Syntactic: Reaction Time Graphs

morgan moyer

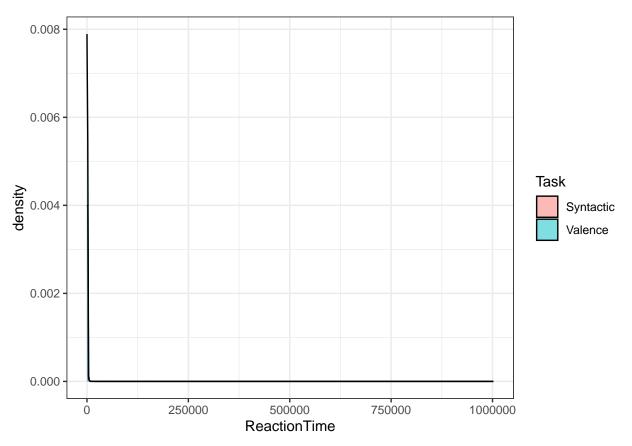
2025-05-04

Looking at overall Log ReactionTime for the data

Before removing outliers

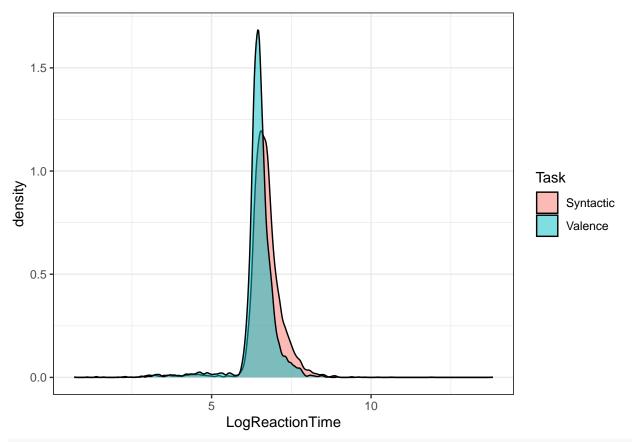
Summary Stats

```
agr <- d %>%
 group_by(Task) %>%
 summarize(MeanRT = mean(ReactionTime),
           SD = sd(ReactionTime),
           MeanLogRT = mean(LogReactionTime))
print(agr)
## # A tibble: 2 x 4
   Task MeanRT
                        SD MeanLogRT
   <chr>
              <dbl> <dbl>
                               <dbl>
## 1 Syntactic 1201. 15018.
                                6.69
## 2 Valence
              739.
                       457.
                                6.47
ggplot(d, aes(ReactionTime, fill=Task)) +
 geom_density(alpha = .5)
```



Long tail justifies outlier removal?

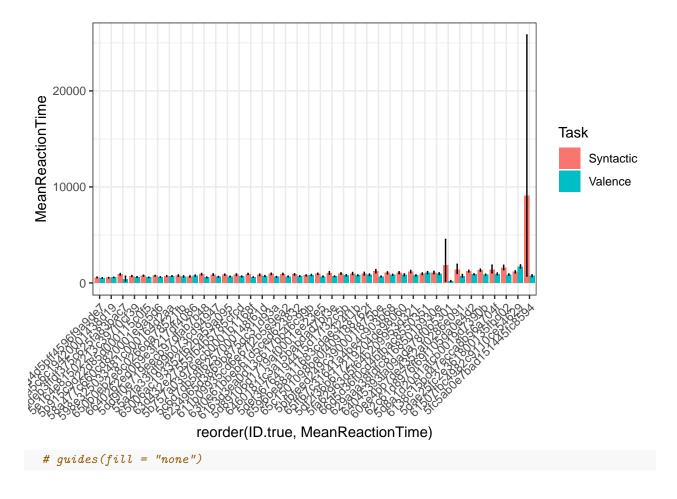
```
ggplot(d, aes(LogReactionTime, fill=Task)) +
geom_density(alpha = .5)
```



summary(d\$LogReactionTime)

`summarise()` has grouped output by 'ID.true'. You can override using the
`.groups` argument.

```
dodge = position_dodge(.9)
ggplot(data=agr, aes(x=reorder(ID.true,MeanReactionTime),y=MeanReactionTime,fill=Task)) +
    geom_bar(position=dodge,stat="identity") +
    # facet_wrap(~Word) +
    geom_errorbar(aes(ymin=YMin,ymax=YMax),width=.25,position=position_dodge(0.9)) +
    theme(axis.text.x = element_text(angle = 45, hjust = 1))
```



