

Format Cetaceans Data

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Creation - jeremy.andreoletti@ens.fr - 13/04/2020

Load occurrence dataset

```
## occurrence_no      record_type      reid_no      flags      collection_no
## Min.      : 68135    occ:4476    Min.      :11942    Mode:logical    Min.      : 4868
## 1st Qu.: 540497                1st Qu.:18135    NA's:4476      1st Qu.: 48884
## Median : 725033                Median :21552                Median : 67030
## Mean   : 804995                Mean   :22595                Mean   : 82570
## 3rd Qu.:1021108              3rd Qu.:27547              3rd Qu.: 99584
## Max.   :1396622              Max.   :34473              Max.   :192401
##                                     NA's   :4191
##                                     identified_name      identified_rank identified_no
## Cetacea indet.      : 271    species      :2438    Min.      : 36652
## Mysticeti indet.    : 140    genus        : 675    1st Qu.: 42951
## Odontoceti indet.   : 124    family       : 648    Median : 63219
## Delphinidae indet.  : 96     suborder     : 297    Mean   : 67516
## Balaenopteridae indet.: 92    unranked clade: 287    3rd Qu.: 65078
## Balaena mysticetus  : 79     superfamily  : 71     Max.   :367667
## (Other)             :3674    (Other)      : 60
##                                     difference      accepted_name      accepted_rank
##                                     :3557    Cetacea      : 294    species      :2057
## recombined as       : 302    Odontoceti   : 240    family       : 761
## nomen dubium        : 301    Mysticeti    : 208    genus        : 741
## subjective synonym of: 118    Balaenopteridae : 156    suborder     : 471
## invalid subgroup of : 56     Delphinidae   : 107    unranked clade: 310
## corrected to        : 30     Balaena mysticetus: 84    superfamily  : 74
## (Other)             : 112    (Other)      :3387    (Other)      : 62
## accepted_no         early_interval      late_interval
```

##	Min.	: 36652	Holocene	: 626		:3967
##	1st Qu.:	42937	Langhian	: 355	Late Pliocene	: 80
##	Median	: 62924	Burdigalian	: 310	Langhian	: 60
##	Mean	: 64432	Zanclean	: 309	Messinian	: 59
##	3rd Qu.:	64626	Tortonian	: 267	Serravallian	: 58
##	Max.	:367667	Serravallian:	239	Early Pleistocene:	34
##			(Other)	:2370	(Other)	: 218
##	max_ma		min_ma		reference_no	
##	Min.	: 0.0117	Min.	: 0.000	Min.	: 289
##	1st Qu.:	3.6000	1st Qu.:	2.588	1st Qu.:	12813
##	Median	:11.6200	Median	: 5.333	Median	:23666
##	Mean	:13.9433	Mean	:10.117	Mean	:27322
##	3rd Qu.:	20.4400	3rd Qu.:	13.820	3rd Qu.:	38452
##	Max.	:66.0000	Max.	:47.800	Max.	:65107
##						

##	occurrence_no	record_type	reid_no	flags	collection_no
##	1	68135	occ	NA	4868
##	2	137494	occ	NA	11601
##	3	141404	occ	NA	12121
##	4	147937	occ	NA	13063
##	5	147938	occ	NA	13064
##	6	148079	occ	NA	13078
##	7	148335	occ	NA	13090
##	8	148353	occ	NA	13092
##	9	148356	occ	NA	13096
##	10	148358	occ	NA	13098
##	11	148360	occ	NA	13100
##	12	148363	occ	NA	13102
##	13	148364	occ	NA	13102
##	14	148365	occ	19615	13103
##	15	150826	occ	NA	11596
##	16	150827	occ	NA	13103
##	17	150828	occ	NA	13402
##	18	150829	occ	NA	13402
##	19	150830	occ	NA	13403
##	20	150831	occ	NA	13403

##		identified_name	identified_rank	identified_no
##	1	n. gen. Georgiacetus n. sp. vogtlensis	species	63123
##	2	Argyroctetus joaquinensis	species	69897
##	3	n. gen. Kharthlidelphis n. sp. diceros	species	53161
##	4	n. gen. Pinocetus n. sp. polonicus	species	53140
##	5	n. gen. Basiloterus n. sp. hussaini	species	53165
##	6	n. gen. Sachalinocetus n. sp. cholmicus	species	63225
##	7	n. gen. Praekogia n. sp. cedrosensis	species	53139
##	8	Aulophyseter n. sp. rionegrensensis	species	53106
##	9	Microcetus n. sp. sharkovi	species	53137
##	10	n. gen. Mixocetus n. sp. elysius	species	64432
##	11	n. gen. Austrosqualodon n. sp. trirrhizodonta	species	63212
##	12	Basilosaurus isis	species	53287
##	13	Dorudon atrox	species	53288
##	14	Dorudon atrox	species	53288
##	15	Basilosaurus n. sp. drazindai	species	53163
##	16	Basilosaurus isis	species	53287

## 17		Basilosaurus isis	species	53287	
## 18		Dorudon atrox	species	53288	
## 19		Basilosaurus isis	species	53287	
## 20		Dorudon atrox	species	53288	
##	difference	accepted_name	accepted_rank	accepted_no	
## 1		Georgiacetus vogtlensis	species	63123	
## 2		Argyrosetus joaquinensis	species	69897	
## 3	nomen dubium	Kharthlidelphis	genus	53160	
## 4		Pinocetus polonicus	species	53140	
## 5		Basiloterus hussaini	species	53165	
## 6		Sachalinocetus cholmicus	species	63225	
## 7		Praekogia cedrosensis	species	53139	
## 8	invalid subgroup of	Physeteroidea	superfamily	53105	
## 9		Microcetus sharkovi	species	53137	
## 10		Mixocetus elysius	species	64432	
## 11		Austrosqualodon trirhizodonta	species	63212	
## 12		Basilosaurus isis	species	62984	
## 13		Dorudon atrox	species	62994	
## 14		Dorudon atrox	species	62994	
## 15		Basilosaurus drazindai	species	53163	
## 16		Basilosaurus isis	species	62984	
## 17		Basilosaurus isis	species	62984	
## 18		Dorudon atrox	species	62994	
## 19		Basilosaurus isis	species	62984	
## 20		Dorudon atrox	species	62994	
##	early_interval	late_interval	max_ma	min_ma	reference_no
## 1	Lutetian		47.800	41.300	289
## 2	Aquitania		23.030	20.440	4175
## 3	Chattian		28.100	23.030	6018
## 4	Langhian		15.970	13.820	4344
## 5	Bartonian		41.300	38.000	6010
## 6	Early Miocene	Middle Miocene	23.030	11.608	4357
## 7	Messinian		7.246	5.333	4361
## 8	Messinian		7.246	5.333	4362
## 9	Chattian		28.100	23.030	4365
## 10	Tortonian		11.620	7.246	10152
## 11	Duntroonian		27.300	25.200	4368
## 12	Priabonian		38.000	33.900	6026
## 13	Priabonian		38.000	33.900	6026
## 14	Priabonian		38.000	33.900	10457
## 15	Bartonian		41.300	38.000	6010
## 16	Priabonian		38.000	33.900	6026
## 17	Priabonian		38.000	33.900	6026
## 18	Priabonian		38.000	33.900	6026
## 19	Priabonian		38.000	33.900	6026
## 20	Priabonian		38.000	33.900	6026

Reorder accepted ranks according to classification standard.

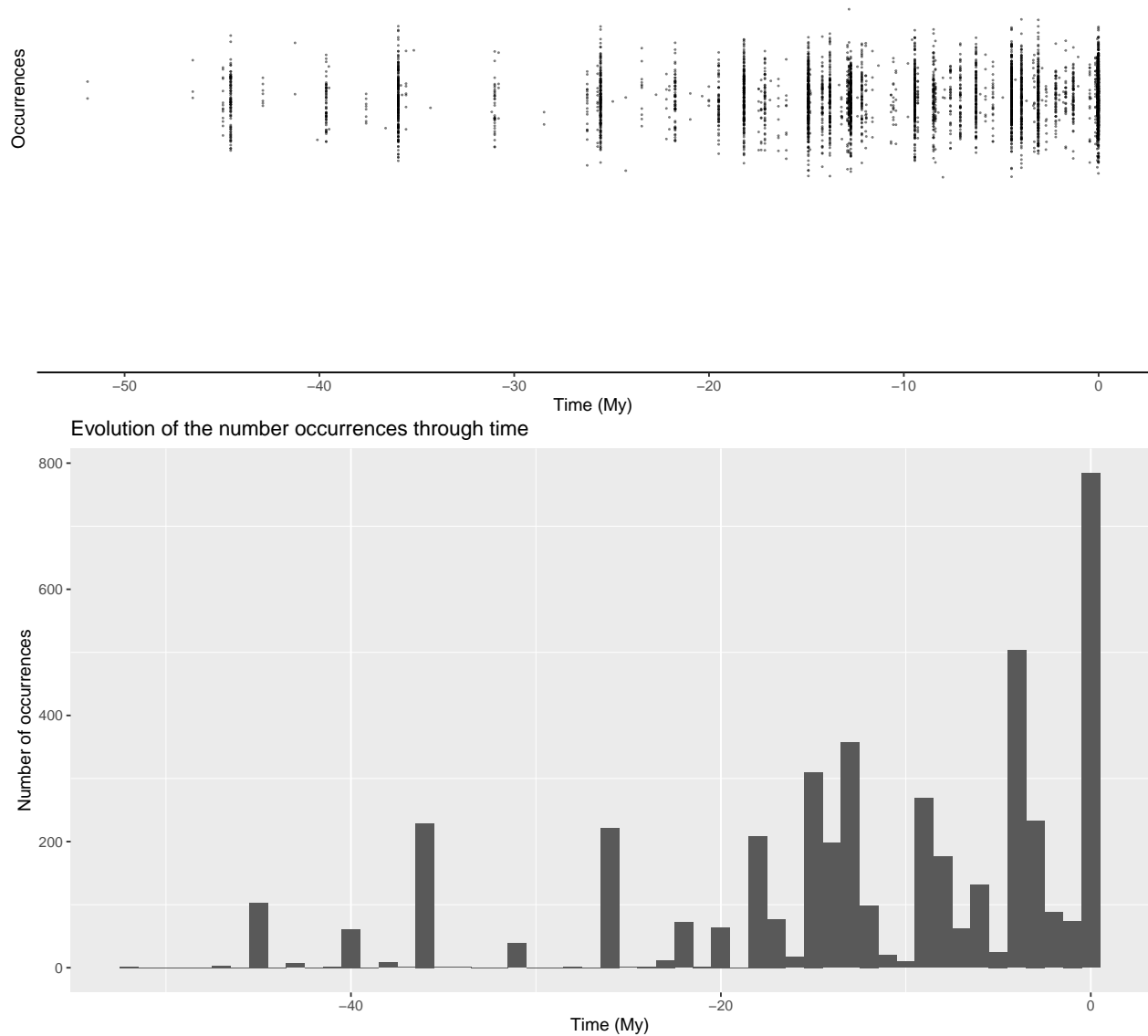
## [1]	"family"	"genus"	"infraorder"	"species"
## [5]	"subfamily"	"suborder"	"subspecies"	"superfamily"
## [9]	"unranked clade"			

Explore the dataset

Repartition through time

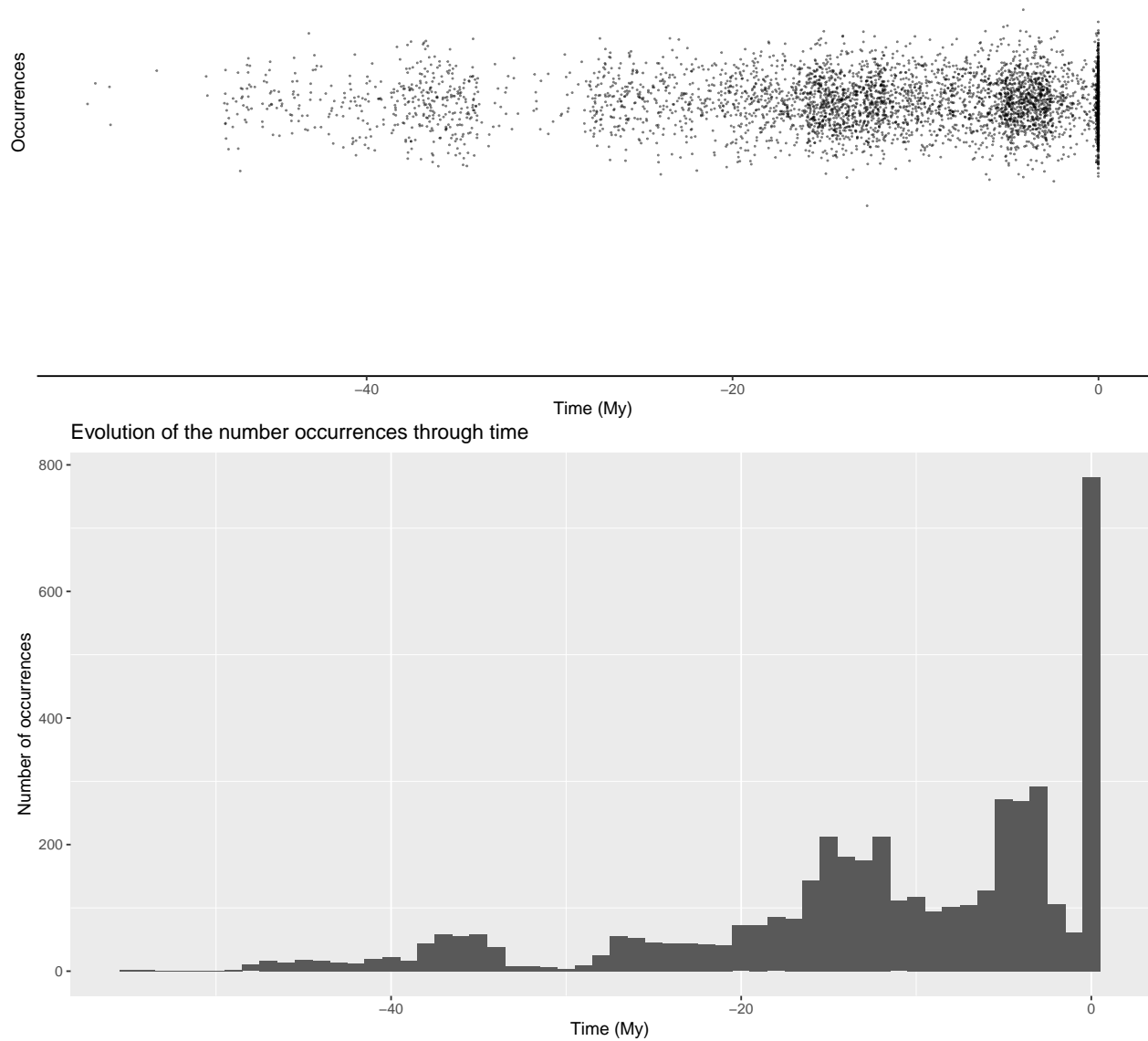
Full fossil record

Repartition of 4476 recorded occurrences through time



→ Numerous occurrences seem to have the same age interval so in order to avoid clusters let's draw them uniformly in their stratigraphic range rather than taking the mean.

Repartition of 4476 recorded occurrences through time

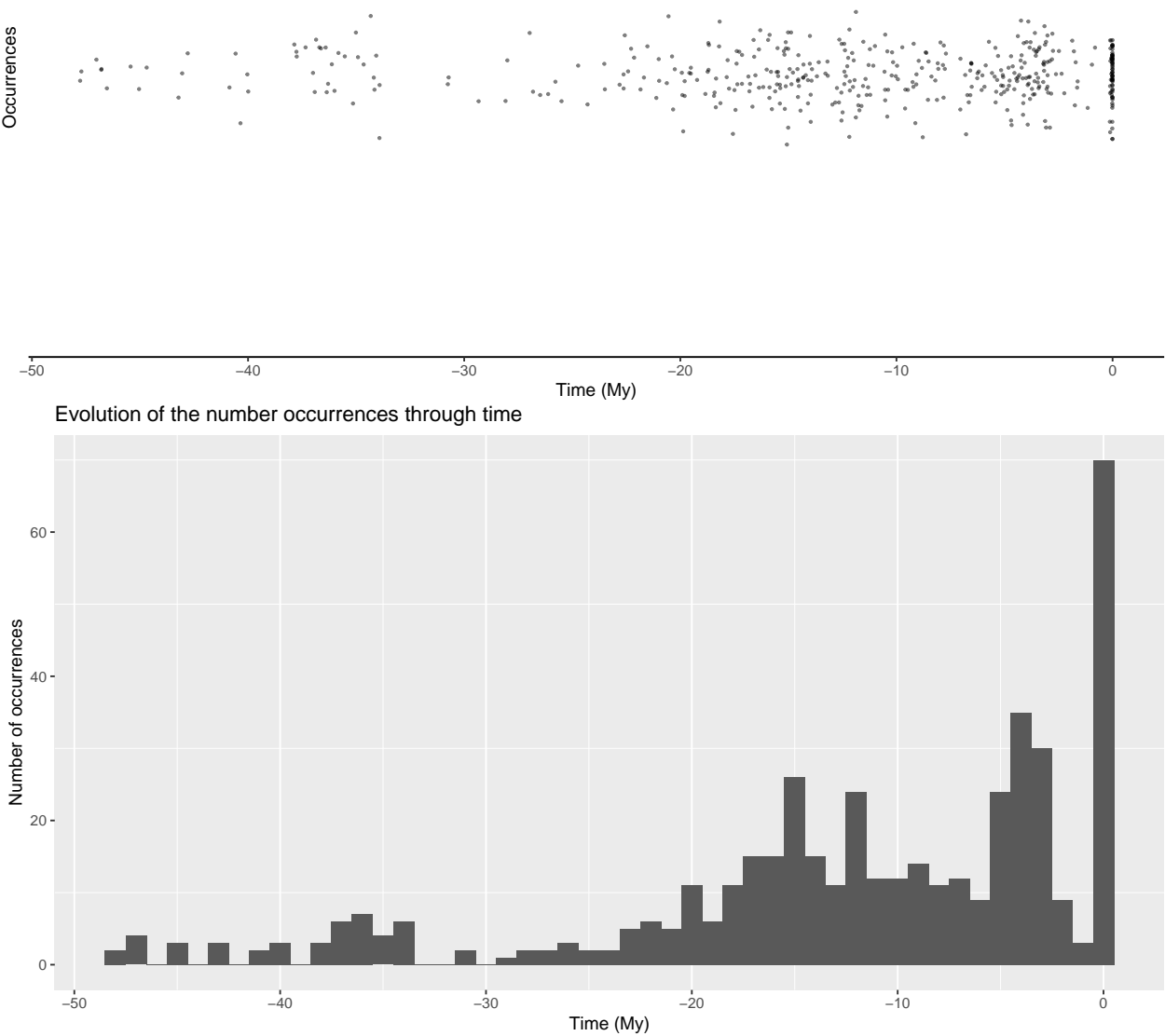


→ The repartition seems much smoother now.

Subsampling

These occurrences are too numerous for our current implementation, let's subsample a fraction of them for now.

