

ManyPrimates: Establishing an Infrastructure for Collaboration in Primate Cognition Research

Elizabeth Warren and Drew Altschul on behalf of “ManyPrimates”

SPRG @ The Burn

11/05/2019

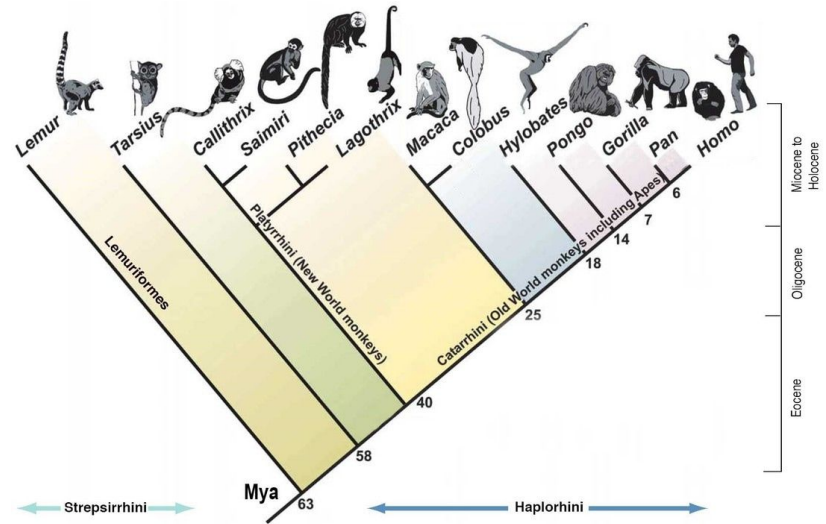
ManyPrimates contributors

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Challenges to primate cognition research

Important questions are under-studied due to lack of infrastructure:

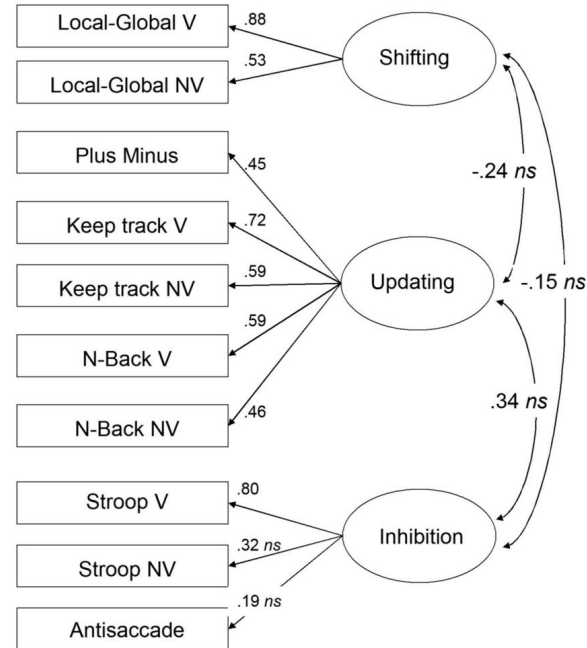
- How does cognition evolve?



Challenges to primate cognition research

Important questions are under-studied due to lack of infrastructure:

- How does cognition evolve?
- How is cognition structured?



Challenges to primate cognition research

Important questions are under-studied due to lack of infrastructure:

- How does cognition evolve?
- How is cognition structured?
- How does cognition develop?



Challenges to primate cognition research

Answering these questions needs:

- Large and diverse samples
- Pooling of resources across labs
- Infrastructure to support studies

ManyPrimates

Network to connect researchers, plan and conduct collaborative studies

Collaboratively deciding on research agenda

Open to all interested in primate cognition

Inspired by:

- Open Science Collaboration // ManyLabs // ManyBabies

Pilot study

Study the phylogeny of a fundamental cognitive ability:

