

Baby_BP_pilot

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To help doctors get the most out of working with a biostats hospital hire, here are some template thoughts...

For project XXX and its main objectives, can you provide me with..

1. descriptive figures to summarize data/get-to-know variables. Ok to send before main objective results.

i.e. distribution on continuous variables and counts for categorical. this should be a baseline. doctors dont always know what to ask for and biostats person, as part of their job, should be providing more than whats requested (otherwise biostats person is no better than AI and can easily be replaced).

2. your interpretation of all results/plots-in simple terms tell me what the plot shows and what you concluded solely based on the figure

this is especially important for main objective plots. citations related to plot type should be provided with biostats report (i.e. citation for bp plot)

3. your concerns with study design and/or results, and any suggestions for improvements. You are the expert and i would greatly appreciate any feedback!

not all concerns can be addressed but doctors should at least have an idea of what biostats person thinks-is the study design sufficient to address main objective? are their limitations? if some adjustments can be made to better address project aims, is it going to require an IRB update? is it reasonable within a timeline discussed for project? i.e. not having enough power to make claim like wrist cuffs are correlated with gold standard-a power analysis can tell you if the current study design will be sufficient to address objectives. i.e. a requested plot had different number of data points (26 vs 29 for all other correlation plots)-why? following # 2 point of template will prevent questions like this in future

4. responses to any of my questions within 48 hours. need more time and will get back to you soon is valid response.

doctors are likely interested in what biostats person makes, so it should be expected that more questions will follow. however, if the biostats person follows template 1-3 points, less questions and discussion may be required because everything is included in report! its a good point for them to invest in doing template points 1-3, if they don't, then template point 4 should absolutely be required. weeks to send any results and being slow to respond to questions from doctors who have looked over results hinders doctors ability to apply their expertise to analysis, get more thorough conclusions from data they have invested time and effort in collecting, and ultimately limits the ability for studies to improve quality of patient care. no excuse.

This markdown is an example of following template points 1-3. it includes basic data summary figures, redundancy of main objective related results (same point made with multiple plot types), and short interpretation following a result (what i thought about table or figure result).

Time spent on this markdown = 1 day.

note - I am not a formally trained biostats person. i would expect the analysis provided by biostats person for hospital to do a MUCH better/more thorough job than this. please consider this a benchmark for minimum effort.

```
## made df for histogram by combining sys and dia into single column. made from Baby_
blood_pressure.csv by copy/paste in excel.
df2 <- read.csv("Baby_blood_pressure_redundant.csv", stringsAsFactors=FALSE)
df2$BabyID <- factor(df2$BabyID)
df2$Session <- factor(df2$Session)
df2[['Device']] <- factor(df2[['Device']], levels = c(
  'Gold',
  'Wristcuff'
))
df2[['Rater']] <- factor(df2[['Rater']], levels = c(
  'RN',
  'Gibbs',
  'Not Gibbs'
))
df2$Device <- df2$Rater
df2[['Device']] <- factor(df2[['Device']],
                        labels = c("Gold standard", "Wrist cuff 1", "Wrist cuff 2")
)
df2[['Type']] <- factor(df2[['Type']], levels = c(
  'Systolic',
```

```

'Diastolic'
))
df2$Disposition <- df2$Dispo
df2[['Disposition']] <- factor(df2[['Disposition']],
                              levels = c(0,1,2,3),
                              labels = c("sleeping", "calm", "active", "crying")
)

## df for all other plots and stats
df <- read.csv("Baby_blood_pressure.csv", stringsAsFactors=FALSE)
df$BabyID <- factor(df$BabyID)
df$Session <- factor(df$Session)
df[['Device']] <- factor(df[['Device']], levels = c(
  'Gold',
  'Wristcuff'
))
df[['Rater']] <- factor(df[['Rater']], levels = c(
  'RN',
  'Gibbs',
  'Not Gibbs'
))
df$Device <- df$Rater
df[['Device']] <- factor(df[['Device']],
                        labels = c("Gold standard", "Wrist cuff 1", "Wrist cuff 2")
)
df$Disposition <- df$Dispo
df[['Disposition']] <- factor(df[['Disposition']],
                              levels = c(0,1,2,3),
                              labels = c("sleeping", "calm", "active", "crying")
)
drops <- c("Dispo")
df <- df[ , !(names(df) %in% drops)]

# CONSTANTS used in cells below
#rater_colors = c("gold", "#d88c80", "#9c81ff")

wes_palettes <- list(
  BottleRocket1 = c("#A42820", "#5F5647", "#9B110E", "#3F5151", "#4E2A1E", "#550307",
"#0C1707"),
  BottleRocket2 = c("#FAD510", "#CB2314", "#273046", "#354823", "#1E1E1E"),
  Rushmore1 = c("#E1BD6D", "#EABE94", "#0B775E", "#35274A", "#F2300F"),
  Rushmore = c("#E1BD6D", "#EABE94", "#0B775E", "#35274A", "#F2300F"),
  Royal1 = c("#899DA4", "#C93312", "#FAEFD1", "#DC863B"),
  Royal2 = c("#9A8822", "#F5CDB4", "#F8AFA8", "#FDDDA0", "#74A089"),
  Zissoul = c("#3B9AB2", "#78B7C5", "#EBCC2A", "#E1AF00", "#F21A00"),
  Darjeeling1 = c("#FF0000", "#00A08A", "#F2AD00", "#F98400", "#5BB6D6"),
  Darjeeling2 = c("#ECCBAE", "#046C9A", "#D69C4E", "#ABDDDE", "#000000"),
  Chevalier1 = c("#446455", "#FDD262", "#D3DDDC", "#C7B19C"),
  FantasticFox1 = c("#DD8D29", "#E2D200", "#46ACC8", "#E58601", "#B40F20"),

```

```

Moonrise1 = c("#F3DF6C", "#CEAB07", "#D5D5D3", "#24281A"),
Moonrise2 = c("#798E87", "#C27D38", "#CCC591", "#29211F"),
Moonrise3 = c("#85D4E3", "#F4B5BD", "#9C964A", "#CDC08C", "#FAD77B"),
Cavalcantil = c("#D8B70A", "#02401B", "#A2A475", "#81A88D", "#972D15"),
GrandBudapest1 = c("#F1BB7B", "#FD6467", "#5B1A18", "#D67236"),
GrandBudapest2 = c("#E6A0C4", "#C6CDF7", "#D8A499", "#7294D4"),
IsleofDogs1 = c("#9986A5", "#79402E", "#CCBA72", "#0F0D0E", "#D9D0D3", "#8D8680"),
IsleofDogs2 = c("#EAD3BF", "#AA9486", "#B6854D", "#39312F", "#1C1718"),
FrenchDispatch = c("#90D4CC", "#BD3027", "#B0AFA2", "#7FC0C6", "#9D9C85")
)

rater_colors = c("#DD8D29", "#E2D200", "#46ACC8") #c("#F2AD00", "#90D4CC", "#FD6467")

dispo_colors = c("#7294D4", "#C6CDF7", "#FD6467", "#5B1A18")

bp_colors = c("#FF0000", "#08519c")

compare_dispo <- list(c("sleeping", "calm"), c("sleeping", "active"), c("calm", "active"))
compare_raters <- list(c("RN", "Gibbs"), c("RN", "Not Gibbs"), c("Gibbs", "Not Gibbs"))
compare_devices <- list(c("Gold standard", "Wrist cuff 1"),
                        c("Gold standard", "Wrist cuff 2"),
                        c("Wrist cuff 1", "Wrist cuff 2"))

```

```
summary(df)
```

```

##      BabyID  Session      Device      Rater      Sys
##  2      : 9   1:39   Gold standard:29   RN      :29   Min.    : 49.00
##  3      : 9   2:30   Wrist cuff 1 :29   Gibbs    :29   1st Qu.: 63.50
##  4      : 9   3:18   Wrist cuff 2 :29   Not Gibbs:29   Median : 71.00
##  8      : 9                                     Mean    : 73.18
## 10      : 9                                     3rd Qu.: 82.00
## 12      : 9                                     Max.    :108.00
## (Other):33
##      Dia      Disposition
## Min.    :26.00   sleeping:19
## 1st Qu.:37.00   calm    :51
## Median :41.00   active  :16
## Mean    :42.74   crying  : 1
## 3rd Qu.:46.50
## Max.    :71.00
##

```

- result: babies 2,3,4,8,10,12 have 3 sessions. each rater has 29 data points for systolic and 29 for diastolic, total of 87 measurement points.

To assess device/rater measurement mean differences, i first used histograms to understand the variable distribution. this matters for picking the best stats test method to use to compare group means.

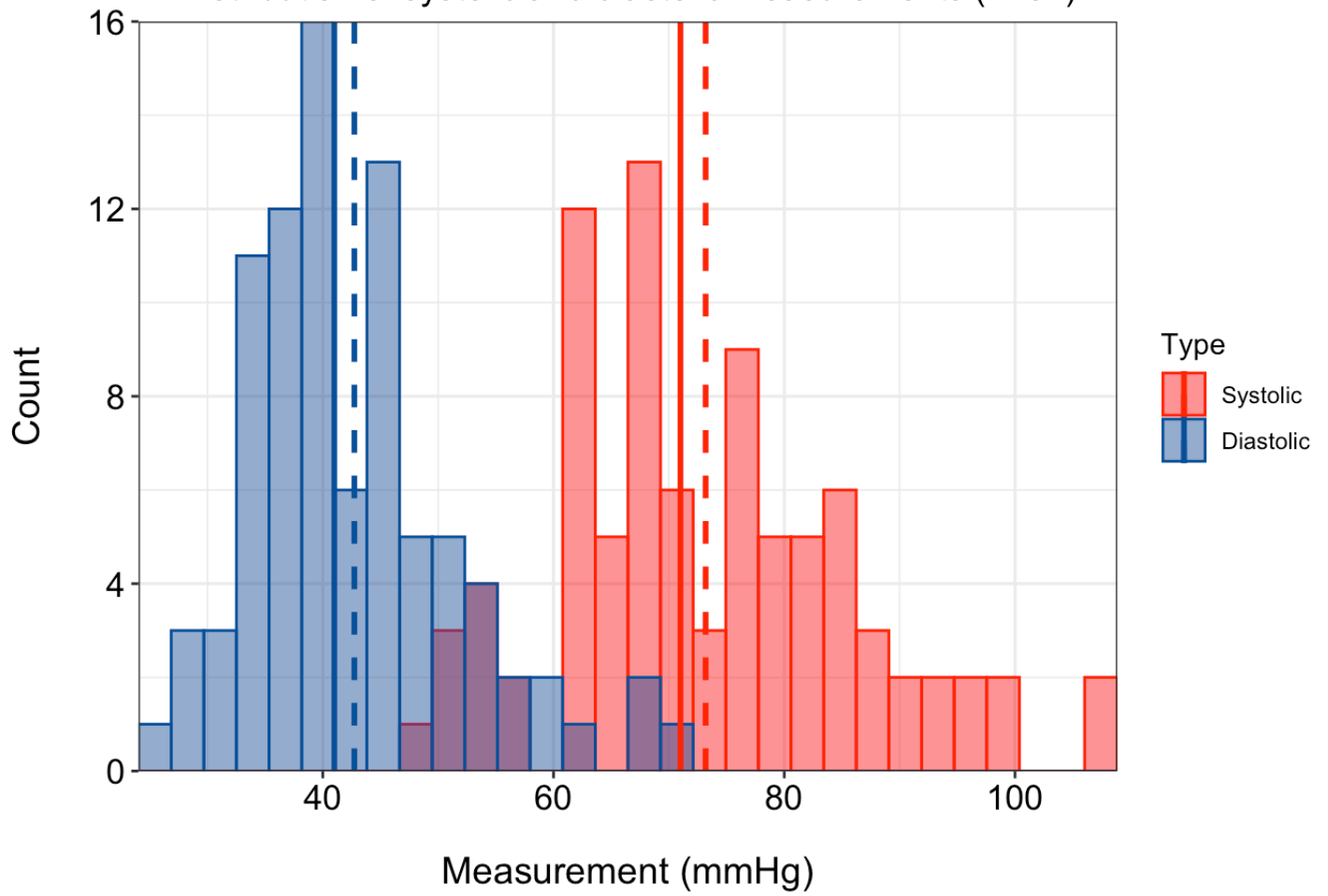
```
summary_mean_median <- ddply(df2, "Type", summarise,  
                             Mean=mean(Measurement),  
                             Median=median(Measurement))  
  
summary_mean_median
```

```
##           Type      Mean Median  
## 1 Systolic 73.18391      71  
## 2 Diastolic 42.73563      41
```

```
normalized_dist_check <- ggplot(df2, aes(x = Measurement, color=Type, fill = Type))  
  
hist_all <- normalized_dist_check +  
  geom_histogram(position="identity", alpha=0.5) + #bins=100, binwidth = 0.02 ,  
  geom_vline(data=summary_mean_median, aes(xintercept=Mean, color=Type), size=1, lin  
etype="dashed") +  
  geom_vline(data=summary_mean_median, aes(xintercept=Median, color=Type), size=1, li  
netype="solid") +  
  theme_bw() +  
  scale_fill_manual(values=bp_colors) +  
  scale_color_manual(values=bp_colors) +  
  scale_y_continuous(expand = c(0, 0)) +  
  scale_x_continuous(expand = c(0, 0)) +  
  ylab("Count") +  
  xlab("Measurement (mmHg)") +  
  ggtitle("Distribution of systolic and diastolic measurements (n=87)") +  
  theme(legend.position="right",  
        plot.title = element_text(hjust=0.5),  
        axis.title.x = element_text(size=14, color="black",  
                                     margin=margin(t=15, b=5)),  
        axis.title.y = element_text(size=14, color="black", margin=margin(t=0, r=15,  
b=0)),  
        axis.text=element_text(size=13,vjust=0.5, color="black", margin=margin(t=20,  
b=20)))  
hist_all
```

```
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```

Distribution of systolic and diastolic measurements (n=87)

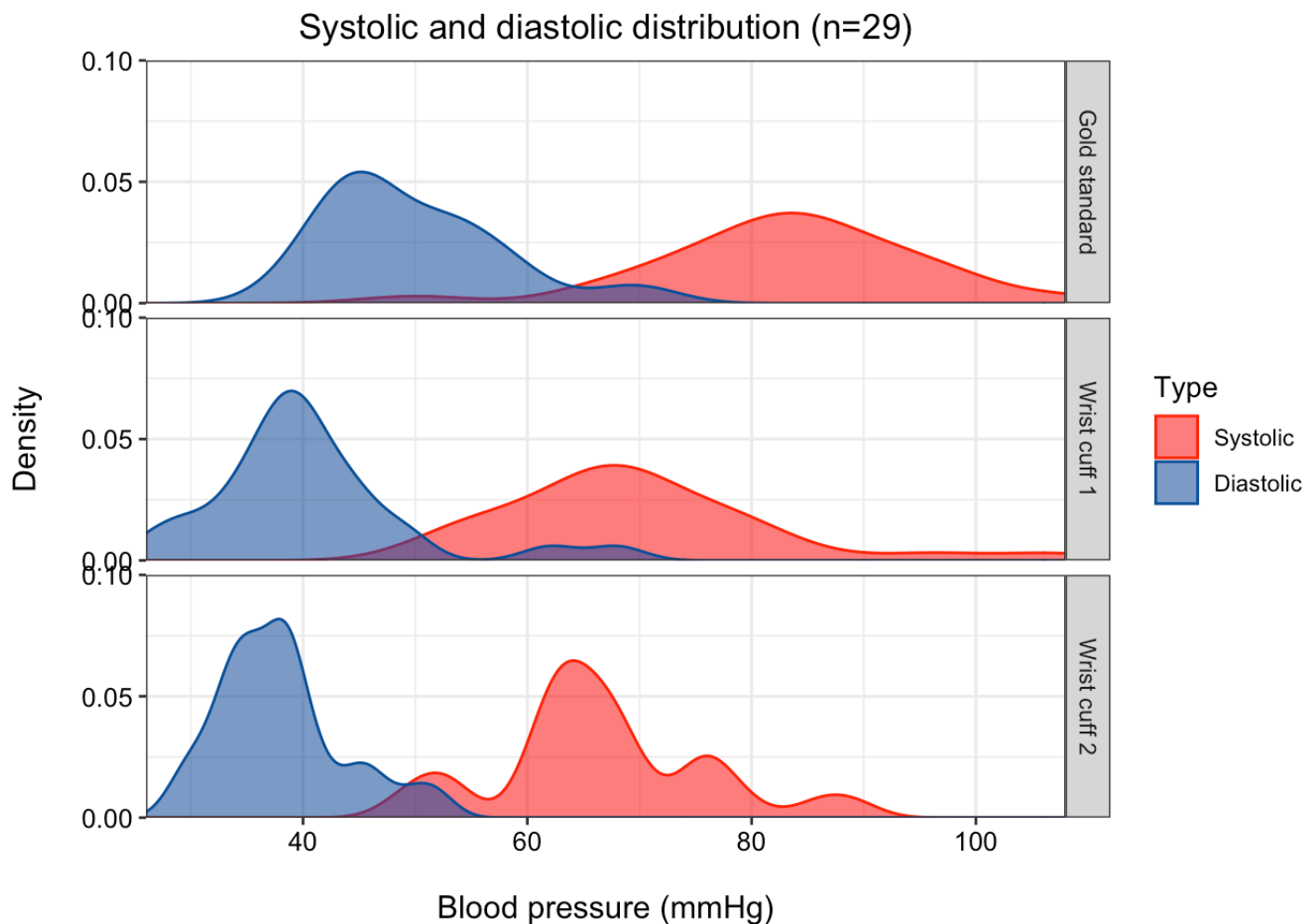


```

hist_wrap <- normalized_dist_check +
  geom_density(alpha=0.6) +
  #geom_histogram(bins=10, binwidth =1 ,position="identity", alpha=0.5) + #
  #geom_vline(data=summary_mean_median, aes(xintercept=Mean, color=Type), size=1, li
netype="dashed") +
  #geom_vline(data=summary_mean_median, aes(xintercept=Median, color=Type), size=1, l
inetype="solid") +
  theme_bw() +
  scale_fill_manual(values=bp_colors) +
  scale_color_manual(values=bp_colors) +
  scale_y_continuous(expand = c(0, 0), limits = c(0,0.1), n.breaks = 4) +
  scale_x_continuous(expand = c(0, 0)) +
  ylab("Density") +
  xlab("Blood pressure (mmHg)") +
  ggtitle("Systolic and diastolic distribution (n=29)") +
  theme(legend.position="right",
        plot.title = element_text(hjust=0.5),
        axis.title.x = element_text(size=12, color="black",
                                     margin=margin(t=15, b=5)),
        axis.title.y = element_text(size=12, color="black", margin=margin(t=0, r=15,
b=0)),
        axis.text=element_text(size=10,vjust=0.5, color="black", margin=margin(t=20,
b=20))) + facet_grid(Device ~ .)

hist_wrap

```



```
ggsave("density_systolic_diastolic_facetgrid_device.png", width=8, height=6, dpi=300)
```

- result: slight right skew in sys and dia distribution. mean (dashed line) slightly higher than median (solid line).
- question of whether to use parametric or nonparametric test- for all data points (n=87)?
- note on central limit theorem- if the sample size is large enough ($n > 30$), we can ignore the distribution of the data and use parametric tests. The central limit theorem tells us that no matter what distribution things have, the sampling distribution tends to be normal if the sample is large enough ($n > 30$).
- but since data is skewed, for comparisons of groups with $n < 30$, is parametric test ok?