Repartition of 448 recorded occurrences through time

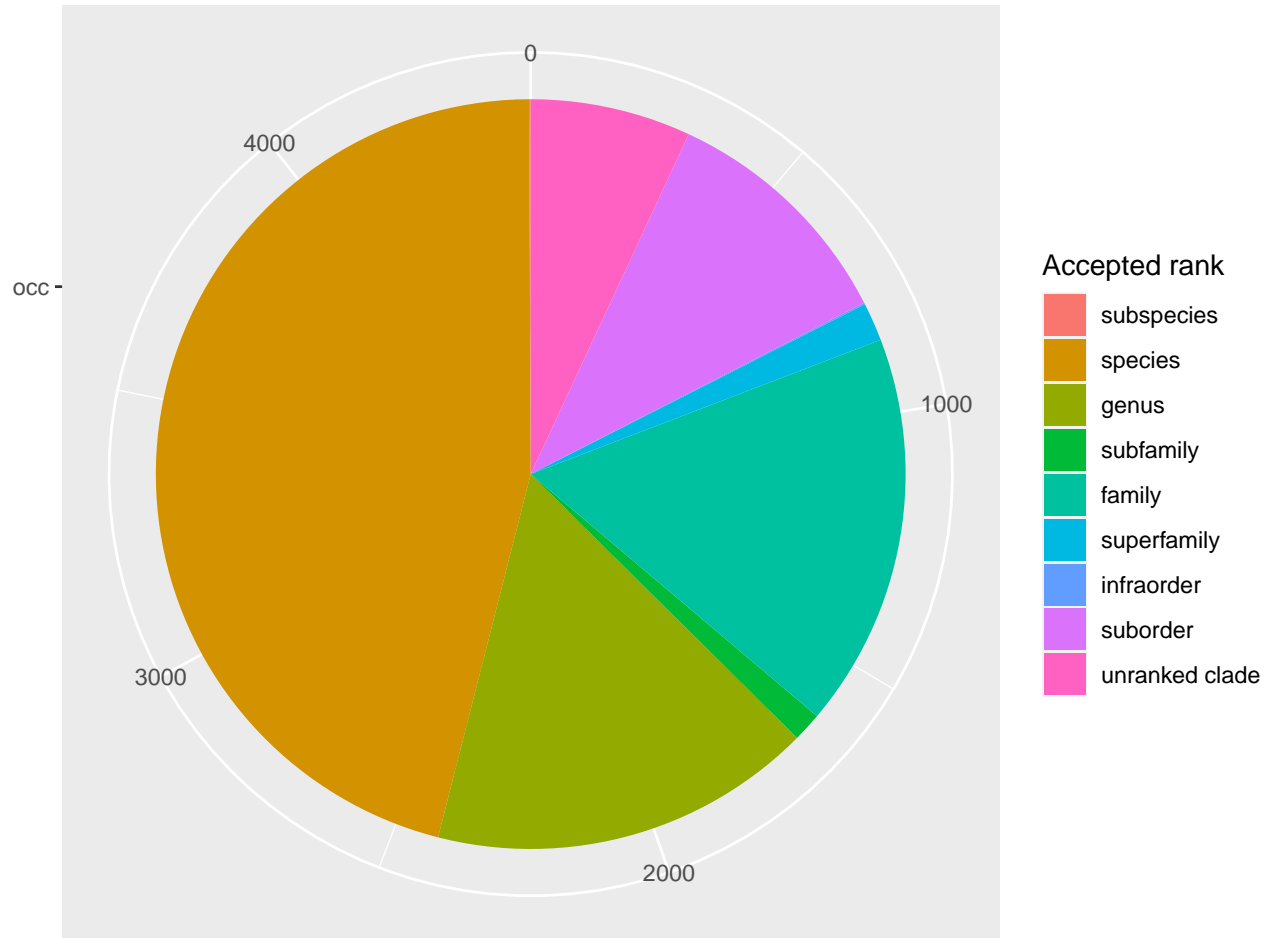


→ The distribution looks similar, with some noise due to higher variance with smaller sample.

Repartition among accepted ranks

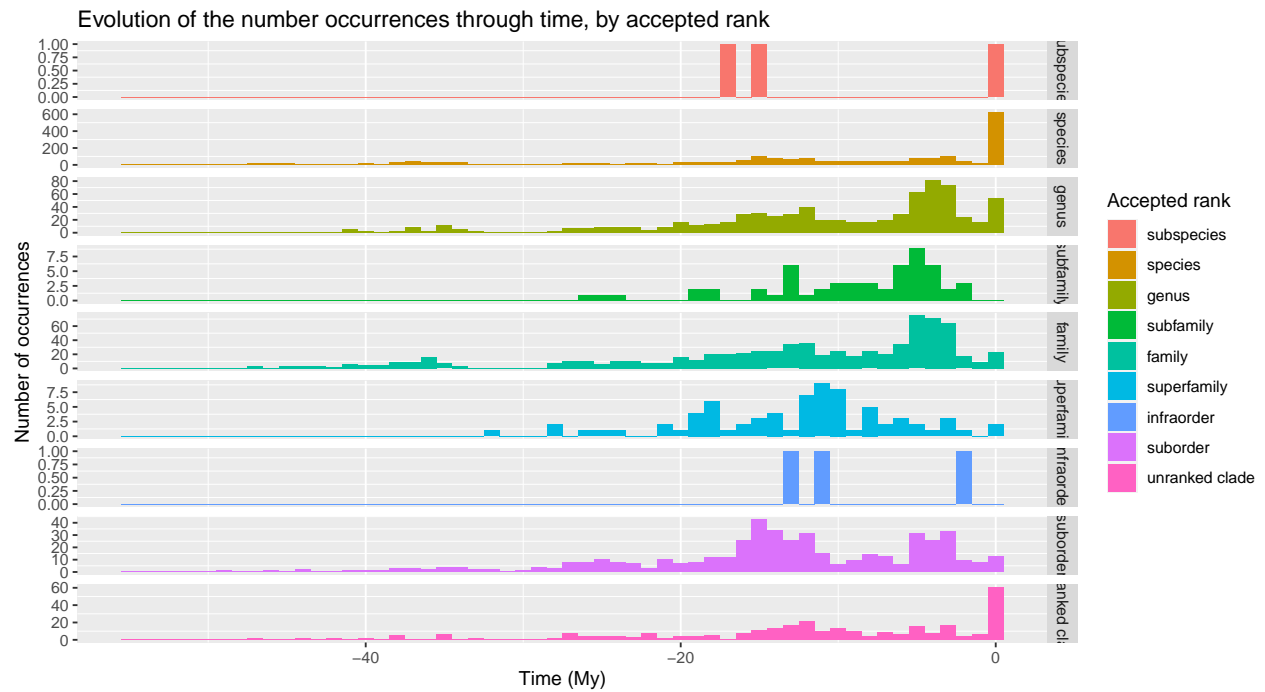
Pie chart

Repartition of occurrences among accepted ranks



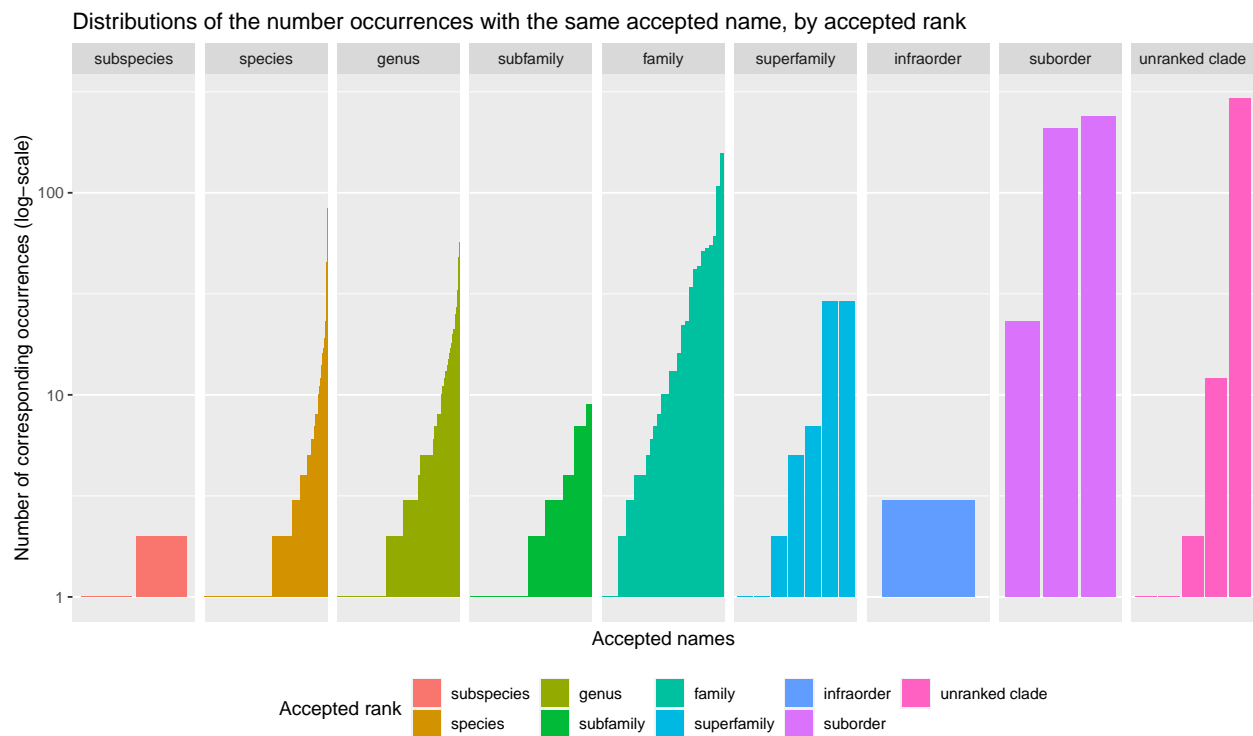
→ Half of the occurrences are identified at the level of the species and 1/3 at the genus or family. Very few occurrences for subspecies/infraorder, maybe fuse with species and suborder or remove ?

Time repartition by rank



→ Similar trends with peaks at ~15My and ~5My and a lot of them around 0 (artefact ?).

Redundancy of occurrences with the same accepted name



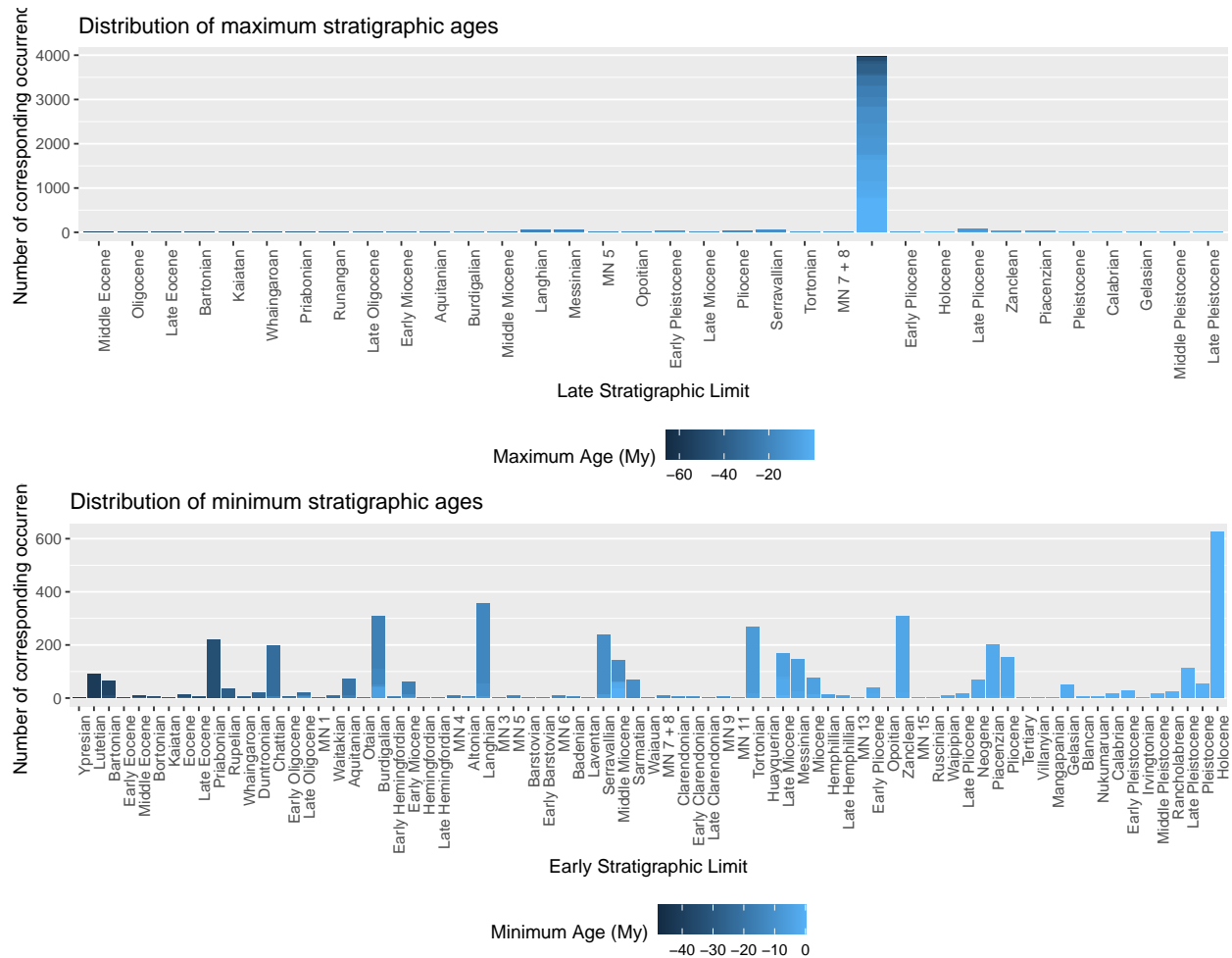
→ ~Half of species/genera/subfamilies have only one specimen by accepted name, but it could go up to ~50

within the same species and ~200 occurrences within the same suborder. **Those differences will have to be corrected because in our model all species are supposed to have the same abundance (identical sampling rates among branches).**

⇒ Our goal now will be to correct this abundance bias.

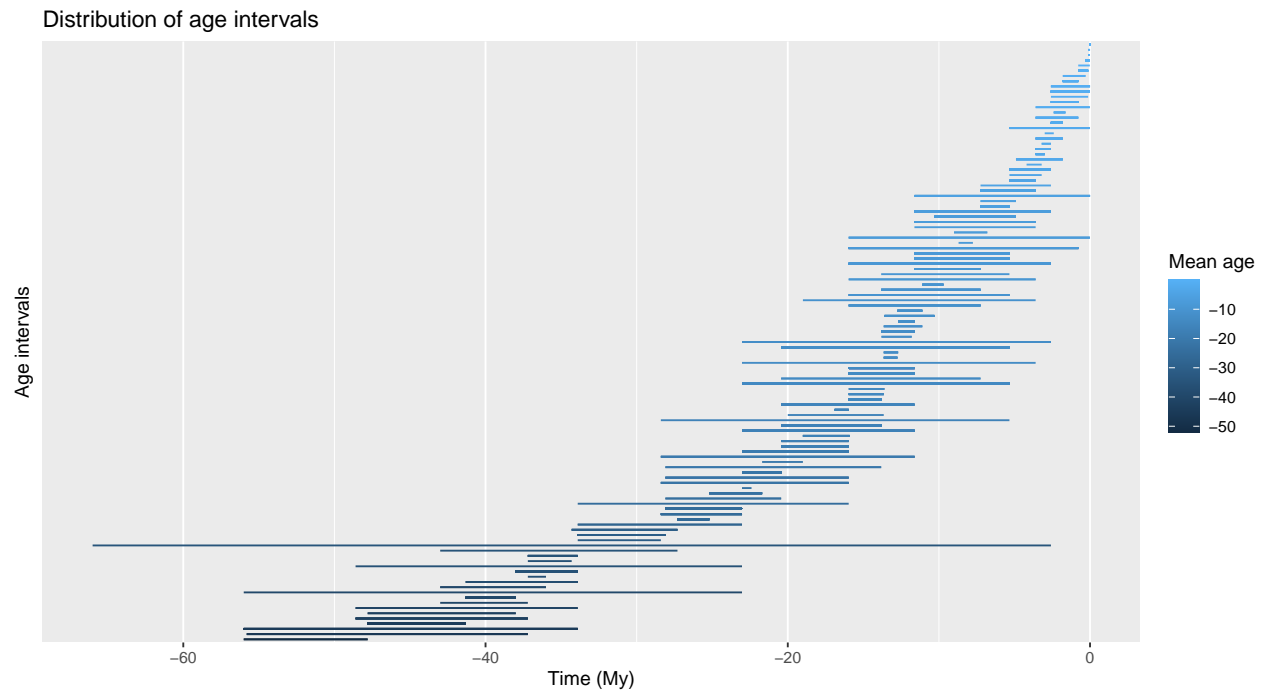
Time intervals = stratigraphic age uncertainty

Minimum and maximum stratigraphic limits



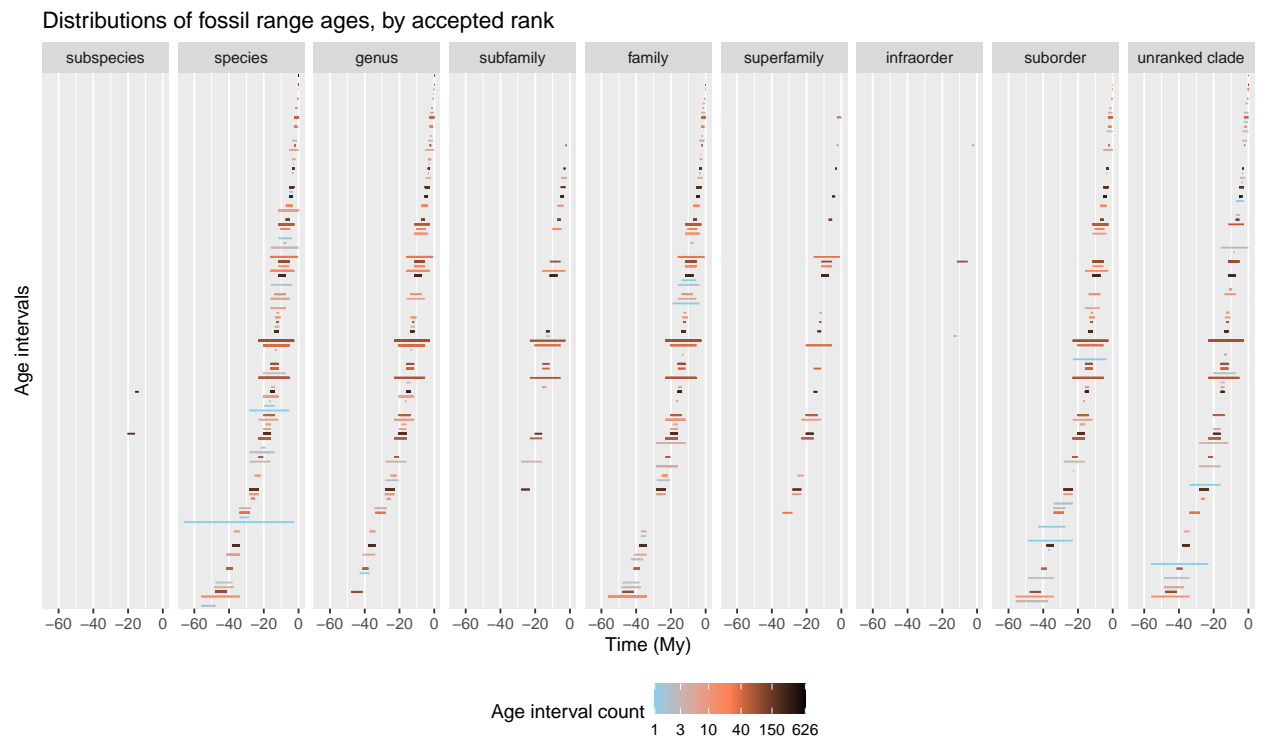
→ Most species have a early but not a late stratigraphic limit.

