summary stats SA1

library(tidyverse)

## ── Attaching packages ─────────────────────────────────────── tidyverse 1.3.2 ──  
## ✔ ggplot2 3.3.6 ✔ purrr 0.3.4  
## ✔ tibble 3.1.8 ✔ dplyr 1.0.9  
## ✔ tidyr 1.2.0 ✔ stringr 1.4.0  
## ✔ readr 2.1.2 ✔ forcats 0.5.1  
## ── Conflicts ────────────────────────────────────────── tidyverse\_conflicts() ──  
## ✖ dplyr::filter() masks stats::filter()  
## ✖ dplyr::lag() masks stats::lag()

library(flextable)

##   
## Attaching package: 'flextable'  
##   
## The following object is masked from 'package:purrr':  
##   
## compose

library(readxl)  
library(extrafont)

## Registering fonts with R

library(forcats)  
library(writexl)

#times new roman tables  
my\_ft\_theme <- function(ft, ...) {  
 # Remove vertical cell padding  
 ft <- padding(ft, padding.top = 0, padding.bottom = 0, part = "all")  
   
 # Change font to TNR 11  
 ft <- font(ft, fontname = "Times New Roman", part = "all")  
 ft <- fontsize(ft, part = "all", size = 12)  
 ft  
}

headscan\_full<-read\_excel("C:\\Users\\19177\\OneDrive - Colostate\\Desktop\\Dissertation\\headscan\_dissertation\\headscan\_full.xlsx")  
headscan\_full$coder <- as.factor(headscan\_full$coder)  
headscan\_full$gender <- as.factor(headscan\_full$gender)  
headscan\_full$race\_eth <- as.factor(headscan\_full$race\_eth)  
headscan\_full$age\_group <- as.factor(headscan\_full$age\_group)  
#reordering factors by frequency  
headscan\_full$race\_eth <- factor(headscan\_full$race\_eth, levels = c("white", "Black", "LatinX", "Asian", "Other", "AIAN", "PTNS", "NHOPI"))  
str(headscan\_full)

## tibble [2,017 × 33] (S3: tbl\_df/tbl/data.frame)  
## $ ID : chr [1:2017] "400-20201012-002" "400-20201012-003" "400-20201012-004" "400-20201012-005" ...  
## $ AA\_C : num [1:2017] 6.5 5.5 7 5.8 6.7 6 5.9 5.9 6.5 6.5 ...  
## $ BGl\_C : num [1:2017] 31.5 28.9 29.3 31.3 28.8 30.6 32 NA 30 27.7 ...  
## $ BiW\_C : num [1:2017] 13 12.7 14.3 14 13.7 13 14.1 13.8 14.3 15 ...  
## $ BiW\_L : num [1:2017] 11.5 10.8 12.1 10.9 10.4 10.6 10.9 11.1 11.3 11.6 ...  
## $ ChCh\_C : num [1:2017] 6.2 6.4 6.8 7 7 7 6.7 6.9 6.7 6.3 ...  
## $ GoSub\_C : num [1:2017] 9.3 9.3 11.5 9.3 10.3 10 7.9 10.6 8.5 10.2 ...  
## $ NRB\_L : num [1:2017] 1.7 1.8 1.9 2.1 1.9 1.4 1.7 1.8 1.6 1.7 ...  
## $ ProA\_L : num [1:2017] 2.8 2.5 3.1 2.3 2.8 2.8 2.6 2.7 3.2 2.8 ...  
## $ ProA\_C : num [1:2017] 3.1 2.7 3.3 2.7 3.1 2.9 2.7 2.9 3.4 3.1 ...  
## $ ProS\_C : num [1:2017] 1.8 2 1.4 1.3 2.2 2.2 1.9 1.4 2.6 2.4 ...  
## $ ProS\_L : num [1:2017] 1.7 1.8 1.4 1.3 2 2 1.8 1.2 2.4 2.2 ...  
## $ SelP\_C : num [1:2017] 4.2 4.1 5.1 4.5 4.7 4.8 4.6 4.2 4.7 4.4 ...  
## $ SelP\_L : num [1:2017] 4.2 4.1 5.1 4.4 4.7 4.8 4.6 4.1 4.6 4.4 ...  
## $ SelDH\_C : num [1:2017] 1.5 0.9 0.9 1.1 1.3 1.5 0.9 0.9 1.2 1.4 ...  
## $ SelM\_L : num [1:2017] 12.2 9.9 13 11.5 11.9 12.6 11.7 11.2 11.7 11.7 ...  
## $ SnasM\_C : num [1:2017] 8.2 5.5 8.4 7.4 7.3 8 7.8 7.6 6.4 7.5 ...  
## $ SmanM\_C : num [1:2017] 5.9 5.1 4.5 4.3 3.3 3.4 5.5 3.7 6.1 4.1 ...  
## $ SmanM\_L : num [1:2017] 5.5 5 4.5 4.2 3.3 3.4 5 3.6 5.9 4 ...  
## $ SnasM\_L : num [1:2017] 7.5 5.3 7.8 6.9 6.7 7.6 6.9 7.1 6.2 6.9 ...  
## $ TrHO\_C : num [1:2017] 17.9 16.3 16.9 16.6 15.9 16.2 16.9 NA 16.7 16.6 ...  
## $ TrEJ\_C : num [1:2017] 4 3.2 3.9 2.9 4.6 4.2 2.9 3.2 2.9 3.3 ...  
## $ TrGo\_C : num [1:2017] 8.4 5.7 7 6.1 6.8 7 7.5 6.1 6.7 6.4 ...  
## $ TrSel\_C : num [1:2017] 14.9 13.8 15 13.3 14 15.1 14 13.8 15.6 14.3 ...  
## $ TrSman\_C : num [1:2017] 17.7 14.5 17.8 14.7 15.7 16.4 14.9 15.9 15.1 16 ...  
## $ TrSnas\_C : num [1:2017] 16.3 14.2 16.7 14.5 15.2 15.7 14.8 14.9 15.7 NA ...  
## $ TrTr\_C : num [1:2017] 29.6 27.6 29.2 27.3 27.9 30 28.3 27.5 30.7 28.6 ...  
## $ TrTr\_L : num [1:2017] 15.5 14.1 15.6 14.9 14.6 14.6 14.7 15.1 15.7 14.4 ...  
## $ coder : Factor w/ 4 levels "Chandler","Isabel",..: 4 4 4 4 4 4 4 4 4 4 ...  
## $ age : num [1:2017] 31 49 49 34 49 55 26 18 25 27 ...  
## $ gender : Factor w/ 4 levels "Female","Male",..: 2 1 2 2 2 2 2 2 2 2 ...  
## $ race\_eth : Factor w/ 8 levels "white","Black",..: 2 1 1 1 1 1 2 2 1 1 ...  
## $ age\_group: Factor w/ 3 levels "18-36","37-54",..: 1 2 2 1 2 3 1 1 1 1 ...

headscan\_full$AA\_C <- headscan\_full$AA\_C \*10  
headscan\_full$BGl\_C <- headscan\_full$BGl\_C \* 10  
headscan\_full$BiW\_C <- headscan\_full$BiW\_C \*10  
headscan\_full$BiW\_L <- headscan\_full$BiW\_L \*10  
headscan\_full$ChCh\_C <- headscan\_full$ChCh\_C \*10  
headscan\_full$GoSub\_C <- headscan\_full$GoSub\_C \*10  
headscan\_full$NRB\_L <- headscan\_full$NRB\_L \*10  
headscan\_full$ProA\_L <- headscan\_full$ProA\_L \*10  
headscan\_full$ProA\_C <- headscan\_full$ProA\_C \*10  
headscan\_full$ProS\_C <- headscan\_full$ProS\_C \*10  
headscan\_full$ProS\_L <- headscan\_full$ProS\_L \*10  
headscan\_full$SelP\_C <- headscan\_full$SelP\_C \*10  
headscan\_full$SelP\_L <- headscan\_full$SelP\_L \*10  
headscan\_full$SelDH\_C <- headscan\_full$SelDH\_C \*10  
headscan\_full$SelM\_L <- headscan\_full$SelM\_L \*10  
headscan\_full$SnasM\_C <- headscan\_full$SnasM\_C \*10  
headscan\_full$SmanM\_C <- headscan\_full$SmanM\_C \*10  
headscan\_full$SmanM\_L <- headscan\_full$SmanM\_L \*10  
headscan\_full$SnasM\_L <- headscan\_full$SnasM\_L \*10  
headscan\_full$TrHO\_C <- headscan\_full$TrHO\_C \*10  
headscan\_full$TrEJ\_C <- headscan\_full$TrEJ\_C \*10  
headscan\_full$TrGo\_C <- headscan\_full$TrGo\_C \*10  
headscan\_full$TrSel\_C <- headscan\_full$TrSel\_C \*10  
headscan\_full$TrSman\_C <- headscan\_full$TrSman\_C \*10  
headscan\_full$TrSnas\_C <- headscan\_full$TrSnas\_C \*10  
headscan\_full$TrTr\_C <- headscan\_full$TrTr\_C \*10  
headscan\_full$TrTr\_L <- headscan\_full$TrTr\_L \*10  
  
str(headscan\_full)

## tibble [2,017 × 33] (S3: tbl\_df/tbl/data.frame)  
## $ ID : chr [1:2017] "400-20201012-002" "400-20201012-003" "400-20201012-004" "400-20201012-005" ...  
## $ AA\_C : num [1:2017] 65 55 70 58 67 60 59 59 65 65 ...  
## $ BGl\_C : num [1:2017] 315 289 293 313 288 306 320 NA 300 277 ...  
## $ BiW\_C : num [1:2017] 130 127 143 140 137 130 141 138 143 150 ...  
## $ BiW\_L : num [1:2017] 115 108 121 109 104 106 109 111 113 116 ...  
## $ ChCh\_C : num [1:2017] 62 64 68 70 70 70 67 69 67 63 ...  
## $ GoSub\_C : num [1:2017] 93 93 115 93 103 100 79 106 85 102 ...  
## $ NRB\_L : num [1:2017] 17 18 19 21 19 14 17 18 16 17 ...  
## $ ProA\_L : num [1:2017] 28 25 31 23 28 28 26 27 32 28 ...  
## $ ProA\_C : num [1:2017] 31 27 33 27 31 29 27 29 34 31 ...  
## $ ProS\_C : num [1:2017] 18 20 14 13 22 22 19 14 26 24 ...  
## $ ProS\_L : num [1:2017] 17 18 14 13 20 20 18 12 24 22 ...  
## $ SelP\_C : num [1:2017] 42 41 51 45 47 48 46 42 47 44 ...  
## $ SelP\_L : num [1:2017] 42 41 51 44 47 48 46 41 46 44 ...  
## $ SelDH\_C : num [1:2017] 15 9 9 11 13 15 9 9 12 14 ...  
## $ SelM\_L : num [1:2017] 122 99 130 115 119 126 117 112 117 117 ...  
## $ SnasM\_C : num [1:2017] 82 55 84 74 73 80 78 76 64 75 ...  
## $ SmanM\_C : num [1:2017] 59 51 45 43 33 34 55 37 61 41 ...  
## $ SmanM\_L : num [1:2017] 55 50 45 42 33 34 50 36 59 40 ...  
## $ SnasM\_L : num [1:2017] 75 53 78 69 67 76 69 71 62 69 ...  
## $ TrHO\_C : num [1:2017] 179 163 169 166 159 162 169 NA 167 166 ...  
## $ TrEJ\_C : num [1:2017] 40 32 39 29 46 42 29 32 29 33 ...  
## $ TrGo\_C : num [1:2017] 84 57 70 61 68 70 75 61 67 64 ...  
## $ TrSel\_C : num [1:2017] 149 138 150 133 140 151 140 138 156 143 ...  
## $ TrSman\_C : num [1:2017] 177 145 178 147 157 164 149 159 151 160 ...  
## $ TrSnas\_C : num [1:2017] 163 142 167 145 152 157 148 149 157 NA ...  
## $ TrTr\_C : num [1:2017] 296 276 292 273 279 300 283 275 307 286 ...  
## $ TrTr\_L : num [1:2017] 155 141 156 149 146 146 147 151 157 144 ...  
## $ coder : Factor w/ 4 levels "Chandler","Isabel",..: 4 4 4 4 4 4 4 4 4 4 ...  
## $ age : num [1:2017] 31 49 49 34 49 55 26 18 25 27 ...  
## $ gender : Factor w/ 4 levels "Female","Male",..: 2 1 2 2 2 2 2 2 2 2 ...  
## $ race\_eth : Factor w/ 8 levels "white","Black",..: 2 1 1 1 1 1 2 2 1 1 ...  
## $ age\_group: Factor w/ 3 levels "18-36","37-54",..: 1 2 2 1 2 3 1 1 1 1 ...

#HTML Table format   
#AACrace\_sumstats %>%   
 #kbl(caption = "Alare to Alare Contour SumStats by Race/Ethnicity") %>%   
 #kable\_styling(bootstrap\_options = c("striped", "hover", "condensed"), full\_width = TRUE)

#exploring NA values for measurements  
measureNAsums <- colSums(is.na(headscan\_full))  
  
measureNAprops <- colMeans(is.na(headscan\_full))  
  
measureNAprops1 <- as.data.frame(measureNAprops)  
measureNAprops1 <- rownames\_to\_column(measureNAprops1, "measure\_name")  
measureNAprops1 <- measureNAprops1 %>% slice(-c(1, 29:33))  
  
measureNAsums1 <- as.data.frame(measureNAsums)  
measureNAsums1 <- rownames\_to\_column(measureNAsums1, "measure\_name")  
measureNAsums1 <- measureNAsums1 %>% slice(-c(1, 29:33))  
  
measureNAs <- inner\_join(measureNAprops1, measureNAsums1, by = "measure\_name", desc)  
  
measureNAs$measureNAprops<- round(measureNAs$measureNAprops, digits=4)  
  
measureNAs$measure\_name <- fct\_reorder(measureNAs$measure\_name, measureNAs$measureNAsums, .desc=TRUE)  
  
str(measureNAs$measure\_name)

## Factor w/ 27 levels "BGl\_C","TrHO\_C",..: 19 1 20 21 12 9 24 23 22 16 ...

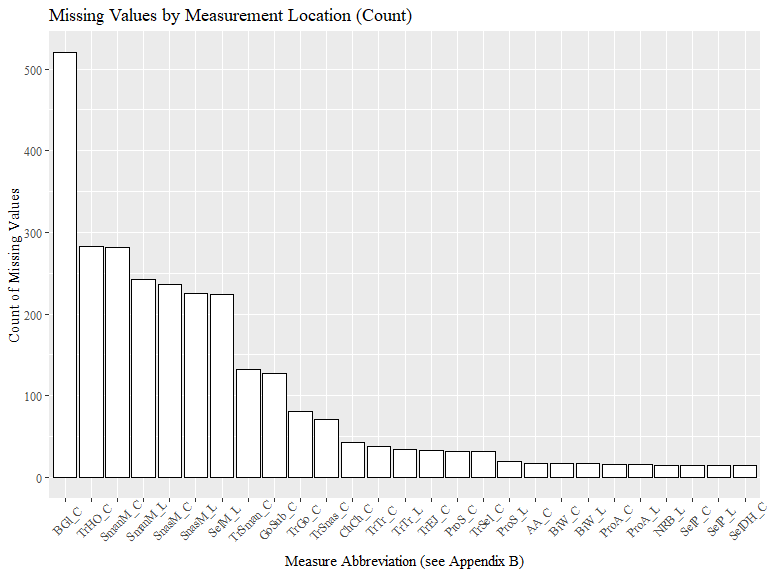
#Size 12 Table TNR  
flextable(measureNAs) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("NA values for each Measurement Location") %>%   
 autofit() %>%   
 set\_header\_labels(values = list(measure\_name = "Measurement Location",  
 measureNAprops = "Proportion of NA values",  
 measureNAsums = "Count of NA values"))

**Table** : NA values for each Measurement Location

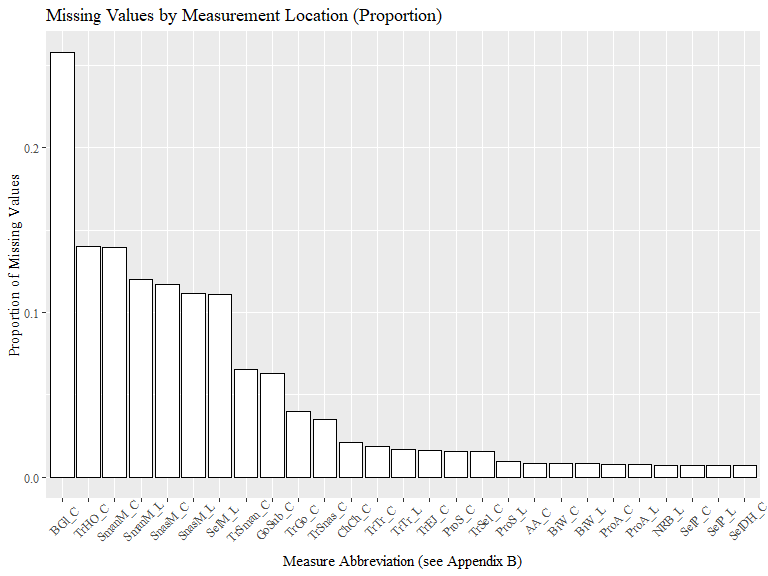
| **Measurement Location** | **Proportion of NA values** | **Count of NA values** |
| --- | --- | --- |
| AA\_C | 0.0084 | 17 |
| BGl\_C | 0.2578 | 520 |
| BiW\_C | 0.0084 | 17 |
| BiW\_L | 0.0084 | 17 |
| ChCh\_C | 0.0213 | 43 |
| GoSub\_C | 0.0630 | 127 |
| NRB\_L | 0.0074 | 15 |
| ProA\_L | 0.0079 | 16 |
| ProA\_C | 0.0079 | 16 |
| ProS\_C | 0.0154 | 31 |
| ProS\_L | 0.0094 | 19 |
| SelP\_C | 0.0074 | 15 |
| SelP\_L | 0.0074 | 15 |
| SelDH\_C | 0.0069 | 14 |
| SelM\_L | 0.1111 | 224 |
| SnasM\_C | 0.1170 | 236 |
| SmanM\_C | 0.1393 | 281 |
| SmanM\_L | 0.1200 | 242 |
| SnasM\_L | 0.1116 | 225 |
| TrHO\_C | 0.1403 | 283 |
| TrEJ\_C | 0.0164 | 33 |
| TrGo\_C | 0.0397 | 80 |
| TrSel\_C | 0.0154 | 31 |
| TrSman\_C | 0.0654 | 132 |
| TrSnas\_C | 0.0352 | 71 |
| TrTr\_C | 0.0188 | 38 |
| TrTr\_L | 0.0169 | 34 |

#boxplot  
#ggplot(data=measureNAs, aes(y=measureNAsums, x=measure\_name))+  
 #geom\_boxplot(color= "black", fill = "white")+  
 #theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 45, vjust=0.7))+  
 #labs(title="Missing Values by Measurement Location",  
 #y="Count of Missing Values",  
 #x="Measure Abbreviation")

#bar chart with counts  
ggplot(data=measureNAs, aes(x=measure\_name, y=measureNAsums))+  
 geom\_bar(stat= "identity", color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 45, vjust=0.7))+  
 labs(title="Missing Values by Measurement Location (Count)",  
 y="Count of Missing Values",  
 x="Measure Abbreviation (see Appendix B)")



#bar chart with proportions  
ggplot(data=measureNAs, aes(x=measure\_name, y=measureNAprops))+  
 geom\_bar(stat= "identity", color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 45, vjust=0.7))+  
 labs(title="Missing Values by Measurement Location (Proportion)",  
 y="Proportion of Missing Values",  
 x="Measure Abbreviation (see Appendix B)")

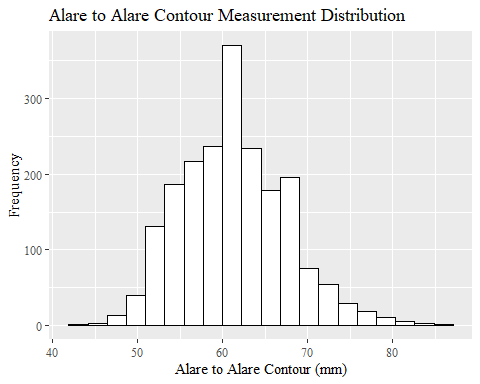


write\_xlsx(measureNAs, "C:\\Users\\19177\\OneDrive - Colostate\\Desktop\\Dissertation\\headscan\_dissertation\\measureNAs.xlsx")

AA\_C race/eth, gender, age group sumstats

#histogram of all AA\_C values  
ggplot(data=headscan\_full, aes(x=AA\_C))+  
 geom\_bar(stat="bin", bins=20, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Alare to Alare Contour Measurement Distribution",  
 y="Frequency",  
 x="Alare to Alare Contour (mm)")

## Warning: Removed 17 rows containing non-finite values (stat\_bin).



#AA\_C race\_eth sumstats  
AA\_Crace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(AA\_C, na.rm = TRUE),  
 max = max(AA\_C, na.rm = TRUE),  
 mean = mean(AA\_C, na.rm = TRUE),  
 mdn = median(AA\_C, na.rm = TRUE),  
 sd = sd(AA\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(AA\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(AA\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(AA\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(AA\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(AA\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(AA\_C)))   
  
AA\_Crace\_sumstats <- AA\_Crace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(AA\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Alare to Alare Contour SumStats by Race/Ethnicity")

**Table** : Alare to Alare Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 44 | 83 | 61.09 | 61 | 6.05 | 0.17 | 52.00 | 57.00 | 61 | 65 | 72.00 | 9 |
| Black | 548 | 47 | 87 | 61.93 | 61 | 6.51 | 0.28 | 52.00 | 57.00 | 61 | 66 | 73.90 | 5 |
| LatinX | 100 | 47 | 75 | 61.54 | 62 | 6.28 | 0.63 | 51.90 | 57.00 | 62 | 66 | 72.10 | 1 |
| Asian | 91 | 50 | 77 | 59.14 | 58 | 6.91 | 0.72 | 50.00 | 53.25 | 58 | 63 | 72.55 | 1 |
| Other | 21 | 45 | 73 | 60.81 | 62 | 6.28 | 1.37 | 50.00 | 60.00 | 62 | 63 | 68.00 | 0 |
| AIAN | 8 | 52 | 69 | 60.86 | 61 | 5.79 | 2.05 | 53.50 | 58.00 | 61 | 64 | 68.40 | 1 |
| PTNS | 5 | 59 | 67 | 61.60 | 61 | 3.13 | 1.40 | 59.20 | 60.00 | 61 | 61 | 65.80 | 0 |
| NHOPI | 4 | 56 | 65 | 59.75 | 59 | 4.11 | 2.06 | 56.15 | 56.75 | 59 | 62 | 64.40 | 0 |

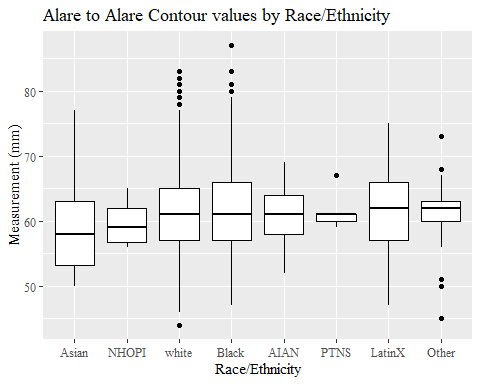
#%>% set\_header\_labels(values = list(AA\_C = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(AA\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Alare to Alare Contour SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Alare to Alare Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 44 | 83 | 61.09 | 61 | 6.05 | 0.17 | 52.00 | 57.00 | 61 | 65 | 72.00 | 9 |
| Black | 548 | 47 | 87 | 61.93 | 61 | 6.51 | 0.28 | 52.00 | 57.00 | 61 | 66 | 73.90 | 5 |
| LatinX | 100 | 47 | 75 | 61.54 | 62 | 6.28 | 0.63 | 51.90 | 57.00 | 62 | 66 | 72.10 | 1 |
| Asian | 91 | 50 | 77 | 59.14 | 58 | 6.91 | 0.72 | 50.00 | 53.25 | 58 | 63 | 72.55 | 1 |
| Other | 21 | 45 | 73 | 60.81 | 62 | 6.28 | 1.37 | 50.00 | 60.00 | 62 | 63 | 68.00 | 0 |
| AIAN | 8 | 52 | 69 | 60.86 | 61 | 5.79 | 2.05 | 53.50 | 58.00 | 61 | 64 | 68.40 | 1 |
| PTNS | 5 | 59 | 67 | 61.60 | 61 | 3.13 | 1.40 | 59.20 | 60.00 | 61 | 61 | 65.80 | 0 |
| NHOPI | 4 | 56 | 65 | 59.75 | 59 | 4.11 | 2.06 | 56.15 | 56.75 | 59 | 62 | 64.40 | 0 |

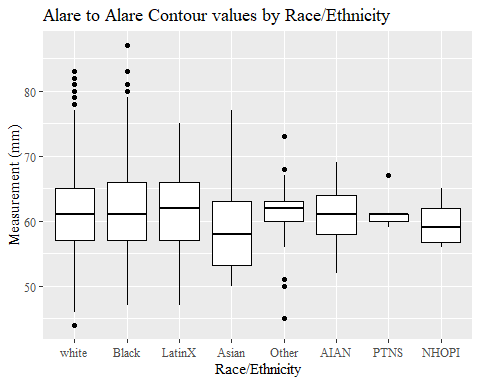
#%>% set\_header\_labels(values = list(AA\_C = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, AA\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=AA\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Alare to Alare Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 17 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=AA\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Alare to Alare Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 17 rows containing non-finite values (stat\_boxplot).



#AA\_C gender sumstats  
AA\_Cgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(AA\_C, na.rm = TRUE),  
 max = max(AA\_C, na.rm = TRUE),  
 mean = mean(AA\_C, na.rm = TRUE),  
 mdn = median(AA\_C, na.rm = TRUE),  
 sd = sd(AA\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(AA\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(AA\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(AA\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(AA\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(AA\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(AA\_C)))  
  
AA\_Cgender\_sumstats <- AA\_Cgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(AA\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Alare to Alare Contour SumStats by Gender")

**Table** : Alare to Alare Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 44 | 77 | 58.31 | 58 | 5.00 | 0.15 | 51.0 | 55.00 | 58 | 62.00 | 67.00 | 12 |
| Male | 939 | 47 | 87 | 64.58 | 64 | 5.85 | 0.19 | 56.0 | 60.00 | 64 | 68.00 | 74.35 | 5 |
| Non-binary or Other | 5 | 52 | 68 | 60.80 | 62 | 5.76 | 2.58 | 53.6 | 60.00 | 62 | 62.00 | 66.80 | 0 |
| Prefer not to say | 1 | 61 | 61 | 61.00 | 61 |  |  | 61.0 | 61.00 | 61 | 61.00 | 61.00 | 0 |
|  | 8 | 53 | 66 | 58.62 | 58 | 4.31 | 1.52 | 53.7 | 55.75 | 58 | 60.75 | 64.95 | 0 |

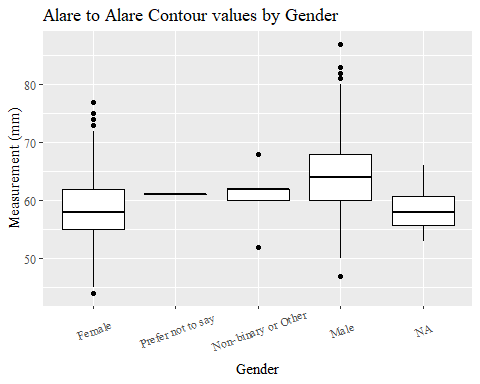
#%>% set\_header\_labels(values = list(AA\_C = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(AA\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Alare to Alare Contour SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Alare to Alare Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 44 | 77 | 58.31 | 58 | 5.00 | 0.15 | 51.0 | 55.00 | 58 | 62.00 | 67.00 | 12 |
| Male | 939 | 47 | 87 | 64.58 | 64 | 5.85 | 0.19 | 56.0 | 60.00 | 64 | 68.00 | 74.35 | 5 |
| Non-binary or Other | 5 | 52 | 68 | 60.80 | 62 | 5.76 | 2.58 | 53.6 | 60.00 | 62 | 62.00 | 66.80 | 0 |
| Prefer not to say | 1 | 61 | 61 | 61.00 | 61 |  |  | 61.0 | 61.00 | 61 | 61.00 | 61.00 | 0 |
|  | 8 | 53 | 66 | 58.62 | 58 | 4.31 | 1.52 | 53.7 | 55.75 | 58 | 60.75 | 64.95 | 0 |

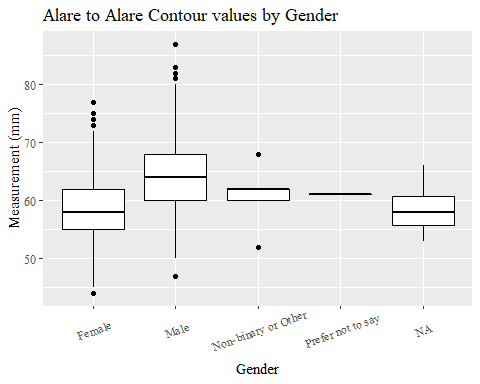
#%>% set\_header\_labels(values = list(AA\_C = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, AA\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=AA\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Alare to Alare Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 17 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=AA\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Alare to Alare Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 17 rows containing non-finite values (stat\_boxplot).



#AA\_C age group sumstats  
AA\_Cage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(AA\_C, na.rm = TRUE),  
 max = max(AA\_C, na.rm = TRUE),  
 mean = mean(AA\_C, na.rm = TRUE),  
 mdn = median(AA\_C, na.rm = TRUE),  
 sd = sd(AA\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(AA\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(AA\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(AA\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(AA\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(AA\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(AA\_C)))  
  
AA\_Cage\_sumstats <- AA\_Cage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(AA\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Alare to Alare Contour SumStats by Age Group")

**Table** : Alare to Alare Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 46 | 83 | 60.33 | 60 | 5.93 | 0.19 | 51.25 | 56 | 60 | 64 | 70 | 6 |
| 37-54 | 940 | 44 | 81 | 61.98 | 62 | 6.35 | 0.21 | 52.00 | 57 | 62 | 66 | 73 | 10 |
| 55-72 | 84 | 50 | 87 | 63.88 | 63 | 6.92 | 0.76 | 54.10 | 59 | 63 | 67 | 74 | 1 |
|  | 1 | 64 | 64 | 64.00 | 64 |  |  | 64.00 | 64 | 64 | 64 | 64 | 0 |

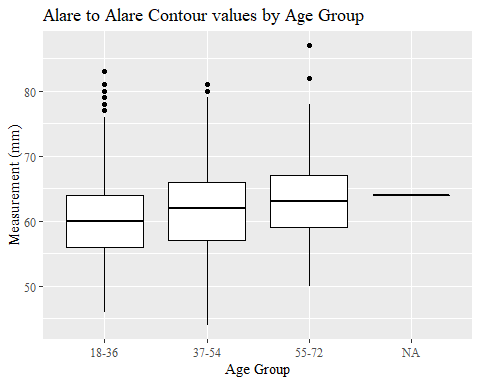
#%>% set\_header\_labels(values = list(AA\_C = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(AA\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Alare to Alare Contour SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Alare to Alare Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 46 | 83 | 60.33 | 60 | 5.93 | 0.19 | 51.25 | 56 | 60 | 64 | 70 | 6 |
| 37-54 | 940 | 44 | 81 | 61.98 | 62 | 6.35 | 0.21 | 52.00 | 57 | 62 | 66 | 73 | 10 |
| 55-72 | 84 | 50 | 87 | 63.88 | 63 | 6.92 | 0.76 | 54.10 | 59 | 63 | 67 | 74 | 1 |
|  | 1 | 64 | 64 | 64.00 | 64 |  |  | 64.00 | 64 | 64 | 64 | 64 | 0 |

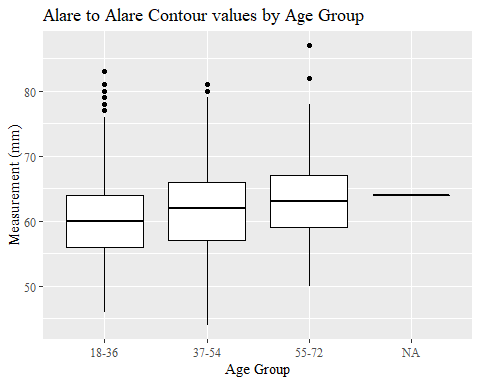
#%>% set\_header\_labels(values = list(AA\_C = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, AA\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=AA\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Alare to Alare Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 17 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=AA\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Alare to Alare Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

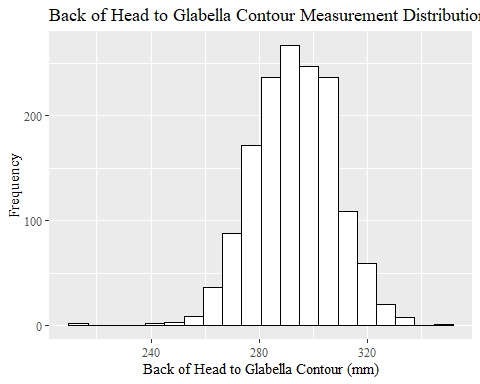
## Warning: Removed 17 rows containing non-finite values (stat\_boxplot).



