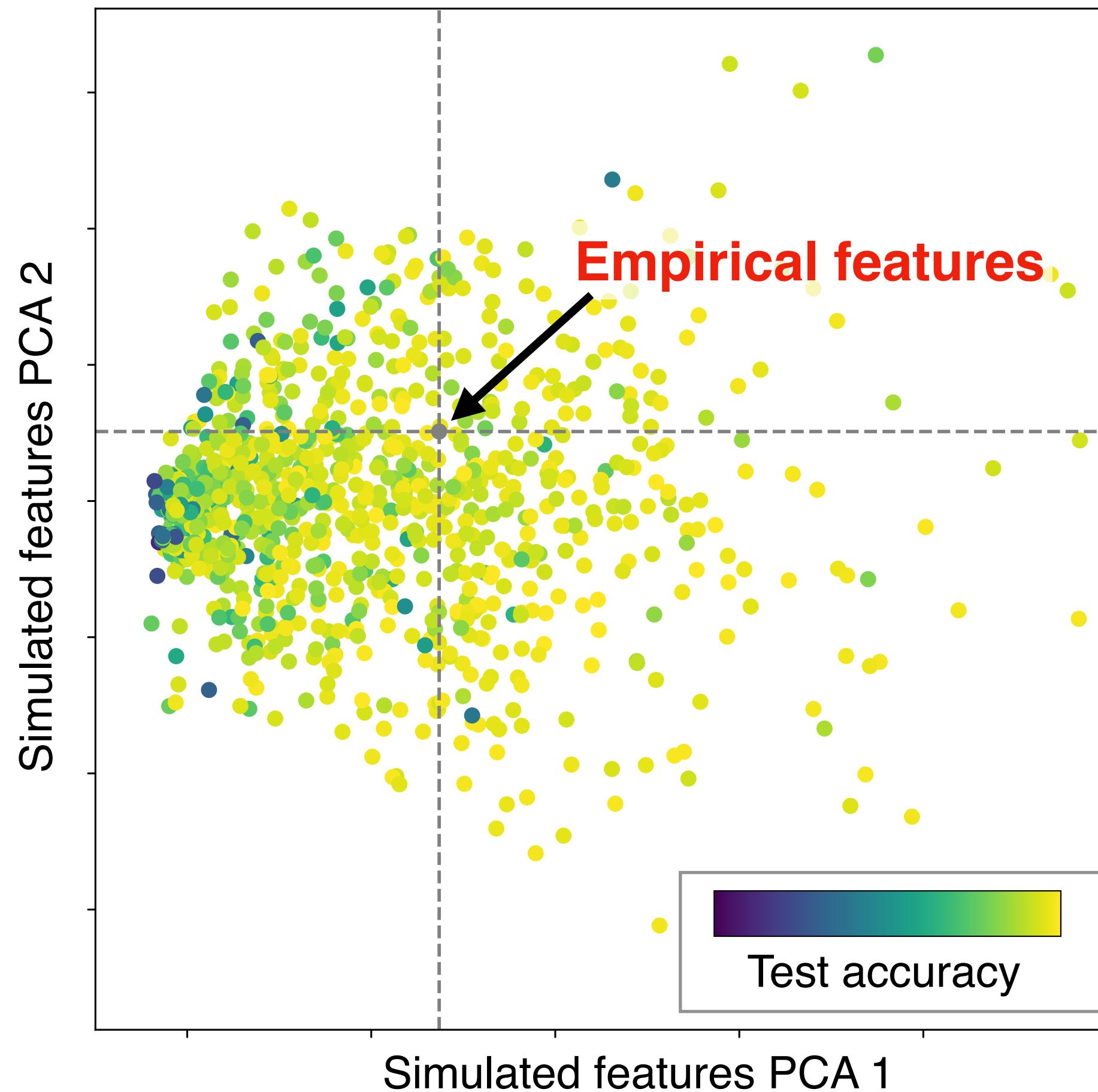
Checking that the simulations are in the right ballpark



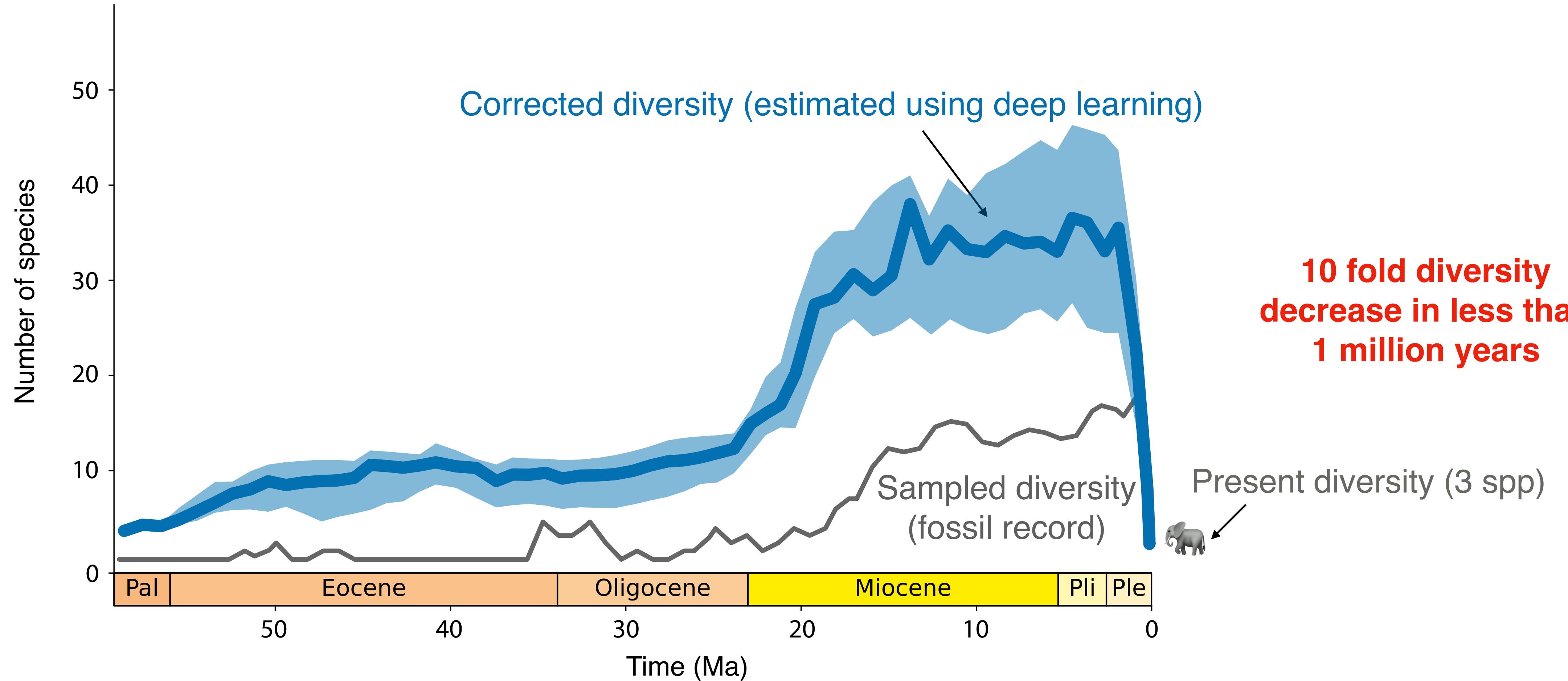
Features of the simulated datasets



Checking that the simulations are in the right ballpark

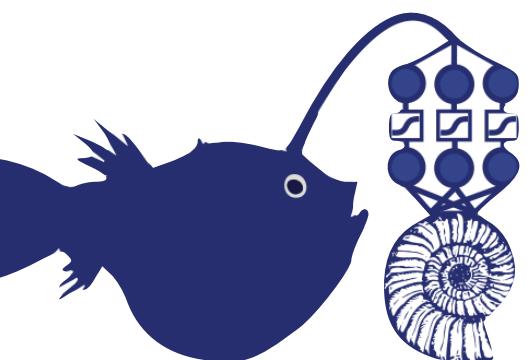


The rise and fall of elephants



R Cooper

github.com/DeepDive-project
Cooper et al. 2024 Nature Comms



How did palms diversify through time?