```
df2_sys <- df2 %>% filter(Type == "Systolic")

summary_mean_median_sys <- ddpoly(df2_sys, "Device", summarise,
  Mean=mean(Measurement),
  Median=median(Measurement))

summary_mean_median_sys
```


##	Device	Mean	Median
## 1	Gold standard	83.17241	83
## 2	Wrist cuff 1	69.79310	68
## 3	Wrist cuff 2	66.58621	65

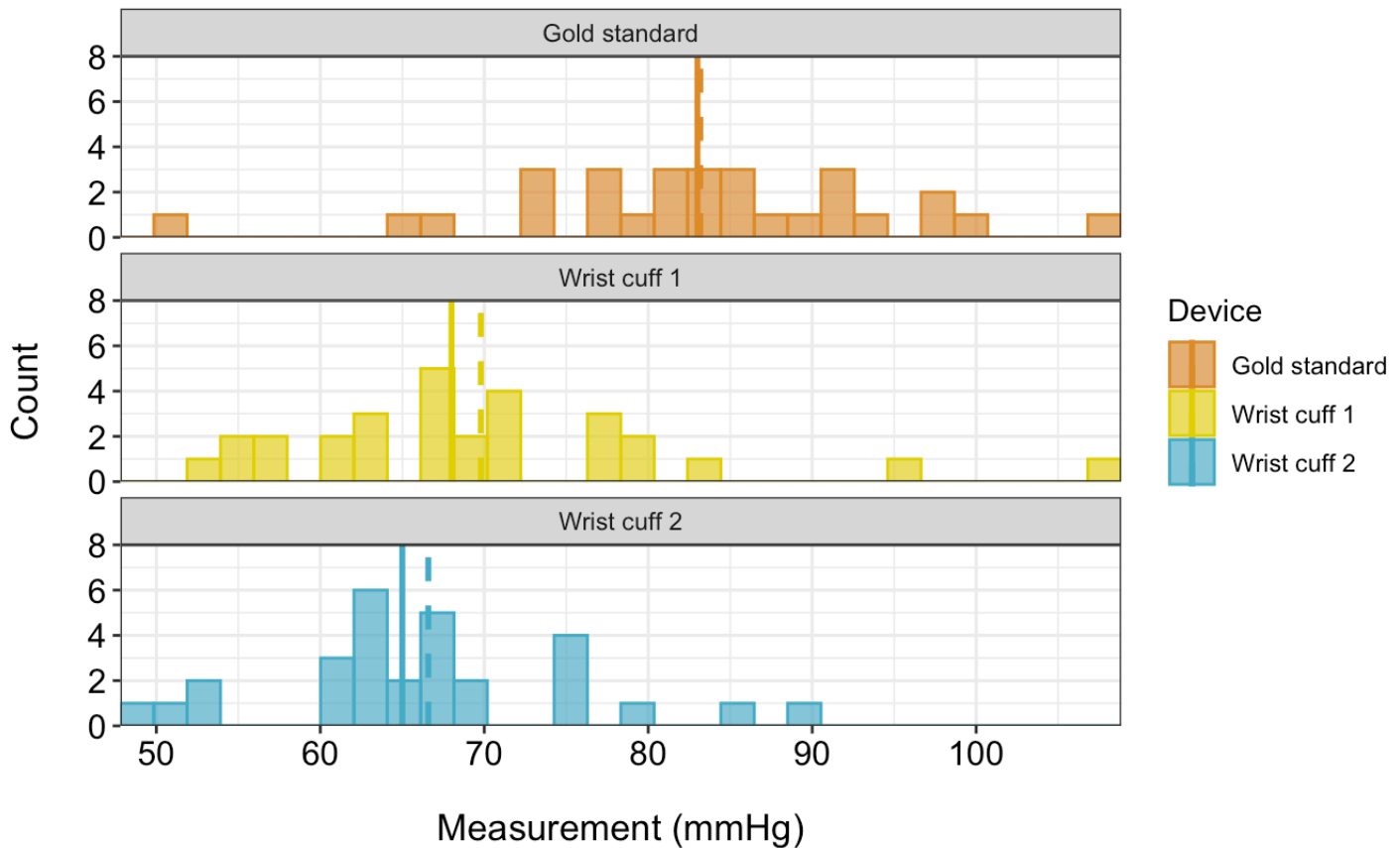
```
hist_sys_data <- ggplot(df2_sys, aes(x = Measurement, color=Device, fill = Device))

hist_raters <- hist_sys_data +
  geom_histogram(position="identity", alpha=0.7) +
  geom_vline(data=summary_mean_median_sys, aes(xintercept=Mean, color=Device), size=
1, linetype="dashed") +
  geom_vline(data=summary_mean_median_sys, aes(xintercept=Median, color=Device), size
=1, linetype="solid") +
  theme_bw() +
  scale_fill_manual(values=rater_colors) +
  scale_color_manual(values=rater_colors) +
  scale_y_continuous(expand = c(0, 0), limits = c(0,8), n.breaks = 5) +
  scale_x_continuous(expand = c(0, 0), n.breaks = 9) +
  labs(y="Count", x="Measurement (mmHg)",
    title="Systolic measurement counts",
    subtitle = "n = 29 measurements per device group",
    caption = "(dashed line marks mean, solid line marks median)") +
  theme(legend.position="right",
    #plot.title = element_text(hjust=0.5),
    axis.title.x = element_text(size=13, color="black",
      margin=margin(t=15, b=5)),
    axis.title.y = element_text(size=13, color="black", margin=margin(t=0, r=15,
b=0)),
    axis.text=element_text(size=12,vjust=0.5, color="black", margin=margin(t=20,
b=20)))
hist_raters = hist_raters + facet_wrap(~ Device, ncol=1)
hist_raters
```

```
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```

Systolic measurement counts

n = 29 measurements per device group



(dashed line marks mean, solid line marks median)

```
ggsave("histogram_systolic_per_device.png", width = 8, height = 6)
```

```
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```

- result = systolic RN/gold standard device is normally distributed (median = mean). systolic wrist cuff measurements skewed.

```
df2_sys <- df2 %>% filter(Type == "Diastolic")

summary_mean_median_sys <- dplyr::summarise(df2_sys, "Device", summarise,
  Mean=mean(Measurement),
  Median=median(Measurement))

summary_mean_median_sys
```

```
##           Device      Mean Median
## 1 Gold standard 49.79310     47
## 2 Wrist cuff 1 40.41379     39
## 3 Wrist cuff 2 38.00000     38
```

```

hist_sys_data <- ggplot(df2_sys, aes(x = Measurement, color=Device, fill = Device))

hist_raters <- hist_sys_data +
  geom_histogram(position="identity", alpha=0.7) +
  geom_vline(data=summary_mean_median_sys, aes(xintercept=Mean, color=Device), size=
1, linetype="dashed") +
  geom_vline(data=summary_mean_median_sys, aes(xintercept=Median, color=Device), size
=1, linetype="solid") +
  theme_bw() +
  scale_fill_manual(values=rater_colors) +
  scale_color_manual(values=rater_colors) +
  scale_y_continuous(expand = c(0, 0), limits = c(0,8), n.breaks = 5) +
  scale_x_continuous(expand = c(0, 0), n.breaks = 9) +
  labs(y="Count", x="Measurement (mmHg)",
       title="Diastolic measurement counts",
       subtitle = "n = 29 measurements per device group",
       caption = "(dashed line marks mean, solid line marks median)") +
  theme(legend.position="right",
        #plot.title = element_text(hjust=0.5),
        axis.title.x = element_text(size=13, color="black",
                                     margin=margin(t=15, b=5)),
        axis.title.y = element_text(size=13, color="black", margin=margin(t=0, r=15,
b=0)),
        axis.text=element_text(size=12,vjust=0.5, color="black", margin=margin(t=20,
b=20)))
hist_raters = hist_raters + facet_wrap(~ Device, ncol=1)
hist_raters

```

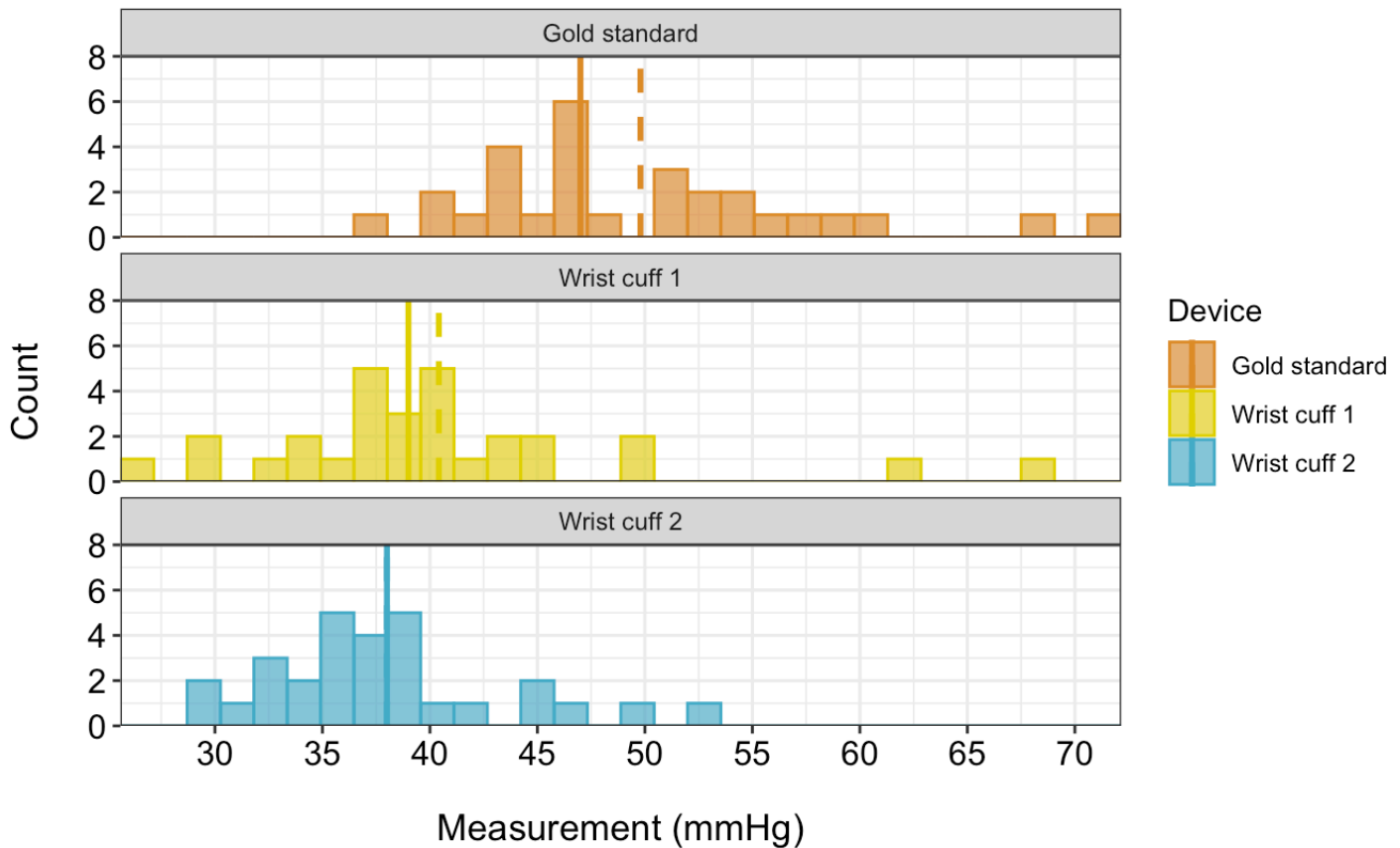
```

## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.

```

Diastolic measurement counts

n = 29 measurements per device group



(dashed line marks mean, solid line marks median)

```
ggsave("histogram_diastolic_per_device.png", width = 8, height = 6)
```

```
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```

- result: similar to systolic result for RN, the not gibbs diastolic result seems normally distributed (mean=median). largest skew observed for RN gold standard measurements for diastolic.

```
shapiro.test(df[['Sys']])
```

```

boxplot_3_comparison <- function(df, xname, yname, fillname, ylabel, yrange, titlename,
subname, capname, colorname, stat_table, stat_label) {
  bxplot <- ggboxplot(df, x = xname, y = yname, fill = fillname, alpha=0.8, notch=FALSE) +
  stat_pvalue_manual(stat_table, label = stat_label) +
  scale_fill_manual(values=colorname) +
  scale_y_continuous(n.breaks = 9, limits=yrange) +
  theme_bw() +
  labs(x=xname,
       y= ylabel,
       title = titlename,
       subtitle = subname,
       caption = capname) +
  theme(panel.grid.minor = element_blank()) +
  theme(legend.position='right') +
  theme(
    axis.title.x = element_text(size=14, color="black", margin=margin(t=15, b=
5)),
    axis.title.y = element_text(size=14, color="black", margin=margin(t=0, r=15,
b=0)),
    axis.text=element_text(size=12, color="black", margin=margin(t=20, b=20)))
  return(bxplot)
}

```

```

sys_test <- compare_means(Sys ~ Device,
                          data=df,
                          method="t.test",
                          p.adjust.method = "fdr",
                          paired=FALSE,
                          comparisons = compare_devices)
sys_test <- sys_test %>% mutate(y.position = c(105, 115, 110))
dia_test <- compare_means(Dia ~ Device,
                          data=df,
                          method="t.test",
                          p.adjust.method = "fdr",
                          paired=FALSE,
                          comparisons = compare_devices)
dia_test <- dia_test %>% mutate(y.position = c(68, 78, 73))
bind_rows(sys_test, dia_test)

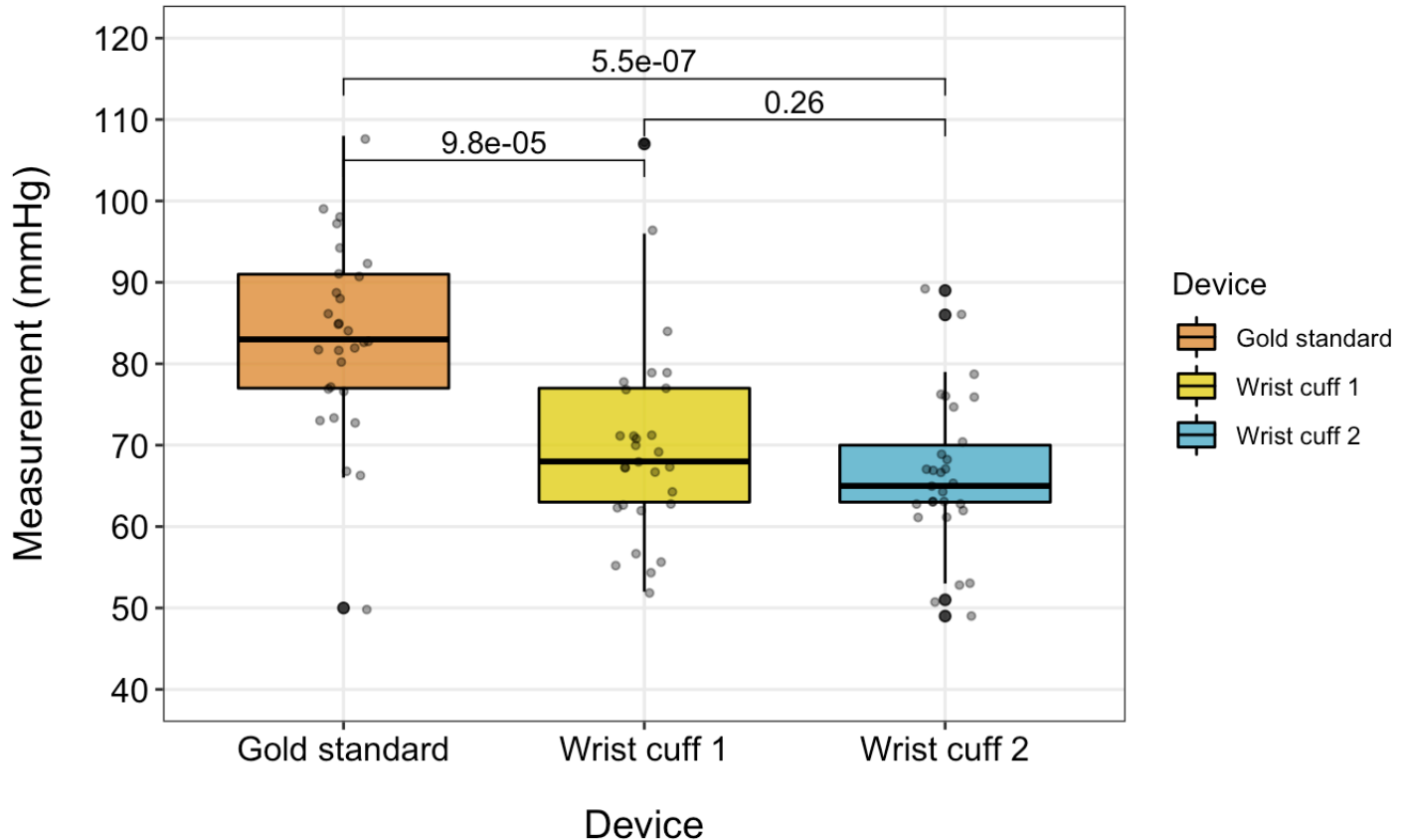
```

```
## # A tibble: 6 × 9
##   .y.    group1      group2      p    p.adj p.format p.signif method y.position
##   <chr> <chr>      <chr>      <dbl>  <dbl> <chr>      <chr>      <chr>      <dbl>
## 1 Sys    Gold standard Wrist ... 6.55e-5 9.8e-5 6.6e-05 ****    T-test      105
## 2 Sys    Gold standard Wrist ... 1.83e-7 5.5e-7 1.8e-07 ****    T-test      115
## 3 Sys    Wrist cuff 1  Wrist ... 2.63e-1 2.6e-1 0.26     ns      T-test      110
## 4 Dia    Gold standard Wrist ... 7.72e-5 1.2e-4 7.7e-05 ****    T-test      68
## 5 Dia    Gold standard Wrist ... 3.14e-8 9.4e-8 3.1e-08 ****    T-test      78
## 6 Dia    Wrist cuff 1  Wrist ... 2.18e-1 2.2e-1 0.22     ns      T-test      73
```

```
sys_all_box1 <- boxplot_3_comparison(df, "Device", "Sys", "Device",
                                   "Measurement (mmHg)",
                                   c(40,120),
                                   "Systolic pressure mean difference",
                                   "n = 29 measurements per device group",
                                   "(method: unpaired t.test with FDR p.adjustmen
t)",
                                   rater_colors,
                                   sys_test,
                                   "p.adj"
                                   )
sys_all_box1 + geom_jitter(size=1, color="black", width=0.1, alpha=0.4)
```

Systolic pressure mean difference

n = 29 measurements per device group



(method: unpaired t.test with FDR p.adjustment)

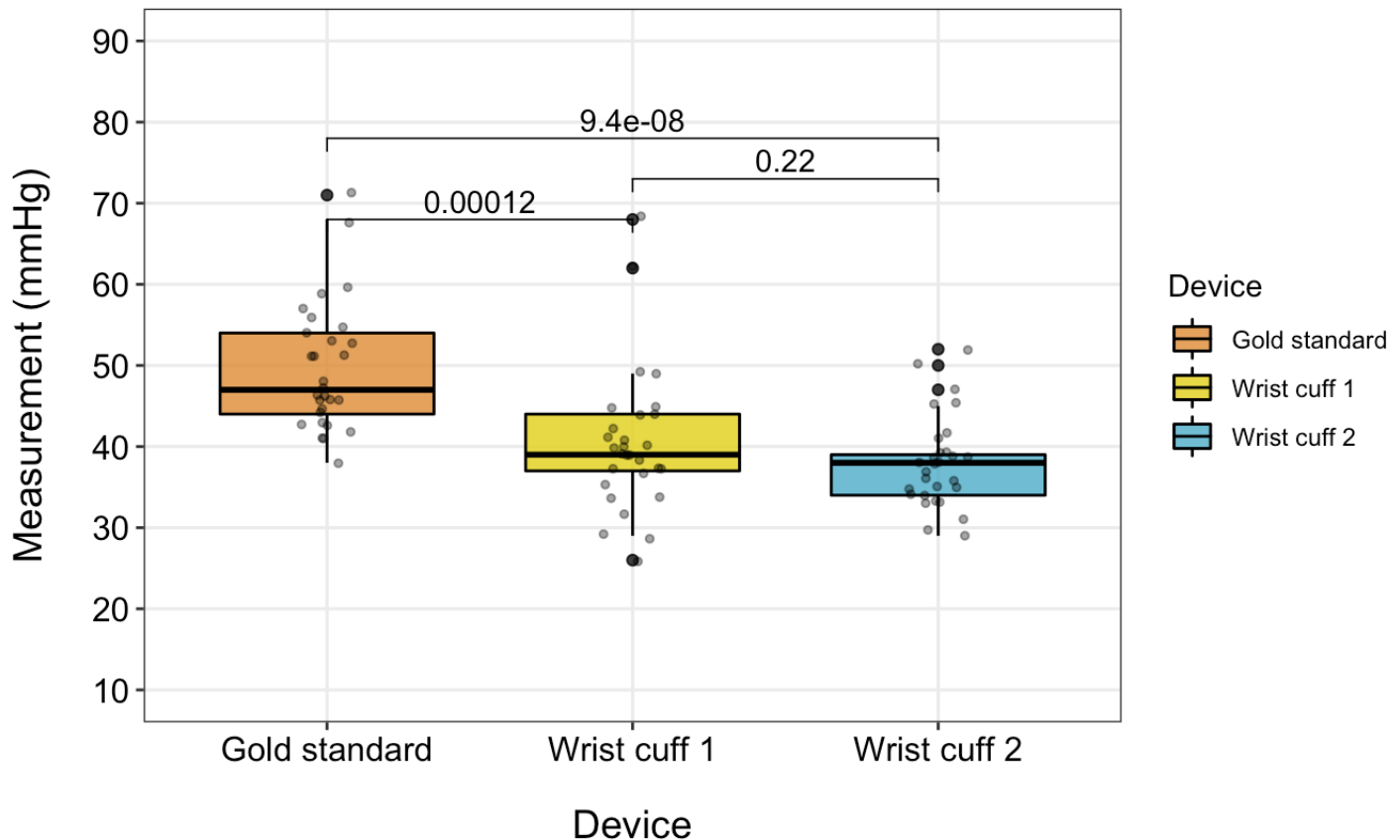
```
ggsave("boxplot_systolic_device.png", width=8, height=6, dpi=300)

dia_all_box1 <- boxplot_3_comparison(df, "Device", "Dia", "Device",
  "Measurement (mmHg)",
  c(10,90),
  "Diastolic pressure mean difference",
  "n = 29 measurements per device group",
  "(method: unpaired t.test with FDR p.adjustment)",
  rater_colors,
  dia_test,
  "p.adj"
)

dia_all_box1 + geom_jitter(size=1, color="black", width=0.1, alpha=0.4)
```

Diastolic pressure mean difference

n = 29 measurements per device group



(method: unpaired t.test with FDR p.adjustment)

```
ggsave("boxplot_diastolic_device.png", width=8, height=6, dpi=300)
```