BGl\_C race/eth, gender, age group sumstats

#histogram of all BGl\_C values  
ggplot(data=headscan\_full, aes(x=BGl\_C))+  
 geom\_bar(stat="bin", bins=20, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Back of Head to Glabella Contour Measurement Distribution",  
 y="Frequency",  
 x="Back of Head to Glabella Contour (mm)")

## Warning: Removed 520 rows containing non-finite values (stat\_bin).



#BGl\_C race/eth sumstats  
BGl\_Crace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(BGl\_C, na.rm = TRUE),  
 max = max(BGl\_C, na.rm = TRUE),  
 mean = mean(BGl\_C, na.rm = TRUE),  
 mdn = median(BGl\_C, na.rm = TRUE),  
 sd = sd(BGl\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(BGl\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(BGl\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(BGl\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(BGl\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(BGl\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(BGl\_C)))   
  
BGl\_Crace\_sumstats <- BGl\_Crace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(BGl\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Back of Head to Glabella Contour SumStats by Race/Ethnicity")

**Table** : Back of Head to Glabella Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 216 | 333 | 291.98 | 292.0 | 14.76 | 0.42 | 269.00 | 282.00 | 292.0 | 303.0 | 316.00 | 271 |
| Black | 548 | 243 | 350 | 297.17 | 296.0 | 14.95 | 0.64 | 274.00 | 285.00 | 296.0 | 308.0 | 323.00 | 197 |
| LatinX | 100 | 262 | 331 | 290.70 | 290.0 | 13.97 | 1.40 | 268.70 | 281.00 | 290.0 | 299.0 | 316.10 | 21 |
| Asian | 91 | 250 | 320 | 285.55 | 285.5 | 14.14 | 1.48 | 264.30 | 276.00 | 285.5 | 296.0 | 306.05 | 17 |
| Other | 21 | 215 | 334 | 288.40 | 289.0 | 26.86 | 5.86 | 252.80 | 279.00 | 289.0 | 301.5 | 324.20 | 6 |
| AIAN | 8 | 284 | 307 | 294.75 | 294.0 | 9.46 | 3.35 | 285.35 | 290.75 | 294.0 | 298.0 | 305.20 | 4 |
| PTNS | 5 | 294 | 294 | 294.00 | 294.0 |  |  | 294.00 | 294.00 | 294.0 | 294.0 | 294.00 | 4 |
| NHOPI | 4 | 296 | 319 | 306.75 | 306.0 | 9.46 | 4.73 | 297.35 | 302.75 | 306.0 | 310.0 | 317.20 | 0 |

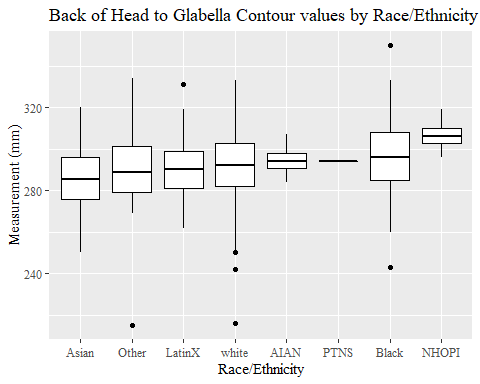
#%>% set\_header\_labels(values = list(BGl\_C = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(BGl\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Back of Head to Glabella Contour SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Back of Head to Glabella Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 216 | 333 | 291.98 | 292.0 | 14.76 | 0.42 | 269.00 | 282.00 | 292.0 | 303.0 | 316.00 | 271 |
| Black | 548 | 243 | 350 | 297.17 | 296.0 | 14.95 | 0.64 | 274.00 | 285.00 | 296.0 | 308.0 | 323.00 | 197 |
| LatinX | 100 | 262 | 331 | 290.70 | 290.0 | 13.97 | 1.40 | 268.70 | 281.00 | 290.0 | 299.0 | 316.10 | 21 |
| Asian | 91 | 250 | 320 | 285.55 | 285.5 | 14.14 | 1.48 | 264.30 | 276.00 | 285.5 | 296.0 | 306.05 | 17 |
| Other | 21 | 215 | 334 | 288.40 | 289.0 | 26.86 | 5.86 | 252.80 | 279.00 | 289.0 | 301.5 | 324.20 | 6 |
| AIAN | 8 | 284 | 307 | 294.75 | 294.0 | 9.46 | 3.35 | 285.35 | 290.75 | 294.0 | 298.0 | 305.20 | 4 |
| PTNS | 5 | 294 | 294 | 294.00 | 294.0 |  |  | 294.00 | 294.00 | 294.0 | 294.0 | 294.00 | 4 |
| NHOPI | 4 | 296 | 319 | 306.75 | 306.0 | 9.46 | 4.73 | 297.35 | 302.75 | 306.0 | 310.0 | 317.20 | 0 |

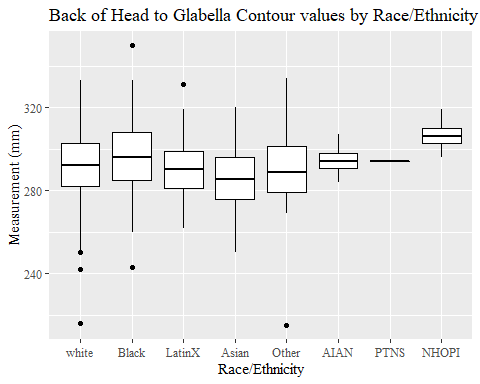
#%>% set\_header\_labels(values = list(BGl\_C = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, BGl\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=BGl\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Back of Head to Glabella Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 520 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=BGl\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Back of Head to Glabella Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 520 rows containing non-finite values (stat\_boxplot).



#BGl\_C gender sumstats  
BGl\_Cgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(BGl\_C, na.rm = TRUE),  
 max = max(BGl\_C, na.rm = TRUE),  
 mean = mean(BGl\_C, na.rm = TRUE),  
 mdn = median(BGl\_C, na.rm = TRUE),  
 sd = sd(BGl\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(BGl\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(BGl\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(BGl\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(BGl\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(BGl\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(BGl\_C)))

## Warning in min(BGl\_C, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf

## Warning in max(BGl\_C, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf

BGl\_Cgender\_sumstats <- BGl\_Cgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(BGl\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Back of Head to Glabella Contour SumStats by Gender")

**Table** : Back of Head to Glabella Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 242 | 332 | 286.39 | 286 | 13.71 | 0.42 | 264.45 | 278.00 | 286 | 295.00 | 310.00 | 454 |
| Male | 939 | 215 | 350 | 297.45 | 298 | 14.36 | 0.47 | 275.00 | 288.00 | 298 | 307.00 | 320.00 | 61 |
| Non-binary or Other | 5 | 268 | 289 | 277.60 | 273 | 9.69 | 4.33 | 268.60 | 271.00 | 273 | 287.00 | 288.60 | 0 |
| Prefer not to say | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |
|  | 8 | 269 | 281 | 275.50 | 276 | 5.92 | 2.09 | 269.45 | 271.25 | 276 | 280.25 | 280.85 | 4 |

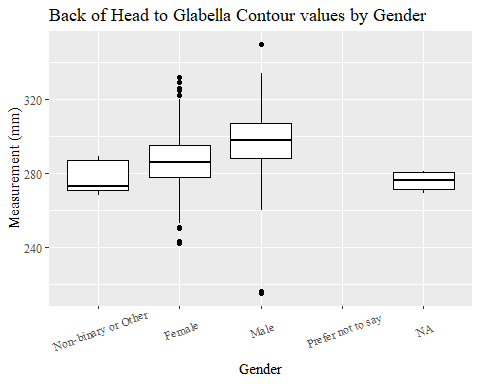
#%>% set\_header\_labels(values = list(BGl\_C = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(BGl\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Back of Head to Glabella Contour SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Back of Head to Glabella Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 242 | 332 | 286.39 | 286 | 13.71 | 0.42 | 264.45 | 278.00 | 286 | 295.00 | 310.00 | 454 |
| Male | 939 | 215 | 350 | 297.45 | 298 | 14.36 | 0.47 | 275.00 | 288.00 | 298 | 307.00 | 320.00 | 61 |
| Non-binary or Other | 5 | 268 | 289 | 277.60 | 273 | 9.69 | 4.33 | 268.60 | 271.00 | 273 | 287.00 | 288.60 | 0 |
| Prefer not to say | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |
|  | 8 | 269 | 281 | 275.50 | 276 | 5.92 | 2.09 | 269.45 | 271.25 | 276 | 280.25 | 280.85 | 4 |

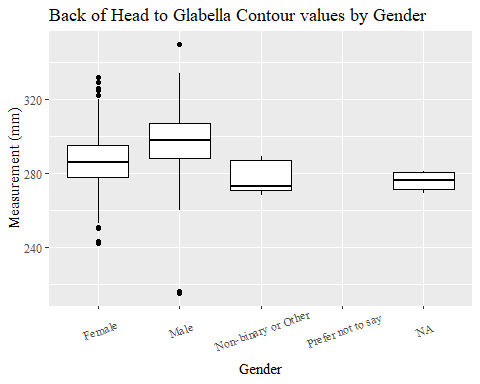
#%>% set\_header\_labels(values = list(BGl\_C = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, BGl\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=BGl\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Back of Head to Glabella Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 520 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=BGl\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Back of Head to Glabella Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 520 rows containing non-finite values (stat\_boxplot).



#BGl\_C age group sumstats  
BGl\_Cage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(BGl\_C, na.rm = TRUE),  
 max = max(BGl\_C, na.rm = TRUE),  
 mean = mean(BGl\_C, na.rm = TRUE),  
 mdn = median(BGl\_C, na.rm = TRUE),  
 sd = sd(BGl\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(BGl\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(BGl\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(BGl\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(BGl\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(BGl\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(BGl\_C)))  
  
BGl\_Cage\_sumstats <- BGl\_Cage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(BGl\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Back of Head to Glabella Contour SumStats by Age Group")

**Table** : Back of Head to Glabella Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 215 | 350 | 293.37 | 294 | 15.30 | 0.49 | 269.35 | 283.00 | 294 | 304 | 318.00 | 264 |
| 37-54 | 940 | 243 | 333 | 291.96 | 292 | 14.96 | 0.49 | 269.00 | 281.75 | 292 | 302 | 317.05 | 240 |
| 55-72 | 84 | 242 | 330 | 295.69 | 295 | 14.57 | 1.59 | 277.00 | 286.50 | 295 | 305 | 320.95 | 16 |
|  | 1 | 299 | 299 | 299.00 | 299 |  |  | 299.00 | 299.00 | 299 | 299 | 299.00 | 0 |

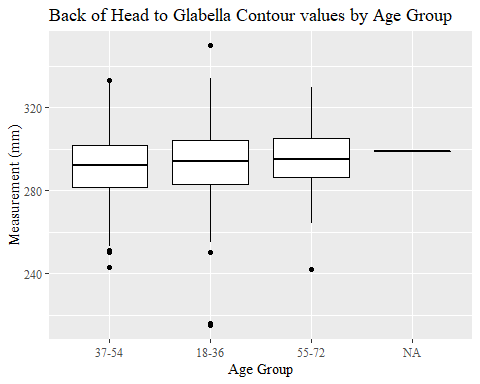
#%>% set\_header\_labels(values = list(BGl\_C = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(BGl\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Back of Head to Glabella Contour SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Back of Head to Glabella Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 215 | 350 | 293.37 | 294 | 15.30 | 0.49 | 269.35 | 283.00 | 294 | 304 | 318.00 | 264 |
| 37-54 | 940 | 243 | 333 | 291.96 | 292 | 14.96 | 0.49 | 269.00 | 281.75 | 292 | 302 | 317.05 | 240 |
| 55-72 | 84 | 242 | 330 | 295.69 | 295 | 14.57 | 1.59 | 277.00 | 286.50 | 295 | 305 | 320.95 | 16 |
|  | 1 | 299 | 299 | 299.00 | 299 |  |  | 299.00 | 299.00 | 299 | 299 | 299.00 | 0 |

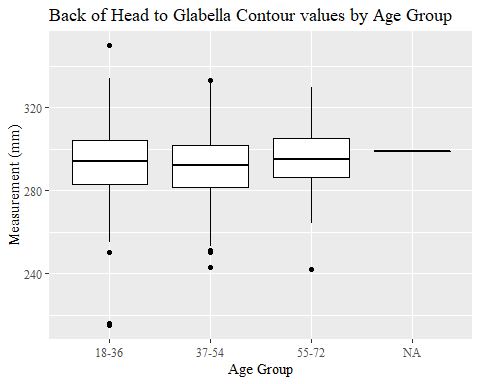
#%>% set\_header\_labels(values = list(BGl\_C = "Alare/AlareCont"))  
  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, BGl\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=BGl\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Back of Head to Glabella Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 520 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=BGl\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Back of Head to Glabella Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

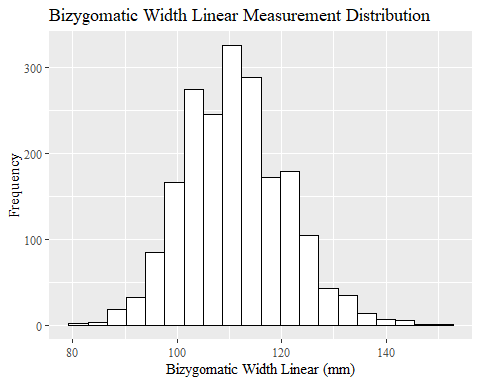
## Warning: Removed 520 rows containing non-finite values (stat\_boxplot).



BiW\_L

#histogram of all BiW\_L values  
ggplot(data=headscan\_full, aes(x=BiW\_L))+  
 geom\_bar(stat="bin", bins=20, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Bizygomatic Width Linear Measurement Distribution",  
 y="Frequency",  
 x="Bizygomatic Width Linear (mm)")

## Warning: Removed 17 rows containing non-finite values (stat\_bin).



#BiW\_L race/eth sumstats  
BiW\_Lrace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(BiW\_L, na.rm = TRUE),  
 max = max(BiW\_L, na.rm = TRUE),  
 mean = mean(BiW\_L, na.rm = TRUE),  
 mdn = median(BiW\_L, na.rm = TRUE),  
 sd = sd(BiW\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(BiW\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(BiW\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(BiW\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(BiW\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(BiW\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(BiW\_L)))   
  
BiW\_Lrace\_sumstats <- BiW\_Lrace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(BiW\_Lrace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Bizygomatic Width Linear SumStats by Race/Ethnicity")

**Table** : Bizygomatic Width Linear SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 82 | 152 | 109.75 | 109 | 9.42 | 0.27 | 95.0 | 103.0 | 109 | 116 | 126.0 | 8 |
| Black | 548 | 91 | 148 | 114.55 | 114 | 9.87 | 0.42 | 100.0 | 108.0 | 114 | 121 | 132.0 | 5 |
| LatinX | 100 | 87 | 138 | 111.60 | 111 | 9.66 | 0.97 | 96.9 | 104.5 | 111 | 119 | 126.3 | 1 |
| Asian | 91 | 92 | 140 | 110.88 | 111 | 9.64 | 1.01 | 94.0 | 105.0 | 111 | 118 | 125.0 | 2 |
| Other | 21 | 97 | 129 | 108.29 | 109 | 9.67 | 2.11 | 97.0 | 98.0 | 109 | 113 | 126.0 | 0 |
| AIAN | 8 | 101 | 117 | 108.00 | 106 | 6.30 | 2.23 | 101.3 | 104.0 | 106 | 112 | 116.7 | 1 |
| PTNS | 5 | 102 | 125 | 113.40 | 113 | 10.11 | 4.52 | 102.6 | 105.0 | 113 | 122 | 124.4 | 0 |
| NHOPI | 4 | 110 | 122 | 116.50 | 117 | 5.00 | 2.50 | 110.9 | 114.5 | 117 | 119 | 121.4 | 0 |

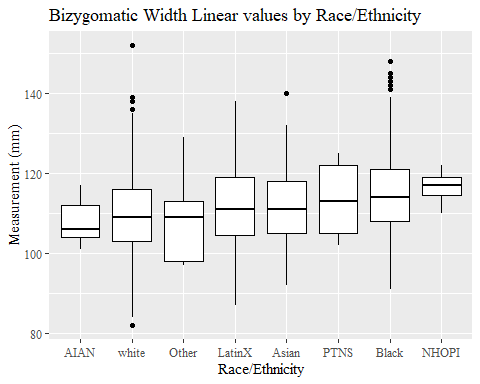
#%>% set\_header\_labels(values = list(BiW\_L = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(BiW\_Lrace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Bizygomatic Width Linear SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Bizygomatic Width Linear SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 82 | 152 | 109.75 | 109 | 9.42 | 0.27 | 95.0 | 103.0 | 109 | 116 | 126.0 | 8 |
| Black | 548 | 91 | 148 | 114.55 | 114 | 9.87 | 0.42 | 100.0 | 108.0 | 114 | 121 | 132.0 | 5 |
| LatinX | 100 | 87 | 138 | 111.60 | 111 | 9.66 | 0.97 | 96.9 | 104.5 | 111 | 119 | 126.3 | 1 |
| Asian | 91 | 92 | 140 | 110.88 | 111 | 9.64 | 1.01 | 94.0 | 105.0 | 111 | 118 | 125.0 | 2 |
| Other | 21 | 97 | 129 | 108.29 | 109 | 9.67 | 2.11 | 97.0 | 98.0 | 109 | 113 | 126.0 | 0 |
| AIAN | 8 | 101 | 117 | 108.00 | 106 | 6.30 | 2.23 | 101.3 | 104.0 | 106 | 112 | 116.7 | 1 |
| PTNS | 5 | 102 | 125 | 113.40 | 113 | 10.11 | 4.52 | 102.6 | 105.0 | 113 | 122 | 124.4 | 0 |
| NHOPI | 4 | 110 | 122 | 116.50 | 117 | 5.00 | 2.50 | 110.9 | 114.5 | 117 | 119 | 121.4 | 0 |

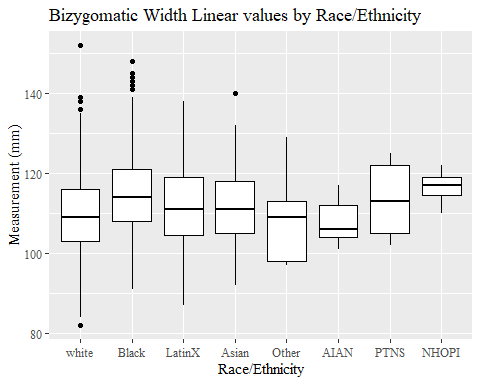
#%>% set\_header\_labels(values = list(BiW\_L = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, BiW\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=BiW\_L, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Bizygomatic Width Linear values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 17 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=BiW\_L, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Bizygomatic Width Linear values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 17 rows containing non-finite values (stat\_boxplot).



#BiW\_L gender sumstats  
BiW\_Lgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(BiW\_L, na.rm = TRUE),  
 max = max(BiW\_L, na.rm = TRUE),  
 mean = mean(BiW\_L, na.rm = TRUE),  
 mdn = median(BiW\_L, na.rm = TRUE),  
 sd = sd(BiW\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(BiW\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(BiW\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(BiW\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(BiW\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(BiW\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(BiW\_L)))  
  
BiW\_Lgender\_sumstats <- BiW\_Lgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(BiW\_Lgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Bizygomatic Width Linear SumStats by Gender")

**Table** : Bizygomatic Width Linear SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 82 | 142 | 109.08 | 108 | 9.46 | 0.29 | 94.00 | 102 | 108 | 116.0 | 125.00 | 12 |
| Male | 939 | 86 | 152 | 113.66 | 113 | 9.54 | 0.31 | 99.00 | 107 | 113 | 119.0 | 130.35 | 5 |
| Non-binary or Other | 5 | 91 | 107 | 98.60 | 99 | 6.07 | 2.71 | 91.80 | 95 | 99 | 101.0 | 105.80 | 0 |
| Prefer not to say | 1 | 113 | 113 | 113.00 | 113 |  |  | 113.00 | 113 | 113 | 113.0 | 113.00 | 0 |
|  | 8 | 96 | 122 | 110.25 | 113 | 8.78 | 3.10 | 97.75 | 104 | 113 | 115.5 | 120.25 | 0 |

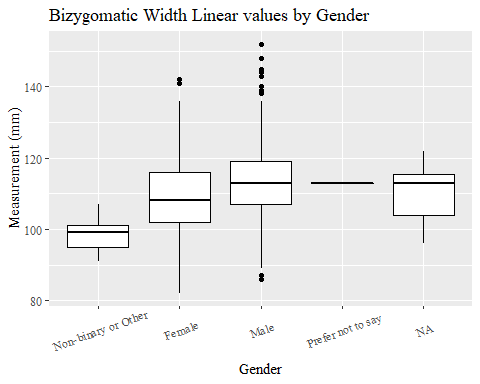
#%>% set\_header\_labels(values = list(BiW\_L = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(BiW\_Lgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Bizygomatic Width Linear SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Bizygomatic Width Linear SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 82 | 142 | 109.08 | 108 | 9.46 | 0.29 | 94.00 | 102 | 108 | 116.0 | 125.00 | 12 |
| Male | 939 | 86 | 152 | 113.66 | 113 | 9.54 | 0.31 | 99.00 | 107 | 113 | 119.0 | 130.35 | 5 |
| Non-binary or Other | 5 | 91 | 107 | 98.60 | 99 | 6.07 | 2.71 | 91.80 | 95 | 99 | 101.0 | 105.80 | 0 |
| Prefer not to say | 1 | 113 | 113 | 113.00 | 113 |  |  | 113.00 | 113 | 113 | 113.0 | 113.00 | 0 |
|  | 8 | 96 | 122 | 110.25 | 113 | 8.78 | 3.10 | 97.75 | 104 | 113 | 115.5 | 120.25 | 0 |

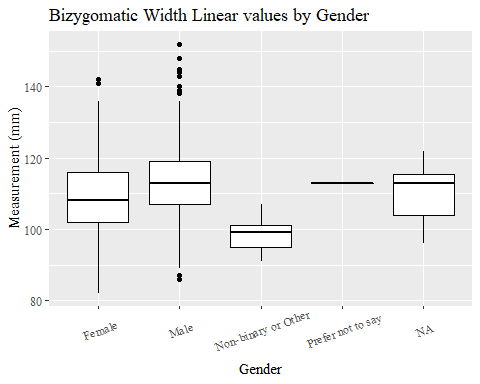
#%>% set\_header\_labels(values = list(BiW\_L = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, BiW\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=BiW\_L, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Bizygomatic Width Linear values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 17 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=BiW\_L, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Bizygomatic Width Linear values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 17 rows containing non-finite values (stat\_boxplot).



#BiW\_L age group sumstats  
BiW\_Lage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(BiW\_L, na.rm = TRUE),  
 max = max(BiW\_L, na.rm = TRUE),  
 mean = mean(BiW\_L, na.rm = TRUE),  
 mdn = median(BiW\_L, na.rm = TRUE),  
 sd = sd(BiW\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(BiW\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(BiW\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(BiW\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(BiW\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(BiW\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(BiW\_L)))  
  
BiW\_Lage\_sumstats <- BiW\_Lage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(BiW\_Lage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Bizygomatic Width Linear SumStats by Age Group")

**Table** : Bizygomatic Width Linear SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 84 | 145 | 111.56 | 111 | 9.83 | 0.31 | 97.00 | 104 | 111 | 118.0 | 129.0 | 6 |
| 37-54 | 940 | 82 | 152 | 110.98 | 111 | 9.74 | 0.32 | 95.00 | 105 | 111 | 118.0 | 126.5 | 9 |
| 55-72 | 84 | 87 | 134 | 109.51 | 109 | 9.30 | 1.02 | 95.05 | 103 | 109 | 115.5 | 126.0 | 2 |
|  | 1 | 103 | 103 | 103.00 | 103 |  |  | 103.00 | 103 | 103 | 103.0 | 103.0 | 0 |

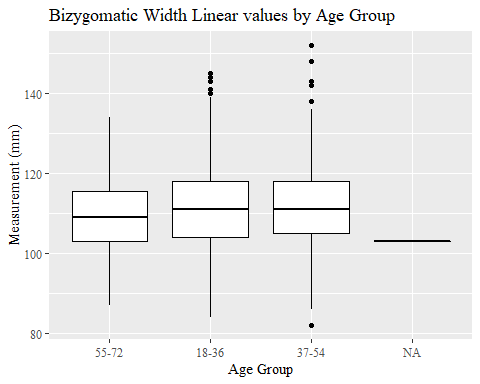
#%>% set\_header\_labels(values = list(BiW\_L = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(BiW\_Lage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Bizygomatic Width Linear SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Bizygomatic Width Linear SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 84 | 145 | 111.56 | 111 | 9.83 | 0.31 | 97.00 | 104 | 111 | 118.0 | 129.0 | 6 |
| 37-54 | 940 | 82 | 152 | 110.98 | 111 | 9.74 | 0.32 | 95.00 | 105 | 111 | 118.0 | 126.5 | 9 |
| 55-72 | 84 | 87 | 134 | 109.51 | 109 | 9.30 | 1.02 | 95.05 | 103 | 109 | 115.5 | 126.0 | 2 |
|  | 1 | 103 | 103 | 103.00 | 103 |  |  | 103.00 | 103 | 103 | 103.0 | 103.0 | 0 |

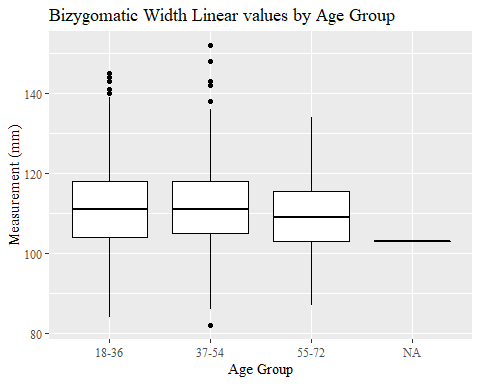
#%>% set\_header\_labels(values = list(BiW\_L = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, BiW\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=BiW\_L, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Bizygomatic Width Linear values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 17 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=BiW\_L, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Bizygomatic Width Linear values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

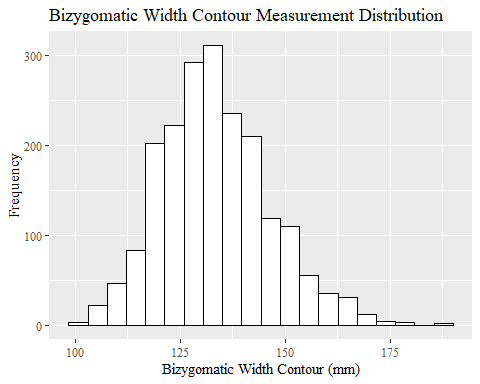
## Warning: Removed 17 rows containing non-finite values (stat\_boxplot).



BiW\_C

#histogram of all BiW\_C values  
ggplot(data=headscan\_full, aes(x=BiW\_C))+  
 geom\_bar(stat="bin", bins=20, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Bizygomatic Width Contour Measurement Distribution",  
 y="Frequency",  
 x="Bizygomatic Width Contour (mm)")

## Warning: Removed 17 rows containing non-finite values (stat\_bin).



#BiW\_C race/eth sumstats  
BiW\_Crace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(BiW\_C, na.rm = TRUE),  
 max = max(BiW\_C, na.rm = TRUE),  
 mean = mean(BiW\_C, na.rm = TRUE),  
 mdn = median(BiW\_C, na.rm = TRUE),  
 sd = sd(BiW\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(BiW\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(BiW\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(BiW\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(BiW\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(BiW\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(BiW\_C)))   
  
BiW\_Crace\_sumstats <- BiW\_Crace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(BiW\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Bizygomatic Width Contour SumStats by Race/Ethnicity")

**Table** : Bizygomatic Width Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 101 | 188 | 133.19 | 133.0 | 12.78 | 0.36 | 114.00 | 124.00 | 133.0 | 141.00 | 155.00 | 8 |
| Black | 548 | 106 | 187 | 134.80 | 134.0 | 13.32 | 0.57 | 116.00 | 125.00 | 134.0 | 143.00 | 160.00 | 5 |
| LatinX | 100 | 106 | 171 | 133.88 | 131.0 | 13.00 | 1.30 | 119.00 | 124.00 | 131.0 | 142.00 | 156.10 | 1 |
| Asian | 91 | 105 | 156 | 128.69 | 130.0 | 11.41 | 1.20 | 111.00 | 120.00 | 130.0 | 135.00 | 148.00 | 2 |
| Other | 21 | 114 | 156 | 130.76 | 130.0 | 12.12 | 2.64 | 116.00 | 120.00 | 130.0 | 138.00 | 150.00 | 0 |
| AIAN | 8 | 115 | 145 | 128.43 | 124.0 | 9.74 | 3.45 | 117.70 | 124.00 | 124.0 | 133.50 | 142.00 | 1 |
| PTNS | 5 | 125 | 145 | 135.40 | 138.0 | 8.20 | 3.67 | 125.80 | 129.00 | 138.0 | 140.00 | 144.00 | 0 |
| NHOPI | 4 | 120 | 155 | 141.00 | 144.5 | 16.06 | 8.03 | 122.55 | 132.75 | 144.5 | 152.75 | 154.55 | 0 |

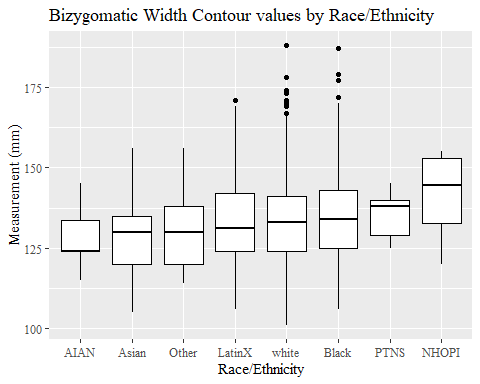
#%>% set\_header\_labels(values = list(BiW\_C = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(BiW\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Bizygomatic Width Contour SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Bizygomatic Width Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 101 | 188 | 133.19 | 133.0 | 12.78 | 0.36 | 114.00 | 124.00 | 133.0 | 141.00 | 155.00 | 8 |
| Black | 548 | 106 | 187 | 134.80 | 134.0 | 13.32 | 0.57 | 116.00 | 125.00 | 134.0 | 143.00 | 160.00 | 5 |
| LatinX | 100 | 106 | 171 | 133.88 | 131.0 | 13.00 | 1.30 | 119.00 | 124.00 | 131.0 | 142.00 | 156.10 | 1 |
| Asian | 91 | 105 | 156 | 128.69 | 130.0 | 11.41 | 1.20 | 111.00 | 120.00 | 130.0 | 135.00 | 148.00 | 2 |
| Other | 21 | 114 | 156 | 130.76 | 130.0 | 12.12 | 2.64 | 116.00 | 120.00 | 130.0 | 138.00 | 150.00 | 0 |
| AIAN | 8 | 115 | 145 | 128.43 | 124.0 | 9.74 | 3.45 | 117.70 | 124.00 | 124.0 | 133.50 | 142.00 | 1 |
| PTNS | 5 | 125 | 145 | 135.40 | 138.0 | 8.20 | 3.67 | 125.80 | 129.00 | 138.0 | 140.00 | 144.00 | 0 |
| NHOPI | 4 | 120 | 155 | 141.00 | 144.5 | 16.06 | 8.03 | 122.55 | 132.75 | 144.5 | 152.75 | 154.55 | 0 |

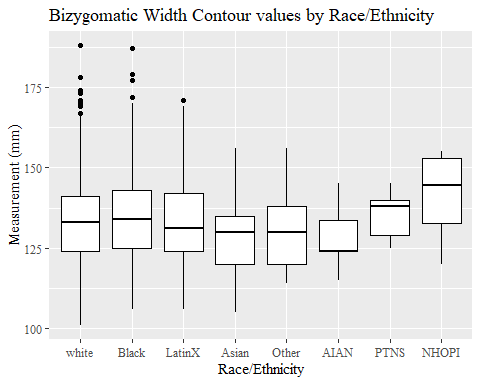
#%>% set\_header\_labels(values = list(BiW\_C = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, BiW\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=BiW\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Bizygomatic Width Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 17 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=BiW\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Bizygomatic Width Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 17 rows containing non-finite values (stat\_boxplot).



#BiW\_C gender sumstats  
BiW\_Cgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(BiW\_C, na.rm = TRUE),  
 max = max(BiW\_C, na.rm = TRUE),  
 mean = mean(BiW\_C, na.rm = TRUE),  
 mdn = median(BiW\_C, na.rm = TRUE),  
 sd = sd(BiW\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(BiW\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(BiW\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(BiW\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(BiW\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(BiW\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(BiW\_C)))  
  
BiW\_Cgender\_sumstats <- BiW\_Cgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(BiW\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Bizygomatic Width Contour SumStats by Gender")

**Table** : Bizygomatic Width Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 101 | 165 | 129.00 | 127 | 11.86 | 0.36 | 111.0 | 121.00 | 127 | 137 | 150.00 | 12 |
| Male | 939 | 105 | 188 | 138.55 | 137 | 12.15 | 0.40 | 121.0 | 131.00 | 137 | 145 | 162.00 | 5 |
| Non-binary or Other | 5 | 116 | 125 | 117.80 | 116 | 4.02 | 1.80 | 116.0 | 116.00 | 116 | 116 | 123.20 | 0 |
| Prefer not to say | 1 | 140 | 140 | 140.00 | 140 |  |  | 140.0 | 140.00 | 140 | 140 | 140.00 | 0 |
|  | 8 | 110 | 151 | 130.25 | 130 | 14.78 | 5.23 | 112.8 | 118.75 | 130 | 138 | 150.65 | 0 |

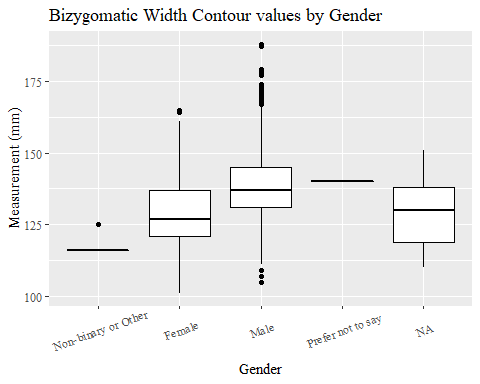
#%>% set\_header\_labels(values = list(BiW\_C = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(BiW\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Bizygomatic Width Contour SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Bizygomatic Width Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 101 | 165 | 129.00 | 127 | 11.86 | 0.36 | 111.0 | 121.00 | 127 | 137 | 150.00 | 12 |
| Male | 939 | 105 | 188 | 138.55 | 137 | 12.15 | 0.40 | 121.0 | 131.00 | 137 | 145 | 162.00 | 5 |
| Non-binary or Other | 5 | 116 | 125 | 117.80 | 116 | 4.02 | 1.80 | 116.0 | 116.00 | 116 | 116 | 123.20 | 0 |
| Prefer not to say | 1 | 140 | 140 | 140.00 | 140 |  |  | 140.0 | 140.00 | 140 | 140 | 140.00 | 0 |
|  | 8 | 110 | 151 | 130.25 | 130 | 14.78 | 5.23 | 112.8 | 118.75 | 130 | 138 | 150.65 | 0 |

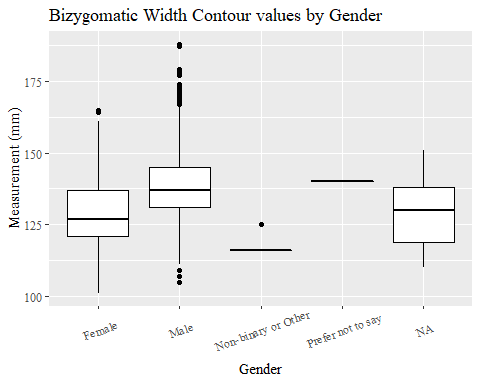
#%>% set\_header\_labels(values = list(BiW\_C = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, BiW\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=BiW\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Bizygomatic Width Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 17 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=BiW\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Bizygomatic Width Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 17 rows containing non-finite values (stat\_boxplot).



