Primate sympatry shapes the evolution of their brain architecture

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Abstract | The main hypotheses on the evolution of animal cognition emphasise the role of conspecifics in affecting the socio-ecological environment shaping cognition. Yet, space is often simultaneously occupied by multiple species from the same ecological guild. These sympatric species can compete for food, which may thereby stimulate or hamper cognition. Considering brain size as a proxy for cognition, we tested whether species sympatry impacted the evolution of cognition in frugivorous primates. We first retraced the evolutionary history of sympatry between frugivorous primate lineages. We then fitted phylogenetic models of the evolution of the size of several brain areas in frugivorous primates, considering or not species sympatry. We found that the whole brain or brain areas used in immediate information processing were best fitted by models not considering sympatry. By contrast, models considering species sympatry best predicted the evolution of brain areas related to a long-term memory of interactions with the social or ecological environment, with a decrease of their size the higher the sympatry. We speculate that species sympatry, by generating intense food depletion, leads to an over-complexification of resource spatio-temporality that may counteract the benefits of high cognitive abilities, drives niche partitioning and specialisation, and thereby induces lower brain area sizes. In addition, we reported

28   that primate species in sympatry diversify more slowly. This comparative study suggests that

29   species sympatry significantly contributes to shaping primate cognition and diversification.

30   **Short title:** Sympatry shapes primates' brain architecture

31   **Keywords:** Brain size - Cognition - Diversification - Frugivory - Primates - Sympatry

INTRODUCTION

Cognition evolution is shaped by the balance between socio-ecological drivers promoting cognitive abilities (González-Forero and Gardner 2018) and physiological and energetic constraints limiting them (Navarrete, Schaik, and Isler 2011). Primates are pivotal species for cognitive studies (Byrne 2000) because their cognition is thought to be promoted by interactions of individuals with conspecifics within the social unit (Byrne 2018; Dunbar and Shultz 2017), among generations (Wilson 1991; Whiten and Schaik 2007; Reader and Laland 2002; Herrmann et al. 2007; Tomasello 2019; Schaik and Burkart 2011), between social units (Ashton, Kennedy, and Radford 2020), or with the rest of their environment (Clutton-Brock and Harvey 1980; Milton 1981; Rosati 2017). However, space is often occupied by many primate species, including species sharing the same diet. Because of direct and indirect interactions linked to resource availability, the presence of heterospecifics is also likely to shape the evolution of species cognition.

Retracing the evolutionary history of cognitive abilities proves to be challenging because there is still no consensual measurement for cognition applicable across all species. Up to now, a raw approximation consists in considering the (relative) brain size as a proxy for cognitive abilities, with larger sizes considered equivalent to more advanced cognitive abilities (mammalian carnivores: Benson-Amram et al. 2016; primates: MacLean et al. 2014). Although the relevance of this assumption is heavily limited within species, in part because of plasticity (Gonda, Herczeg, and Merilä 2013), this holds true when comparing different species (e.g. in primates, Reader and Laland 2002). In addition, instead of considering the brain as a whole, the multifaceted aspect of animal cognition is then more precisely depicted by appreciating the mosaic nature of the brain (Barton and Harvey 2000). For instance, variations in the size of some specific brain areas have been robustly associated with variations in cognition related to the functions of these areas (Healy and

Rowe 2007). The brain is therefore a patchwork of areas cognitively specialised that may follow

different evolutionary trajectories.

Because the coexistence of primate species in the same biogeographic area (henceforth referred to

as sympatry) can affect multiple aspects of their social-ecological environment, brain areas may be

affected differently by sympatry. Considering a simplistic functional picture of the primate brain,

one can hypothesise how sympatry might, through direct or indirect competition or cooperation,

influence the relative benefits of cognition (i.e. the balance between its benefits and its costs).

First, sympatric primate species may have dietary overlaps and compete for the same food

resources (Kamilar and Ledogar 2011). Broad dietary guilds (frugivores, folivores, nectarivores,

granivores, insectivores, etc.) are generally considered based on the main type of food they

consume. Having several sympatric species competing for the same food resource often leads to an

increase in food depletion of the shared resource compared with an environment with only one

foraging species (Minot 1981). Such depletion may therefore complexify the pattern of resource

distribution and availability in space and time. This complexification may in turn affect the selective

pressures upon brain areas involved in the storing of spatio-temporal information, such as the

hippocampus (Burgess, Maguire, and O'Keefe 2002, Hypothesis 1: the long term memory is affected

by sympatry). This complexification may also occur when the dietary overlap is inexistent, as

foragers may rely on phenological cross-correlations between plant species (e.g. if plant species A

produces fruits earlier in the season than plant species B, thus the availability of B fruits can be

predicted based on the availability of A fruits, even though A is not consumed by the forager, Robira

et al. 2021). Food depletion may thus weaken these cross-correlations and make food availability

predictions less reliable.

Second, sympatric species may enrich the landscape of visual, olfactory or acoustic cues usable to

locate available food (e.g. Avarguès-Weber, Dawson, and Chittka 2013; Kashetsky, Avgar, and Dukas

2021). Consequently, it may impact the selective pressures upon brain areas involved in processing

more immediate sensory information, such as the Main Olfactory Bulb (MOB), the cerebellum

(Koziol et al. 2014; Sokolov, Miall, and Ivry 2017), and the neocortex (Wiltgen et al. 2004)

(Hypothesis 2: cue processing is affected by sympatry).

Third, besides indirect interactions through foraging, cognition can also be triggered by direct

"social" interactions with other individuals (Byrne 2018; Dunbar and Shultz 2017). The striatum, a

brain area stimulated during social interactions (Báez-Mendoza and Schultz 2013), or to a lesser

extent, the hippocampus (Todorov et al. 2019), may therefore be affected by the increase of direct

social interactions between heterospecifics (Hypothesis 3: sociality is affected by sympatry).

Under these (non-exclusive) hypotheses, sympatry could stimulate or hamper cognition evolution

by affecting the relative benefits that cognition provides. Indeed, memory stands as a valuable tool

to infer food availability and location when food is rare and ephemeral but predictable (Milton

1981; Rosati 2017), but is energetically costly. On the one hand, having a better memory should be

advantageous under reasonable food depletion. In addition, competition for the shared resource

between species should promote anticipatory behaviour, hence high cognition, as expected for

within-species competition (Ashton, Kennedy, and Radford 2020). In this case, the size of the

hippocampus (reflecting long-term memory abilities) should be larger the higher the sympatry

intensity (Prediction 1.1). On the other hand, intense depletion also increases environmental

unpredictability. In the case of a frugivore searching for fruit, for instance, the perceived synchrony

in fruit production between trees of the same or different species, used to infer food availability

(Janmaat et al. 2012), can be lowered by depletion, therefore limiting the benefits of memory

(Robira et al. 2021). Due to the energy constraints of maintaining a large brain, the hippocampus

size could thus be smaller in species experiencing high levels of sympatry (Prediction 1.2).