- result = when looking at all data points, the mean difference between two wrist cuff groups is insignificant. significant mean difference observed for all wrist cuff to gold standard comparisons, with the gold standard on average having higher measurement values than the wrist cuff.

```
rater_dispo_percent <- df %>%  
  dplyr::group_by(Device, Disposition) %>%  
  dplyr::tally() %>%  
  dplyr::mutate(Percent = round(n/sum(n) * 100, 1))  
rater_dispo_percent
```



```
## # A tibble: 10 × 4
## # Groups:   Device [3]
##   Device      Disposition      n Percent
##   <fct>      <fct>      <int>  <dbl>
## 1 Gold standard sleeping        3    10.3
## 2 Gold standard calm          18    62.1
## 3 Gold standard active         7    24.1
## 4 Gold standard crying         1     3.4
## 5 Wrist cuff 1 sleeping         7    24.1
## 6 Wrist cuff 1 calm          17    58.6
## 7 Wrist cuff 1 active          5    17.2
## 8 Wrist cuff 2 sleeping         9     31
## 9 Wrist cuff 2 calm          16    55.2
## 10 Wrist cuff 2 active          4    13.8
```

```

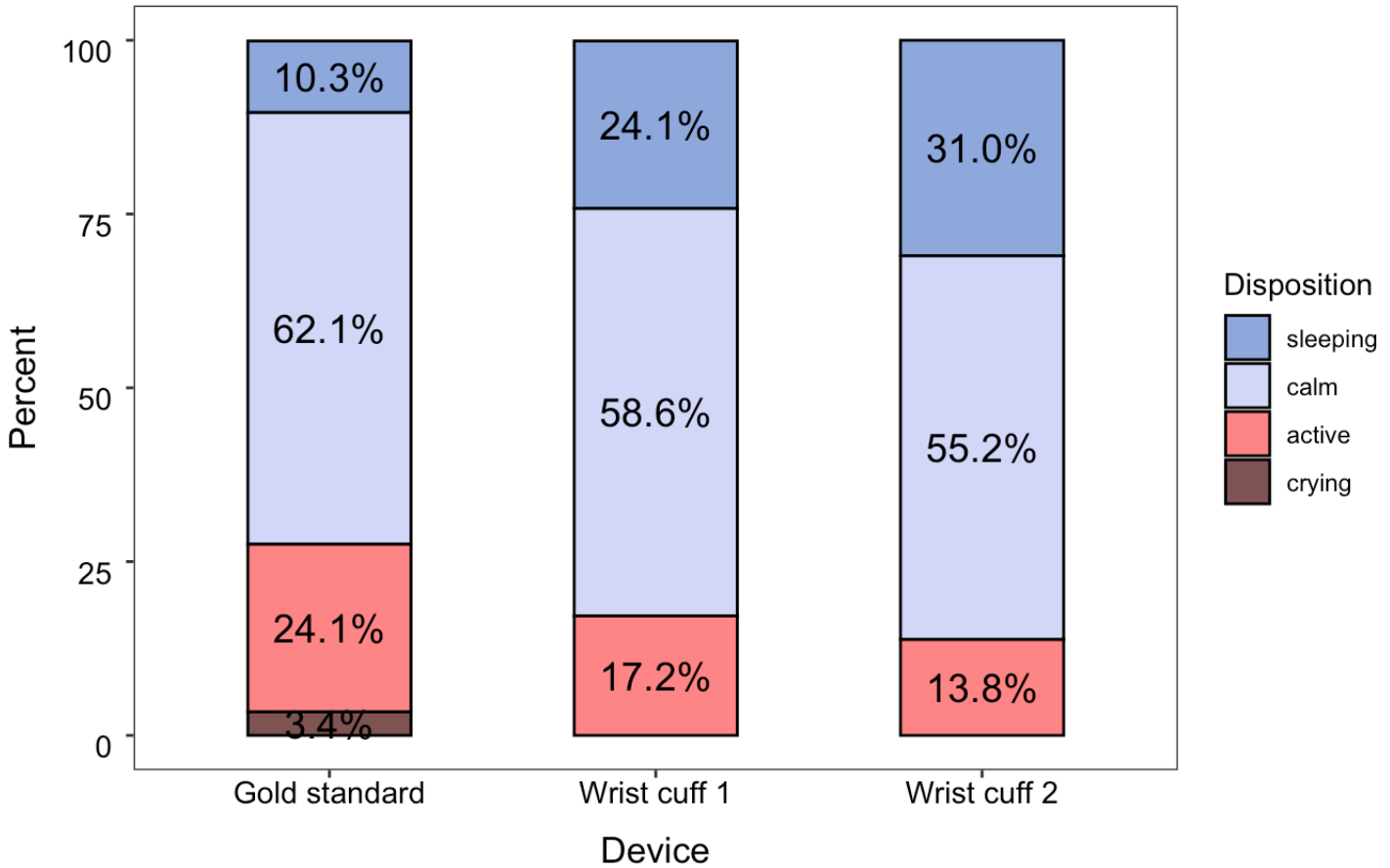
bar_dispo_percentnt <- ggplot(rater_dispo_percent, aes(x=Device,
              y=Percent,
              fill=Disposition)) +
  geom_bar(stat="identity", width=0.5, colour="black", alpha=0.8) +
  geom_text(aes(label=paste0(sprintf("%1.1f", Percent), "%"),
              position=position_stack(vjust=0.5),
              colour="black",
              size = 5,
              check_overlap = TRUE) +
  scale_fill_manual(values = dispo_colors) +
  labs(fill='Disposition') +
  theme_bw() +
  theme(panel.grid.minor = element_blank(),
        panel.grid.major = element_blank(),
        panel.background = element_blank(),
        axis.line = element_blank()) +
        labs(x="Device",
              y= "Percent",
              title="Disposition proportions per device group",
              subtitle = "n = 29 measurements per group") +
        theme(legend.position='right') +

        theme(axis.title.x = element_text(size=13, color="black", margin=
margin(t=10, b=5)),
              axis.title.y = element_text(vjust = 3, size=13, color="black", margin=margin(t=5, b=5, r=1, l=1)),
              axis.text.y = element_text(vjust = 1, size=11, color="black", margin = margin(t = 5, r=5, b = 5)),
              axis.text.x = element_text(size=11, color="black", margin
= margin(t = 1)),
              axis.text = element_text(size=13, color="black", margin = margin(t = 1))
        )
bar_dispo_percentnt

```

Disposition proportions per device group

n = 29 measurements per group



```
ggsave("barstack_disposition_proportions_per_device.png", width = 8, height = 6, dpi=300)
```

- result: larger % of babies are sleeping during measurements taken with wrist cuff (24% and 31%) than with gold standard (10%). this makes me think that the lower measurements seen for wrist cuff vs gold standard are related to baby disposition, and interpretation of above results is confounded by this variable. Most babies were calm during measurements for all 3 groups. are the differences significant when only looking at calm disposition measurements?

```
dispo_count <- df %>%  
  dplyr::group_by(Disposition) %>%  
  dplyr::tally() %>%  
  dplyr::mutate(Percent.Dispo = round(n/sum(n) * 100, 1))  
dispo_count
```

```
## # A tibble: 4 × 3
##   Disposition      n Percent.Dispo
##   <fct>         <int>         <dbl>
## 1 sleeping         19          21.8
## 2 calm            51          58.6
## 3 active          16          18.4
## 4 crying           1           1.1
```

```
nocry <- df %>% filter(Disposition != "crying")

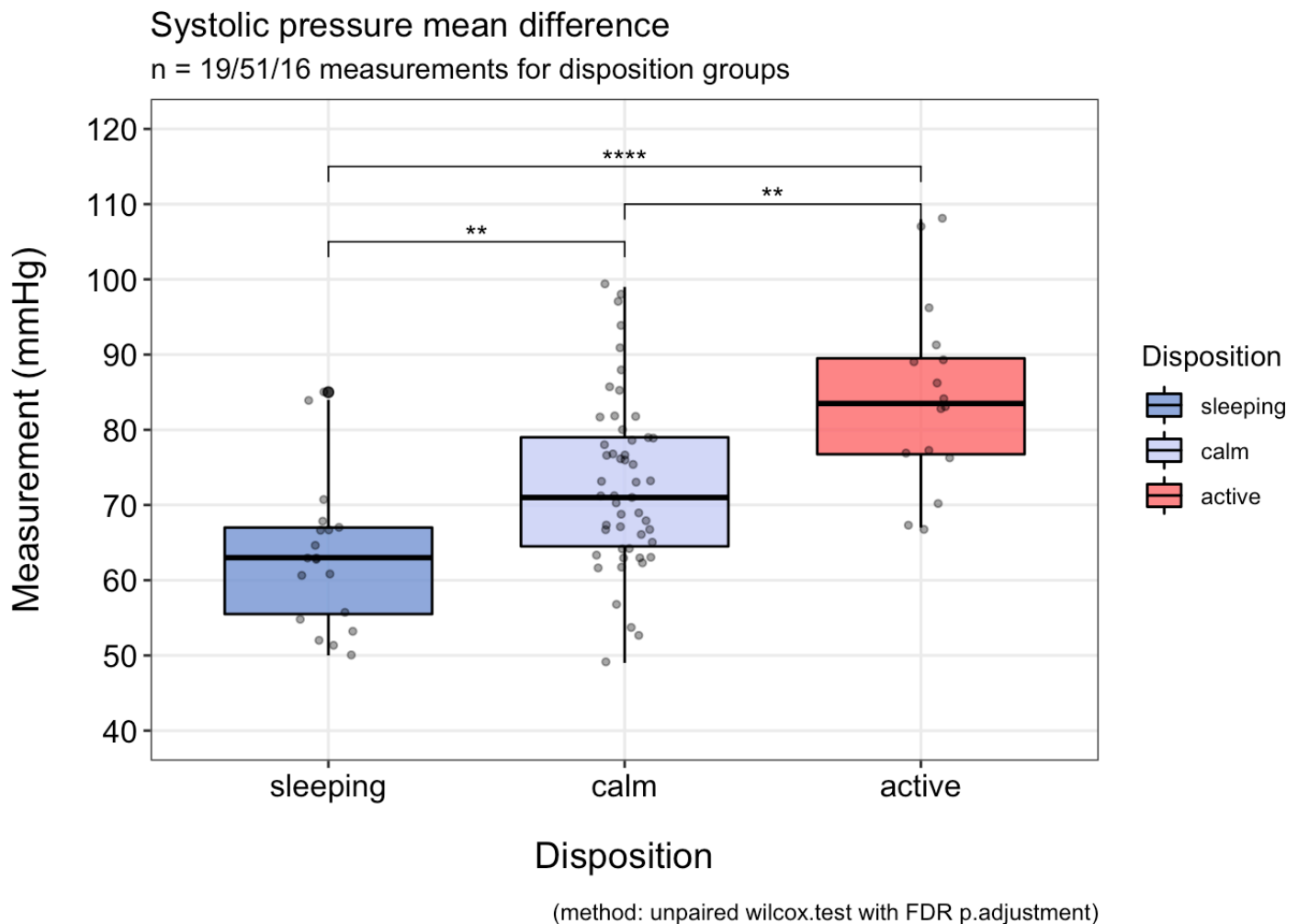
sys_test <- compare_means(Sys ~ Disposition,
                          data=nocry,
                          method="wilcox.test",
                          p.adjust.method = "fdr",
                          paired=FALSE,
                          comparisons = compare_dispo)
sys_test <- sys_test %>% mutate(y.position = c(105, 115, 110))

dia_test <- compare_means(Dia ~ Disposition,
                          data=nocry,
                          method="wilcox.test",
                          p.adjust.method = "fdr",
                          paired=FALSE,
                          comparisons = compare_dispo)
dia_test <- dia_test %>% mutate(y.position = c(68, 78, 73))

bind_rows(sys_test, dia_test)
```

```
## # A tibble: 6 × 9
##   .y.   group1 group2      p    p.adj p.format p.signif method y.position
##   <chr> <chr>   <chr>    <dbl>  <dbl> <chr>   <chr>   <chr>    <dbl>
## 1 Sys   sleeping calm    0.00128  0.0019  0.0013  **      Wilcox...  105
## 2 Sys   sleeping active  0.0000291  0.000087  2.9e-05  ****     Wilcox...  115
## 3 Sys   calm    active  0.00220  0.0022  0.0022  **      Wilcox...  110
## 4 Dia   sleeping calm    0.000455  0.00046  0.00046  ***     Wilcox...   68
## 5 Dia   sleeping active  0.00000378  0.000011  3.8e-06  ****     Wilcox...   78
## 6 Dia   calm    active  0.0000114  0.000017  1.1e-05  ****     Wilcox...   73
```

```
# plot with p adjust value
sys_all_box1 <- boxplot_3_comparison(nocry, "Disposition", "Sys", "Disposition",
                                   "Measurement (mmHg)",
                                   c(40,120),
                                   "Systolic pressure mean difference",
                                   "n = 19/51/16 measurements for disposition group
s",
                                   "(method: unpaired wilcox.test with FDR p.adjust
ment)",
                                   dispo_colors,
                                   sys_test,
                                   "p.signif"
                                   )
sys_all_box1+ geom_jitter(size=1, color="black", alpha=0.4, width=0.1)
```

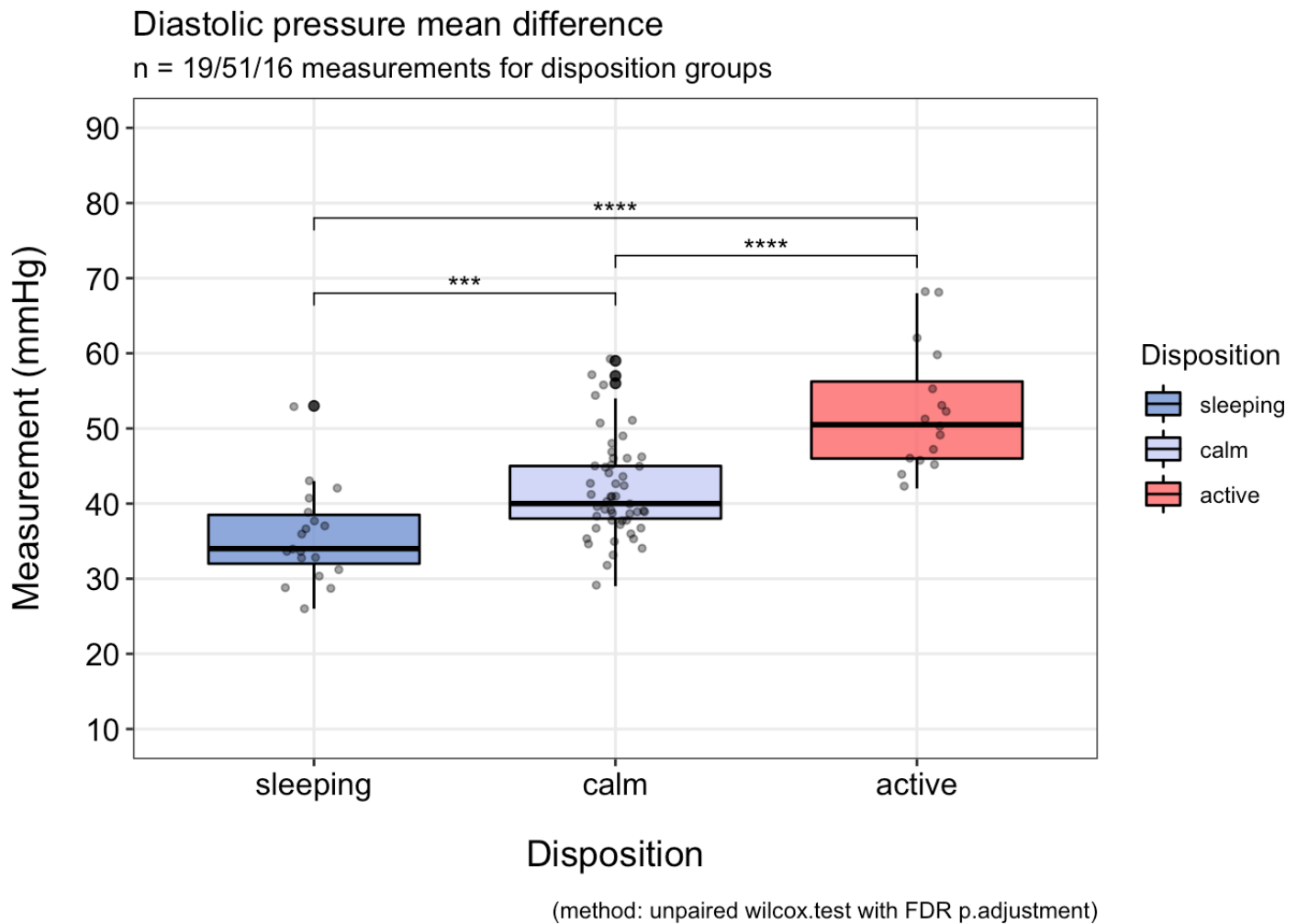


```

ggsave("boxplot_systolic_disposition.png", width=8, height=6, dpi=300)

# plot with p adjust value
dia_all_box1 <- boxplot_3_comparison(nocry, "Disposition", "Dia", "Disposition",
  "Measurement (mmHg)",
  c(10,90),
  "Diastolic pressure mean difference",
  "n = 19/51/16 measurements for disposition group
s",
  "(method: unpaired wilcox.test with FDR p.adjust
ment)",
  dispo_colors,
  dia_test,
  "p.signif"
)
dia_all_box1 + geom_jitter(size=1, color="black", alpha=0.4, width=0.1)

```



```

ggsave("boxplot_diastolic_disposition.png", width=8, height=6, dpi=300)

```

```

calm <- df %>% filter(Disposition == 'calm')

sys_test <- compare_means(Sys ~ Device,
                          data=calm,
                          method="wilcox.test",
                          p.adjust.method = "fdr",
                          paired=FALSE,
                          comparisons = compare_devices)
sys_test <- sys_test %>% mutate(y.position = c(105, 115, 110))

dia_test <- compare_means(Dia ~ Device,
                          data=calm,
                          method="wilcox.test",
                          p.adjust.method = "fdr",
                          paired=FALSE,
                          comparisons = compare_devices)
dia_test <- dia_test %>% mutate(y.position = c(68, 78, 73))

bind_rows(sys_test, dia_test)

```

```

## # A tibble: 6 × 9
##   .y.    group1      group2      p  p.adj p.format p.signif method y.position
##   <chr> <chr>      <chr>    <dbl> <dbl> <chr>    <chr>    <chr>    <dbl>
## 1 Sys   Gold standard Wrist ... 5.15e-5 7.7e-5 5.2e-05 ****    Wilco...    105
## 2 Sys   Gold standard Wrist ... 2.71e-5 7.7e-5 2.7e-05 ****    Wilco...    115
## 3 Sys   Wrist cuff 1   Wrist ... 4.27e-1 4.3e-1 0.43     ns      Wilco...    110
## 4 Dia   Gold standard Wrist ... 1.53e-4 2.3e-4 0.00015 ***     Wilco...     68
## 5 Dia   Gold standard Wrist ... 1.30e-5 3.9e-5 1.3e-05 ****    Wilco...     78
## 6 Dia   Wrist cuff 1   Wrist ... 2.69e-1 2.7e-1 0.26857 ns      Wilco...     73

```

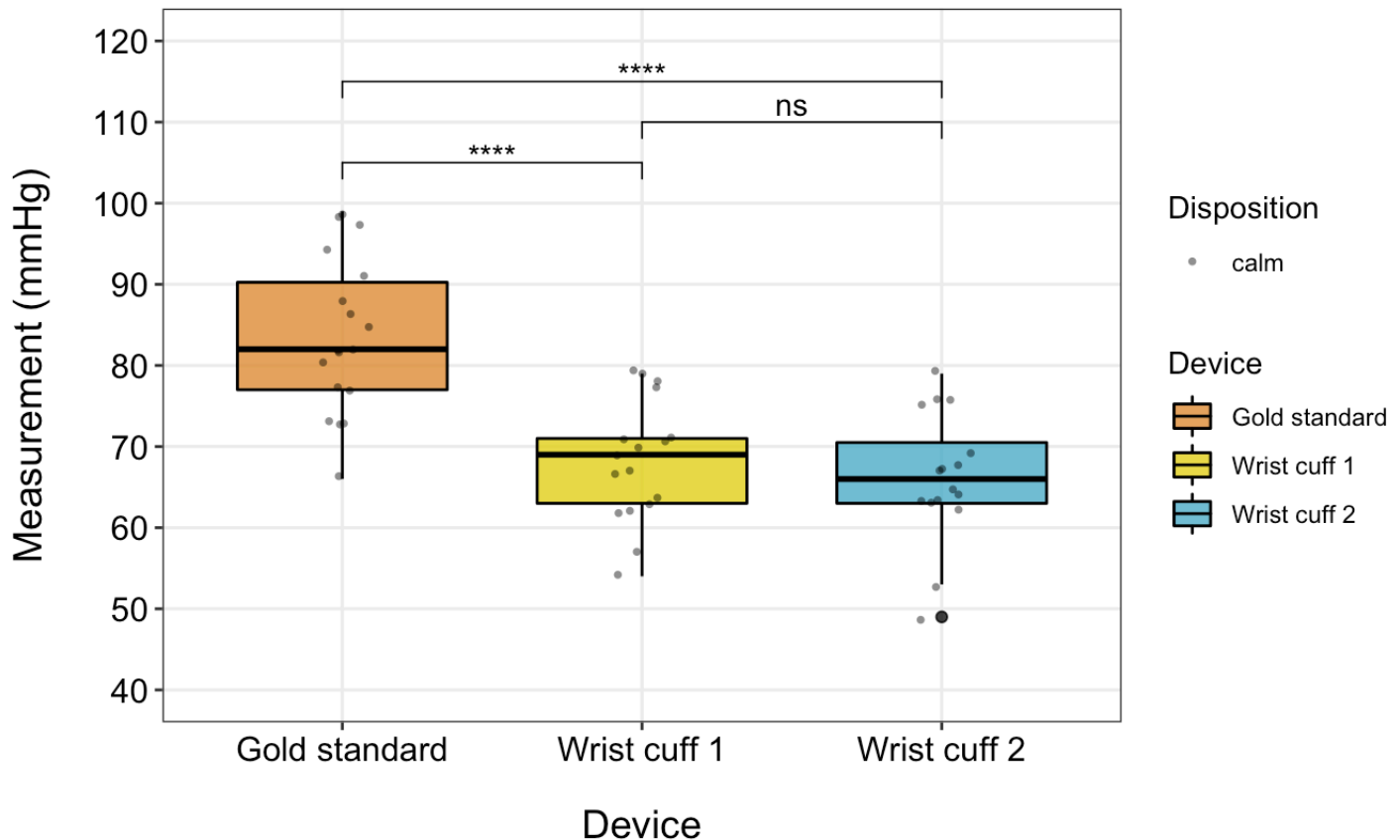
```

# plot with p adjust value
sys_all_box1 <- boxplot_3_comparison(calm, "Device", "Sys", "Device",
                                   "Measurement (mmHg)",
                                   c(40,120),
                                   "Systolic pressure mean difference",
                                   "n = 18/17/16 measurements for calm disposition",
                                   "(method: unpaired wilcox.test with FDR p.adjust
ment)",
                                   rater_colors,
                                   sys_test,
                                   "p.signif"
                                   )
sys_all_box1 + geom_jitter(aes(shape=Disposition),size=1, color="black", alpha=0.5, w
idth=0.1)