#BiW\_C age group sumstats  
BiW\_Cage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(BiW\_C, na.rm = TRUE),  
 max = max(BiW\_C, na.rm = TRUE),  
 mean = mean(BiW\_C, na.rm = TRUE),  
 mdn = median(BiW\_C, na.rm = TRUE),  
 sd = sd(BiW\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(BiW\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(BiW\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(BiW\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(BiW\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(BiW\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(BiW\_C)))  
  
BiW\_Cage\_sumstats <- BiW\_Cage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(BiW\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Bizygomatic Width Contour SumStats by Age Group")

**Table** : Bizygomatic Width Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 102 | 179 | 133.73 | 133 | 12.77 | 0.41 | 115.00 | 125 | 133 | 141 | 157.00 | 6 |
| 37-54 | 940 | 101 | 188 | 133.22 | 132 | 13.06 | 0.43 | 113.00 | 124 | 132 | 141 | 156.00 | 9 |
| 55-72 | 84 | 107 | 164 | 132.61 | 131 | 13.12 | 1.43 | 114.05 | 123 | 131 | 139 | 158.85 | 2 |
|  | 1 | 124 | 124 | 124.00 | 124 |  |  | 124.00 | 124 | 124 | 124 | 124.00 | 0 |

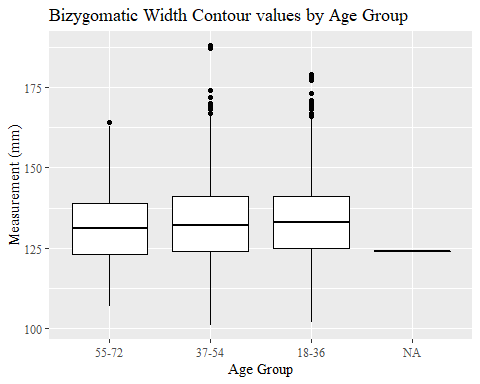
#%>% set\_header\_labels(values = list(BiW\_C = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(BiW\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Bizygomatic Width Contour SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Bizygomatic Width Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 102 | 179 | 133.73 | 133 | 12.77 | 0.41 | 115.00 | 125 | 133 | 141 | 157.00 | 6 |
| 37-54 | 940 | 101 | 188 | 133.22 | 132 | 13.06 | 0.43 | 113.00 | 124 | 132 | 141 | 156.00 | 9 |
| 55-72 | 84 | 107 | 164 | 132.61 | 131 | 13.12 | 1.43 | 114.05 | 123 | 131 | 139 | 158.85 | 2 |
|  | 1 | 124 | 124 | 124.00 | 124 |  |  | 124.00 | 124 | 124 | 124 | 124.00 | 0 |

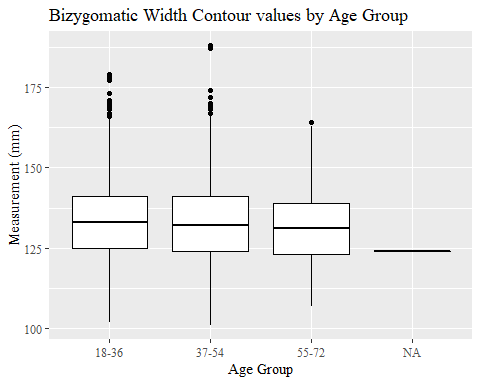
#%>% set\_header\_labels(values = list(BiW\_C = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, BiW\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=BiW\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Bizygomatic Width Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 17 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=BiW\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Bizygomatic Width Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

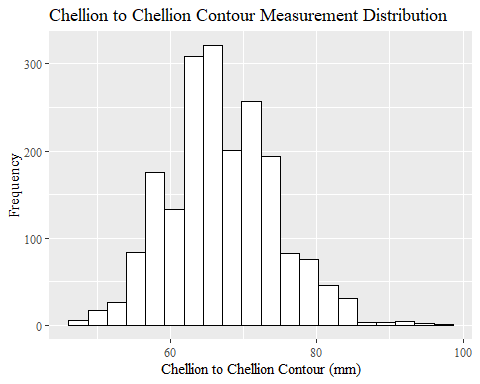
## Warning: Removed 17 rows containing non-finite values (stat\_boxplot).



ChCh\_C

#histogram of all ChCh\_C values  
ggplot(data=headscan\_full, aes(x=ChCh\_C))+  
 geom\_bar(stat="bin", bins=20, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Chellion to Chellion Contour Measurement Distribution",  
 y="Frequency",  
 x="Chellion to Chellion Contour (mm)")

## Warning: Removed 43 rows containing non-finite values (stat\_bin).



#ChCh\_C race/eth sumstats  
ChCh\_Crace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(ChCh\_C, na.rm = TRUE),  
 max = max(ChCh\_C, na.rm = TRUE),  
 mean = mean(ChCh\_C, na.rm = TRUE),  
 mdn = median(ChCh\_C, na.rm = TRUE),  
 sd = sd(ChCh\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(ChCh\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(ChCh\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(ChCh\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(ChCh\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(ChCh\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(ChCh\_C)))   
  
ChCh\_Crace\_sumstats <- ChCh\_Crace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(ChCh\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Chellion Chellion Contour SumStats by Race/Ethnicity")

**Table** : Chellion Chellion Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 47 | 95 | 65.30 | 65.0 | 6.89 | 0.20 | 54.00 | 61.00 | 65.0 | 70.00 | 77.00 | 33 |
| Black | 548 | 51 | 97 | 71.23 | 71.0 | 7.14 | 0.31 | 60.00 | 66.00 | 71.0 | 76.00 | 83.00 | 6 |
| LatinX | 100 | 48 | 84 | 67.14 | 67.0 | 6.50 | 0.65 | 58.00 | 62.00 | 67.0 | 72.00 | 78.00 | 2 |
| Asian | 91 | 48 | 84 | 64.84 | 65.0 | 7.09 | 0.74 | 53.45 | 60.00 | 65.0 | 69.00 | 75.55 | 1 |
| Other | 21 | 56 | 84 | 67.33 | 67.0 | 6.51 | 1.42 | 59.00 | 64.00 | 67.0 | 70.00 | 78.00 | 0 |
| AIAN | 8 | 61 | 81 | 67.57 | 65.0 | 6.97 | 2.47 | 61.00 | 63.00 | 65.0 | 70.00 | 77.70 | 1 |
| PTNS | 5 | 58 | 84 | 68.60 | 67.0 | 9.86 | 4.41 | 59.00 | 63.00 | 67.0 | 71.00 | 81.40 | 0 |
| NHOPI | 4 | 61 | 76 | 68.50 | 68.5 | 8.10 | 4.05 | 61.15 | 61.75 | 68.5 | 75.25 | 75.85 | 0 |

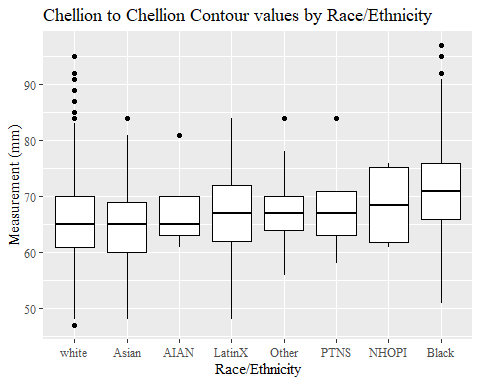
#%>% set\_header\_labels(values = list(ChCh\_C = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(ChCh\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Chellion Chellion Contour SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Chellion Chellion Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 47 | 95 | 65.30 | 65.0 | 6.89 | 0.20 | 54.00 | 61.00 | 65.0 | 70.00 | 77.00 | 33 |
| Black | 548 | 51 | 97 | 71.23 | 71.0 | 7.14 | 0.31 | 60.00 | 66.00 | 71.0 | 76.00 | 83.00 | 6 |
| LatinX | 100 | 48 | 84 | 67.14 | 67.0 | 6.50 | 0.65 | 58.00 | 62.00 | 67.0 | 72.00 | 78.00 | 2 |
| Asian | 91 | 48 | 84 | 64.84 | 65.0 | 7.09 | 0.74 | 53.45 | 60.00 | 65.0 | 69.00 | 75.55 | 1 |
| Other | 21 | 56 | 84 | 67.33 | 67.0 | 6.51 | 1.42 | 59.00 | 64.00 | 67.0 | 70.00 | 78.00 | 0 |
| AIAN | 8 | 61 | 81 | 67.57 | 65.0 | 6.97 | 2.47 | 61.00 | 63.00 | 65.0 | 70.00 | 77.70 | 1 |
| PTNS | 5 | 58 | 84 | 68.60 | 67.0 | 9.86 | 4.41 | 59.00 | 63.00 | 67.0 | 71.00 | 81.40 | 0 |
| NHOPI | 4 | 61 | 76 | 68.50 | 68.5 | 8.10 | 4.05 | 61.15 | 61.75 | 68.5 | 75.25 | 75.85 | 0 |

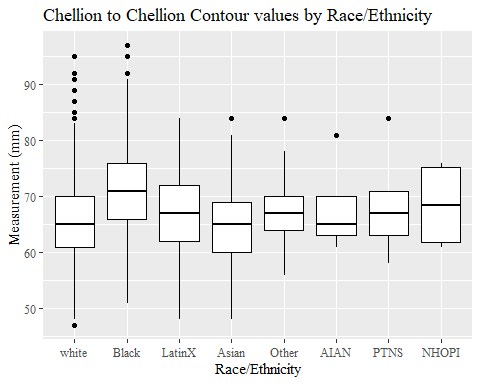
#%>% set\_header\_labels(values = list(ChCh\_C = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, ChCh\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=ChCh\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Chellion to Chellion Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 43 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=ChCh\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Chellion to Chellion Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 43 rows containing non-finite values (stat\_boxplot).



#ChCh\_C gender sumstats  
ChCh\_Cgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(ChCh\_C, na.rm = TRUE),  
 max = max(ChCh\_C, na.rm = TRUE),  
 mean = mean(ChCh\_C, na.rm = TRUE),  
 mdn = median(ChCh\_C, na.rm = TRUE),  
 sd = sd(ChCh\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(ChCh\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(ChCh\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(ChCh\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(ChCh\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(ChCh\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(ChCh\_C)))  
  
ChCh\_Cgender\_sumstats <- ChCh\_Cgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(ChCh\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Chellion to Chellion Contour SumStats by Gender")

**Table** : Chellion to Chellion Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 47 | 97 | 65.21 | 65 | 7.24 | 0.22 | 54.0 | 60.0 | 65 | 70.00 | 78.0 | 19 |
| Male | 939 | 48 | 95 | 69.13 | 69 | 7.09 | 0.23 | 58.0 | 64.0 | 69 | 74.00 | 81.0 | 24 |
| Non-binary or Other | 5 | 54 | 74 | 67.40 | 72 | 8.47 | 3.79 | 56.0 | 64.0 | 72 | 73.00 | 73.8 | 0 |
| Prefer not to say | 1 | 67 | 67 | 67.00 | 67 |  |  | 67.0 | 67.0 | 67 | 67.00 | 67.0 | 0 |
|  | 8 | 58 | 80 | 67.12 | 68 | 7.14 | 2.52 | 58.7 | 61.5 | 68 | 69.75 | 77.2 | 0 |

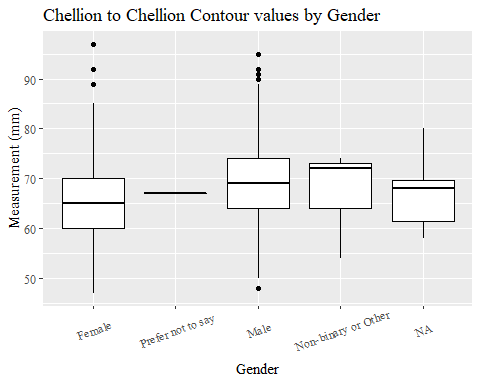
#%>% set\_header\_labels(values = list(ChCh\_C = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(ChCh\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Chellion to Chellion Contour SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Chellion to Chellion Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 47 | 97 | 65.21 | 65 | 7.24 | 0.22 | 54.0 | 60.0 | 65 | 70.00 | 78.0 | 19 |
| Male | 939 | 48 | 95 | 69.13 | 69 | 7.09 | 0.23 | 58.0 | 64.0 | 69 | 74.00 | 81.0 | 24 |
| Non-binary or Other | 5 | 54 | 74 | 67.40 | 72 | 8.47 | 3.79 | 56.0 | 64.0 | 72 | 73.00 | 73.8 | 0 |
| Prefer not to say | 1 | 67 | 67 | 67.00 | 67 |  |  | 67.0 | 67.0 | 67 | 67.00 | 67.0 | 0 |
|  | 8 | 58 | 80 | 67.12 | 68 | 7.14 | 2.52 | 58.7 | 61.5 | 68 | 69.75 | 77.2 | 0 |

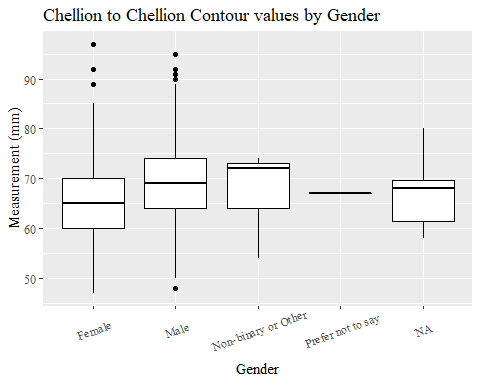
#%>% set\_header\_labels(values = list(ChCh\_C = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, ChCh\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=ChCh\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Chellion to Chellion Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 43 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=ChCh\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Chellion to Chellion Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 43 rows containing non-finite values (stat\_boxplot).



#ChCh\_C age group sumstats  
ChCh\_Cage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(ChCh\_C, na.rm = TRUE),  
 max = max(ChCh\_C, na.rm = TRUE),  
 mean = mean(ChCh\_C, na.rm = TRUE),  
 mdn = median(ChCh\_C, na.rm = TRUE),  
 sd = sd(ChCh\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(ChCh\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(ChCh\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(ChCh\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(ChCh\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(ChCh\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(ChCh\_C)))  
  
ChCh\_Cage\_sumstats <- ChCh\_Cage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(ChCh\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Chellion to Chellion Contour SumStats by Age Group")

**Table** : Chellion to Chellion Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 47 | 85 | 65.82 | 66.0 | 6.88 | 0.22 | 55.00 | 61 | 66.0 | 70 | 78.00 | 17 |
| 37-54 | 940 | 47 | 97 | 68.08 | 67.5 | 7.67 | 0.25 | 56.00 | 63 | 67.5 | 73 | 81.00 | 24 |
| 55-72 | 84 | 54 | 91 | 69.99 | 70.0 | 8.34 | 0.91 | 56.05 | 64 | 70.0 | 76 | 82.95 | 2 |
|  | 1 | 75 | 75 | 75.00 | 75.0 |  |  | 75.00 | 75 | 75.0 | 75 | 75.00 | 0 |

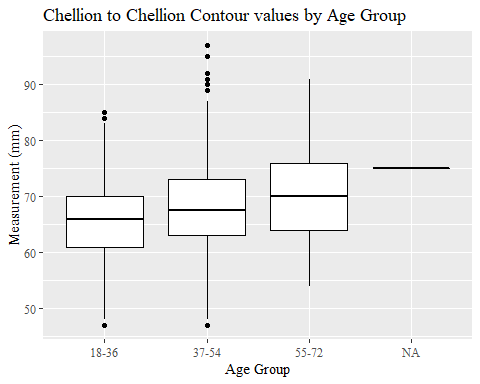
#%>% set\_header\_labels(values = list(ChCh\_C = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(ChCh\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Chellion to Chellion Contour SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Chellion to Chellion Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 47 | 85 | 65.82 | 66.0 | 6.88 | 0.22 | 55.00 | 61 | 66.0 | 70 | 78.00 | 17 |
| 37-54 | 940 | 47 | 97 | 68.08 | 67.5 | 7.67 | 0.25 | 56.00 | 63 | 67.5 | 73 | 81.00 | 24 |
| 55-72 | 84 | 54 | 91 | 69.99 | 70.0 | 8.34 | 0.91 | 56.05 | 64 | 70.0 | 76 | 82.95 | 2 |
|  | 1 | 75 | 75 | 75.00 | 75.0 |  |  | 75.00 | 75 | 75.0 | 75 | 75.00 | 0 |

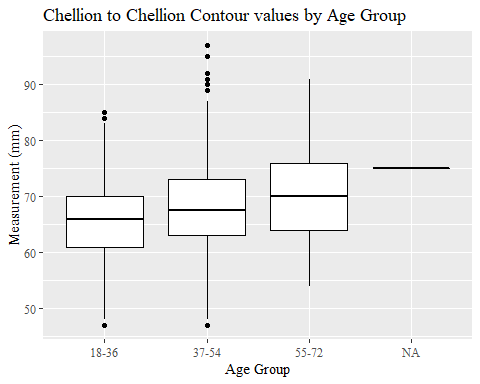
#%>% set\_header\_labels(values = list(ChCh\_C = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, ChCh\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=ChCh\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Chellion to Chellion Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 43 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=ChCh\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Chellion to Chellion Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

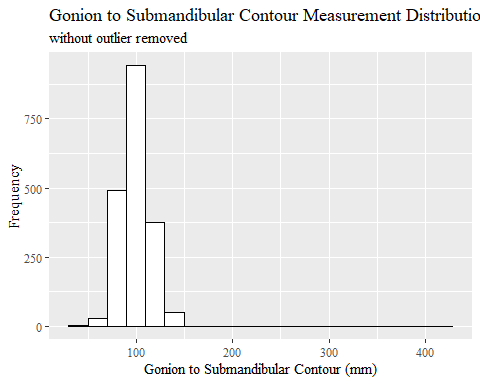
## Warning: Removed 43 rows containing non-finite values (stat\_boxplot).



GoSub\_C WITHOUT Outlier Removed!!!!

#histogram of all GoSub\_C values  
ggplot(data=headscan\_full, aes(x=GoSub\_C))+  
 geom\_bar(stat="bin", bins=20, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Gonion to Submandibular Contour Measurement Distribution",  
 subtitle= "without outlier removed",  
 y="Frequency",  
 x="Gonion to Submandibular Contour (mm)")

## Warning: Removed 127 rows containing non-finite values (stat\_bin).



#GoSub\_C race/eth sumstats  
GoSub\_Crace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(GoSub\_C, na.rm = TRUE),  
 max = max(GoSub\_C, na.rm = TRUE),  
 mean = mean(GoSub\_C, na.rm = TRUE),  
 mdn = median(GoSub\_C, na.rm = TRUE),  
 sd = sd(GoSub\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(GoSub\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(GoSub\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(GoSub\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(GoSub\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(GoSub\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(GoSub\_C)))   
  
GoSub\_Crace\_sumstats <- GoSub\_Crace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(GoSub\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Gonion to Submandibular Contour SumStats by Race/Ethnicity")

**Table** : Gonion to Submandibular Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 45 | 424 | 98.12 | 97 | 18.53 | 0.53 | 74.0 | 87 | 97 | 107.0 | 124.6 | 71 |
| Black | 548 | 63 | 146 | 102.33 | 102 | 14.29 | 0.61 | 78.0 | 93 | 102 | 112.0 | 125.6 | 39 |
| LatinX | 100 | 67 | 132 | 97.47 | 98 | 13.65 | 1.37 | 76.0 | 88 | 98 | 104.0 | 119.0 | 9 |
| Asian | 91 | 71 | 137 | 95.61 | 94 | 14.40 | 1.51 | 74.8 | 85 | 94 | 106.0 | 120.2 | 2 |
| Other | 21 | 73 | 126 | 100.00 | 96 | 14.35 | 3.13 | 78.6 | 90 | 96 | 109.0 | 121.2 | 4 |
| AIAN | 8 | 71 | 111 | 89.86 | 94 | 14.02 | 4.96 | 73.1 | 79 | 94 | 97.5 | 107.4 | 1 |
| PTNS | 5 | 69 | 134 | 93.20 | 91 | 24.93 | 11.15 | 70.8 | 78 | 91 | 94.0 | 126.0 | 0 |
| NHOPI | 4 | 75 | 96 | 86.67 | 89 | 10.69 | 5.35 | 76.4 | 82 | 89 | 92.5 | 95.3 | 1 |

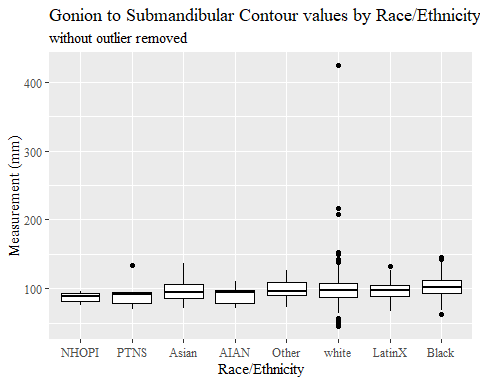
#%>% set\_header\_labels(values = list(GoSub\_C = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(GoSub\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Gonion to Submandibular Contour SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Gonion to Submandibular Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 45 | 424 | 98.12 | 97 | 18.53 | 0.53 | 74.0 | 87 | 97 | 107.0 | 124.6 | 71 |
| Black | 548 | 63 | 146 | 102.33 | 102 | 14.29 | 0.61 | 78.0 | 93 | 102 | 112.0 | 125.6 | 39 |
| LatinX | 100 | 67 | 132 | 97.47 | 98 | 13.65 | 1.37 | 76.0 | 88 | 98 | 104.0 | 119.0 | 9 |
| Asian | 91 | 71 | 137 | 95.61 | 94 | 14.40 | 1.51 | 74.8 | 85 | 94 | 106.0 | 120.2 | 2 |
| Other | 21 | 73 | 126 | 100.00 | 96 | 14.35 | 3.13 | 78.6 | 90 | 96 | 109.0 | 121.2 | 4 |
| AIAN | 8 | 71 | 111 | 89.86 | 94 | 14.02 | 4.96 | 73.1 | 79 | 94 | 97.5 | 107.4 | 1 |
| PTNS | 5 | 69 | 134 | 93.20 | 91 | 24.93 | 11.15 | 70.8 | 78 | 91 | 94.0 | 126.0 | 0 |
| NHOPI | 4 | 75 | 96 | 86.67 | 89 | 10.69 | 5.35 | 76.4 | 82 | 89 | 92.5 | 95.3 | 1 |

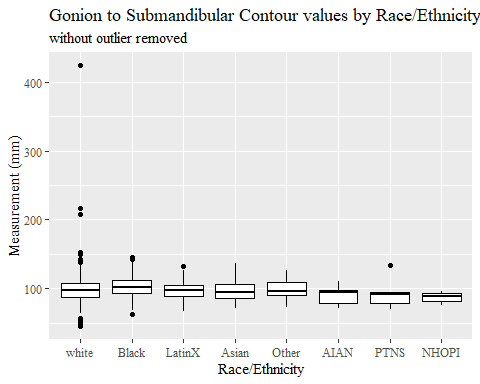
#%>% set\_header\_labels(values = list(GoSub\_C = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, GoSub\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=GoSub\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Gonion to Submandibular Contour values by Race/Ethnicity",  
 subtitle= "without outlier removed",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 127 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=GoSub\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Gonion to Submandibular Contour values by Race/Ethnicity",  
 subtitle= "without outlier removed",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 127 rows containing non-finite values (stat\_boxplot).



#GoSub\_C gender sumstats  
GoSub\_Cgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(GoSub\_C, na.rm = TRUE),  
 max = max(GoSub\_C, na.rm = TRUE),  
 mean = mean(GoSub\_C, na.rm = TRUE),  
 mdn = median(GoSub\_C, na.rm = TRUE),  
 sd = sd(GoSub\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(GoSub\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(GoSub\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(GoSub\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(GoSub\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(GoSub\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(GoSub\_C)))  
  
GoSub\_Cgender\_sumstats <- GoSub\_Cgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(GoSub\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Gonion to Submandibular Contour SumStats by Gender")

**Table** : Gonion to Submandibular Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 45 | 208 | 93.68 | 93.5 | 13.92 | 0.43 | 72.00 | 84 | 93.5 | 103.00 | 117.00 | 26 |
| Male | 939 | 52 | 424 | 105.70 | 105.0 | 18.46 | 0.60 | 83.00 | 96 | 105.0 | 114.00 | 130.00 | 101 |
| Non-binary or Other | 5 | 72 | 114 | 97.20 | 104.0 | 16.10 | 7.20 | 76.00 | 92 | 104.0 | 104.00 | 112.00 | 0 |
| Prefer not to say | 1 | 134 | 134 | 134.00 | 134.0 |  |  | 134.00 | 134 | 134.0 | 134.00 | 134.00 | 0 |
|  | 8 | 82 | 112 | 98.12 | 97.0 | 9.43 | 3.34 | 85.85 | 93 | 97.0 | 104.75 | 110.25 | 0 |

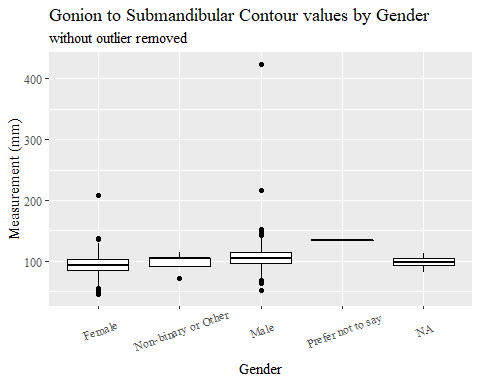
#%>% set\_header\_labels(values = list(GoSub\_C = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(GoSub\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Gonion to Submandibular Contour SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Gonion to Submandibular Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 45 | 208 | 93.68 | 93.5 | 13.92 | 0.43 | 72.00 | 84 | 93.5 | 103.00 | 117.00 | 26 |
| Male | 939 | 52 | 424 | 105.70 | 105.0 | 18.46 | 0.60 | 83.00 | 96 | 105.0 | 114.00 | 130.00 | 101 |
| Non-binary or Other | 5 | 72 | 114 | 97.20 | 104.0 | 16.10 | 7.20 | 76.00 | 92 | 104.0 | 104.00 | 112.00 | 0 |
| Prefer not to say | 1 | 134 | 134 | 134.00 | 134.0 |  |  | 134.00 | 134 | 134.0 | 134.00 | 134.00 | 0 |
|  | 8 | 82 | 112 | 98.12 | 97.0 | 9.43 | 3.34 | 85.85 | 93 | 97.0 | 104.75 | 110.25 | 0 |

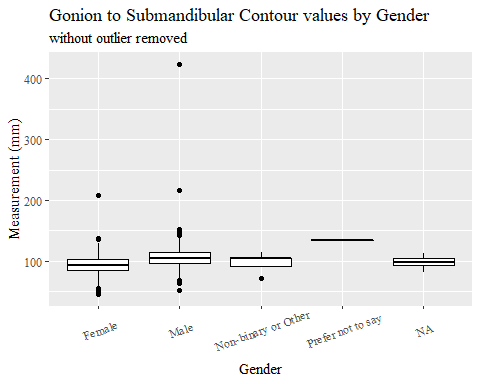
#%>% set\_header\_labels(values = list(GoSub\_C = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, GoSub\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=GoSub\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Gonion to Submandibular Contour values by Gender",  
 subtitle= "without outlier removed",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 127 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=GoSub\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Gonion to Submandibular Contour values by Gender",  
 subtitle= "without outlier removed",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 127 rows containing non-finite values (stat\_boxplot).



#GoSub\_C age group sumstats  
GoSub\_Cage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(GoSub\_C, na.rm = TRUE),  
 max = max(GoSub\_C, na.rm = TRUE),  
 mean = mean(GoSub\_C, na.rm = TRUE),  
 mdn = median(GoSub\_C, na.rm = TRUE),  
 sd = sd(GoSub\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(GoSub\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(GoSub\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(GoSub\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(GoSub\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(GoSub\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(GoSub\_C)))  
  
GoSub\_Cage\_sumstats <- GoSub\_Cage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(GoSub\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Gonion to Submandibular Contour SumStats by Age Group")

**Table** : Gonion to Submandibular Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 49 | 146 | 95.18 | 95 | 14.33 | 0.46 | 73.0 | 85 | 95 | 105 | 121.0 | 59 |
| 37-54 | 940 | 45 | 424 | 102.55 | 102 | 19.13 | 0.62 | 77.7 | 93 | 102 | 112 | 127.3 | 65 |
| 55-72 | 84 | 65 | 145 | 106.01 | 106 | 14.14 | 1.54 | 84.0 | 97 | 106 | 116 | 129.0 | 3 |
|  | 1 | 101 | 101 | 101.00 | 101 |  |  | 101.0 | 101 | 101 | 101 | 101.0 | 0 |

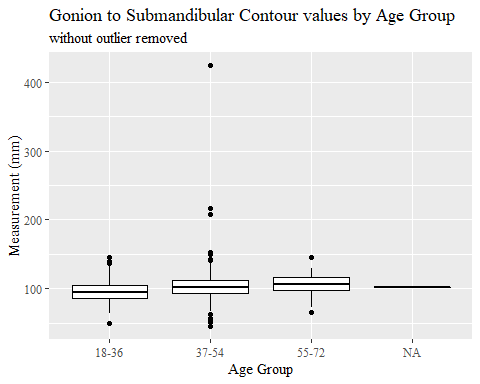
#%>% set\_header\_labels(values = list(GoSub\_C = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(GoSub\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Gonion to Submandibular Contour SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Gonion to Submandibular Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 49 | 146 | 95.18 | 95 | 14.33 | 0.46 | 73.0 | 85 | 95 | 105 | 121.0 | 59 |
| 37-54 | 940 | 45 | 424 | 102.55 | 102 | 19.13 | 0.62 | 77.7 | 93 | 102 | 112 | 127.3 | 65 |
| 55-72 | 84 | 65 | 145 | 106.01 | 106 | 14.14 | 1.54 | 84.0 | 97 | 106 | 116 | 129.0 | 3 |
|  | 1 | 101 | 101 | 101.00 | 101 |  |  | 101.0 | 101 | 101 | 101 | 101.0 | 0 |

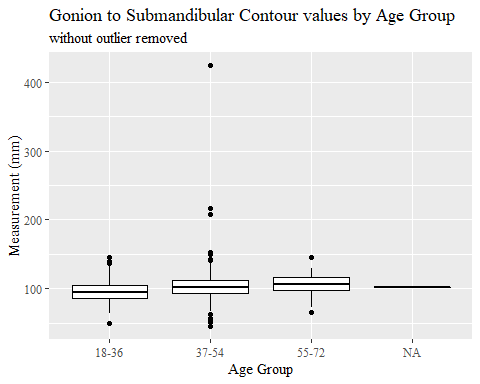
#%>% set\_header\_labels(values = list(GoSub\_C = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, GoSub\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=GoSub\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Gonion to Submandibular Contour values by Age Group",  
 subtitle= "without outlier removed",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 127 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=GoSub\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Gonion to Submandibular Contour values by Age Group",  
 subtitle= "without outlier removed",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 127 rows containing non-finite values (stat\_boxplot).



GoSub\_C WITH Outlier Removed!!!!