Furthermore, the competition between sympatric species from a given dietary guild may be lowered

by specialising in different food resources, i.e. niche partitioning. While a species might specialise in

food that is difficult to access and requires high cognition (e.g. through the use of tools),

specialisation is generally associated with reduced flexibility and thus lower cognitive abilities

(Henke-von der Malsburg, Kappeler, and Fichtel 2020). A strong association between niche

partitioning and cognitive specialisation has been previously raised in primates (Aristide et al.

2016). From this perspective, a high level of sympatry could be enabled by high cognitive

specialisation toward specific food types, requiring less long-term memory abilities, and could also

result in smaller hippocampus sizes in highly (specialised) sympatric species.

Meanwhile, cues left out by heterospecifics and usable to locate available food might also add to

environmental ones already available. Hence, sympatry could be associated with larger sizes of the

MOB, the cerebellum, or the neocortex (Prediction 2).

Finally, an increase in direct interactions between species, such as with the formation of mixed-

group species (Goodale et al. 2010), should imply an upsurge of social stimuli leading to a larger size

of the striatum or hippocampus in sympatry (Prediction 3).

In any case, the evolution of brain size in primates likely impacted their dynamic of species

diversification. Larger brain sizes are indeed found to be associated with higher diversification rates

in birds (Sayol et al. 2019) and similar patterns have been suggested in primates (Melchionna et al.

2020). However, it remains unclear how brain size and diversification are interlinked in the context

of sympatry.

Here, we investigated whether species sympatry affected the evolution of cognition using

frugivorous primates as a study example. Frugivorous primates are an interesting group for such a

question because fruit is the archetype of a hard-to-find resource yet predictable (Janmaat et al.

2016), for which cognition considerably shapes the foraging strategy (Trapanese et al. 2019). To

infer the effect of species sympatry on cognition in frugivorous primates, we evaluated the support

for phylogenetic models of brain size evolution accounting or not for species sympatry, and

investigated the directionality of the selection induced by sympatry on brain size evolution. Finally,

we tested for correlative patterns between brain size or current sympatry and species

diversification in all primates, to better understand the impact of cognition and interactions
between primates on their evolutionary success.

METHODS

133 Data processing, analyses, and plots were computed with R software (v.4.1.2, R Core Team 2020).

Data Collection

Phylogeny

We used chronogram trees of the primate taxon of the 10kTrees project (downloaded on May 2021,
version 3), as well as a consensus tree of 1000 trees for the subsequent phylogenetic analyses. The
trees contain 301 primate species (Figure 2). Note that in all these analyses, we discarded *Homo*
sapiens and *Macaca sylvanus*. The latter was discarded because of its complete geographic isolation
and repeated intervention of humans in population maintenance (Modolo, Salzburger, and Martin
2005). A summary of available data per species is presented in Supplementary Figure S4.

Trait data

Data were pooled from previous literature surveys (see Supplementary Material "Data availability").
Brain data were obtained from DeCasien and Higham (2019) for the whole brain and all other areas
mentioned (cerebellum, hippocampus, Main Olfactory Bulb (MOB), neocortex, striatum) from
Powell, Isler, and Barton (2017) and from Powell, Barton, and Street (2019) for the whole brain,
cerebellum and neocortex size, from Todorov et al. (2019) for hippocampus, and neocortex size,
from Grueter (2015) for the whole brain size, and from Navarrete et al. (2018) for the whole brain,
cerebellum, hippocampus, and striatum size. They were freely available in the main manuscript or
supplementary material. For each primate species, the percentage of frugivory and/or folivory was
obtained based on a freely available dataset from DeCasien, Williams, and Higham (2017), from

Powell, Isler, and Barton (2017), and from Willems, Hellriegel, and Schaik (2013). The availability of

trait and distribution range for the 301 primate species is depicted in Supplementary Figure S4.

From the global endocranial brain volume, we obtained the Encephalization Quotient (EQ, $N_{EQ,max}$ =

182) as follows (DeCasien, Williams, and Higham 2017)

$$EQ = 1.036 \times \text{Brain volume}/(0.085 \times \text{Body mass}^{0.775})$$

with the brain volume in cm^3, 1.036 g/cm^3 being the assumed homogeneous brain density, and the

body mass in g. EQ indicates whether the brain size ranges above (> 1) or below (< 1) what is

expected given the body mass. Body mass was obtained from DeCasien, Williams, and Higham

(2017), from Powell, Isler, and Barton (2017), from Grueter (2015), and from Pearce et al. (2013).

The sub-parts of the brain were chosen because they were involved in immediate sensory

information processing (MOB, $N_{MOB,max}$ = 39), in movement and/or general information processing

and retention (neocortex, $N_{Neocortex,max}$ = 69, Wiltgen et al. 2004; cerebellum, $N_{Cerebellum,max}$ = 70,

Koziol et al. 2014; Sokolov, Miall, and Ivry 2017), or short-term working memory and long-term

spatio-temporal memory (hippocampus, $N_{Hippocampus,max}$ = 63, Burgess, Maguire, and O'Keefe

2002). The striatum ($N_{Striatum,max}$ = 63) supports information processing during social interaction,

reward assessment, planning or goal-oriented behaviours (Báez-Mendoza and Schultz 2013;

Johnson, Meer, and Redish 2007). To investigate their evolutionary history, we first used the ratio

between their volume and body mass to maximize comparability. As such, the use of specific area

sizes relative to the body mass and not raw sizes depicts the evolution of cognitive abilities in terms

of allocation rather than abilities per se (but see discussion in Deaner, Nunn, and Schaik 2000).

Second, we repeated the analyses considering the ratio between the brain area volume and the

whole brain volume, as this might reflect within brain energy reallocation (i.e. mosaic evolution of

the brain).

Ranging Data

The current biogeographic range of each primate species was assessed using ranging maps

provided by the IUCN red list (IUCN 2021). Ranging data were available for 249 species among the

301 represented in the 10kTrees primate phylogeny.

Retracing past sympatry between primate species

Based on the biogeographic distribution of each extant primate species, we first reconstructed the

history of past sympatry between primate lineages. To do so, we followed Drury et al. (2018) and

first reconstructed the biogeographic history of each primate lineage to then retrace which pairs of

primate lineages were likely to be simultaneously present at the same place. Leaning on Kamilar

(2009), we considered that the biogeography of primates can be described by 12 discrete

biogeographic areas with highly similar community structures shaped by both the environment

geography and climatic correlates. These geographic areas, mapped using Google earth professional

(v7.3.3), are represented in Figure 1. One to multiple biogeographic areas were assigned to each

species as soon as 10% of their current distribution range overlapped on the surface with a given

biogeographic area. We also replicated these biogeographic assignations by using instead a larger

threshold of 30%. This upper threshold was chosen because, as such, a species could initially occupy

a maximum of three areas, thus at best would occupy one continent at present (Figure 1). Overlaps

of primate current ranges with biogeographic areas were calculated with the "gIntersection"

function from the *rgeos* package (Bivand and Rundel 2021) applied to Mercator-projected data to

get the overlapping contour, and the "area" function from the *geosphere* package (Hijmans 2021),

applied directly on unprojected longitudinal-latitudinal data for area size calculation.