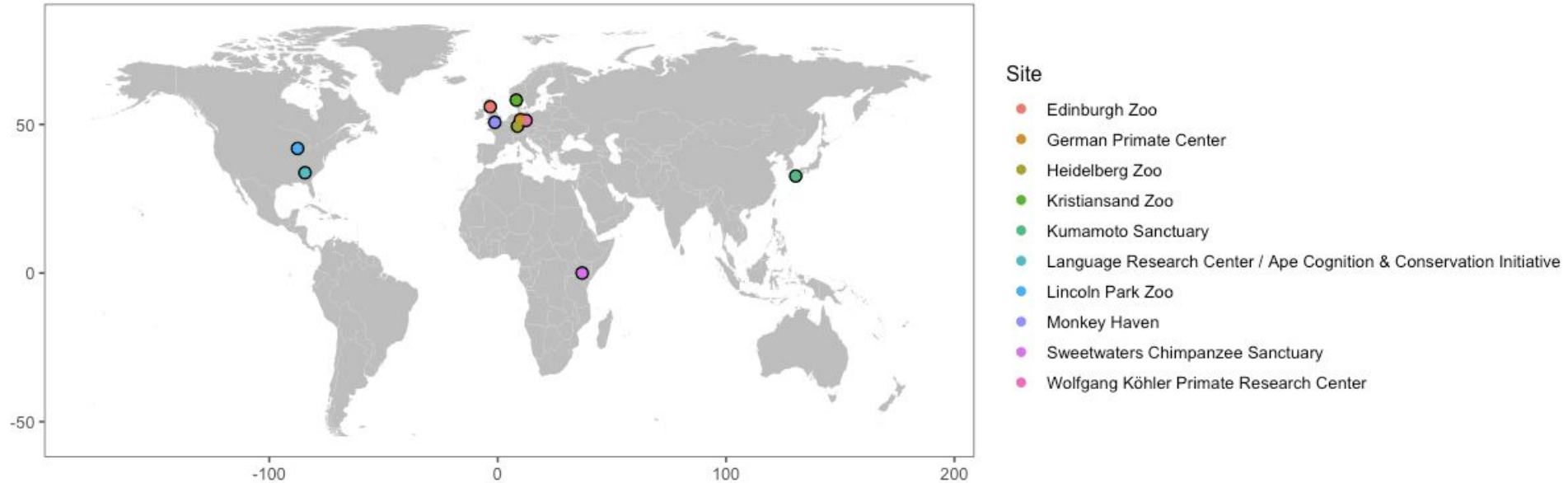
Short-term memory

Build basic infrastructure for future projects

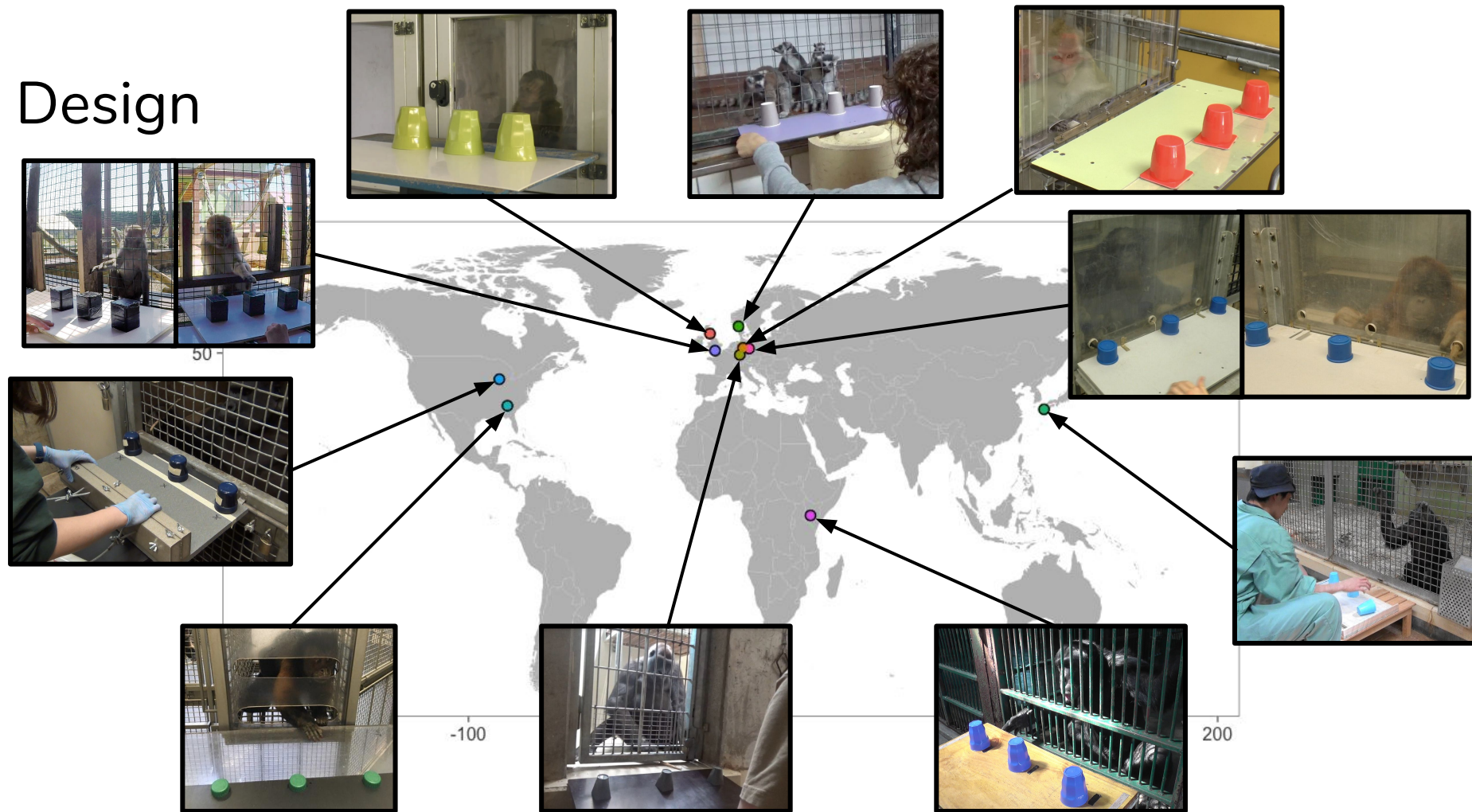
Timeline

December '17:	Initial contact
January '18:	Decision on topic for pilot study
February '18:	Pre-registration of design and analysis
March - July '18:	Data collection
August - February '19:	Data analysis and writing
March '19:	Manuscript submitted to PLOS One