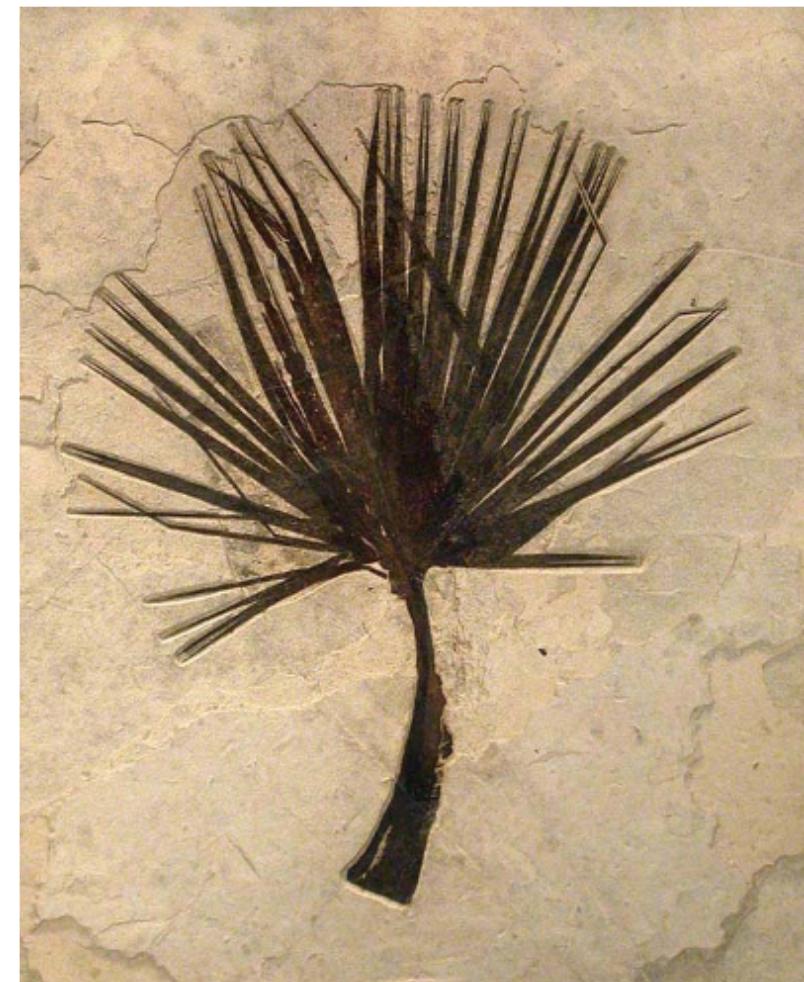


Image from Lim et al 2021 GEB

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

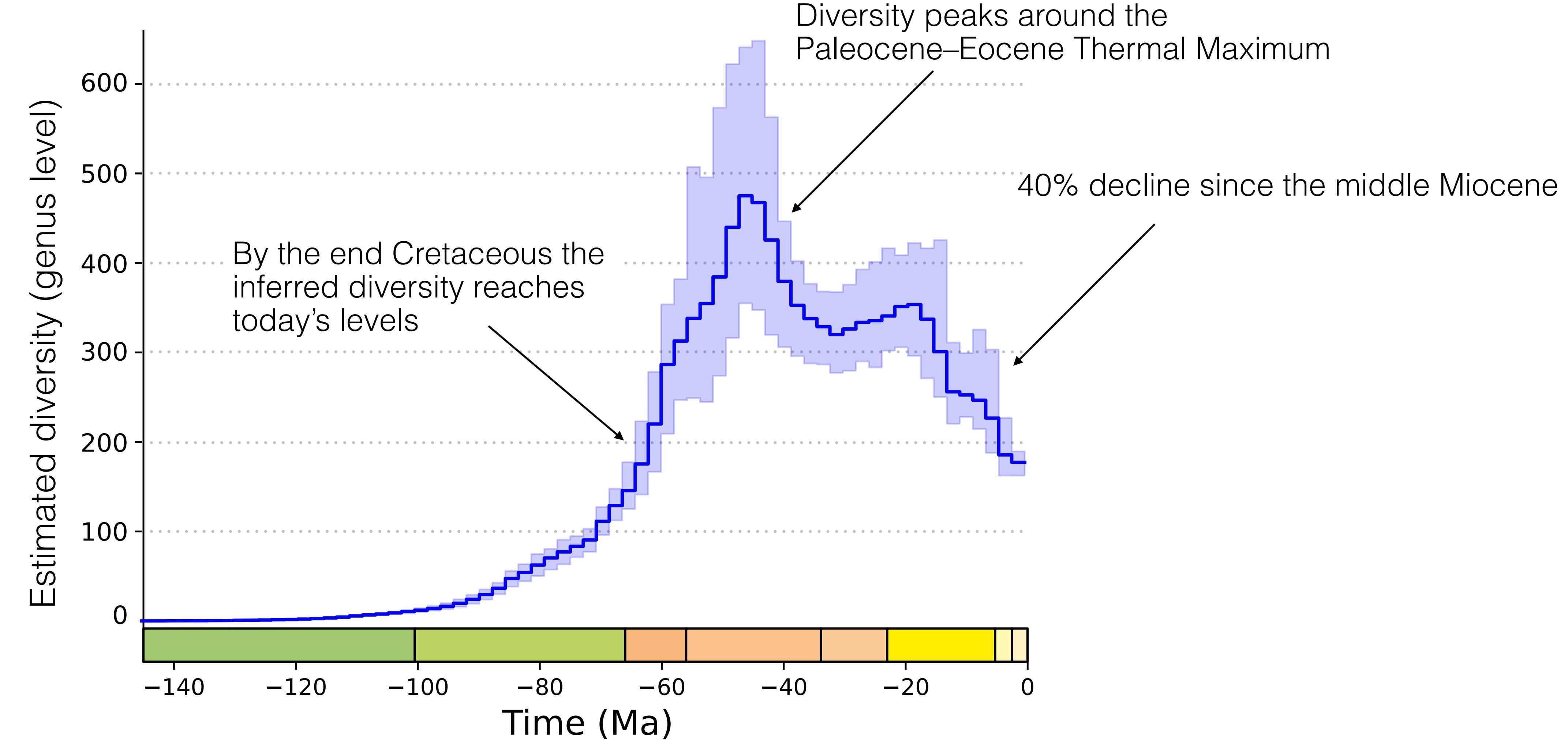
A team (ongoing) effort

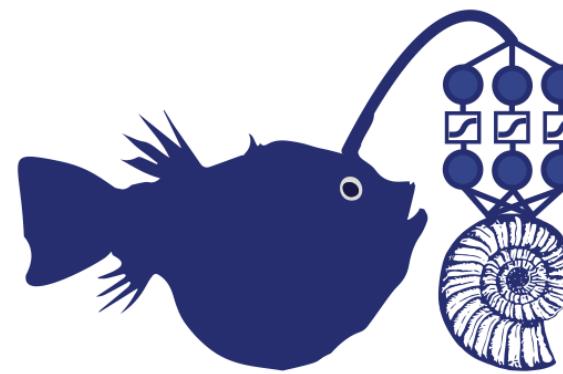
Rosane Collevatti
Carina Hoorn
Huasheng Huang
Viktoria Keller