Given these 12 biogeographic areas, we retraced the biogeographic history of primates with the

BioGeoBEARS package (Matzke 2013), using the biogeographic stochastic mapping algorithm

(Matzke 2016). We fitted non-time-stratified dispersal-extinction-cladogenesis (DEC) models

specifically suiting analyses of range data since it accounts for spatially explicit processes of

cladogenetic and anagenetic events (see Matzke (2013) for further details on these events). We

fixed to three biogeographic areas the maximum number of areas that a lineage can simultaneously

occupy since it offers the possibility to occupy a complete mainland continent while keeping

computational time reasonable. DEC models were independently fitted when considering either a

10% or a 30% threshold of range overlaps. Finally, to account for the uncertainty in biogeographic

reconstructions, we sampled 10 histories of primate biogeographic ranges. We assumed that two

primate lineages were in sympatry at a given time whenever two lineages simultaneously occupied

the same biogeographic area.

Inferring past diets of primate lineages

Next, we retraced the evolutionary history of frugivorous lineages in primates. We first classified

extant species as either "frugivorous" or "folivorous" based on the availability of percentage of

frugivory and folivory, prioritizing frugivory over folivory. A species was classified as frugivorous if

the percentage of frugivory was at least above 20%. If this was not the case, or if the percentage of

frugivory was unavailable, a species could be classified as folivorous if the percentage of folivory

was at least above 40%. Otherwise, DeCasien, Williams, and Higham (2017) gave a binary

classification of diet, species being categorised as frugivorous or folivorous, partly based on

anatomical criteria. Whenever the percentages were not available, we referred to this classification.

In any other cases (e.g. species feeding on invertebrates or tree exudates), the species were

discarded. We also replicated these diet assignments by considering a threshold of 40% for

frugivory and 60% for folivory.

Second, considering diet as a binary variable (frugivory versus folivory), we retraced the

evolutionary history of such discrete traits based on a continuous Markovian process (extended Mk

models) using Bayesian inference (Bollback 2006), with the "simmap" function of the *phytools*

package (Revell 2012) and internally setting up the prior probability of trait, but with no prior on

the transition matrix. Ancestral diet reconstructions were performed using both combinations of

dietary thresholds (20/40% and 40/60% for frugivory and folivory, respectively). To account for

the uncertainty in the reconstructions, we sampled 10 stochastic mapping of ancestral diet

reconstructions. These ancestral reconstructions were used in combination with the histories of

primate biogeographic ranges to assess whether a pair of frugivorous species were in sympatry or

not (i.e. we obtained reconstructions of the evolutionary history of sympatry between frugivorous

primate lineages).

Phylogenetic models

We assessed the effect of sympatry on primate brain evolution using two approaches. First, we used

phylogenetic models of trait evolution to assess the role of sympatry in the evolution of brain size.

Second, we investigated how sympatry has influenced brain size evolution (i.e. selection towards

smaller or larger brain sizes) by evaluating correlations between current levels of sympatry and

brain sizes, using linear models independently for each brain area. Although it does not explicitly

consider the potential independence between brain areas, we modelled independently each brain

area to preserve the maximum power (i.e. the largest sample size) in each analysis. Finally, we also

checked for correlative patterns between primate diversification rate and brain size or levels of

sympatry to better understand the impact of cognition and interactions between primates on their

evolutionary success.

Phylogenetic models of trait evolution: does species sympatry shape brain size evolution?

(a)   Fitting models of trait evolution

We restricted the analyses to frugivorous species to test whether species sympatry has impacted

the evolution of cognition, depicted either by the whole brain (using the Encephalization Quotient,

EQ), or the size of the aforementioned specific brain areas relative (i) to the whole body mass

(Figure 3) or (ii) to the whole brain size. Models of trait evolution testing the effect of species

sympatry on one brain area relative to the whole body mass (models (i)) test whether the total

allocation towards specific brain area is affected by species sympatry. In contrast, models of trait

evolution testing the effect of species sympatry on one brain area relative to the whole brain size

(models (ii)) test whether the within-brain energy reallocation (i.e. mosaic evolution) is affected by

species sympatry.

Considering the size of the different brain areas, we independently fitted phylogenetic models of

trait evolution (Drury et al. 2016). For models accounting for species sympatry, following Drury et

al. (2016), we used our ancestral reconstructions of sympatry between frugivorous lineages. In

practice, we obtained a series of interaction matrices (i.e. lines and columns correspond to

frugivorous species, and each cell indicates whether a given species pair is in sympatry (value of 1)

or not (value of 0)), along the phylogenetic tree of frugivorous primates (see Drury et al. 2016). We

fitted three models of brain size evolution accounting for species sympatry: the matching

competition (MC) model (Nuismer and Harmon 2015) and density-dependent models (DD$_{lin}$ and

DD$_{exp}$, Drury et al. 2016). Specifically, these models expand classical Brownian motion (BM) models

of stochastic evolution, by including an additional variable related to the current brain size of

sympatric species (MC), or by considering density-dependent evolutionary rate (DD models). The

MC model considers the repulsion of traits of sympatric lineages from the same dietary guild due to

competition (character displacement), that is $z_i(t + dt) = z_i(t) + S(\mu(t) - z_i(t))dt + \sigma dB_i$ where z

is the brain size of a species i at time t, μ is the mean value of the trait of sympatric species, S

reflects the strength of the effect of species sympatry and σdB_i is the drift with a constant

evolutionary rate σ (Drury et al. 2016). Here, S is constrained to be negative, which means that

sympatric species would tend to divergently evolve either lower, or higher, EQ or relative brain size.

Linear (DD_{lin}) or exponential (DD_{exp}) density-dependence (Drury et al. 2016; Weir and Mursleen

2013) assumes that the evolutionary rate, σ, of trait change, varies either positively or negatively as

a function f of the number of frugivorous sympatric lineages, such as

$$\sigma_l = f_{lin}(l) = \sigma_0 + rl$$

$$\sigma_l = f_{exp}(l) = \sigma_0 \exp(rl)$$

where σ_0 corresponds to the value of the initial ancestor, l indicates the number of lineages at a

given time, r allows for modelling the speed and direction of the dependency to lineage number

($r > 0$ leads to an increase of trait changes, while $r < 0$ leads to a decline of the trait changes). In

particular, DD models have been extensively used for testing hypotheses linked with adaptive

radiations ($r < 0$), yet, by their mathematic constructions, they can accomodate much broader

scenarios of trait evolution. We fitted models considering species sympatry using the "fit_t_comp"

function from the *RPANDA* package (Morlon et al. 2016).

Depending on the brain area and the frugivory threshold we considered, the models were fitted on

different sample sizes: EQ: 148 to 182, striatum: 56 to 63, MOB: 34 to 39, neocortex: 61 to 69,

hippocampus: 56 to 63, cerebellum: 62 to 70 frugivorous species. For a given set of models

(i.e. within a brain area), the sample was strictly identical, allowing within-set model comparisons.

Prior to fitting, trait parameters were log-transformed to reach more symmetrical distributions.

