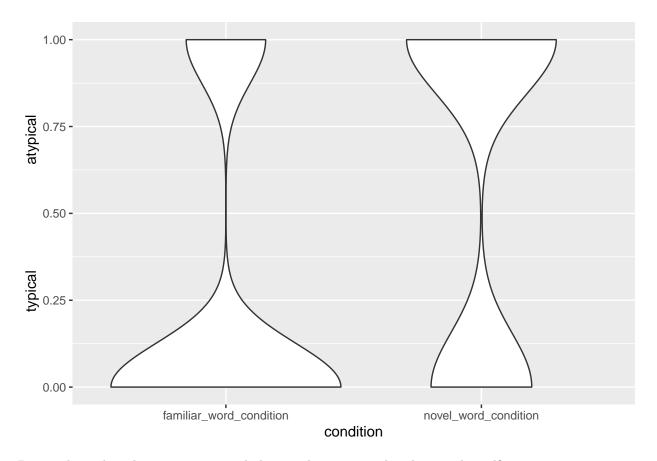
pilot analysis

2025-03-12

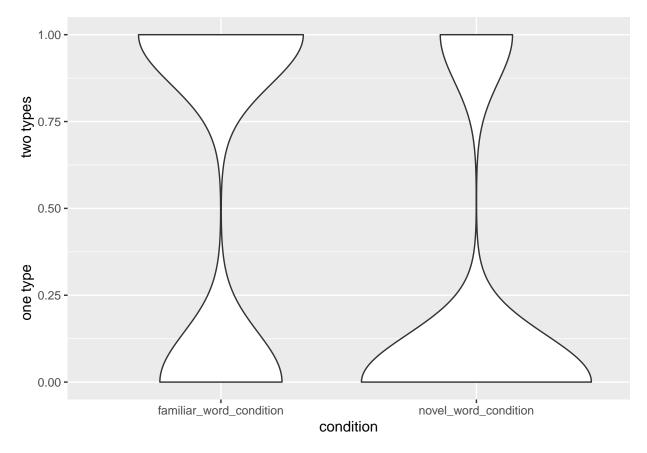
```
raw_data <- list.files( path=here("../data/pilot_raw_data"), full.names=TRUE ) %>%
  map_dfr( read_csv )
data <- raw_data %>%
  filter("study_id" != "UNSAVED_STUDY",
         !is.na(study_id)) %>%
  select(-prolific_id)
write_csv(data, here("../data/pilot_data.csv"))
data <- read_csv(here("../data/pilot_data.csv")) %>%
  mutate(stimulus_subclass = str_remove(image_name, "^[^_]*_"),
         stimulus_subclass = str_remove(stimulus_subclass, "_.*")) %>%
  mutate(typicality = typicality - 1)
data %>%
  ggplot(aes(x = condition, y = typicality)) +
  geom_violin() +
 ylab("typical
                                         atypical")
```



Do people tend to choose more atypical objects when prompted with a novel word?

Participants' choices on familiar word (e.g., "leaf") vs. novel word (e.g., "neft") trials. Higher values indicate they chose more atypical items. Stimuli typicality scored by experimenter on a binary scale. People choose more atypical exemplars when prompted with a novel compared to a familiar word. They also strongly prefer typical exemplars when prompted with familiar words.