Luis Palazzi
Shalani Parmar
Vandana Prasad
David Sunderlin

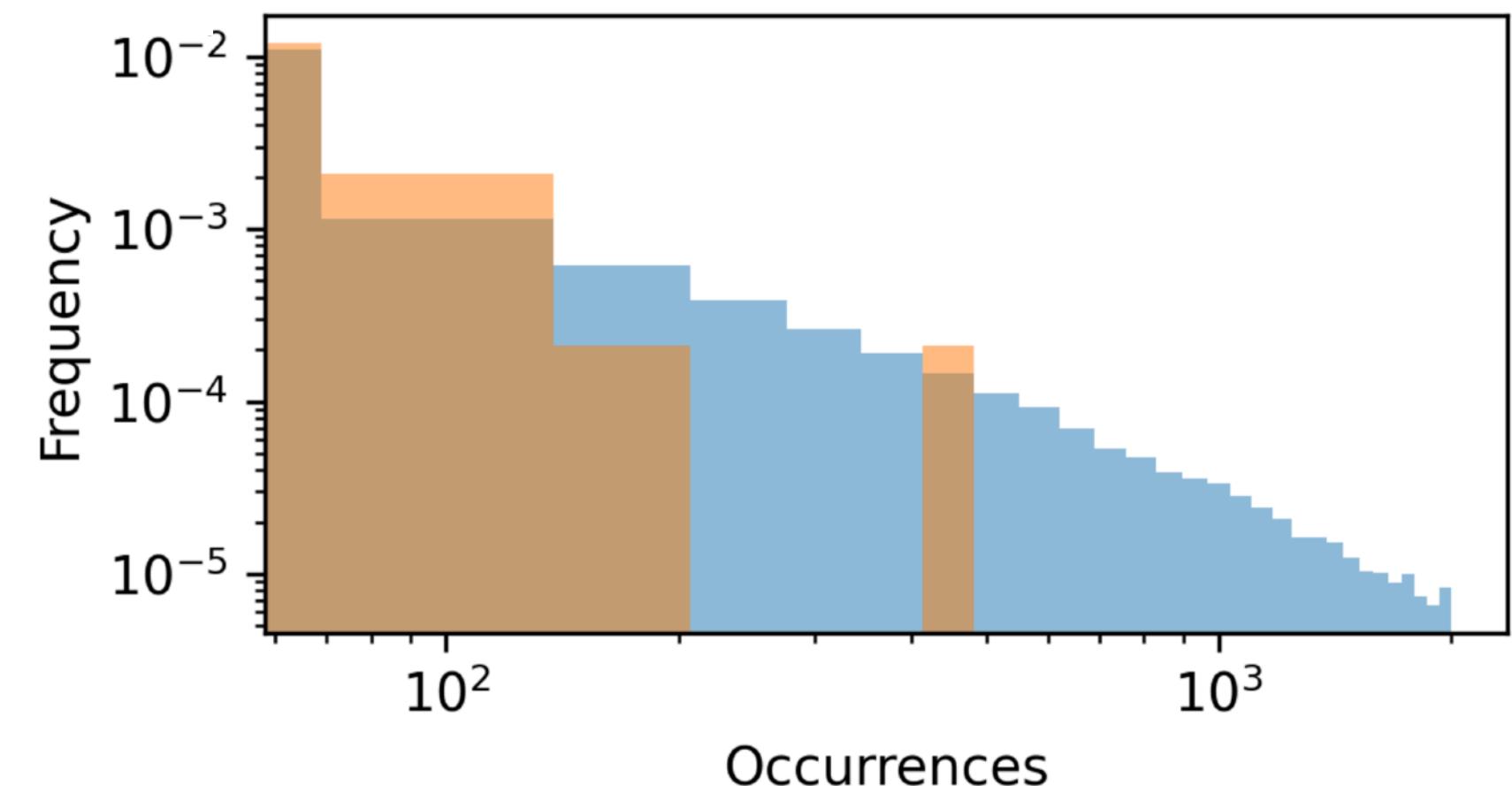
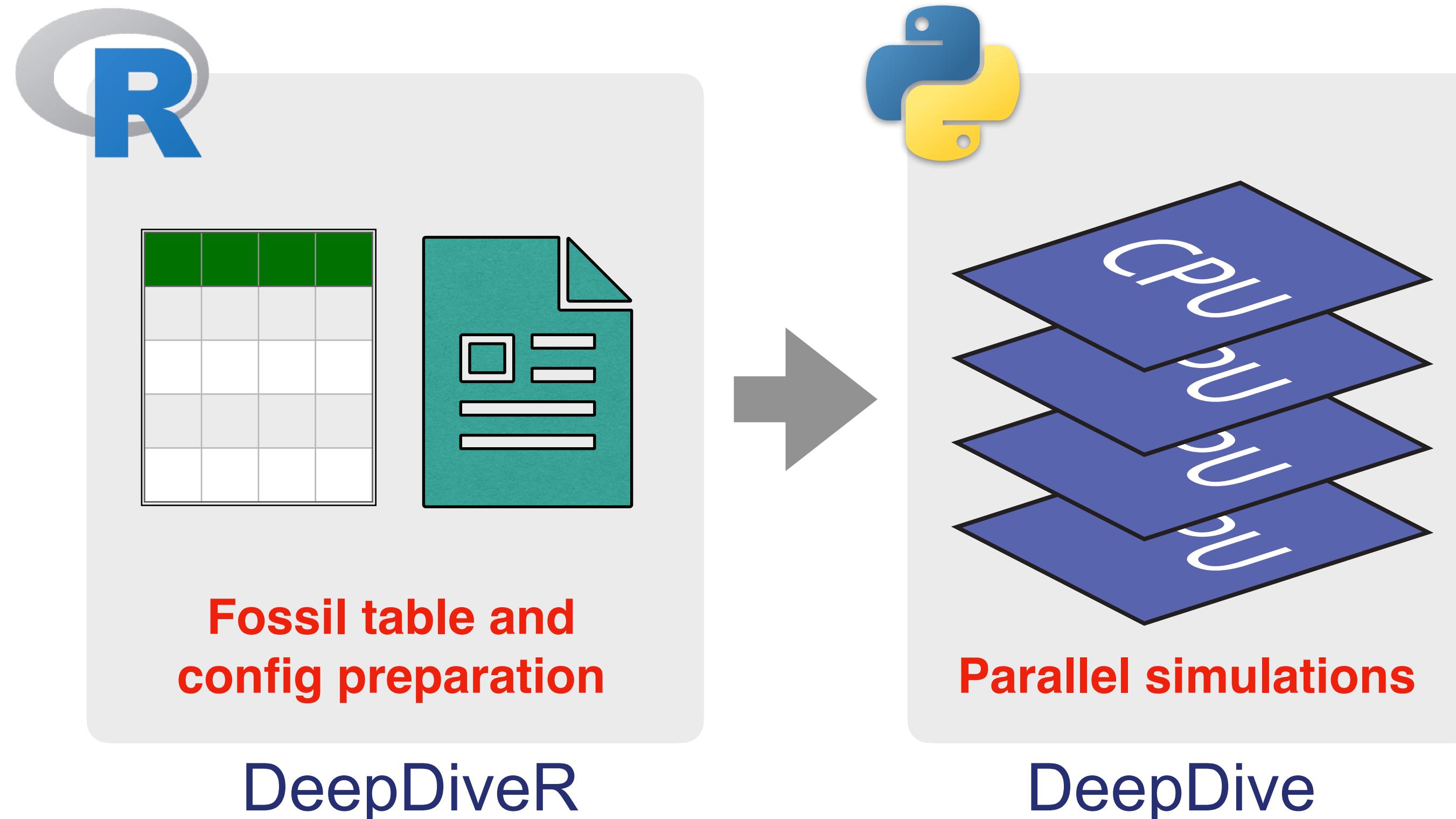
Kelly Matsunaga
Robert Morley
Yaowu Xing

The inferred palm evolutionary trajectory



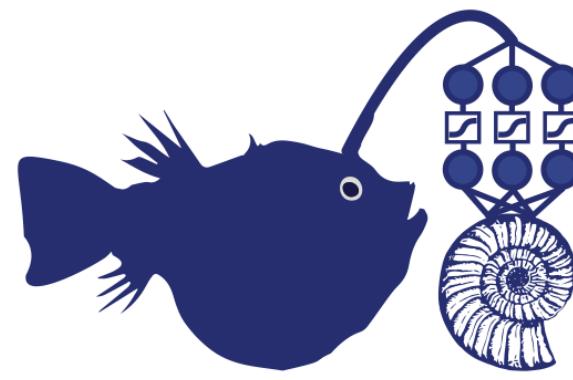


How DeepDive works

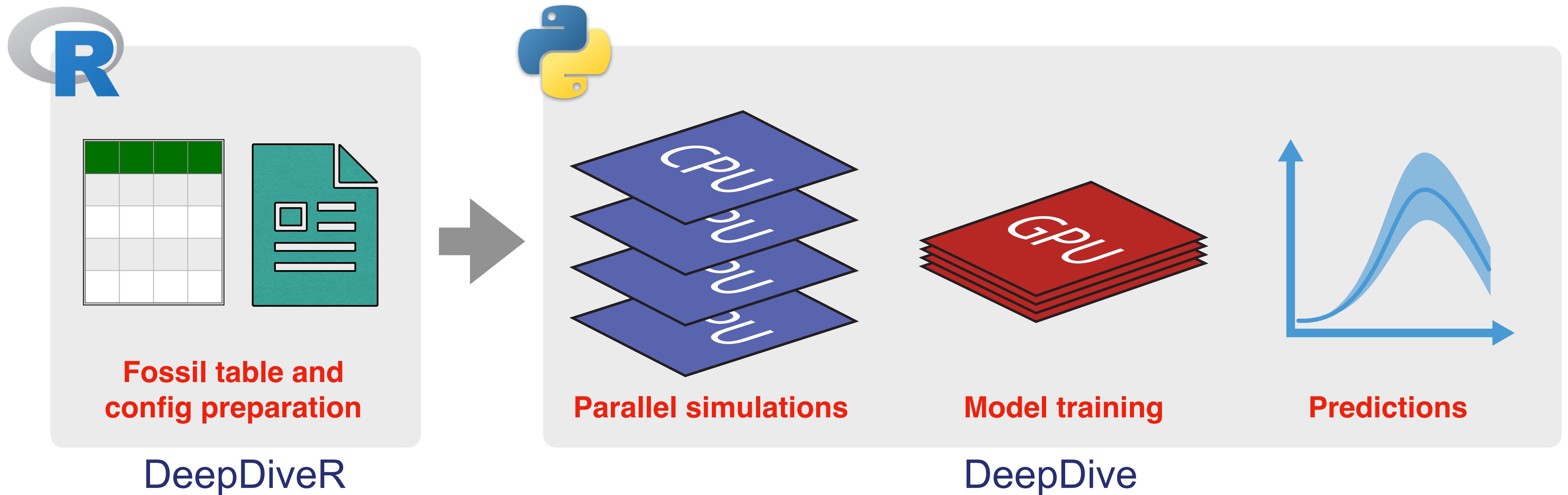


Simulations settings can
be adjusted manually but
are otherwise automatically
calibrated to match the
empirical data

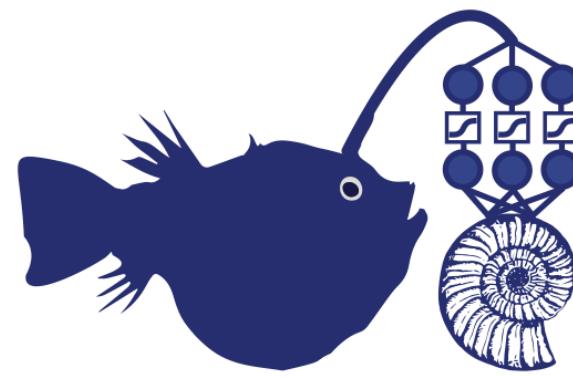
github.com/DeepDive-project



How DeepDive works



github.com/DeepDive-project



Setting up an analysis using DeepDiveR

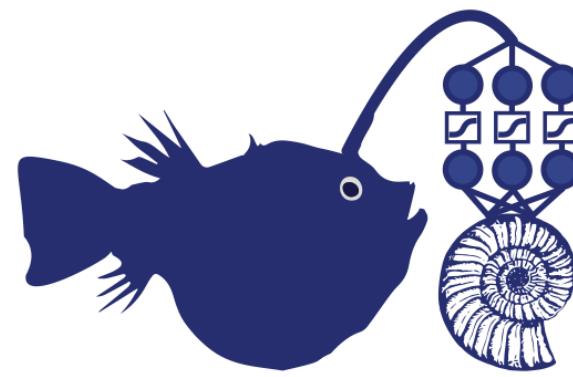
Taxon	Area	MinAge	MaxAge	Locality
<i>Ailurus</i>	Asia	0.6	1.3	loc_1
<i>Ailurus</i>	Asia	9.5	10.35	loc_2
<i>Alopecocyon</i>	Asia	9.5	10.35	loc_2
<i>Alopecocyon</i>	Europe	9.7	11.11	loc_3
...