Minimum and maximum ages

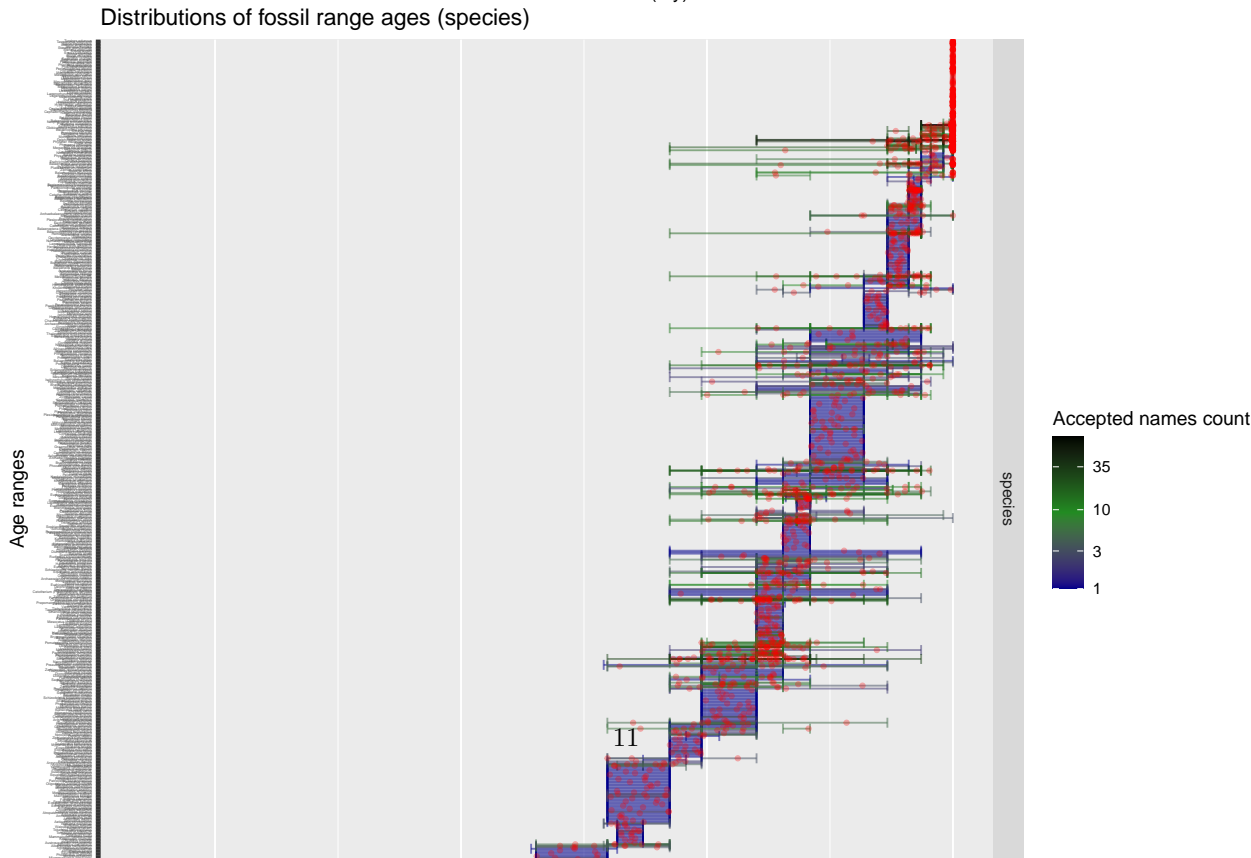
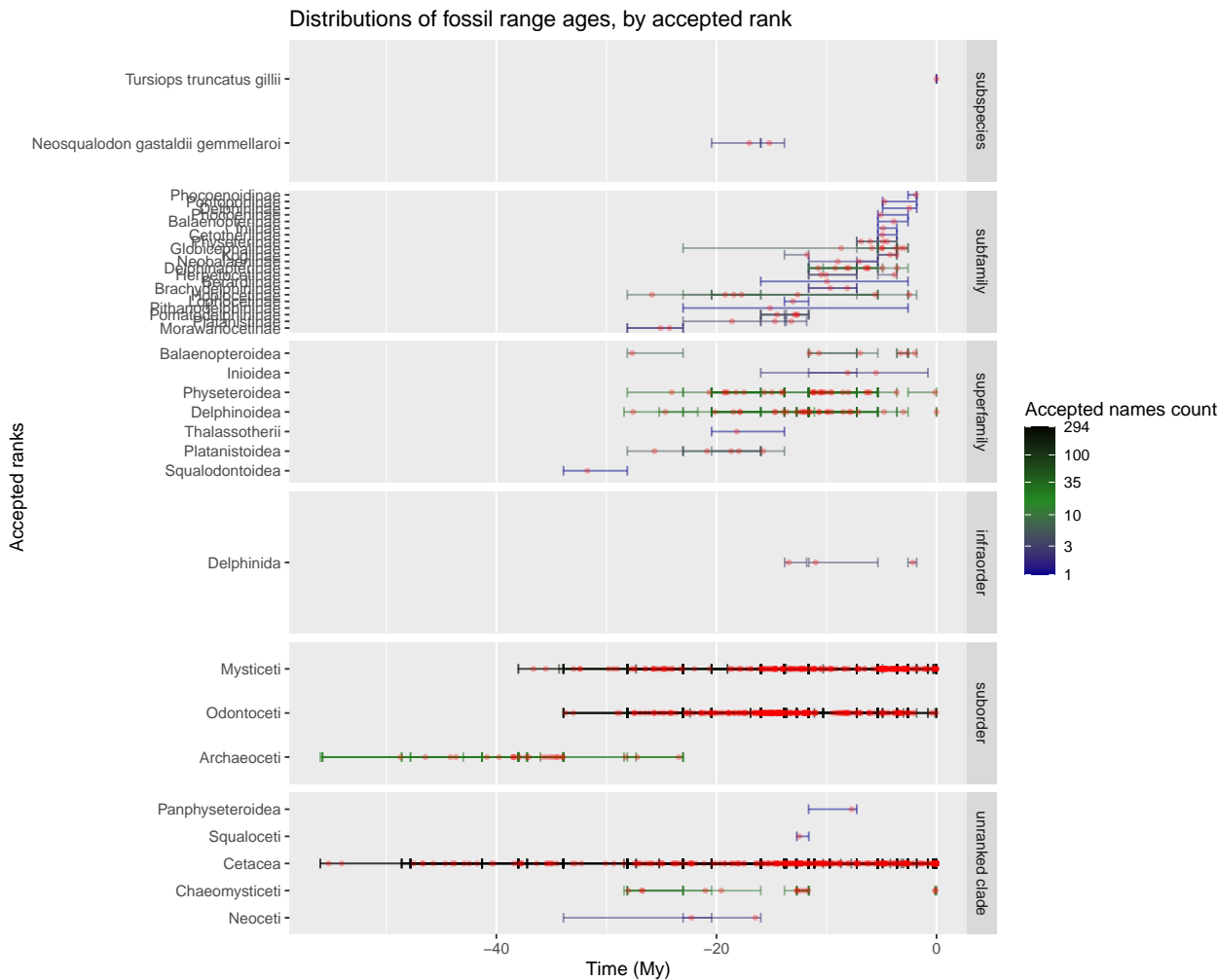


Time ranges = duration of the time intervals

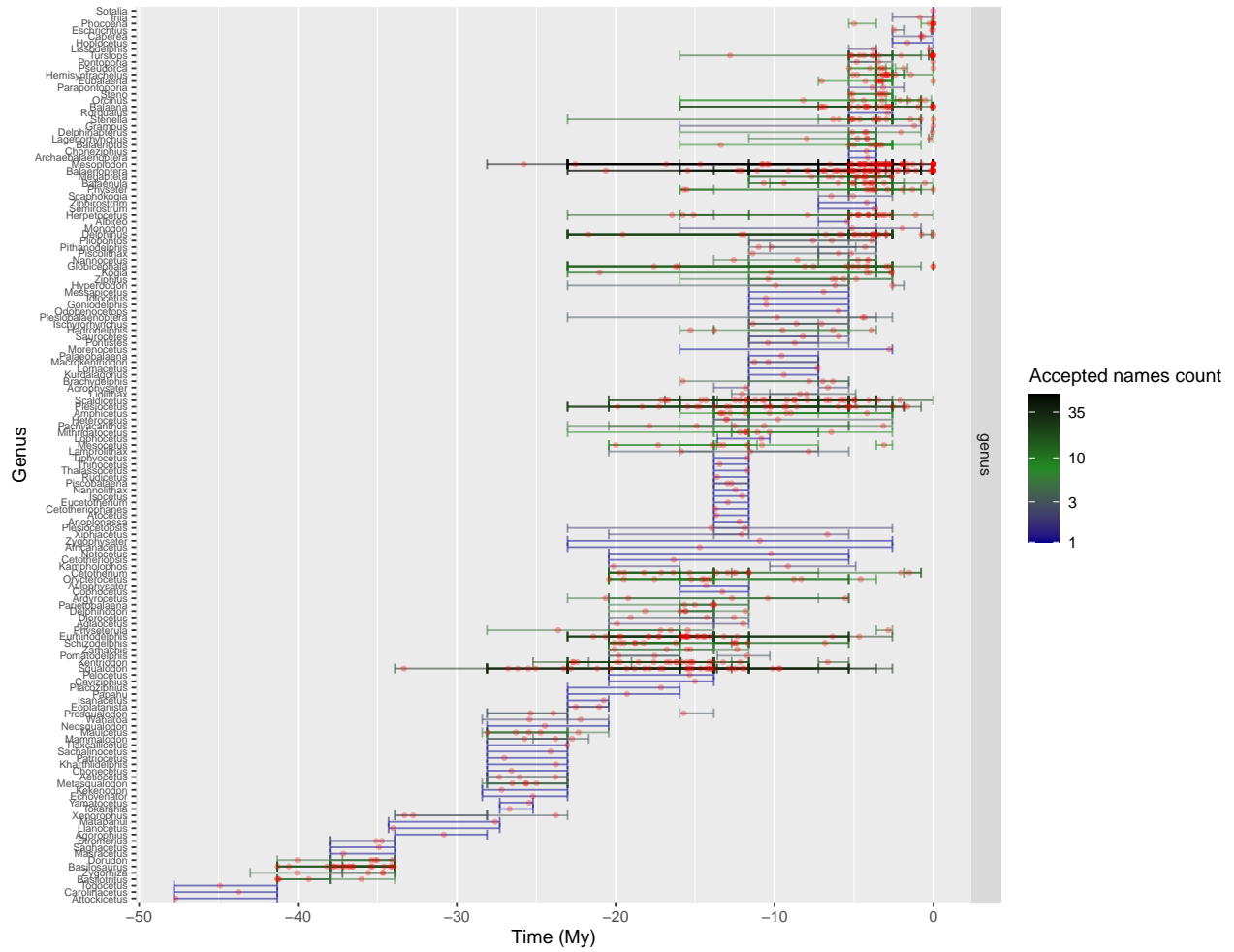
Count occurrences by age interval

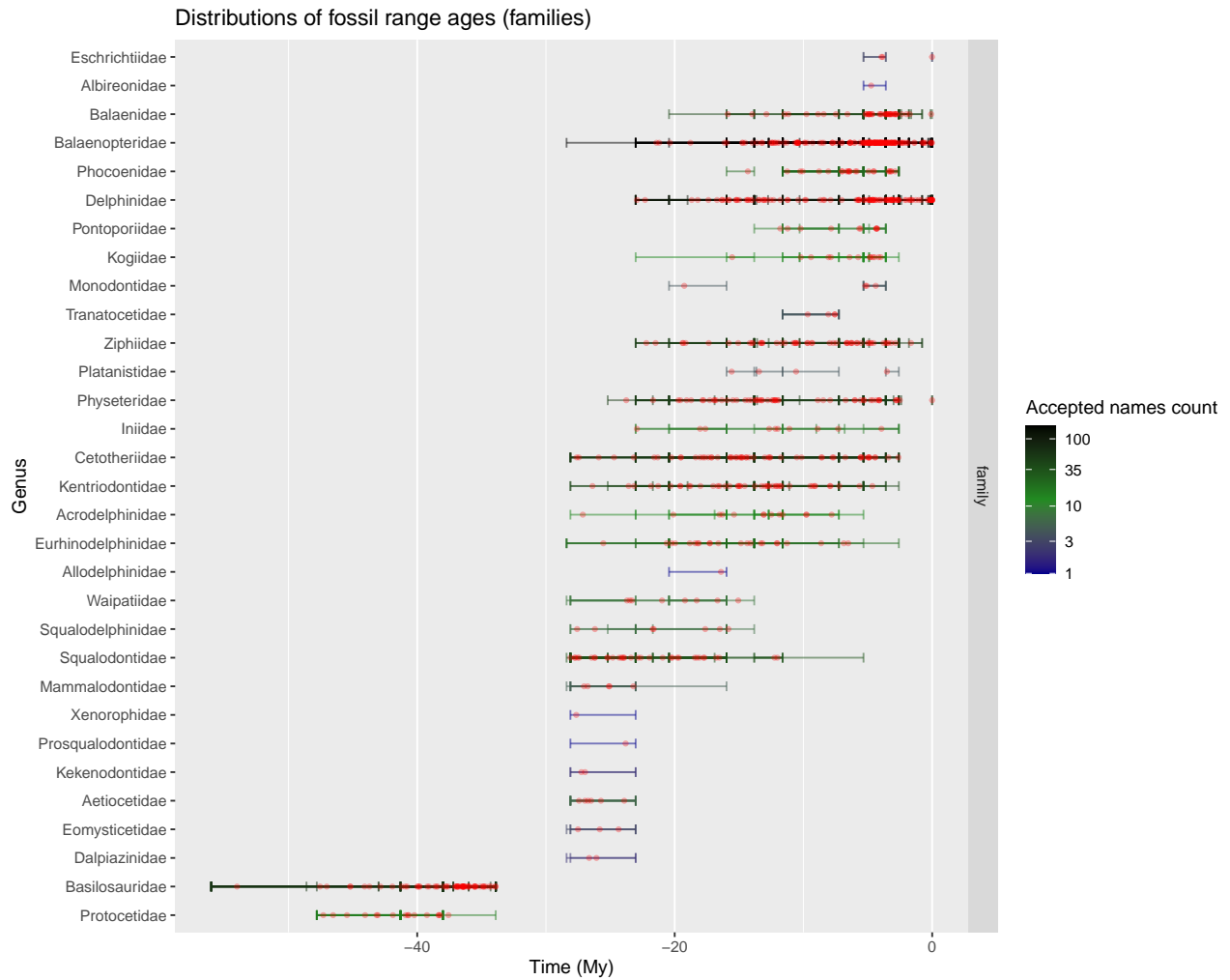


Count occurrences by accepted name



Distributions of fossil range ages (genera)

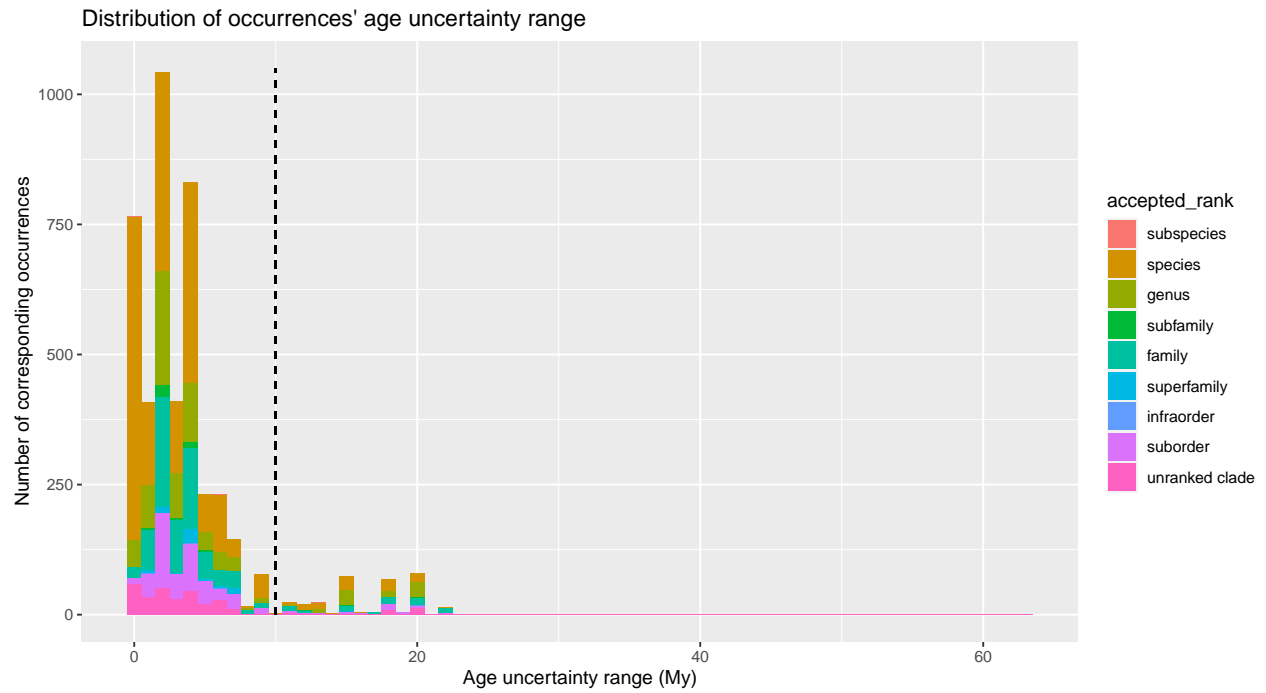




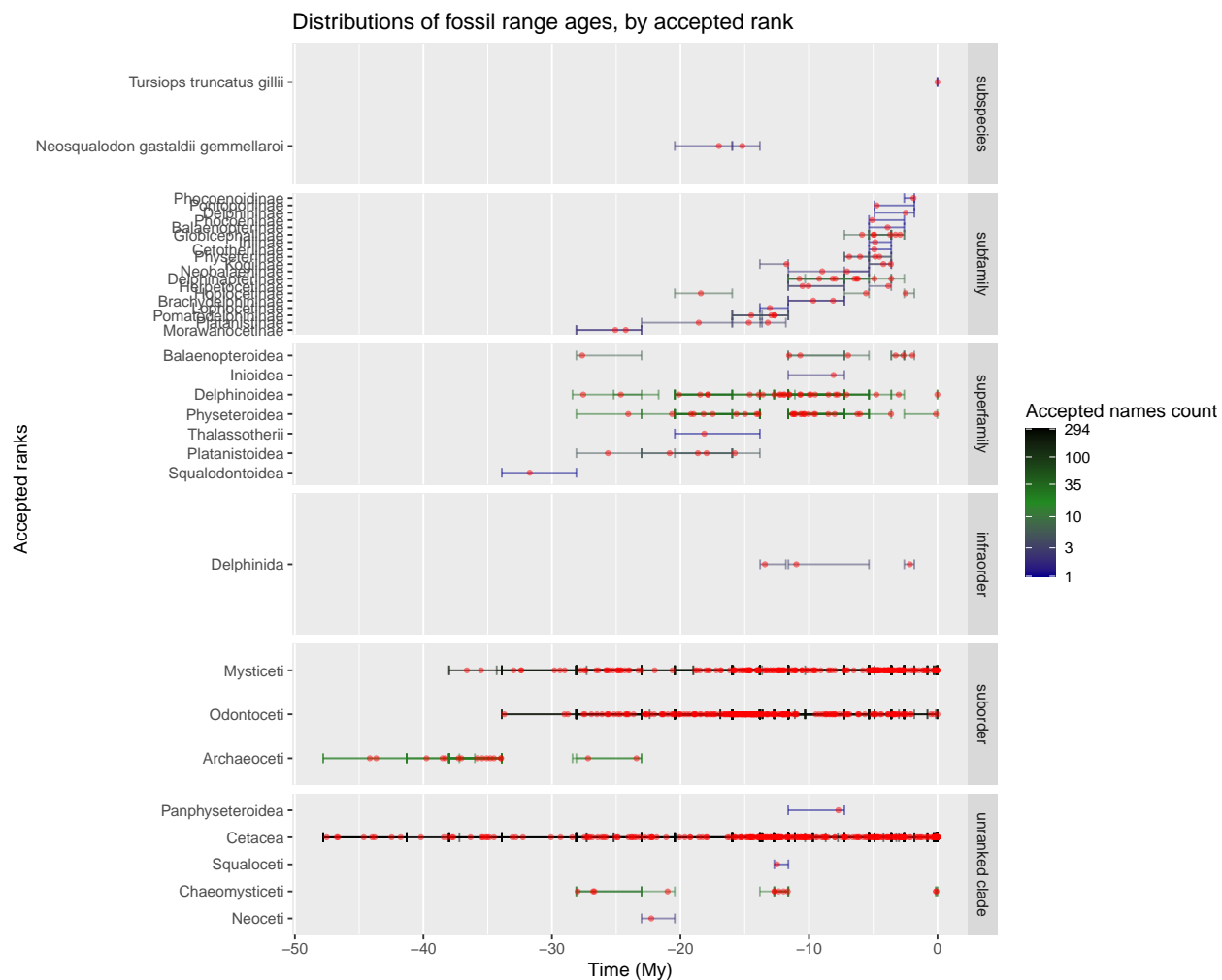
→ Some occurrences have too much age uncertainty, they could be removed because they are not very informative.