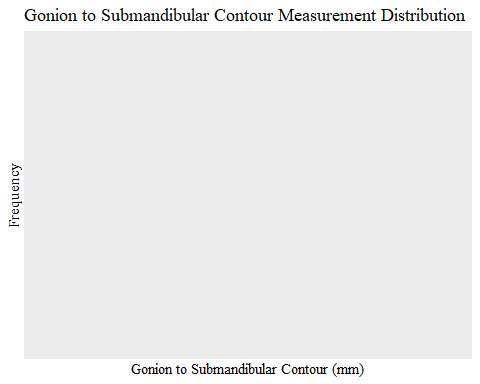
#REMOVING OUTLIER GoSubC  
  
max(headscan\_full$GoSub\_C, na.rm = TRUE)

## [1] 424

#400-20210129-009  
  
#CODE TO REMOVE OUTLIER, use later to remove from headscan\_full  
GoSub\_no\_out <- headscan\_full %>% mutate(GoSub\_C = replace(GoSub\_C, GoSub\_C>40, NA))

#histogram of all GoSub\_C values  
ggplot(data=GoSub\_no\_out, aes(x=GoSub\_C))+  
 geom\_bar(stat="bin", bins=20, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Gonion to Submandibular Contour Measurement Distribution",  
 y="Frequency",  
 x="Gonion to Submandibular Contour (mm)")

## Warning: Removed 2017 rows containing non-finite values (stat\_bin).



#GoSub\_C race/eth sumstats  
GoSub\_Crace\_sumstats1 <- GoSub\_no\_out %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(GoSub\_C, na.rm = TRUE),  
 max = max(GoSub\_C, na.rm = TRUE),  
 mean = mean(GoSub\_C, na.rm = TRUE),  
 mdn = median(GoSub\_C, na.rm = TRUE),  
 sd = sd(GoSub\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(GoSub\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(GoSub\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(GoSub\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(GoSub\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(GoSub\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(GoSub\_C)))

## Warning in min(GoSub\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(GoSub\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(GoSub\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(GoSub\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(GoSub\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(GoSub\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(GoSub\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(GoSub\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf

## Warning in max(GoSub\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(GoSub\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(GoSub\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(GoSub\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(GoSub\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(GoSub\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(GoSub\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(GoSub\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf

GoSub\_Crace\_sumstats1 <- GoSub\_Crace\_sumstats1 %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(GoSub\_Crace\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Gonion to Submandibular Contour SumStats by Race/Ethnicity")

**Table** : Gonion to Submandibular Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1,240 |
| Black | 548 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 548 |
| LatinX | 100 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 100 |
| Asian | 91 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 91 |
| Other | 21 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 21 |
| AIAN | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |
| PTNS | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| NHOPI | 4 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 4 |

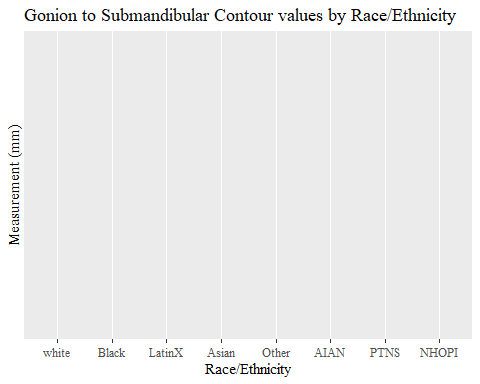
#%>% set\_header\_labels(values = list(GoSub\_C = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(GoSub\_Crace\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Gonion to Submandibular Contour SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Gonion to Submandibular Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1,240 |
| Black | 548 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 548 |
| LatinX | 100 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 100 |
| Asian | 91 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 91 |
| Other | 21 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 21 |
| AIAN | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |
| PTNS | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| NHOPI | 4 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 4 |

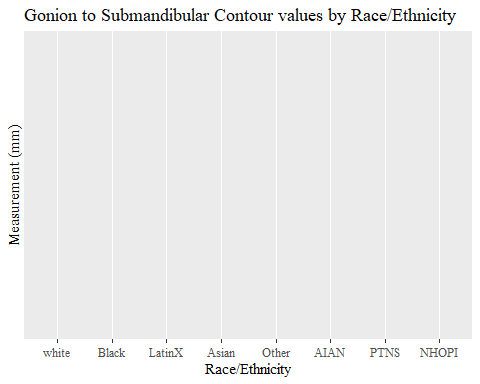
#%>% set\_header\_labels(values = list(GoSub\_C = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
GoSub\_no\_out %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, GoSub\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=GoSub\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Gonion to Submandibular Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=GoSub\_no\_out, aes(y=GoSub\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Gonion to Submandibular Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#GoSub\_C gender sumstats  
GoSub\_Cgender\_sumstats1 <- GoSub\_no\_out %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(GoSub\_C, na.rm = TRUE),  
 max = max(GoSub\_C, na.rm = TRUE),  
 mean = mean(GoSub\_C, na.rm = TRUE),  
 mdn = median(GoSub\_C, na.rm = TRUE),  
 sd = sd(GoSub\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(GoSub\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(GoSub\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(GoSub\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(GoSub\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(GoSub\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(GoSub\_C)))

## Warning in min(GoSub\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(GoSub\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(GoSub\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(GoSub\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(GoSub\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf

## Warning in max(GoSub\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(GoSub\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(GoSub\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(GoSub\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(GoSub\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf

GoSub\_Cgender\_sumstats1 <- GoSub\_Cgender\_sumstats1 %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(GoSub\_Cgender\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Gonion to Submandibular Contour SumStats by Gender")

**Table** : Gonion to Submandibular Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1,064 |
| Male | 939 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 939 |
| Non-binary or Other | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| Prefer not to say | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |
|  | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |

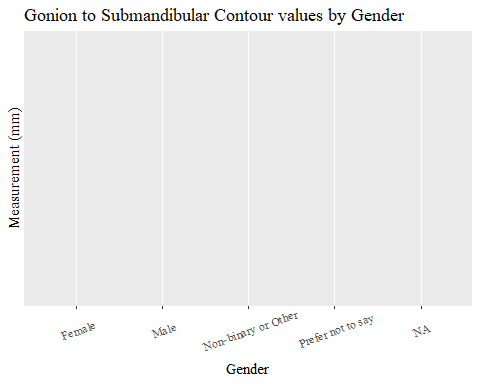
#%>% set\_header\_labels(values = list(GoSub\_C = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(GoSub\_Cgender\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Gonion to Submandibular Contour SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Gonion to Submandibular Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1,064 |
| Male | 939 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 939 |
| Non-binary or Other | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| Prefer not to say | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |
|  | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |

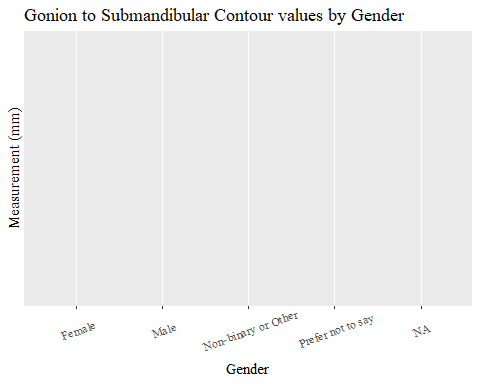
#%>% set\_header\_labels(values = list(GoSub\_C = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
GoSub\_no\_out %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, GoSub\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=GoSub\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Gonion to Submandibular Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=GoSub\_no\_out, aes(y=GoSub\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Gonion to Submandibular Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#GoSub\_C age group sumstats  
GoSub\_Cage\_sumstats1 <- GoSub\_no\_out %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(GoSub\_C, na.rm = TRUE),  
 max = max(GoSub\_C, na.rm = TRUE),  
 mean = mean(GoSub\_C, na.rm = TRUE),  
 mdn = median(GoSub\_C, na.rm = TRUE),  
 sd = sd(GoSub\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(GoSub\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(GoSub\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(GoSub\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(GoSub\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(GoSub\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(GoSub\_C)))

## Warning in min(GoSub\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(GoSub\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(GoSub\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(GoSub\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf

## Warning in max(GoSub\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(GoSub\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(GoSub\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(GoSub\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf

GoSub\_Cage\_sumstats1 <- GoSub\_Cage\_sumstats1 %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(GoSub\_Cage\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Gonion to Submandibular Contour SumStats by Age Group")

**Table** : Gonion to Submandibular Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 992 |
| 37-54 | 940 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 940 |
| 55-72 | 84 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 84 |
|  | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |

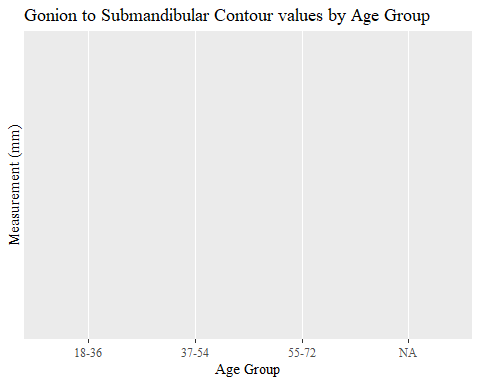
#%>% set\_header\_labels(values = list(GoSub\_C = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(GoSub\_Cage\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Gonion to Submandibular Contour SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Gonion to Submandibular Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 992 |
| 37-54 | 940 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 940 |
| 55-72 | 84 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 84 |
|  | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |

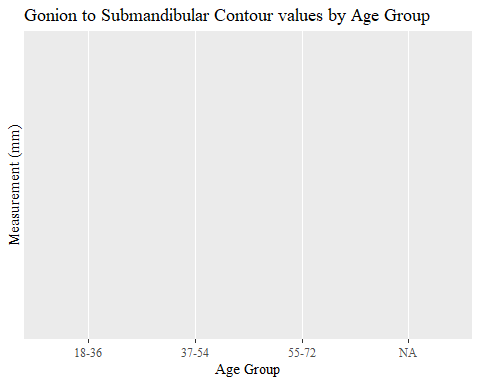
#%>% set\_header\_labels(values = list(GoSub\_C = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
GoSub\_no\_out %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, GoSub\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=GoSub\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Gonion to Submandibular Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=GoSub\_no\_out, aes(y=GoSub\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Gonion to Submandibular Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

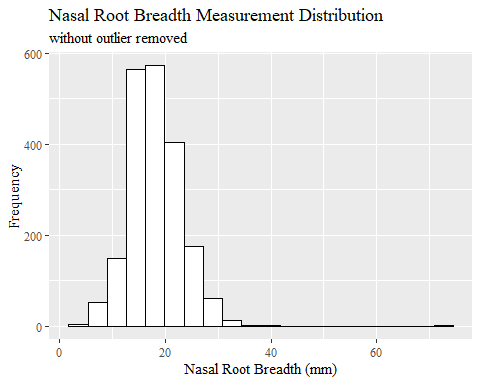
## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



NRB\_L WIHTOUT outlier removed

#histogram of all NRB\_L values  
ggplot(data=headscan\_full, aes(x=NRB\_L))+  
 geom\_bar(stat="bin", bins=20, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Nasal Root Breadth Measurement Distribution",  
 subtitle= "without outlier removed",  
 y="Frequency",  
 x="Nasal Root Breadth (mm)")

## Warning: Removed 15 rows containing non-finite values (stat\_bin).



#NRB\_L race/eth sumstats  
NRB\_Lrace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(NRB\_L, na.rm = TRUE),  
 max = max(NRB\_L, na.rm = TRUE),  
 mean = mean(NRB\_L, na.rm = TRUE),  
 mdn = median(NRB\_L, na.rm = TRUE),  
 sd = sd(NRB\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(NRB\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(NRB\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(NRB\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(NRB\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(NRB\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(NRB\_L)))   
  
NRB\_Lrace\_sumstats <- NRB\_Lrace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(NRB\_Lrace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Nasal Root Breadth SumStats by Race/Ethnicity")

**Table** : Nasal Root Breadth SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 3 | 72 | 17.31 | 17 | 4.69 | 0.13 | 10.0 | 14.0 | 17 | 20.0 | 25.0 | 8 |
| Black | 548 | 7 | 35 | 19.75 | 19 | 4.92 | 0.21 | 12.0 | 17.0 | 19 | 23.0 | 29.0 | 4 |
| LatinX | 100 | 8 | 29 | 17.48 | 17 | 4.41 | 0.44 | 11.0 | 14.0 | 17 | 20.0 | 25.1 | 1 |
| Asian | 91 | 8 | 31 | 16.61 | 16 | 4.98 | 0.52 | 10.0 | 13.0 | 16 | 19.0 | 26.0 | 1 |
| Other | 21 | 8 | 30 | 20.00 | 20 | 5.47 | 1.19 | 14.0 | 17.0 | 20 | 21.0 | 29.0 | 0 |
| AIAN | 8 | 8 | 28 | 18.14 | 18 | 6.20 | 2.19 | 10.1 | 15.5 | 18 | 21.0 | 25.9 | 1 |
| PTNS | 5 | 13 | 25 | 20.60 | 21 | 4.98 | 2.23 | 14.2 | 19.0 | 21 | 25.0 | 25.0 | 0 |
| NHOPI | 4 | 11 | 21 | 15.00 | 14 | 4.32 | 2.16 | 11.3 | 12.5 | 14 | 16.5 | 20.1 | 0 |

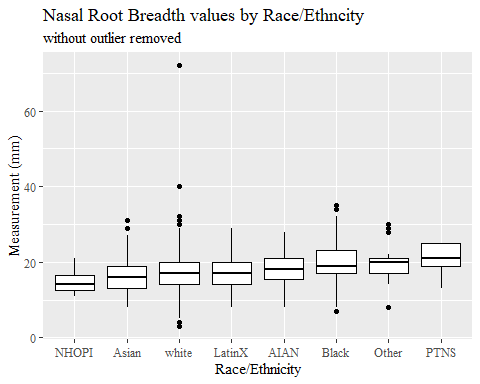
#%>% set\_header\_labels(values = list(NRB\_L = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(NRB\_Lrace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Nasal Root Breadth SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Nasal Root Breadth SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 3 | 72 | 17.31 | 17 | 4.69 | 0.13 | 10.0 | 14.0 | 17 | 20.0 | 25.0 | 8 |
| Black | 548 | 7 | 35 | 19.75 | 19 | 4.92 | 0.21 | 12.0 | 17.0 | 19 | 23.0 | 29.0 | 4 |
| LatinX | 100 | 8 | 29 | 17.48 | 17 | 4.41 | 0.44 | 11.0 | 14.0 | 17 | 20.0 | 25.1 | 1 |
| Asian | 91 | 8 | 31 | 16.61 | 16 | 4.98 | 0.52 | 10.0 | 13.0 | 16 | 19.0 | 26.0 | 1 |
| Other | 21 | 8 | 30 | 20.00 | 20 | 5.47 | 1.19 | 14.0 | 17.0 | 20 | 21.0 | 29.0 | 0 |
| AIAN | 8 | 8 | 28 | 18.14 | 18 | 6.20 | 2.19 | 10.1 | 15.5 | 18 | 21.0 | 25.9 | 1 |
| PTNS | 5 | 13 | 25 | 20.60 | 21 | 4.98 | 2.23 | 14.2 | 19.0 | 21 | 25.0 | 25.0 | 0 |
| NHOPI | 4 | 11 | 21 | 15.00 | 14 | 4.32 | 2.16 | 11.3 | 12.5 | 14 | 16.5 | 20.1 | 0 |

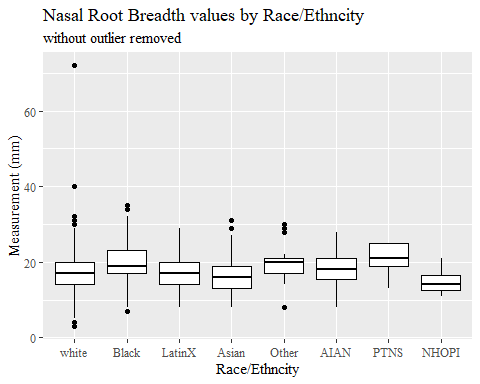
#%>% set\_header\_labels(values = list(NRB\_L = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, NRB\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=NRB\_L, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Nasal Root Breadth values by Race/Ethncity",  
 subtitle= "without outlier removed",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 15 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=NRB\_L, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Nasal Root Breadth values by Race/Ethncity",  
 subtitle= "without outlier removed",  
 y="Measurement (mm)",  
 x="Race/Ethncity")

## Warning: Removed 15 rows containing non-finite values (stat\_boxplot).



#NRB\_L gender sumstats  
NRB\_Lgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(NRB\_L, na.rm = TRUE),  
 max = max(NRB\_L, na.rm = TRUE),  
 mean = mean(NRB\_L, na.rm = TRUE),  
 mdn = median(NRB\_L, na.rm = TRUE),  
 sd = sd(NRB\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(NRB\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(NRB\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(NRB\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(NRB\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(NRB\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(NRB\_L)))  
  
NRB\_Lgender\_sumstats <- NRB\_Lgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(NRB\_Lgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Nasal Root Breadth SumStats by Gender")

**Table** : Nasal Root Breadth SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 4 | 32 | 18.18 | 18 | 4.76 | 0.15 | 10.0 | 15.00 | 18 | 21 | 27.00 | 9 |
| Male | 939 | 3 | 72 | 17.79 | 17 | 5.04 | 0.16 | 11.0 | 15.00 | 17 | 20 | 27.00 | 6 |
| Non-binary or Other | 5 | 8 | 20 | 14.20 | 13 | 5.17 | 2.31 | 8.6 | 11.00 | 13 | 19 | 19.80 | 0 |
| Prefer not to say | 1 | 19 | 19 | 19.00 | 19 |  |  | 19.0 | 19.00 | 19 | 19 | 19.00 | 0 |
|  | 8 | 14 | 19 | 17.00 | 17 | 1.51 | 0.53 | 14.7 | 16.75 | 17 | 18 | 18.65 | 0 |

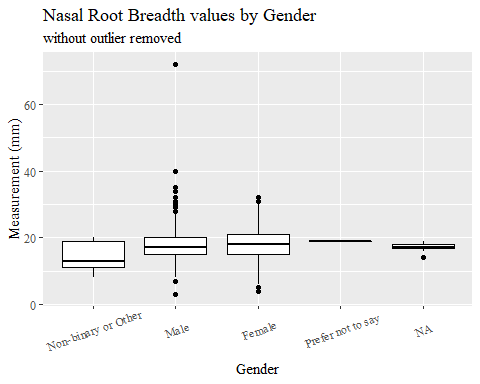
#%>% set\_header\_labels(values = list(NRB\_L = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(NRB\_Lgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Nasal Root Breadth SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Nasal Root Breadth SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 4 | 32 | 18.18 | 18 | 4.76 | 0.15 | 10.0 | 15.00 | 18 | 21 | 27.00 | 9 |
| Male | 939 | 3 | 72 | 17.79 | 17 | 5.04 | 0.16 | 11.0 | 15.00 | 17 | 20 | 27.00 | 6 |
| Non-binary or Other | 5 | 8 | 20 | 14.20 | 13 | 5.17 | 2.31 | 8.6 | 11.00 | 13 | 19 | 19.80 | 0 |
| Prefer not to say | 1 | 19 | 19 | 19.00 | 19 |  |  | 19.0 | 19.00 | 19 | 19 | 19.00 | 0 |
|  | 8 | 14 | 19 | 17.00 | 17 | 1.51 | 0.53 | 14.7 | 16.75 | 17 | 18 | 18.65 | 0 |

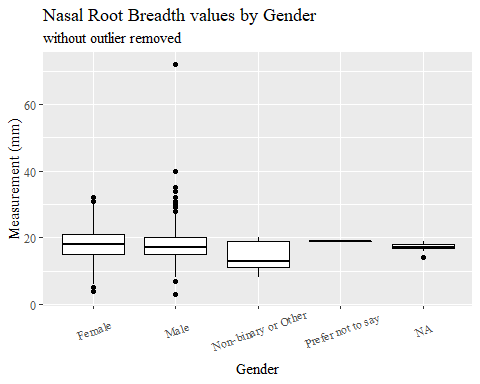
#%>% set\_header\_labels(values = list(NRB\_L = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, NRB\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=NRB\_L, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Nasal Root Breadth values by Gender",  
 subtitle= "without outlier removed",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 15 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=NRB\_L, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Nasal Root Breadth values by Gender",  
 subtitle= "without outlier removed",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 15 rows containing non-finite values (stat\_boxplot).



#NRB\_L age group sumstats  
NRB\_Lage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(NRB\_L, na.rm = TRUE),  
 max = max(NRB\_L, na.rm = TRUE),  
 mean = mean(NRB\_L, na.rm = TRUE),  
 mdn = median(NRB\_L, na.rm = TRUE),  
 sd = sd(NRB\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(NRB\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(NRB\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(NRB\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(NRB\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(NRB\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(NRB\_L)))  
  
NRB\_Lage\_sumstats <- NRB\_Lage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(NRB\_Lage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Nasal Root Breadth SumStats by Age Group")

**Table** : Nasal Root Breadth SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 4 | 72 | 18.20 | 18 | 5.04 | 0.16 | 11.0 | 15 | 18 | 21.0 | 27 | 6 |
| 37-54 | 940 | 3 | 35 | 17.82 | 18 | 4.69 | 0.15 | 10.0 | 15 | 18 | 20.0 | 26 | 8 |
| 55-72 | 84 | 8 | 32 | 17.42 | 17 | 5.26 | 0.57 | 9.1 | 14 | 17 | 19.5 | 28 | 1 |
|  | 1 | 13 | 13 | 13.00 | 13 |  |  | 13.0 | 13 | 13 | 13.0 | 13 | 0 |

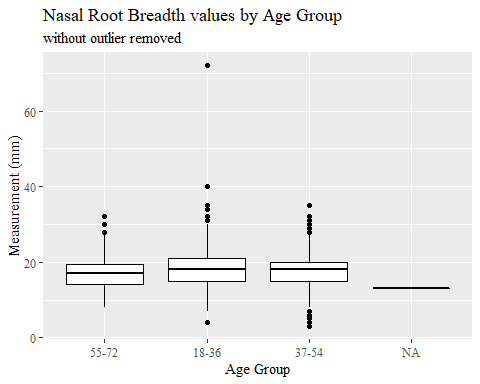
#%>% set\_header\_labels(values = list(NRB\_L = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(NRB\_Lage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Nasal Root Breadth SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Nasal Root Breadth SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 4 | 72 | 18.20 | 18 | 5.04 | 0.16 | 11.0 | 15 | 18 | 21.0 | 27 | 6 |
| 37-54 | 940 | 3 | 35 | 17.82 | 18 | 4.69 | 0.15 | 10.0 | 15 | 18 | 20.0 | 26 | 8 |
| 55-72 | 84 | 8 | 32 | 17.42 | 17 | 5.26 | 0.57 | 9.1 | 14 | 17 | 19.5 | 28 | 1 |
|  | 1 | 13 | 13 | 13.00 | 13 |  |  | 13.0 | 13 | 13 | 13.0 | 13 | 0 |

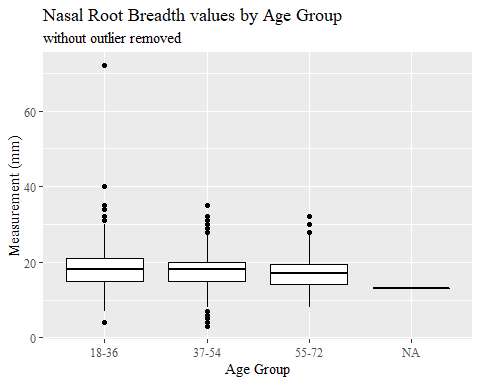
#%>% set\_header\_labels(values = list(NRB\_L = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, NRB\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=NRB\_L, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Nasal Root Breadth values by Age Group",  
 subtitle= "without outlier removed",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 15 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=NRB\_L, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Nasal Root Breadth values by Age Group",  
 subtitle= "without outlier removed",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 15 rows containing non-finite values (stat\_boxplot).



NRB\_L WITH Outlier removed

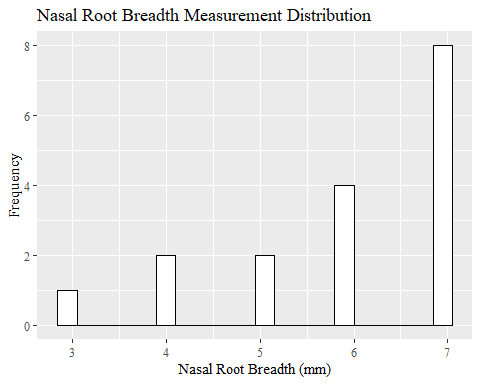
#REMOVING OUTLIER NRB\_L  
  
max(headscan\_full$NRB\_L, na.rm = TRUE)

## [1] 72

#CODE TO REMOVE OUTLIER, use later to remove from headscan\_full  
NRB\_L\_no\_out <- headscan\_full %>% mutate(NRB\_L = replace(NRB\_L, NRB\_L>7, NA))

#histogram of all NRB\_L values  
ggplot(data=NRB\_L\_no\_out, aes(x=NRB\_L))+  
 geom\_bar(stat="bin", bins=20, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Nasal Root Breadth Measurement Distribution",  
 y="Frequency",  
 x="Nasal Root Breadth (mm)")

## Warning: Removed 2000 rows containing non-finite values (stat\_bin).



#NRB\_L race/eth sumstats  
NRB\_Lrace\_sumstats1 <- NRB\_L\_no\_out %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(NRB\_L, na.rm = TRUE),  
 max = max(NRB\_L, na.rm = TRUE),  
 mean = mean(NRB\_L, na.rm = TRUE),  
 mdn = median(NRB\_L, na.rm = TRUE),  
 sd = sd(NRB\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(NRB\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(NRB\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(NRB\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(NRB\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(NRB\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(NRB\_L)))

## Warning in min(NRB\_L, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf  
  
## Warning in min(NRB\_L, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf  
  
## Warning in min(NRB\_L, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf  
  
## Warning in min(NRB\_L, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf  
  
## Warning in min(NRB\_L, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf  
  
## Warning in min(NRB\_L, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf

## Warning in max(NRB\_L, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf  
  
## Warning in max(NRB\_L, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf  
  
## Warning in max(NRB\_L, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf  
  
## Warning in max(NRB\_L, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf  
  
## Warning in max(NRB\_L, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf  
  
## Warning in max(NRB\_L, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf

NRB\_Lrace\_sumstats1 <- NRB\_Lrace\_sumstats1 %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(NRB\_Lrace\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Nasal Root Breadth SumStats by Race/Ethnicity")

**Table** : Nasal Root Breadth SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 3 | 7 | 5.88 | 6 | 1.31 | 0.04 | 3.75 | 5 | 6 | 7 | 7 | 1,224 |
| Black | 548 | 7 | 7 | 7.00 | 7 |  |  | 7.00 | 7 | 7 | 7 | 7 | 547 |
| LatinX | 100 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 100 |
| Asian | 91 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 91 |
| Other | 21 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 21 |
| AIAN | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |
| PTNS | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| NHOPI | 4 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 4 |

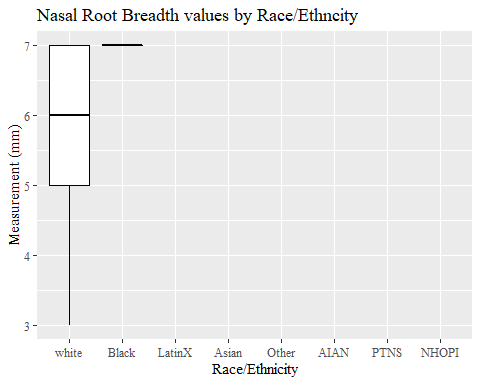
#%>% set\_header\_labels(values = list(NRB\_L = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(NRB\_Lrace\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Nasal Root Breadth SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Nasal Root Breadth SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 3 | 7 | 5.88 | 6 | 1.31 | 0.04 | 3.75 | 5 | 6 | 7 | 7 | 1,224 |
| Black | 548 | 7 | 7 | 7.00 | 7 |  |  | 7.00 | 7 | 7 | 7 | 7 | 547 |
| LatinX | 100 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 100 |
| Asian | 91 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 91 |
| Other | 21 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 21 |
| AIAN | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |
| PTNS | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| NHOPI | 4 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 4 |

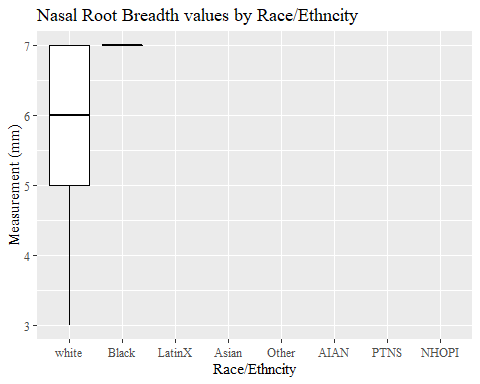
#%>% set\_header\_labels(values = list(NRB\_L = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
NRB\_L\_no\_out %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, NRB\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=NRB\_L, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Nasal Root Breadth values by Race/Ethncity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 2000 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=NRB\_L\_no\_out, aes(y=NRB\_L, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Nasal Root Breadth values by Race/Ethncity",  
 y="Measurement (mm)",  
 x="Race/Ethncity")

## Warning: Removed 2000 rows containing non-finite values (stat\_boxplot).



#NRB\_L gender sumstats  
NRB\_Lgender\_sumstats1 <- NRB\_L\_no\_out %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(NRB\_L, na.rm = TRUE),  
 max = max(NRB\_L, na.rm = TRUE),  
 mean = mean(NRB\_L, na.rm = TRUE),  
 mdn = median(NRB\_L, na.rm = TRUE),  
 sd = sd(NRB\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(NRB\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(NRB\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(NRB\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(NRB\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(NRB\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(NRB\_L)))

## Warning in min(NRB\_L, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf  
  
## Warning in min(NRB\_L, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf  
  
## Warning in min(NRB\_L, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf

## Warning in max(NRB\_L, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf  
  
## Warning in max(NRB\_L, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf  
  
## Warning in max(NRB\_L, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf

NRB\_Lgender\_sumstats1 <- NRB\_Lgender\_sumstats1 %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(NRB\_Lgender\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Nasal Root Breadth SumStats by Gender")

**Table** : Nasal Root Breadth SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 4 | 7 | 5.83 | 6 | 1.11 | 0.03 | 4.0 | 5 | 6 | 7 | 7 | 1,052 |
| Male | 939 | 3 | 7 | 6.20 | 7 | 1.79 | 0.06 | 3.8 | 7 | 7 | 7 | 7 | 934 |
| Non-binary or Other | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| Prefer not to say | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |
|  | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |

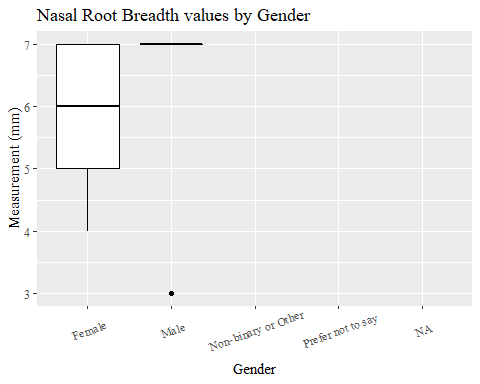
#%>% set\_header\_labels(values = list(NRB\_L = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(NRB\_Lgender\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Nasal Root Breadth SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Nasal Root Breadth SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 4 | 7 | 5.83 | 6 | 1.11 | 0.03 | 4.0 | 5 | 6 | 7 | 7 | 1,052 |
| Male | 939 | 3 | 7 | 6.20 | 7 | 1.79 | 0.06 | 3.8 | 7 | 7 | 7 | 7 | 934 |
| Non-binary or Other | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| Prefer not to say | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |
|  | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |

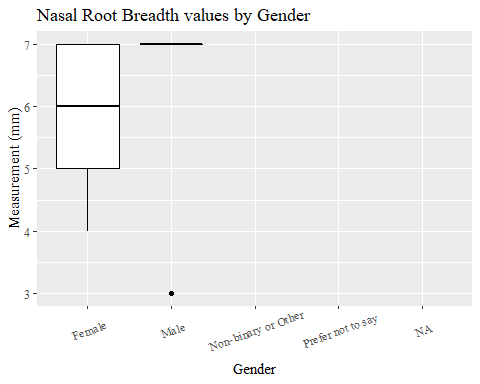
#%>% set\_header\_labels(values = list(NRB\_L = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
NRB\_L\_no\_out %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, NRB\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=NRB\_L, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Nasal Root Breadth values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 2000 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=NRB\_L\_no\_out, aes(y=NRB\_L, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Nasal Root Breadth values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 2000 rows containing non-finite values (stat\_boxplot).



#NRB\_L age group sumstats  
NRB\_Lage\_sumstats1 <- NRB\_L\_no\_out %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(NRB\_L, na.rm = TRUE),  
 max = max(NRB\_L, na.rm = TRUE),  
 mean = mean(NRB\_L, na.rm = TRUE),  
 mdn = median(NRB\_L, na.rm = TRUE),  
 sd = sd(NRB\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(NRB\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(NRB\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(NRB\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(NRB\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(NRB\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(NRB\_L)))

## Warning in min(NRB\_L, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf  
  
## Warning in min(NRB\_L, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf

## Warning in max(NRB\_L, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf  
  
## Warning in max(NRB\_L, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf

NRB\_Lage\_sumstats1 <- NRB\_Lage\_sumstats1 %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(NRB\_Lage\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Nasal Root Breadth SumStats by Age Group")

**Table** : Nasal Root Breadth SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 4 | 7 | 6.00 | 7 | 1.73 | 0.05 | 4.30 | 5.50 | 7 | 7 | 7 | 989 |
| 37-54 | 940 | 3 | 7 | 5.93 | 6 | 1.27 | 0.04 | 3.65 | 5.25 | 6 | 7 | 7 | 926 |
| 55-72 | 84 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 84 |
|  | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |

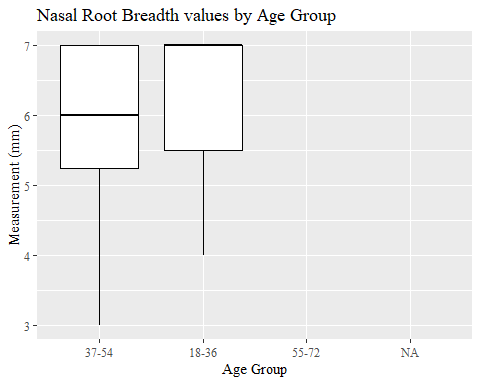
#%>% set\_header\_labels(values = list(NRB\_L = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(NRB\_Lage\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Nasal Root Breadth SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Nasal Root Breadth SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 4 | 7 | 6.00 | 7 | 1.73 | 0.05 | 4.30 | 5.50 | 7 | 7 | 7 | 989 |
| 37-54 | 940 | 3 | 7 | 5.93 | 6 | 1.27 | 0.04 | 3.65 | 5.25 | 6 | 7 | 7 | 926 |
| 55-72 | 84 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 84 |
|  | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |

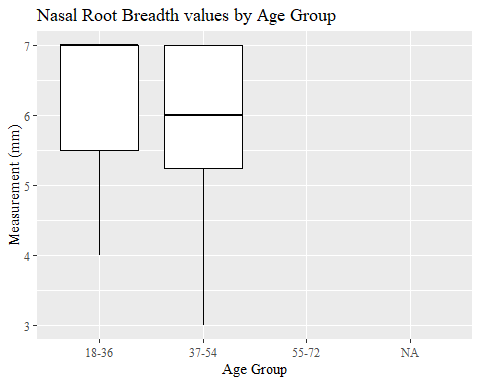
#%>% set\_header\_labels(values = list(NRB\_L = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
NRB\_L\_no\_out %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, NRB\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=NRB\_L, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Nasal Root Breadth values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 2000 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=NRB\_L\_no\_out, aes(y=NRB\_L, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Nasal Root Breadth values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

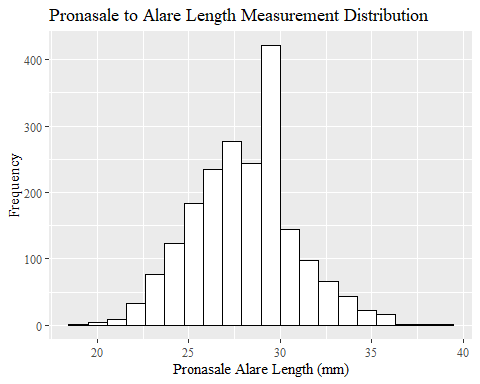
## Warning: Removed 2000 rows containing non-finite values (stat\_boxplot).