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") ← Define timelines and
4
5 # define time bins (here of 1 myr) ← number of age
6 bins <- seq(23, 0, by=-1) ← randomizations
7
8 # prep DeepDive-formatted fossil dataset
9 prep_dd_input(fossil_tbl, bins=bins, output_file="fossil_input.csv", r=100)
```



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



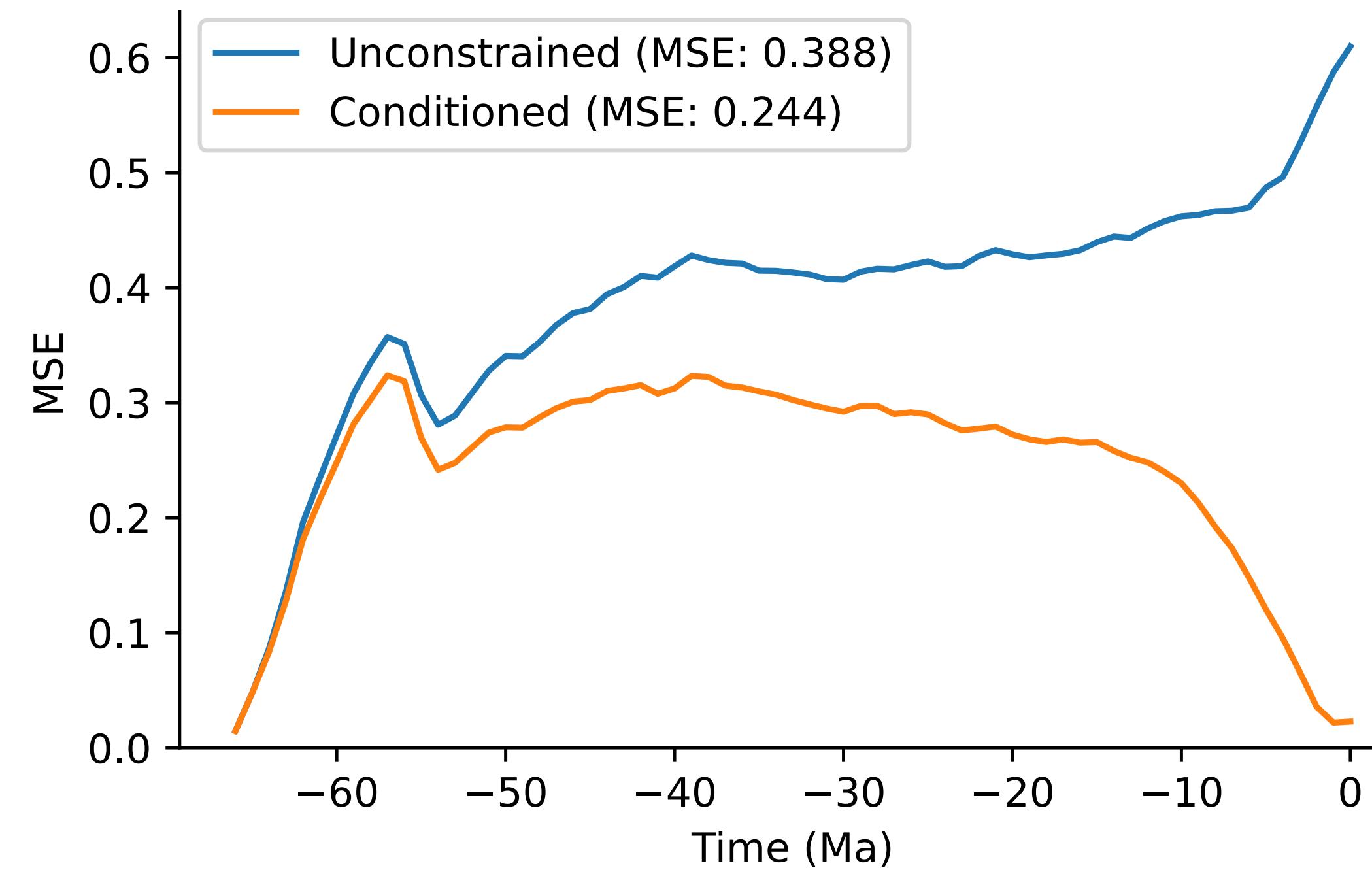
Parse fossil table and prepare config file

```
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
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")
```

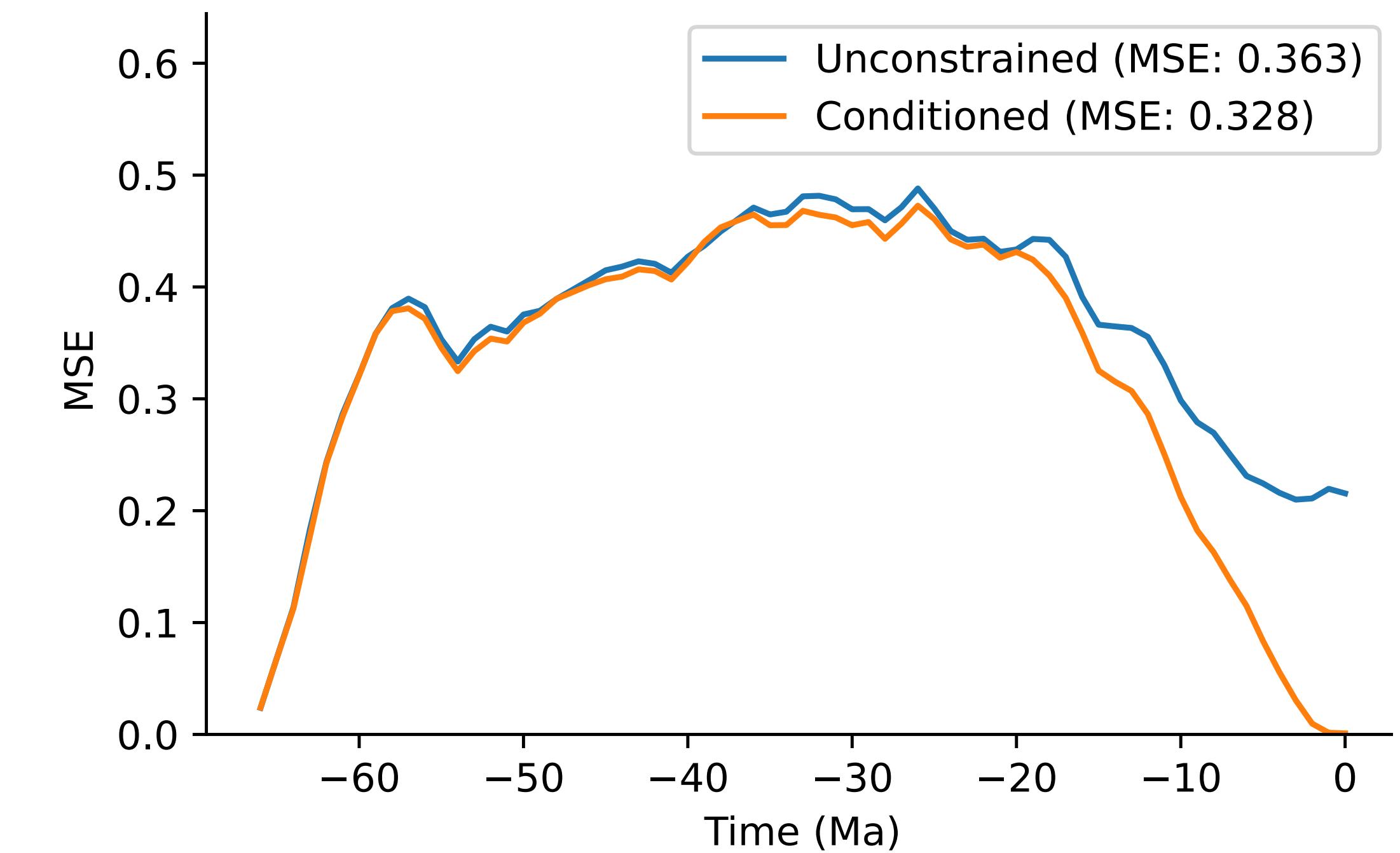
Include present diversity (if applicable)

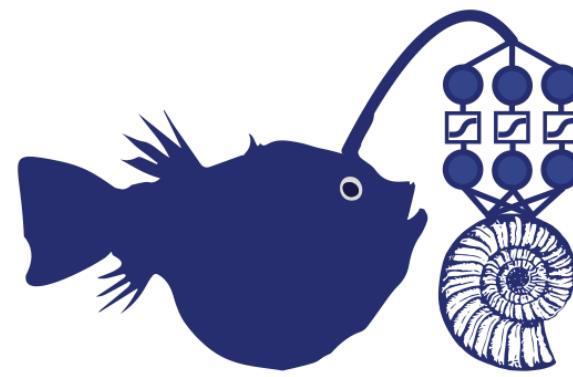
Conditioning on present diversity

Extant clades



Extinct clades



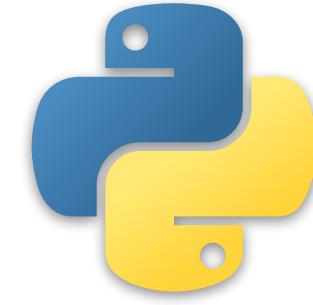
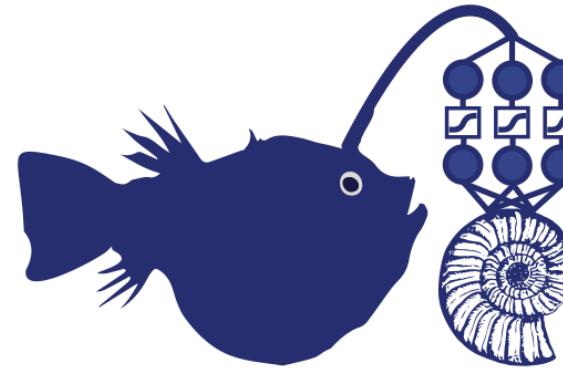


Adding biogeographic information (timing and availability of areas)

Setting up an analysis using DeepDiveR