Data collection sites



Design

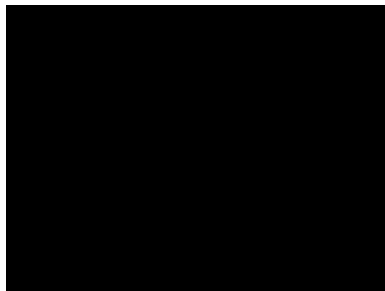


Design

Rhesus Macaque - Medium Delay



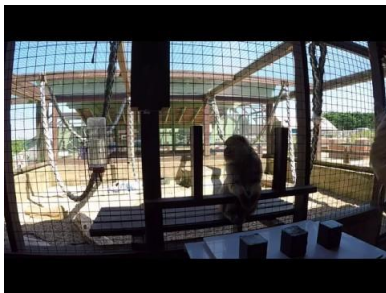
Chimpanzee - Medium Delay



Capuchin Monkey - Medium Delay



Barbary Macaque - Medium Delay

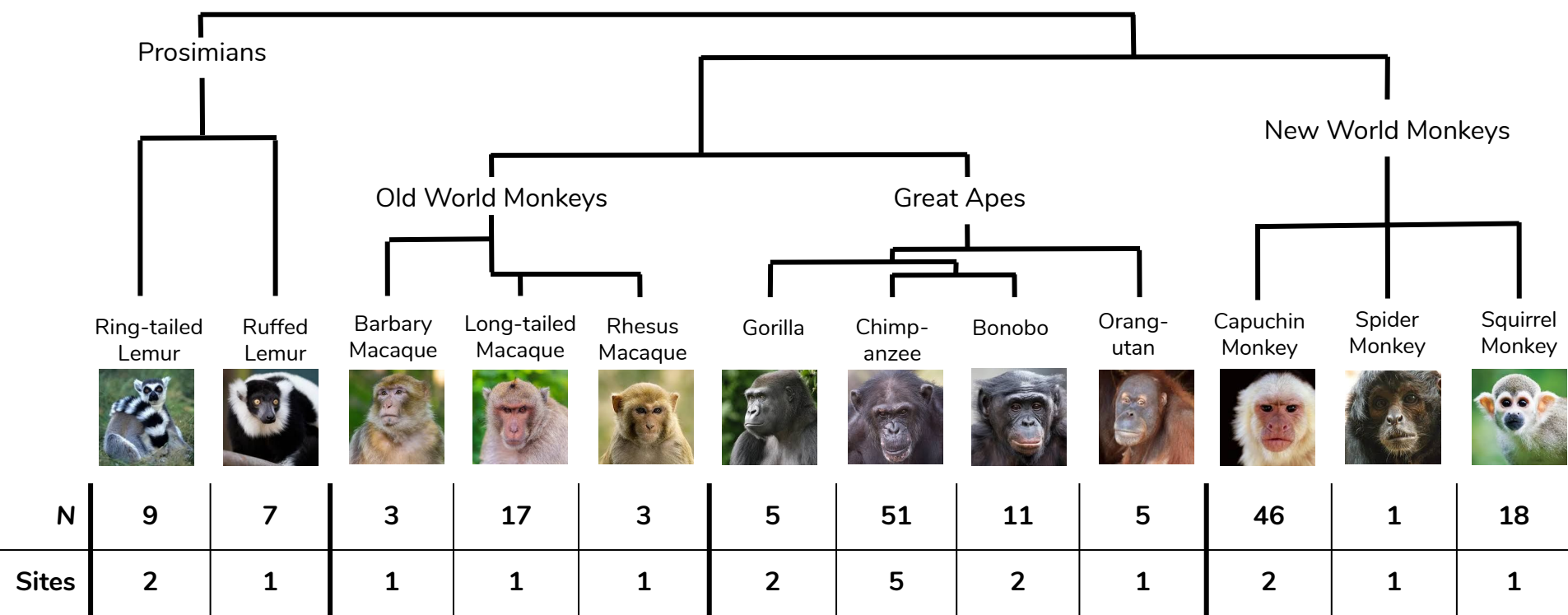


Squirrel Monkey - Short Delay



Sample

Total Species	12
Total Sites	11
Total N	176



Prosimians

New World Monkeys

Old World Monkeys

Great Apes

Ring-tailed
Lemur

Ruffed
Lemur

Barbary
Macaque

Long-tailed
Macaque

Rhesus
Macaque

Gorilla

Chimp-
anzee

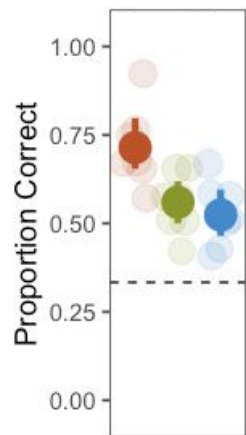
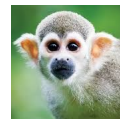
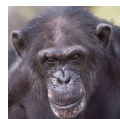
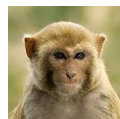
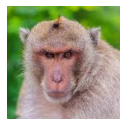
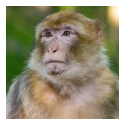
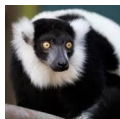
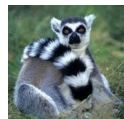
Bonobo