```

Systolic pressure mean difference

n = 18/17/16 measurements for calm disposition



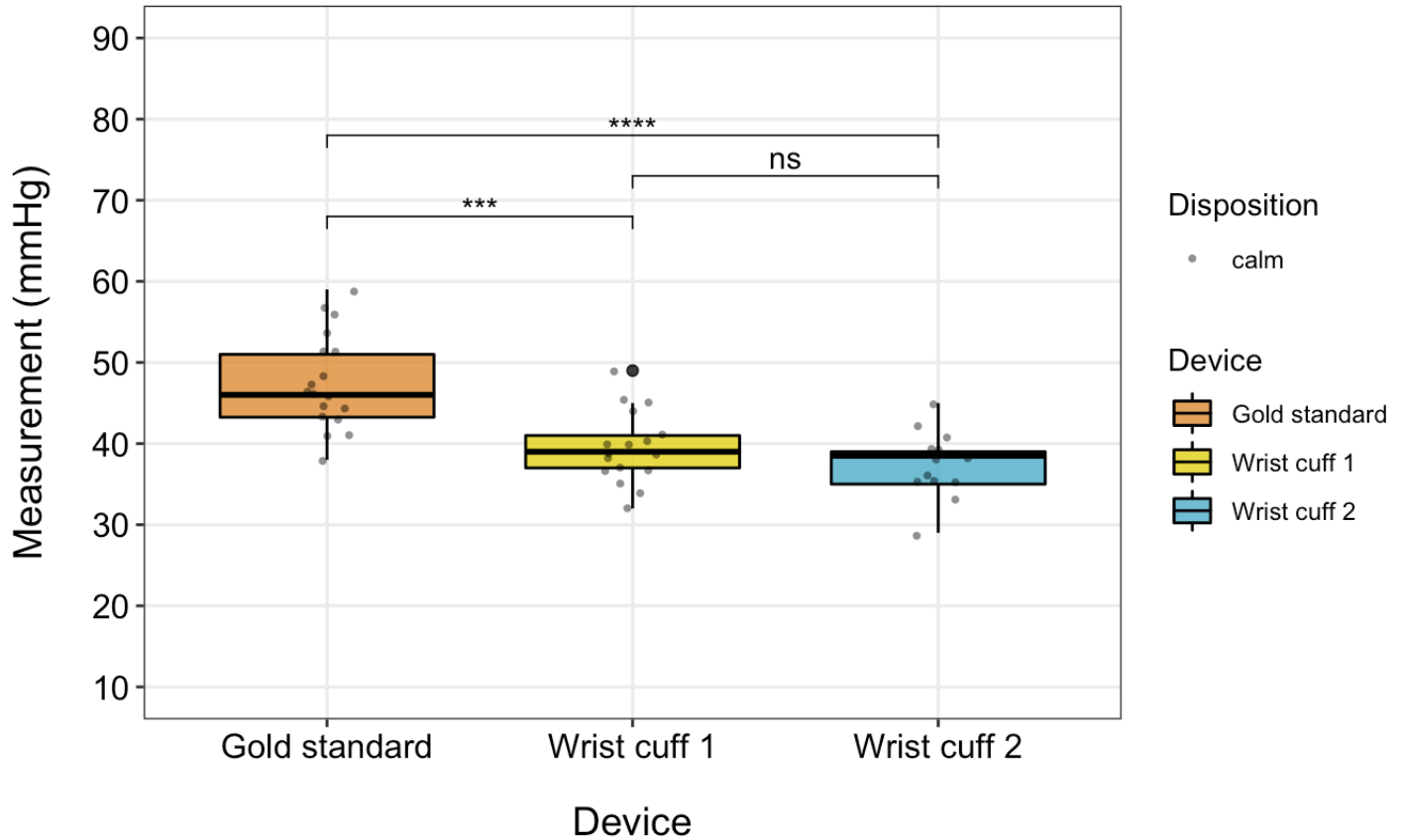
(method: unpaired wilcox.test with FDR p.adjustment)

```
ggsave("boxplot_systolic_calm_disposition.png", width=8, height=6, dpi=300)

# plot with p adjust value
dia_all_box1 <- boxplot_3_comparison(calm, "Device", "Dia", "Device",
                                     "Measurement (mmHg)",
                                     c(10,90),
                                     "Diastolic pressure mean difference",
                                     "n = 18/17/16 measurements for calm dispositio
n",
                                     "(method: unpaired wilcox.test with FDR p.adjust
ment)",
                                     rater_colors,
                                     dia_test,
                                     "p.signif"
                                     )
dia_all_box1 + geom_jitter(aes(shape=Disposition),size=1, color="black", alpha=0.5, width=0.1)
```


Diastolic pressure mean difference

n = 18/17/16 measurements for calm disposition



(method: unpaired wilcox.test with FDR p.adjustment)

```
ggsave("boxplot_diastolic_calm_disposition.png", width=8, height=6, dpi=300)
```

- result: used wilcox non parametric test because each group has different number of measurements (18 vs 17 vs 16) and paired=false because multiple babies included. calm only results replicate all data point boxplot result.

```

sleep <- df %>% filter(Disposition == 'sleeping')

sys_test <- compare_means(Sys ~ Device,
                          data=sleep,
                          method="wilcox.test",
                          p.adjust.method = "fdr",
                          paired=FALSE,
                          comparisons = compare_devices)
sys_test <- sys_test %>% mutate(y.position = c(105, 115, 110))

dia_test <- compare_means(Dia ~ Device,
                          data=sleep,
                          method="wilcox.test",
                          p.adjust.method = "fdr",
                          paired=FALSE,
                          comparisons = compare_devices)
dia_test <- dia_test %>% mutate(y.position = c(68, 78, 73))

bind_rows(sys_test, dia_test)

```

```

## # A tibble: 6 × 9
##   .y.    group1      group2      p p.adj p.format p.signif method y.position
##   <chr> <chr>      <chr>      <dbl> <dbl> <chr>      <chr>      <chr>      <dbl>
## 1 Sys   Gold standard Wrist cu... 0.517  0.67  0.52      ns      Wilco...    105
## 2 Sys   Gold standard Wrist cu... 0.457  0.67  0.46      ns      Wilco...    115
## 3 Sys   Wrist cuff 1   Wrist cu... 0.670  0.67  0.67      ns      Wilco...    110
## 4 Dia   Gold standard Wrist cu... 0.0222 0.033 0.022      *      Wilco...     68
## 5 Dia   Gold standard Wrist cu... 0.0159 0.033 0.016      *      Wilco...     78
## 6 Dia   Wrist cuff 1   Wrist cu... 0.958  0.96  0.958     ns      Wilco...     73

```

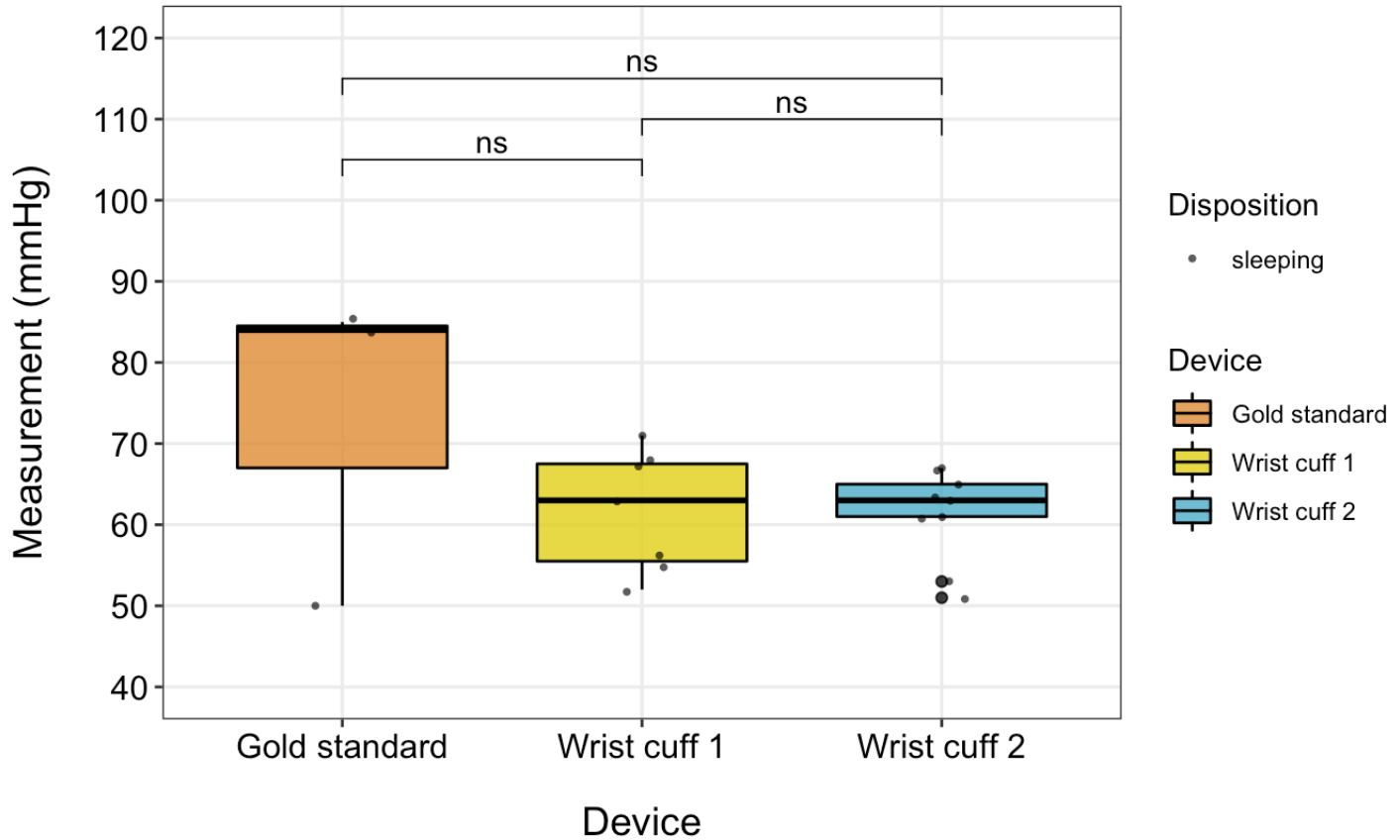
```

# plot with p adjust value
sys_all_box1 <- boxplot_3_comparison(sleep, "Device", "Sys", "Device",
                                   "Measurement (mmHg)",
                                   c(40,120),
                                   "Systolic pressure mean difference",
                                   "n = 3/7/9 measurements for sleeping dispositio
n",
                                   "(method: unpaired wilcox.test with FDR p.adjust
ment)",
                                   rater_colors,
                                   sys_test,
                                   "p.signif"
                                   )
sys_all_box1 + geom_jitter(aes(shape=Disposition),size=1, width=0.1, color="black", a
lpha=0.7)

```

Systolic pressure mean difference

n = 3/7/9 measurements for sleeping disposition



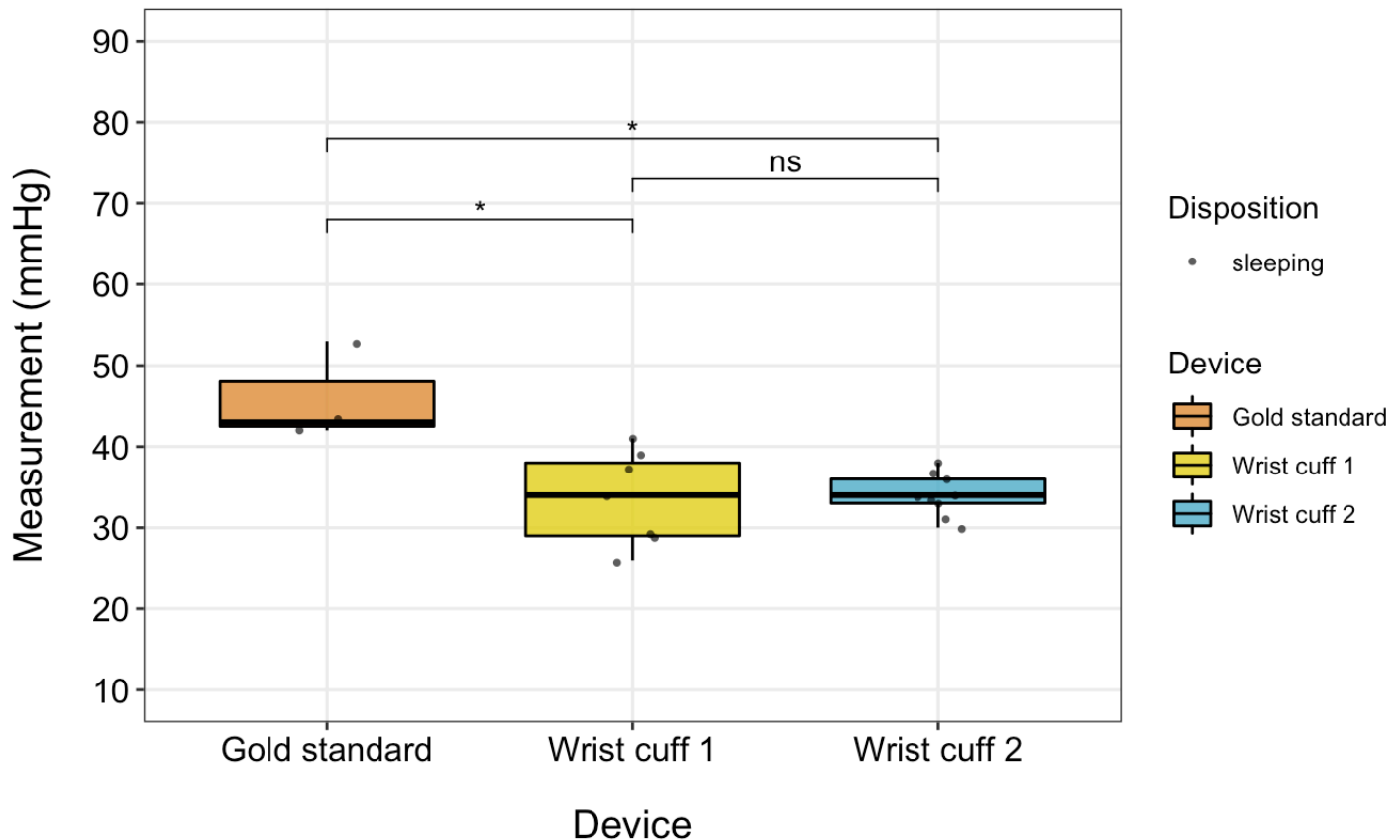
(method: unpaired wilcox.test with FDR p.adjustment)

```
ggsave("boxplot_systolic_sleeping_disposition.png", width=8, height=6, dpi=300)

# plot with p adjust value
dia_all_box1 <- boxplot_3_comparison(sleep, "Device", "Dia", "Device",
                                     "Measurement (mmHg)",
                                     c(10,90),
                                     "Diastolic pressure mean difference",
                                     "n = 3/7/9 measurements for sleeping dispositio
n",
                                     "(method: unpaired wilcox.test with FDR p.adjust
ment)",
                                     rater_colors,
                                     dia_test,
                                     "p.signif"
                                     )
dia_all_box1 + geom_jitter(aes(shape=Disposition),size=1, width=0.1, color="black", a
lpha=0.7)
```

Diastolic pressure mean difference

n = 3/7/9 measurements for sleeping disposition



(method: unpaired wilcox.test with FDR p.adjustment)

```
ggsave("boxplot_diastolic_sleeping_disposition.png", width=8, height=6, dpi=300)
```

- result: used wilcox non parametric test because each group has different number of measurements (3 vs 7 vs 9) and paired=false. unlike calm result, the sleeping only results did not replicate significance seen for data points boxplot result. either babies are less variable and devices therefore show similar readings (no significant mean diff for sys and slight significant diff for diastolic) when they are sleeping, or the gold standard having the lowest number of data points confounds the differences thats possible to detect between devices.

following these results for all data points and select disposition data points, i wondered if measurement differences were significant when controlling for baby by looking at data points taken of same baby. for stats test i used paired = True, method = t.test, selected babies with 3 sessions because they have more data points available. * question for biostats expert- is t.test best method to use for single baby measurements? wilcox was first used because all data was not normally distributed but none of the values were significantly different which seemed odd given some box plots for gold vs wristcuff are clearly different (non overlapping interquartile ranges)

```

baby <- df %>% filter(BabyID == '2')

sys_test <- compare_means(Sys ~ Device,
                          data=baby,
                          method="t.test",
                          p.adjust.method = "fdr",
                          paired=TRUE,
                          comparisons = compare_devices)
sys_test <- sys_test %>% mutate(y.position = c(105, 115, 110))
dia_test <- compare_means(Dia ~ Device,
                          data=baby,
                          method="t.test",
                          p.adjust.method = "fdr",
                          paired=TRUE,
                          comparisons = compare_devices)
dia_test <- dia_test %>% mutate(y.position = c(68, 78, 73))
bind_rows(sys_test, dia_test)

```

```

## # A tibble: 6 × 9
##   .y.   group1      group2      p p.adj p.format p.signif method y.position
##   <chr> <chr>      <chr>    <dbl> <dbl> <chr>      <chr>    <chr>      <dbl>
## 1 Sys   Gold standard Wrist cu... 0.174  0.26  0.174      ns      T-test      105
## 2 Sys   Gold standard Wrist cu... 0.0192 0.058 0.019      *      T-test      115
## 3 Sys   Wrist cuff 1   Wrist cu... 0.572  0.57  0.572      ns      T-test      110
## 4 Dia   Gold standard Wrist cu... 0.876  0.88  0.88       ns      T-test       68
## 5 Dia   Gold standard Wrist cu... 0.156  0.47  0.16       ns      T-test       78
## 6 Dia   Wrist cuff 1   Wrist cu... 0.564  0.85  0.56       ns      T-test       73

```

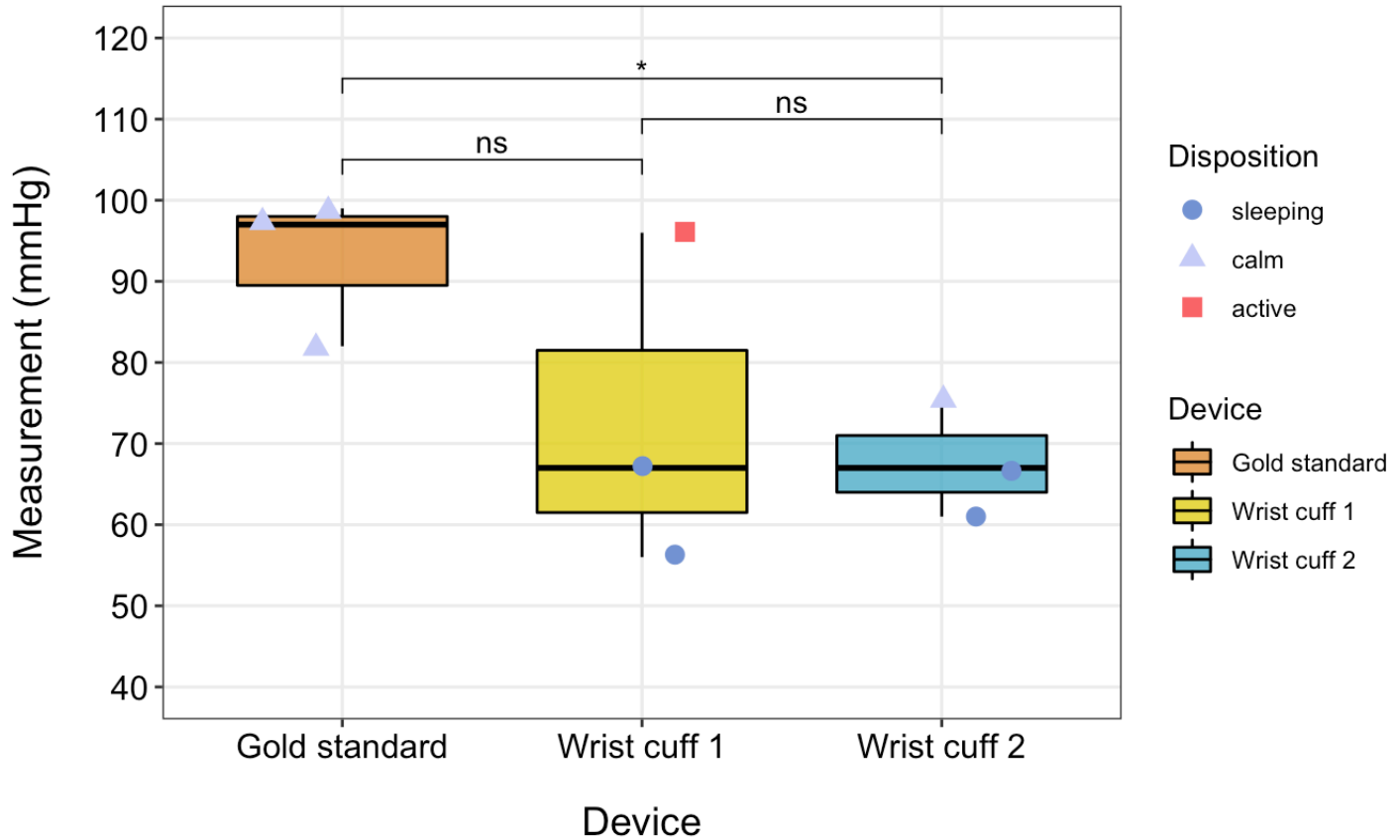
```

# plot sys
sys_baby_box2 <- boxplot_3_comparison(baby, "Device", "Sys", "Device",
                                     "Measurement (mmHg)",
                                     c(40,120),
                                     "Baby 2 Systolic pressure mean difference",
                                     "n = 9 measurements for babyID 2",
                                     "(method: paired t.test with FDR p.adjustment)",
                                     rater_colors,
                                     sys_test,
                                     "p.signif"
                                     )
sys_baby_box2 + geom_jitter(aes(shape=Disposition, colour=Disposition),size=3, alpha=
1) + scale_colour_manual(values = dispo_colors)