```
1 area_ages <- rbind(  
2   c(0, 0), # Africa  
3   c(30, 40), # Antarctica  
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 Antarctica inhabitable
after ice sheet forms



Run in an interactive
Python console

Running a DeepDive analysis from config

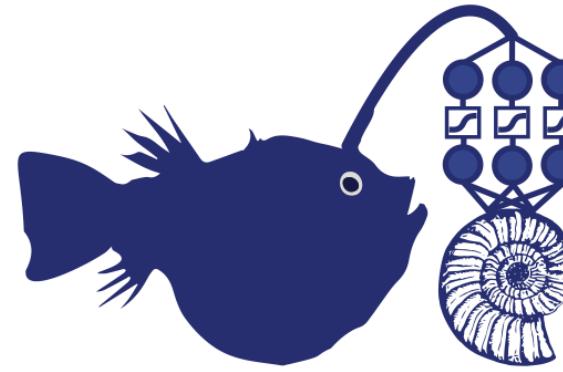
```
1 # import the DeepDive python library
2 from deepdive import config_runner
3
4 wd = "my_working_dir"
5
6 # launch the analysis (simulations, training, predictions)
7 config_runner.run_config("config_file.ini", wd=wd, CPU=64)
8
```



Parallel simulations

Model training

Predictions



Running a DeepDive analysis from config



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

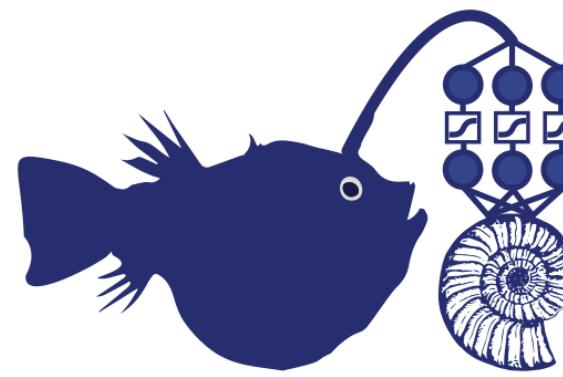
```
mbp:~ $ python run_dd_config.py my_path/config_file.ini -cpu 10 -wd my_path
```