ProA\_L

#histogram of all ProA\_L values  
ggplot(data=headscan\_full, aes(x=ProA\_L))+  
 geom\_bar(stat="bin", bins=20, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Pronasale to Alare Length Measurement Distribution",  
 y="Frequency",  
 x="Pronasale Alare Length (mm)")

## Warning: Removed 16 rows containing non-finite values (stat\_bin).



#ProA\_L race/eth sumstats  
ProA\_Lrace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(ProA\_L, na.rm = TRUE),  
 max = max(ProA\_L, na.rm = TRUE),  
 mean = mean(ProA\_L, na.rm = TRUE),  
 mdn = median(ProA\_L, na.rm = TRUE),  
 sd = sd(ProA\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(ProA\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(ProA\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(ProA\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(ProA\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(ProA\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(ProA\_L)))   
  
ProA\_Lrace\_sumstats <- ProA\_Lrace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(ProA\_Lrace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Pronasale to Alare Length SumStats by Race/Ethnicity")

**Table** : Pronasale to Alare Length SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 19 | 37 | 27.82 | 28.0 | 2.96 | 0.08 | 23.0 | 26.0 | 28.0 | 30.00 | 33.00 | 8 |
| Black | 548 | 20 | 39 | 28.34 | 28.0 | 3.13 | 0.13 | 23.1 | 26.0 | 28.0 | 30.00 | 34.00 | 5 |
| LatinX | 100 | 22 | 34 | 28.08 | 28.0 | 3.05 | 0.30 | 23.0 | 26.0 | 28.0 | 30.00 | 33.10 | 1 |
| Asian | 91 | 22 | 36 | 27.12 | 27.0 | 3.18 | 0.33 | 23.0 | 25.0 | 27.0 | 29.00 | 33.00 | 1 |
| Other | 21 | 21 | 32 | 27.81 | 28.0 | 2.94 | 0.64 | 21.0 | 27.0 | 28.0 | 30.00 | 31.00 | 0 |
| AIAN | 8 | 22 | 32 | 27.29 | 27.0 | 3.04 | 1.07 | 23.2 | 26.5 | 27.0 | 28.50 | 31.10 | 1 |
| PTNS | 5 | 26 | 32 | 28.00 | 27.0 | 2.55 | 1.14 | 26.0 | 26.0 | 27.0 | 29.00 | 31.40 | 0 |
| NHOPI | 4 | 26 | 30 | 27.25 | 26.5 | 1.89 | 0.95 | 26.0 | 26.0 | 26.5 | 27.75 | 29.55 | 0 |

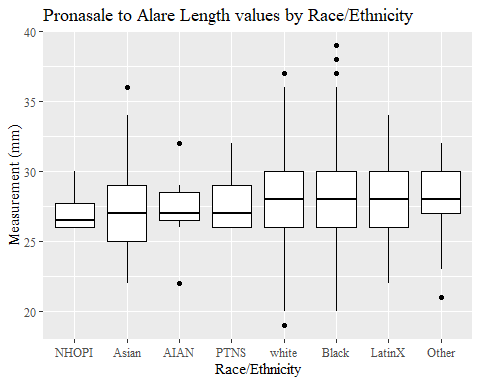
#%>% set\_header\_labels(values = list(ProA\_L = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(ProA\_Lrace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Pronasale to Alare Length SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Pronasale to Alare Length SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 19 | 37 | 27.82 | 28.0 | 2.96 | 0.08 | 23.0 | 26.0 | 28.0 | 30.00 | 33.00 | 8 |
| Black | 548 | 20 | 39 | 28.34 | 28.0 | 3.13 | 0.13 | 23.1 | 26.0 | 28.0 | 30.00 | 34.00 | 5 |
| LatinX | 100 | 22 | 34 | 28.08 | 28.0 | 3.05 | 0.30 | 23.0 | 26.0 | 28.0 | 30.00 | 33.10 | 1 |
| Asian | 91 | 22 | 36 | 27.12 | 27.0 | 3.18 | 0.33 | 23.0 | 25.0 | 27.0 | 29.00 | 33.00 | 1 |
| Other | 21 | 21 | 32 | 27.81 | 28.0 | 2.94 | 0.64 | 21.0 | 27.0 | 28.0 | 30.00 | 31.00 | 0 |
| AIAN | 8 | 22 | 32 | 27.29 | 27.0 | 3.04 | 1.07 | 23.2 | 26.5 | 27.0 | 28.50 | 31.10 | 1 |
| PTNS | 5 | 26 | 32 | 28.00 | 27.0 | 2.55 | 1.14 | 26.0 | 26.0 | 27.0 | 29.00 | 31.40 | 0 |
| NHOPI | 4 | 26 | 30 | 27.25 | 26.5 | 1.89 | 0.95 | 26.0 | 26.0 | 26.5 | 27.75 | 29.55 | 0 |

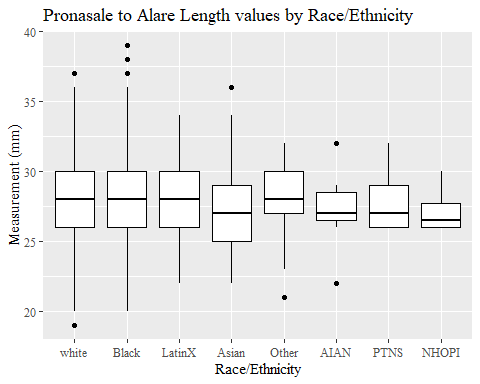
#%>% set\_header\_labels(values = list(ProA\_L = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, ProA\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=ProA\_L, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Pronasale to Alare Length values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 16 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=ProA\_L, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Pronasale to Alare Length values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 16 rows containing non-finite values (stat\_boxplot).



#ProA\_L gender sumstats  
ProA\_Lgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(ProA\_L, na.rm = TRUE),  
 max = max(ProA\_L, na.rm = TRUE),  
 mean = mean(ProA\_L, na.rm = TRUE),  
 mdn = median(ProA\_L, na.rm = TRUE),  
 sd = sd(ProA\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(ProA\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(ProA\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(ProA\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(ProA\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(ProA\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(ProA\_L)))  
  
ProA\_Lgender\_sumstats <- ProA\_Lgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(ProA\_Lgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Pronasale to Alare Length SumStats by Gender")

**Table** : Pronasale to Alare Length SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 19 | 36 | 26.47 | 26 | 2.48 | 0.08 | 23.0 | 25.00 | 26 | 28.00 | 31.00 | 11 |
| Male | 939 | 23 | 39 | 29.62 | 30 | 2.70 | 0.09 | 26.0 | 28.00 | 30 | 31.00 | 34.00 | 5 |
| Non-binary or Other | 5 | 23 | 28 | 26.80 | 28 | 2.17 | 0.97 | 23.8 | 27.00 | 28 | 28.00 | 28.00 | 0 |
| Prefer not to say | 1 | 26 | 26 | 26.00 | 26 |  |  | 26.0 | 26.00 | 26 | 26.00 | 26.00 | 0 |
|  | 8 | 22 | 30 | 25.50 | 25 | 2.33 | 0.82 | 22.7 | 24.75 | 25 | 26.25 | 28.95 | 0 |

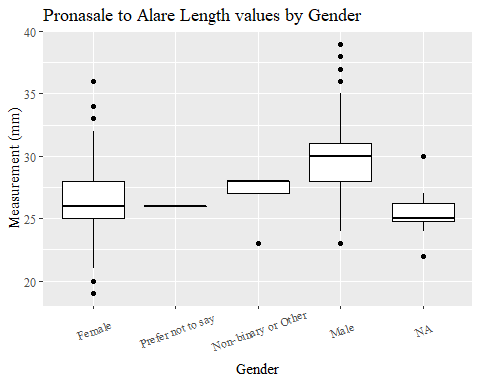
#%>% set\_header\_labels(values = list(ProA\_L = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(ProA\_Lgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Pronasale to Alare Length SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Pronasale to Alare Length SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 19 | 36 | 26.47 | 26 | 2.48 | 0.08 | 23.0 | 25.00 | 26 | 28.00 | 31.00 | 11 |
| Male | 939 | 23 | 39 | 29.62 | 30 | 2.70 | 0.09 | 26.0 | 28.00 | 30 | 31.00 | 34.00 | 5 |
| Non-binary or Other | 5 | 23 | 28 | 26.80 | 28 | 2.17 | 0.97 | 23.8 | 27.00 | 28 | 28.00 | 28.00 | 0 |
| Prefer not to say | 1 | 26 | 26 | 26.00 | 26 |  |  | 26.0 | 26.00 | 26 | 26.00 | 26.00 | 0 |
|  | 8 | 22 | 30 | 25.50 | 25 | 2.33 | 0.82 | 22.7 | 24.75 | 25 | 26.25 | 28.95 | 0 |

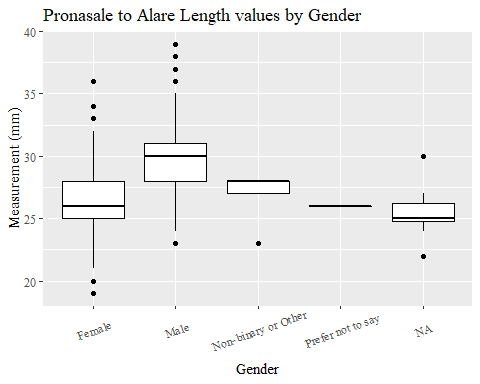
#%>% set\_header\_labels(values = list(ProA\_L = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, ProA\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=ProA\_L, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Pronasale to Alare Length values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 16 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=ProA\_L, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Pronasale to Alare Length values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 16 rows containing non-finite values (stat\_boxplot).



#ProA\_L age group sumstats  
ProA\_Lage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(ProA\_L, na.rm = TRUE),  
 max = max(ProA\_L, na.rm = TRUE),  
 mean = mean(ProA\_L, na.rm = TRUE),  
 mdn = median(ProA\_L, na.rm = TRUE),  
 sd = sd(ProA\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(ProA\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(ProA\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(ProA\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(ProA\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(ProA\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(ProA\_L)))  
  
ProA\_Lage\_sumstats <- ProA\_Lage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(ProA\_Lage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Pronasale Alare Length SumStats by Age Group")

**Table** : Pronasale Alare Length SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 20 | 37 | 27.55 | 27 | 2.87 | 0.09 | 23.0 | 26 | 27 | 29.75 | 32.0 | 6 |
| 37-54 | 940 | 19 | 38 | 28.26 | 28 | 3.12 | 0.10 | 23.5 | 26 | 28 | 30.00 | 34.0 | 9 |
| 55-72 | 84 | 22 | 39 | 28.99 | 29 | 3.20 | 0.35 | 25.0 | 27 | 29 | 31.00 | 34.9 | 1 |
|  | 1 | 25 | 25 | 25.00 | 25 |  |  | 25.0 | 25 | 25 | 25.00 | 25.0 | 0 |

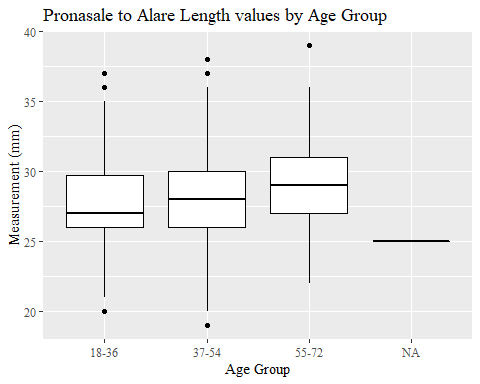
#%>% set\_header\_labels(values = list(ProA\_L = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(ProA\_Lage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Pronasale to Alare Length SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Pronasale to Alare Length SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 20 | 37 | 27.55 | 27 | 2.87 | 0.09 | 23.0 | 26 | 27 | 29.75 | 32.0 | 6 |
| 37-54 | 940 | 19 | 38 | 28.26 | 28 | 3.12 | 0.10 | 23.5 | 26 | 28 | 30.00 | 34.0 | 9 |
| 55-72 | 84 | 22 | 39 | 28.99 | 29 | 3.20 | 0.35 | 25.0 | 27 | 29 | 31.00 | 34.9 | 1 |
|  | 1 | 25 | 25 | 25.00 | 25 |  |  | 25.0 | 25 | 25 | 25.00 | 25.0 | 0 |

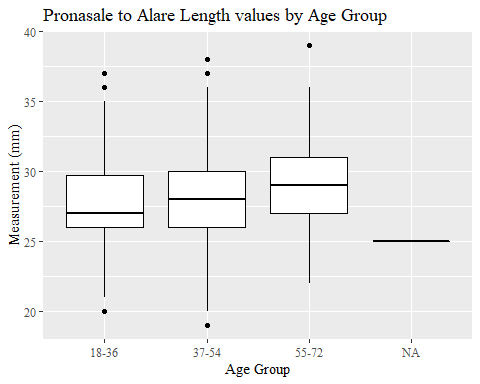
#%>% set\_header\_labels(values = list(ProA\_L = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, ProA\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=ProA\_L, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Pronasale to Alare Length values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 16 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=ProA\_L, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Pronasale to Alare Length values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

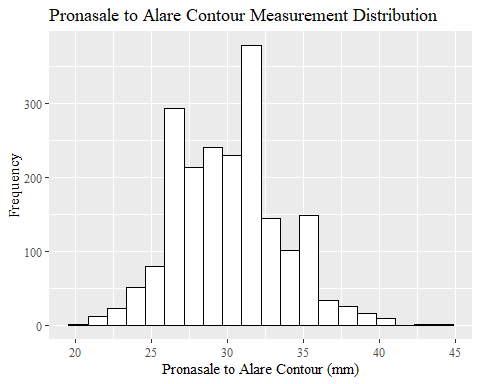
## Warning: Removed 16 rows containing non-finite values (stat\_boxplot).



ProA\_C

#histogram of all ProA\_C values  
ggplot(data=headscan\_full, aes(x=ProA\_C))+  
 geom\_bar(stat="bin", bins=20, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Pronasale to Alare Contour Measurement Distribution",  
 y="Frequency",  
 x="Pronasale to Alare Contour (mm)")

## Warning: Removed 16 rows containing non-finite values (stat\_bin).



#ProA\_C race/eth sumstats  
ProA\_Crace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(ProA\_C, na.rm = TRUE),  
 max = max(ProA\_C, na.rm = TRUE),  
 mean = mean(ProA\_C, na.rm = TRUE),  
 mdn = median(ProA\_C, na.rm = TRUE),  
 sd = sd(ProA\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(ProA\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(ProA\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(ProA\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(ProA\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(ProA\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(ProA\_C)))   
  
ProA\_Crace\_sumstats <- ProA\_Crace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(ProA\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Pronasale to Alare Contour SumStats by Race/Ethnicity")

**Table** : Pronasale to Alare Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 20 | 41 | 30.09 | 30.0 | 3.43 | 0.10 | 25.00 | 28.00 | 30.0 | 32.00 | 36.00 | 8 |
| Black | 548 | 21 | 44 | 30.34 | 30.0 | 3.64 | 0.16 | 25.00 | 28.00 | 30.0 | 33.00 | 37.00 | 5 |
| LatinX | 100 | 24 | 38 | 30.25 | 30.0 | 3.41 | 0.34 | 24.90 | 28.00 | 30.0 | 33.00 | 36.00 | 1 |
| Asian | 91 | 22 | 40 | 29.08 | 28.5 | 3.62 | 0.38 | 24.00 | 26.00 | 28.5 | 31.00 | 35.00 | 1 |
| Other | 21 | 22 | 35 | 29.76 | 30.0 | 3.55 | 0.77 | 22.00 | 29.00 | 30.0 | 32.00 | 35.00 | 0 |
| AIAN | 8 | 23 | 34 | 29.29 | 29.0 | 3.55 | 1.25 | 24.50 | 28.00 | 29.0 | 31.50 | 33.40 | 1 |
| PTNS | 5 | 27 | 35 | 30.60 | 30.0 | 3.05 | 1.36 | 27.40 | 29.00 | 30.0 | 32.00 | 34.40 | 0 |
| NHOPI | 4 | 27 | 34 | 29.50 | 28.5 | 3.11 | 1.55 | 27.15 | 27.75 | 28.5 | 30.25 | 33.25 | 0 |

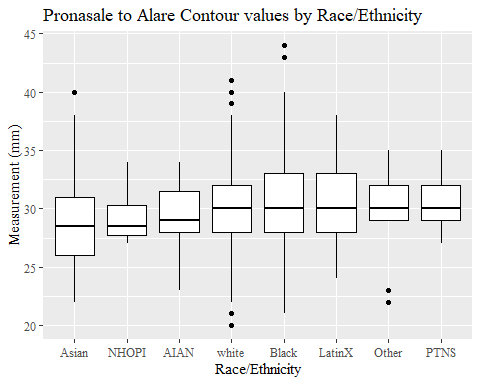
#%>% set\_header\_Cabels(values = list(ProA\_C = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(ProA\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Pronasale to Alare Contour SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Pronasale to Alare Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 20 | 41 | 30.09 | 30.0 | 3.43 | 0.10 | 25.00 | 28.00 | 30.0 | 32.00 | 36.00 | 8 |
| Black | 548 | 21 | 44 | 30.34 | 30.0 | 3.64 | 0.16 | 25.00 | 28.00 | 30.0 | 33.00 | 37.00 | 5 |
| LatinX | 100 | 24 | 38 | 30.25 | 30.0 | 3.41 | 0.34 | 24.90 | 28.00 | 30.0 | 33.00 | 36.00 | 1 |
| Asian | 91 | 22 | 40 | 29.08 | 28.5 | 3.62 | 0.38 | 24.00 | 26.00 | 28.5 | 31.00 | 35.00 | 1 |
| Other | 21 | 22 | 35 | 29.76 | 30.0 | 3.55 | 0.77 | 22.00 | 29.00 | 30.0 | 32.00 | 35.00 | 0 |
| AIAN | 8 | 23 | 34 | 29.29 | 29.0 | 3.55 | 1.25 | 24.50 | 28.00 | 29.0 | 31.50 | 33.40 | 1 |
| PTNS | 5 | 27 | 35 | 30.60 | 30.0 | 3.05 | 1.36 | 27.40 | 29.00 | 30.0 | 32.00 | 34.40 | 0 |
| NHOPI | 4 | 27 | 34 | 29.50 | 28.5 | 3.11 | 1.55 | 27.15 | 27.75 | 28.5 | 30.25 | 33.25 | 0 |

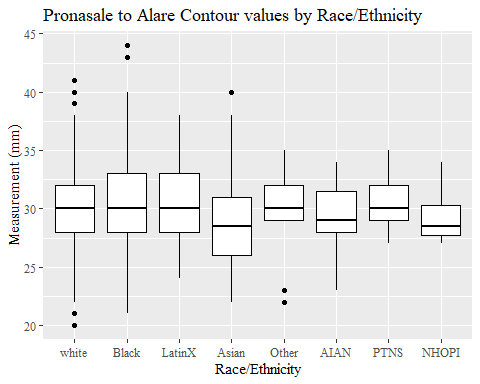
#%>% set\_header\_Cabels(values = list(ProA\_C = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, ProA\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=ProA\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Pronasale to Alare Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 16 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=ProA\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Pronasale to Alare Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 16 rows containing non-finite values (stat\_boxplot).



#ProA\_C gender sumstats  
ProA\_Cgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(ProA\_C, na.rm = TRUE),  
 max = max(ProA\_C, na.rm = TRUE),  
 mean = mean(ProA\_C, na.rm = TRUE),  
 mdn = median(ProA\_C, na.rm = TRUE),  
 sd = sd(ProA\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(ProA\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(ProA\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(ProA\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(ProA\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(ProA\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(ProA\_C)))  
  
ProA\_Cgender\_sumstats <- ProA\_Cgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(ProA\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Pronasale to Alare Contour SumStats by Gender")

**Table** : Pronasale to Alare Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 20 | 41 | 28.51 | 28.0 | 2.94 | 0.09 | 24.00 | 27 | 28.0 | 30.00 | 34.0 | 11 |
| Male | 939 | 23 | 44 | 31.95 | 32.0 | 3.17 | 0.10 | 27.00 | 30 | 32.0 | 34.00 | 38.0 | 5 |
| Non-binary or Other | 5 | 25 | 31 | 29.20 | 30.0 | 2.49 | 1.11 | 25.80 | 29 | 30.0 | 31.00 | 31.0 | 0 |
| Prefer not to say | 1 | 27 | 27 | 27.00 | 27.0 |  |  | 27.00 | 27 | 27.0 | 27.00 | 27.0 | 0 |
|  | 8 | 23 | 33 | 27.50 | 27.5 | 2.88 | 1.02 | 24.05 | 26 | 27.5 | 28.25 | 31.6 | 0 |

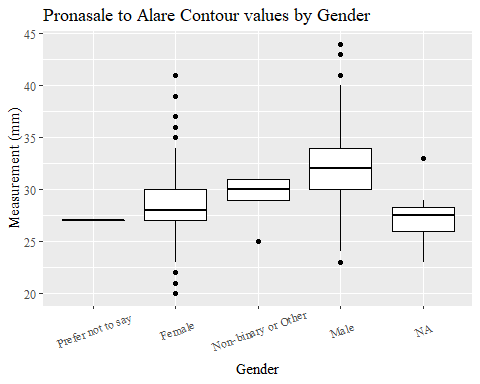
#%>% set\_header\_Cabels(values = list(ProA\_C = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(ProA\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Pronasale to Alare Contour SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Pronasale to Alare Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 20 | 41 | 28.51 | 28.0 | 2.94 | 0.09 | 24.00 | 27 | 28.0 | 30.00 | 34.0 | 11 |
| Male | 939 | 23 | 44 | 31.95 | 32.0 | 3.17 | 0.10 | 27.00 | 30 | 32.0 | 34.00 | 38.0 | 5 |
| Non-binary or Other | 5 | 25 | 31 | 29.20 | 30.0 | 2.49 | 1.11 | 25.80 | 29 | 30.0 | 31.00 | 31.0 | 0 |
| Prefer not to say | 1 | 27 | 27 | 27.00 | 27.0 |  |  | 27.00 | 27 | 27.0 | 27.00 | 27.0 | 0 |
|  | 8 | 23 | 33 | 27.50 | 27.5 | 2.88 | 1.02 | 24.05 | 26 | 27.5 | 28.25 | 31.6 | 0 |

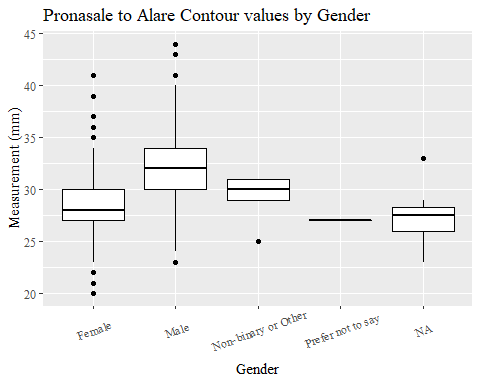
#%>% set\_header\_Cabels(values = list(ProA\_C = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, ProA\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=ProA\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Pronasale to Alare Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 16 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=ProA\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Pronasale to Alare Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 16 rows containing non-finite values (stat\_boxplot).



#ProA\_C age group sumstats  
ProA\_Cage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(ProA\_C, na.rm = TRUE),  
 max = max(ProA\_C, na.rm = TRUE),  
 mean = mean(ProA\_C, na.rm = TRUE),  
 mdn = median(ProA\_C, na.rm = TRUE),  
 sd = sd(ProA\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(ProA\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(ProA\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(ProA\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(ProA\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(ProA\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(ProA\_C)))  
  
ProA\_Cage\_sumstats <- ProA\_Cage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(ProA\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Pronasale to Alare Contour SumStats by Age Group")

**Table** : Pronasale to Alare Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 21 | 41 | 29.65 | 29 | 3.34 | 0.11 | 24 | 27 | 29 | 32.0 | 35 | 6 |
| 37-54 | 940 | 20 | 43 | 30.51 | 30 | 3.59 | 0.12 | 25 | 28 | 30 | 33.0 | 37 | 9 |
| 55-72 | 84 | 23 | 44 | 31.13 | 31 | 3.60 | 0.39 | 26 | 29 | 31 | 33.5 | 37 | 1 |
|  | 1 | 27 | 27 | 27.00 | 27 |  |  | 27 | 27 | 27 | 27.0 | 27 | 0 |

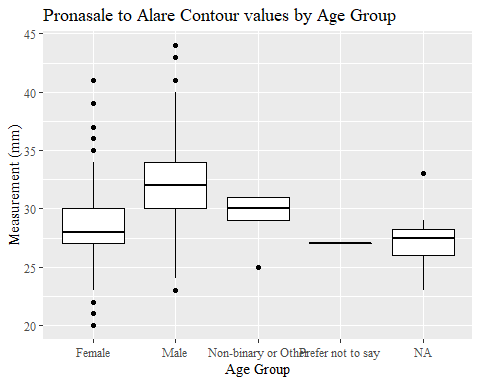
#%>% set\_header\_Cabels(values = list(ProA\_C = "Alare/AlareCont"))  
  
#Autofit Width Table TNR  
flextable(ProA\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Pronasale to Alare Contour SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Pronasale to Alare Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 21 | 41 | 29.65 | 29 | 3.34 | 0.11 | 24 | 27 | 29 | 32.0 | 35 | 6 |
| 37-54 | 940 | 20 | 43 | 30.51 | 30 | 3.59 | 0.12 | 25 | 28 | 30 | 33.0 | 37 | 9 |
| 55-72 | 84 | 23 | 44 | 31.13 | 31 | 3.60 | 0.39 | 26 | 29 | 31 | 33.5 | 37 | 1 |
|  | 1 | 27 | 27 | 27.00 | 27 |  |  | 27 | 27 | 27 | 27.0 | 27 | 0 |

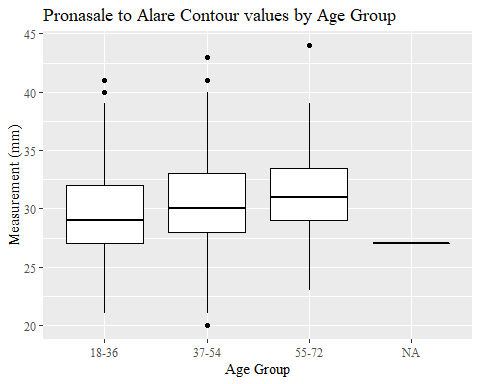
#%>% set\_header\_Cabels(values = list(ProA\_C = "Alare/AlareCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, ProA\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=ProA\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Pronasale to Alare Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 16 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=ProA\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Pronasale to Alare Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

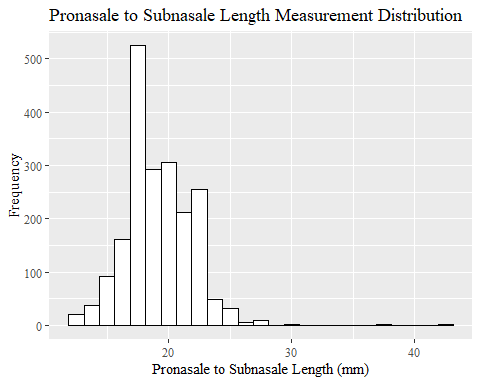
## Warning: Removed 16 rows containing non-finite values (stat\_boxplot).



ProS\_L

#histogram of all ProS\_L values  
ggplot(data=headscan\_full, aes(x=ProS\_L))+  
 geom\_bar(stat="bin", bins=25, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Pronasale to Subnasale Length Measurement Distribution",  
 y="Frequency",  
 x="Pronasale to Subnasale Length (mm)")

## Warning: Removed 19 rows containing non-finite values (stat\_bin).



#ProS\_L race/eth sumstats  
ProS\_Lrace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(ProS\_L, na.rm = TRUE),  
 max = max(ProS\_L, na.rm = TRUE),  
 mean = mean(ProS\_L, na.rm = TRUE),  
 mdn = median(AA\_C, na.rm = TRUE),  
 sd = sd(ProS\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(ProS\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(ProS\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(ProS\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(ProS\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(ProS\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(ProS\_L)))   
  
ProS\_Lrace\_sumstats <- ProS\_Lrace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(ProS\_Lrace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Pronasale to Subnasale Length SumStats by Race/Ethnicity")

**Table** : Pronasale to Subnasale Length SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 13 | 30 | 19.70 | 61 | 2.54 | 0.07 | 16 | 18.0 | 20 | 21.0 | 24.0 | 11 |
| Black | 548 | 12 | 42 | 18.19 | 61 | 2.73 | 0.12 | 14 | 16.5 | 18 | 20.0 | 22.0 | 5 |
| LatinX | 100 | 13 | 25 | 18.78 | 62 | 2.31 | 0.23 | 15 | 17.0 | 19 | 20.0 | 22.0 | 1 |
| Asian | 91 | 13 | 42 | 18.41 | 58 | 3.26 | 0.34 | 15 | 17.0 | 18 | 20.0 | 22.0 | 1 |
| Other | 21 | 12 | 22 | 18.05 | 62 | 2.58 | 0.56 | 13 | 16.0 | 19 | 19.0 | 22.0 | 0 |
| AIAN | 8 | 17 | 19 | 18.00 | 61 | 0.82 | 0.29 | 17 | 17.5 | 18 | 18.5 | 19.0 | 1 |
| PTNS | 5 | 16 | 19 | 17.00 | 61 | 1.22 | 0.55 | 16 | 16.0 | 17 | 17.0 | 18.6 | 0 |
| NHOPI | 4 | 16 | 24 | 19.00 | 59 | 3.83 | 1.91 | 16 | 16.0 | 18 | 21.0 | 23.4 | 0 |

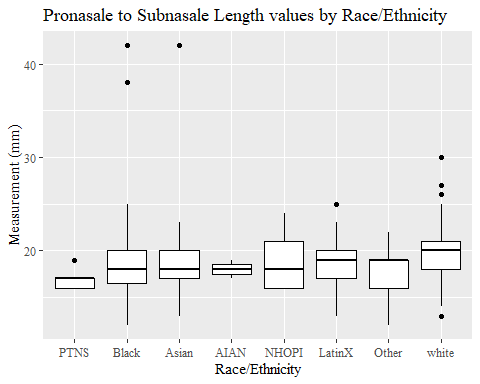
#%>% set\_header\_labels(values = list(ProS\_L = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(ProS\_Lrace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Pronasale to Subnasale Length SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Pronasale to Subnasale Length SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 13 | 30 | 19.70 | 61 | 2.54 | 0.07 | 16 | 18.0 | 20 | 21.0 | 24.0 | 11 |
| Black | 548 | 12 | 42 | 18.19 | 61 | 2.73 | 0.12 | 14 | 16.5 | 18 | 20.0 | 22.0 | 5 |
| LatinX | 100 | 13 | 25 | 18.78 | 62 | 2.31 | 0.23 | 15 | 17.0 | 19 | 20.0 | 22.0 | 1 |
| Asian | 91 | 13 | 42 | 18.41 | 58 | 3.26 | 0.34 | 15 | 17.0 | 18 | 20.0 | 22.0 | 1 |
| Other | 21 | 12 | 22 | 18.05 | 62 | 2.58 | 0.56 | 13 | 16.0 | 19 | 19.0 | 22.0 | 0 |
| AIAN | 8 | 17 | 19 | 18.00 | 61 | 0.82 | 0.29 | 17 | 17.5 | 18 | 18.5 | 19.0 | 1 |
| PTNS | 5 | 16 | 19 | 17.00 | 61 | 1.22 | 0.55 | 16 | 16.0 | 17 | 17.0 | 18.6 | 0 |
| NHOPI | 4 | 16 | 24 | 19.00 | 59 | 3.83 | 1.91 | 16 | 16.0 | 18 | 21.0 | 23.4 | 0 |

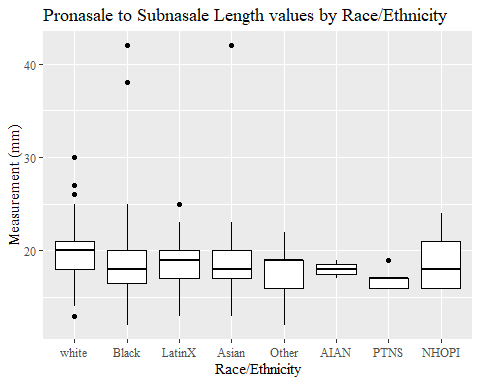
#%>% set\_header\_labels(values = list(ProS\_L = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, ProS\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=ProS\_L, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Pronasale to Subnasale Length values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 19 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=ProS\_L, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Pronasale to Subnasale Length values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 19 rows containing non-finite values (stat\_boxplot).



#ProS\_L gender sumstats  
ProS\_Lgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(ProS\_L, na.rm = TRUE),  
 max = max(ProS\_L, na.rm = TRUE),  
 mean = mean(ProS\_L, na.rm = TRUE),  
 mdn = median(ProS\_L, na.rm = TRUE),  
 sd = sd(ProS\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(ProS\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(ProS\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(ProS\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(ProS\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(ProS\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(ProS\_L)))  
  
ProS\_Lgender\_sumstats <- ProS\_Lgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(ProS\_Lgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Pronasale to Subnasale Length SumStats by Gender")

**Table** : Pronasale to Subnasale Length SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 12 | 42 | 18.74 | 19 | 2.74 | 0.08 | 15.00 | 17.00 | 19 | 20 | 23.0 | 11 |
| Male | 939 | 12 | 30 | 19.65 | 20 | 2.58 | 0.08 | 16.00 | 18.00 | 20 | 21 | 24.0 | 8 |
| Non-binary or Other | 5 | 15 | 20 | 17.80 | 18 | 1.79 | 0.80 | 15.60 | 18.00 | 18 | 18 | 19.6 | 0 |
| Prefer not to say | 1 | 17 | 17 | 17.00 | 17 |  |  | 17.00 | 17.00 | 17 | 17 | 17.0 | 0 |
|  | 8 | 14 | 21 | 17.75 | 17 | 2.87 | 1.01 | 14.35 | 15.75 | 17 | 21 | 21.0 | 0 |

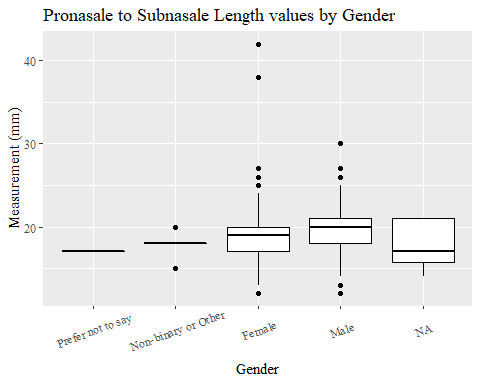
#%>% set\_header\_labels(values = list(ProS\_L = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(ProS\_Lgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Pronasale to Subnasale Length SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Pronasale to Subnasale Length SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 12 | 42 | 18.74 | 19 | 2.74 | 0.08 | 15.00 | 17.00 | 19 | 20 | 23.0 | 11 |
| Male | 939 | 12 | 30 | 19.65 | 20 | 2.58 | 0.08 | 16.00 | 18.00 | 20 | 21 | 24.0 | 8 |
| Non-binary or Other | 5 | 15 | 20 | 17.80 | 18 | 1.79 | 0.80 | 15.60 | 18.00 | 18 | 18 | 19.6 | 0 |
| Prefer not to say | 1 | 17 | 17 | 17.00 | 17 |  |  | 17.00 | 17.00 | 17 | 17 | 17.0 | 0 |
|  | 8 | 14 | 21 | 17.75 | 17 | 2.87 | 1.01 | 14.35 | 15.75 | 17 | 21 | 21.0 | 0 |

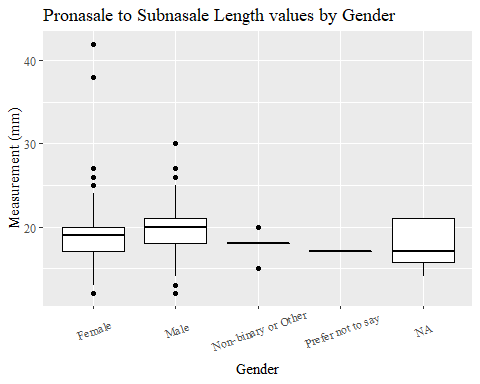
#%>% set\_header\_labels(values = list(ProS\_L = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, ProS\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=ProS\_L, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Pronasale to Subnasale Length values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 19 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=ProS\_L, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Pronasale to Subnasale Length values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 19 rows containing non-finite values (stat\_boxplot).