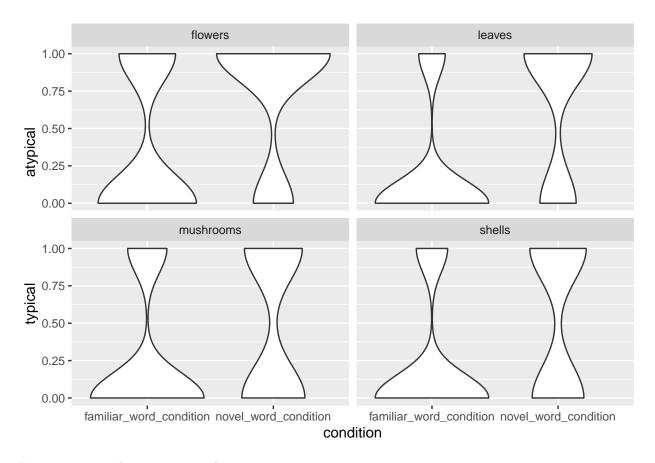
```
glmer(typicality ~ condition + (1|participant_id), family = "binomial",
      data = data) %>%
  tidy()
## # A tibble: 3 x 7
##
     effect
              group
                             term
                                              estimate std.error statistic
                                                                              p.value
     <chr>
              <chr>
                              <chr>
                                                 <dbl>
                                                           <dbl>
                                                                      <dbl>
                                                                                <dbl>
## 1 fixed
              <NA>
                              (Intercept)
                                                -1.16
                                                                      -6.54 6.28e-11
                                                            0.177
## 2 fixed
              <NA>
                             conditionnovel~
                                                 1.59
                                                           0.271
                                                                       5.85 5.00e- 9
## 3 ran_pars participant_id sd__(Intercept)
                                                 0.616
                                                          NA
                                                                      NA
                                                                            NA
data <- data %>%
  group_by(participant_id, trial_number) %>%
  mutate(distinct_types = n_distinct(stimulus_subclass) - 1) %>%
  ungroup()
data %>%
  ggplot(aes(x = condition, y = distinct_types)) +
  geom_violin() +
 ylab("one type
                                           two types")
```



Do people tend to choose more varied objects when prompted with familiar vs. novel words? Another way to tell whether people think the meaning of a novel word is narrower is to test whether they choose one subtype (e.g., bean-shaped leaves) vs. multiple subtypes (e.g., teardrop-shaped and oak-shaped leaves).

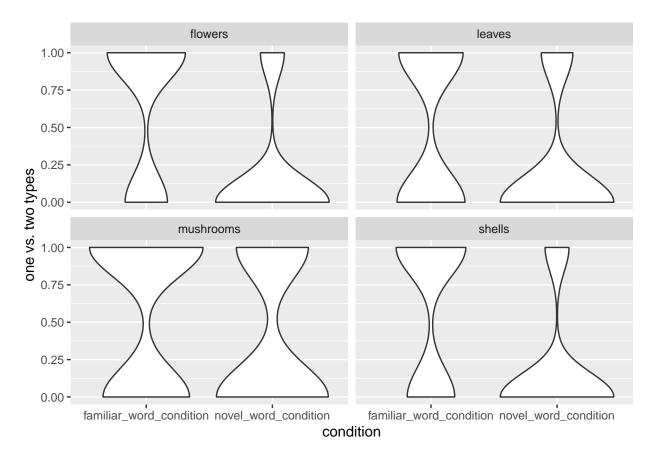
Above, participants' choices of exemplars; they have two choices on each trial. A score of 0 reflects that they chose all of one subtype, and a score of 1 reflects that they chose two distinct subtypes. People tend to choose fewer distinct subtypes when prompted with novel words.

```
glmer(distinct_types ~ condition + (1|participant_id), family = "binomial",
      data = data) %>%
  tidy()
## # A tibble: 3 x 7
     effect
              group
                              term
                                                estimate std.error statistic
                                                                               p.value
##
     <chr>>
              <chr>>
                              <chr>>
                                                   <dbl>
                                                              <dbl>
                                                                        <dbl>
                                                                                  <dbl>
## 1 fixed
              <NA>
                              (Intercept)
                                                   0.511
                                                             0.498
                                                                         1.03
                                                                               3.05e-1
              <NA>
                                                                        -3.29 9.97e-4
## 2 fixed
                              conditionnovel_~
                                                  -2.95
                                                             0.897
## 3 ran_pars participant_id sd__(Intercept)
                                                   2.70
                                                            NA
                                                                        NA
                                                                              NA
data %>%
  ggplot(aes(x = condition, y = typicality)) +
  geom_violin() +
  facet_wrap(~category) +
                                           atypical")
  ylab("typical
```



Examining typicality across stimulus types.

```
data %>%
   ggplot(aes(x = condition, y = distinct_types)) +
   geom_violin() +
   facet_wrap(~category) +
   ylab("one vs. two types")
```



Examining distinct type choices across stimulus types.