Parallel simulations

Model training

Predictions



Check out our tutorials on Github

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

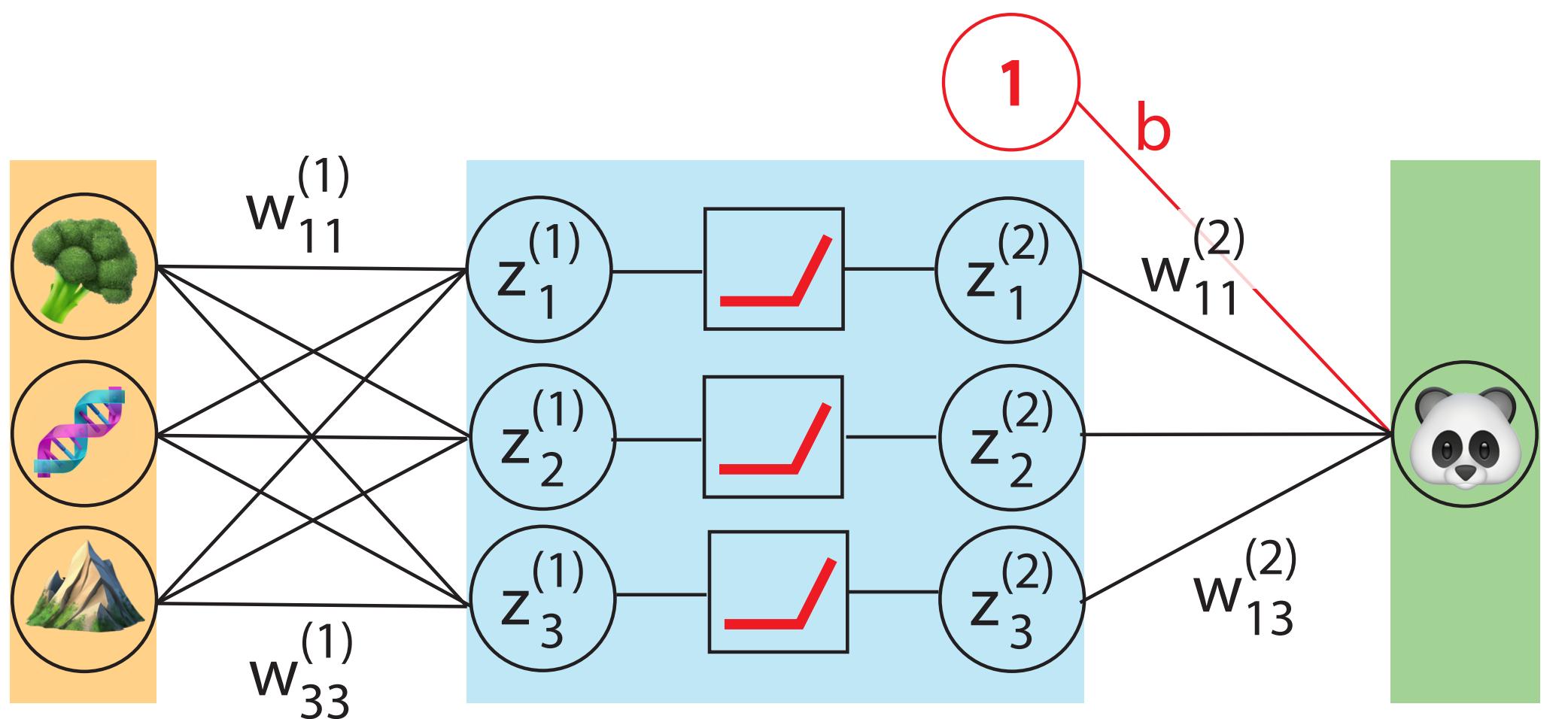


Parallel simulations

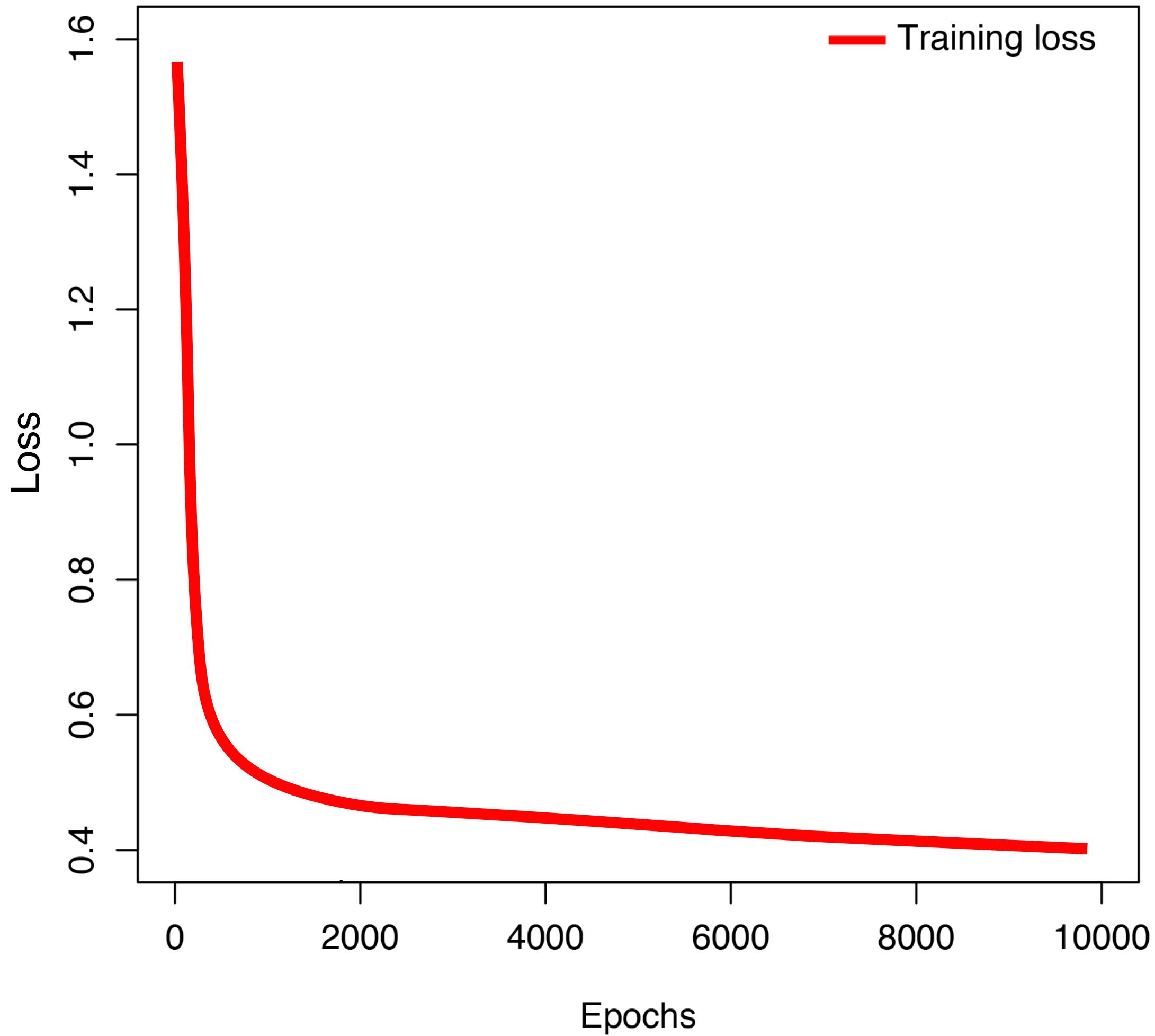
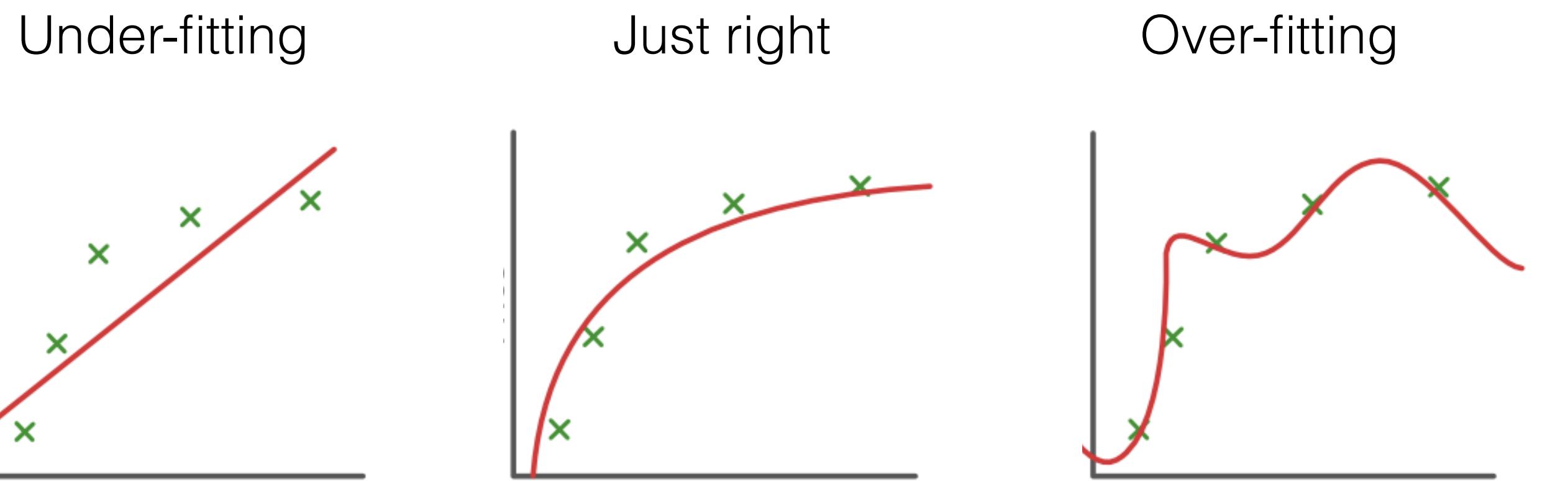
Model training