Orang-
utan

Capuchin
Monkey

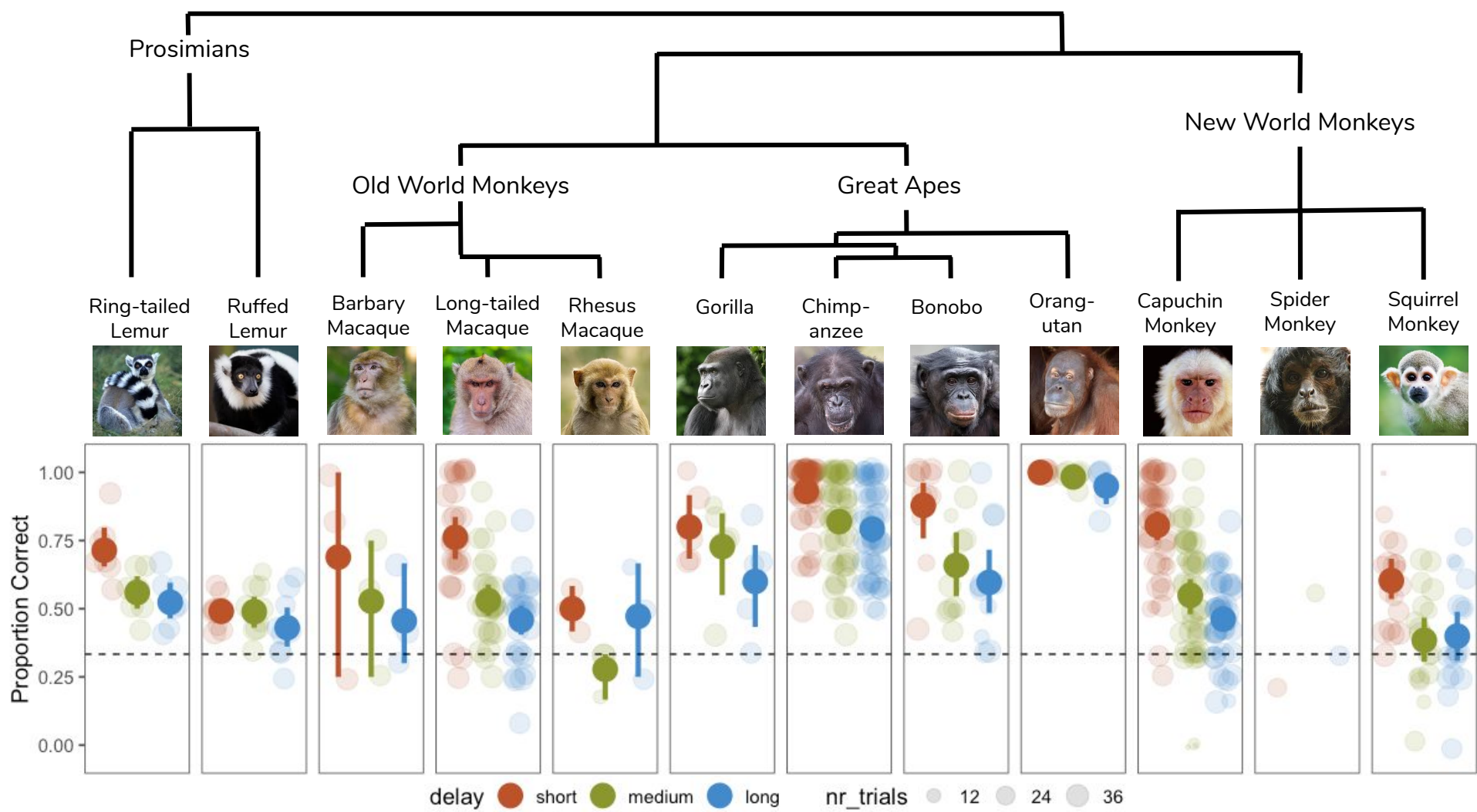
Spider
Monkey

Squirrel
Monkey



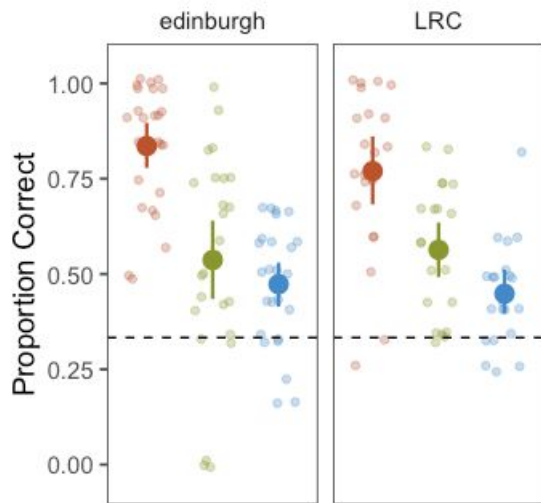
delay short medium long

nr_trials 12 24 36

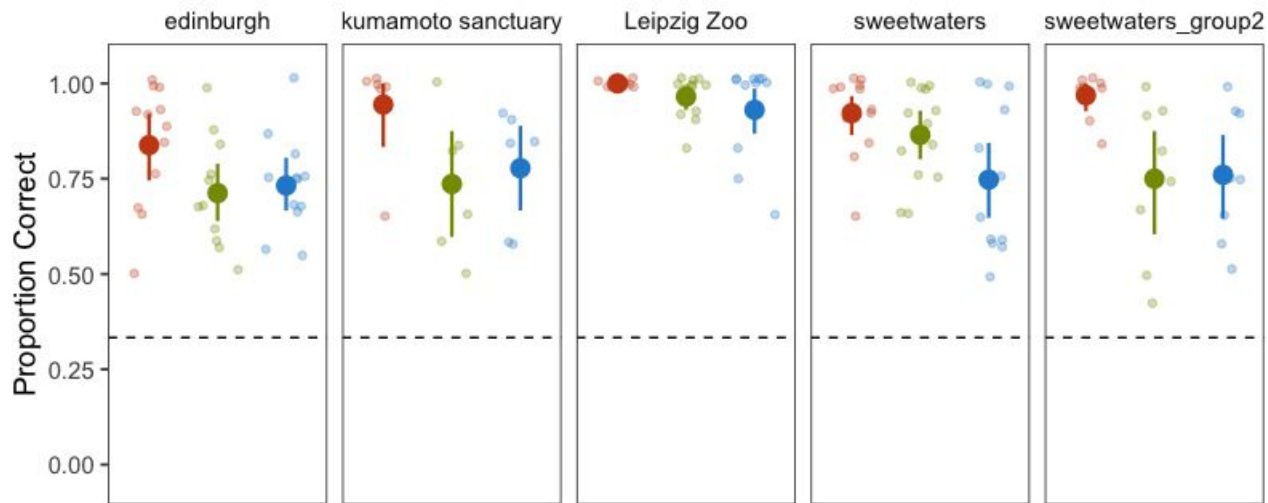


Variation across sites

Capuchin Monkeys

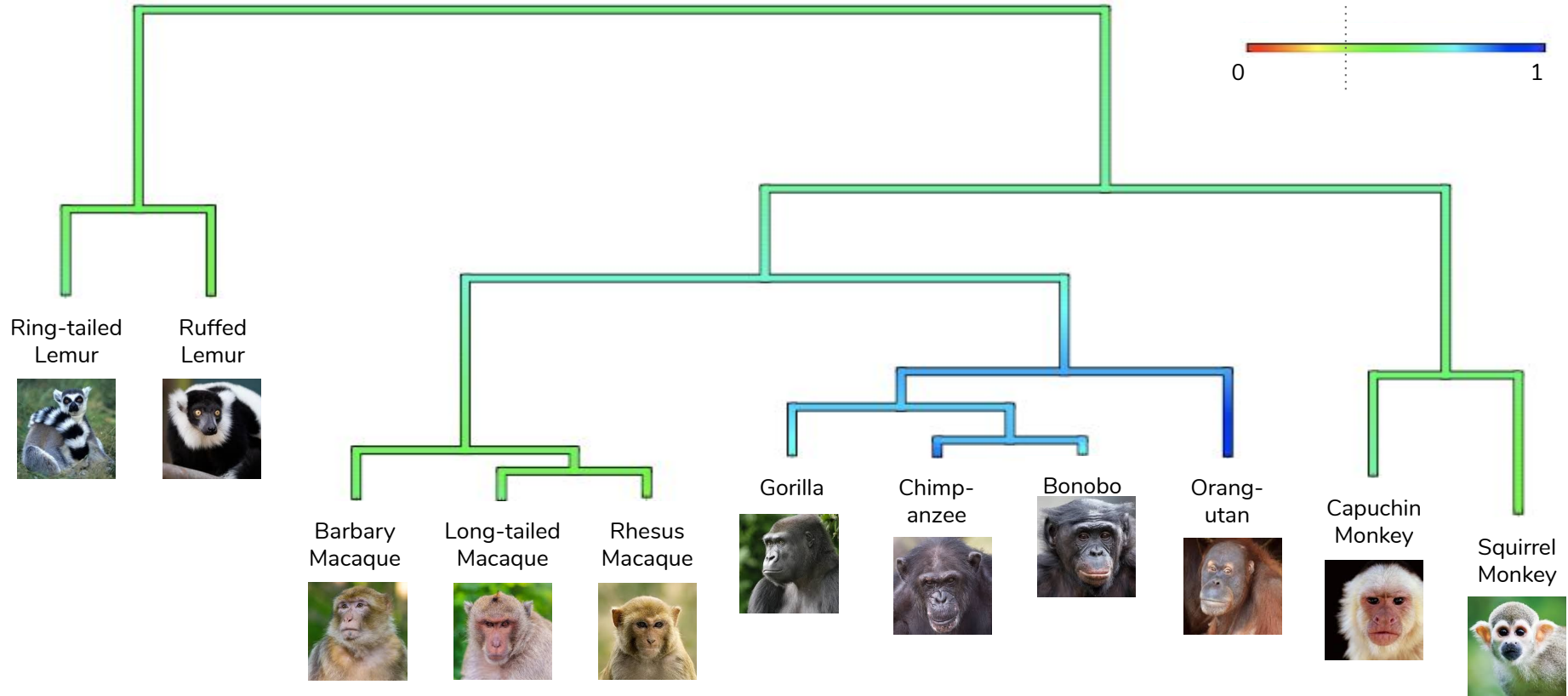


Chimpanzees



delay short medium long

Phylogenetic analysis



Contributions

Study planning

- Generating ideas // designing studies // coordination

Data collection

- Data collection // coding & reliability

Analysis and publication

- Data analysis // writing // public outreach

Outlook

- Spread the word and get more people involved
- Continue with short-term memory study
 - Currently working on pre-registration for ManyPrimates 1
- Collect ideas for future studies
- Further diversification (e.g. including non-captive samples)

Acknowledgements

Cog Etho Lab (DPZ): to Lukas Schad and Carolin Kade

Seed, Call, Völter: RZSS Edinburgh Zoo

Joly, Waller: to Charlotte Gurney-Read and the Monkey Haven, Isle of Wight

Beran, Flessert: Language Research Center and ACCI (especially Amanda Epping)

Sánchez-Amaro, Hanus: WKPRC Leipzig Zoo

Hernandez-Aguilar, Motes-Rodrigo: Dyreparken Kristiansand (especially Helene Axelsen and Tanya Michin)

Herrmann, Melis, Duguid, Haux: Sweetwaters Chimpanzee Sanctuary

Thank you!