```

Baby 2 Systolic pressure mean difference

n = 9 measurements for babyID 2



(method: paired t.test with FDR p.adjustment)

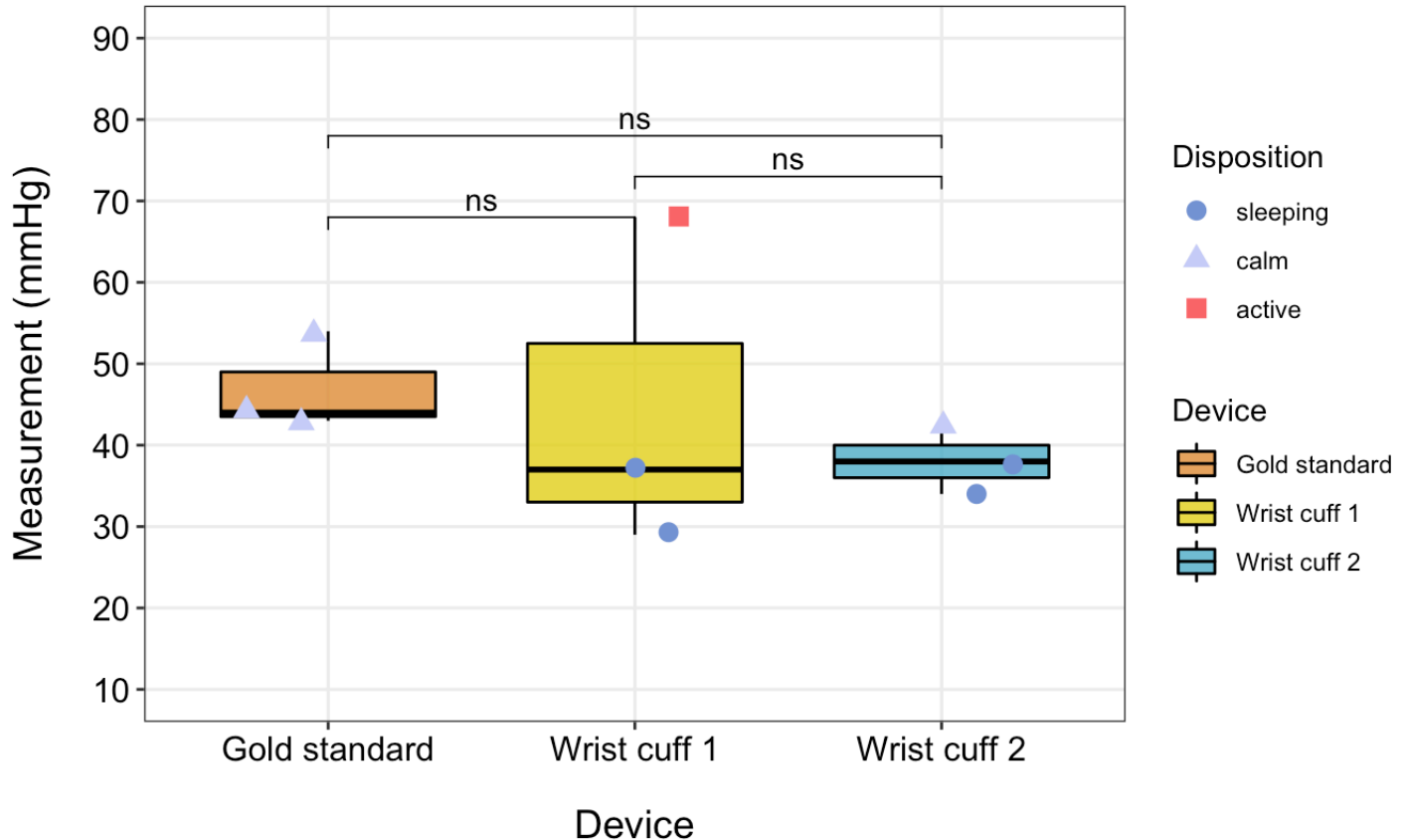
```
ggsave("baby2_systolic.png", width=8, height=6, dpi=300)

dia_baby_box2 <- boxplot_3_comparison(baby, "Device", "Dia", "Device",
                                     "Measurement (mmHg)",
                                     c(10,90),
                                     "Baby 2 Diastolic pressure mean difference",
                                     "n = 9 measurements",
                                     "(method: paired t.test with FDR p.adjustment)",
                                     rater_colors,
                                     dia_test,
                                     "p.signif"
                                   )

dia_baby_box2 + geom_jitter(aes(shape=Disposition, colour=Disposition),size=3, alpha=
1) + scale_colour_manual(values = dispo_colors)
```

Baby 2 Diastolic pressure mean difference

n = 9 measurements



(method: paired t.test with FDR p.adjustment)

```
ggsave("baby2_diastolic.png", width=8, height=6, dpi=300)
```

```

baby <- df %>% filter(BabyID == '3')

sys_test <- compare_means(Sys ~ Device,
                          data=baby,
                          method="t.test",
                          p.adjust.method = "fdr",
                          paired=TRUE,
                          comparisons = compare_devices)
sys_test <- sys_test %>% mutate(y.position = c(105, 115, 110))
dia_test <- compare_means(Dia ~ Device,
                          data=baby,
                          method="t.test",
                          p.adjust.method = "fdr",
                          paired=TRUE,
                          comparisons = compare_devices)
dia_test <- dia_test %>% mutate(y.position = c(68, 78, 73))
bind_rows(sys_test, dia_test)
    
```

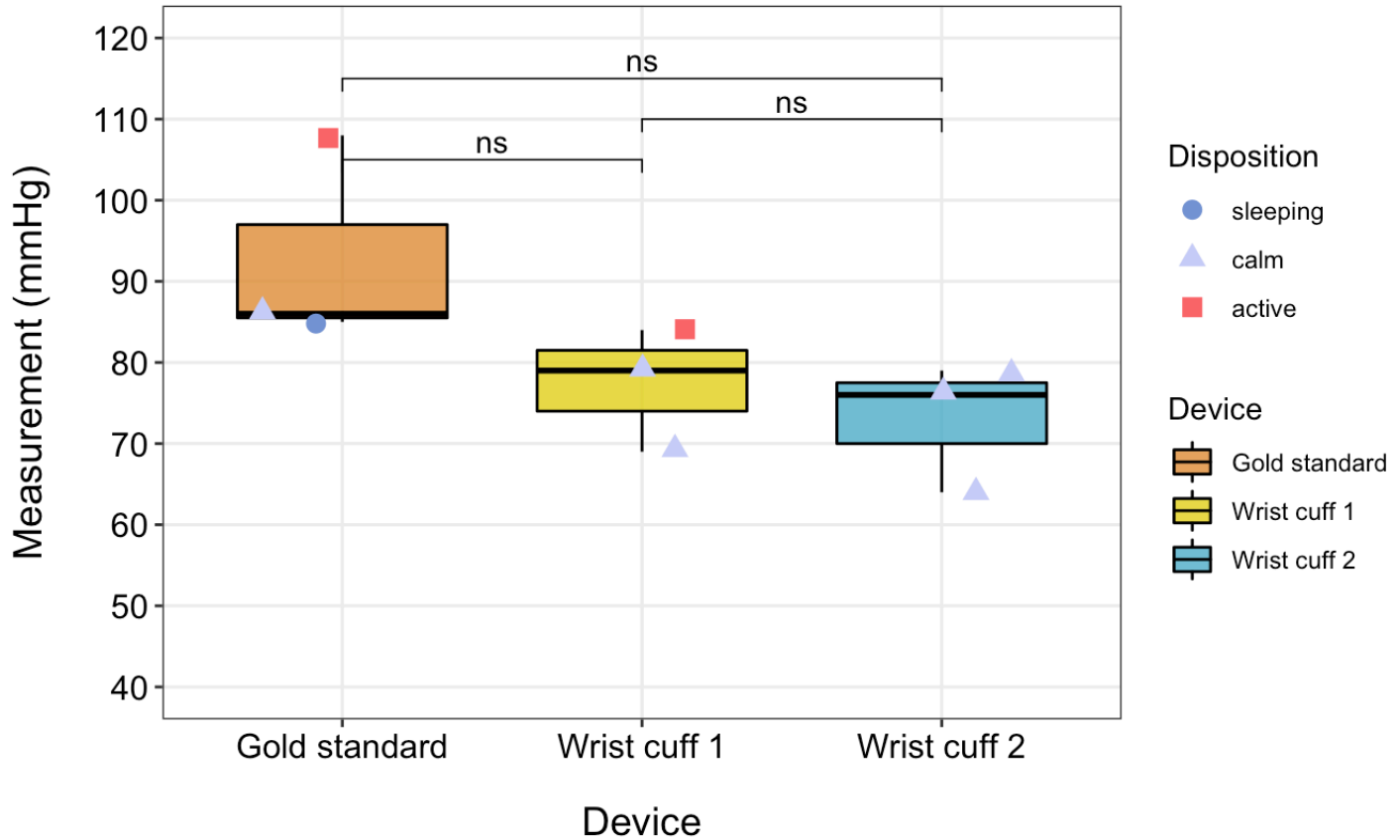
```
## # A tibble: 6 × 9
```

	.y.	group1	group2	p	p.adj	p.format	p.signif	method	y.position
	<chr>	<chr>	<chr>	<dbl>	<dbl>	<chr>	<chr>	<chr>	<dbl>
## 1	Sys	Gold standard	Wrist cu...	0.182	0.2	0.182	ns	T-test	105
## 2	Sys	Gold standard	Wrist cu...	0.0682	0.2	0.068	ns	T-test	115
## 3	Sys	Wrist cuff 1	Wrist cu...	0.204	0.2	0.204	ns	T-test	110
## 4	Dia	Gold standard	Wrist cu...	0.310	0.31	0.310	ns	T-test	68
## 5	Dia	Gold standard	Wrist cu...	0.123	0.18	0.123	ns	T-test	78
## 6	Dia	Wrist cuff 1	Wrist cu...	0.0171	0.051	0.017	*	T-test	73

```
# plot sys
sys_baby_box2 <- boxplot_3_comparison(baby, "Device", "Sys", "Device",
                                     "Measurement (mmHg)",
                                     c(40,120),
                                     "Baby 3 Systolic pressure mean difference",
                                     "n = 9 measurements",
                                     "(method: paired t.test with FDR p.adjustment)",
                                     rater_colors,
                                     sys_test,
                                     "p.signif"
                                   )
sys_baby_box2 + geom_jitter(aes(shape=Disposition, colour=Disposition),size=3, alpha=
1) + scale_colour_manual(values = dispo_colors)
```


Baby 3 Systolic pressure mean difference

n = 9 measurements



(method: paired t.test with FDR p.adjustment)

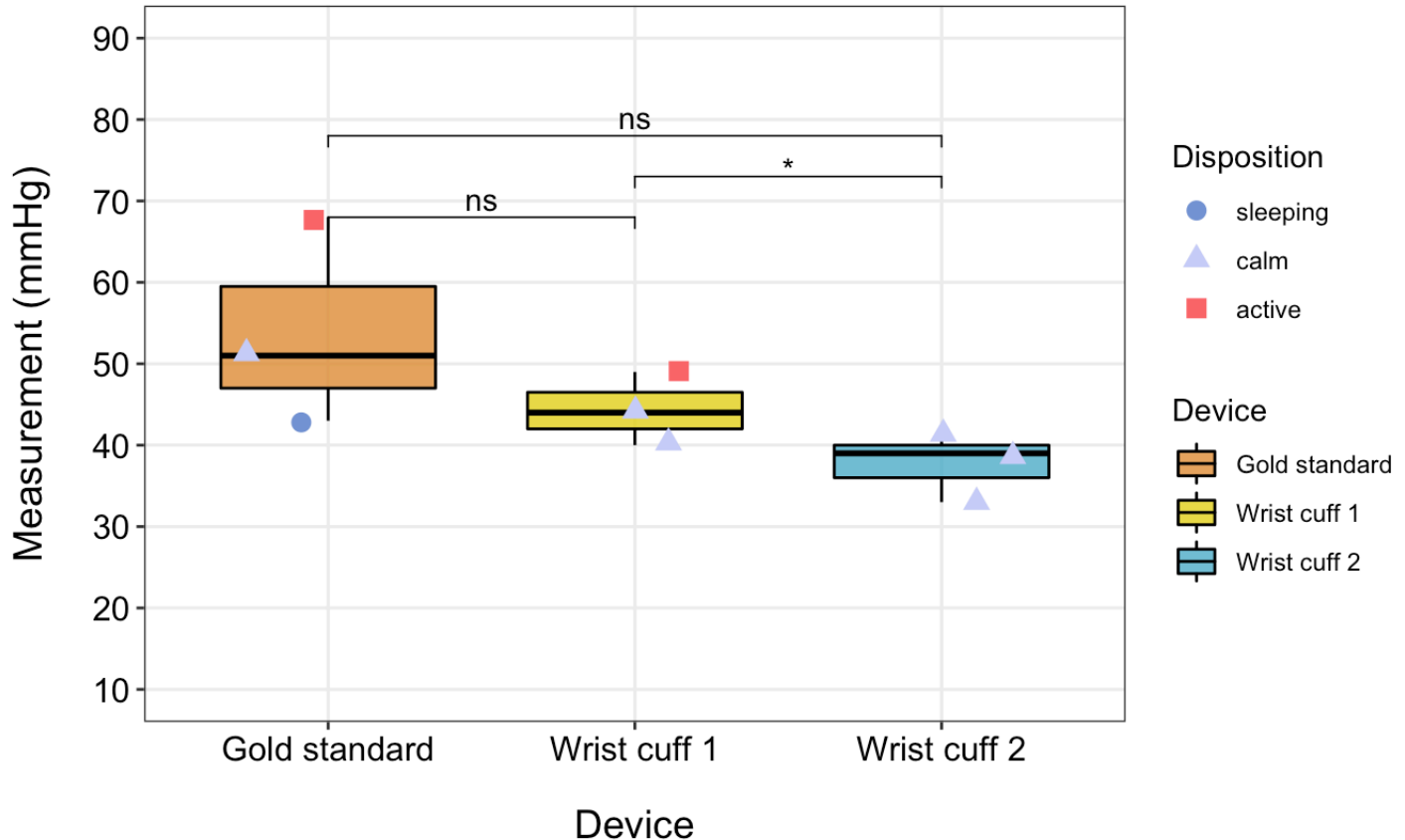
```
ggsave("baby3_systolic.png", width=8, height=6, dpi=300)

# plot dia
dia_baby_box2 <- boxplot_3_comparison(baby, "Device", "Dia", "Device",
                                     "Measurement (mmHg)",
                                     c(10,90),
                                     "Baby 3 Diastolic pressure mean difference",
                                     "n = 9 measurements",
                                     "(method: paired t.test with FDR p.adjustment)",
                                     rater_colors,
                                     dia_test,
                                     "p.signif"
                                     )

dia_baby_box2 + geom_jitter(aes(shape=Disposition, colour=Disposition),size=3, alpha=
1) + scale_colour_manual(values = dispo_colors)
```

Baby 3 Diastolic pressure mean difference

n = 9 measurements



(method: paired t.test with FDR p.adjustment)

```
ggsave("baby3_diastolic.png", width=8, height=6, dpi=300)
```

```

baby <- df %>% filter(BabyID == '4')

sys_test <- compare_means(Sys ~ Device,
                          data=baby,
                          method="t.test",
                          p.adjust.method = "fdr",
                          paired=TRUE,
                          comparisons = compare_devices)
sys_test <- sys_test %>% mutate(y.position = c(105, 115, 110))
dia_test <- compare_means(Dia ~ Device,
                          data=baby,
                          method="t.test",
                          p.adjust.method = "fdr",
                          paired=TRUE,
                          comparisons = compare_devices)
dia_test <- dia_test %>% mutate(y.position = c(68, 78, 73))
bind_rows(sys_test, dia_test)
    
```

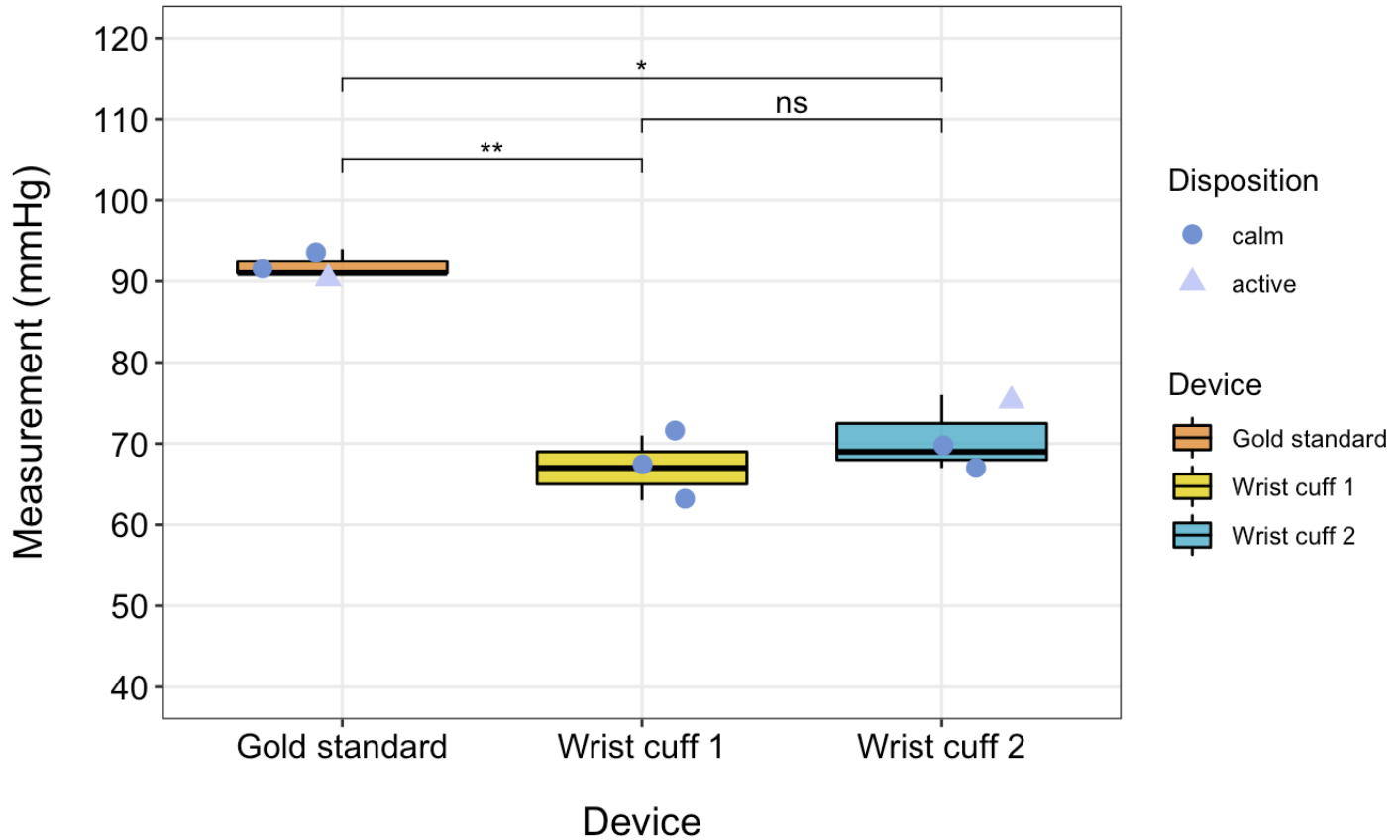
```
## # A tibble: 6 × 9
```

	.y.	group1	group2	p	p.adj	p.format	p.signif	method	y.position
	<chr>	<chr>	<chr>	<dbl>	<dbl>	<chr>	<chr>	<chr>	<dbl>
## 1	Sys	Gold standard	Wrist c...	0.00371	0.011	0.0037	**	T-test	105
## 2	Sys	Gold standard	Wrist c...	0.0256	0.038	0.0256	*	T-test	115
## 3	Sys	Wrist cuff 1	Wrist c...	0.449	0.45	0.4493	ns	T-test	110
## 4	Dia	Gold standard	Wrist c...	0.0886	0.16	0.089	ns	T-test	68
## 5	Dia	Gold standard	Wrist c...	0.107	0.16	0.107	ns	T-test	78
## 6	Dia	Wrist cuff 1	Wrist c...	0.293	0.29	0.293	ns	T-test	73

```
# plot sys
sys_baby_box2 <- boxplot_3_comparison(baby, "Device", "Sys", "Device",
                                     "Measurement (mmHg)",
                                     c(40,120),
                                     "Baby 4 Systolic pressure mean difference",
                                     "n = 9 measurements",
                                     "(method: paired t.test with FDR p.adjustment)",
                                     rater_colors,
                                     sys_test,
                                     "p.signif"
                                   )
sys_baby_box2 + geom_jitter(aes(shape=Disposition, colour=Disposition),size=3, alpha=
1) + scale_colour_manual(values = dispo_colors)
```