#ProS\_L age group sumstats  
ProS\_Lage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(ProS\_L, na.rm = TRUE),  
 max = max(ProS\_L, na.rm = TRUE),  
 mean = mean(ProS\_L, na.rm = TRUE),  
 mdn = median(ProS\_L, na.rm = TRUE),  
 sd = sd(ProS\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(ProS\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(ProS\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(ProS\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(ProS\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(ProS\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(ProS\_L)))  
  
ProS\_Lage\_sumstats <- ProS\_Lage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(ProS\_Lage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Pronasale Subnasale Length SumStats by Age Group")

**Table** : Pronasale Subnasale Length SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 12 | 30 | 18.73 | 19 | 2.45 | 0.08 | 15 | 17 | 19 | 20 | 23 | 6 |
| 37-54 | 940 | 12 | 42 | 19.54 | 19 | 2.91 | 0.09 | 15 | 18 | 19 | 21 | 24 | 11 |
| 55-72 | 84 | 13 | 26 | 19.80 | 20 | 2.44 | 0.27 | 16 | 18 | 20 | 21 | 24 | 2 |
|  | 1 | 23 | 23 | 23.00 | 23 |  |  | 23 | 23 | 23 | 23 | 23 | 0 |

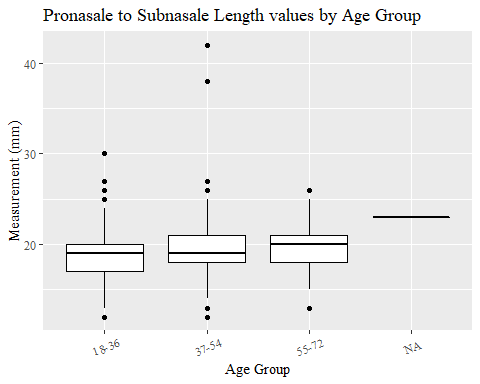
#%>% set\_header\_labels(values = list(ProS\_L = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(ProS\_Lage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Pronasale Subnasale Length SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Pronasale Subnasale Length SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 12 | 30 | 18.73 | 19 | 2.45 | 0.08 | 15 | 17 | 19 | 20 | 23 | 6 |
| 37-54 | 940 | 12 | 42 | 19.54 | 19 | 2.91 | 0.09 | 15 | 18 | 19 | 21 | 24 | 11 |
| 55-72 | 84 | 13 | 26 | 19.80 | 20 | 2.44 | 0.27 | 16 | 18 | 20 | 21 | 24 | 2 |
|  | 1 | 23 | 23 | 23.00 | 23 |  |  | 23 | 23 | 23 | 23 | 23 | 0 |

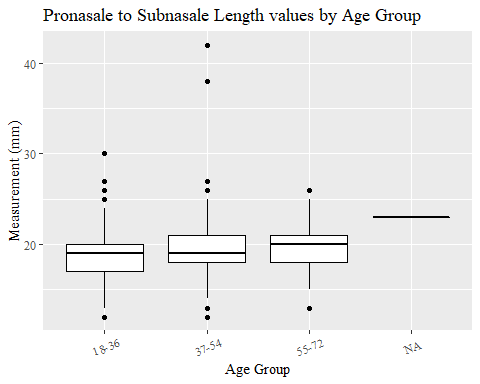
#%>% set\_header\_labels(values = list(ProS\_L = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, ProS\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=ProS\_L, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Pronasale to Subnasale Length values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 19 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=ProS\_L, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Pronasale to Subnasale Length values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

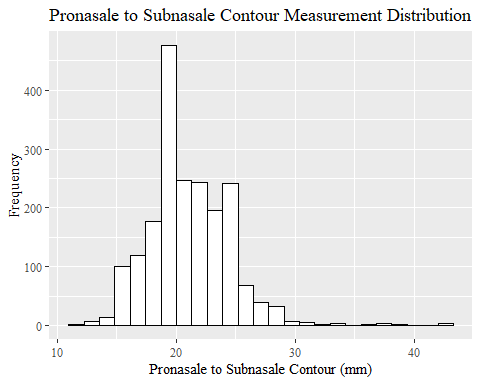
## Warning: Removed 19 rows containing non-finite values (stat\_boxplot).



ProS\_C

#histogram of all ProS\_C values  
ggplot(data=headscan\_full, aes(x=ProS\_C))+  
 geom\_bar(stat="bin", bins=25, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Pronasale to Subnasale Contour Measurement Distribution",  
 y="Frequency",  
 x="Pronasale to Subnasale Contour (mm)")

## Warning: Removed 31 rows containing non-finite values (stat\_bin).



#ProS\_C race/eth sumstats  
ProS\_Crace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(ProS\_C, na.rm = TRUE),  
 max = max(ProS\_C, na.rm = TRUE),  
 mean = mean(ProS\_C, na.rm = TRUE),  
 mdn = median(ProS\_C, na.rm = TRUE),  
 sd = sd(ProS\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(ProS\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(ProS\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(ProS\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(ProS\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(ProS\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(ProS\_C)))   
  
ProS\_Crace\_sumstats <- ProS\_Crace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(ProS\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Pronasale to Subnasale Contour SumStats by Race/Ethnicity")

**Table** : Pronasale to Subnasale Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 13 | 43 | 21.64 | 22.0 | 3.11 | 0.09 | 17.00 | 19.00 | 22.0 | 24.0 | 27.00 | 21 |
| Black | 548 | 12 | 42 | 20.20 | 20.0 | 3.64 | 0.16 | 15.00 | 18.00 | 20.0 | 22.0 | 25.00 | 7 |
| LatinX | 100 | 14 | 29 | 20.70 | 20.0 | 2.88 | 0.29 | 16.00 | 19.00 | 20.0 | 22.0 | 25.00 | 1 |
| Asian | 91 | 14 | 42 | 20.18 | 20.0 | 3.73 | 0.39 | 16.00 | 18.00 | 20.0 | 22.0 | 25.55 | 1 |
| Other | 21 | 13 | 26 | 19.62 | 21.0 | 3.12 | 0.68 | 13.00 | 18.00 | 21.0 | 21.0 | 24.00 | 0 |
| AIAN | 8 | 19 | 23 | 20.29 | 20.0 | 1.25 | 0.44 | 19.30 | 20.00 | 20.0 | 20.0 | 22.10 | 1 |
| PTNS | 5 | 17 | 21 | 18.40 | 18.0 | 1.52 | 0.68 | 17.20 | 18.00 | 18.0 | 18.0 | 20.40 | 0 |
| NHOPI | 4 | 17 | 29 | 21.75 | 20.5 | 5.50 | 2.75 | 17.15 | 17.75 | 20.5 | 24.5 | 28.10 | 0 |

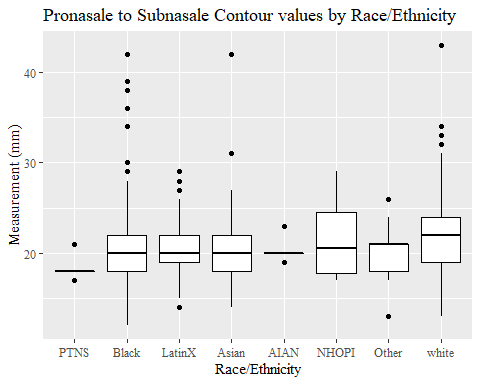
#%>% set\_header\_Cabels(values = list(ProS\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(ProS\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Pronasale to Subnasale Contour SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Pronasale to Subnasale Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 13 | 43 | 21.64 | 22.0 | 3.11 | 0.09 | 17.00 | 19.00 | 22.0 | 24.0 | 27.00 | 21 |
| Black | 548 | 12 | 42 | 20.20 | 20.0 | 3.64 | 0.16 | 15.00 | 18.00 | 20.0 | 22.0 | 25.00 | 7 |
| LatinX | 100 | 14 | 29 | 20.70 | 20.0 | 2.88 | 0.29 | 16.00 | 19.00 | 20.0 | 22.0 | 25.00 | 1 |
| Asian | 91 | 14 | 42 | 20.18 | 20.0 | 3.73 | 0.39 | 16.00 | 18.00 | 20.0 | 22.0 | 25.55 | 1 |
| Other | 21 | 13 | 26 | 19.62 | 21.0 | 3.12 | 0.68 | 13.00 | 18.00 | 21.0 | 21.0 | 24.00 | 0 |
| AIAN | 8 | 19 | 23 | 20.29 | 20.0 | 1.25 | 0.44 | 19.30 | 20.00 | 20.0 | 20.0 | 22.10 | 1 |
| PTNS | 5 | 17 | 21 | 18.40 | 18.0 | 1.52 | 0.68 | 17.20 | 18.00 | 18.0 | 18.0 | 20.40 | 0 |
| NHOPI | 4 | 17 | 29 | 21.75 | 20.5 | 5.50 | 2.75 | 17.15 | 17.75 | 20.5 | 24.5 | 28.10 | 0 |

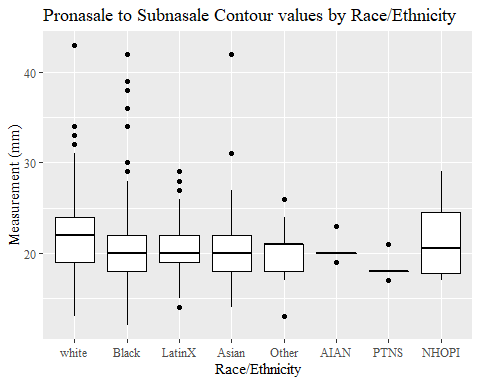
#%>% set\_header\_Cabels(values = list(ProS\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, ProS\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=ProS\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Pronasale to Subnasale Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 31 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=ProS\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Pronasale to Subnasale Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 31 rows containing non-finite values (stat\_boxplot).



#ProS\_C gender sumstats  
ProS\_Cgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(ProS\_C, na.rm = TRUE),  
 max = max(ProS\_C, na.rm = TRUE),  
 mean = mean(ProS\_C, na.rm = TRUE),  
 mdn = median(ProS\_C, na.rm = TRUE),  
 sd = sd(ProS\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(ProS\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(ProS\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(ProS\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(ProS\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(ProS\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(ProS\_C)))  
  
ProS\_Cgender\_sumstats <- ProS\_Cgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(ProS\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Pronasale to Subnasale Contour SumStats by Gender")

**Table** : Pronasale to Subnasale Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 13 | 42 | 20.46 | 20 | 3.16 | 0.10 | 16.00 | 18.00 | 20 | 22 | 25.00 | 12 |
| Male | 939 | 12 | 43 | 21.86 | 22 | 3.41 | 0.11 | 17.00 | 20.00 | 22 | 24 | 27.00 | 19 |
| Non-binary or Other | 5 | 17 | 20 | 19.00 | 19 | 1.22 | 0.55 | 17.40 | 19.00 | 19 | 20 | 20.00 | 0 |
| Prefer not to say | 1 | 18 | 18 | 18.00 | 18 |  |  | 18.00 | 18.00 | 18 | 18 | 18.00 | 0 |
|  | 8 | 15 | 24 | 19.50 | 19 | 3.46 | 1.22 | 15.35 | 16.75 | 19 | 23 | 23.65 | 0 |

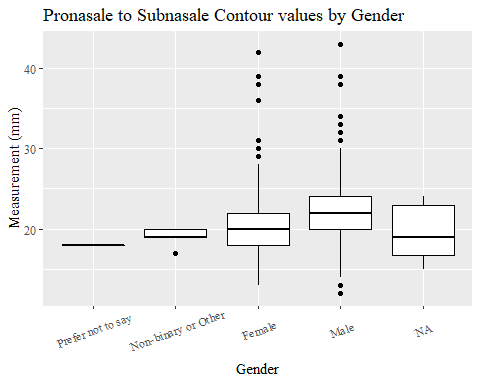
#%>% set\_header\_Cabels(values = list(ProS\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(ProS\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Pronasale to Subnasale Contour SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Pronasale to Subnasale Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 13 | 42 | 20.46 | 20 | 3.16 | 0.10 | 16.00 | 18.00 | 20 | 22 | 25.00 | 12 |
| Male | 939 | 12 | 43 | 21.86 | 22 | 3.41 | 0.11 | 17.00 | 20.00 | 22 | 24 | 27.00 | 19 |
| Non-binary or Other | 5 | 17 | 20 | 19.00 | 19 | 1.22 | 0.55 | 17.40 | 19.00 | 19 | 20 | 20.00 | 0 |
| Prefer not to say | 1 | 18 | 18 | 18.00 | 18 |  |  | 18.00 | 18.00 | 18 | 18 | 18.00 | 0 |
|  | 8 | 15 | 24 | 19.50 | 19 | 3.46 | 1.22 | 15.35 | 16.75 | 19 | 23 | 23.65 | 0 |

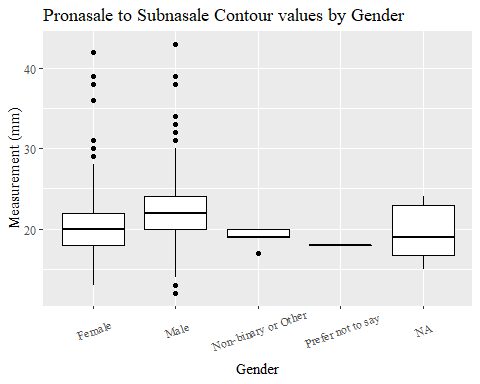
#%>% set\_header\_Cabels(values = list(ProS\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, ProS\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=ProS\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Pronasale to Subnasale Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 31 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=ProS\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Pronasale to Subnasale Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 31 rows containing non-finite values (stat\_boxplot).



#ProS\_C age group sumstats  
ProS\_Cage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(ProS\_C, na.rm = TRUE),  
 max = max(ProS\_C, na.rm = TRUE),  
 mean = mean(ProS\_C, na.rm = TRUE),  
 mdn = median(ProS\_C, na.rm = TRUE),  
 sd = sd(ProS\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(ProS\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(ProS\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(ProS\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(ProS\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(ProS\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(ProS\_C)))  
  
ProS\_Cage\_sumstats <- ProS\_Cage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(ProS\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Pronasale to Subnasale Contour SumStats by Age Group")

**Table** : Pronasale to Subnasale Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 13 | 43 | 20.57 | 20 | 3.12 | 0.10 | 16 | 18 | 20 | 22 | 25 | 12 |
| 37-54 | 940 | 12 | 42 | 21.60 | 21 | 3.52 | 0.11 | 16 | 19 | 21 | 23 | 27 | 16 |
| 55-72 | 84 | 15 | 32 | 21.80 | 22 | 2.93 | 0.32 | 17 | 20 | 22 | 24 | 27 | 3 |
|  | 1 | 25 | 25 | 25.00 | 25 |  |  | 25 | 25 | 25 | 25 | 25 | 0 |

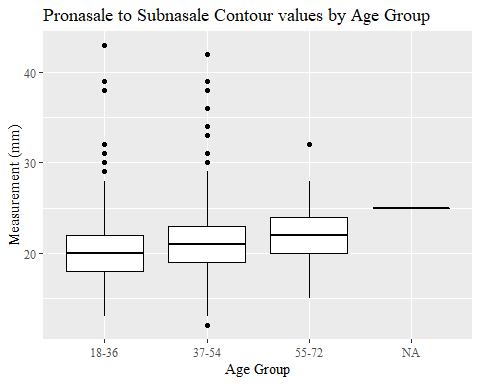
#%>% set\_header\_Cabels(values = list(ProS\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(ProS\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Pronasale to Subnasale Contour SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Pronasale to Subnasale Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 13 | 43 | 20.57 | 20 | 3.12 | 0.10 | 16 | 18 | 20 | 22 | 25 | 12 |
| 37-54 | 940 | 12 | 42 | 21.60 | 21 | 3.52 | 0.11 | 16 | 19 | 21 | 23 | 27 | 16 |
| 55-72 | 84 | 15 | 32 | 21.80 | 22 | 2.93 | 0.32 | 17 | 20 | 22 | 24 | 27 | 3 |
|  | 1 | 25 | 25 | 25.00 | 25 |  |  | 25 | 25 | 25 | 25 | 25 | 0 |

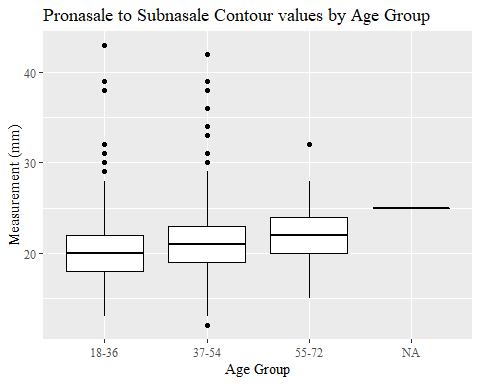
#%>% set\_header\_Cabels(values = list(ProS\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, ProS\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=ProS\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Pronasale to Subnasale Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 31 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=ProS\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Pronasale to Subnasale Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

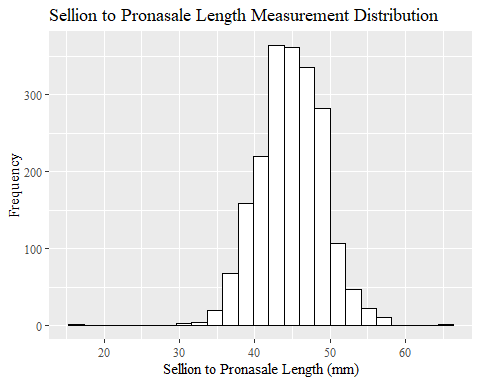
## Warning: Removed 31 rows containing non-finite values (stat\_boxplot).



SelP\_L

#histogram of all SelP\_L values  
ggplot(data=headscan\_full, aes(x=SelP\_L))+  
 geom\_bar(stat="bin", bins=25, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Sellion to Pronasale Length Measurement Distribution",  
 y="Frequency",  
 x="Sellion to Pronasale Length (mm)")

## Warning: Removed 15 rows containing non-finite values (stat\_bin).



#SelP\_L race/eth sumstats  
SelP\_Lrace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(SelP\_L, na.rm = TRUE),  
 max = max(SelP\_L, na.rm = TRUE),  
 mean = mean(SelP\_L, na.rm = TRUE),  
 mdn = median(SelP\_L, na.rm = TRUE),  
 sd = sd(SelP\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(SelP\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(SelP\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(SelP\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(SelP\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(SelP\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(SelP\_L)))   
  
SelP\_Lrace\_sumstats <- SelP\_Lrace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(SelP\_Lrace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Pronasale Length SumStats by Race/Ethnicity")

**Table** : Sellion to Pronasale Length SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 16 | 65 | 45.07 | 45 | 4.47 | 0.13 | 38.00 | 42.00 | 45 | 48.0 | 53.0 | 7 |
| Black | 548 | 30 | 57 | 43.26 | 43 | 4.19 | 0.18 | 36.10 | 40.50 | 43 | 46.0 | 50.0 | 5 |
| LatinX | 100 | 34 | 54 | 45.30 | 45 | 3.88 | 0.39 | 38.90 | 43.00 | 45 | 48.0 | 52.0 | 1 |
| Asian | 91 | 36 | 54 | 43.73 | 44 | 3.92 | 0.41 | 38.45 | 41.00 | 44 | 46.0 | 51.0 | 1 |
| Other | 21 | 37 | 55 | 45.57 | 46 | 3.85 | 0.84 | 41.00 | 44.00 | 46 | 47.0 | 51.0 | 0 |
| AIAN | 8 | 37 | 52 | 45.57 | 47 | 5.26 | 1.86 | 37.90 | 43.00 | 47 | 48.5 | 51.1 | 1 |
| PTNS | 5 | 40 | 45 | 42.40 | 42 | 1.82 | 0.81 | 40.40 | 42.00 | 42 | 43.0 | 44.6 | 0 |
| NHOPI | 4 | 39 | 48 | 43.75 | 44 | 3.69 | 1.84 | 39.75 | 42.75 | 44 | 45.0 | 47.4 | 0 |

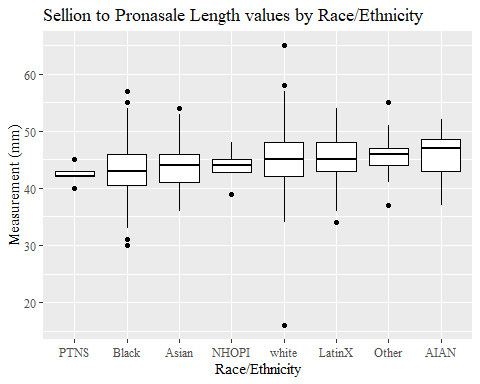
#%>% set\_header\_Cabels(values = list(SelP\_L = "Sellion/SellionCont"))  
  
#Autofit Width Table TNR  
flextable(SelP\_Lrace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Pronasale Length SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Sellion to Pronasale Length SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 16 | 65 | 45.07 | 45 | 4.47 | 0.13 | 38.00 | 42.00 | 45 | 48.0 | 53.0 | 7 |
| Black | 548 | 30 | 57 | 43.26 | 43 | 4.19 | 0.18 | 36.10 | 40.50 | 43 | 46.0 | 50.0 | 5 |
| LatinX | 100 | 34 | 54 | 45.30 | 45 | 3.88 | 0.39 | 38.90 | 43.00 | 45 | 48.0 | 52.0 | 1 |
| Asian | 91 | 36 | 54 | 43.73 | 44 | 3.92 | 0.41 | 38.45 | 41.00 | 44 | 46.0 | 51.0 | 1 |
| Other | 21 | 37 | 55 | 45.57 | 46 | 3.85 | 0.84 | 41.00 | 44.00 | 46 | 47.0 | 51.0 | 0 |
| AIAN | 8 | 37 | 52 | 45.57 | 47 | 5.26 | 1.86 | 37.90 | 43.00 | 47 | 48.5 | 51.1 | 1 |
| PTNS | 5 | 40 | 45 | 42.40 | 42 | 1.82 | 0.81 | 40.40 | 42.00 | 42 | 43.0 | 44.6 | 0 |
| NHOPI | 4 | 39 | 48 | 43.75 | 44 | 3.69 | 1.84 | 39.75 | 42.75 | 44 | 45.0 | 47.4 | 0 |

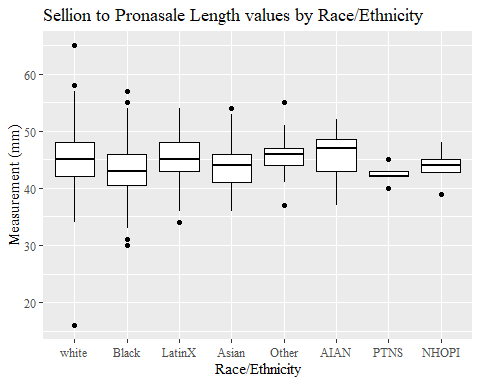
#%>% set\_header\_Cabels(values = list(SelP\_L = "Sellion/SellionCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, SelP\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=SelP\_L, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Pronasale Length values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 15 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=SelP\_L, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Pronasale Length values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 15 rows containing non-finite values (stat\_boxplot).



#SelP\_L gender sumstats  
SelP\_Lgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(SelP\_L, na.rm = TRUE),  
 max = max(SelP\_L, na.rm = TRUE),  
 mean = mean(SelP\_L, na.rm = TRUE),  
 mdn = median(SelP\_L, na.rm = TRUE),  
 sd = sd(SelP\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(SelP\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(SelP\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(SelP\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(SelP\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(SelP\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(SelP\_L)))  
  
SelP\_Lgender\_sumstats <- SelP\_Lgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(SelP\_Lgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Pronasale Length SumStats by Gender")

**Table** : Sellion to Pronasale Length SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 16 | 57 | 42.85 | 43 | 3.87 | 0.12 | 37.00 | 40.0 | 43 | 45.0 | 49.0 | 10 |
| Male | 939 | 31 | 65 | 46.48 | 46 | 4.15 | 0.14 | 40.00 | 44.0 | 46 | 49.0 | 53.0 | 5 |
| Non-binary or Other | 5 | 36 | 52 | 40.60 | 38 | 6.47 | 2.89 | 36.40 | 38.0 | 38 | 39.0 | 49.4 | 0 |
| Prefer not to say | 1 | 40 | 40 | 40.00 | 40 |  |  | 40.00 | 40.0 | 40 | 40.0 | 40.0 | 0 |
|  | 8 | 37 | 51 | 41.25 | 40 | 4.33 | 1.53 | 37.35 | 39.5 | 40 | 41.5 | 48.2 | 0 |

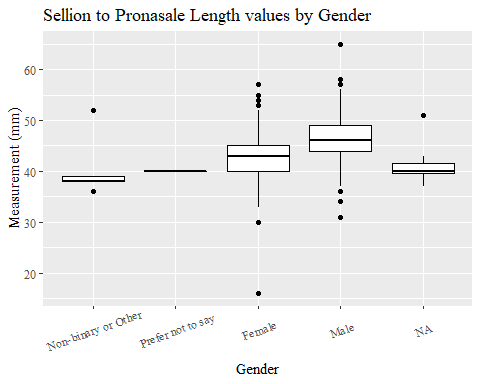
#%>% set\_header\_Cabels(values = list(SelP\_L = "Sellion/SellionCont"))  
  
#Autofit Width Table TNR  
flextable(SelP\_Lgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Pronasale Length SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Sellion to Pronasale Length SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 16 | 57 | 42.85 | 43 | 3.87 | 0.12 | 37.00 | 40.0 | 43 | 45.0 | 49.0 | 10 |
| Male | 939 | 31 | 65 | 46.48 | 46 | 4.15 | 0.14 | 40.00 | 44.0 | 46 | 49.0 | 53.0 | 5 |
| Non-binary or Other | 5 | 36 | 52 | 40.60 | 38 | 6.47 | 2.89 | 36.40 | 38.0 | 38 | 39.0 | 49.4 | 0 |
| Prefer not to say | 1 | 40 | 40 | 40.00 | 40 |  |  | 40.00 | 40.0 | 40 | 40.0 | 40.0 | 0 |
|  | 8 | 37 | 51 | 41.25 | 40 | 4.33 | 1.53 | 37.35 | 39.5 | 40 | 41.5 | 48.2 | 0 |

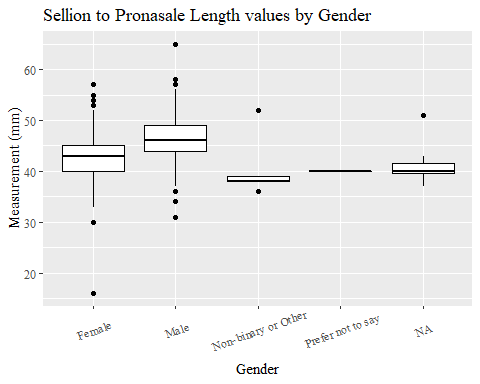
#%>% set\_header\_Cabels(values = list(SelP\_L = "Sellion/SellionCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, SelP\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=SelP\_L, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Pronasale Length values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 15 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=SelP\_L, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Pronasale Length values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 15 rows containing non-finite values (stat\_boxplot).



#SelP\_L age group sumstats  
SelP\_Lage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(SelP\_L, na.rm = TRUE),  
 max = max(SelP\_L, na.rm = TRUE),  
 mean = mean(SelP\_L, na.rm = TRUE),  
 mdn = median(SelP\_L, na.rm = TRUE),  
 sd = sd(SelP\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(SelP\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(SelP\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(SelP\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(SelP\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(SelP\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(SelP\_L)))  
  
SelP\_Lage\_sumstats <- SelP\_Lage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(SelP\_Lage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Pronasale Length SumStats by Age Group")

**Table** : Sellion to Pronasale Length SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 31 | 57 | 44.35 | 44 | 4.21 | 0.13 | 38 | 42 | 44 | 47 | 51 | 5 |
| 37-54 | 940 | 16 | 65 | 44.65 | 45 | 4.60 | 0.15 | 37 | 42 | 45 | 47 | 53 | 9 |
| 55-72 | 84 | 37 | 58 | 45.31 | 45 | 4.37 | 0.48 | 40 | 42 | 45 | 47 | 53 | 1 |
|  | 1 | 43 | 43 | 43.00 | 43 |  |  | 43 | 43 | 43 | 43 | 43 | 0 |

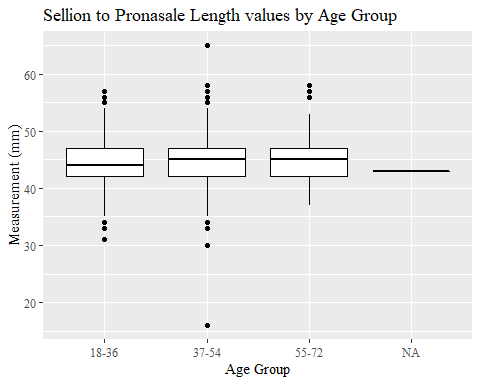
#%>% set\_header\_Cabels(values = list(SelP\_L = "Sellion/SellionCont"))  
  
#Autofit Width Table TNR  
flextable(SelP\_Lage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Pronasale Length SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Sellion to Pronasale Length SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 31 | 57 | 44.35 | 44 | 4.21 | 0.13 | 38 | 42 | 44 | 47 | 51 | 5 |
| 37-54 | 940 | 16 | 65 | 44.65 | 45 | 4.60 | 0.15 | 37 | 42 | 45 | 47 | 53 | 9 |
| 55-72 | 84 | 37 | 58 | 45.31 | 45 | 4.37 | 0.48 | 40 | 42 | 45 | 47 | 53 | 1 |
|  | 1 | 43 | 43 | 43.00 | 43 |  |  | 43 | 43 | 43 | 43 | 43 | 0 |

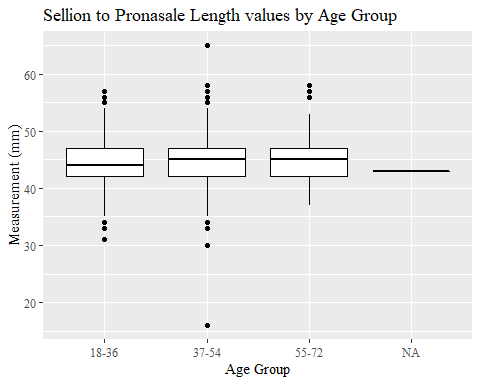
#%>% set\_header\_Cabels(values = list(SelP\_L = "Sellion/SellionCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, SelP\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=SelP\_L, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Pronasale Length values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 15 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=SelP\_L, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Pronasale Length values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

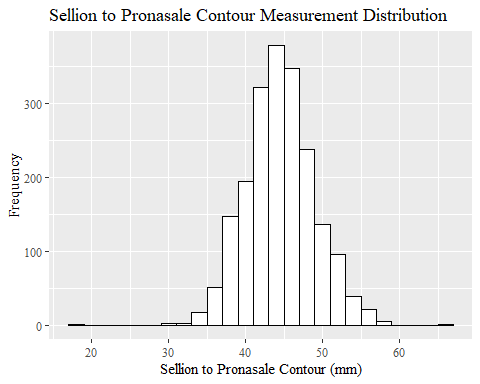
## Warning: Removed 15 rows containing non-finite values (stat\_boxplot).



SelP\_C

#histogram of all SelP\_C values  
ggplot(data=headscan\_full, aes(x=SelP\_C ))+  
 geom\_bar(stat="bin", bins=25, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Sellion to Pronasale Contour Measurement Distribution",  
 y="Frequency",  
 x="Sellion to Pronasale Contour (mm)")

## Warning: Removed 15 rows containing non-finite values (stat\_bin).



#SelP\_C race/eth sumstats  
SelP\_Crace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(SelP\_C, na.rm = TRUE),  
 max = max(SelP\_C, na.rm = TRUE),  
 mean = mean(SelP\_C, na.rm = TRUE),  
 mdn = median(SelP\_C, na.rm = TRUE),  
 sd = sd(SelP\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(SelP\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(SelP\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(SelP\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(SelP\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(SelP\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(SelP\_C)))   
  
SelP\_Crace\_sumstats <- SelP\_Crace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(SelP\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Pronasale Contour SumStats by Race/Ethnicity")

**Table** : Sellion to Pronasale Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 18 | 66 | 45.52 | 45 | 4.56 | 0.13 | 38.0 | 43.0 | 45 | 48.00 | 53.0 | 7 |
| Black | 548 | 30 | 58 | 43.78 | 44 | 4.28 | 0.18 | 37.0 | 41.0 | 44 | 47.00 | 51.0 | 5 |
| LatinX | 100 | 34 | 55 | 45.83 | 46 | 3.96 | 0.40 | 39.0 | 43.0 | 46 | 49.00 | 52.1 | 1 |
| Asian | 91 | 36 | 55 | 44.21 | 44 | 4.09 | 0.43 | 39.0 | 41.0 | 44 | 46.75 | 51.0 | 1 |
| Other | 21 | 38 | 55 | 46.19 | 46 | 3.94 | 0.86 | 41.0 | 44.0 | 46 | 48.00 | 52.0 | 0 |
| AIAN | 8 | 37 | 52 | 46.14 | 48 | 5.52 | 1.95 | 37.9 | 43.5 | 48 | 49.50 | 51.4 | 1 |
| PTNS | 5 | 41 | 46 | 43.00 | 43 | 1.87 | 0.84 | 41.2 | 42.0 | 43 | 43.00 | 45.4 | 0 |
| NHOPI | 4 | 40 | 48 | 44.00 | 44 | 3.27 | 1.63 | 40.6 | 43.0 | 44 | 45.00 | 47.4 | 0 |

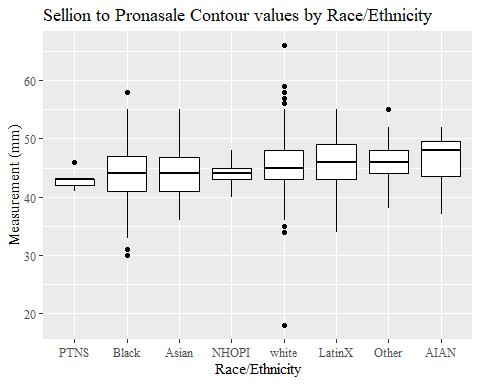
#%>% set\_header\_Cabels(values = list(SelP\_C = "Sellion/SellionCont"))  
  
#Autofit Width Table TNR  
flextable(SelP\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Pronasale Contour SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Sellion to Pronasale Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 18 | 66 | 45.52 | 45 | 4.56 | 0.13 | 38.0 | 43.0 | 45 | 48.00 | 53.0 | 7 |
| Black | 548 | 30 | 58 | 43.78 | 44 | 4.28 | 0.18 | 37.0 | 41.0 | 44 | 47.00 | 51.0 | 5 |
| LatinX | 100 | 34 | 55 | 45.83 | 46 | 3.96 | 0.40 | 39.0 | 43.0 | 46 | 49.00 | 52.1 | 1 |
| Asian | 91 | 36 | 55 | 44.21 | 44 | 4.09 | 0.43 | 39.0 | 41.0 | 44 | 46.75 | 51.0 | 1 |
| Other | 21 | 38 | 55 | 46.19 | 46 | 3.94 | 0.86 | 41.0 | 44.0 | 46 | 48.00 | 52.0 | 0 |
| AIAN | 8 | 37 | 52 | 46.14 | 48 | 5.52 | 1.95 | 37.9 | 43.5 | 48 | 49.50 | 51.4 | 1 |
| PTNS | 5 | 41 | 46 | 43.00 | 43 | 1.87 | 0.84 | 41.2 | 42.0 | 43 | 43.00 | 45.4 | 0 |
| NHOPI | 4 | 40 | 48 | 44.00 | 44 | 3.27 | 1.63 | 40.6 | 43.0 | 44 | 45.00 | 47.4 | 0 |

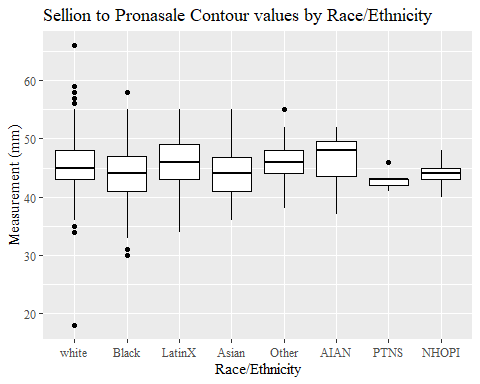
#%>% set\_header\_Cabels(values = list(SelP\_C = "Sellion/SellionCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, SelP\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=SelP\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Pronasale Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 15 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=SelP\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Pronasale Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 15 rows containing non-finite values (stat\_boxplot).