ManyPrimates website:

https://github.com/ManyPrimates/mp_pilot

Code and data:

https://github.com/ManyPrimates/mp_pilot

Preprint:

<https://psyarxiv.com/3xu7q/>

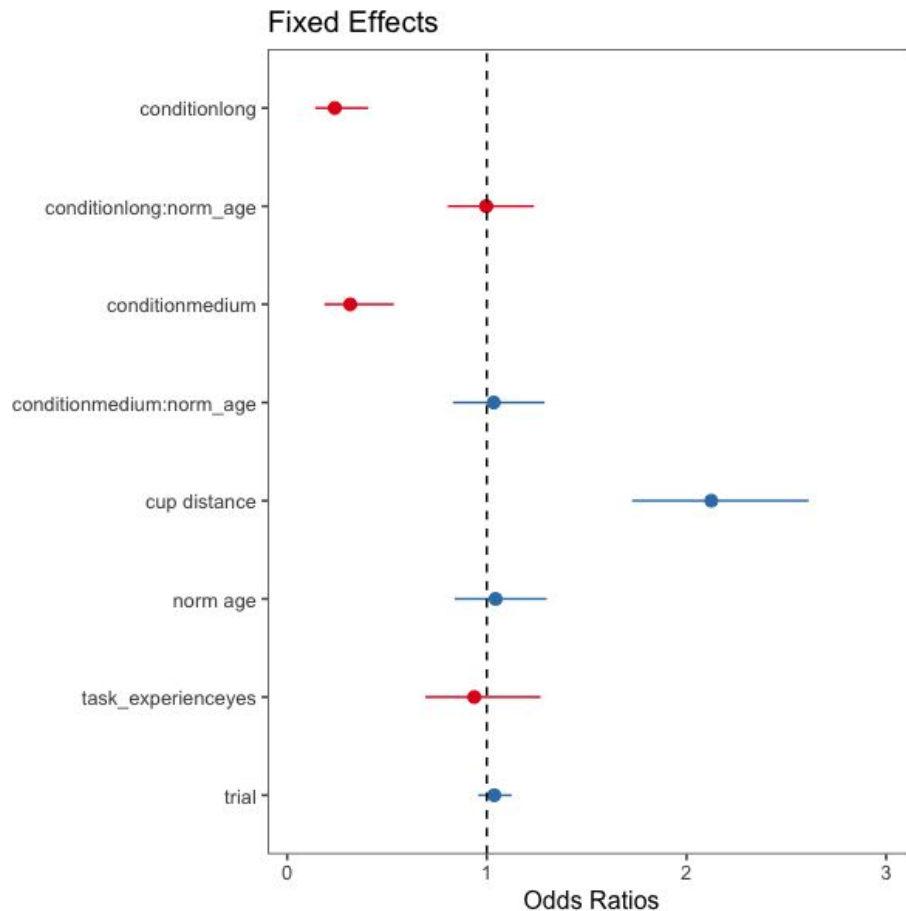
Results

Generalized linear mixed model:

```
correct ~ condition * age +
```

```
  task_experience + cup_distance +  
    board_size + trial +
```

```
(1 + condition + trial | site/subject) +  
  (1 + condition | species)
```



Results

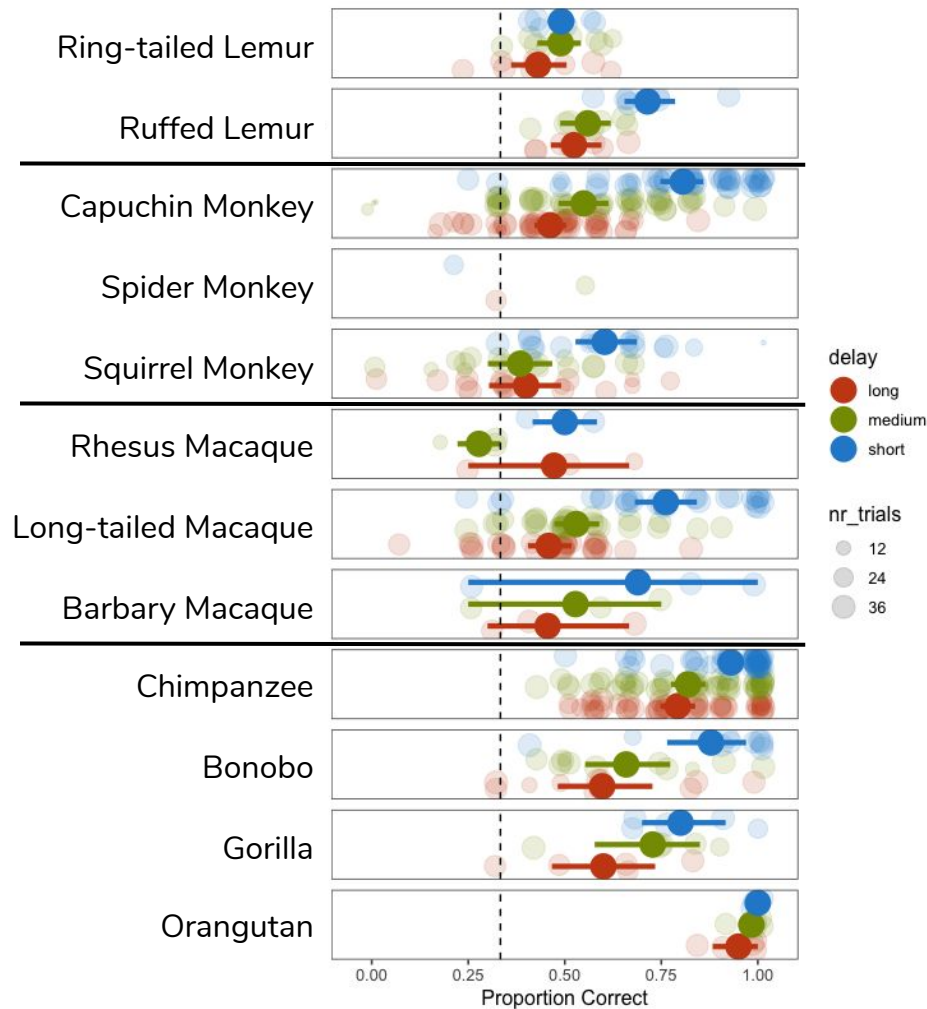
Generalized linear mixed model:

```
correct ~ condition * age +
```

```
task_experience + cup_distance +  
board_size + trial +
```