Baby 4 Systolic pressure mean difference

n = 9 measurements



(method: paired t.test with FDR p.adjustment)

```
ggsave("baby4_systolic.png", width=8, height=6, dpi=300)
```

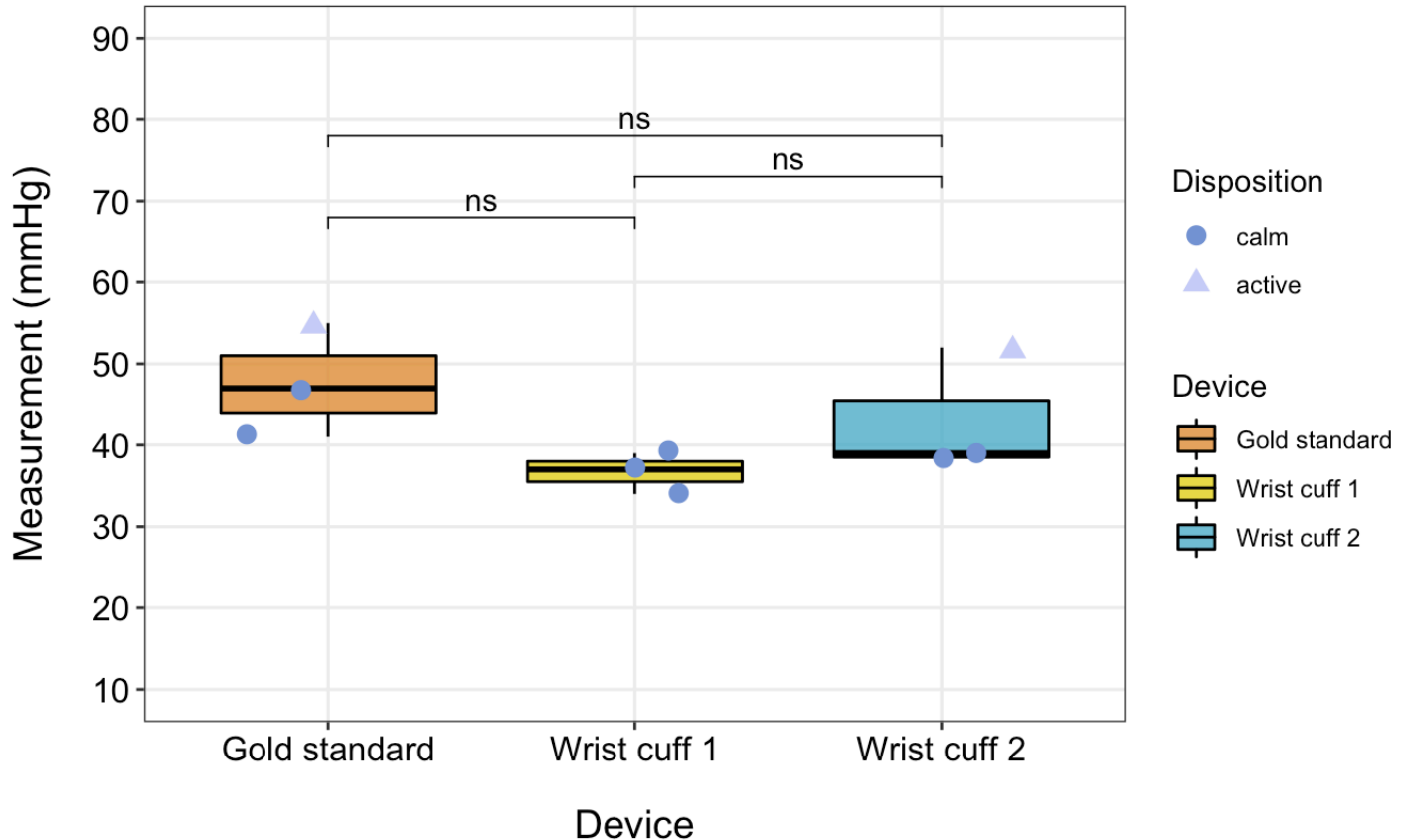
```
# plot dia
```

```
dia_baby_box2 <- boxplot_3_comparison(baby, "Device", "Dia", "Device",
                                     "Measurement (mmHg)",
                                     c(10,90),
                                     "Baby 4 Diastolic pressure mean difference",
                                     "n = 9 measurements",
                                     "(method: paired t.test with FDR p.adjustment)",
                                     rater_colors,
                                     dia_test,
                                     "p.signif"
                                   )
```

```
dia_baby_box2 + geom_jitter(aes(shape=Disposition, colour=Disposition),size=3, alpha=
1) + scale_colour_manual(values = dispo_colors)
```

Baby 4 Diastolic pressure mean difference

n = 9 measurements



(method: paired t.test with FDR p.adjustment)

```
ggsave("baby4_diastolic.png", width=8, height=6, dpi=300)
```

```

baby <- df %>% filter(BabyID == '8')

sys_test <- compare_means(Sys ~ Device,
                          data=baby,
                          method="t.test",
                          p.adjust.method = "fdr",
                          paired=TRUE,
                          comparisons = compare_devices)
sys_test <- sys_test %>% mutate(y.position = c(105, 115, 110))
dia_test <- compare_means(Dia ~ Device,
                          data=baby,
                          method="t.test",
                          p.adjust.method = "fdr",
                          paired=TRUE,
                          comparisons = compare_devices)
dia_test <- dia_test %>% mutate(y.position = c(68, 78, 73))
bind_rows(sys_test, dia_test)
    
```

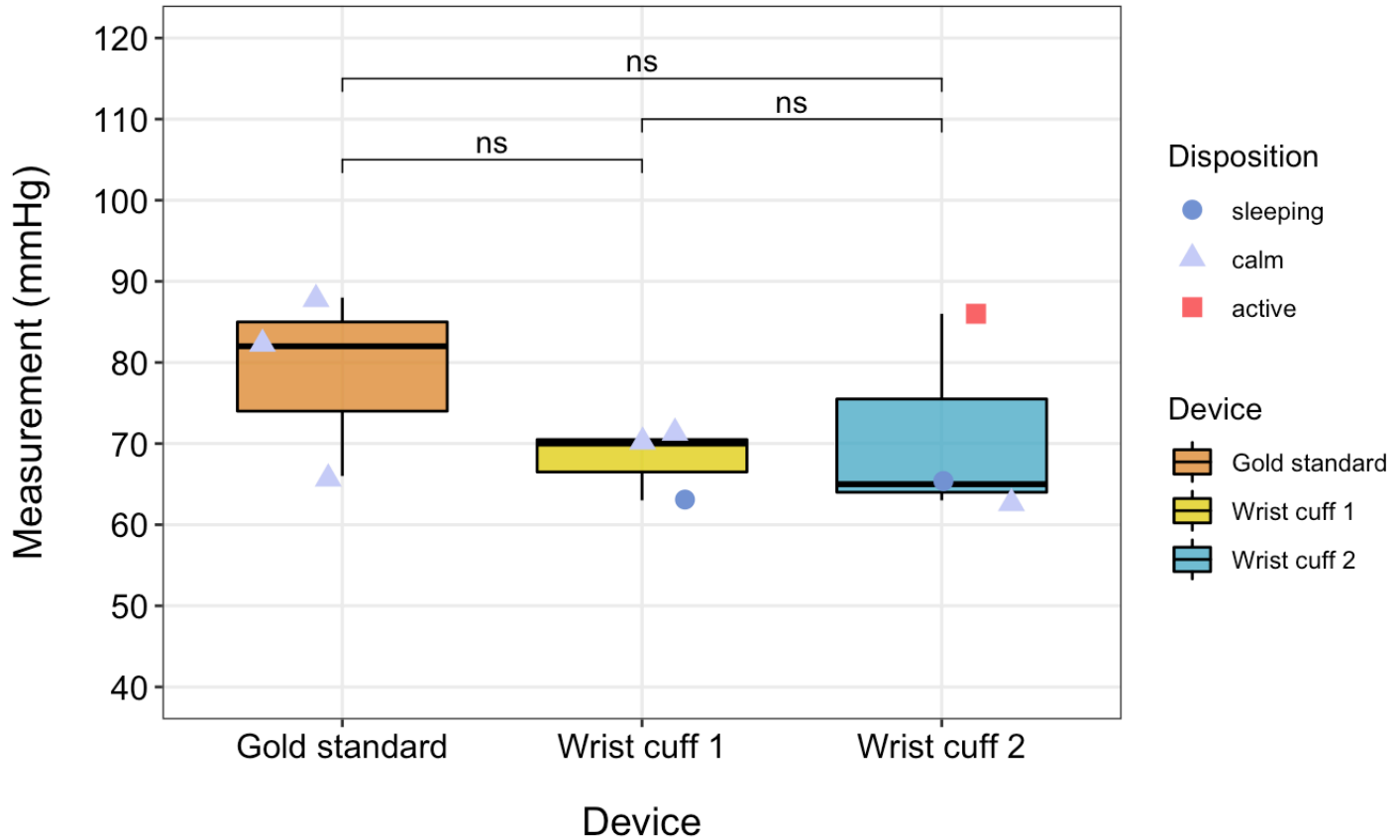
```
## # A tibble: 6 × 9
```

	.y.	group1	group2	p	p.adj	p.format	p.signif	method	y.position
	<chr>	<chr>	<chr>	<dbl>	<dbl>	<chr>	<chr>	<chr>	<dbl>
## 1	Sys	Gold standard	Wrist cuf...	0.284	0.43	0.28	ns	T-test	105
## 2	Sys	Gold standard	Wrist cuf...	0.269	0.43	0.27	ns	T-test	115
## 3	Sys	Wrist cuff 1	Wrist cuf...	0.654	0.65	0.65	ns	T-test	110
## 4	Dia	Gold standard	Wrist cuf...	0.715	0.71	0.71	ns	T-test	68
## 5	Dia	Gold standard	Wrist cuf...	0.549	0.71	0.55	ns	T-test	78
## 6	Dia	Wrist cuff 1	Wrist cuf...	0.225	0.68	0.23	ns	T-test	73

```
# plot sys
sys_baby_box2 <- boxplot_3_comparison(baby, "Device", "Sys", "Device",
                                     "Measurement (mmHg)",
                                     c(40,120),
                                     "Baby 8 Systolic pressure mean difference",
                                     "n = 9 measurements",
                                     "(method: paired t.test with FDR p.adjustment)",
                                     rater_colors,
                                     sys_test,
                                     "p.signif"
                                   )
sys_baby_box2+ geom_jitter(aes(shape=Disposition, colour=Disposition),size=3, alpha=
1) + scale_colour_manual(values = dispo_colors)
```

Baby 8 Systolic pressure mean difference

n = 9 measurements



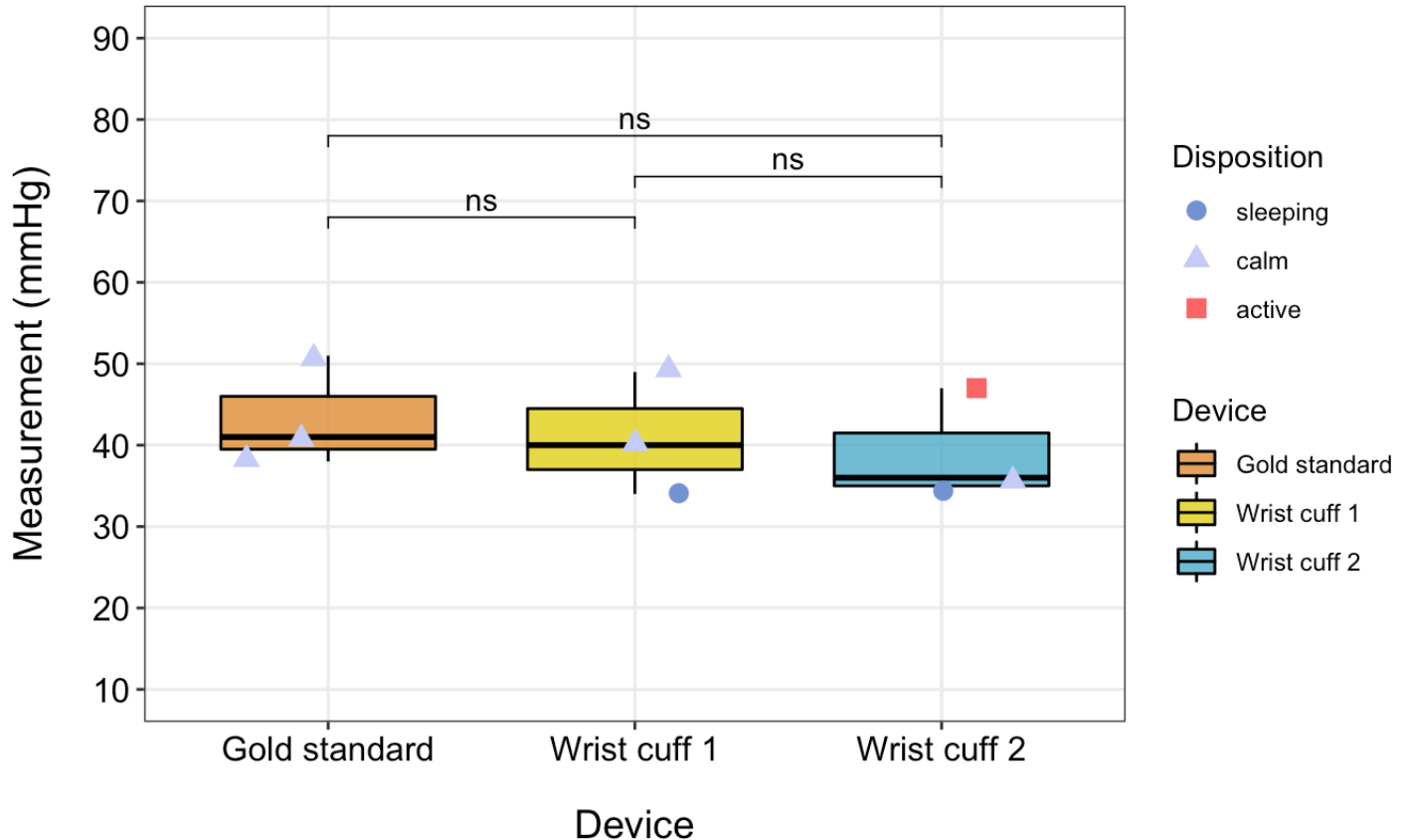
(method: paired t.test with FDR p.adjustment)

```
ggsave("baby8_systolic.png", width=8, height=6, dpi=300)

# plot dia
dia_baby_box2 <- boxplot_3_comparison(baby, "Device", "Dia", "Device",
                                     "Measurement (mmHg)",
                                     c(10,90),
                                     "Baby 8 Diastolic pressure mean difference",
                                     "n = 9 measurements",
                                     "(method: paired t.test with FDR p.adjustment)",
                                     rater_colors,
                                     dia_test,
                                     "p.signif"
                                     )
dia_baby_box2+ geom_jitter(aes(shape=Disposition, colour=Disposition),size=3, alpha=
1) + scale_colour_manual(values = dispo_colors)
```

Baby 8 Diastolic pressure mean difference

n = 9 measurements



(method: paired t.test with FDR p.adjustment)

```
ggsave("baby8_diastolic.png", width=8, height=6, dpi=300)
```

```

baby <- df %>% filter(BabyID == '10')

sys_test <- compare_means(Sys ~ Device,
                          data=baby,
                          method="t.test",
                          p.adjust.method = "fdr",
                          paired=TRUE,
                          comparisons = compare_devices)
sys_test <- sys_test %>% mutate(y.position = c(105, 115, 110))
dia_test <- compare_means(Dia ~ Device,
                          data=baby,
                          method="t.test",
                          p.adjust.method = "fdr",
                          paired=TRUE,
                          comparisons = compare_devices)
dia_test <- dia_test %>% mutate(y.position = c(68, 78, 73))
bind_rows(sys_test, dia_test)
    
```



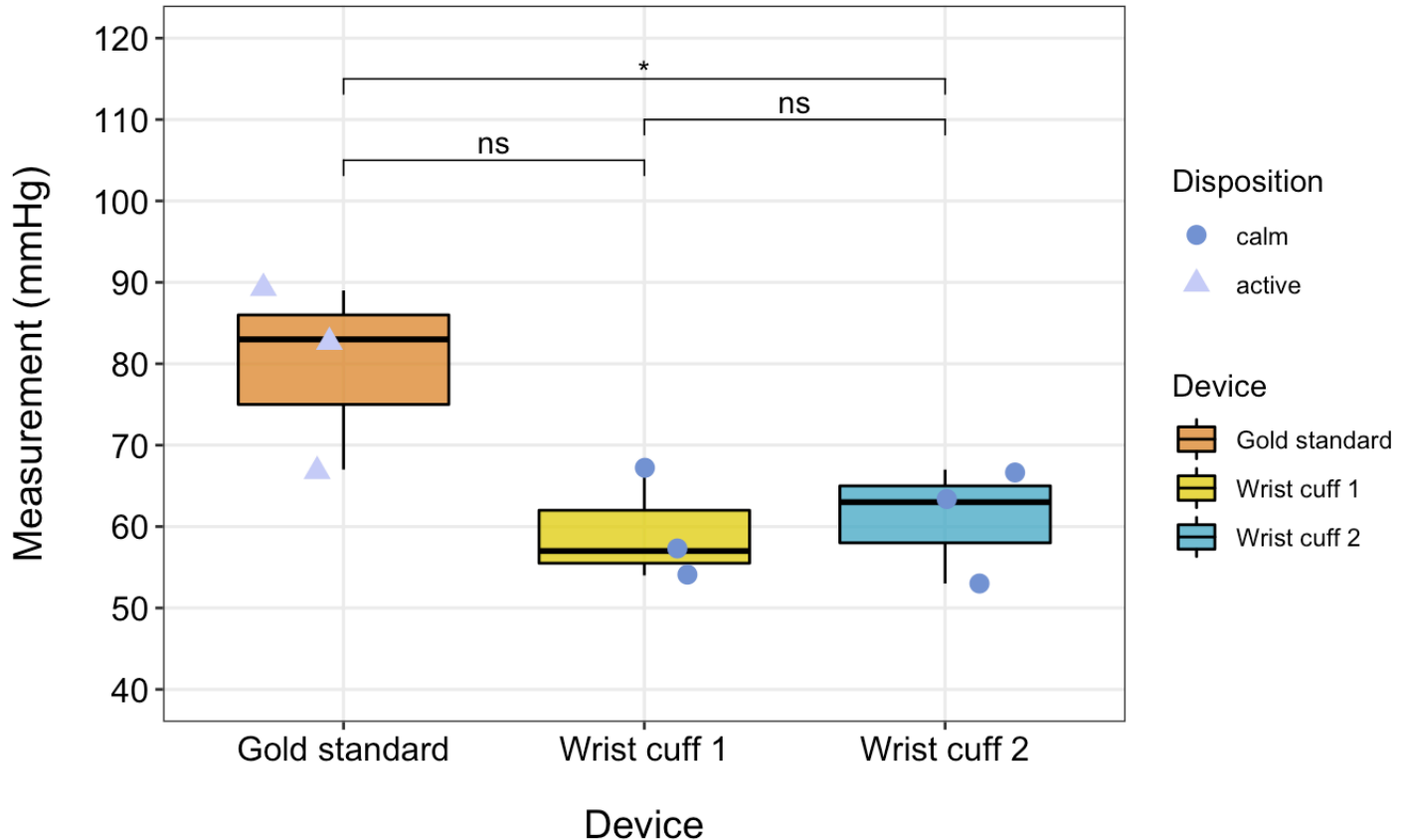
```
## # A tibble: 6 × 9
```

	.y.	group1	group2	p	p.adj	p.format	p.signif	method	y.position
	<chr>	<chr>	<chr>	<dbl>	<dbl>	<chr>	<chr>	<chr>	<dbl>
## 1	Sys	Gold standard	Wrist cu...	0.114	0.17	0.114	ns	T-test	105
## 2	Sys	Gold standard	Wrist cu...	0.0373	0.11	0.037	*	T-test	115
## 3	Sys	Wrist cuff 1	Wrist cu...	0.707	0.71	0.707	ns	T-test	110
## 4	Dia	Gold standard	Wrist cu...	0.0342	0.062	0.034	*	T-test	68
## 5	Dia	Gold standard	Wrist cu...	0.0414	0.062	0.041	*	T-test	78
## 6	Dia	Wrist cuff 1	Wrist cu...	0.624	0.62	0.624	ns	T-test	73

```
# plot sys
sys_baby_box2 <- boxplot_3_comparison(baby, "Device", "Sys", "Device",
                                     "Measurement (mmHg)",
                                     c(40,120),
                                     "Baby 10 Systolic pressure mean difference",
                                     "n = 9 measurements",
                                     "(method: paired t.test with FDR p.adjustment)",
                                     rater_colors,
                                     sys_test,
                                     "p.signif"
                                   )
sys_baby_box2 + geom_jitter(aes(shape=Disposition, colour=Disposition),size=3, alpha=
1) + scale_colour_manual(values = dispo_colors)
```