First Remove participants who aren't super , aggregating over Task

```
length(unique(d$ID.true))
## [1] 38
inacc.parts <- d %>%
  group_by(ID.true,Task) %>%
  summarise(MeanAccuracy = mean(Accuracy)) %>%
  filter(MeanAccuracy < .75)</pre>
## `summarise()` has grouped output by 'ID.true'. You can override using the
## `.groups` argument.
# How many participants have Accuracy < .75?
length(unique(inacc.parts$ID.true))
## [1] 12
d.inaccurate.removed <- d %>%
  anti_join(inacc.parts, by = "ID.true")
# Sanity check
length(unique(d.inaccurate.removed$ID.true))
## [1] 26
```

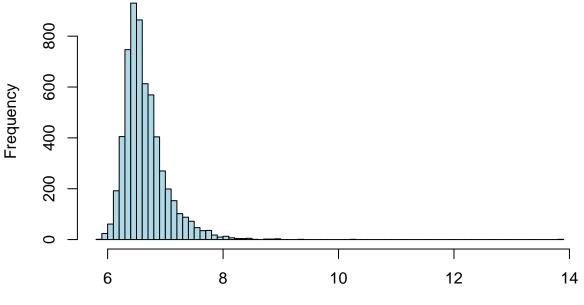
By Participant

```
agr <- d.inaccurate.removed %>%
   group_by(ID.true,Task) %>%
   summarize(MeanReactionTime = mean(ReactionTime),
            CILow = ci.low(ReactionTime),
            CIHigh = ci.high(ReactionTime)) %>%
   mutate(YMin = MeanReactionTime - CILow,
         YMax = MeanReactionTime + CIHigh)
## `summarise()` has grouped output by 'ID.true'. You can override using the
## `.groups` argument.
dodge = position dodge(.9)
ggplot(data=agr, aes(x=reorder(ID.true,MeanReactionTime),y=MeanReactionTime,fill=Task)) +
 geom_bar(position=dodge,stat="identity") +
 # facet_wrap(~Word) +
 geom_errorbar(aes(ymin=YMin,ymax=YMax),width=.25,position=position_dodge(0.9)) +
 theme(axis.text.x = element text(angle = 45, hjust = 1))
   20000
MeanReactionTime
                                                                     Task
                                                                         Syntactic
                                                                         Valence
    0000
                                MeanRer
                    reorder
                            reorder(ID.true, MeanReactionTime)
 # guides(fill = "none")
```

remove all inaccurate trials

```
orig <- nrow(d.inaccurate.removed)
d.inaccurate.removed <- d.inaccurate.removed %>%
```

```
filter(Accuracy == 1)
nrow(d.inaccurate.removed)/orig*100
## [1] 94.29487
# Remove subjects with ReactionTime higher than 3x IQR
summary(d.inaccurate.removed$LogReactionTime)
##
     Min. 1st Qu. Median
                             Mean 3rd Qu.
                                             Max.
    5.855
            6.404
                    6.567
                            6.647
                                    6.801 13.818
 # Min. 1st Qu. Median
                            Mean 3rd Qu.
                                            Max.
                                   7.579 10.008
           7.328
                   7.436
                           7.479
range(d.inaccurate.removed$LogReactionTime)
## [1] 5.855072 13.817820
hist(d.inaccurate.removed$LogReactionTime, breaks=100, col="lightblue", xlab="LogReactionTime (ms)",
       main="Histogram with Normal Curve")
                           Histogram with Normal Curve
```



```
LogReactionTime (ms)

quantile(d.inaccurate.removed$LogReactionTime, na.rm = TRUE)

## 0% 25% 50% 75% 100%

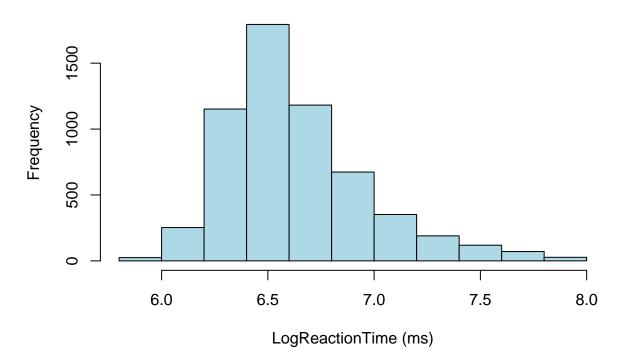
## 5.855072 6.403574 6.566672 6.801283 13.817820

IQR(d.inaccurate.removed$LogReactionTime, na.rm = TRUE)*3 # 0.7526289

## [1] 1.193127
```