$$\beta_{\text{medium}} = -1.153, p > .0001$$

$$\beta_{\text{long}} = -1.433, p > .0001$$



Results

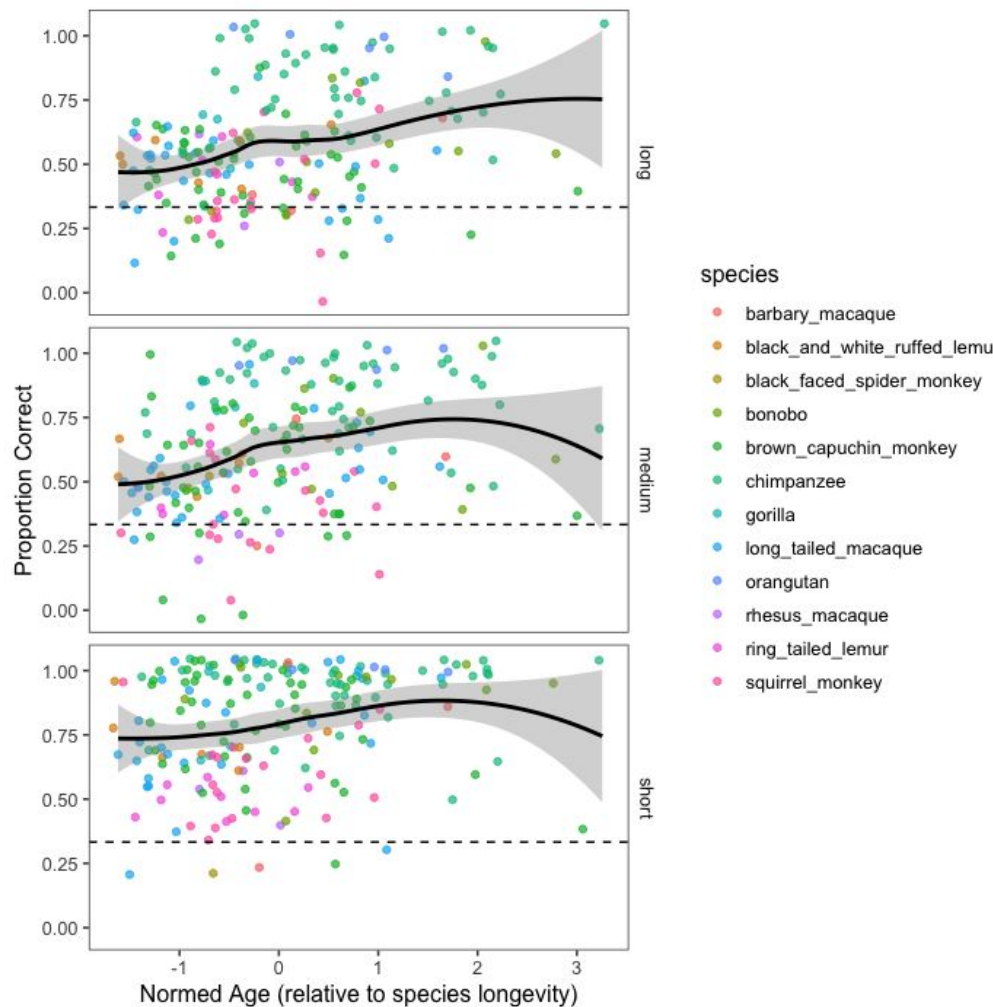
Generalized linear mixed model:

```
correct ~ condition * age +
```

```
  task_experience + cup_distance +  
    board_size + trial
```

$$\beta_{\text{medium*age}} = 0.034, p = .76$$

$$\beta_{\text{long*age}} = -0.003, p = .98$$



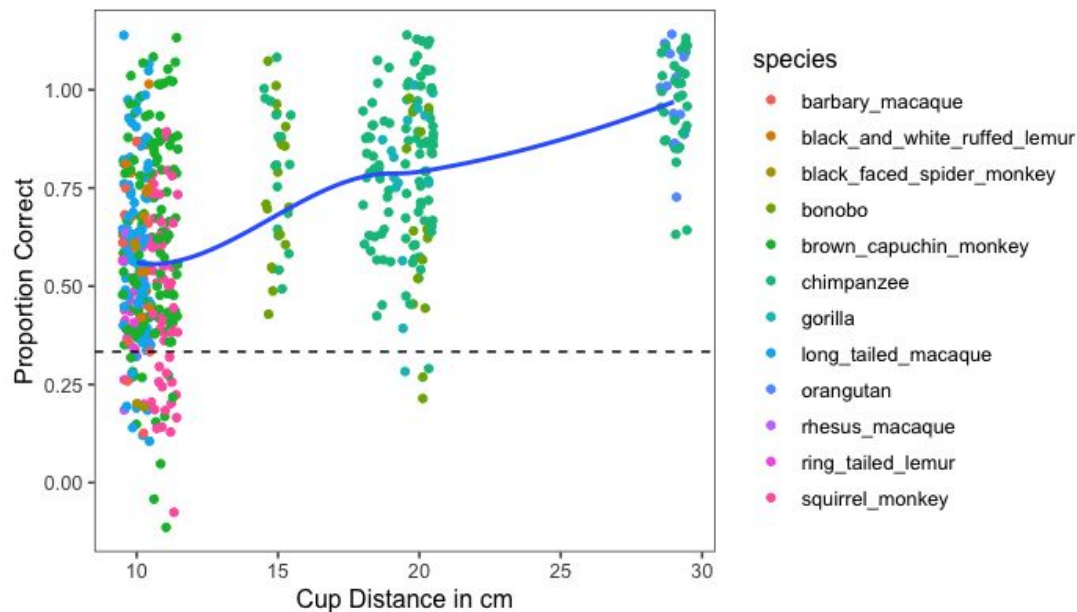
Results

Generalized linear mixed model:

```
correct ~ condition * age +
```

```
    task_experience + cup_distance +  
    board_size + trial +
```

$\beta_{\text{medium}} = 0.753$, CI [:], $p > .0001$



Phylogenetic analysis

Based on data averaged across conditions

Phylogenetic signal (Lambda - λ):

- Do values cluster as expected due to phylogenetic relatedness (range: 0 - 1)
 - Based on performance means
 - Based on phylogenetic t-tests (updated through comparison to chance level)