Predictions

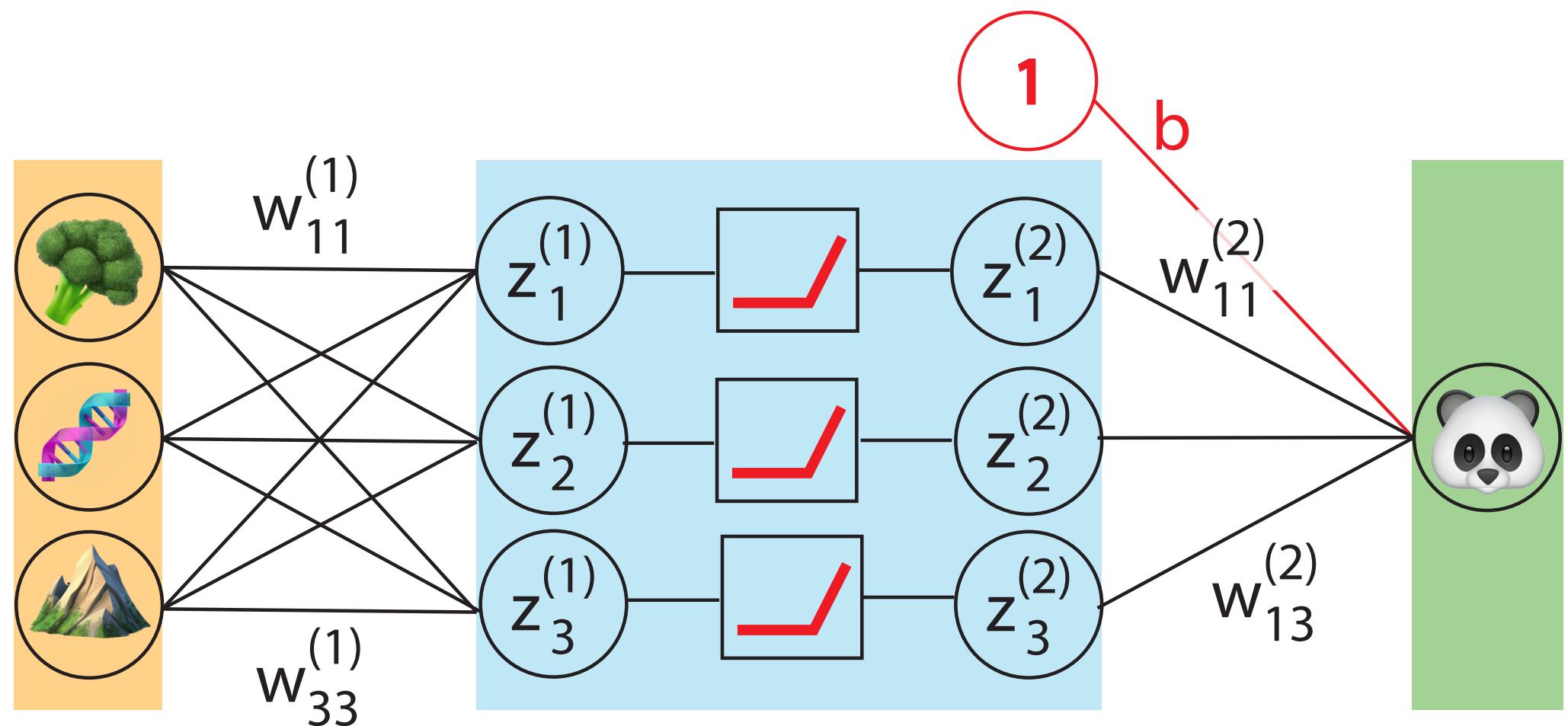
Optimization of a NN model



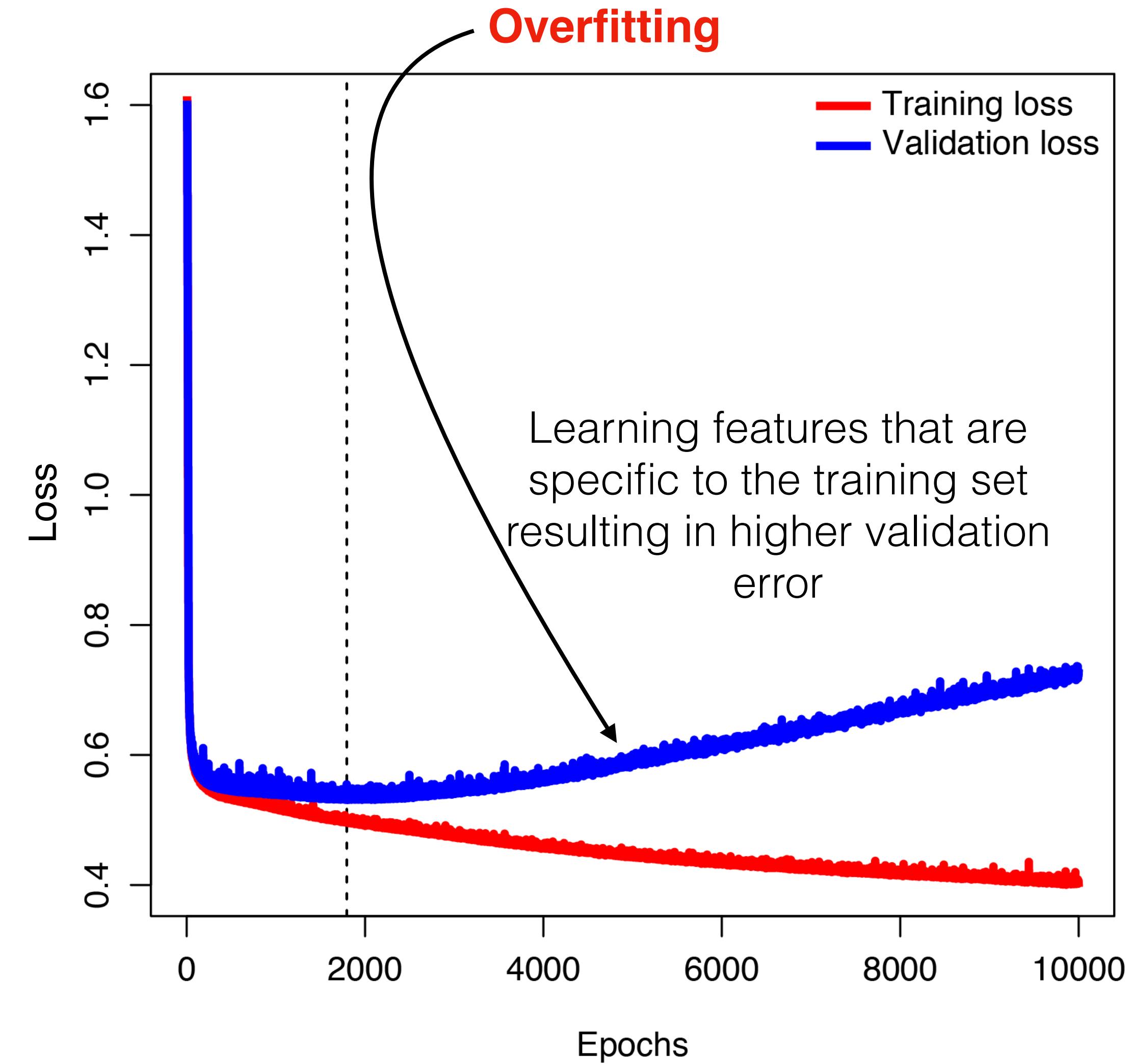
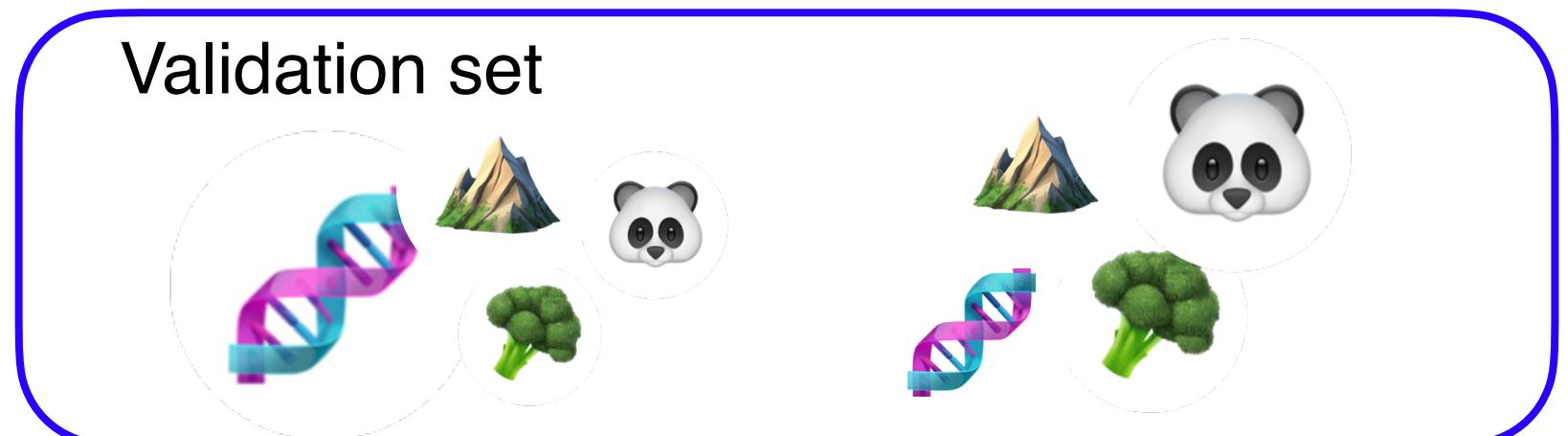
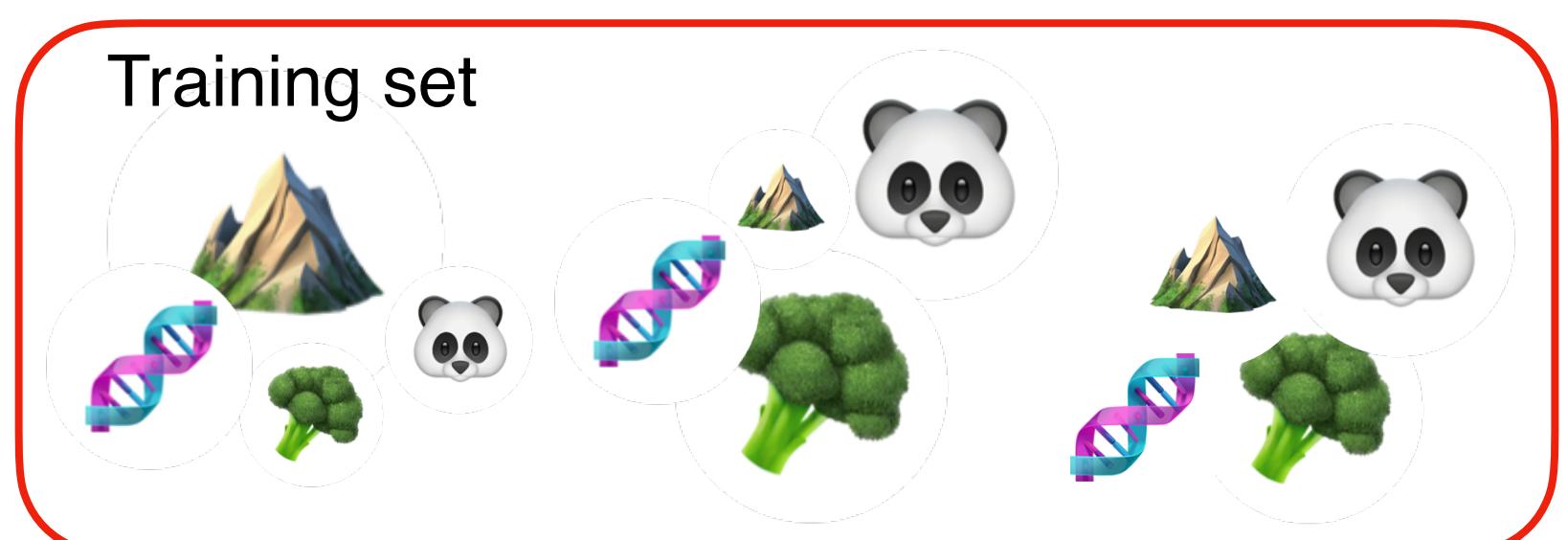
A maximum likelihood optimization of an NN would inevitably lead to over-fitting because NNs are over-parameterized