cutoff.high <- quantile(d.inaccurate.removed\$LogReactionTime, na.rm = TRUE)[4] + IQR(d.inaccurate.removecutoff.low <- quantile(d.inaccurate.removed\$LogReactionTime, na.rm = TRUE)[2] - IQR(d.inaccurate.removecutoff.low <- quantile(d.inaccurate.removed\$LogReactionTime, na.rm = TRUE)[2] - IQR(d.inaccurate.removecutoff.low <- quantile(d.inaccurate.removed\$LogReactionTime, na.rm = TRUE)[2] - IQR(d.inaccurate.removecutoff.low <- quantile(d.inaccurate.removed\$LogReactionTime, na.rm = TRUE)[4] + IQR(d.inaccurate.removed\$LogReactionTime, na.rm = TRUE)[4] + IQR(d.inaccurate.removed\$Log

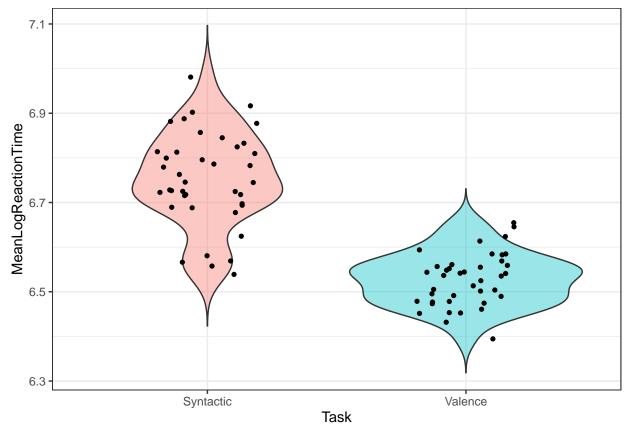
Histogram with Normal Curve



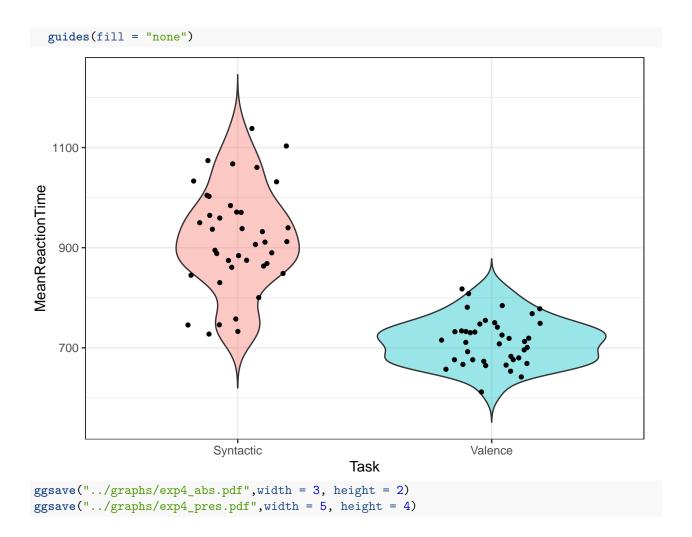
Summary Stats

```
agr <- d.inaccurate.removed %>%
  group_by(Task) %>%
  summarize(MeanRT = mean(ReactionTime),
            SD = sd(ReactionTime),
            MeanLogRT = mean(LogReactionTime))
print(agr)
## # A tibble: 2 x 4
##
     Task
               MeanRT
                           SD MeanLogRT
     <chr>>
                <dbl>
                                  <dbl>
                       <dbl>
## 1 Syntactic 1314. 18701.
                                   6.77
## 2 Valence
                 720.
                         308.
                                   6.53
```