Recreation of ancestral state

- Estimate likely performance level of common ancestor

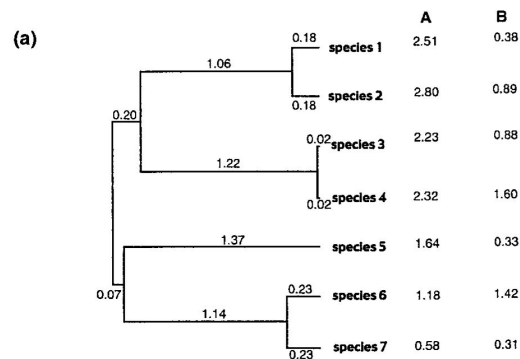
Phylogenetic analysis

Based on data averaged across conditions

Phylogenetic signal (Lambda - λ):

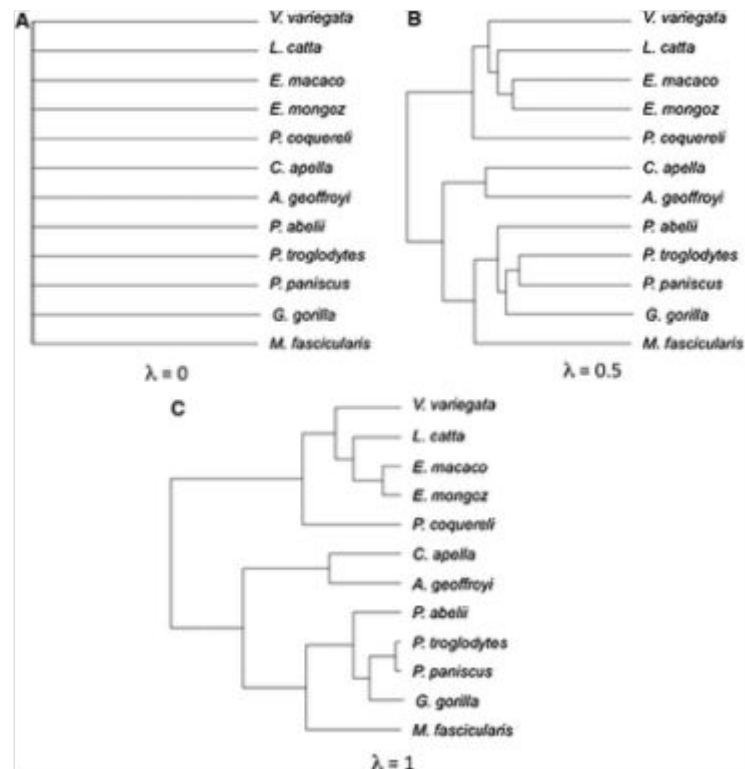
- Base λ : 0.74, $p = .26$
- Updated λ : 0.79, $p = .02$

More about λ

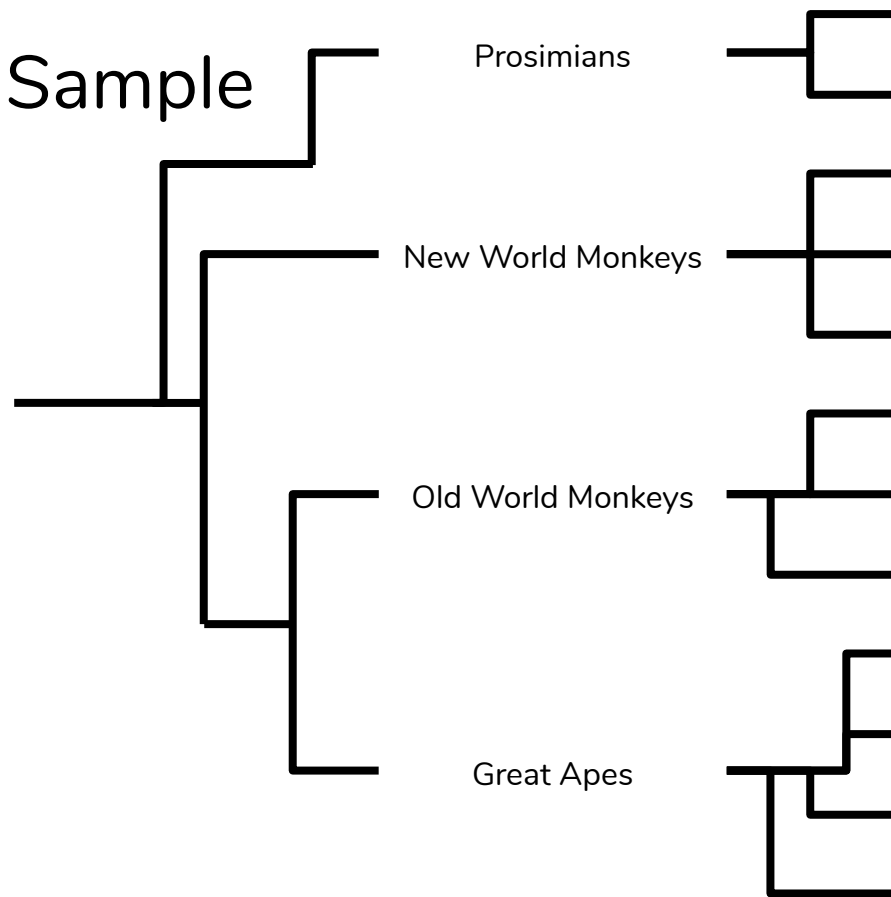


(b)

	species 1	species 2	species 3	species 4	species 5	species 6	species 7
species 1	1.44	1.26	0.20	0.20	0	0	0
species 2	1.26	1.44	0.20	0.20	0	0	0
species 3	0.20	0.20	1.44	1.42	0	0	0
species 4	0.20	0.20	1.44	1.44	0	0	0
species 5	0	0	0	0	1.44	0.07	0.07
species 6	0	0	0	0	0.07	1.44	1.21
species 7	0	0	0	0	0.07	1.21	1.44