Baby 10 Systolic pressure mean difference

n = 9 measurements



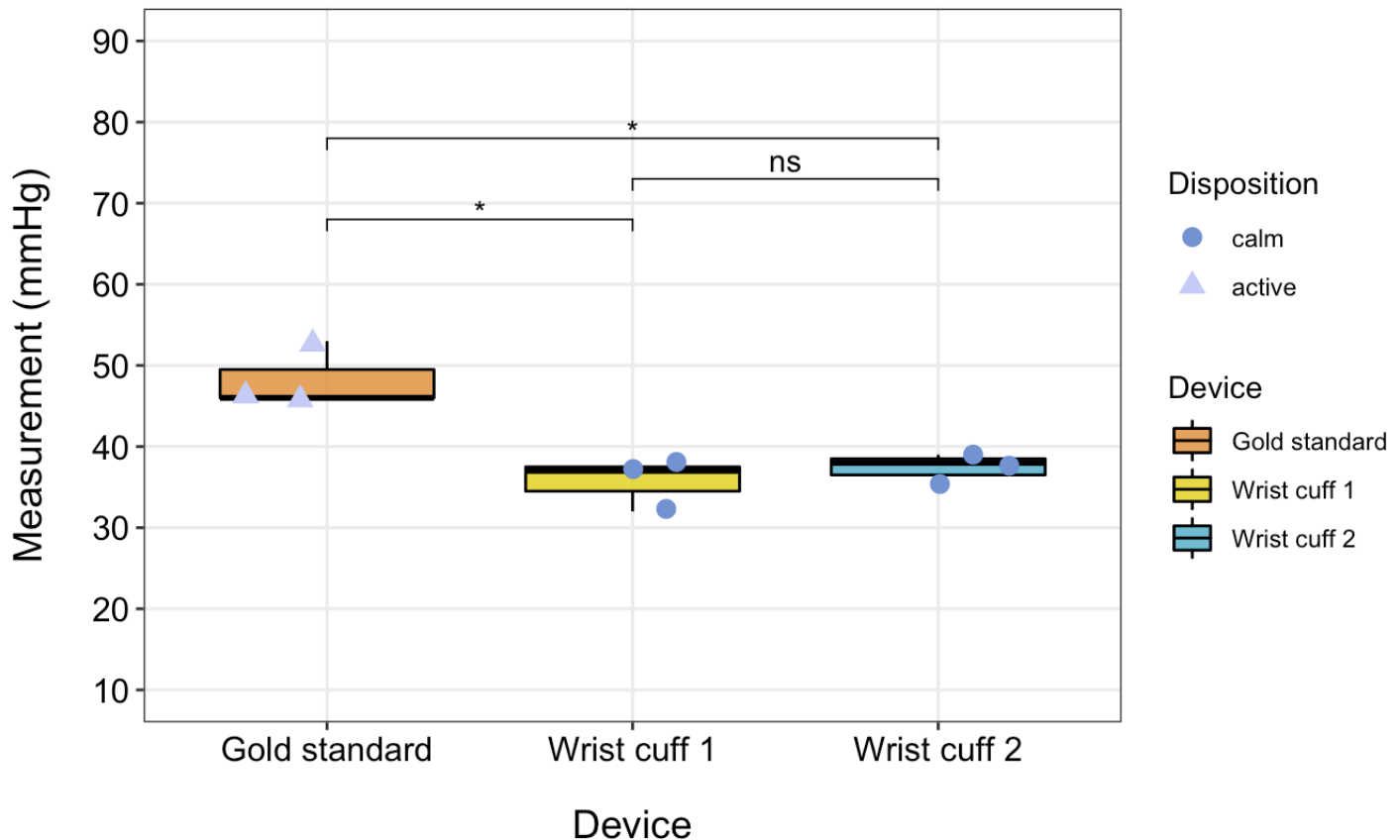
(method: paired t.test with FDR p.adjustment)

```
ggsave("baby10_systolic.png", width=8, height=6, dpi=300)

# plot dia
dia_baby_box2 <- boxplot_3_comparison(baby, "Device", "Dia", "Device",
                                     "Measurement (mmHg)",
                                     c(10,90),
                                     "Baby 10 Diastolic pressure mean difference",
                                     "n = 9 measurements",
                                     "(method: paired t.test with FDR p.adjustment)",
                                     rater_colors,
                                     dia_test,
                                     "p.signif"
                                     )
dia_baby_box2+ geom_jitter(aes(shape=Disposition, colour=Disposition),size=3, alpha=
1) + scale_colour_manual(values = dispo_colors)
```

Baby 10 Diastolic pressure mean difference

n = 9 measurements



(method: paired t.test with FDR p.adjustment)

```
ggsave("baby10_diastolic.png", width=8, height=6, dpi=300)
```

```

baby <- df %>% filter(BabyID == '12')

sys_test <- compare_means(Sys ~ Device,
                          data=baby,
                          method="t.test",
                          p.adjust.method = "fdr",
                          paired=TRUE,
                          comparisons = compare_devices)
sys_test <- sys_test %>% mutate(y.position = c(105, 115, 110))
dia_test <- compare_means(Dia ~ Device,
                          data=baby,
                          method="t.test",
                          p.adjust.method = "fdr",
                          paired=TRUE,
                          comparisons = compare_devices)
dia_test <- dia_test %>% mutate(y.position = c(68, 78, 73))
bind_rows(sys_test, dia_test)
    
```

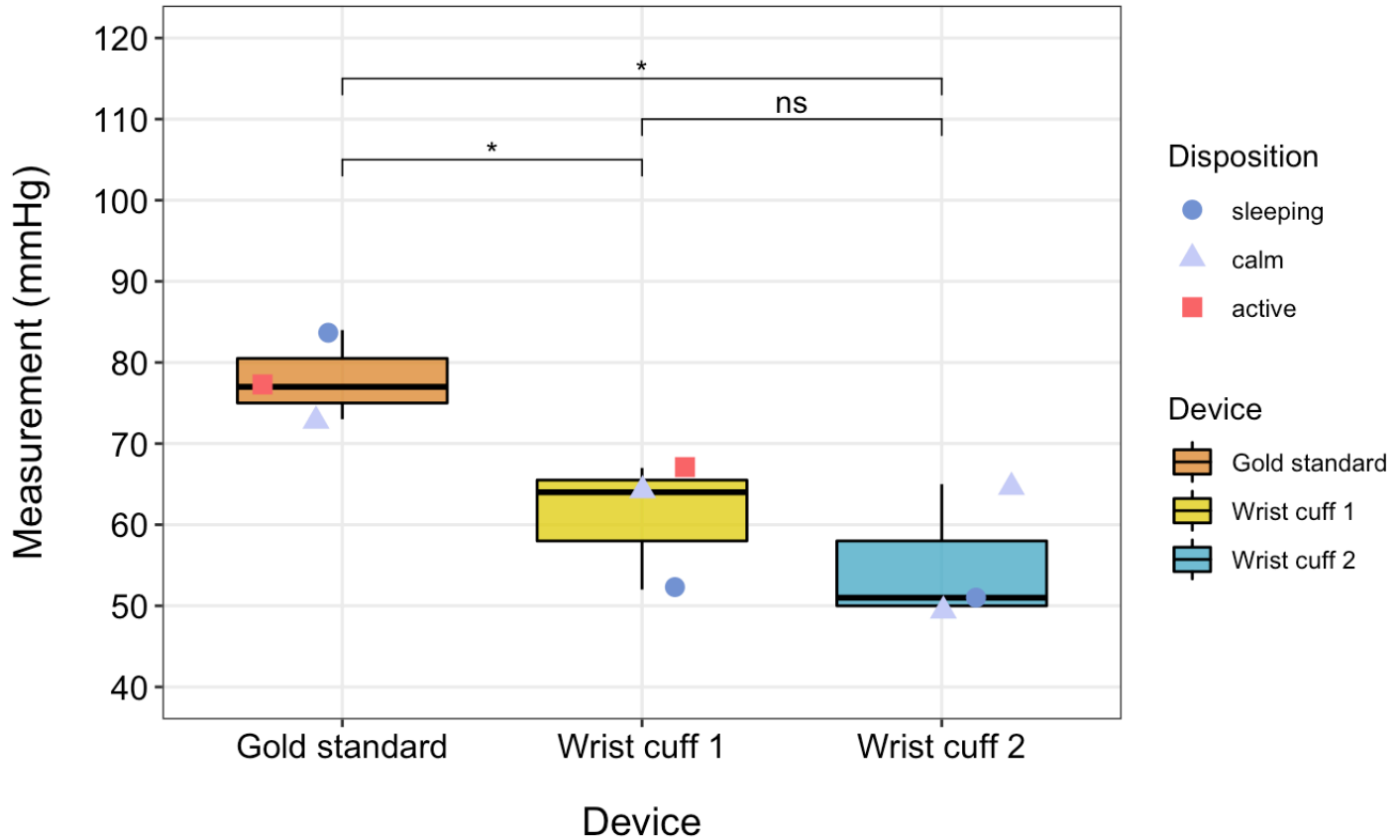
```
## # A tibble: 6 × 9
```

	.y.	group1	group2	p	p.adj	p.format	p.signif	method	y.position
	<chr>	<chr>	<chr>	<dbl>	<dbl>	<chr>	<chr>	<chr>	<dbl>
## 1	Sys	Gold standard	Wrist cu...	0.0401	0.06	0.040	*	T-test	105
## 2	Sys	Gold standard	Wrist cu...	0.0130	0.039	0.013	*	T-test	115
## 3	Sys	Wrist cuff 1	Wrist cu...	0.424	0.42	0.424	ns	T-test	110
## 4	Dia	Gold standard	Wrist cu...	0.0430	0.064	0.043	*	T-test	68
## 5	Dia	Gold standard	Wrist cu...	0.0187	0.056	0.019	*	T-test	78
## 6	Dia	Wrist cuff 1	Wrist cu...	0.706	0.71	0.706	ns	T-test	73

```
# plot sys
sys_baby_box2 <- boxplot_3_comparison(baby, "Device", "Sys", "Device",
                                     "Measurement (mmHg)",
                                     c(40,120),
                                     "Baby 12 Systolic pressure mean difference",
                                     "n = 9 measurements",
                                     "(method: paired t.test with FDR p.adjustment)",
                                     rater_colors,
                                     sys_test,
                                     "p.signif"
                                   )
sys_baby_box2 + geom_jitter(aes(shape=Disposition, colour=Disposition),size=3, alpha=
1) + scale_colour_manual(values = dispo_colors)
```

Baby 12 Systolic pressure mean difference

n = 9 measurements



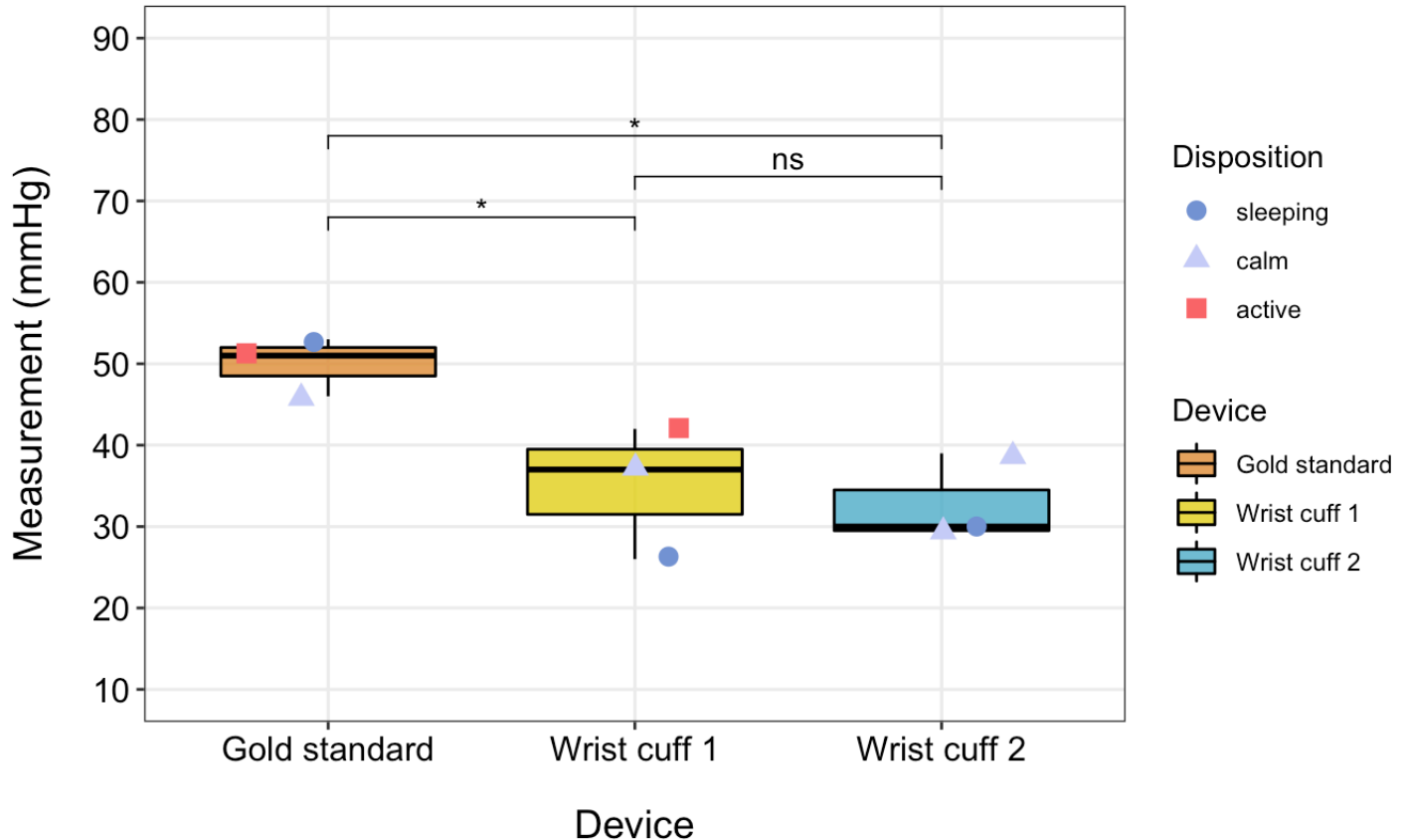
(method: paired t.test with FDR p.adjustment)

```
ggsave("baby12_systolic.png", width=8, height=6, dpi=300)

# plot dia
dia_baby_box2 <- boxplot_3_comparison(baby, "Device", "Dia", "Device",
                                     "Measurement (mmHg)",
                                     c(10,90),
                                     "Baby 12 Diastolic pressure mean difference",
                                     "n = 9 measurements",
                                     "(method: paired t.test with FDR p.adjustment)",
                                     rater_colors,
                                     dia_test,
                                     "p.signif"
                                     )
dia_baby_box2 + geom_jitter(aes(shape=Disposition, colour=Disposition),size=3, alpha=
1) + scale_colour_manual(values = dispo_colors)
```

Baby 12 Diastolic pressure mean difference

n = 9 measurements



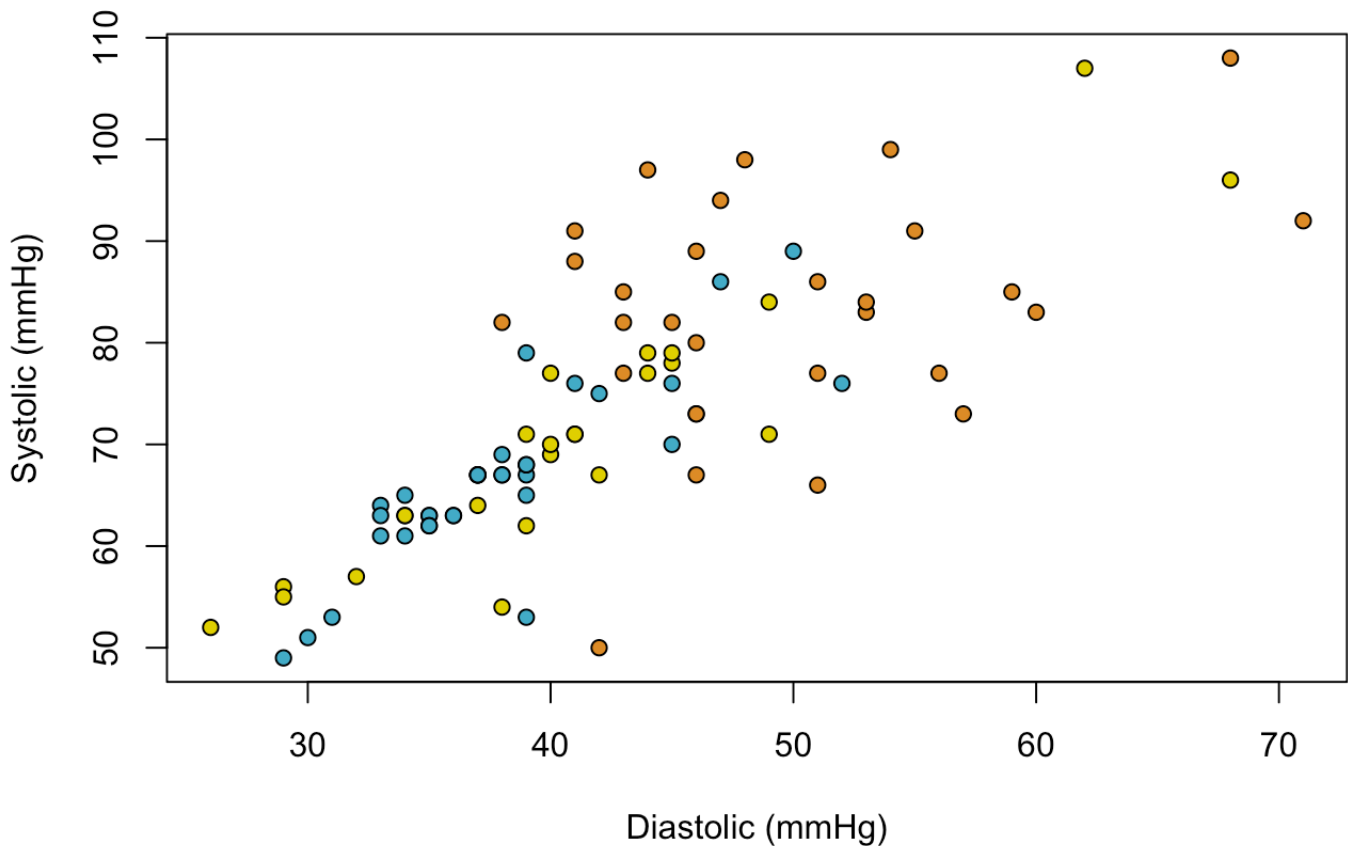
(method: paired t.test with FDR p.adjustment)

```
ggsave("baby12_diastolic.png", width=8, height=6, dpi=300)
```

regression analysis can be used to infer causal relationships between the independent and dependent variables.

```
plot(df$Dia, df$Sys, pch=21, bg=rater_colors[unclass(df$Device)],  
     main="Baby BP measurements", xlab="Diastolic (mmHg)", ylab="Systolic (mmHg)")
```

Baby BP measurements



```
# some dependency between axis variables  
# gap in between gold points means data will not be normal
```

[Testing correlations] - captures relationship between two random variables

- pearson is magnitude based
- spearman is rank based - spearman test, cant get exact p value, math behind it based on if rank of A bigger than rank of B, p value of spearman assumes you have n coin flips, non parametric because it can convert everything to coin flip. BUT if two values are the same, it can not calculate exact pvalue. But p value is close enough...

```
# r_p=cor(df$Sys, df$Dia, method = c("pearson"))  
# r_p  
# r_s=cor(df$Sys, df$Dia, method = c("spearman"))  
# r_s  
cor.test(df$Sys,df$Dia,method="pearson")
```

```
##
## Pearson's product-moment correlation
##
## data: df$Sys and df$Dia
## t = 11.46, df = 85, p-value < 2.2e-16
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
## 0.6801394 0.8502551
## sample estimates:
## cor
## 0.7791576
```

```
cor.test(df$Sys,df$Dia,method="spearman")
```

```
## Warning in cor.test.default(df$Sys, df$Dia, method = "spearman"): Cannot compute
## exact p-value with ties
```

```
##
## Spearman's rank correlation rho
##
## data: df$Sys and df$Dia
## S = 19907, p-value < 2.2e-16
## alternative hypothesis: true rho is not equal to 0
## sample estimates:
## rho
## 0.8185903
```

```
# function from file:///Users/mariapalafox/Desktop/TOOLBOXR/19_04_22_RregressionStats
_facetWrap.html
```

```
regression=function(df){
  reg_fun<-lm(formula = df$Sys ~ df$Disposition)
  #getting the slope, intercept, R square and adjusted R squared of
  #the regression function (with 3 decimals).
  slope<-round(coef(reg_fun)[2],3)
  intercept<-round(coef(reg_fun)[1],3)
  R2<-round(as.numeric(summary(reg_fun)[8]),3)
  R2.Adj<-round(as.numeric(summary(reg_fun)[9]),3)
  c(slope,intercept,R2,R2.Adj)
}
# funx call
regressions_data <- ddply(df,"Device", regression)
colnames(regressions_data) <-c("Device","slope","intercept","R2","R2.Adj")

regressions_data
```


##		Device	slope	intercept	R2	R2.Adj
## 1	Gold standard	10.500	73.000	0.113	0.007	
## 2	Wrist cuff 1	6.580	61.714	0.456	0.414	
## 3	Wrist cuff 2	4.965	61.222	0.408	0.363	