Remove occurrences with highly uncertain datation (range > 10My)

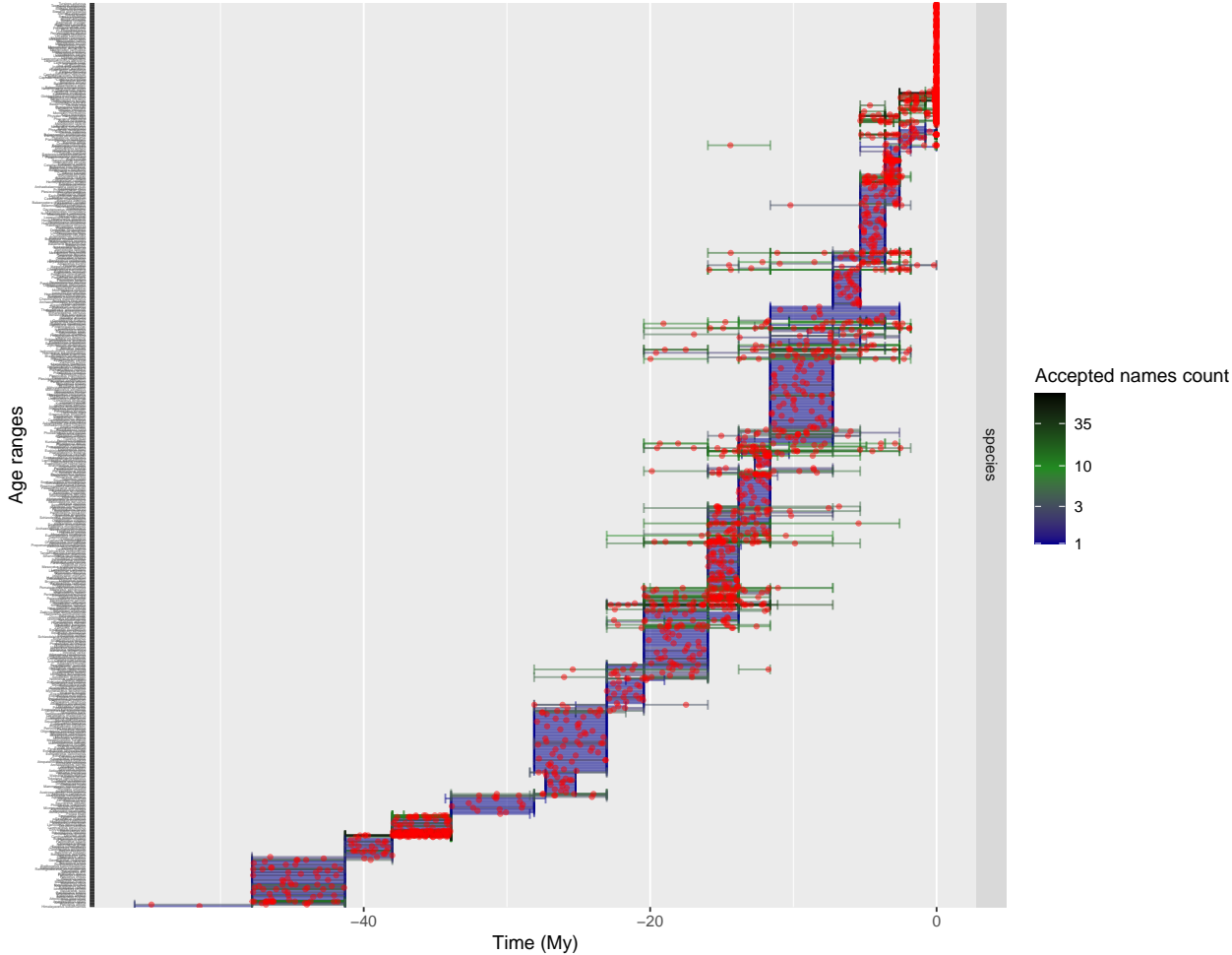
[1] 4157 23



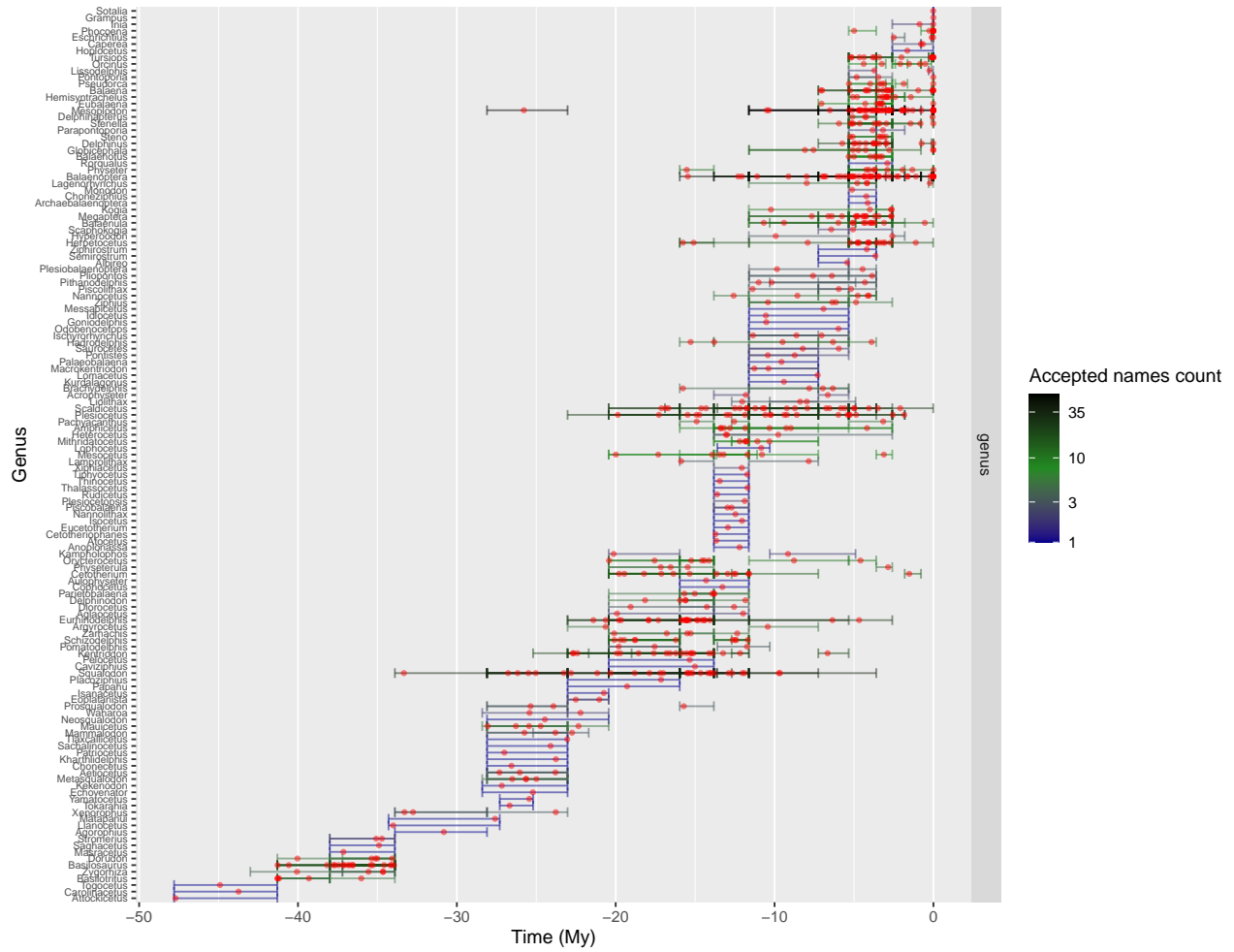
Most of occurrences (4157) show less than 10 My age uncertainty, let's keep only these ones.

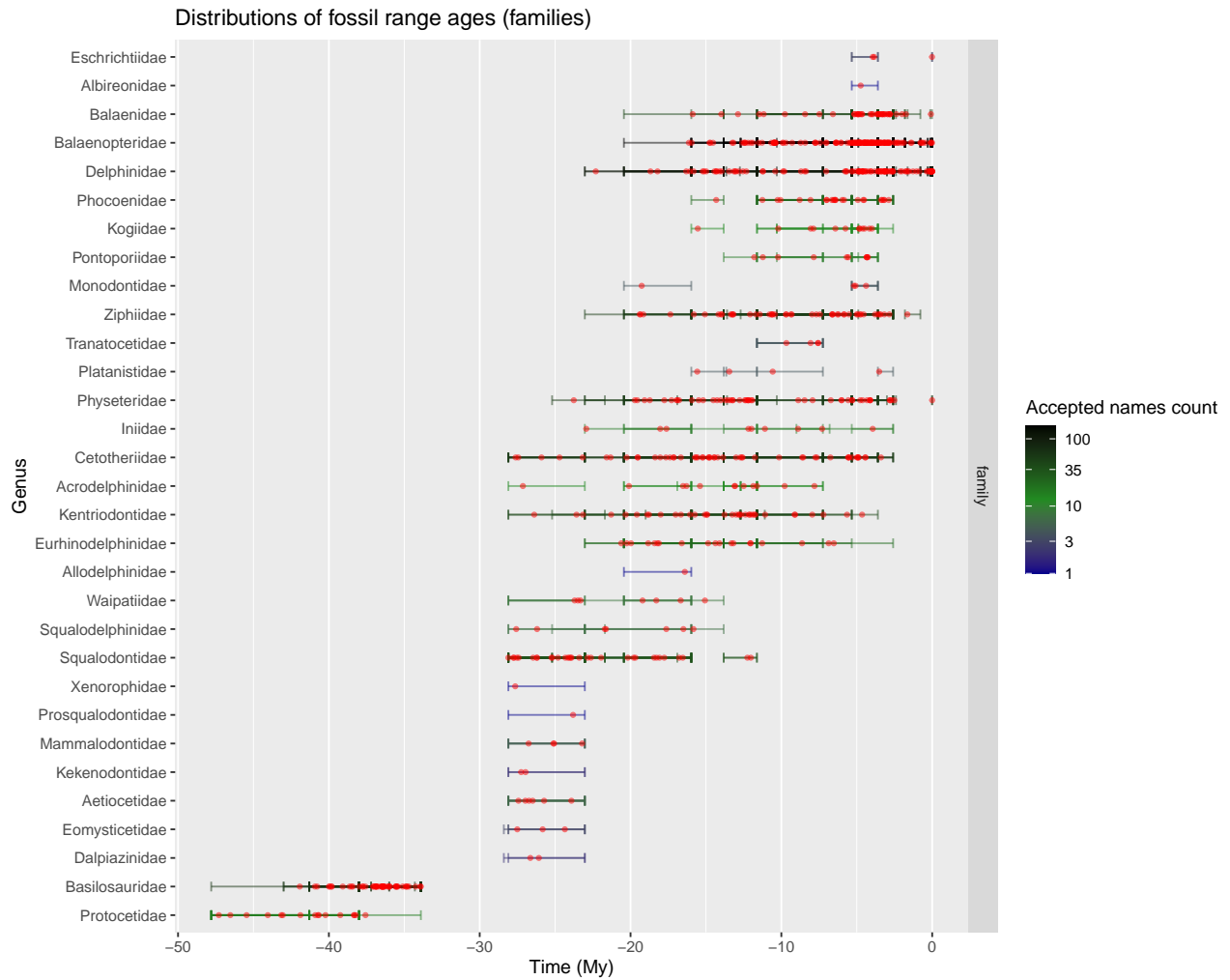


Distributions of fossil range ages (species)



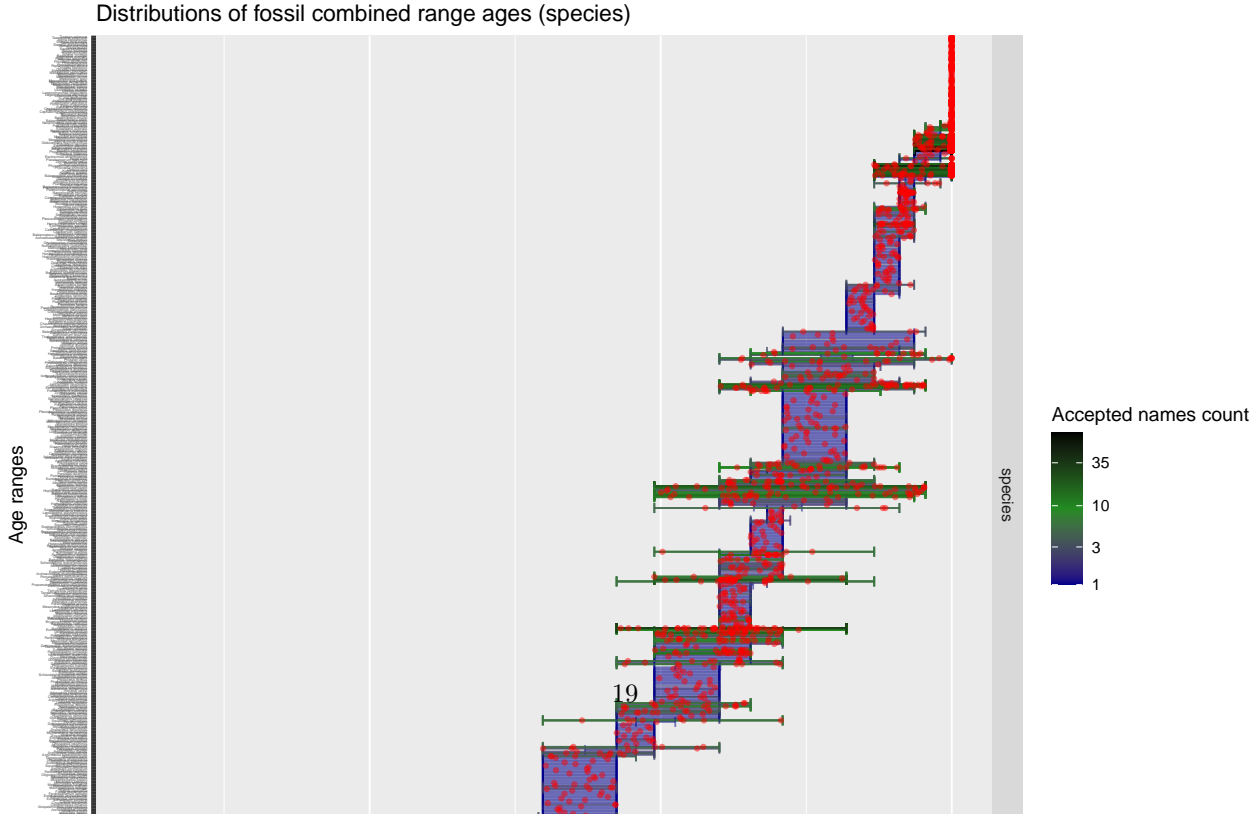
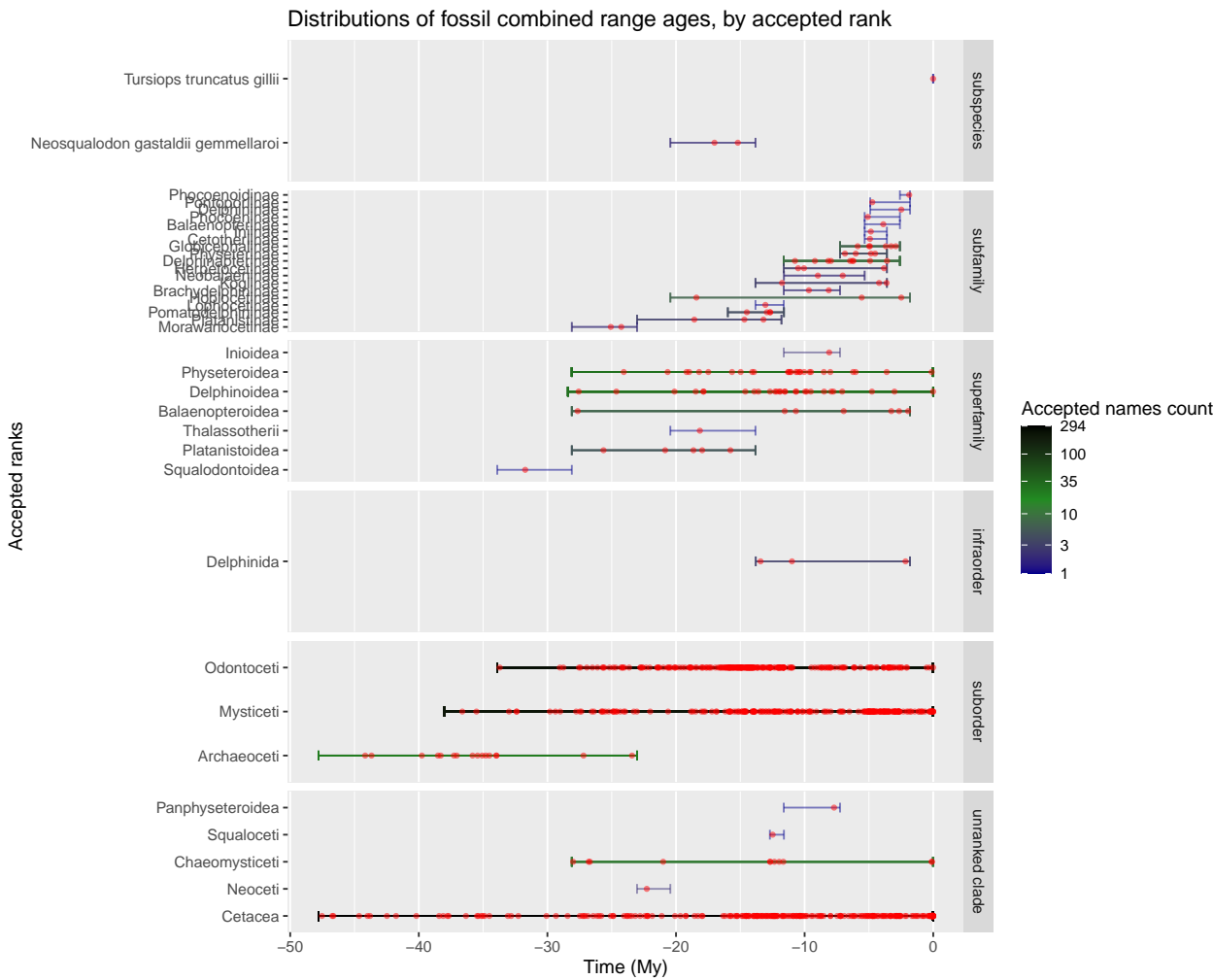
Distributions of fossil range ages (genera)



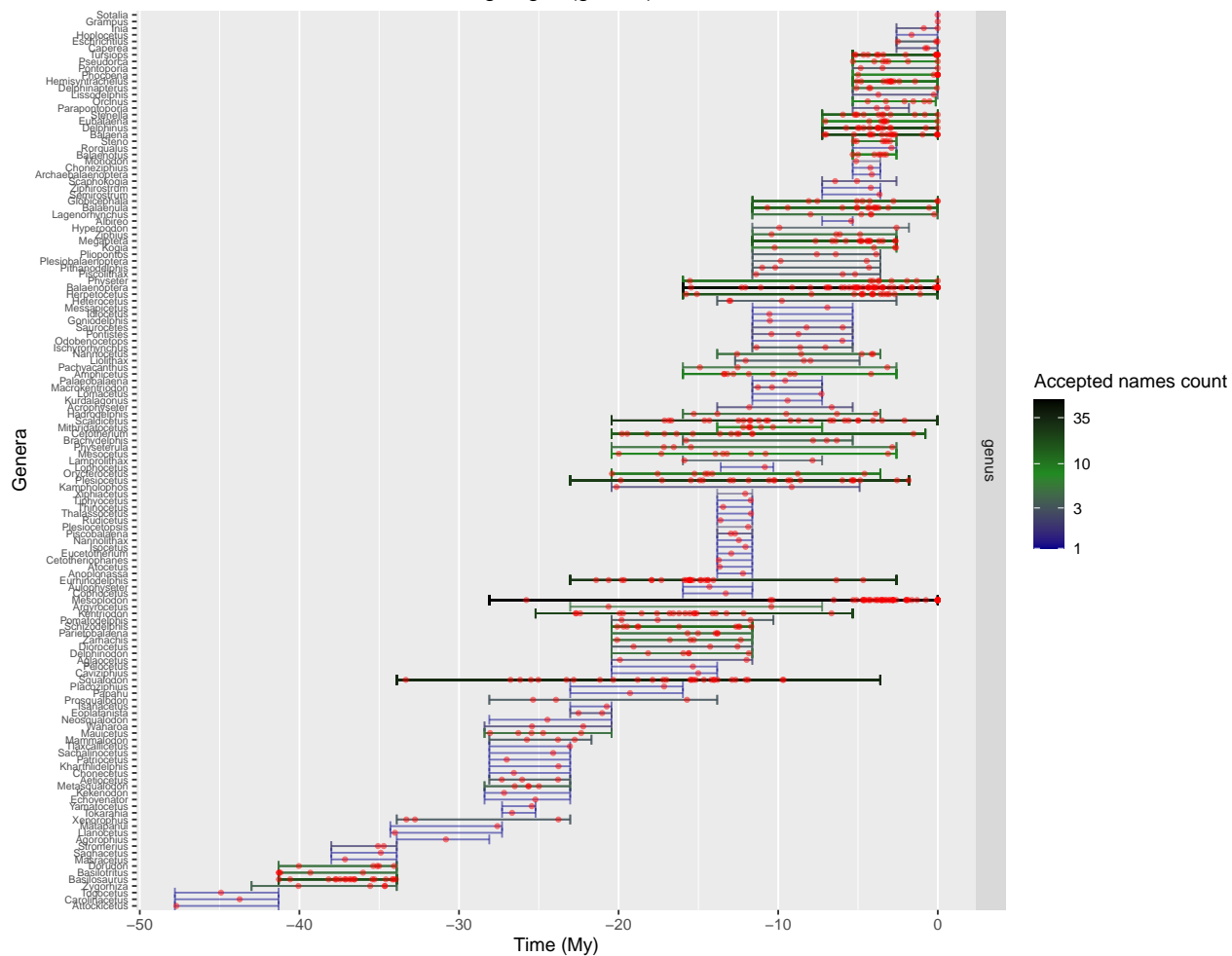


→ Some species (or other ranks) have several occurrences with several time ranges, let's combine them into a unique range covering all the other.