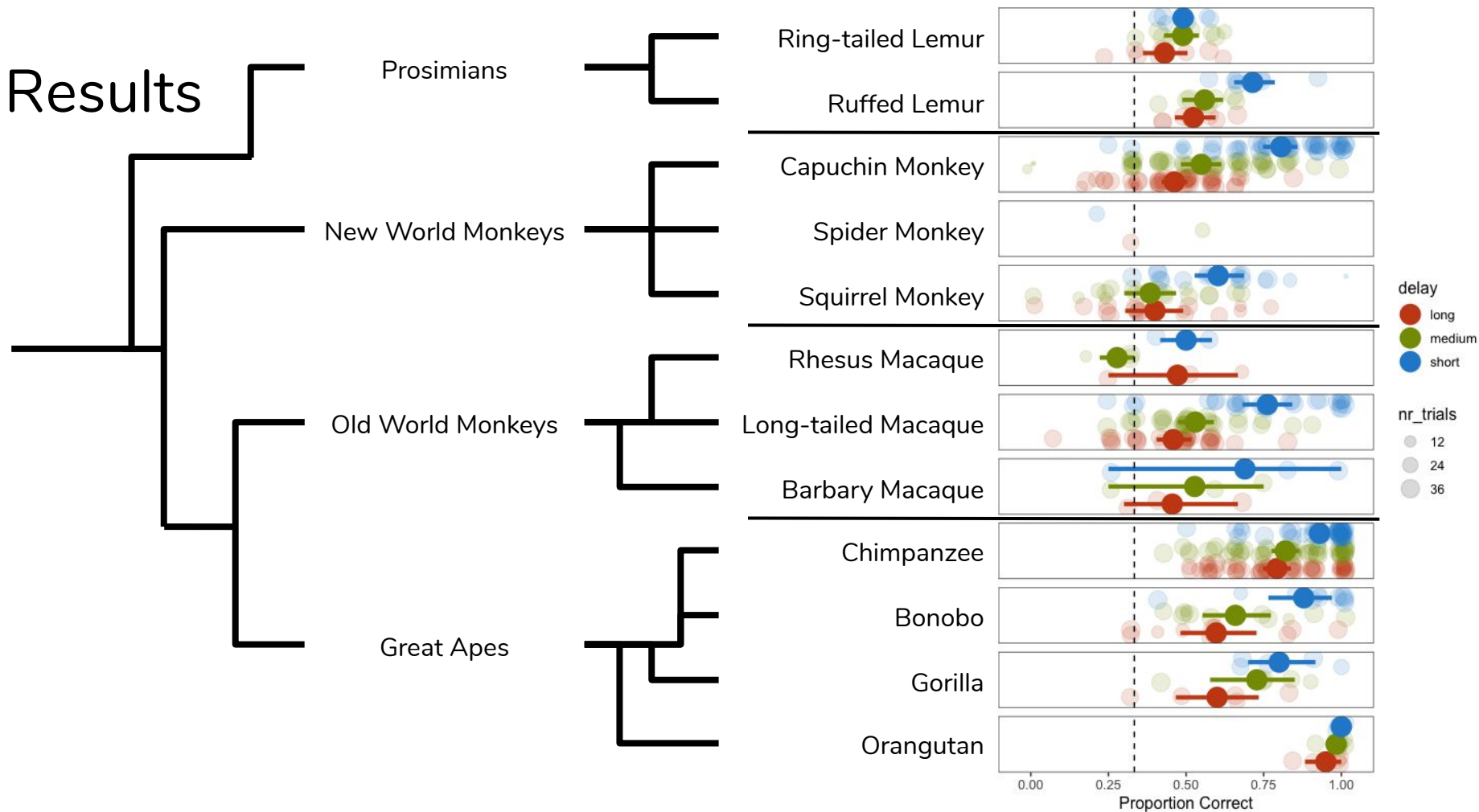


Sample

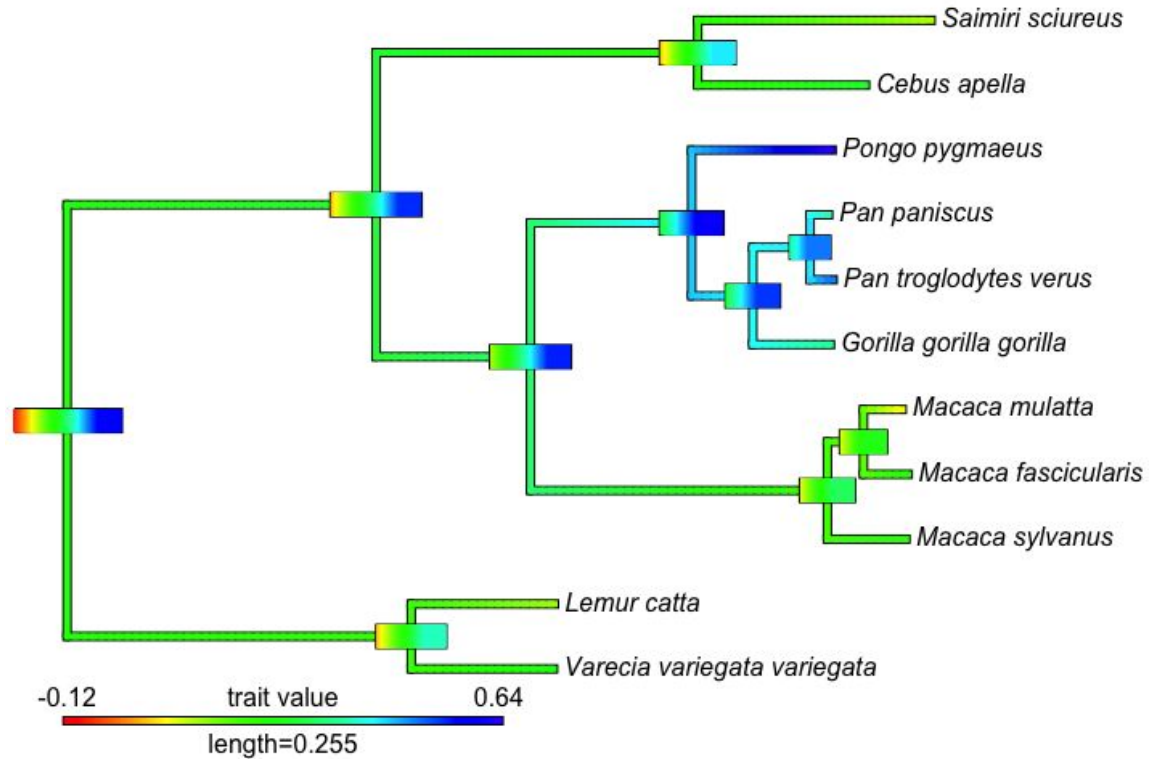


Species	N	Sites
Ring-tailed Lemur	9	2
Ruffed Lemur	7	1
Capuchin Monkey	46	2
Spider Monkey	1	1
Squirrel Monkey	18	1
Rhesus Macaque	3	1
Long-tailed Macaque	28	1
Barbary Macaque	3	1
Chimpanzee	51	5
Bonobo	11	2
Gorilla	5	2
Orangutan	5	1

Results



Phylogenetic analysis



ManyPrimates

Open for people without direct access to primates

- Organize / support studies
- Contribute to design, analysis and writing
- ECR especially encouraged to join!

Follow open science practises

- Pre-register methods and analysis
- Sharing data and code
- Publish open access

Challenges

Adjustment of test layout to site/species specific needs

- Statistical covariation: Species not a privileged variable

→ *Solution*: Phylogenetic analysis

Ad-hoc decisions (gaps in pre-registration)

→ *Solution*: Pilot data collection / Thinking hard in advance!