We compared the support of models considering species sympatry to the support of simpler

models assuming no effect of species sympatry on the evolution of brain sizes: the Brownian motion

(BM), the Ornstein-Uhlenbeck process (OU, a model with an optimum value, see Blomberg,

Rathnayake, and Moreau (2020) for a review), or the Early-Burst model (EB), for assessing a time-

dependence of the evolutionary rate, irrespectively of the intensity of species sympatry (Blomberg,

Garland, and Ives 2003). These models not accounting for species sympatry were fitted using the

"fitContinuous" function from the *geiger* package (Slater et al. 2012; Pennell et al. 2014). All these

models were repeated 10 times, using 10 different combinations for the ancestral reconstruction of

primate biogeography and diet. They were then compared within an information-theoretic

framework (Burnham and Anderson 2002), based on the weights of Akaike Information Criteria

corrected for small samples (AICc) when considering all six models (MC, DD_{lin}, DD_{exp}, BM, OU, EB).

The model weight depicts how well the model fits the observed data compared with the other tested

models.

Each model represents the average of 10 (uncertainty on diet/ranging evolution) x 10 (uncertainty

in brain/diet rate data) x 2 (geographic overlap threshold) x 2 (frugivory threshold) x 2 (folivory

threshold) = 800 sub-models. We stopped computations when the calculation of the likelihood was

excessively long (> 1 week). The final sample size thus was 730 models.

(b)   Determining the effect of sympatry on brain sizes

If diversity-dependent models of traits evolution considering species sympatry inform whether

species sympatry has impacted the evolution of the brain size by increasing or decreasing the tempo

of trait evolution, they are not informative on the directionality of the effect (i.e. are brain sizes in

frugivorous sympatric primates increasing or decreasing?). To determine whether species sympatry

positively or negatively affected the sizes of brain areas, we independently fitted Gaussian Pagel's

lambda phylogenetic regressions for each brain area of extant frugivorous species. This model is a

derivative of the BM model, where the phylogenetic variance-covariance matrix has all coefficients,

but its diagonal ones, multiplied by lambda: it thus relaxes the hypothesis of BM since we included

brain areas for which the evolutionary history was best described by models considering sympatry

(see Results). To fit these models, we used a frequentist-based approach with the "phylolm" function

from the *phylolm* package (Ho and Ane 2014). We considered the least stringent frugivory

assessment, with the frugivory threshold fixed at 20% and the folivory threshold fixed at 40%. If

due to data variability, a species did not robustly fit into the categorical classification "frugivorous

versus folivorous" (i.e. could be either of the two), it was considered as frugivorous nonetheless.

Again, insectivorous species, species feeding on tree exudates, etc., were not considered.

The response variable was the relative size of each brain area (relative to the whole body mass or

to the whole brain size). Due to data variability, we took the mean of the possible values given the

different datasets and assessed the sensitivity using non-averaged values (see Supplementary

Material "Phylogenetic regressions: results, stability, and assumption"). In this model, we used as

covariates (i.e. continuous predictors) two explicit measures of the level of species sympatry for

each extant frugivorous species: (1) the number of frugivorous sympatric species (square-rooted to

reach symmetrical distribution) and (2) the average percentage of overlapping current range

(assessed based on IUCN data) with other sympatric frugivorous species. For a given species A,

sympatry with another species B was considered when at least 10% of the range of species A

overlaps with the range of species B. This was done to reduce the noise induced by coarse

identification of species range.

To sum up, when assessing the interplay between species sympatry and the evolution of frugivorous

primates' brain architecture, we considered sympatry under different forms. To assess whether it

affected brain size evolution, sympatry was added to classical phylogenetic models of trait evolution

as an additional variable depicting the mean trait value of sympatric species (MC models), or as a

density-dependent term (i.e. the total number of sympatric lineages at a given time; in DD_{lin} and

DD_{exp} models). Then, to assess the directionality of the effect of sympatry on brain sizes, sympatry

was used as a tested predictor in phylogenetic linear regressions, under two forms: the number of

currently sympatric species, and the average range overlap with currently sympatric species.

Body mass and sympatry

To control for a confounding effect due to a relationship between sympatry and body mass, as body mass was used to obtain relative brain size, and is associated with the whole brain volume (Smaers et al. 2021), we repeated all model fitting (models of trait evolution, and PGLS) with body mass as the output variable (see Supplementary Material, Does sympatry shape body mass evolutionary history?).

Models of species diversification

Next, to investigate whether cognition and/or species sympatry have affected primate diversification, we inferred how primates diversified over time and across lineages. Lineage-specific net diversification rates (defined as speciation minus extinction rates) were estimated using an updated version of the *ClaDS* algorithm (Maliet, Hartig, and Morlon 2019) based on data augmentation techniques (Maliet and Morlon 2021). Particularly, we used *ClaDS2*, the model with constant turnover (i.e. constant ratio between extinction and speciation rates; see Supplementary Material "Primate diversification rate over time" for further explanations). We extracted the mean diversification rates through time and the lineage-specific diversification rate of each extant species.

We also fitted Gaussian Pagel's lambda phylogenetic regressions of the different relative brain sizes against the net diversification rates. Because assumptions for a frequentist-based approach were unmet, we performed Bayesian-based inferences instead. We used the "MCMCglmm" function of the *MCMCglmm* package (Hadfield 2010). Each chain had a burn-in period of 5000 iterations, a total length of 50 000 iterations, and was sampled every 50 iterations. We used the least informative priors. Fixed priors were set to default values (Gaussian distribution of mean 0 and variance 10^8). Again, we took the mean of the brain trait values for the main model and assessed the sensitivity by re-running the model several times using non-averaged values.

To determine whether species sympatry was associated with lower or larger diversification rates,

we fitted frequentist-based Gaussian Pagel's lambda phylogenetic regressions with the lineage-

specific diversification rate as the output variable, and used the two metrics for describing sympatry

(the number of frugivorous sympatric species and the average percentage of overlapping range with

other sympatric frugivorous species) as the tested variables, as in (a).

Model implementation and stability

Details on the implementation, stability, and uncertainty of phylogenetic regressions are provided in

Supplementary Material (see "Phylogenetic regressions: results, stability, and assumption"). Prior to

fitting, covariates were transformed so as to reach more symmetrical distributions when adequate.

Necessary assumptions on the normal distribution of residuals and homoscedasticity were visually

assessed and pointed out no violation (see Supplementary Material, Model assumptions). In

addition, we did not observe correlation issues among predictors (Variance Inflation Factor, VIF_{max}

< 2, Mundry 2014).

RESULTS

The database we gathered contained between 34 to 182 frugivorous primate species (depending on

the brain area considered). When pondering by whole body mass, we observed ample variations in

the sizes of the studied brain areas. For instance, the lemuriformes, which are known to prioritize

smell compared with other primate species, have the largest relative MOB size (lemuriformes: mean

$\pm$ SE = 0.23 $\pm$ 0.07, other: 0.12 $\pm$ 0.04, 3). Similarly, the highest relative sizes of the striatum were

found in platyrrhini (platyrrhini: mean $\pm$ SE = 0.91 $\pm$ 0.07, other: 0.59 $\pm$ 0.07, 3), which are known

to form poly-specific associations (callitrichine in particular, Heymann and Buchanan-Smith 2000).