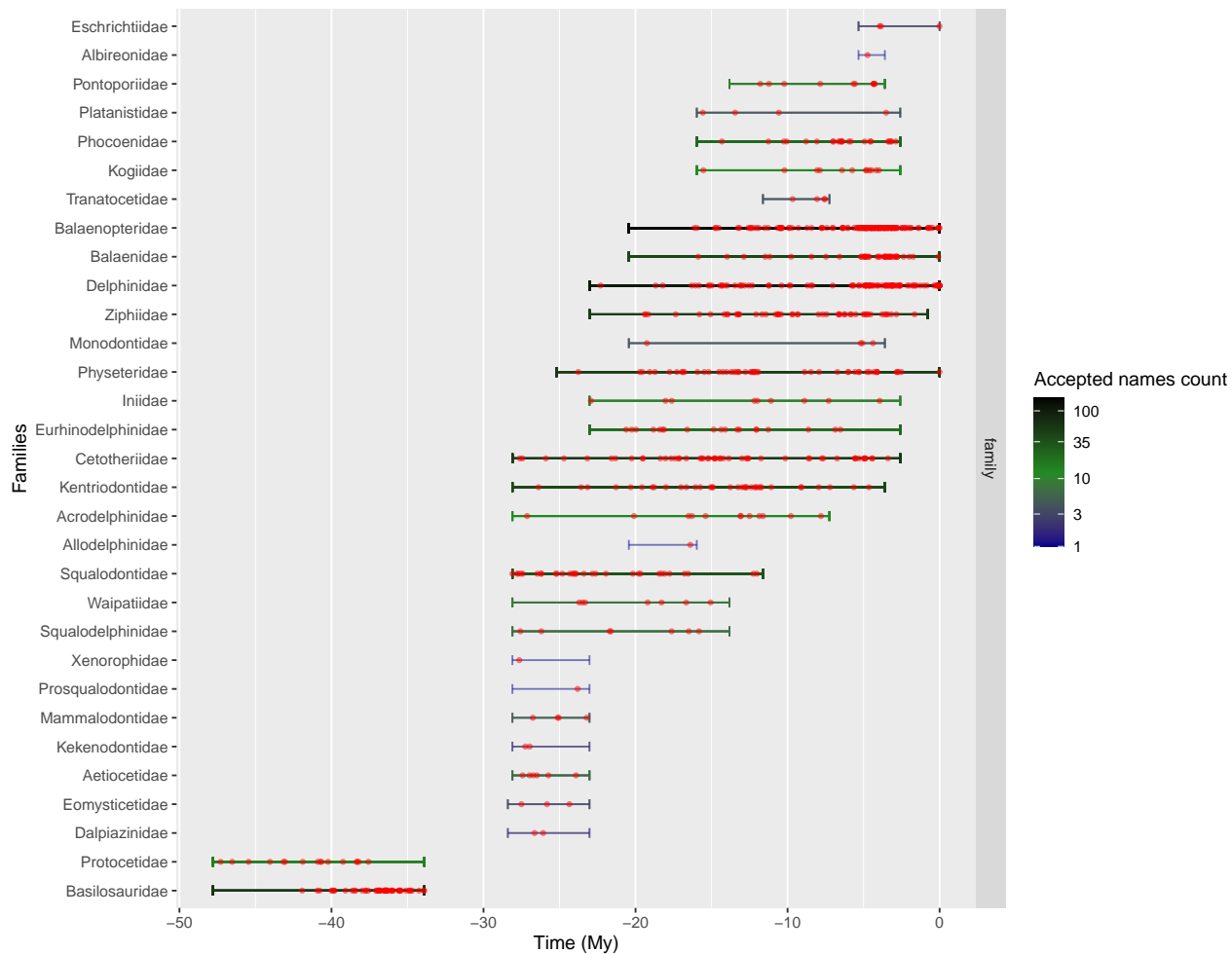
Combined time ranges = unique time range for occurrences with the same name
(without the biggest ones)



Distributions of fossil combined range ages (genera)

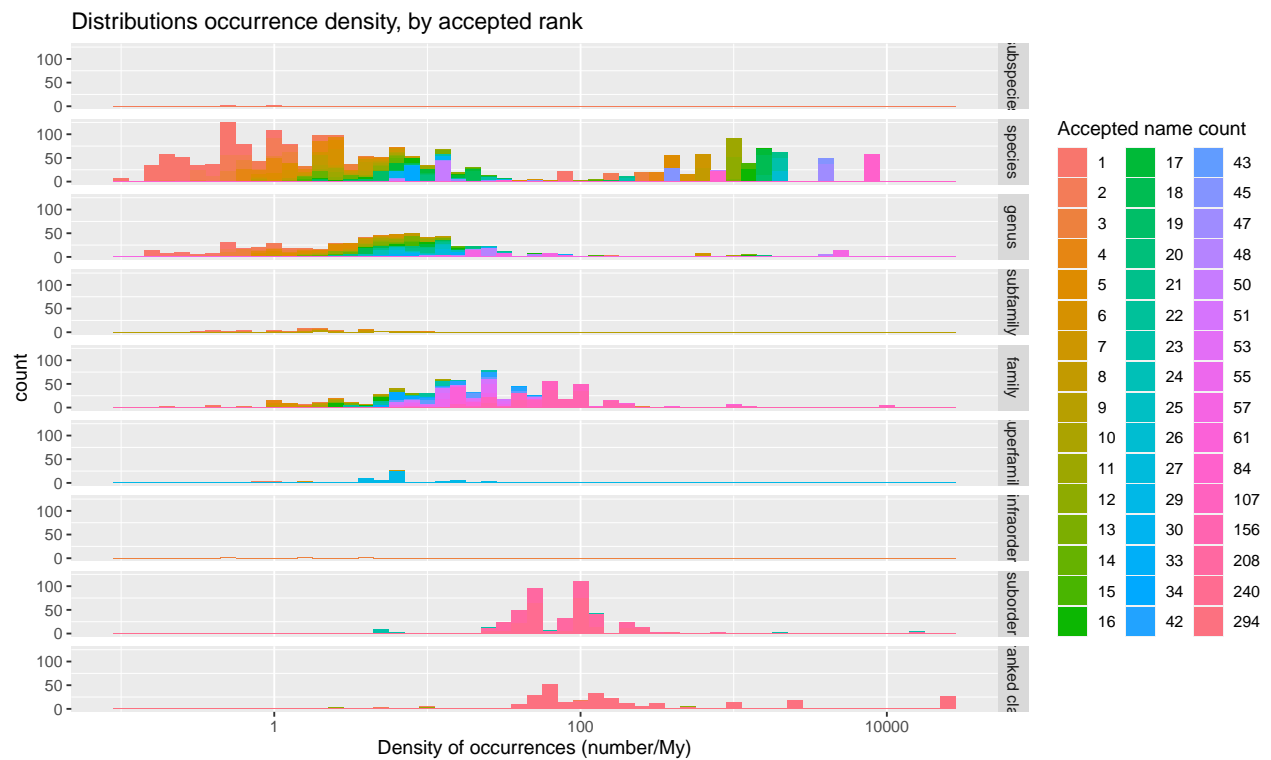


Distributions of fossil combined range ages (families)

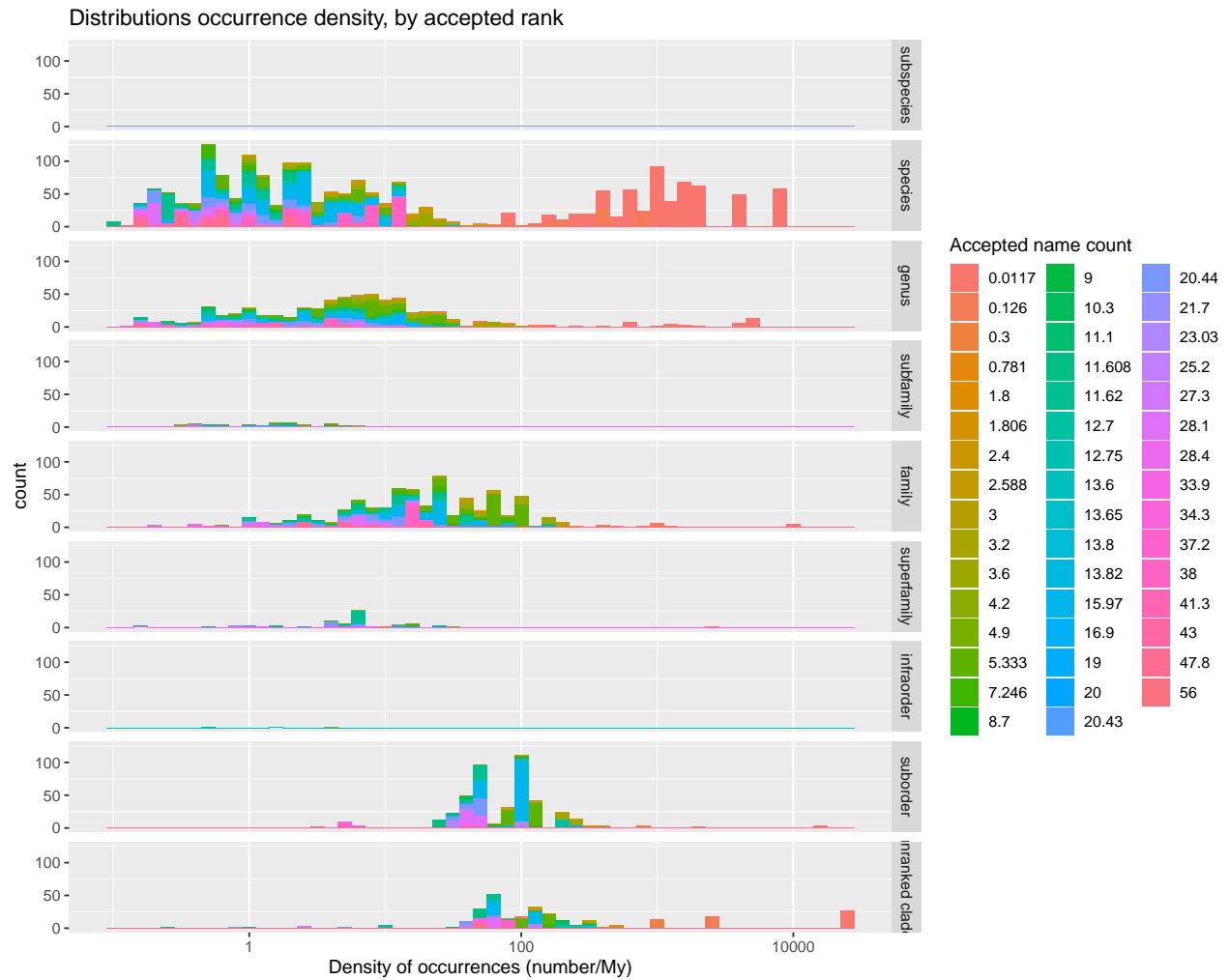


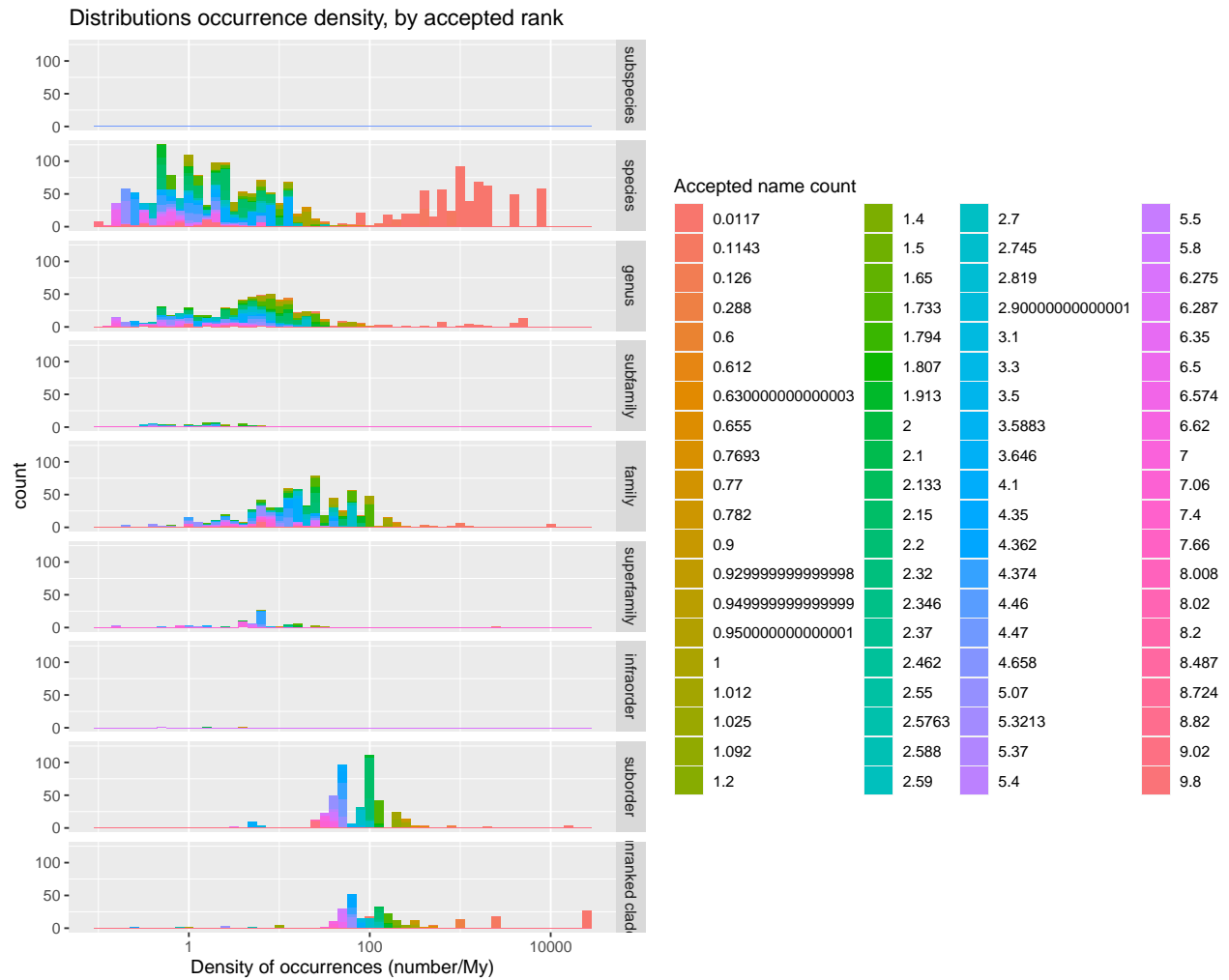
Occurrence density

Density distributions



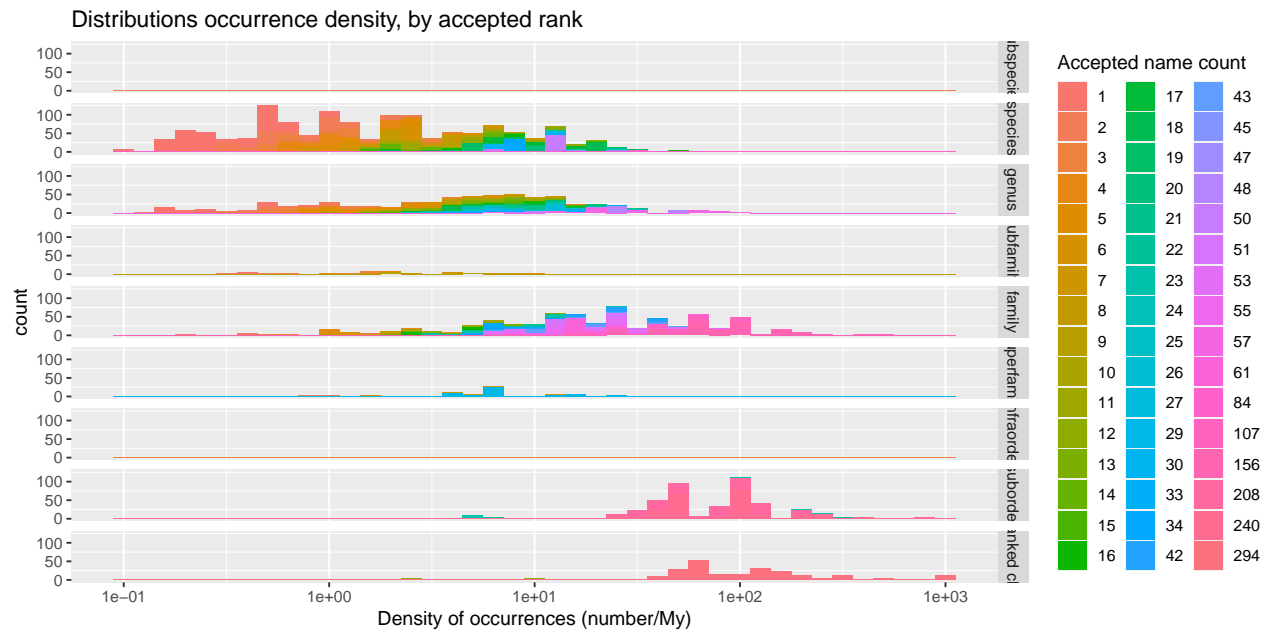
→ Density logically increases as taxa ranks increase, but there is a cluster of high density within the species. Let's identify its origin :





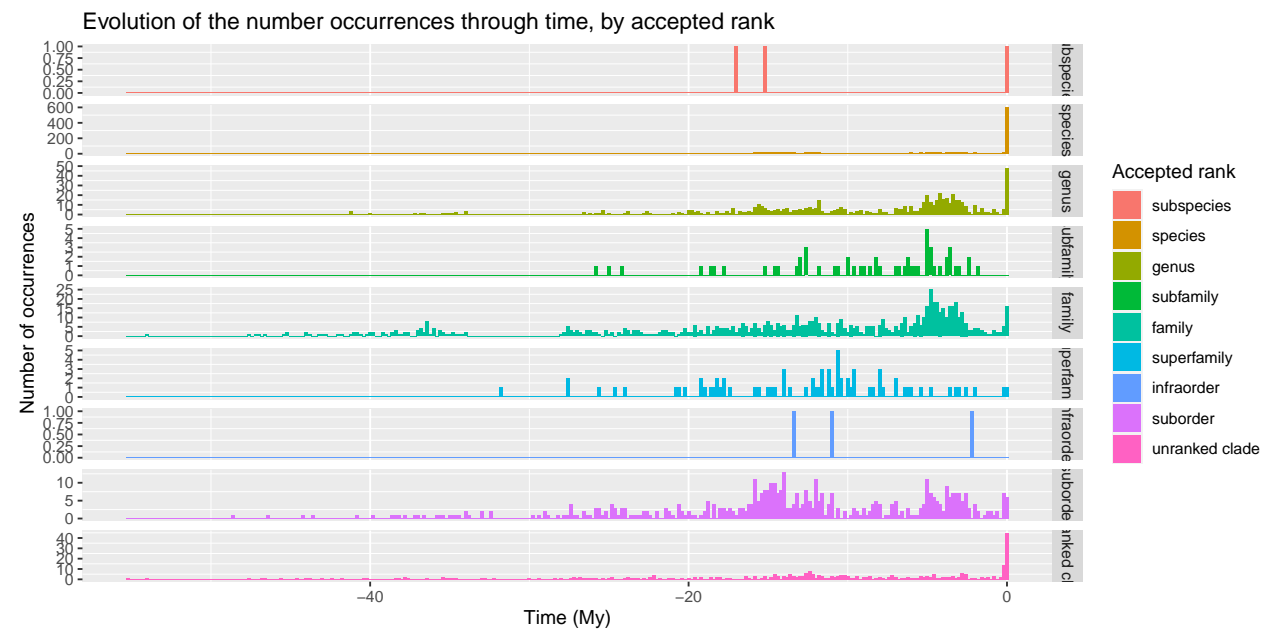
→ This cluster corresponds to recent samples ($< 150,000$ years) with very precise datation (different technique?). Let's try to remove it but later it could be interesting to subsample instead.