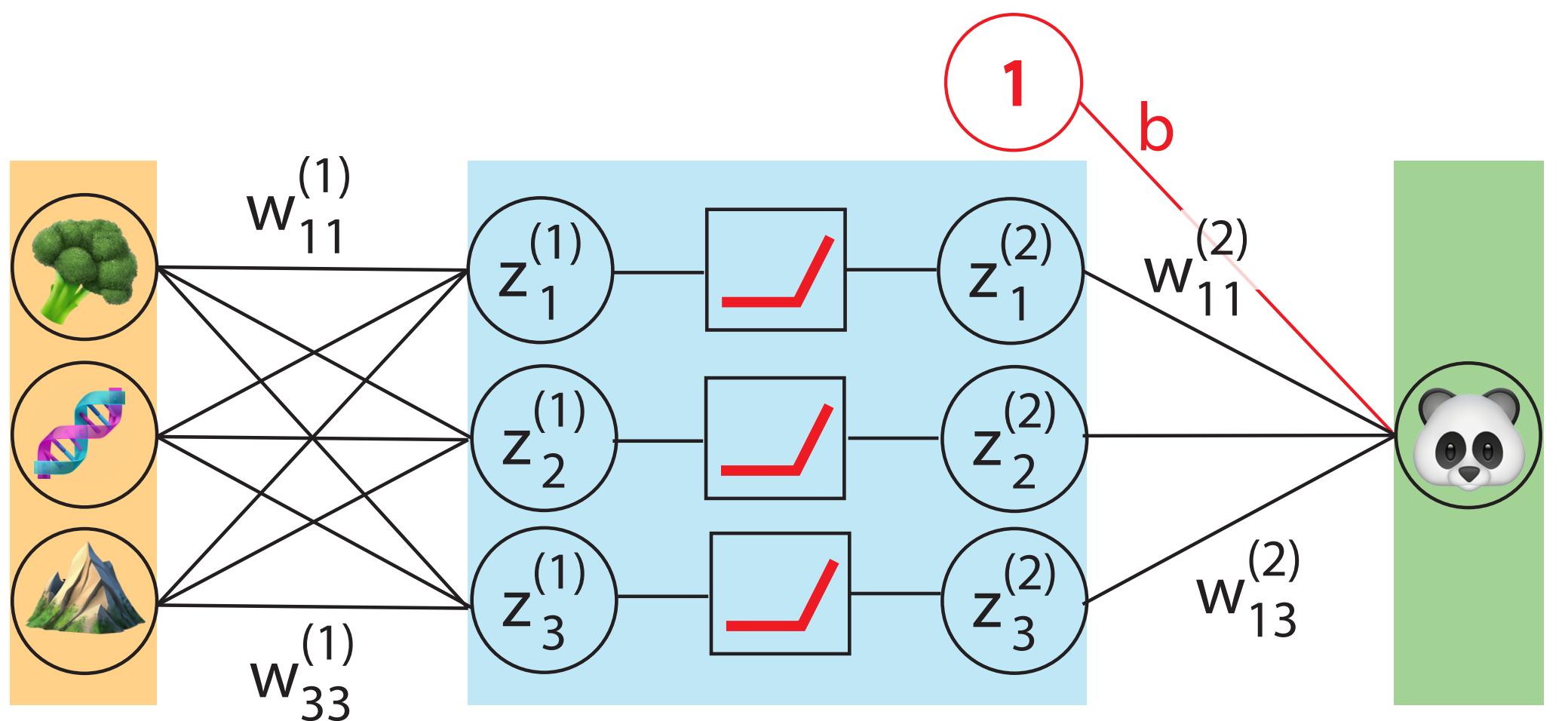
Optimization of a NN model



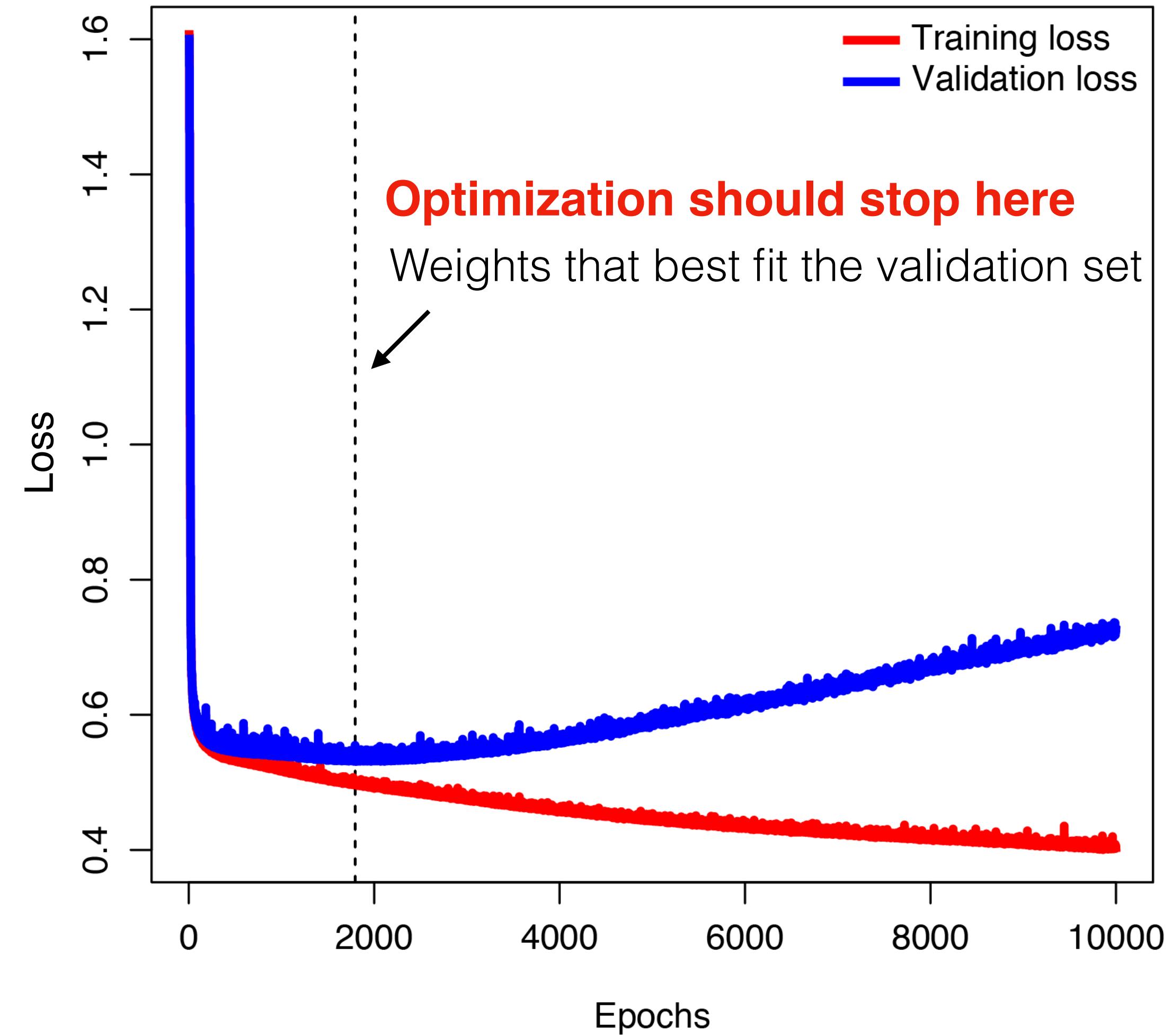
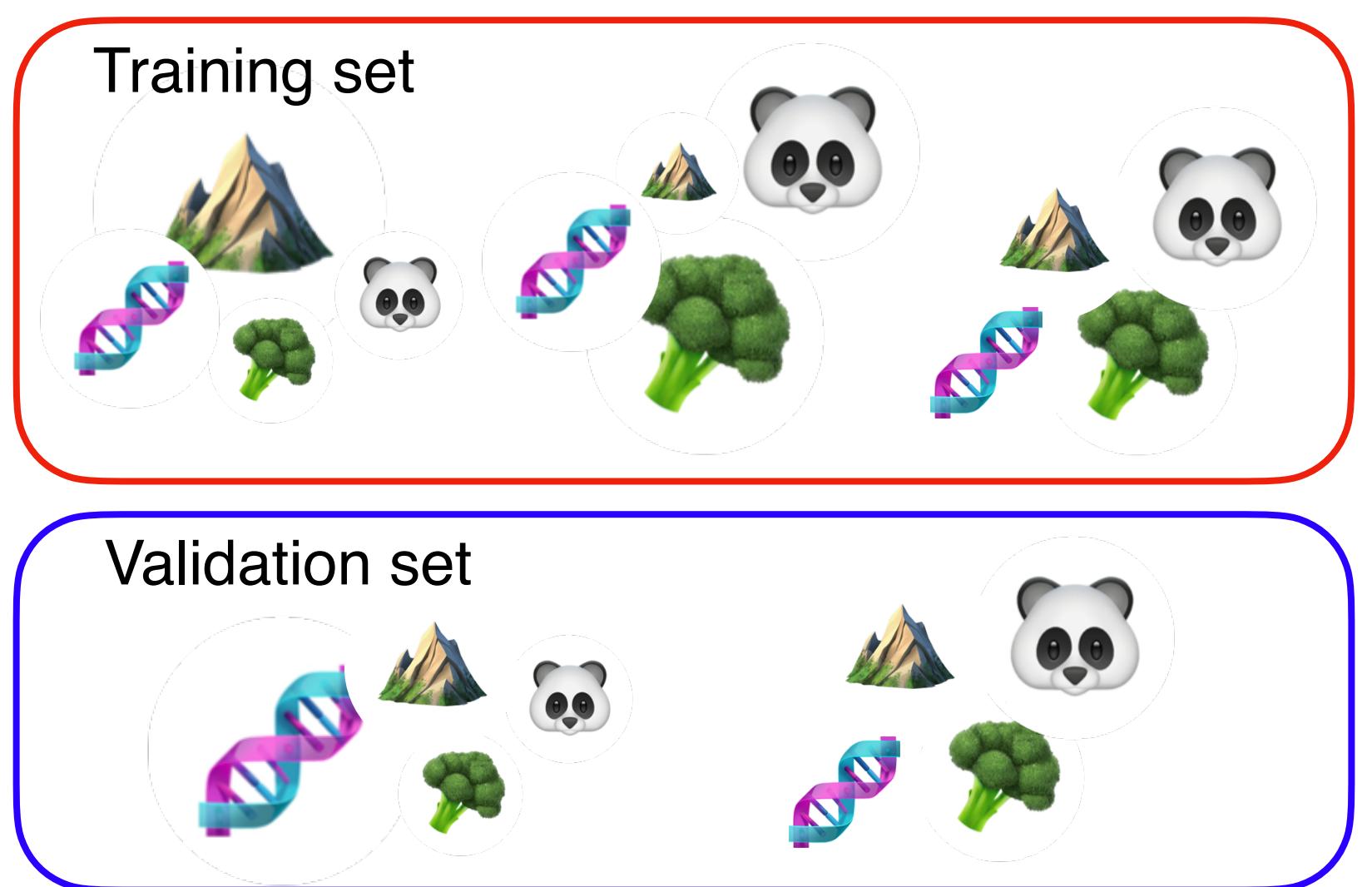
In machine learning a separate validation set is used to stop the optimization process before it starts overfitting

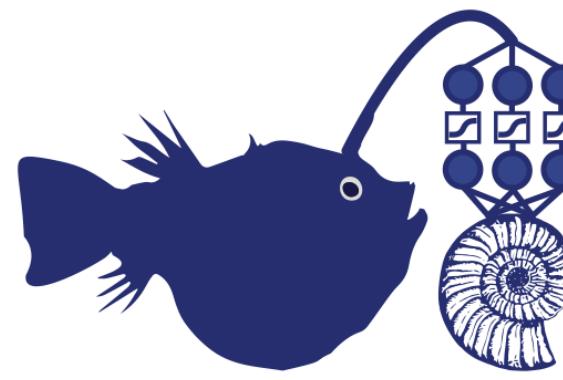


Optimization of a NN model



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)