In terms of the measures of sympatry, we observed that on average ($\pm$ SE), the considered primate

species had 52% ($\pm$ 2) of their range overlapping with other species. That ranged from 0% of

overlap (*Macaca nigra*), to 100% of overlap (*Cercopithecus pogonias, Alouatta pigra, Loris*

tardigradus, Hylobates moloch, Cercocebus galeritus, Presbytis melalophos, Semnopithecus entellus).

In terms of the distribution range, the considered primate species co-occurred on average with 6.38

($\pm$ 0.39) other primate species, ranging from 0 other species to 21.

To retrace past species sympatry between frugivorous lineages, we first reconstructed primate

biogeographic history when considering 12 biogeographic areas (Figure 1, Kamilar 2009) and their

diet evolution. We found that the ancestors of primates were likely to be frugivorous and that

folivory evolved more than 5 times, in particular in cercopithecidae and in memuridae. Estimated

diet history was consistent with fossil evidence (see Supplementary Material, Diet reconstruction,

Figure S1). We then modelled the evolution of the size of the whole brain (EQ), or regionalised areas

(neocortex, cerebellum, MOB, hippocampus, and striatum) when considering species sympatry or

not. Weighting each brain area by the whole body mass (that gives insight into energy allocation

optimisation) and weighting by whole brain size (more likely reflects the mosaic evolution of the

brain) yielded similar results (see Supplementary Material, Weighting the size of brain areas by

whole brain size), therefore we only present the case when weighting by body mass. We found that

models not considering species sympatry best described the evolutionary history of the EQ, the

neocortex, and the cerebellum (Figures 3 and 4), two areas specifically involved in immediate

sensory information processing (Wiltgen et al. 2004; Koziol et al. 2014; Sokolov, Miall, and Ivry

2017), and also in memory consolidation for the neocortex (Wiltgen et al. 2004). The fact that these

biggest areas are best described by the Ornstein-Uhlenbeck process suggests a stabilisation towards

an optimal size, which may illustrate the trade-off between costs and benefits of brain development

(Isler and Schaik 2009). By contrast, density-dependent models considering species sympatry

(DD_{lin} and DD_{exp}) were best supported in the foraging-related and social-related areas respectively:

the hippocampus, specialised in spatio-temporal memory (Burgess, Maguire, and O'Keefe 2002) and

the striatum, involved in social interactions (Báez-Mendoza and Schultz 2013). The fact that we

inferred positive rates r of density-dependence (Figure 4) suggested an acceleration of the

evolutionary tempo of trait evolution together with increased diversity of frugivorous sympatric

lineages for the hippocampus and the striatum. The MOB, the area involved in sensory abilities, also

tended to be best fitted by models considering sympatry as a whole. Yet, Brownian motion (BM)

was as likely as density-dependent or matching competition models, preventing firm conclusions on

whether sympatry affected or not MOB size evolution (Figures 3 and 4), especially since this

coincided with the most reduced sample size we had ($N_{species}$ = 34 to 39).

Next, we assessed whether higher levels of species sympatry lead to "bigger" or "smaller" brain

areas. To do so, we fitted phylogenetic regressions in extant frugivorous species between the

relative sizes of the different brain areas and two measures of sympatry (1) the average percentage

of overlapping range with other frugivorous sympatric species, and (2) the number of such

sympatric frugivorous species across their current entire distribution range. When weighting each

brain area by the whole body mass, the number of sympatric species never significantly influenced

the relative brain sizes (Table 1). Conversely, we found that the average percentage of overlapping

range significantly correlated, or tended to correlate, with the relative size of brain areas that were

better fitted with models considering sympatry: the hippocampus and the striatum (hippocampus: t

= -1.94, p = 0.058; striatum: t = -2.26, p = 0.028). The correlations were all negative (hippocampus:

est. = -0.39, CI95% = [-0.76,-0.01]; striatum: est. = -0.4, CI95% = [-0.77,-0.04]), which means that

higher range overlap between sympatric species associates with lower relative size, insensitive to

data and phylogenetic uncertainties (Supplementary Material Table S1, Supplementary Figure S7).

The acceleration of the tempo of trait evolution with species sympatry ($r > 0$ in the density-

dependent models) suggests that, compared with isolated species, sympatric species are subject to a

positive selection towards smaller brains, and not to a less intense selection for advanced cognitive

abilities. Similar results were obtained when weighting each brain area by the whole brain size

instead of the whole body mass (see Supplementary Material).

Furthermore, we also investigated the influence of sympatry on body mass, which could be a

confounding explanation for the observed patterns (Smaers et al. 2021). We found that the

evolutionary history of body mass is best explained by MC models (Supplementary Figure S6). In

other words, species in sympatry tend to diverge towards being large or small. There is thus no

overall linear relationship of the whole body mass with sympatry, explaining why we found no

effect of sympatry variables on body mass in PGLS; see Supplementary Table S2).

Finally, we investigated the evolutionary consequences of cognition and species sympatry by

evaluating whether brain sizes and sympatry intensities correlated with the lineage-specific net

diversification rates of primates (Figure 5). Overall, species diversification rates, estimated based on

the primate molecular phylogeny, particularly boomed in the early and late Miocene, around 25 and

11 Myr ago (Figure 5). When accounting for phylogenetic dependence, no significant relationship

between the net diversification rate and the relative size of brain areas was found (Table 2,

Supplementary Figure S8; see robustness in Supplementary Material Table S3). Although

diversification was uncorrelated with brain size in frugivorous primates, it was influenced by the

sympatry context. In particular, phylogenetic regressions highlighted a negative effect of the

number of sympatric species on the diversification rate (est. = -5.04e-03, CI95% = [-0.01,1.34e-04], t

= 2.56e-03, p < 0.001, Table 3, Supplementary Figure S9, Supplementary Material Table S5). In other

words, the higher the number of sympatric species, the lower the diversification rate.

DISCUSSION

Sympatry of frugivorous primate species impacts the evolution of their

hippocampus and striatum sizes

The size of the brain is subject to a compromise between the energy it incurs, and the increase of

fitness it allows. This is clearly emphasised by the fact that the evolution of the biggest brain areas,

the cerebellum and the neocortex, as well as the whole brain (Encephalization Quotient; EQ), were

best fitted by the Ornstein-Uhlenbeck process. This may suggest a stabilisation towards an optimal

size resulting from an equilibrium between costs and benefits. Although allometric and

developmental constraints, as well as spatial proximity in the brain, can induce correlation in the

evolution of different brain areas (Gómez-Robles, Hopkins, and Sherwood 2014), brain areas

underpin different cognitive functions and can thus be under different, independent, selective

pressures (Barton and Harvey 2000). The functional regionalisation is for instance evidenced here

by the differences in relative sizes across lineages in the MOB, with larger sizes in the lemuriformes

that mostly rely on smell to forage. The differences in evolutionary trajectories are also highlighted

by the variations in the best fit models of size evolution for the different brain areas. We indeed

show that sympatry is one factor that affects the evolutionary regime under which only some brain

area evolves: although the brain as a whole was not significantly affected by species sympatry, the

latter nonetheless induced a change in the relative size of the hippocampus and the striatum,

independently of whether the relative size was obtained by weighting by whole body mass or whole

brain size (Supplementary Figure S6). Theoretically, the two weighting methods are expected to

give insights into differences in resource allocation between body tissues or differences in resource

allocation within the brain tissue, respectively. Biologically, nonetheless, we observed no

differences in the response to sympatry by weighting by whole body mass or whole brain size,

despite weak to moderate correlations between sizes relative to the whole body mass and sizes

relative to the whole brain size (see Supplementary Figure S5). Therefore, our results suggest that

species sympatry simultaneously impacted between-tissues and within-brain reallocations for the

hippocampus and the striatum.