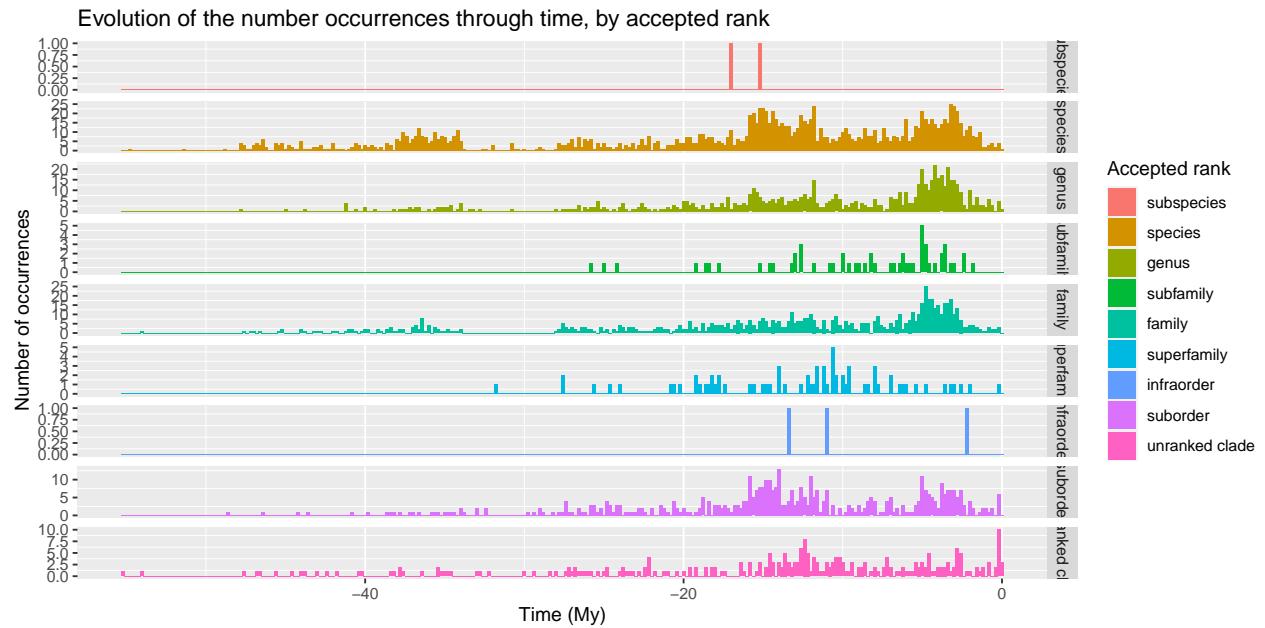
Remove highly concentrated occurrences at present



→ The distributions look normal again.

Compare the number of occurrences through time with or without the concentrated ones at present

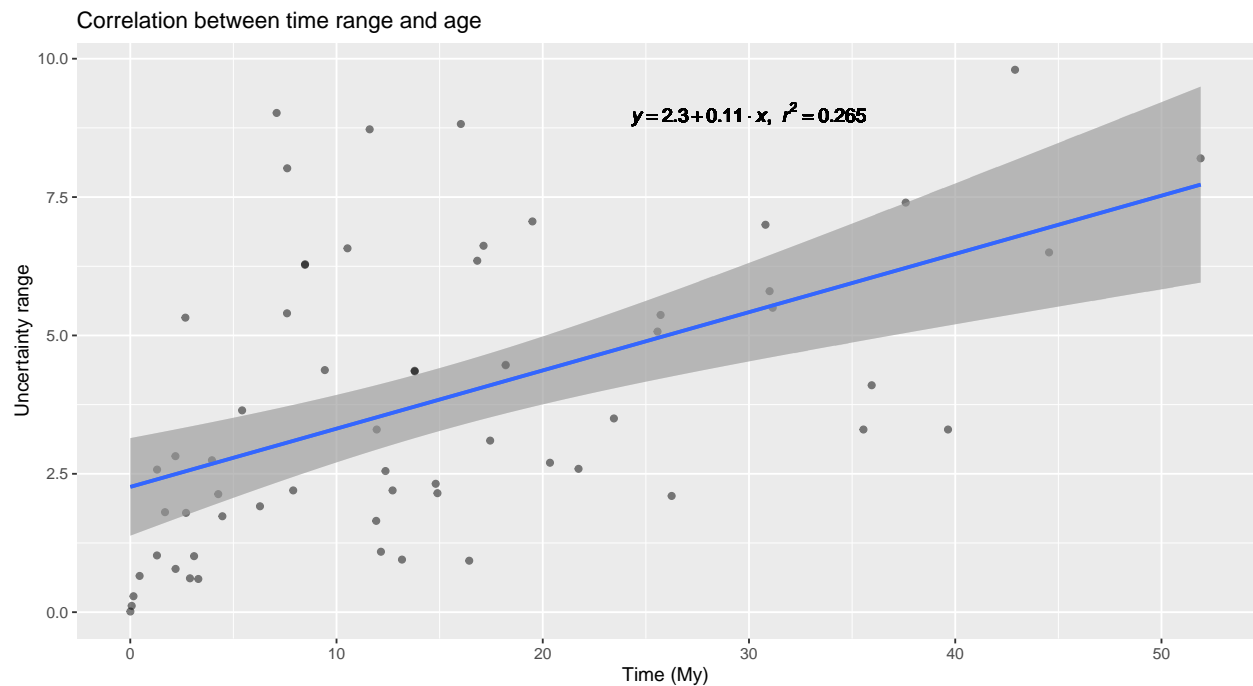




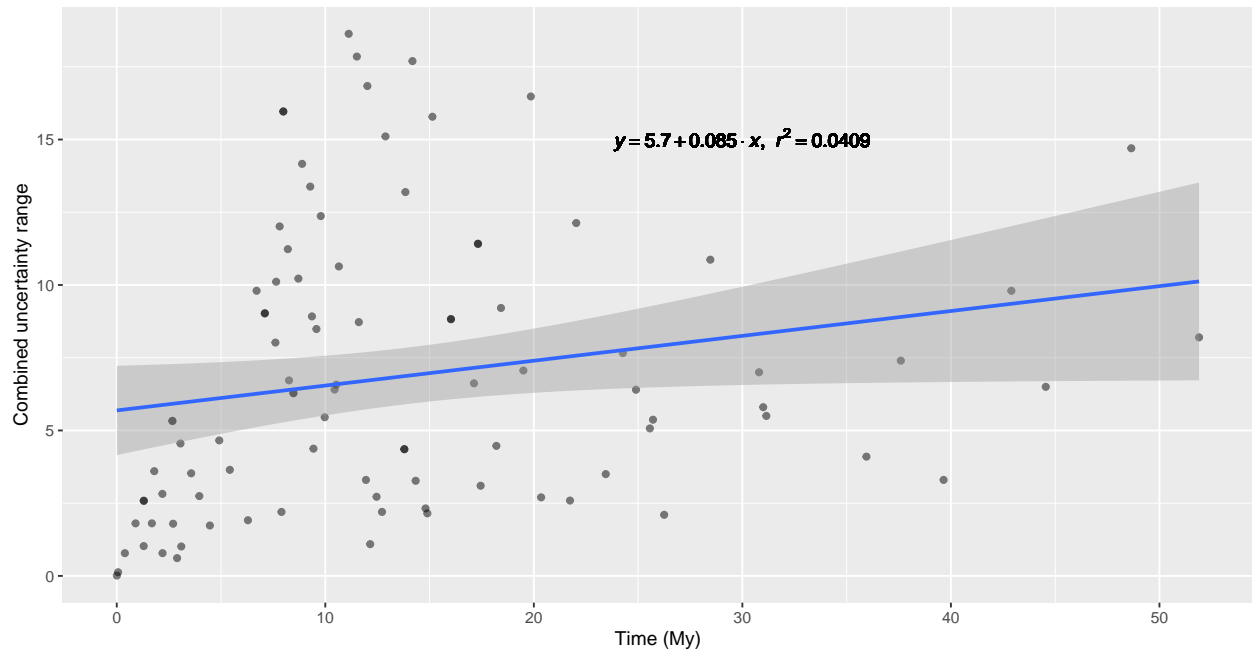
→ The removal gives much clearer occurrence distributions at the species and genera levels.

Correlation between time range and age

If we want to correct species abundance differences based on the number of occurrences in the time range (“density”), those factors should not depend on time in order to avoid penalizing periods with bigger ranges.



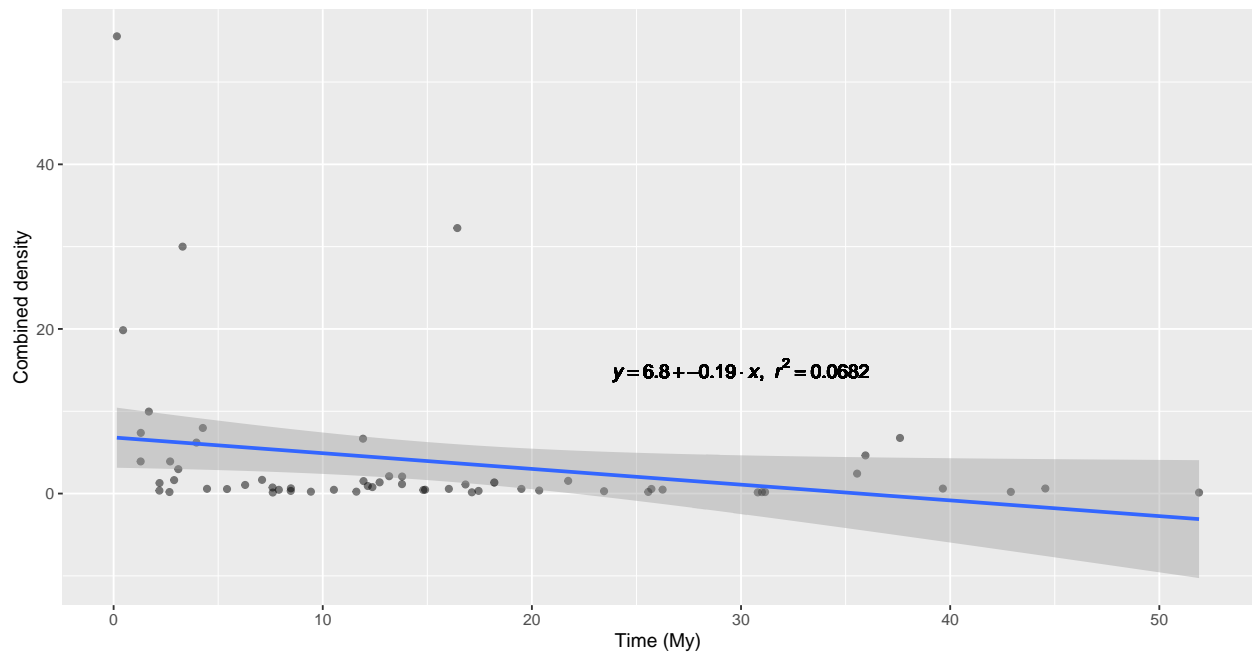
Correlation between combined time range and age

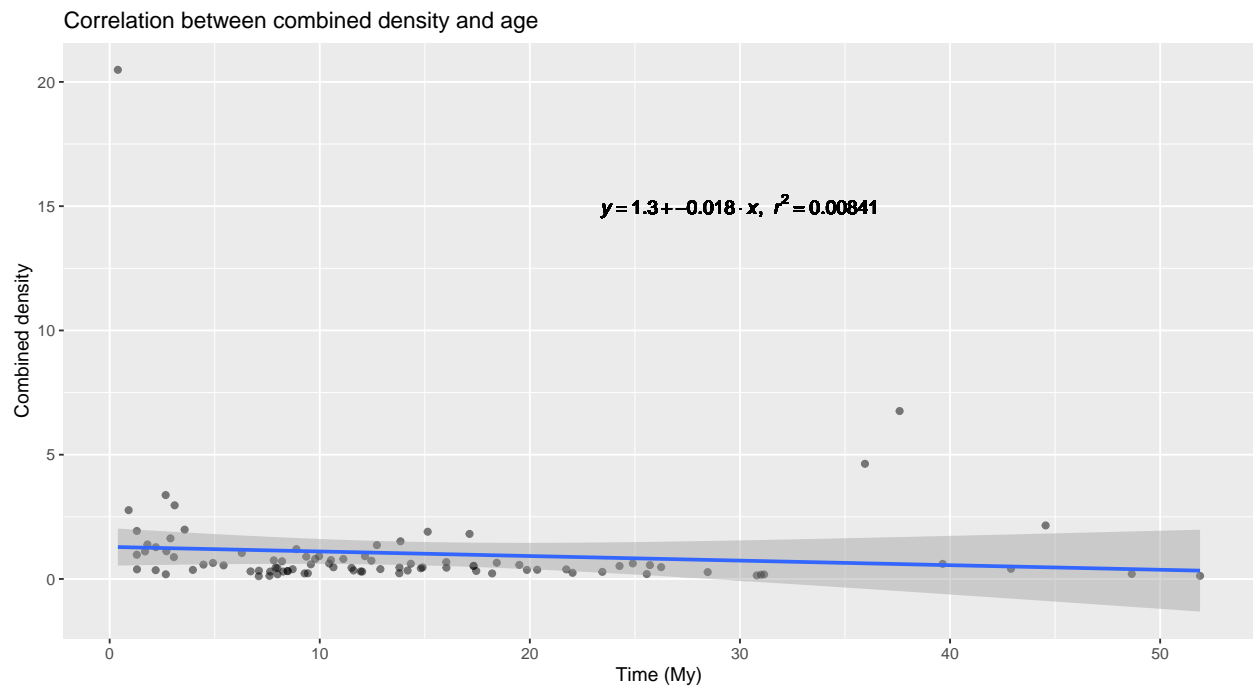


→ It seems that age range is much less correlated with time when we take the full combined range into account. However this plot is quite ugly (“triangle” instead of a nice point cloud) so this correlation may not be very meaningful. Nevertheless, we will use those ranges for normalising the occurrence density because we don’t want to penalize older specimens.

Let’s look at the density directly, because this is what is interesting us directly.

Correlation between density and age



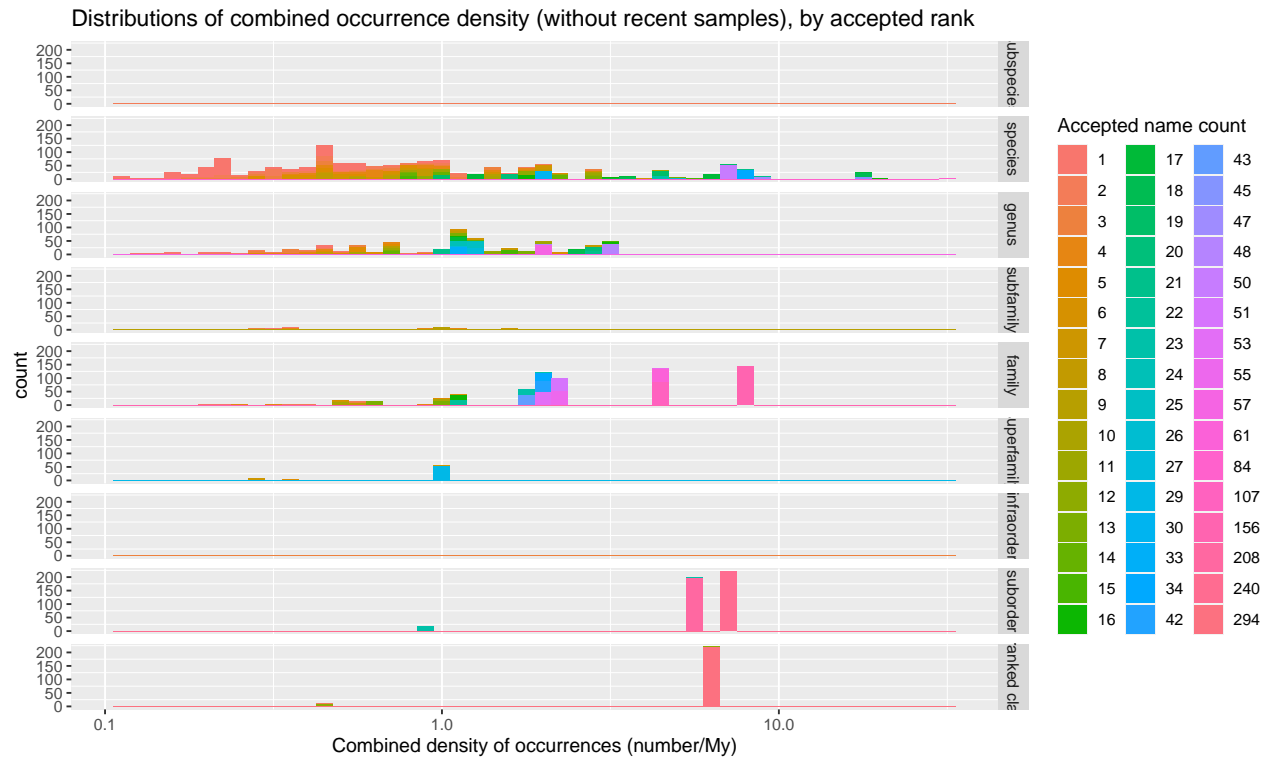


→ Again the combined version is less correlated with time.

Sub-sampling of occurrences with a normalized density along the combined ranges

Compare densities for single vs. combined ranges.

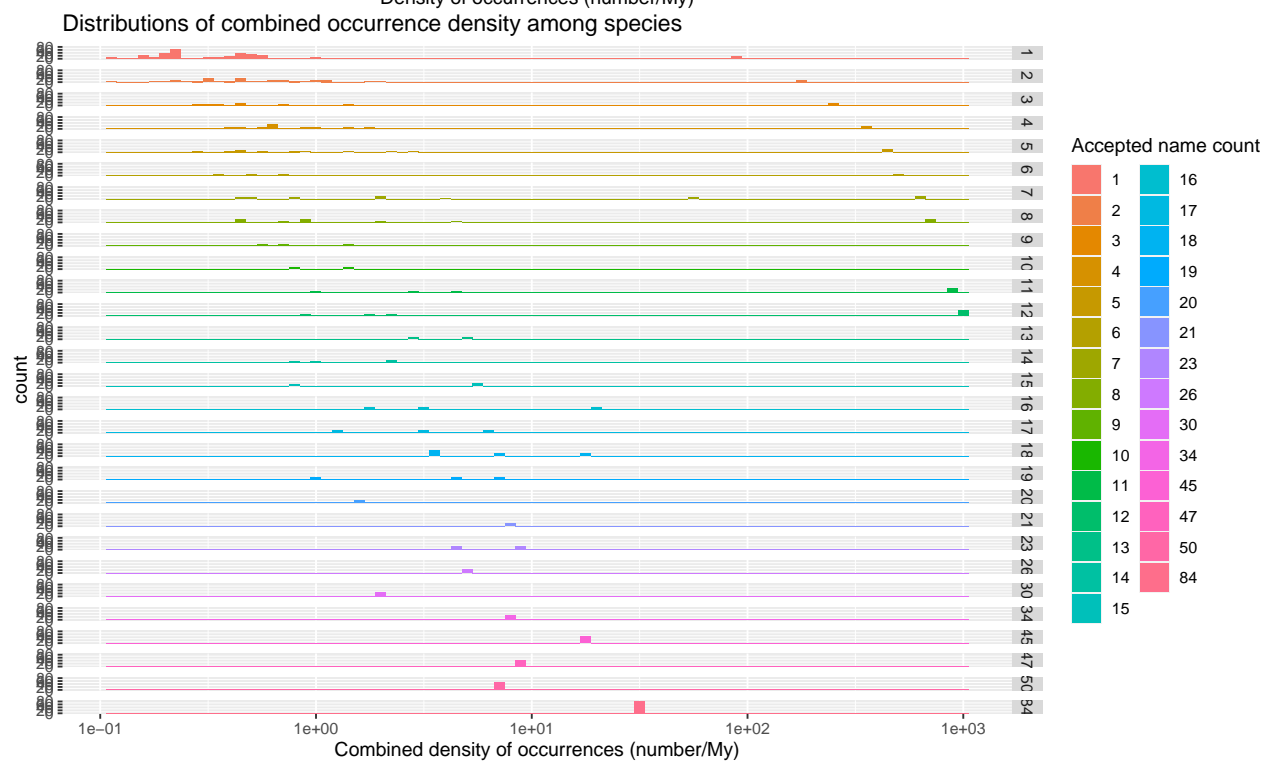
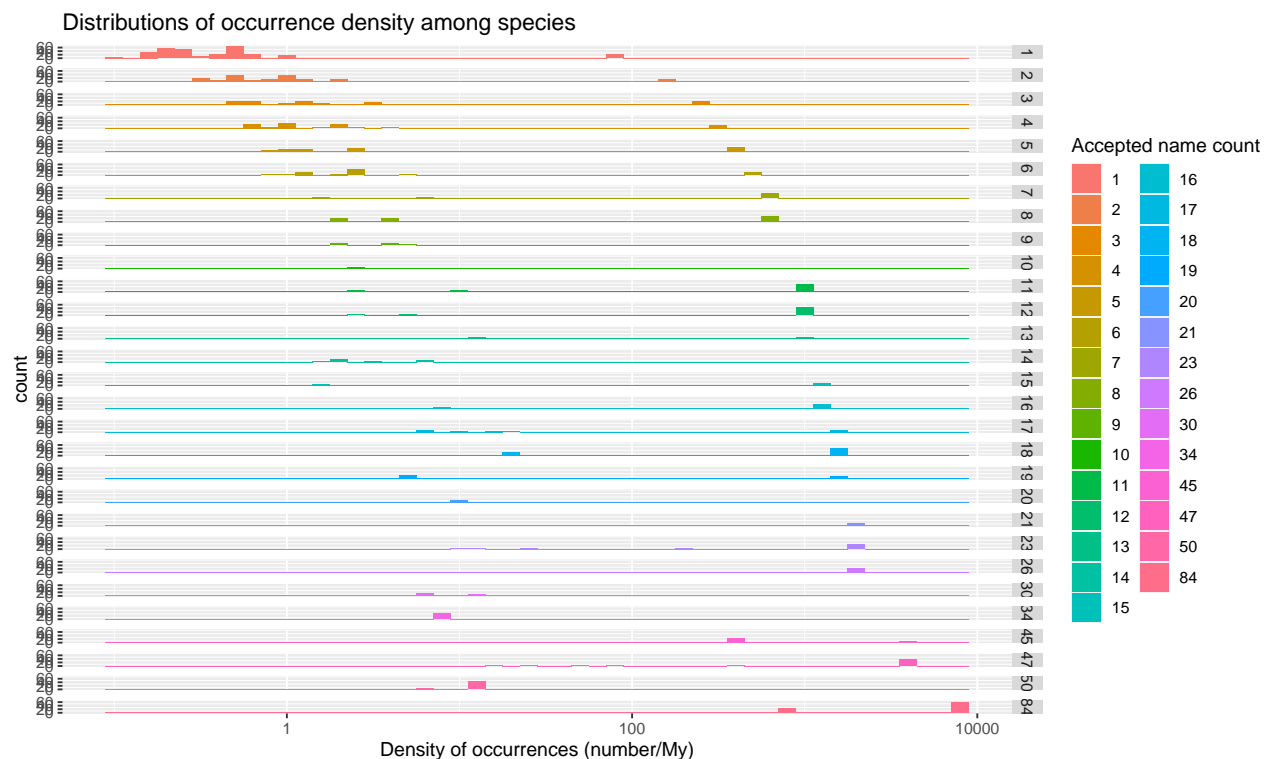