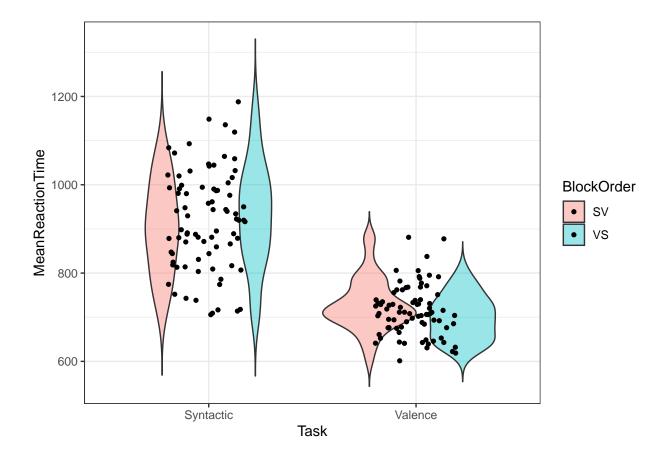
LogReactionTime by Task



ReactionTime by Task



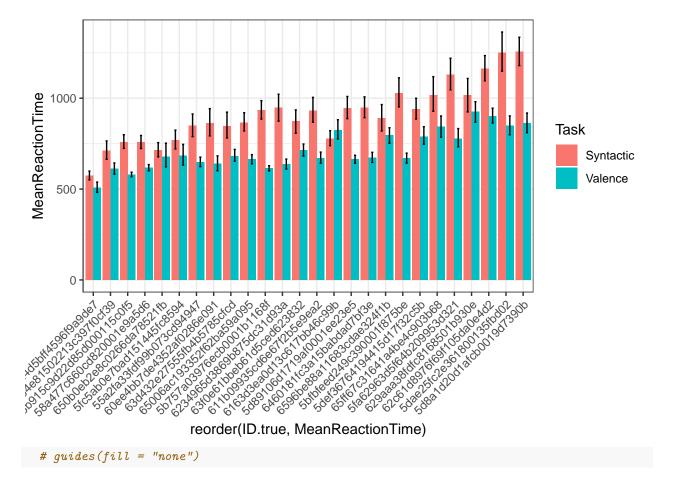
ReactionTime by BlockOrder and Task



By Item

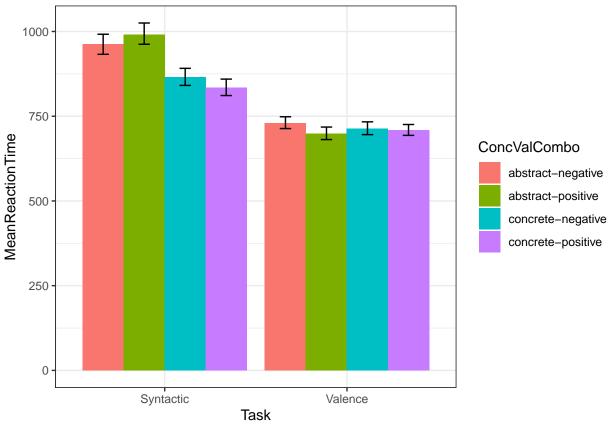


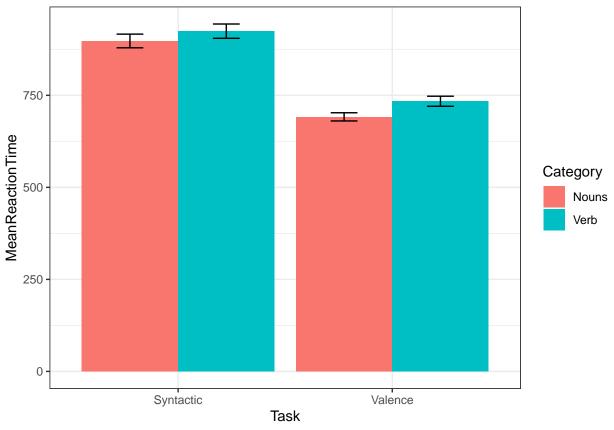
By Participant



By ConcValCombo category and Task

Mean Raw ReactionTime and Effects of Word Valence/Syntacticness





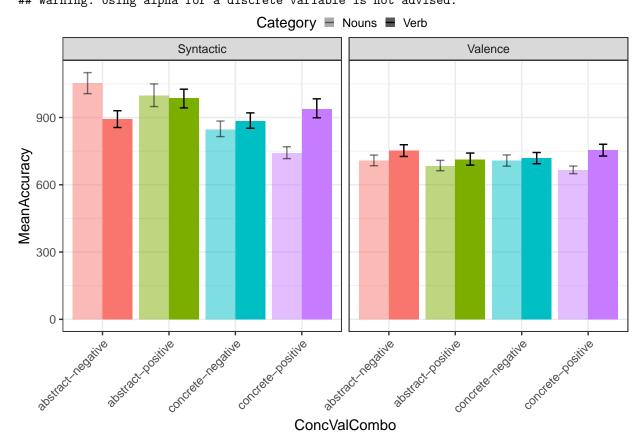
```
# theme(axis.text.x = element_text(angle = 45, hjust = 1))
# guides(fill = "none")
```

```
agr <- df.outliers.removed %>%
   group_by(Task,ConcValCombo,Category) %>%
   reframe(MeanReactionTime = mean(ReactionTime),
            CILow = ci.low(ReactionTime),
            CIHigh = ci.high(ReactionTime)) %>%
   mutate(YMin = MeanReactionTime - CILow,
           YMax = MeanReactionTime + CIHigh)
ggplot(agr,aes(x=ConcValCombo, y=MeanReactionTime, alpha=Category, fill=ConcValCombo)) +
  geom_bar(position="dodge",stat="identity") +
  geom_errorbar(aes(ymin=YMin,ymax=YMax),width=.25,position=position_dodge(0.9)) +
  facet_wrap(~Task, ncol=10) +
  xlab("ConcValCombo") +
  ylab("MeanAccuracy") +