The hippocampus and the striatum are areas involved in individual-based and social-based

information processing, pinpointing that these two components might be under strong selection in

primates (DeCasien, Williams, and Higham 2017; Powell, Isler, and Barton 2017; González-Forero

and Gardner 2018). The hippocampus in particular, may have played a considerable role in the

evolution of primate-like behaviours (Schilder, Petry, and Hof 2020), driven by the changes that

primates faced in their ecological environment (e.g. the spatio-temporal distribution of the food,

DeCasien and Higham 2019), or by the social environment that they faced (e.g. the number of

conspecifics to interact with, Todorov et al. 2019; DeCasien and Higham 2019).

Overall, the fact that the hippocampus, particularly relevant to process and memorise spatio-

temporal information, is sensitive to sympatry, is consistent with the idea of an effect of sympatric

species on resource spatio-temporality (Hypothesis 1). Competition is generally the first-thought

mechanism to describe community structures (de Almeida Rocha et al. 2015) because it might affect

the environment in which species evolve. We show that a higher intensity of sympatry is actually

associated with smaller sizes of the hippocampus (in accordance with Prediction 1.2). This suggests

that indirect competition for food might contribute to convoluting the environment, and such an

over-complexification of the resource spatio-temporality may render cognitive foraging not

advantageous anymore. As a result, it might even generate a selection for smaller brain areas

involved in foraging. In parallel with the complexification of their environment, species might have

also narrowed frugivorous primates' niche, which might have synergistically affected their

cognition. Indeed, the support for matching competition when modelling the evolution of the whole

body mass of frugivorous primates indicates that sympatric primate species tended to diverge in

terms of body mass, evolving towards lower or higher body mass in sympatry. This is consistent

with the idea of niche partitioning in sympatry, e.g. where low body mass primate species would

occupy the canopy layer, while heavier primate species would occupy the ground. Such niche

partitioning may be accompanied by dietary specialisation (Schreier et al. 2009) and impact

cognitive abilities (Aristide et al. 2016), as dietary specialisation often requires lower cognitive

abilities and thus smaller brain area sizes. Agent-based modelling (Railsback and Grimm 2019)

could greatly contribute to improving the understanding of the proximal mechanisms involved and

in particular the link between resource spatio-temporal complexification, niche partitioning, and

cognition.

In contrast to direct/indirect competition, potential indirect facilitation between species due to

"social" cues (Hypothesis 2), is ruled out by the absence of an effect of sympatry on brain areas

involved in immediate sensory information processing (e.g. cerebellum or neocortex). This absence

of effect can stem from two possibilities. Either foragers do not exploit cues left out by sympatric

heterospecifics, or it has been shown that foragers tend to use social information over

environmental (i.e. personal) information, in particular in non-perfectly predictable environments

(Rafacz and Templeton 2003; Dunlap et al. 2016). Thus, if environmental complexity increases too

much, "social" cues provided by heterospecifics might replace environmental ones. As such,

stimulation intensity of the MOB, the cerebellum, or the neocortex would somehow remain

equivalent when in sympatry or not. Further work should explicitly test for these possibilities.

As expected (Hypothesis 3), the striatum size was relatively larger in callitrichines, particularly

known to form mixed-species groups (Heymann and Hsia 2015). Yet, overall, the striatum size was

negatively affected by sympatry. This puzzle might take root in secondary, but key, functions

supported by the striatum, namely reward expectation, goal-directed behaviour, and planning

abilities (Johnson, Meer, and Redish 2007). These three functions might as well be advantageous

when foraging. As for the hippocampus, then, the increase in the environmental unpredictability

could diminish the benefits of these future-oriented skills.

Sympatry of frugivorous primate species is associated with slower diversification

Given the context-dependence of the direction of selection (towards bigger sizes when sympatry is

low, smaller sizes otherwise), there is no surprise that we do not observe a correlation between the

net diversification rate and the three brain areas affected by species sympatry. Surprisingly

however, we found no positive association between the net diversification rate and the EQ, the

cerebellum, or the neocortex, which were insensitive to species sympatry. By contrast, a positive

association between brain size and diversification was also found in birds (Sayol et al. 2019) given

that bigger brains act as a buffer to environmental challenges (Sol et al. 2007). A visual inspection of

the regressions clearly evidenced a positive trend if not considering phylogeny (EQ and neocortex,

Figure S7). Sudden encephalisation in primates is clearly associated with a limited number of

closely-related species (DeCasien, Williams, and Higham 2017; Melchionna et al. 2020). Thus, this

clearly limits the statistical power of our phylogenetically-corrected analyses, as we cannot decipher

whether larger brain size and faster species diversification result from a true biological link or

appeared simultaneously but independently. This means that a positive association between brain

size and species diversification remains a likely possibility (as previously suggested in primates,

Melchionna et al. 2020). Species sympatry, however, induced a significant slowdown in primate

diversification, a density-dependence trend frequently observed in many tetrapod clades

(Condamine, Rolland, and Morlon 2019). This frames coherently with a competitive scenario, where

the tempo of species diversification decreases when ecological niches are progressively filled up

(Rabosky and Lovette 2008). Species competing for resources are thought to contribute to limiting

competitors' range (Price and Kirkpatrick 2009), hence constraining population size and

diversification rate (Pigot and Tobias 2013).

Limits

The use of brain size as a proxy for cognition is a central debate with no optimal solution (see

grounded criticism from Deaner, Nunn, and Schaik 2000; Healy and Rowe 2007; Logan et al. 2018;

Van Schaik et al. 2021). The current flourishment of consortia, allowing for much more detailed and

standardised anatomical measurements (e.g. in primates: Milham et al. 2018), or with standardised

behaviourally explicit comparisons (e.g. on captive, Altschul et al. 2019; or wild primates, Janmaat et

al. 2021), might alleviate biases stemming from brain size analysis, but this will take time to

generate large-enough datasets. In the meanwhile, brain size is a proxy much appreciated in

practice, because of its easy accessibility for a "large" number of species, while the multifaceted

aspect of cognition can simply be taken into account by considering the brain as a mosaic of singular

and independent regionalised areas that are cognitively specialised. Yet, this approach implies

collating datasets from different sources, which can induce a high intra- and inter-species

variability, as the methodology has changed over time (discussed in Navarrete et al. 2018). We

therefore sampled measures among the available datasets and repeated the analysis several times,

in order to account for the variability of the data sets. The results remained robust to this approach.