#SelP\_C gender sumstats  
SelP\_Cgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(SelP\_C, na.rm = TRUE),  
 max = max(SelP\_C, na.rm = TRUE),  
 mean = mean(SelP\_C, na.rm = TRUE),  
 mdn = median(SelP\_C, na.rm = TRUE),  
 sd = sd(SelP\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(SelP\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(SelP\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(SelP\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(SelP\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(SelP\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(SelP\_C)))  
  
SelP\_Cgender\_sumstats <- SelP\_Cgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(SelP\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Pronasale Contour SumStats by Gender")

**Table** : Sellion to Pronasale Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 18 | 58 | 43.26 | 43 | 3.93 | 0.12 | 37.0 | 41.0 | 43 | 46.00 | 49.0 | 10 |
| Male | 939 | 31 | 66 | 47.03 | 47 | 4.22 | 0.14 | 40.0 | 44.0 | 47 | 50.00 | 54.0 | 5 |
| Non-binary or Other | 5 | 36 | 53 | 41.20 | 39 | 6.72 | 3.01 | 36.6 | 39.0 | 39 | 39.00 | 50.2 | 0 |
| Prefer not to say | 1 | 41 | 41 | 41.00 | 41 |  |  | 41.0 | 41.0 | 41 | 41.00 | 41.0 | 0 |
|  | 8 | 38 | 51 | 41.50 | 40 | 4.21 | 1.49 | 38.0 | 39.5 | 40 | 42.25 | 48.2 | 0 |

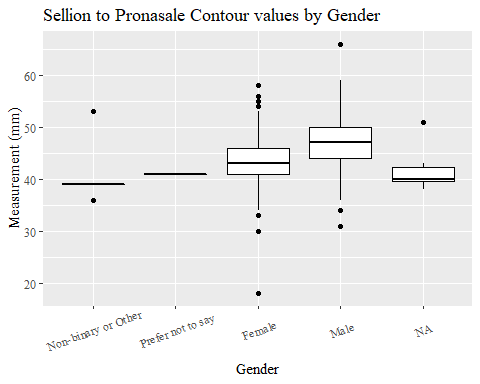
#%>% set\_header\_Cabels(values = list(SelP\_C = "Sellion/SellionCont"))  
  
#Autofit Width Table TNR  
flextable(SelP\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Pronasale Contour SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Sellion to Pronasale Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 18 | 58 | 43.26 | 43 | 3.93 | 0.12 | 37.0 | 41.0 | 43 | 46.00 | 49.0 | 10 |
| Male | 939 | 31 | 66 | 47.03 | 47 | 4.22 | 0.14 | 40.0 | 44.0 | 47 | 50.00 | 54.0 | 5 |
| Non-binary or Other | 5 | 36 | 53 | 41.20 | 39 | 6.72 | 3.01 | 36.6 | 39.0 | 39 | 39.00 | 50.2 | 0 |
| Prefer not to say | 1 | 41 | 41 | 41.00 | 41 |  |  | 41.0 | 41.0 | 41 | 41.00 | 41.0 | 0 |
|  | 8 | 38 | 51 | 41.50 | 40 | 4.21 | 1.49 | 38.0 | 39.5 | 40 | 42.25 | 48.2 | 0 |

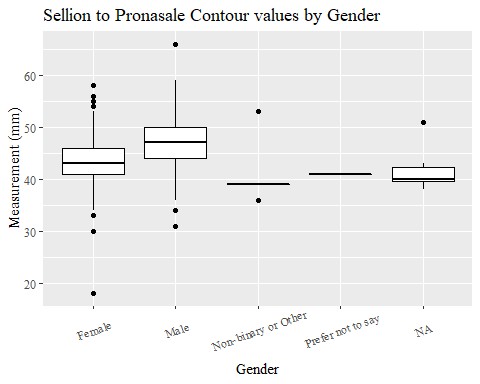
#%>% set\_header\_Cabels(values = list(SelP\_C = "Sellion/SellionCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, SelP\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=SelP\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Pronasale Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 15 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=SelP\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Pronasale Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 15 rows containing non-finite values (stat\_boxplot).



#SelP\_C age group sumstats  
SelP\_Cage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(SelP\_C, na.rm = TRUE),  
 max = max(SelP\_C, na.rm = TRUE),  
 mean = mean(SelP\_C, na.rm = TRUE),  
 mdn = median(SelP\_C, na.rm = TRUE),  
 sd = sd(SelP\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(SelP\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(SelP\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(SelP\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(SelP\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(SelP\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(SelP\_C)))  
  
SelP\_Cage\_sumstats <- SelP\_Cage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(SelP\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Pronasale Contour SumStats by Age Group")

**Table** : Sellion to Pronasale Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 31 | 58 | 44.81 | 45 | 4.31 | 0.14 | 38.0 | 42 | 45 | 48 | 52.0 | 5 |
| 37-54 | 940 | 18 | 66 | 45.14 | 45 | 4.68 | 0.15 | 38.0 | 42 | 45 | 48 | 53.0 | 9 |
| 55-72 | 84 | 38 | 58 | 45.89 | 45 | 4.40 | 0.48 | 40.1 | 43 | 45 | 48 | 53.9 | 1 |
|  | 1 | 43 | 43 | 43.00 | 43 |  |  | 43.0 | 43 | 43 | 43 | 43.0 | 0 |

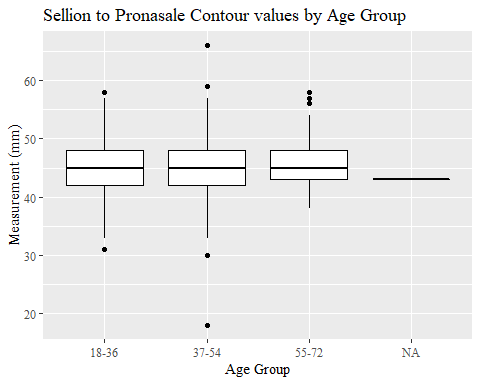
#%>% set\_header\_Cabels(values = list(SelP\_C = "Sellion/SellionCont"))  
  
#Autofit Width Table TNR  
flextable(SelP\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Pronasale Contour SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Sellion to Pronasale Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 31 | 58 | 44.81 | 45 | 4.31 | 0.14 | 38.0 | 42 | 45 | 48 | 52.0 | 5 |
| 37-54 | 940 | 18 | 66 | 45.14 | 45 | 4.68 | 0.15 | 38.0 | 42 | 45 | 48 | 53.0 | 9 |
| 55-72 | 84 | 38 | 58 | 45.89 | 45 | 4.40 | 0.48 | 40.1 | 43 | 45 | 48 | 53.9 | 1 |
|  | 1 | 43 | 43 | 43.00 | 43 |  |  | 43.0 | 43 | 43 | 43 | 43.0 | 0 |

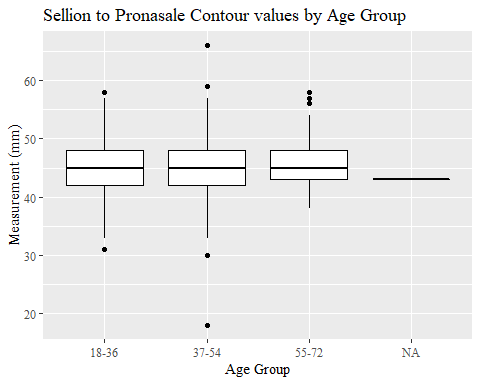
#%>% set\_header\_Cabels(values = list(SelP\_C = "Sellion/SellionCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, SelP\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=SelP\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Pronasale Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 15 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=SelP\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Pronasale Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

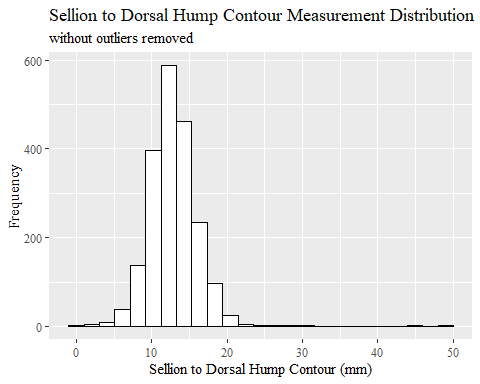
## Warning: Removed 15 rows containing non-finite values (stat\_boxplot).



SelDH\_C with outliers

#histogram of all SelDH\_C values  
ggplot(data=headscan\_full, aes(x=SelDH\_C ))+  
 geom\_bar(stat="bin", bins=25, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Sellion to Dorsal Hump Contour Measurement Distribution",  
 subtitle = "without outliers removed",  
 y="Frequency",  
 x="Sellion to Dorsal Hump Contour (mm)")

## Warning: Removed 14 rows containing non-finite values (stat\_bin).



#SelDH\_C race/eth sumstats  
SelDH\_Crace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(SelDH\_C, na.rm = TRUE),  
 max = max(SelDH\_C, na.rm = TRUE),  
 mean = mean(SelDH\_C, na.rm = TRUE),  
 mdn = median(SelDH\_C, na.rm = TRUE),  
 sd = sd(SelDH\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(SelDH\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(SelDH\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(SelDH\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(SelDH\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(SelDH\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(SelDH\_C)))   
  
SelDH\_Crace\_sumstats <- SelDH\_Crace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(SelDH\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Dorsal Hump Contour SumStats by Race/Ethnicity")

**Table** : Sellion to Dorsal Hump Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 1 | 50 | 13.32 | 13.0 | 3.14 | 0.09 | 9.00 | 11.0 | 13.0 | 15.0 | 18.0 | 7 |
| Black | 548 | 3 | 24 | 12.46 | 12.0 | 2.82 | 0.12 | 8.00 | 11.0 | 12.0 | 14.0 | 17.0 | 4 |
| LatinX | 100 | 7 | 45 | 13.75 | 13.0 | 4.20 | 0.42 | 8.90 | 12.0 | 13.0 | 15.0 | 18.0 | 1 |
| Asian | 91 | 8 | 20 | 13.03 | 13.0 | 2.62 | 0.27 | 9.45 | 11.0 | 13.0 | 14.0 | 18.0 | 1 |
| Other | 21 | 8 | 28 | 12.86 | 12.0 | 4.30 | 0.94 | 9.00 | 10.0 | 12.0 | 15.0 | 16.0 | 0 |
| AIAN | 8 | 2 | 15 | 11.43 | 14.0 | 4.76 | 1.68 | 3.80 | 10.5 | 14.0 | 14.0 | 14.7 | 1 |
| PTNS | 5 | 4 | 15 | 10.80 | 12.0 | 4.09 | 1.83 | 5.40 | 11.0 | 12.0 | 12.0 | 14.4 | 0 |
| NHOPI | 4 | 12 | 17 | 14.00 | 13.5 | 2.45 | 1.22 | 12.00 | 12.0 | 13.5 | 15.5 | 16.7 | 0 |

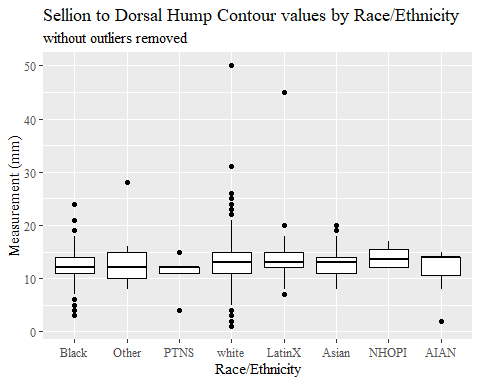
#%>% set\_header\_Cabels(values = list(SelDH\_C = "Sellion/SellionCont"))  
  
#Autofit Width Table TNR  
flextable(SelDH\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Dorsal Hump Contour SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Sellion to Dorsal Hump Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 1 | 50 | 13.32 | 13.0 | 3.14 | 0.09 | 9.00 | 11.0 | 13.0 | 15.0 | 18.0 | 7 |
| Black | 548 | 3 | 24 | 12.46 | 12.0 | 2.82 | 0.12 | 8.00 | 11.0 | 12.0 | 14.0 | 17.0 | 4 |
| LatinX | 100 | 7 | 45 | 13.75 | 13.0 | 4.20 | 0.42 | 8.90 | 12.0 | 13.0 | 15.0 | 18.0 | 1 |
| Asian | 91 | 8 | 20 | 13.03 | 13.0 | 2.62 | 0.27 | 9.45 | 11.0 | 13.0 | 14.0 | 18.0 | 1 |
| Other | 21 | 8 | 28 | 12.86 | 12.0 | 4.30 | 0.94 | 9.00 | 10.0 | 12.0 | 15.0 | 16.0 | 0 |
| AIAN | 8 | 2 | 15 | 11.43 | 14.0 | 4.76 | 1.68 | 3.80 | 10.5 | 14.0 | 14.0 | 14.7 | 1 |
| PTNS | 5 | 4 | 15 | 10.80 | 12.0 | 4.09 | 1.83 | 5.40 | 11.0 | 12.0 | 12.0 | 14.4 | 0 |
| NHOPI | 4 | 12 | 17 | 14.00 | 13.5 | 2.45 | 1.22 | 12.00 | 12.0 | 13.5 | 15.5 | 16.7 | 0 |

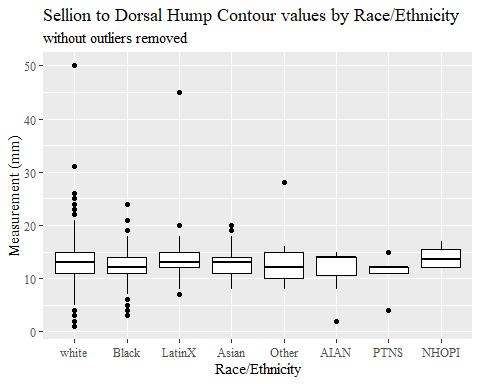
#%>% set\_header\_Cabels(values = list(SelDH\_C = "Sellion/SellionCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, SelDH\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=SelDH\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Dorsal Hump Contour values by Race/Ethnicity",  
 subtitle = "without outliers removed",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 14 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=SelDH\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Dorsal Hump Contour values by Race/Ethnicity",  
 subtitle = "without outliers removed",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 14 rows containing non-finite values (stat\_boxplot).



#SelDH\_C gender sumstats  
SelDH\_Cgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(SelDH\_C, na.rm = TRUE),  
 max = max(SelDH\_C, na.rm = TRUE),  
 mean = mean(SelDH\_C, na.rm = TRUE),  
 mdn = median(SelDH\_C, na.rm = TRUE),  
 sd = sd(SelDH\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(SelDH\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(SelDH\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(SelDH\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(SelDH\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(SelDH\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(SelDH\_C)))  
  
SelDH\_Cgender\_sumstats <- SelDH\_Cgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(SelDH\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Dorsal Hump Contour SumStats by Gender")

**Table** : Sellion to Dorsal Hump Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 1 | 50 | 12.59 | 13.0 | 3.14 | 0.10 | 8.0 | 11 | 13.0 | 14.00 | 17 | 9 |
| Male | 939 | 4 | 31 | 13.69 | 14.0 | 3.02 | 0.10 | 9.0 | 12 | 14.0 | 15.00 | 18 | 5 |
| Non-binary or Other | 5 | 8 | 14 | 9.80 | 9.0 | 2.39 | 1.07 | 8.2 | 9 | 9.0 | 9.00 | 13 | 0 |
| Prefer not to say | 1 | 12 | 12 | 12.00 | 12.0 |  |  | 12.0 | 12 | 12.0 | 12.00 | 12 | 0 |
|  | 8 | 6 | 14 | 9.75 | 8.5 | 2.96 | 1.05 | 6.7 | 8 | 8.5 | 11.75 | 14 | 0 |

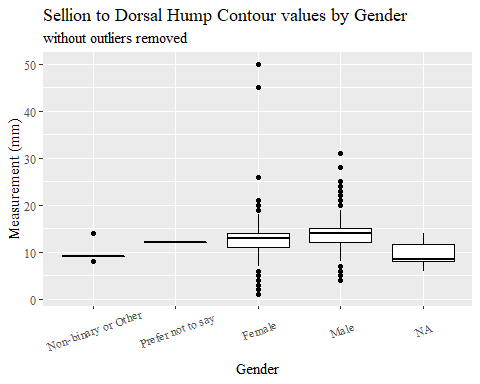
#%>% set\_header\_Cabels(values = list(SelDH\_C = "Sellion/SellionCont"))  
  
#Autofit Width Table TNR  
flextable(SelDH\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Dorsal Hump Contour SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Sellion to Dorsal Hump Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 1 | 50 | 12.59 | 13.0 | 3.14 | 0.10 | 8.0 | 11 | 13.0 | 14.00 | 17 | 9 |
| Male | 939 | 4 | 31 | 13.69 | 14.0 | 3.02 | 0.10 | 9.0 | 12 | 14.0 | 15.00 | 18 | 5 |
| Non-binary or Other | 5 | 8 | 14 | 9.80 | 9.0 | 2.39 | 1.07 | 8.2 | 9 | 9.0 | 9.00 | 13 | 0 |
| Prefer not to say | 1 | 12 | 12 | 12.00 | 12.0 |  |  | 12.0 | 12 | 12.0 | 12.00 | 12 | 0 |
|  | 8 | 6 | 14 | 9.75 | 8.5 | 2.96 | 1.05 | 6.7 | 8 | 8.5 | 11.75 | 14 | 0 |

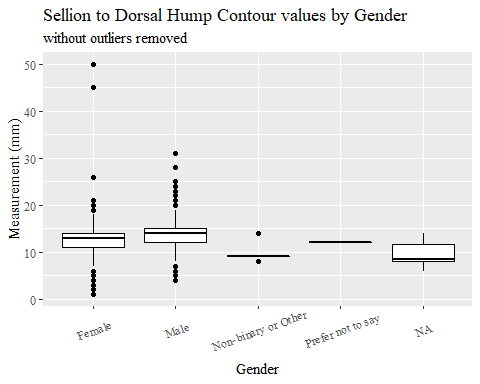
#%>% set\_header\_Cabels(values = list(SelDH\_C = "Sellion/SellionCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, SelDH\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=SelDH\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Dorsal Hump Contour values by Gender",  
 subtitle = "without outliers removed",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 14 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=SelDH\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Dorsal Hump Contour values by Gender",  
 subtitle = "without outliers removed",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 14 rows containing non-finite values (stat\_boxplot).



#SelDH\_C age group sumstats  
SelDH\_Cage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(SelDH\_C, na.rm = TRUE),  
 max = max(SelDH\_C, na.rm = TRUE),  
 mean = mean(SelDH\_C, na.rm = TRUE),  
 mdn = median(SelDH\_C, na.rm = TRUE),  
 sd = sd(SelDH\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(SelDH\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(SelDH\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(SelDH\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(SelDH\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(SelDH\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(SelDH\_C)))  
  
SelDH\_Cage\_sumstats <- SelDH\_Cage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(SelDH\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Dorsal Hump Contour SumStats by Age Group")

**Table** : Sellion to Dorsal Hump Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 2 | 28 | 13.06 | 13 | 2.81 | 0.09 | 9.0 | 11 | 13 | 15 | 18 | 5 |
| 37-54 | 940 | 1 | 50 | 13.10 | 13 | 3.47 | 0.11 | 8.0 | 11 | 13 | 15 | 18 | 8 |
| 55-72 | 84 | 6 | 21 | 13.04 | 13 | 3.16 | 0.34 | 7.1 | 11 | 13 | 15 | 18 | 1 |
|  | 1 | 13 | 13 | 13.00 | 13 |  |  | 13.0 | 13 | 13 | 13 | 13 | 0 |

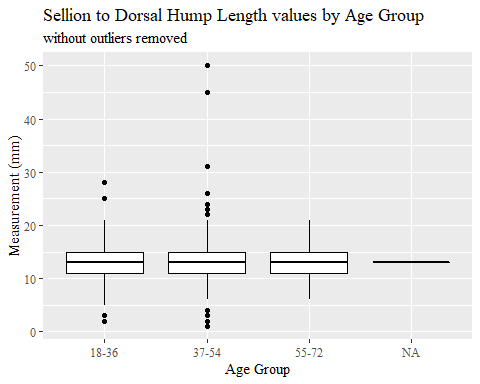
#%>% set\_header\_Cabels(values = list(SelDH\_C = "Sellion/SellionCont"))  
  
#Autofit Width Table TNR  
flextable(SelDH\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Dorsal Hump Contour SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Sellion to Dorsal Hump Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 2 | 28 | 13.06 | 13 | 2.81 | 0.09 | 9.0 | 11 | 13 | 15 | 18 | 5 |
| 37-54 | 940 | 1 | 50 | 13.10 | 13 | 3.47 | 0.11 | 8.0 | 11 | 13 | 15 | 18 | 8 |
| 55-72 | 84 | 6 | 21 | 13.04 | 13 | 3.16 | 0.34 | 7.1 | 11 | 13 | 15 | 18 | 1 |
|  | 1 | 13 | 13 | 13.00 | 13 |  |  | 13.0 | 13 | 13 | 13 | 13 | 0 |

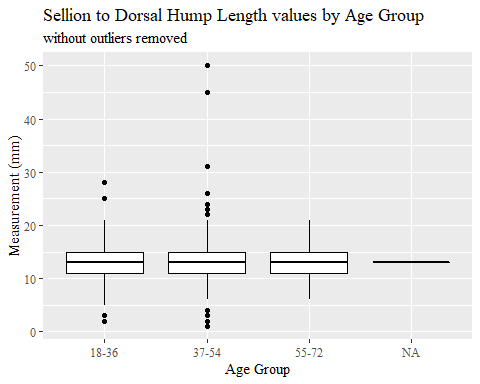
#%>% set\_header\_Cabels(values = list(SelDH\_C = "Sellion/SellionCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, SelDH\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=SelDH\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Dorsal Hump Length values by Age Group",  
 subtitle = "without outliers removed",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 14 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=SelDH\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Dorsal Hump Length values by Age Group",  
 subtitle = "without outliers removed",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 14 rows containing non-finite values (stat\_boxplot).



SelDH\_C without outliers

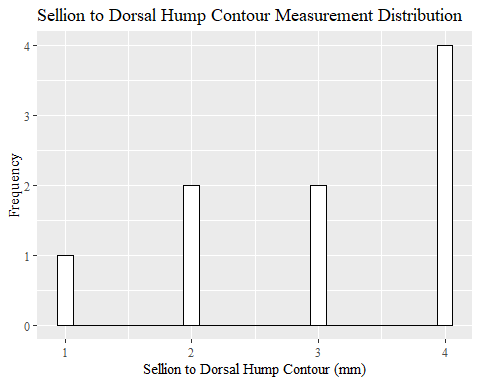
max(headscan\_full$SelDH\_C, na.rm = TRUE)

## [1] 50

#CODE TO REMOVE OUTLIER, use later to remove from headscan\_full  
SelDH\_C\_no\_out <- headscan\_full %>% mutate(SelDH\_C = replace(SelDH\_C, SelDH\_C>4, NA))

#histogram of all SelDH\_C values  
ggplot(data=SelDH\_C\_no\_out, aes(x=SelDH\_C))+  
 geom\_bar(stat="bin", bins=25, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Sellion to Dorsal Hump Contour Measurement Distribution",  
 y="Frequency",  
 x="Sellion to Dorsal Hump Contour (mm)")

## Warning: Removed 2008 rows containing non-finite values (stat\_bin).



#SelDH\_C race/eth sumstats  
SelDH\_Crace\_sumstats1 <- SelDH\_C\_no\_out %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(SelDH\_C, na.rm = TRUE),  
 max = max(SelDH\_C, na.rm = TRUE),  
 mean = mean(SelDH\_C, na.rm = TRUE),  
 mdn = median(SelDH\_C, na.rm = TRUE),  
 sd = sd(SelDH\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(SelDH\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(SelDH\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(SelDH\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(SelDH\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(SelDH\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(SelDH\_C)))

## Warning in min(SelDH\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(SelDH\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(SelDH\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(SelDH\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf

## Warning in max(SelDH\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(SelDH\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(SelDH\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(SelDH\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf

SelDH\_Crace\_sumstats1 <- SelDH\_Crace\_sumstats1 %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(SelDH\_Crace\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Dorsal Hump Contour SumStats by Race/Ethnicity")

**Table** : Sellion to Dorsal Hump Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 1 | 4 | 2.8 | 3.0 | 1.30 | 0.04 | 1.20 | 2.00 | 3.0 | 4.00 | 4.00 | 1,235 |
| Black | 548 | 3 | 4 | 3.5 | 3.5 | 0.71 | 0.03 | 3.05 | 3.25 | 3.5 | 3.75 | 3.95 | 546 |
| LatinX | 100 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 100 |
| Asian | 91 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 91 |
| Other | 21 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 21 |
| AIAN | 8 | 2 | 2 | 2.0 | 2.0 |  |  | 2.00 | 2.00 | 2.0 | 2.00 | 2.00 | 7 |
| PTNS | 5 | 4 | 4 | 4.0 | 4.0 |  |  | 4.00 | 4.00 | 4.0 | 4.00 | 4.00 | 4 |
| NHOPI | 4 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 4 |

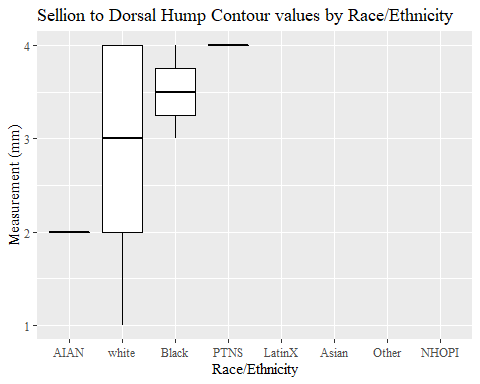
#%>% set\_header\_Cabels(values = list(SelDH\_C = "Sellion/SellionCont"))  
  
#Autofit Width Table TNR  
flextable(SelDH\_Crace\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Dorsal Hump Contour SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Sellion to Dorsal Hump Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 1 | 4 | 2.8 | 3.0 | 1.30 | 0.04 | 1.20 | 2.00 | 3.0 | 4.00 | 4.00 | 1,235 |
| Black | 548 | 3 | 4 | 3.5 | 3.5 | 0.71 | 0.03 | 3.05 | 3.25 | 3.5 | 3.75 | 3.95 | 546 |
| LatinX | 100 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 100 |
| Asian | 91 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 91 |
| Other | 21 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 21 |
| AIAN | 8 | 2 | 2 | 2.0 | 2.0 |  |  | 2.00 | 2.00 | 2.0 | 2.00 | 2.00 | 7 |
| PTNS | 5 | 4 | 4 | 4.0 | 4.0 |  |  | 4.00 | 4.00 | 4.0 | 4.00 | 4.00 | 4 |
| NHOPI | 4 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 4 |

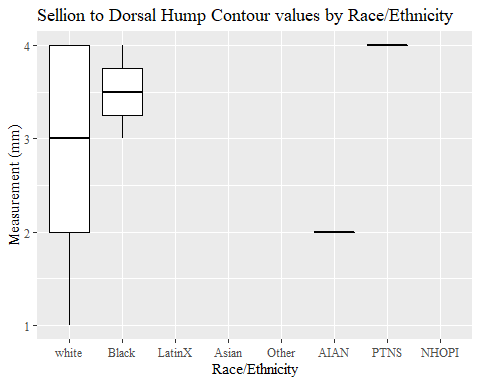
#%>% set\_header\_Cabels(values = list(SelDH\_C = "Sellion/SellionCont"))  
  
#boxplot reorderd by median  
SelDH\_C\_no\_out %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, SelDH\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=SelDH\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Dorsal Hump Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 2008 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=SelDH\_C\_no\_out, aes(y=SelDH\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Dorsal Hump Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 2008 rows containing non-finite values (stat\_boxplot).



#SelDH\_C gender sumstats  
SelDH\_Cgender\_sumstats1 <- SelDH\_C\_no\_out %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(SelDH\_C, na.rm = TRUE),  
 max = max(SelDH\_C, na.rm = TRUE),  
 mean = mean(SelDH\_C, na.rm = TRUE),  
 mdn = median(SelDH\_C, na.rm = TRUE),  
 sd = sd(SelDH\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(SelDH\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(SelDH\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(SelDH\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(SelDH\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(SelDH\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(SelDH\_C)))

## Warning in min(SelDH\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(SelDH\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(SelDH\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf

## Warning in max(SelDH\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(SelDH\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(SelDH\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf

SelDH\_Cgender\_sumstats1 <- SelDH\_Cgender\_sumstats1 %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(SelDH\_Cgender\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Dorsal Hump Contour SumStats by Gender")

**Table** : Sellion to Dorsal Hump Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 1 | 4 | 2.71 | 3 | 1.11 | 0.03 | 1.3 | 2 | 3 | 3.5 | 4 | 1,057 |
| Male | 939 | 4 | 4 | 4.00 | 4 | 0.00 | 0.00 | 4.0 | 4 | 4 | 4.0 | 4 | 937 |
| Non-binary or Other | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| Prefer not to say | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |
|  | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |

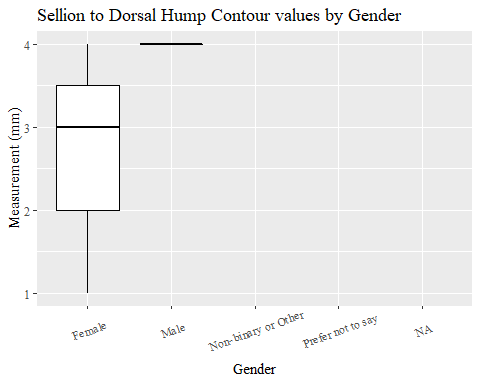
#%>% set\_header\_Cabels(values = list(SelDH\_C = "Sellion/SellionCont"))  
  
#Autofit Width Table TNR  
flextable(SelDH\_Cgender\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Dorsal Hump Contour SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Sellion to Dorsal Hump Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 1 | 4 | 2.71 | 3 | 1.11 | 0.03 | 1.3 | 2 | 3 | 3.5 | 4 | 1,057 |
| Male | 939 | 4 | 4 | 4.00 | 4 | 0.00 | 0.00 | 4.0 | 4 | 4 | 4.0 | 4 | 937 |
| Non-binary or Other | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| Prefer not to say | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |
|  | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |

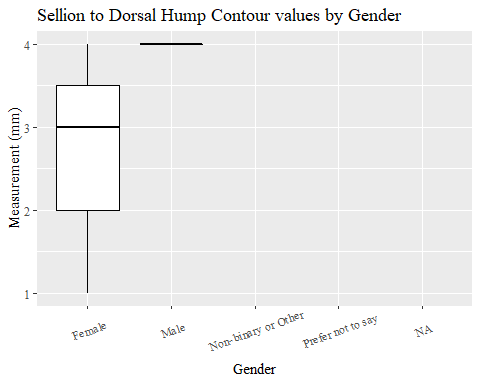
#%>% set\_header\_Cabels(values = list(SelDH\_C = "Sellion/SellionCont"))  
  
#boxplot reorderd by median  
SelDH\_C\_no\_out %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, SelDH\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=SelDH\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Dorsal Hump Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 2008 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=SelDH\_C\_no\_out, aes(y=SelDH\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Dorsal Hump Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 2008 rows containing non-finite values (stat\_boxplot).



