Table 1: Summary of Prey Frequency by Infraorder

prey	haplorrhini	strepsirrhini
amphibia	47	3
aves	147	13
mammalia	156	4
squamata	65	10
teleostei	13	0
vertebrata	3	0

INTRODUCTION

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CARNIVORY ACROSS PRIMATES

Watts (2020) reviewed meat eating across primates.

 $\mbox{\tt \#\#}$ 'summarise()' has grouped output by 'prey'. You can override using the $\mbox{\tt \#\#}$ '.groups' argument.

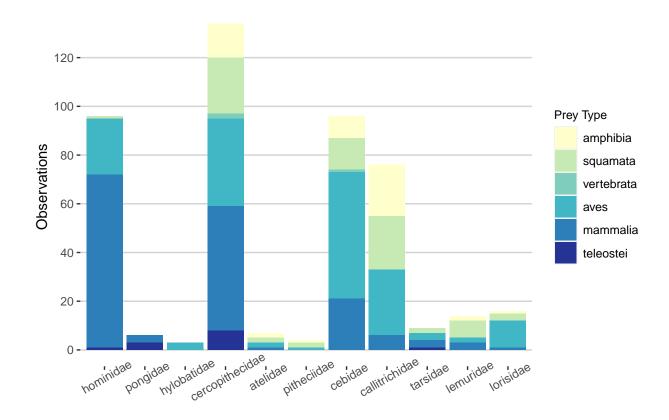


Table 2: Inconsistent Tool Use

use_tools	haplorrhini	strepsirrhini	NA
0	131	45	1
1	37	0	0

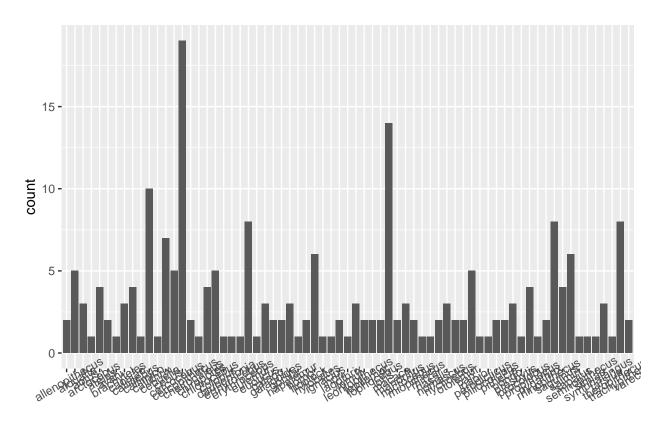
Table 3: Inconsistent Tool Use

use_tools	haplorrhini	strepsirrhini	NA
0	131	45	1
1	37	0	0

TOOL USE ACROSS PRIMATES

variable into a factor?

```
summary_tool <- all_df %>%
  group_by(use_tools, infraorder) %>%
  summarize(users = n()) %>%
  pivot_wider(names_from = infraorder, values_from = users, values_fill = 0)
## 'summarise()' has grouped output by 'use_tools'. You can override using the
## '.groups' argument.
kable(summary_tool, caption = "Inconsistent Tool Use") %>%
  kable_styling(bootstrap_options = "striped")
summary_tool2 <- all_df %>%
  group_by(use_tools, genus) %>%
  summarize(users = n()) %>%
 pivot_wider(names_from = genus, values_from = users, values_fill = 0)
## 'summarise()' has grouped output by 'use_tools'. You can override using the
## '.groups' argument.
kable(summary_tool, caption = "Inconsistent Tool Use") %>%
  kable_styling(bootstrap_options = "striped")
p <- ggplot(data = all_df, aes(x=genus, fill=use_tools)) +</pre>
  theme(axis.text.x = element_text(angle = 30)) +
  geom_bar()
p
## Warning: The following aesthetics were dropped during statistical transformation: fill
## i This can happen when ggplot fails to infer the correct grouping structure in
   the data.
## i Did you forget to specify a 'group' aesthetic or to convert a numerical
```



genus

METHODS

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