This supports the idea that intra-species variation is reduced relative to inter-species variation, a

necessary condition for brain size to be an honest signal of cognition differences across species,

despite differences in measurement protocols.

Not only can the nature of the data be biased, but the methods themselves can suffer from

significant limitations. In our case, these limitations are particularly true for the reconstruction of

diet history, ancestral biogeography, or species diversification that we performed, which infer the

most likely evolutionary histories based on observations at present alone. To verify the realism of

these inferences, we confronted the diet history reconstruction with fossil evidence. Dental

microwear textures in tooth fossils can indeed be used to reconstruct past diet and we found that

our estimates of ancestral primate diets were consistent with available fossil evidence

(Ramdarshan, Merceron, and Marivaux 2012; Merceron et al. 2009, see Supplementary Figure S1).

Similarly, we found that our estimates of species diversification rates from molecular phylogenies

were consistent with estimates of past primate diversity reconstructed from all available fossil data

(Springer et al. 2012). In addition, the identified breakpoints matched estimates from previous

studies (Arbour and Santana 2017; Springer et al. 2012). We did not use fossils for the

biogeographic reconstruction because, although this may help the reconstruction, the overall

benefices are minor (Wisniewski, Lloyd, and Slater 2022) and highly influenced by the fossil data

quality (e.g. spatial coverage), which is still heavily limited and biased. We nevertheless accounted

for the uncertainty in all reconstructions in our analyses by reporting results based on multiple sets

of ancestral reconstructions.

Furthermore, the effectiveness of a method also depends on the proxy and associated definitions,

which may clearly impact the insights of such large-scale analyses. For instance, this issue is clearly

illustrated with group size being used as a proxy of social complexity in social cognition (Dunbar

and Shultz 2007). In this study, brain areas were associated with one main function, an overlap of

diets was considered within the whole frugivorous primate clade, and sympatry was considered to

be based as soon as spatial overlap occured. Instead, each brain area can be considered as a Russian

doll, with possibly redundant sub-functionalization between spatially distant areas, frugivorous

primates can be differentially selective in the fruits they eat (Campera et al. 2019), and species can

avoid each other spatio-temporally at fine scale (Deane et al. 2013). Although we cannot exclude

that more accurate definitions might change our results, it is important to note that such a

simplification is imposed by many constraints: from the sake of theorising to computational time. In

addition, these large definitions also have some advantages. First, they can capture a broad variety

of situations. For instance, simply defining sympatry as co-occurrence in a given biogeographic area

enables considering both, direct and indirect interactions. Second, more stringent definitions might

induce strong initial constraints that would eventually bias inferences of past history. For instance,

the considered range areas to define sympatry are large and are unlikely to capture smaller-scale

spatial segregation between species (e.g. due to habitat-specific segregation or vertical segregation).

Yet, a finer mapping of species range may poorly reflect the "realised niche," as primate populations

suffer intense local extinction in recent years due to human activity (Estrada et al. 2017; Pavoine et

al. 2019). Furthermore, we also took care of varying the stringency of our definitions

(e.g. thresholds used to consider a species frugivorous based on the percentage of frugivory,

thresholds used to consider two species as sympatric based on range overlap), and it highlighted

that results were robust to the threshold choices (see Supplementary Material).

CONCLUSION

We showed that species sympatry is an important factor shaping the evolutionary history of

primates' brains. It now seems crucial to scrutinise more carefully how sympatry fits with other

socio-ecological variables that have been shown to influence brain size evolution in this clade

(e.g. diet, group size, home range etc., see DeCasien, Williams, and Higham 2017; DeCasien and

Higham 2019; Powell, Isler, and Barton 2017). In addition, dietary overlap and food competition

might not only happen between frugivorous primates, but any frugivores in the same area. In fact, it

is very likely that any hypotheses on cognition evolution, generally discussed within species, could

be broadened to a between-species context: foraging facilitation between species does exist (Olupot,

Waser, and Chapman 1998; Havmøller et al. 2021), and so do polyspecific social associations (Porter

2001), as well as inter-species territory defence (Drury, Cowen, and Grether 2020; Losin et al. 2016)

or imitation and copying (Persson, Sauciuc, and Madsen 2018; Pepperberg 2002). Similarly, prey-

predator races could shape selection on cognitive abilities (Shultz and Dunbar 2006). As Alice said

"It's a great huge game of chess that's being played—all over the world" (Carroll 1871, chap. II) and

all individuals are just pieces to play with or against, no matter the species.

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AUTHORS' CONTRIBUTION

634 BR conceived the study, gathered, cleaned and analysed the data, drew the figures, wrote the first

635 version of the manuscript and subsequently revised it. BP-L implemented the ClaDS algorithm for

636 our data, helped with running other analyses and drawing figures, and revised the manuscript

637 multiple times. The authors declare having no conflict of interest. All authors gave final approval for

638 publication and agree to be held accountable for the work performed therein.

DATA ACCESSIBILITY

640 The data and codes that support the findings of this study are openly available at

641 https://github.com/benjaminrobira/Meta_analysis_cognition_primates.

	Est.	CI2.5%	CI97.5%	Sd	t	p-value
EQ (log) (N=127)						
Intercept	-0.17	-0.53	0.22	0.20	-	-
% of overlapped range	0.02	-0.08	0.13	0.05	0.41	0.68
Number of sympatric frugivores (sqrt)	0.02	-0.02	0.05	0.02	1.03	0.31
Lambda	0.98	0.94	1.00			
Hippocampus (/body mass, log) (N=50)						
Intercept	-0.92	-1.95	0.05	0.53	-	-
% of overlapped range	-0.39	-0.76	-0.01	0.20	-1.94	0.06
Number of sympatric frugivores (sqrt)	0.08	-0.06	0.20	0.07	1.21	0.23
Lambda	0.99	0.92	1.00			
Neocortex (/body mass, log) (N=56)						
Intercept	2.07	1.31	2.86	0.41	-	-
% of overlapped range	-0.23	-0.54	0.11	0.16	-1.46	0.15
Number of sympatric frugivores (sqrt)	0.02	-0.08	0.13	0.05	0.48	0.63
Lambda	0.99	0.91	1.00			

Cerebellum (/body mass, log) (N=57)

Intercept	0.60	-0.15	1.35	0.39	-	-
% of overlapped range	-0.08	-0.32	0.17	0.12	-0.7	0.49
Number of sympatric frugivores (sqrt)	-0.01	-0.1	0.07	0.04	-0.34	0.74
Lambda	1.00	0.96	1.00			

Striatum (/body mass, log) (N=50)

Intercept	-0.36	-1.18	0.44	0.44	-	-
% of overlapped range	**-0.40**	**-0.77**	**-0.04**	**0.18**	**-2.26**	**0.03**
Number of sympatric frugivores (sqrt)	0.03	-0.08	0.15	0.06	0.61	0.54
Lambda	0.98	0.85	1.00			

MOB (/body mass, log) (N=31)

Intercept	-2.76	-4.61	-0.93	1.00	-	-
% of overlapped range	-1.20	-2.65	0.35	0.80	-1.49	0.15
Number of sympatric frugivores (sqrt)	0.21	-0.18	0.56	0.19	1.12	0.27
Lambda	1.00	1e-07	1.00			

Table 2: Relative brain sizes do not correlate with primate species diversification in frugivorous primates | Model
estimates and significance of Bayesian phylogenetic regressions to assess the correlation between the net
diversification rates and the relative brain sizes. Est.=Estimate, HDP2.5%=Lower border of the 95% Highest
Posterior Density, HDP97.5%=Upper border of the 95% Highest Posterior Density, Eff. samp.=Effective sample
(adjusted for autocorrelation). The brain areas (as well as the associated sample sizes) are indicated prior to each
list of estimates. The (log) indicates log-transformed variables, while the (/body mass) indicates variables
pondered by body mass.