  guides(fill=FALSE) +
  # guides(alpha=guide_legend(title="Task")) +
  theme(legend.key.size = unit(0.3, "cm"),
        legend.position = "top", # c(.5,1)
        legend.direction = "horizontal",
        legend.margin=margin(0,0,0,0),
        legend.box.margin=margin(0,0,-5,-5),legend.spacing.y = unit(0.001, 'cm')) +
    # scale fill manual(values=cbPalette) +
    # scale_color_manual(values=cbPalette) +
    scale_alpha_discrete(range = c(.5,1)) +
```

```
theme(axis.text.x = element_text(angle = 45, hjust = 1))
```

```
## Warning: The `<scale>` argument of `guides()` cannot be `FALSE`. Use "none" instead as
## of ggplot2 3.3.4.
## This warning is displayed once every 8 hours.
## Call `lifecycle::last_lifecycle_warnings()` to see where this warning was
```

Warning: Using alpha for a discrete variable is not advised.

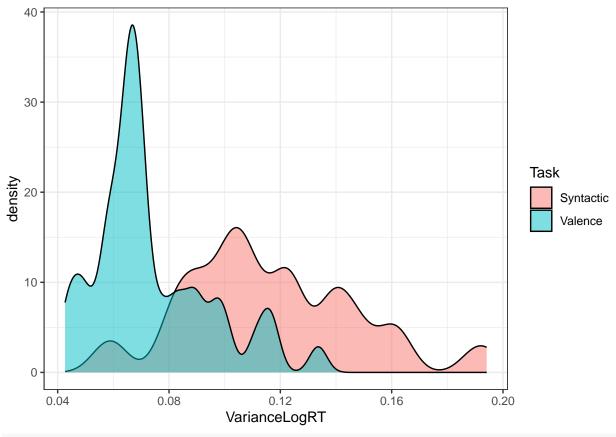


Variance

generated.

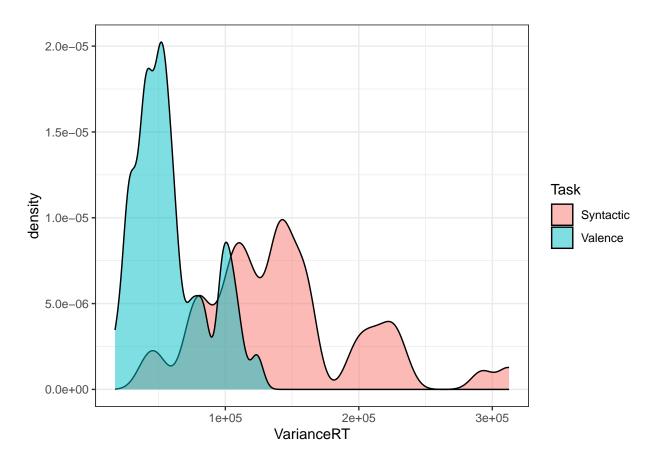
```
agr <- df.outliers.removed %>%
  group_by(Task,Word) %>%
  mutate(VarianceLogRT = var(LogReactionTime))

ggplot(agr, aes(VarianceLogRT, fill=Task)) +
  geom_density(alpha = .5)
```