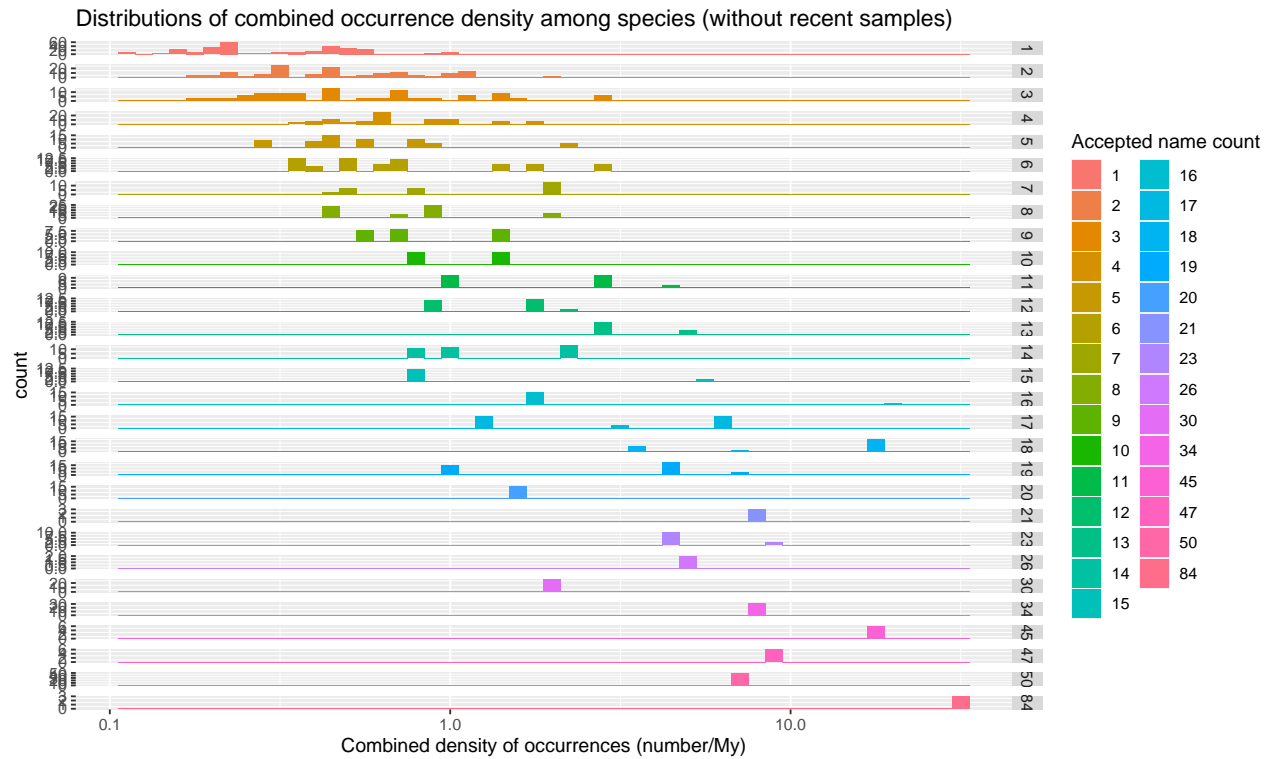


→ Densities are smaller and more concentrated with the combined ranges (larger time span + less ranges in total because of combination).

Compare densities by accepted name count (species only)

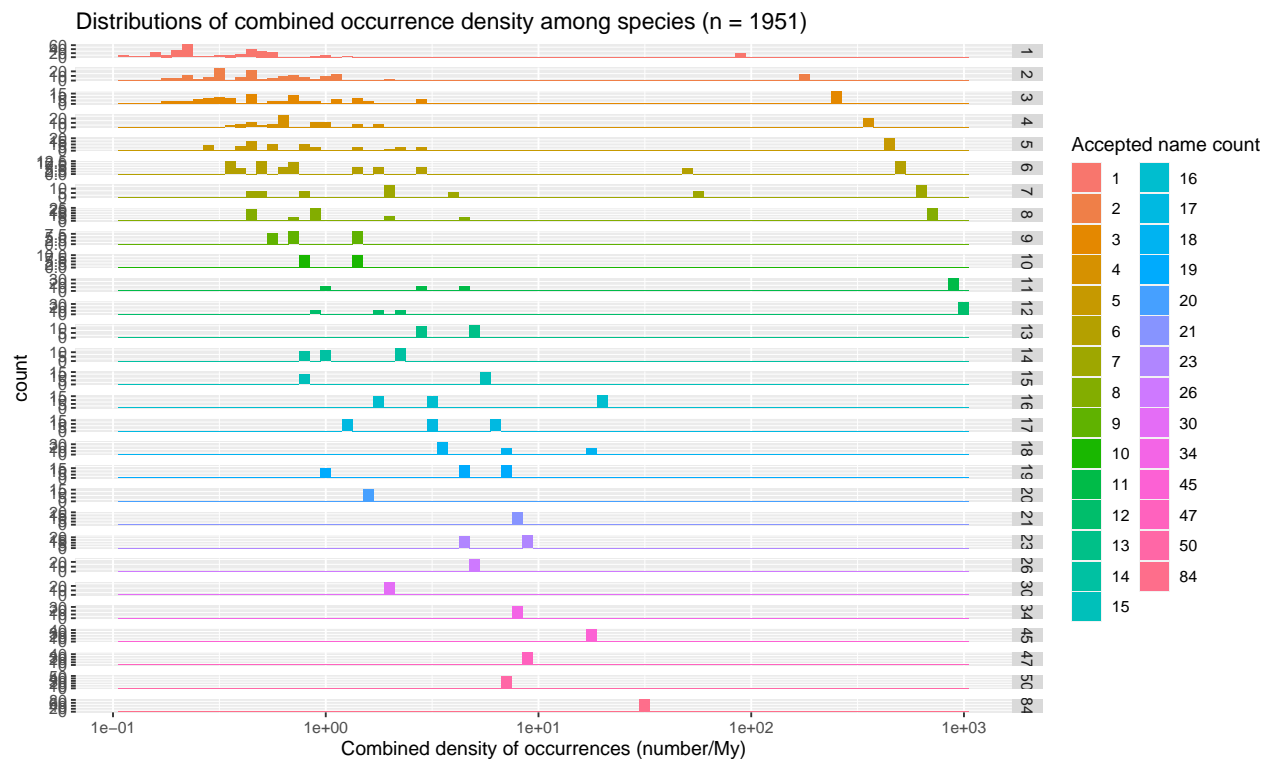
Let's focus now on the occurrences accepted at the species level because they are the one for which we can correct the abundance bias by subsampling the most concentrated combined intervals.



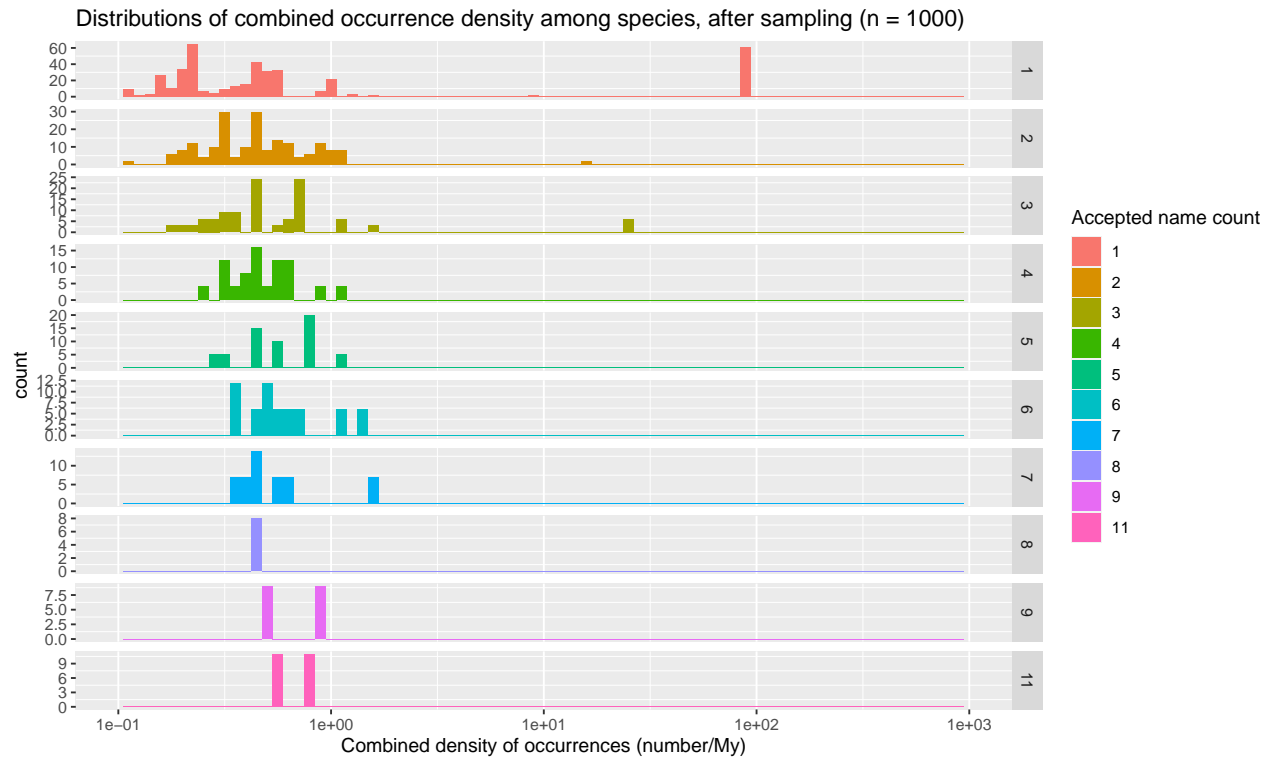


→ There is a huge span of densities driven by the number of occurrences for the same species that we can reduce by subsampling the most concentrated intervals.

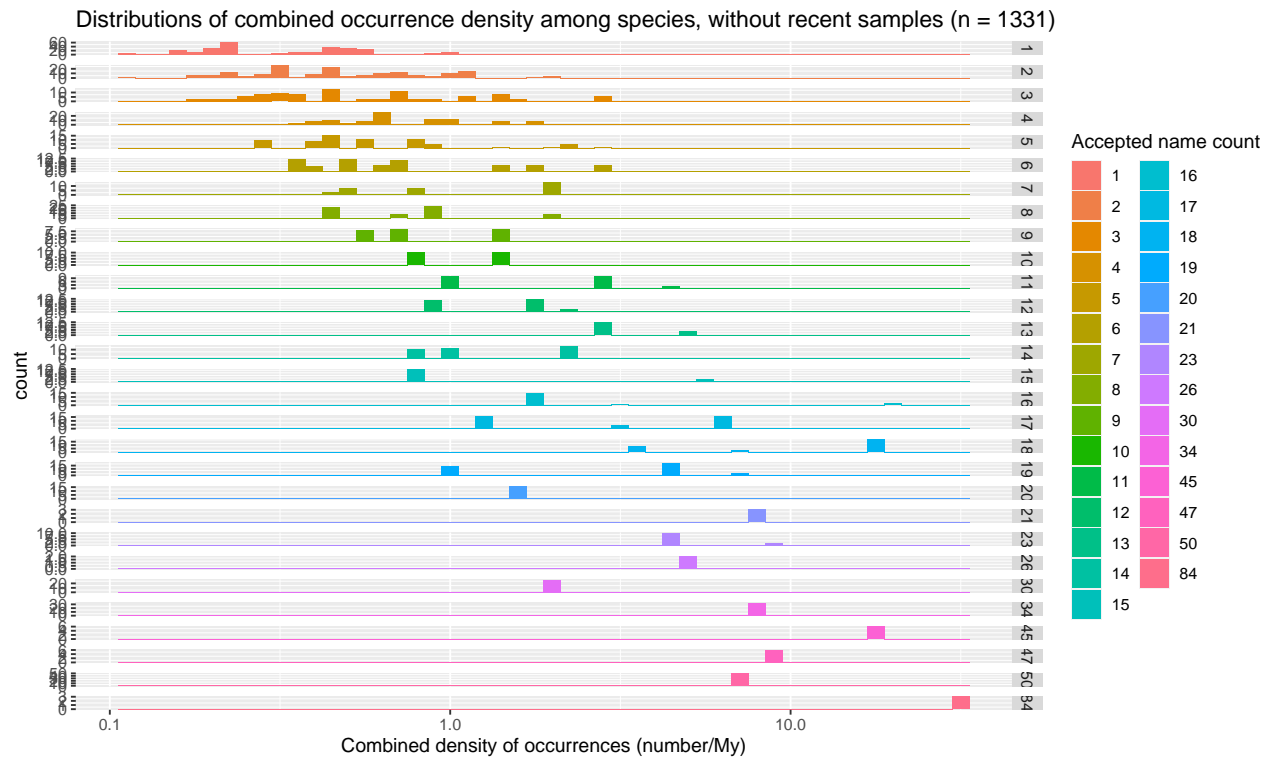
Impact of correcting subsampling on density distributions (species only)



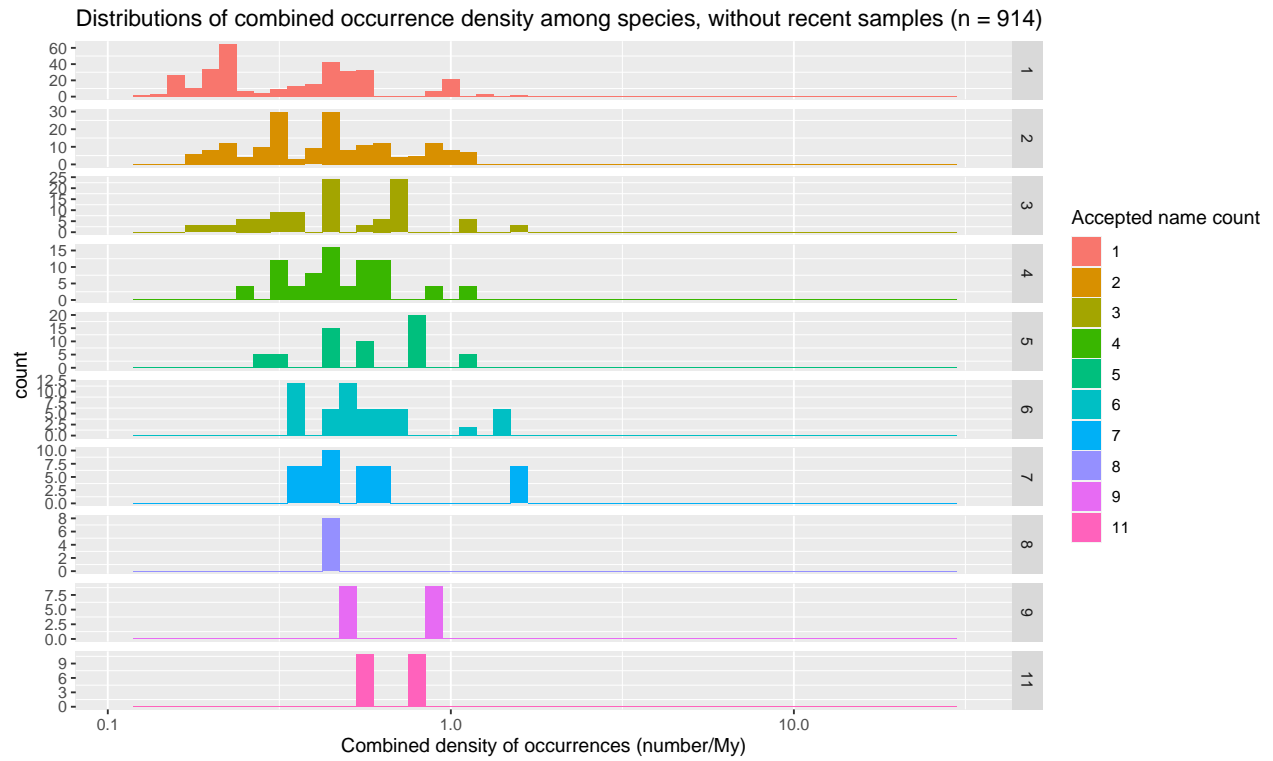
Warning: Removed 20 rows containing missing values (geom_bar).



→ Subsampling successfully reduces the density span from 2 to 1 order of magnitude, apart from the artefactual recent samples that we can hide :



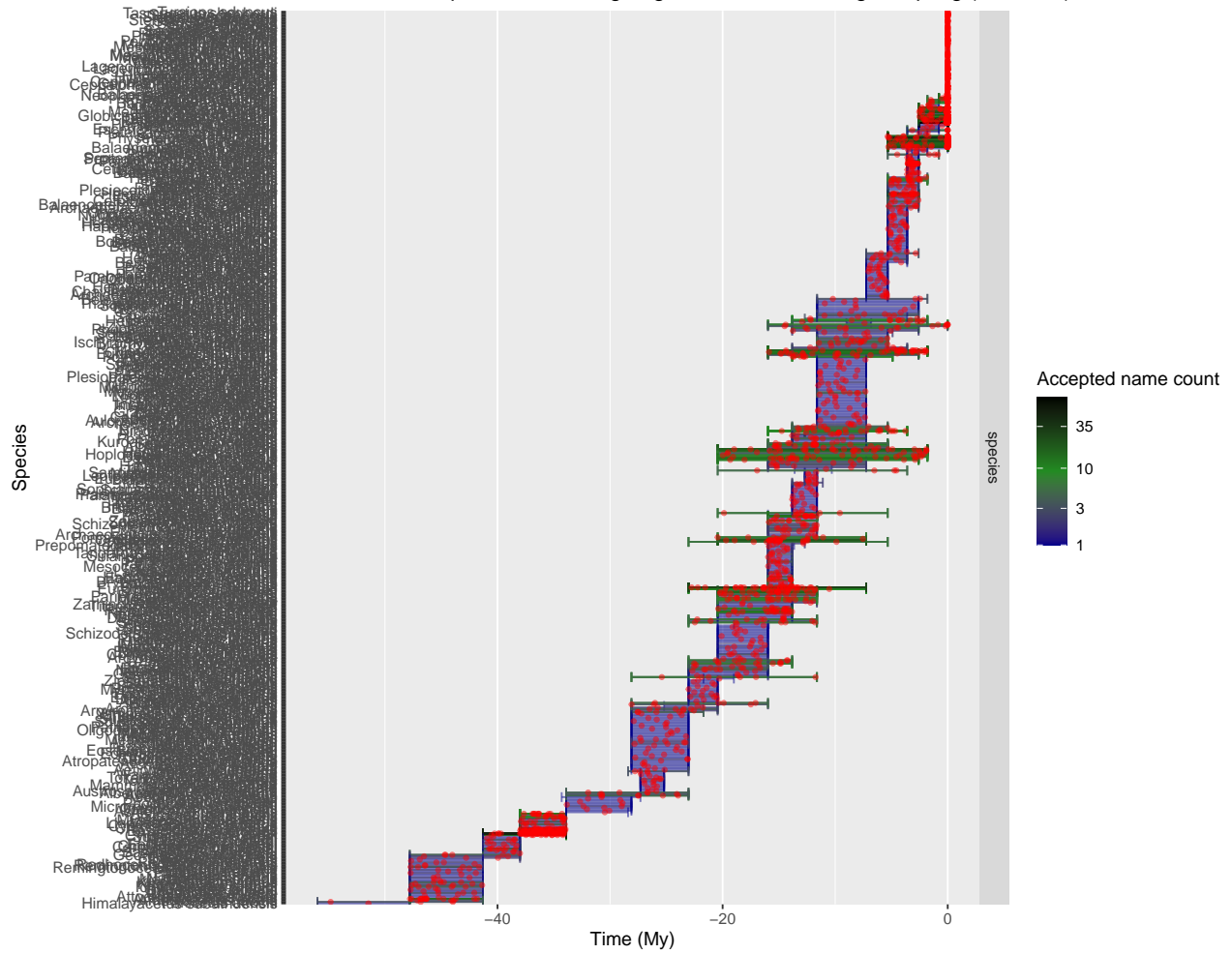
Warning: Removed 20 rows containing missing values (geom_bar).