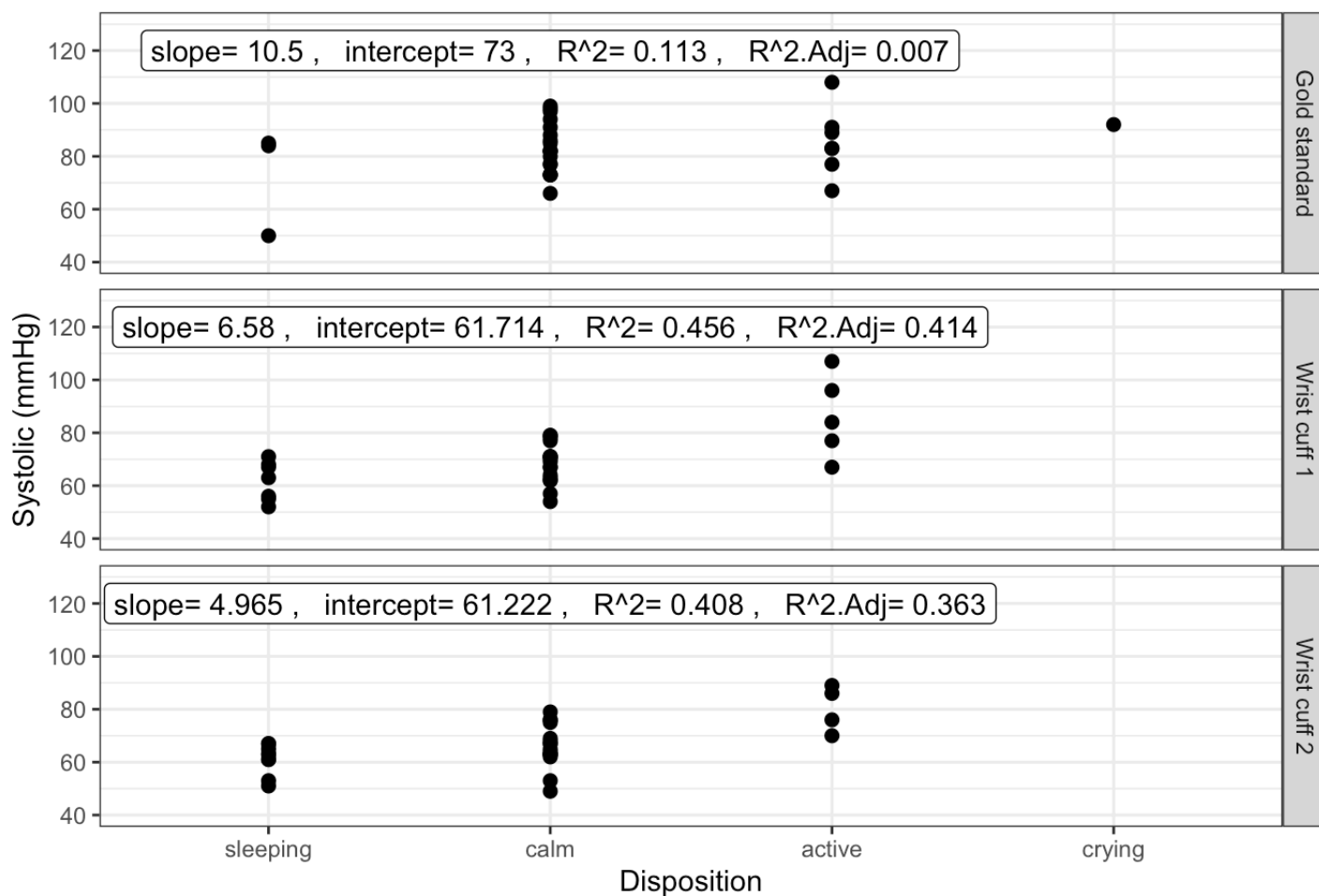
```

qplot(Disposition, Sys, data = df, size = I(2)) +
  geom_smooth(method="lm") +
  facet_grid(Device ~ .) +
  scale_y_continuous(limits = c(40, 130)) +
  # scale_x_continuous(limits = c(20, 80)) +
  labs(x="Disposition", y="Systolic (mmHg)") +
  ggtitle("Regressions") +
  theme_bw() +
  geom_label(data=regressions_data,
            inherit.aes=FALSE,
            aes(x = 2, y = 120,
              label=paste("slope=",slope," "," ", "intercept=",
                intercept," "," ", "R^2=",R2," "," ", "R^2.Adj=",R2.Adj)))

```

```
## `geom_smooth()` using formula 'y ~ x'
```

Regressions



```
regression=function(df){
  reg_fun<-lm(formula = df$Dia ~ df$Disposition) # CHANGE
  slope<-round(coef(reg_fun)[2],3)
  intercept<-round(coef(reg_fun)[1],3)
  R2<-round(as.numeric(summary(reg_fun)[8]),3)
  R2.Adj<-round(as.numeric(summary(reg_fun)[9]),3)
  c(slope,intercept,R2,R2.Adj)
}

regressions_data <- ddply(df,"Device", regression) # CHANGE
colnames(regressions_data) <-c("Device","slope","intercept","R2","R2.Adj")

regressions_data
```

```
##           Device slope intercept    R2 R2.Adj
## 1 Gold standard 1.556    46.000 0.406  0.335
## 2 Wrist cuff 1  5.958    33.571 0.526  0.489
## 3 Wrist cuff 2  3.625    34.000 0.663  0.637
```

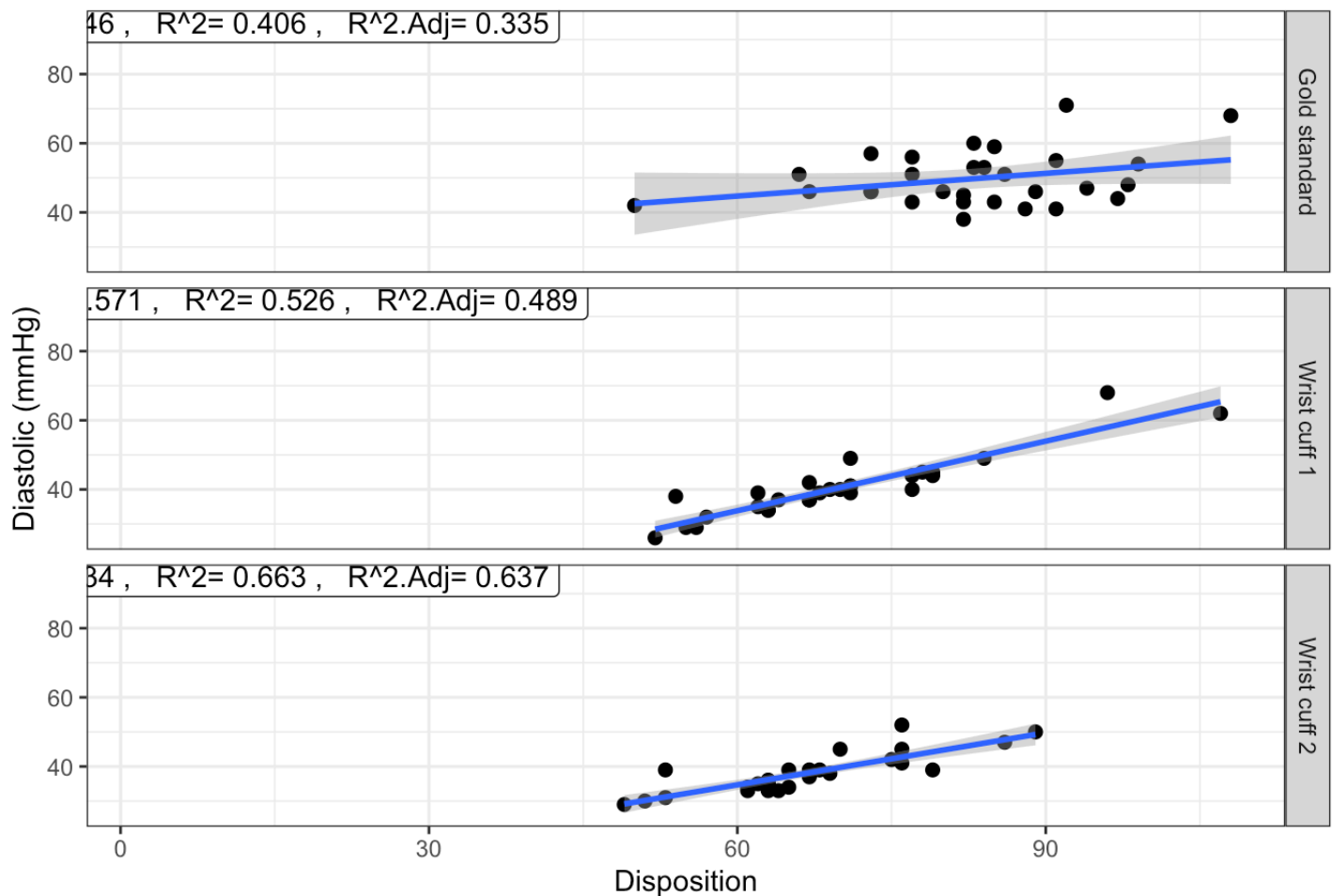
```

qplot(Sys, Dia, data = df, size = I(2)) + # CHANGE
  geom_smooth(method="lm") +
  facet_grid(Device ~ .) +
  #scale_y_continuous(limits = c(20, 100)) +
  # scale_x_continuous(limits = c(20, 80)) +
  labs(x="Disposition", y="Diastolic (mmHg)") +
  theme_bw() +
  ggtitle("Regressions") +
  geom_label(data=regressions_data,
            inherit.aes=FALSE,
            aes(x = 2, y = 95,
                label=paste("slope=",slope," "," ", "intercept=",
                             intercept," "," ", "R^2=",R2," "," ", "R^2.Adj=",R2.Adj)))

```

```
## `geom_smooth()` using formula 'y ~ x'
```

Regressions



```
# -1 says remove intercept term, you need to do this because you have 3 different spe
cies. now every species gets its own off set
gs <- df %>% filter(Device == 'Gold standard')
wc1 <- df %>% filter(Device == 'Wrist cuff 1')
wc2 <- df %>% filter(Device == 'Wrist cuff 2')

lm_gs = lm(Sys ~ Disposition, data=gs) #
summary(lm_gs)
```

```
##
## Call:
## lm(formula = Sys ~ Disposition, data = gs)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -23.00  -6.50  -1.50   7.50  22.57
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)      73.000      6.672  10.942 5.06e-11 ***
## Dispositioncalm    10.500      7.206   1.457   0.158
## Dispositionactive  12.429      7.974   1.559   0.132
## Dispositioncrying  19.000     13.343   1.424   0.167
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 11.56 on 25 degrees of freedom
## Multiple R-squared:  0.1132, Adjusted R-squared:  0.006731
## F-statistic: 1.063 on 3 and 25 DF,  p-value: 0.3824
```

```
# average all wrist col rows together
library(data.table)
```

```
##
## Attaching package: 'data.table'
```

```
## The following objects are masked from 'package:dplyr':
##
##      between, first, last
```

```
## The following object is masked from 'package:purrr':
##
##      transpose
```

```

dat <- df2 %>% select(BabyID, Session, Device, Measurement, Type, Disposition)
keys <- colnames(dat)[!grepl('Measurement', colnames(dat))]
X <- as.data.table(dat)
X[,list(mean_measurement = mean(Measurement)),keys]

```

```

##      BabyID Session      Device      Type Disposition mean_measurement
##  1:         1       1 Gold standard Systolic      calm              98
##  2:         1       1 Wrist cuff 1 Systolic      active             107
##  3:         1       1 Wrist cuff 2 Systolic      active              70
##  4:         2       1 Gold standard Systolic      calm              99
##  5:         2       1 Wrist cuff 1 Systolic      sleeping             67
## ---
## 170:        13       1 Wrist cuff 1 Diastolic      sleeping             41
## 171:        13       1 Wrist cuff 2 Diastolic      sleeping             37
## 172:        13       2 Gold standard Diastolic      calm              56
## 173:        13       2 Wrist cuff 1 Diastolic      active              44
## 174:        13       2 Wrist cuff 2 Diastolic      calm              39

```

```

# doesnt work because i cant average dispo

```

```

# make unique identifier
reg <- unite(df2, col='ID', c('BabyID', 'Session'), sep='-')
reg <- reg %>% select(ID, Device, Measurement, Type, Disposition)

gs <- reg %>% filter(Device == 'Gold standard')
wc1 <- reg %>% filter(Device == 'Wrist cuff 1')
wc2 <- reg %>% filter(Device == 'Wrist cuff 2')

gs

```

```

##      ID      Device Measurement      Type Disposition
##  1  1-1 Gold standard      98 Systolic      calm
##  2  2-1 Gold standard      99 Systolic      calm
##  3  2-2 Gold standard      97 Systolic      calm
##  4  2-3 Gold standard      82 Systolic      calm
##  5  3-1 Gold standard     108 Systolic      active
##  6  3-2 Gold standard      86 Systolic      calm
##  7  3-3 Gold standard      85 Systolic      sleeping
##  8  4-1 Gold standard      91 Systolic      active
##  9  4-2 Gold standard      91 Systolic      calm
## 10  4-3 Gold standard      94 Systolic      calm
## 11  5-1 Gold standard      77 Systolic      calm
## 12  6-1 Gold standard      73 Systolic      calm
## 13  7-1 Gold standard      80 Systolic      calm
## 14  7-2 Gold standard      85 Systolic      calm

```

## 15	8-1	Gold standard	66	Systolic	calm
## 16	8-2	Gold standard	82	Systolic	calm
## 17	8-3	Gold standard	88	Systolic	calm
## 18	9-1	Gold standard	50	Systolic	sleeping
## 19	9-2	Gold standard	73	Systolic	calm
## 20	10-1	Gold standard	83	Systolic	active
## 21	10-2	Gold standard	89	Systolic	active
## 22	10-3	Gold standard	67	Systolic	active
## 23	11-1	Gold standard	83	Systolic	active
## 24	11-2	Gold standard	92	Systolic	crying
## 25	12-1	Gold standard	84	Systolic	sleeping
## 26	12-2	Gold standard	77	Systolic	active
## 27	12-3	Gold standard	73	Systolic	calm
## 28	13-1	Gold standard	82	Systolic	calm
## 29	13-2	Gold standard	77	Systolic	calm
## 30	1-1	Gold standard	48	Diastolic	calm
## 31	2-1	Gold standard	54	Diastolic	calm
## 32	2-2	Gold standard	44	Diastolic	calm
## 33	2-3	Gold standard	43	Diastolic	calm
## 34	3-1	Gold standard	68	Diastolic	active
## 35	3-2	Gold standard	51	Diastolic	calm
## 36	3-3	Gold standard	43	Diastolic	sleeping
## 37	4-1	Gold standard	55	Diastolic	active
## 38	4-2	Gold standard	41	Diastolic	calm
## 39	4-3	Gold standard	47	Diastolic	calm
## 40	5-1	Gold standard	43	Diastolic	calm
## 41	6-1	Gold standard	57	Diastolic	calm
## 42	7-1	Gold standard	46	Diastolic	calm
## 43	7-2	Gold standard	59	Diastolic	calm
## 44	8-1	Gold standard	51	Diastolic	calm
## 45	8-2	Gold standard	38	Diastolic	calm
## 46	8-3	Gold standard	41	Diastolic	calm
## 47	9-1	Gold standard	42	Diastolic	sleeping
## 48	9-2	Gold standard	46	Diastolic	calm
## 49	10-1	Gold standard	53	Diastolic	active
## 50	10-2	Gold standard	46	Diastolic	active
## 51	10-3	Gold standard	46	Diastolic	active
## 52	11-1	Gold standard	60	Diastolic	active
## 53	11-2	Gold standard	71	Diastolic	crying
## 54	12-1	Gold standard	53	Diastolic	sleeping
## 55	12-2	Gold standard	51	Diastolic	active
## 56	12-3	Gold standard	46	Diastolic	calm
## 57	13-1	Gold standard	45	Diastolic	calm
## 58	13-2	Gold standard	56	Diastolic	calm

wc1

##	ID	Device Measurement	Type	Disposition
----	----	--------------------	------	-------------

## 1	1-1 Wrist cuff 1	107	Systolic	active
## 2	2-1 Wrist cuff 1	67	Systolic	sleeping
## 3	2-2 Wrist cuff 1	96	Systolic	active
## 4	2-3 Wrist cuff 1	56	Systolic	sleeping
## 5	3-1 Wrist cuff 1	79	Systolic	calm
## 6	3-2 Wrist cuff 1	84	Systolic	active
## 7	3-3 Wrist cuff 1	69	Systolic	calm
## 8	4-1 Wrist cuff 1	67	Systolic	calm
## 9	4-2 Wrist cuff 1	63	Systolic	calm
## 10	4-3 Wrist cuff 1	71	Systolic	calm
## 11	5-1 Wrist cuff 1	77	Systolic	calm
## 12	6-1 Wrist cuff 1	78	Systolic	calm
## 13	7-1 Wrist cuff 1	71	Systolic	calm
## 14	7-2 Wrist cuff 1	68	Systolic	sleeping
## 15	8-1 Wrist cuff 1	70	Systolic	calm
## 16	8-2 Wrist cuff 1	63	Systolic	sleeping
## 17	8-3 Wrist cuff 1	71	Systolic	calm
## 18	9-1 Wrist cuff 1	55	Systolic	sleeping
## 19	9-2 Wrist cuff 1	62	Systolic	calm
## 20	10-1 Wrist cuff 1	67	Systolic	calm
## 21	10-2 Wrist cuff 1	54	Systolic	calm
## 22	10-3 Wrist cuff 1	57	Systolic	calm
## 23	11-1 Wrist cuff 1	62	Systolic	calm
## 24	11-2 Wrist cuff 1	79	Systolic	calm
## 25	12-1 Wrist cuff 1	64	Systolic	calm
## 26	12-2 Wrist cuff 1	67	Systolic	active
## 27	12-3 Wrist cuff 1	52	Systolic	sleeping
## 28	13-1 Wrist cuff 1	71	Systolic	sleeping
## 29	13-2 Wrist cuff 1	77	Systolic	active
## 30	1-1 Wrist cuff 1	62	Diastolic	active
## 31	2-1 Wrist cuff 1	37	Diastolic	sleeping
## 32	2-2 Wrist cuff 1	68	Diastolic	active
## 33	2-3 Wrist cuff 1	29	Diastolic	sleeping
## 34	3-1 Wrist cuff 1	44	Diastolic	calm
## 35	3-2 Wrist cuff 1	49	Diastolic	active
## 36	3-3 Wrist cuff 1	40	Diastolic	calm
## 37	4-1 Wrist cuff 1	37	Diastolic	calm
## 38	4-2 Wrist cuff 1	34	Diastolic	calm
## 39	4-3 Wrist cuff 1	39	Diastolic	calm
## 40	5-1 Wrist cuff 1	40	Diastolic	calm
## 41	6-1 Wrist cuff 1	45	Diastolic	calm
## 42	7-1 Wrist cuff 1	41	Diastolic	calm
## 43	7-2 Wrist cuff 1	39	Diastolic	sleeping
## 44	8-1 Wrist cuff 1	40	Diastolic	calm
## 45	8-2 Wrist cuff 1	34	Diastolic	sleeping
## 46	8-3 Wrist cuff 1	49	Diastolic	calm
## 47	9-1 Wrist cuff 1	29	Diastolic	sleeping
## 48	9-2 Wrist cuff 1	39	Diastolic	calm
## 49	10-1 Wrist cuff 1	37	Diastolic	calm

```
## 50 10-2 Wrist cuff 1      38 Diastolic      calm
## 51 10-3 Wrist cuff 1      32 Diastolic      calm
## 52 11-1 Wrist cuff 1      35 Diastolic      calm
## 53 11-2 Wrist cuff 1      45 Diastolic      calm
## 54 12-1 Wrist cuff 1      37 Diastolic      calm
## 55 12-2 Wrist cuff 1      42 Diastolic      active
## 56 12-3 Wrist cuff 1      26 Diastolic      sleeping
## 57 13-1 Wrist cuff 1      41 Diastolic      sleeping
## 58 13-2 Wrist cuff 1      44 Diastolic      active
```

```
#reg <- bind_rows
# colnames(gs)[colnames(gs) == "Measurement"] = "Gold standard"
# colnames(wc1)[colnames(wc1) == "Measurement"] = "Wrist cuff 1"
# colnames(wc2)[colnames(wc2) == "Measurement"] = "Wrist cuff 2"
# reg <- list(gs,wc1,wc2) %>% reduce(inner_join, by='ID')
# reg
```

```
regression=function(df){
  #setting the regression function.
  reg_fun<-lm(formula= df$Sys ~ df$Dia)
  #getting the slope, intercept, R square and adjusted R squared of
  #the regression function (with 3 decimals).
  slope<-round(coef(reg_fun)[2],3)
  intercept<-round(coef(reg_fun)[1],3)
  R2<-round(as.numeric(summary(reg_fun)[8]),3)
  R2.Adj<-round(as.numeric(summary(reg_fun)[9]),3)
  c(slope,intercept,R2,R2.Adj)
}

regressions_data <- ddply(df,"Device", regression)

colnames(regressions_data) <-c("Device","slope","intercept","R2","R2.Adj")

qplot(Dia, Sys, data = df, size = I(2)) +
  geom_smooth(method="lm") +
  facet_grid(Device ~ .) +
  scale_y_continuous(limits = c(40, 130)) +
  scale_x_continuous(limits = c(20, 80)) +
  labs(x="Diastolic (mmHg)", y="Systolic (mmHg)") +
  ggtitle("Regressions") +
  geom_label(data=regressions_data,
            inherit.aes=FALSE,
            aes(x = 50, y = 120,
                label=paste("slope=",slope," "," ", "intercept=",
                             intercept," "," ", "R^2=",R2," "," ", "R^2.Adj=",R2.Adj)))
```



```
## `geom_smooth()` using formula 'y ~ x'
```

Regressions