	Est.	HDP2.5%	HDP97.5%	Eff. samp	pMCMC
Diversification EQ (N=148)					
Intercept	0.12	0.08	0.16	900.00	-
EQ (log)	0.02	-7.91e-03	0.05	789.25	0.15
Lambda	0.83	0.76	0.9		
Diversification Hippocampus (N=61)					
Intercept	0.13	0.09	0.18	900.00	-
Hippocampus (/body mass, log)	9.10e-03	-9.48e-03	0.03	900.00	0.34
Lambda	0.73	0.6	0.85		
Diversification Neocortex (N=67)					
Intercept	0.1	0.04	0.17	991.53	-
Neocortex (/body mass, log)	7.26e-03	-0.02	0.03	900.00	0.56
Lambda	0.74	0.6	0.86		
Diversification Cerebellum (N=68)					
Intercept	0.12	0.07	0.16	900.00	-
Cerebellum (/body mass, log)	3.94e-03	-0.02	0.03	989.21	0.76
Lambda	0.74	0.6	0.86		

Diversification Striatum (N=61)					
Intercept	0.12	0.08	0.17	900.00	-
Striatum (/body mass, log)	9.11e-03	-0.01	0.03	900.00	0.44
Lambda	0.73	0.59	0.85		
Diversification MOB (N=37)					
Intercept	0.11	0.05	0.17	900.00	-
MOB (/body mass, log)	-4.79e-03	-0.02	0.01	900.00	0.59
Lambda	0.65	0.46	0.83		

659 **Table 3: Species sympatry slowdowns primate diversification | Model estimates and significance of phylogenetic**
660 **regressions to assess the correlation between diversification rate and species sympatry. Est.=Estimate,**
661 **CI2.5%=Lower border of the CI95%, CI97.5%=Upper border of the CI95%, Sd= Standard deviation, t= Statistics t-**
662 **value. The brain areas (as well as the associated sample sizes) are indicated prior to each list of estimates. The**
663 **transformation (logarithm or square-root) is indicated in parentheses by the abbreviation (log or sqrt).**

	Est.	CI2.5%	CI97.5%	Sd	t	p-value
Diversification (N=128)						
Intercept	0.15	0.10	0.2	0.03	-	-
% of overlapped range	-5.40e-03	-0.02	9.35e-03	8.14e-03	-0.66	0.51
Number of sympatric frugivores (sqrt)	**-5.04e-03**	**-0.01**	**1.34e-04**	**2.56e-03**	**-1.97**	**0.05**
Lambda	0.96	0.89	0.99			

Figure 1: Biogeographic areas used for reconstructing the history of sympatry in frugivorous primates represented on the Mercator projection of the world | Areas were defined as a combination of geographic and environmental criteria relative to the primate taxonomy following results from Kamilar (2009): (1) East Madagascar (2) West Madagascar (3) West Africa (4) Central Africa (5) East/South Africa (6) Central America (7) North South-America (8) South South-America (9) West Asia (10) Central/East Asia (11) South Asia (12) Asian peninsula and islands. Note that the northern Africa and the southern of Europe were discarded because *Macaca sylvanus* was not considered.

Figure 2: The intensity of species sympatry varies across the primate phylogeny | Primate phylogeny from the
consensus tree of the 10kTrees project is depicted in the center, together with abbreviated species names. The
corresponding non-abbreviated names can be found using Supplementary Figure S4. Sympatric frugivorous
(based on a frugivory threshold of 20% and folivory threshold of 40%) species are linked by light grey lines. The
geographic areas occupied by a species are depicted by coloured rectangles. Presence was assessed given an
overlap between the species range and the geographic area of 10%.

Figure 3: Variations in relative brain size areas among extant frugivorous primates | (Left) Circular plot of the relative sizes of the different brain areas. Colours indicate the rows for the different brain areas. The darker background emphasises when values are above average, while the lighter background emphasises when values are below average. The mean value (after scaling and based on one random sampling among possible values, but see Supplementary Figure S3 for visualization of measure variability) for the Encephalization Quotient (EQ) or relative size of brain areas, when available, is depicted by a plain circle for frugivorous species. The frugivorous threshold was fixed to 20% and the folivory threshold to 40%. (Right) The different studied brain areas (human brain as an illustration). In short, the MOB is involved in immediate olfactory information processing, the neocortex and the cerebellum support working memory and memory consolidation of immediate sensory information processing (Wiltgen et al. 2004; Koziol et al. 2014; Sokolov, Miall, and Ivry 2017), and the hippocampus supports a working memory and a long-term spatio-temporal memory (Burgess, Maguire, and O'Keefe 2002). The striatum is involved in social information processing (Báez-Mendoza and Schultz 2013).

Figure 4: The evolution of the hippocampus and striatum in frugivorous primates are best fitted by models of trait
evolution considering species sympatry | Plotted is the AICc weight, a measure of relative support for a given
model, for models not considering species sympatry (BM, OU, EB) or considering species sympatry (MC, DD_{lin},
DD_{exp}). The points represent the average AICc weight obtained (when considering the six models from the same
run), while the vertical bars indicate the standard deviation given all tested conditions (see Phylogenetic models of
trait evolution: does species sympatry shape brain size evolution?).

Figure 5: Species sympatry in primates is negatively associated with species diversification | a) Net diversification rate over time in the primate taxon. The average diversification rate estimated based on an assumed sampling fraction of primate species ranging from 60 to 95% (at a step of 10%; then 5% from 90%) is depicted by the plain line. The grey background depicts the standard deviation. The two breakpoints, depicted by the plain dots and the vertical dotted bars, were estimated based on a three-linear regression segmentation using the *strucchange* package [Zeileis et al. (2002); Zeileis et al. (2003); Zeileis (2006); see the vignette package for statistical details]. The three fitted regressions are displayed by the dashed lines. The choice of two breakpoints was first assessed by choosing the number of breakpoints minimizing the Bayesian Information Criterion. b) The branches of the primate phylogenetic trees are colored according to their speciation rates estimated using ClaDS2: red colors indicate lineages that speciate frequently, whereas blue colors indicate lineages that rarely speciate. At the tips of the tree, we reported for each extant species whether we had data on its biogeography (column 'Data availability') and if so, we indicated the number of sympatric species it has. Using PGLS, we found a significant relationship between lineage-specific diversification rates and numbers of sympatric species at present.

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