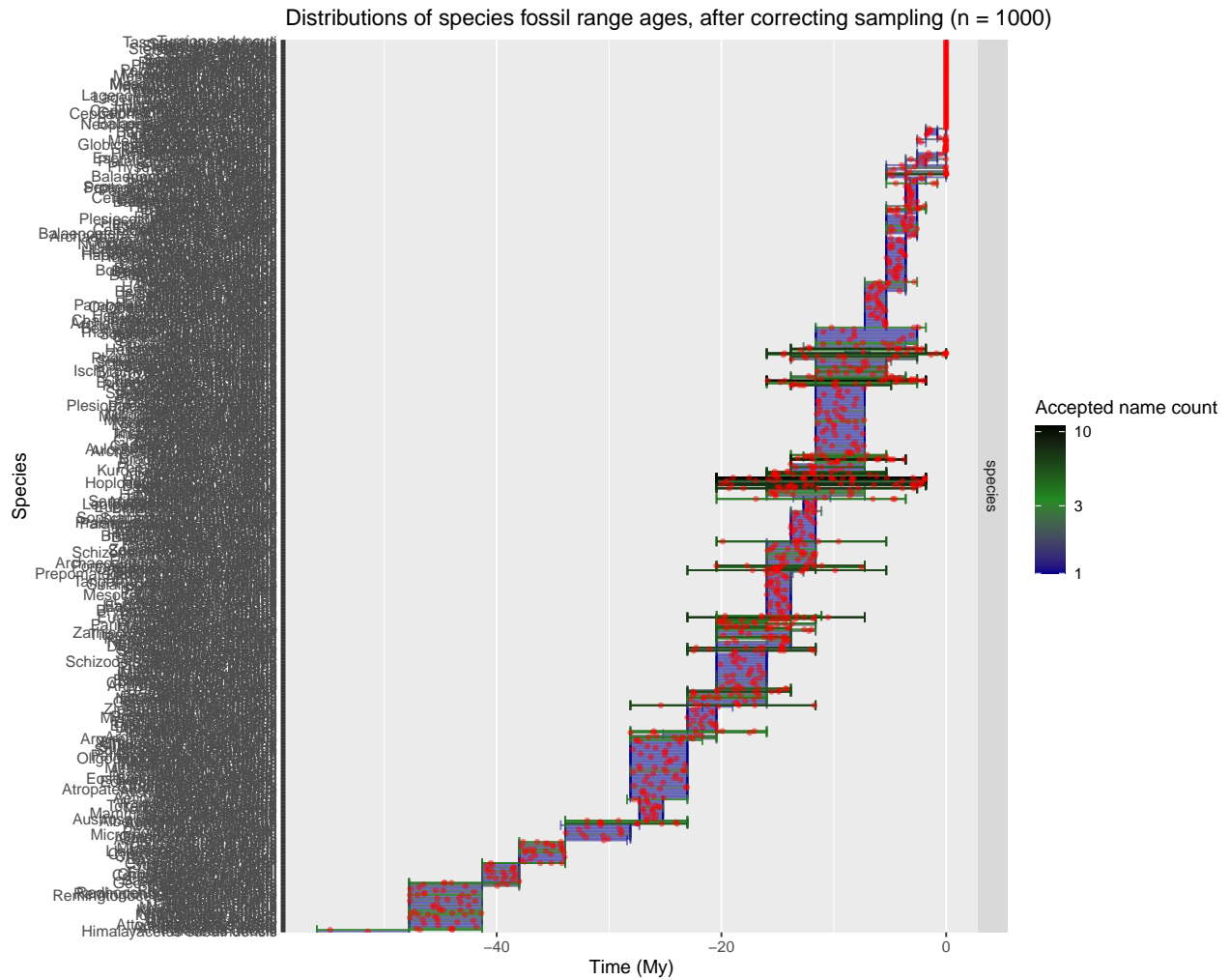


Impact of subsampling on occurrences repartition (species only)

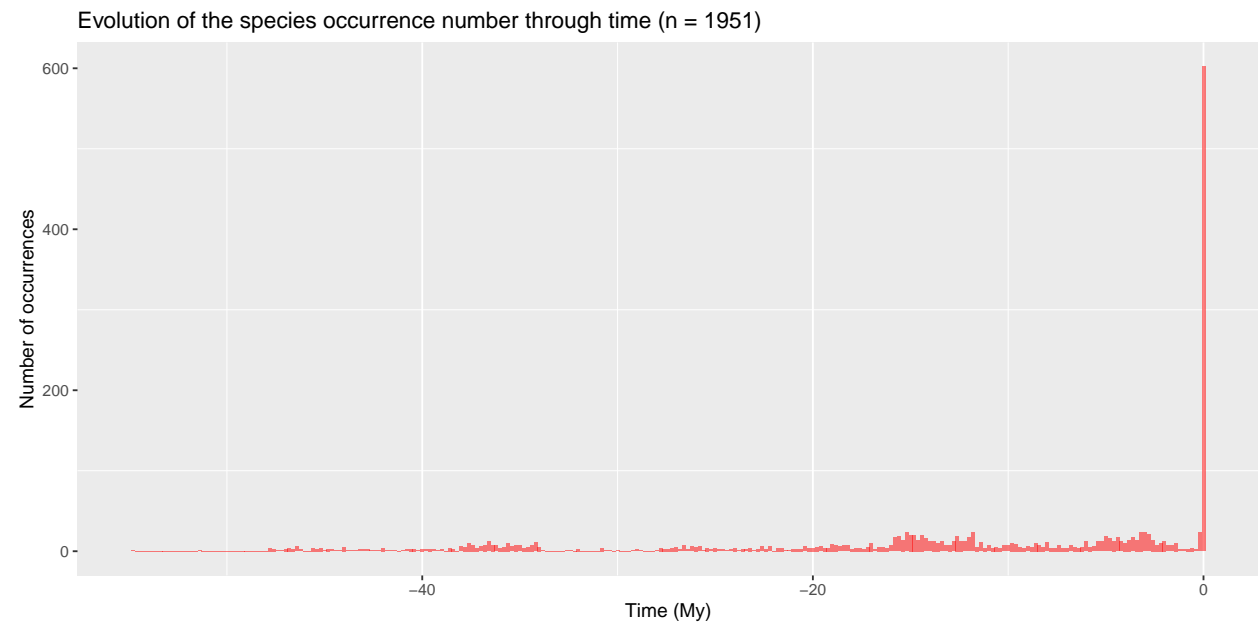
See what our distributions

Distributions of species fossil range ages, before correcting sampling (n = 1951)

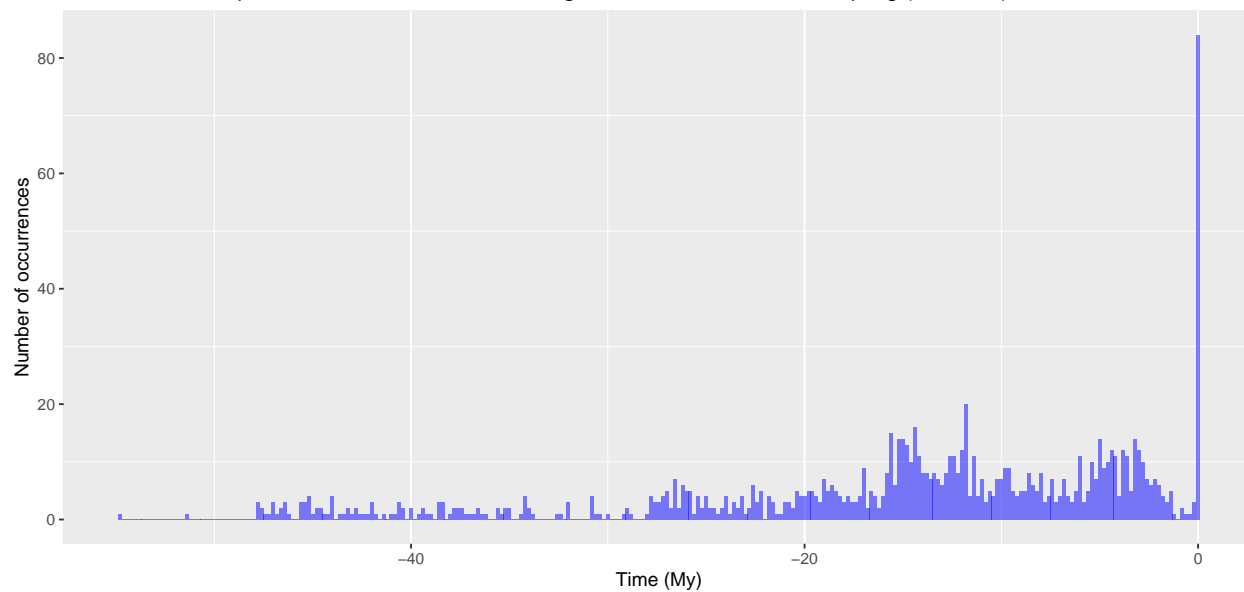




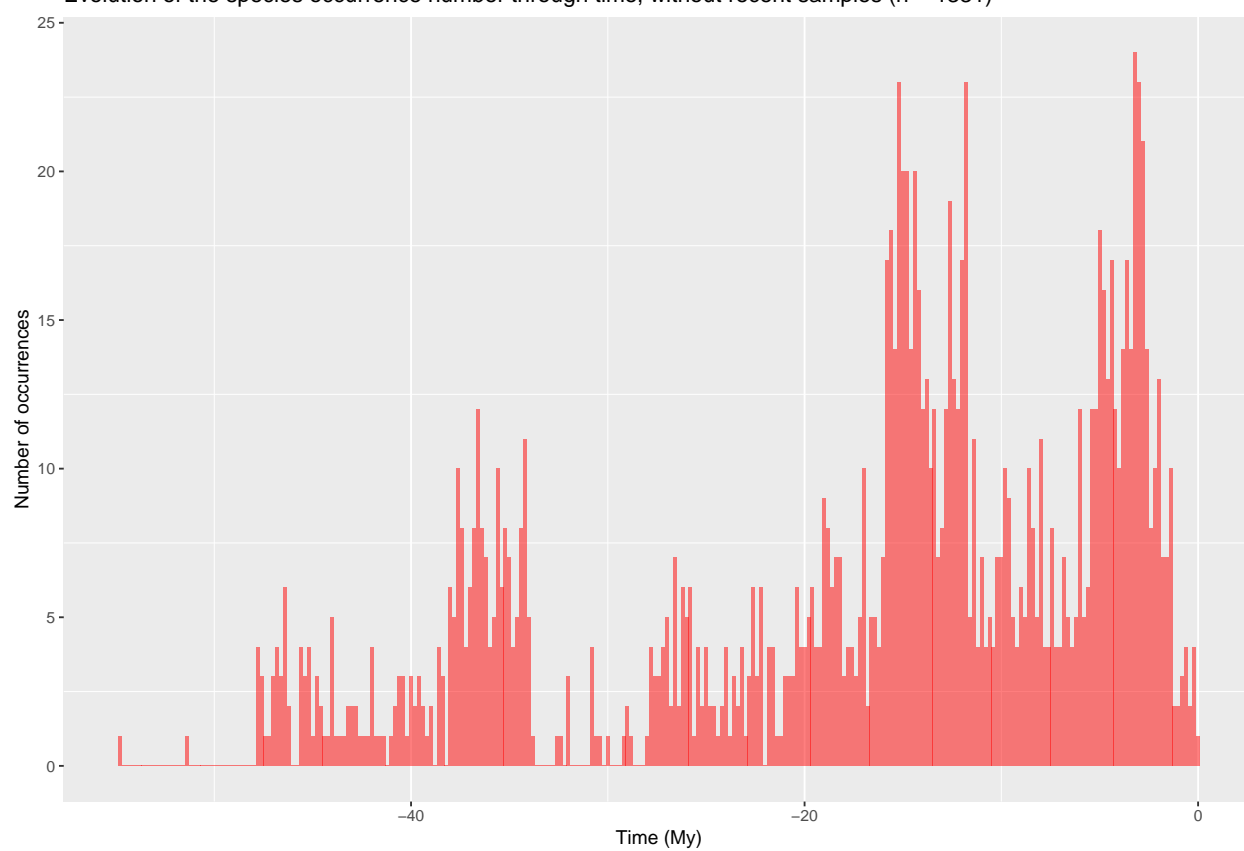
→ Some highly dense cluster became much more similar to the others.

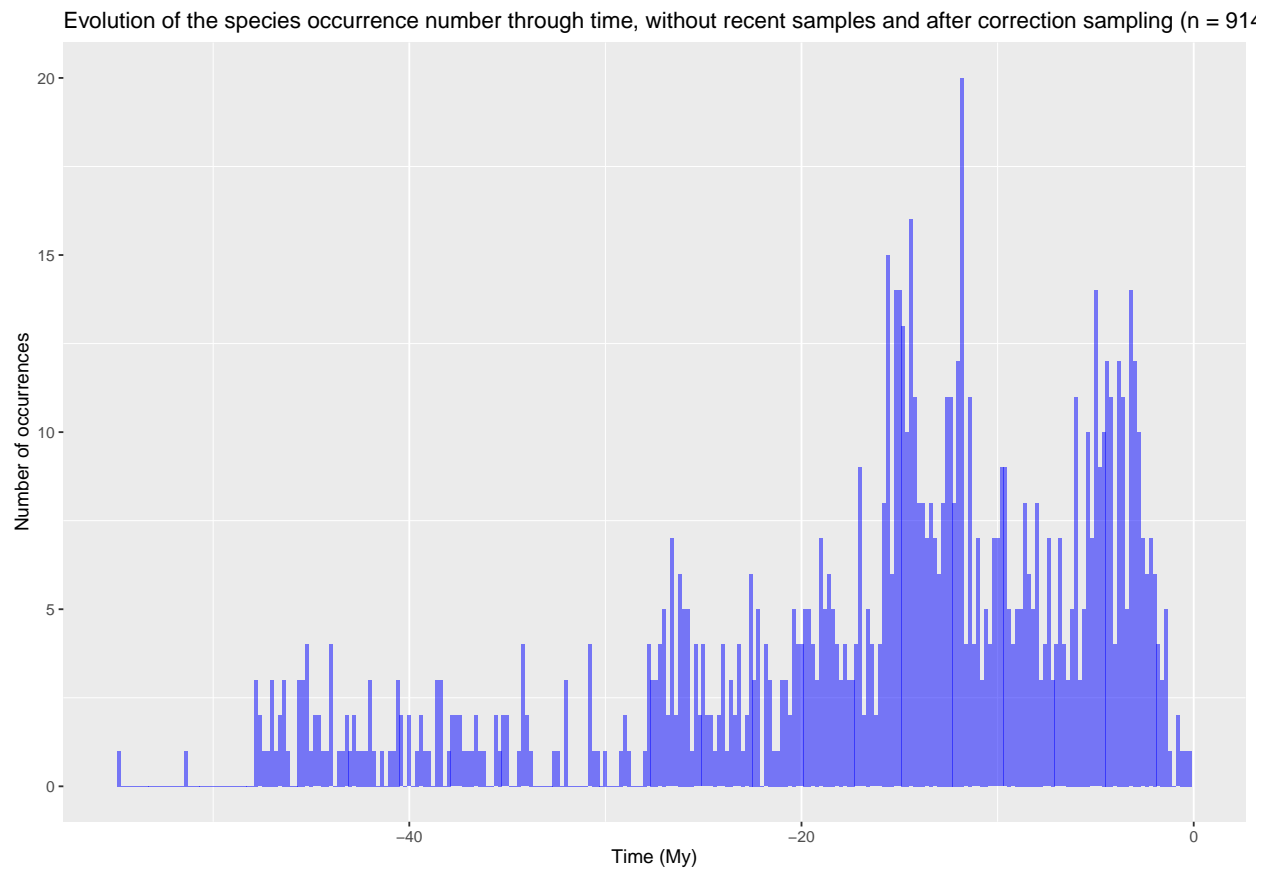


Evolution of the species occurrence number through time, after correction sampling (n = 1000)

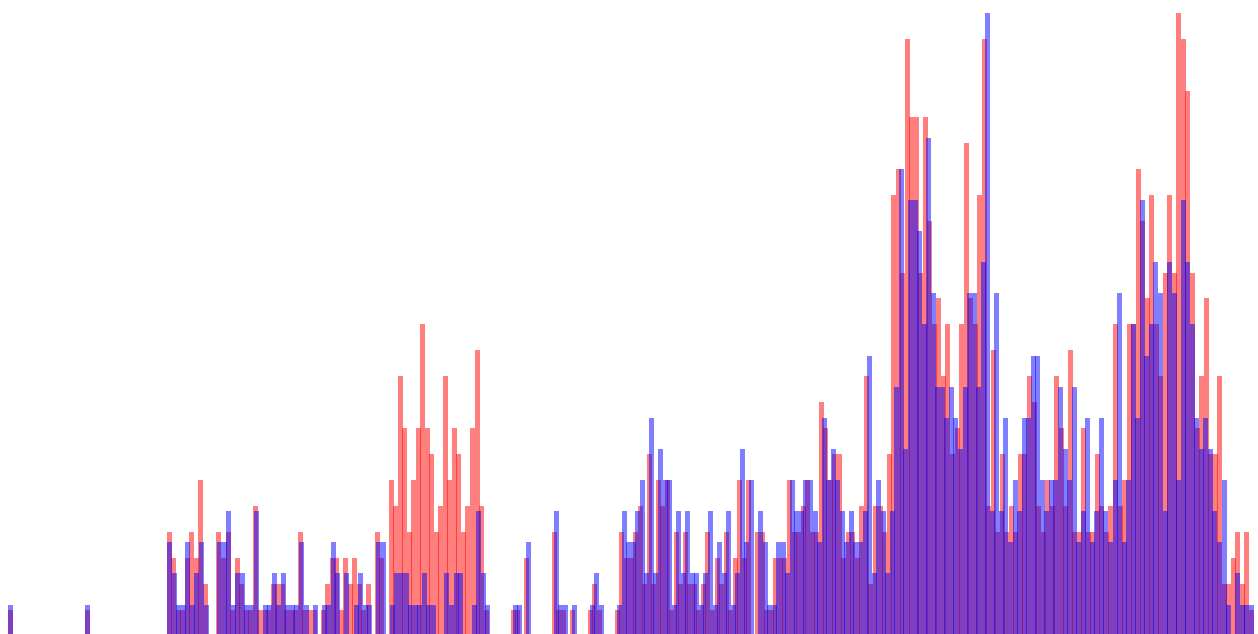


Evolution of the species occurrence number through time, without recent samples (n = 1331)





If we superpose these 2 plots :



→ We get the new species occurrence repartition after subsampling correction, that could be used for doing inference with the occurrence birth-death model.

Conclusions

Achievements :

- It seems possible to adequately reduce the abundance bias by subsampling the most concentrated intervals → **species only**
- Using combined ranges by species appears to be more robust → **to be confirmed**
- Very recent samples may have been dated with a more precise method and contain much more fossils, so they should be removed or treated separately → **additional information needed**

Open questions :

- What about other accepted ranks ?
 1. The problem is that differences in the number of occurrences at higher ranks could be due to differences in individual abundances inside species or due to differences in the number of species inside that group.
 2. A solution could be to look at the number of species by group based on the indicated species, and include it in the bias correction : homogeneizing the *number of occurrences / time unit / number of species in the group* → **additional data required** (ranks classification)
- Why do most occurrences miss a late stratigraphic limit ?
- Some occurrences have very huge time intervals → **Was is a good idea to remove those >10My ? Should we remove more of them (>5My) ?**