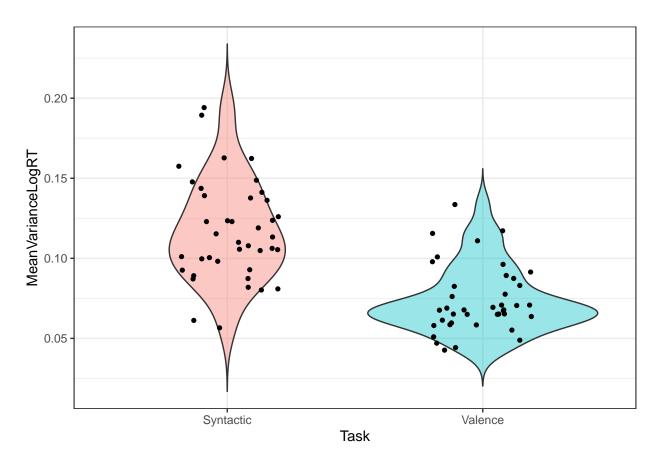


```
agr <- df.outliers.removed %>%
  group_by(Task,Word) %>%
  mutate(VarianceRT = var(ReactionTime))

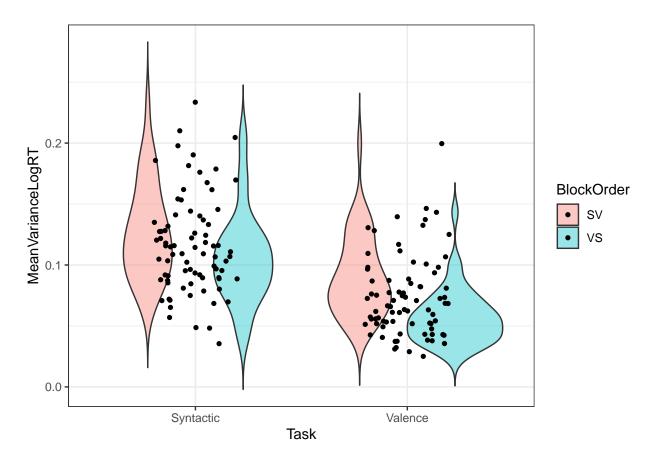
ggplot(agr, aes(VarianceRT, fill=Task)) +
  geom_density(alpha = .5)
```



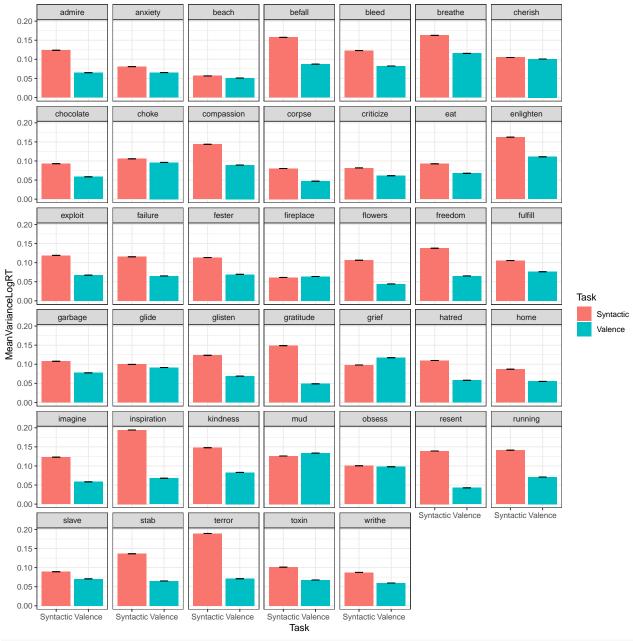
ReactionTime by Task



LogReactionTime by BlockOrder and Task



By Item



```
# theme(axis.text.x = element_text(angle = 45, hjust = 1))
# guides(fill = "none")
```