#SelDH\_C age group sumstats  
SelDH\_Cage\_sumstats1 <- SelDH\_C\_no\_out %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(SelDH\_C, na.rm = TRUE),  
 max = max(SelDH\_C, na.rm = TRUE),  
 mean = mean(SelDH\_C, na.rm = TRUE),  
 mdn = median(SelDH\_C, na.rm = TRUE),  
 sd = sd(SelDH\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(SelDH\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(SelDH\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(SelDH\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(SelDH\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(SelDH\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(SelDH\_C)))

## Warning in min(SelDH\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(SelDH\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf

## Warning in max(SelDH\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(SelDH\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf

SelDH\_Cage\_sumstats1 <- SelDH\_Cage\_sumstats1 %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(SelDH\_Cage\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Dorsal Hump Contour SumStats by Age Group")

**Table** : Sellion to Dorsal Hump Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 2 | 3 | 2.50 | 2.5 | 0.71 | 0.02 | 2.05 | 2.25 | 2.5 | 2.75 | 2.95 | 990 |
| 37-54 | 940 | 1 | 4 | 3.14 | 4.0 | 1.21 | 0.04 | 1.30 | 2.50 | 4.0 | 4.00 | 4.00 | 933 |
| 55-72 | 84 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 84 |
|  | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |

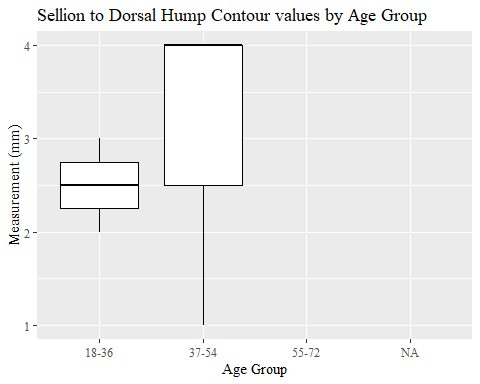
#%>% set\_header\_Cabels(values = list(SelDH\_C = "Sellion/SellionCont"))  
  
#Autofit Width Table TNR  
flextable(SelDH\_Cage\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Dorsal Hump Contour SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Sellion to Dorsal Hump Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 2 | 3 | 2.50 | 2.5 | 0.71 | 0.02 | 2.05 | 2.25 | 2.5 | 2.75 | 2.95 | 990 |
| 37-54 | 940 | 1 | 4 | 3.14 | 4.0 | 1.21 | 0.04 | 1.30 | 2.50 | 4.0 | 4.00 | 4.00 | 933 |
| 55-72 | 84 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 84 |
|  | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |

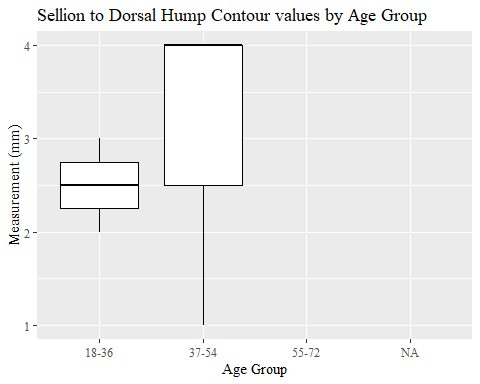
#%>% set\_header\_Cabels(values = list(SelDH\_C = "Sellion/SellionCont"))  
  
#boxplot reorderd by median  
SelDH\_C\_no\_out %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, SelDH\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=SelDH\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Dorsal Hump Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 2008 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=SelDH\_C\_no\_out, aes(y=SelDH\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Dorsal Hump Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

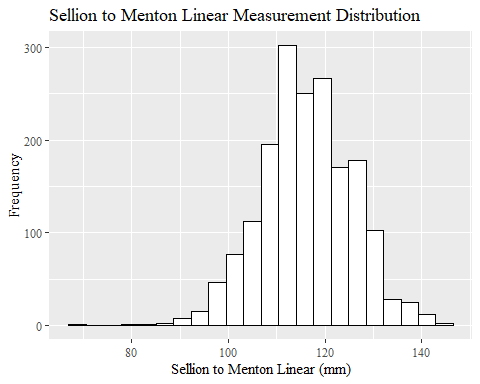
## Warning: Removed 2008 rows containing non-finite values (stat\_boxplot).



SelM\_L

#histogram of all SelM\_L values  
ggplot(data=headscan\_full, aes(x=SelM\_L))+  
 geom\_bar(stat="bin", bins=22, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Sellion to Menton Linear Measurement Distribution",  
 y="Frequency",  
 x="Sellion to Menton Linear (mm)")

## Warning: Removed 224 rows containing non-finite values (stat\_bin).



#SelM\_L race/eth sumstats  
SelM\_Lrace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(SelM\_L, na.rm = TRUE),  
 max = max(SelM\_L, na.rm = TRUE),  
 mean = mean(SelM\_L, na.rm = TRUE),  
 mdn = median(SelM\_L, na.rm = TRUE),  
 sd = sd(SelM\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(SelM\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(SelM\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(SelM\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(SelM\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(SelM\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(SelM\_L)))   
  
SelM\_Lrace\_sumstats <- SelM\_Lrace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(SelM\_Lrace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Menton Length SumStats by Race/Ethnicity")

**Table** : Sellion to Menton Length SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 69 | 145 | 115.68 | 115.0 | 9.33 | 0.26 | 100.00 | 110.00 | 115.0 | 122.00 | 131.00 | 135 |
| Black | 548 | 91 | 142 | 118.51 | 118.0 | 9.00 | 0.38 | 102.00 | 112.00 | 118.0 | 125.00 | 132.00 | 70 |
| LatinX | 100 | 84 | 142 | 115.85 | 116.0 | 10.72 | 1.07 | 98.00 | 109.00 | 116.0 | 122.00 | 134.90 | 8 |
| Asian | 91 | 88 | 132 | 111.86 | 111.0 | 8.66 | 0.91 | 97.00 | 107.50 | 111.0 | 117.50 | 126.00 | 4 |
| Other | 21 | 105 | 126 | 114.44 | 113.0 | 5.84 | 1.27 | 107.25 | 111.00 | 113.0 | 116.75 | 123.75 | 5 |
| AIAN | 8 | 99 | 137 | 115.50 | 111.5 | 13.44 | 4.75 | 101.50 | 109.25 | 111.5 | 122.00 | 134.00 | 2 |
| PTNS | 5 | 102 | 126 | 115.80 | 117.0 | 8.90 | 3.98 | 104.40 | 114.00 | 117.0 | 120.00 | 124.80 | 0 |
| NHOPI | 4 | 98 | 122 | 115.00 | 120.0 | 11.37 | 5.69 | 101.30 | 114.50 | 120.0 | 120.50 | 121.70 | 0 |

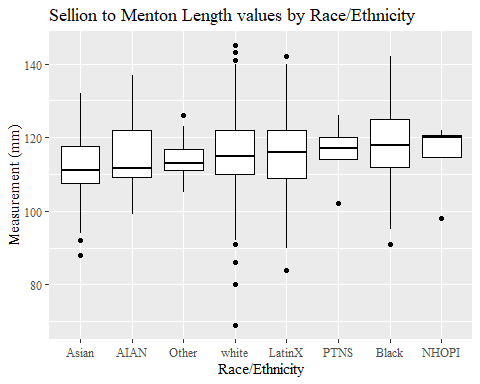
#%>% set\_header\_Cabels(values = list(SelM\_L = "Sellion/SellionCont"))  
  
#Autofit Width Table TNR  
flextable(SelM\_Lrace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Menton Length SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Sellion to Menton Length SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 69 | 145 | 115.68 | 115.0 | 9.33 | 0.26 | 100.00 | 110.00 | 115.0 | 122.00 | 131.00 | 135 |
| Black | 548 | 91 | 142 | 118.51 | 118.0 | 9.00 | 0.38 | 102.00 | 112.00 | 118.0 | 125.00 | 132.00 | 70 |
| LatinX | 100 | 84 | 142 | 115.85 | 116.0 | 10.72 | 1.07 | 98.00 | 109.00 | 116.0 | 122.00 | 134.90 | 8 |
| Asian | 91 | 88 | 132 | 111.86 | 111.0 | 8.66 | 0.91 | 97.00 | 107.50 | 111.0 | 117.50 | 126.00 | 4 |
| Other | 21 | 105 | 126 | 114.44 | 113.0 | 5.84 | 1.27 | 107.25 | 111.00 | 113.0 | 116.75 | 123.75 | 5 |
| AIAN | 8 | 99 | 137 | 115.50 | 111.5 | 13.44 | 4.75 | 101.50 | 109.25 | 111.5 | 122.00 | 134.00 | 2 |
| PTNS | 5 | 102 | 126 | 115.80 | 117.0 | 8.90 | 3.98 | 104.40 | 114.00 | 117.0 | 120.00 | 124.80 | 0 |
| NHOPI | 4 | 98 | 122 | 115.00 | 120.0 | 11.37 | 5.69 | 101.30 | 114.50 | 120.0 | 120.50 | 121.70 | 0 |

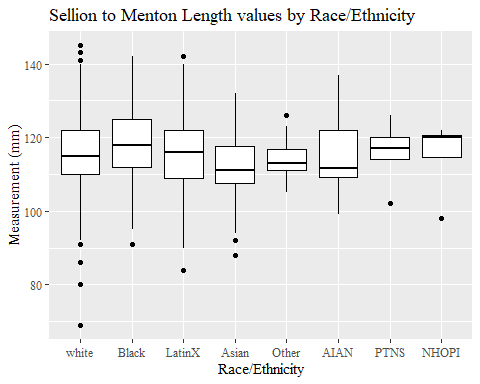
#%>% set\_header\_Cabels(values = list(SelM\_L = "Sellion/SellionCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, SelM\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=SelM\_L, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Menton Length values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 224 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=SelM\_L, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Menton Length values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 224 rows containing non-finite values (stat\_boxplot).



#SelM\_L gender sumstats  
SelM\_Lgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(SelM\_L, na.rm = TRUE),  
 max = max(SelM\_L, na.rm = TRUE),  
 mean = mean(SelM\_L, na.rm = TRUE),  
 mdn = median(SelM\_L, na.rm = TRUE),  
 sd = sd(SelM\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(SelM\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(SelM\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(SelM\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(SelM\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(SelM\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(SelM\_L)))  
  
SelM\_Lgender\_sumstats <- SelM\_Lgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(SelM\_Lgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Menton Length SumStats by Gender")

**Table** : Sellion to Menton Length SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 69 | 140 | 113.10 | 114 | 8.28 | 0.25 | 99.00 | 108.00 | 114 | 118.0 | 126.00 | 17 |
| Male | 939 | 93 | 145 | 120.81 | 121 | 8.98 | 0.29 | 106.00 | 115.00 | 121 | 127.0 | 135.45 | 207 |
| Non-binary or Other | 5 | 80 | 116 | 106.40 | 114 | 15.18 | 6.79 | 85.40 | 107.00 | 114 | 115.0 | 115.80 | 0 |
| Prefer not to say | 1 | 126 | 126 | 126.00 | 126 |  |  | 126.00 | 126.00 | 126 | 126.0 | 126.00 | 0 |
|  | 8 | 103 | 128 | 114.50 | 112 | 8.72 | 3.08 | 104.75 | 108.75 | 112 | 121.5 | 126.25 | 0 |

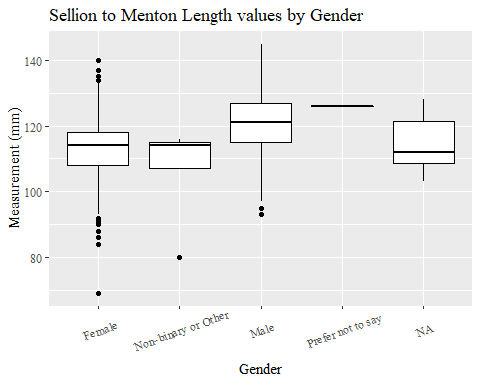
#%>% set\_header\_Cabels(values = list(SelM\_L = "Sellion/SellionCont"))  
  
#Autofit Width Table TNR  
flextable(SelM\_Lgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Menton Length SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Sellion to Menton Length SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 69 | 140 | 113.10 | 114 | 8.28 | 0.25 | 99.00 | 108.00 | 114 | 118.0 | 126.00 | 17 |
| Male | 939 | 93 | 145 | 120.81 | 121 | 8.98 | 0.29 | 106.00 | 115.00 | 121 | 127.0 | 135.45 | 207 |
| Non-binary or Other | 5 | 80 | 116 | 106.40 | 114 | 15.18 | 6.79 | 85.40 | 107.00 | 114 | 115.0 | 115.80 | 0 |
| Prefer not to say | 1 | 126 | 126 | 126.00 | 126 |  |  | 126.00 | 126.00 | 126 | 126.0 | 126.00 | 0 |
|  | 8 | 103 | 128 | 114.50 | 112 | 8.72 | 3.08 | 104.75 | 108.75 | 112 | 121.5 | 126.25 | 0 |

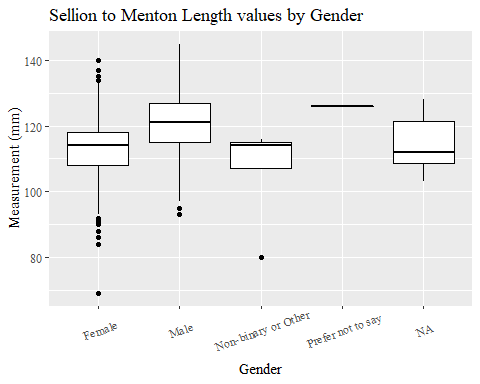
#%>% set\_header\_Cabels(values = list(SelM\_L = "Sellion/SellionCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, SelM\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=SelM\_L, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Menton Length values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 224 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=SelM\_L, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Menton Length values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 224 rows containing non-finite values (stat\_boxplot).



#SelM\_L age group sumstats  
SelM\_Lage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(SelM\_L, na.rm = TRUE),  
 max = max(SelM\_L, na.rm = TRUE),  
 mean = mean(SelM\_L, na.rm = TRUE),  
 mdn = median(SelM\_L, na.rm = TRUE),  
 sd = sd(SelM\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(SelM\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(SelM\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(SelM\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(SelM\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(SelM\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(SelM\_L)))  
  
SelM\_Lage\_sumstats <- SelM\_Lage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(SelM\_Lage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Menton Length SumStats by Age Group")

**Table** : Sellion to Menton Length SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 80 | 143 | 115.74 | 116 | 9.07 | 0.29 | 101.0 | 110 | 116 | 122.0 | 131.0 | 120 |
| 37-54 | 940 | 69 | 142 | 116.80 | 117 | 9.64 | 0.31 | 101.0 | 111 | 117 | 123.0 | 132.0 | 99 |
| 55-72 | 84 | 86 | 145 | 115.89 | 117 | 10.14 | 1.11 | 97.9 | 111 | 117 | 123.5 | 128.2 | 5 |
|  | 1 | 113 | 113 | 113.00 | 113 |  |  | 113.0 | 113 | 113 | 113.0 | 113.0 | 0 |

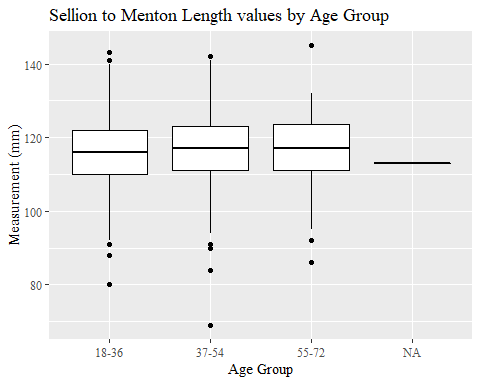
#%>% set\_header\_Cabels(values = list(SelM\_L = "Sellion/SellionCont"))  
  
#Autofit Width Table TNR  
flextable(SelM\_Lage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Sellion to Menton Length SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Sellion to Menton Length SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 80 | 143 | 115.74 | 116 | 9.07 | 0.29 | 101.0 | 110 | 116 | 122.0 | 131.0 | 120 |
| 37-54 | 940 | 69 | 142 | 116.80 | 117 | 9.64 | 0.31 | 101.0 | 111 | 117 | 123.0 | 132.0 | 99 |
| 55-72 | 84 | 86 | 145 | 115.89 | 117 | 10.14 | 1.11 | 97.9 | 111 | 117 | 123.5 | 128.2 | 5 |
|  | 1 | 113 | 113 | 113.00 | 113 |  |  | 113.0 | 113 | 113 | 113.0 | 113.0 | 0 |

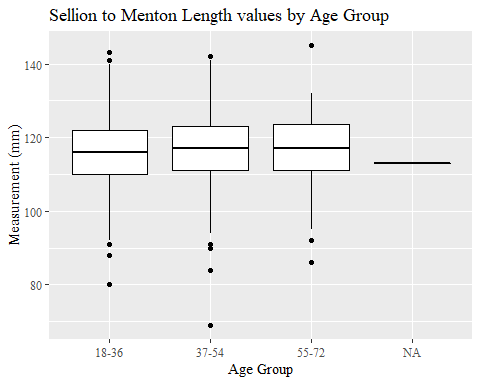
#%>% set\_header\_Cabels(values = list(SelM\_L = "Sellion/SellionCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, SelM\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=SelM\_L, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Menton Length values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 224 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=SelM\_L, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Sellion to Menton Length values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

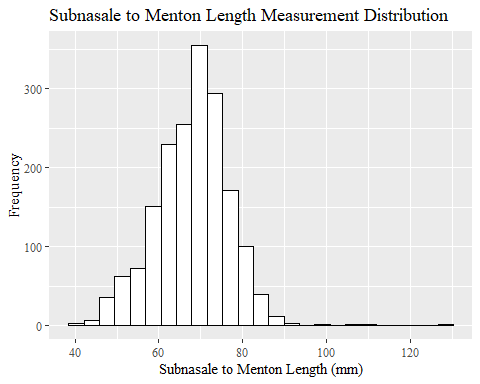
## Warning: Removed 224 rows containing non-finite values (stat\_boxplot).



SnasM\_L

#histogram of all SnasM\_L values  
ggplot(data=headscan\_full, aes(x=SnasM\_L))+  
 geom\_bar(stat="bin", bins=25, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Subnasale to Menton Length Measurement Distribution",  
 y="Frequency",  
 x="Subnasale to Menton Length (mm)")

## Warning: Removed 225 rows containing non-finite values (stat\_bin).



#SnasM\_L race/eth sumstats  
SnasM\_Lrace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(SnasM\_L, na.rm = TRUE),  
 max = max(SnasM\_L, na.rm = TRUE),  
 mean = mean(SnasM\_L, na.rm = TRUE),  
 mdn = median(SnasM\_L, na.rm = TRUE),  
 sd = sd(SnasM\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(SnasM\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(SnasM\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(SnasM\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(SnasM\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(SnasM\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(SnasM\_L)))   
  
SnasM\_Lrace\_sumstats <- SnasM\_Lrace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(SnasM\_Lrace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Subnasale to Menton Length SumStats by Race/Ethnicity")

**Table** : Subnasale to Menton Length SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 41 | 128 | 66.96 | 67.0 | 8.29 | 0.24 | 52.00 | 62.00 | 67.0 | 72.00 | 79.00 | 136 |
| Black | 548 | 46 | 111 | 71.03 | 72.0 | 8.32 | 0.36 | 56.00 | 66.00 | 72.0 | 77.00 | 83.00 | 70 |
| LatinX | 100 | 43 | 90 | 67.30 | 68.0 | 9.65 | 0.96 | 49.55 | 62.00 | 68.0 | 73.25 | 82.45 | 8 |
| Asian | 91 | 40 | 105 | 64.26 | 64.0 | 9.64 | 1.01 | 50.00 | 58.50 | 64.0 | 71.00 | 78.40 | 4 |
| Other | 21 | 52 | 78 | 66.56 | 66.5 | 6.55 | 1.43 | 56.50 | 64.75 | 66.5 | 71.25 | 75.75 | 5 |
| AIAN | 8 | 54 | 83 | 65.00 | 61.5 | 10.77 | 3.81 | 55.00 | 58.25 | 61.5 | 70.00 | 80.25 | 2 |
| PTNS | 5 | 59 | 81 | 70.60 | 72.0 | 8.44 | 3.78 | 60.40 | 66.00 | 72.0 | 75.00 | 79.80 | 0 |
| NHOPI | 4 | 46 | 78 | 65.75 | 69.5 | 14.06 | 7.03 | 49.00 | 61.00 | 69.5 | 74.25 | 77.25 | 0 |

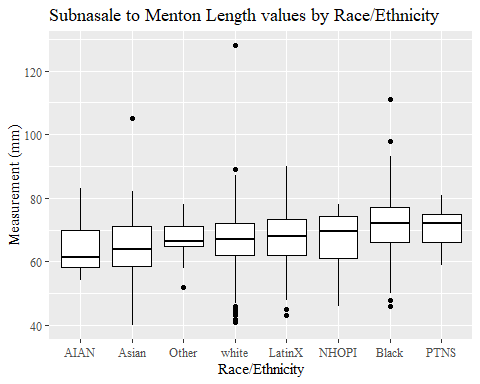
#%>% set\_header\_Cabels(values = list(SnasM\_L = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(SnasM\_Lrace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Subnasale to Menton Length SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Subnasale to Menton Length SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 41 | 128 | 66.96 | 67.0 | 8.29 | 0.24 | 52.00 | 62.00 | 67.0 | 72.00 | 79.00 | 136 |
| Black | 548 | 46 | 111 | 71.03 | 72.0 | 8.32 | 0.36 | 56.00 | 66.00 | 72.0 | 77.00 | 83.00 | 70 |
| LatinX | 100 | 43 | 90 | 67.30 | 68.0 | 9.65 | 0.96 | 49.55 | 62.00 | 68.0 | 73.25 | 82.45 | 8 |
| Asian | 91 | 40 | 105 | 64.26 | 64.0 | 9.64 | 1.01 | 50.00 | 58.50 | 64.0 | 71.00 | 78.40 | 4 |
| Other | 21 | 52 | 78 | 66.56 | 66.5 | 6.55 | 1.43 | 56.50 | 64.75 | 66.5 | 71.25 | 75.75 | 5 |
| AIAN | 8 | 54 | 83 | 65.00 | 61.5 | 10.77 | 3.81 | 55.00 | 58.25 | 61.5 | 70.00 | 80.25 | 2 |
| PTNS | 5 | 59 | 81 | 70.60 | 72.0 | 8.44 | 3.78 | 60.40 | 66.00 | 72.0 | 75.00 | 79.80 | 0 |
| NHOPI | 4 | 46 | 78 | 65.75 | 69.5 | 14.06 | 7.03 | 49.00 | 61.00 | 69.5 | 74.25 | 77.25 | 0 |

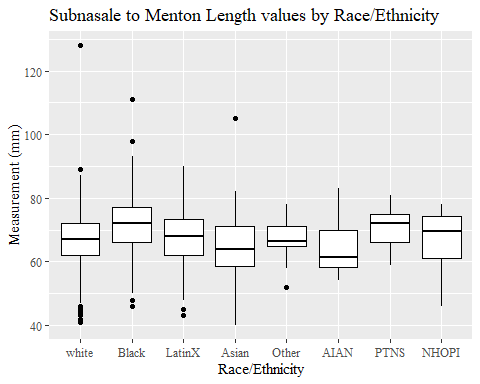
#%>% set\_header\_Cabels(values = list(SnasM\_L = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, SnasM\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=SnasM\_L, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Subnasale to Menton Length values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 225 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=SnasM\_L, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Subnasale to Menton Length values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 225 rows containing non-finite values (stat\_boxplot).



#SnasM\_L gender sumstats  
SnasM\_Lgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(SnasM\_L, na.rm = TRUE),  
 max = max(SnasM\_L, na.rm = TRUE),  
 mean = mean(SnasM\_L, na.rm = TRUE),  
 mdn = median(SnasM\_L, na.rm = TRUE),  
 sd = sd(SnasM\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(SnasM\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(SnasM\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(SnasM\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(SnasM\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(SnasM\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(SnasM\_L)))  
  
SnasM\_Lgender\_sumstats <- SnasM\_Lgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(SnasM\_Lgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Subnasale to Menton Length SumStats by Gender")

**Table** : Subnasale to Menton Length SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 40 | 111 | 66.24 | 67 | 8.08 | 0.25 | 51.00 | 62.0 | 67 | 72.00 | 78.00 | 17 |
| Male | 939 | 42 | 128 | 70.38 | 71 | 8.84 | 0.29 | 55.00 | 65.0 | 71 | 76.00 | 83.00 | 208 |
| Non-binary or Other | 5 | 41 | 74 | 62.20 | 70 | 14.25 | 6.37 | 43.60 | 54.0 | 70 | 72.00 | 73.60 | 0 |
| Prefer not to say | 1 | 81 | 81 | 81.00 | 81 |  |  | 81.00 | 81.0 | 81 | 81.00 | 81.00 | 0 |
|  | 8 | 57 | 78 | 68.00 | 67 | 8.33 | 2.95 | 58.05 | 61.5 | 67 | 76.25 | 77.65 | 0 |

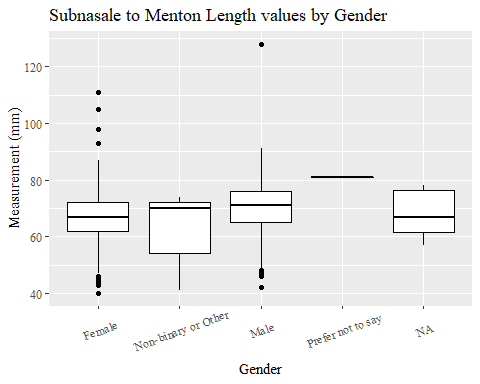
#%>% set\_header\_Cabels(values = list(SnasM\_L = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(SnasM\_Lgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Subnasale to Menton Length SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Subnasale to Menton Length SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 40 | 111 | 66.24 | 67 | 8.08 | 0.25 | 51.00 | 62.0 | 67 | 72.00 | 78.00 | 17 |
| Male | 939 | 42 | 128 | 70.38 | 71 | 8.84 | 0.29 | 55.00 | 65.0 | 71 | 76.00 | 83.00 | 208 |
| Non-binary or Other | 5 | 41 | 74 | 62.20 | 70 | 14.25 | 6.37 | 43.60 | 54.0 | 70 | 72.00 | 73.60 | 0 |
| Prefer not to say | 1 | 81 | 81 | 81.00 | 81 |  |  | 81.00 | 81.0 | 81 | 81.00 | 81.00 | 0 |
|  | 8 | 57 | 78 | 68.00 | 67 | 8.33 | 2.95 | 58.05 | 61.5 | 67 | 76.25 | 77.65 | 0 |

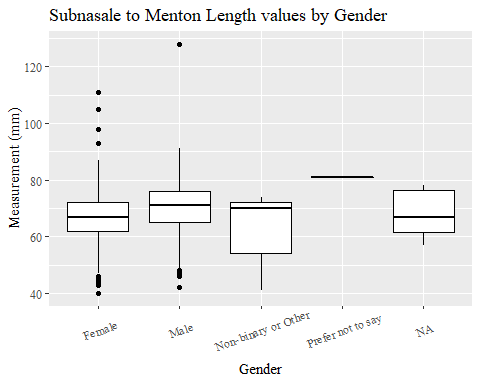
#%>% set\_header\_Cabels(values = list(SnasM\_L = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, SnasM\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=SnasM\_L, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Subnasale to Menton Length values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 225 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=SnasM\_L, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Subnasale to Menton Length values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 225 rows containing non-finite values (stat\_boxplot).



#SnasM\_L age group sumstats  
SnasM\_Lage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(SnasM\_L, na.rm = TRUE),  
 max = max(SnasM\_L, na.rm = TRUE),  
 mean = mean(SnasM\_L, na.rm = TRUE),  
 mdn = median(SnasM\_L, na.rm = TRUE),  
 sd = sd(SnasM\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(SnasM\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(SnasM\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(SnasM\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(SnasM\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(SnasM\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(SnasM\_L)))  
  
SnasM\_Lage\_sumstats <- SnasM\_Lage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(SnasM\_Lage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Subnasale to Menton Length SumStats by Age Group")

**Table** : Subnasale to Menton Length SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 40 | 128 | 67.85 | 68 | 8.68 | 0.28 | 52.0 | 63.00 | 68 | 74 | 81.00 | 120 |
| 37-54 | 940 | 43 | 111 | 68.18 | 68 | 8.65 | 0.28 | 53.0 | 63.00 | 68 | 74 | 81.00 | 99 |
| 55-72 | 84 | 43 | 82 | 66.21 | 67 | 8.56 | 0.93 | 51.7 | 61.25 | 67 | 73 | 77.15 | 6 |
|  | 1 | 64 | 64 | 64.00 | 64 |  |  | 64.0 | 64.00 | 64 | 64 | 64.00 | 0 |

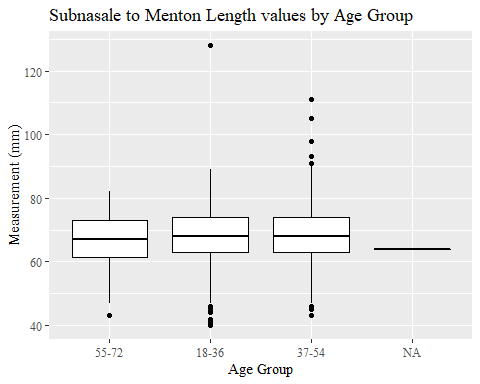
#%>% set\_header\_Cabels(values = list(SnasM\_L = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(SnasM\_Lage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Subnasale to Menton Length SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Subnasale to Menton Length SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 40 | 128 | 67.85 | 68 | 8.68 | 0.28 | 52.0 | 63.00 | 68 | 74 | 81.00 | 120 |
| 37-54 | 940 | 43 | 111 | 68.18 | 68 | 8.65 | 0.28 | 53.0 | 63.00 | 68 | 74 | 81.00 | 99 |
| 55-72 | 84 | 43 | 82 | 66.21 | 67 | 8.56 | 0.93 | 51.7 | 61.25 | 67 | 73 | 77.15 | 6 |
|  | 1 | 64 | 64 | 64.00 | 64 |  |  | 64.0 | 64.00 | 64 | 64 | 64.00 | 0 |

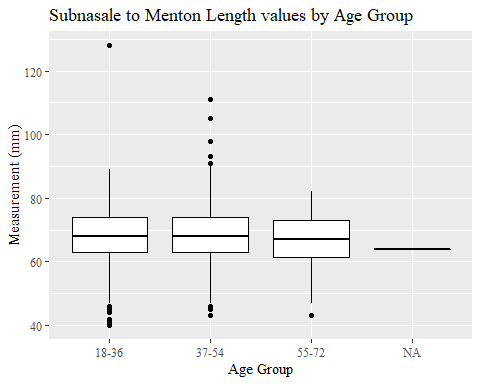
#%>% set\_header\_Cabels(values = list(SnasM\_L = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, SnasM\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=SnasM\_L, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Subnasale to Menton Length values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 225 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=SnasM\_L, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Subnasale to Menton Length values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

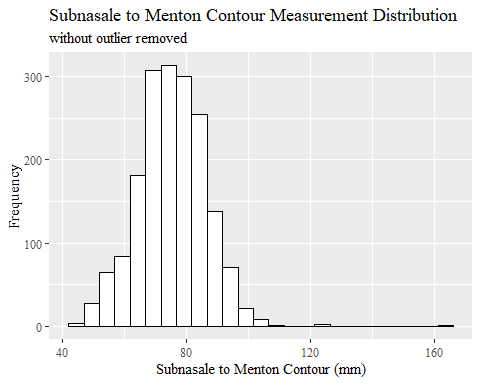
## Warning: Removed 225 rows containing non-finite values (stat\_boxplot).



SnasM\_C without outliers removed

#histogram of all SnasM\_C values  
ggplot(data=headscan\_full, aes(x=SnasM\_C))+  
 geom\_bar(stat="bin", bins=25, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Subnasale to Menton Contour Measurement Distribution",  
 subtitle = "without outlier removed",  
 y="Frequency",  
 x="Subnasale to Menton Contour (mm)")

## Warning: Removed 236 rows containing non-finite values (stat\_bin).



#SnasM\_C race/eth sumstats  
SnasM\_Crace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(SnasM\_C, na.rm = TRUE),  
 max = max(SnasM\_C, na.rm = TRUE),  
 mean = mean(SnasM\_C, na.rm = TRUE),  
 mdn = median(SnasM\_C, na.rm = TRUE),  
 sd = sd(SnasM\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(SnasM\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(SnasM\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(SnasM\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(SnasM\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(SnasM\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(SnasM\_C)))   
  
SnasM\_Crace\_sumstats <- SnasM\_Crace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(SnasM\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Subnasale to Menton Contour SumStats by Race/Ethnicity")

**Table** : Subnasale to Menton Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 44 | 163 | 72.89 | 73.0 | 10.09 | 0.29 | 56.00 | 67.00 | 73.0 | 79.00 | 89.00 | 144 |
| Black | 548 | 54 | 124 | 81.09 | 82.0 | 10.03 | 0.43 | 63.00 | 75.00 | 82.0 | 87.00 | 96.20 | 71 |
| LatinX | 100 | 50 | 97 | 73.64 | 74.0 | 10.98 | 1.10 | 54.50 | 67.00 | 74.0 | 82.00 | 92.50 | 9 |
| Asian | 91 | 46 | 125 | 72.16 | 71.5 | 11.66 | 1.22 | 55.50 | 65.25 | 71.5 | 78.75 | 91.50 | 5 |
| Other | 21 | 59 | 85 | 72.12 | 71.5 | 7.31 | 1.60 | 59.75 | 69.00 | 71.5 | 76.50 | 82.75 | 5 |
| AIAN | 8 | 58 | 93 | 71.50 | 66.0 | 13.85 | 4.90 | 59.00 | 62.50 | 66.0 | 80.00 | 90.75 | 2 |
| PTNS | 5 | 62 | 90 | 79.80 | 86.0 | 11.41 | 5.10 | 64.60 | 75.00 | 86.0 | 86.00 | 89.20 | 0 |
| NHOPI | 4 | 62 | 85 | 74.75 | 76.0 | 10.69 | 5.34 | 63.20 | 68.00 | 76.0 | 82.75 | 84.55 | 0 |

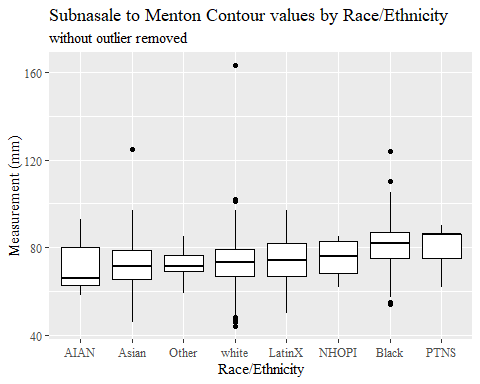
#%>% set\_header\_Cabels(values = list(SnasM\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(SnasM\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Subnasale to Menton Contour SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Subnasale to Menton Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 44 | 163 | 72.89 | 73.0 | 10.09 | 0.29 | 56.00 | 67.00 | 73.0 | 79.00 | 89.00 | 144 |
| Black | 548 | 54 | 124 | 81.09 | 82.0 | 10.03 | 0.43 | 63.00 | 75.00 | 82.0 | 87.00 | 96.20 | 71 |
| LatinX | 100 | 50 | 97 | 73.64 | 74.0 | 10.98 | 1.10 | 54.50 | 67.00 | 74.0 | 82.00 | 92.50 | 9 |
| Asian | 91 | 46 | 125 | 72.16 | 71.5 | 11.66 | 1.22 | 55.50 | 65.25 | 71.5 | 78.75 | 91.50 | 5 |
| Other | 21 | 59 | 85 | 72.12 | 71.5 | 7.31 | 1.60 | 59.75 | 69.00 | 71.5 | 76.50 | 82.75 | 5 |
| AIAN | 8 | 58 | 93 | 71.50 | 66.0 | 13.85 | 4.90 | 59.00 | 62.50 | 66.0 | 80.00 | 90.75 | 2 |
| PTNS | 5 | 62 | 90 | 79.80 | 86.0 | 11.41 | 5.10 | 64.60 | 75.00 | 86.0 | 86.00 | 89.20 | 0 |
| NHOPI | 4 | 62 | 85 | 74.75 | 76.0 | 10.69 | 5.34 | 63.20 | 68.00 | 76.0 | 82.75 | 84.55 | 0 |

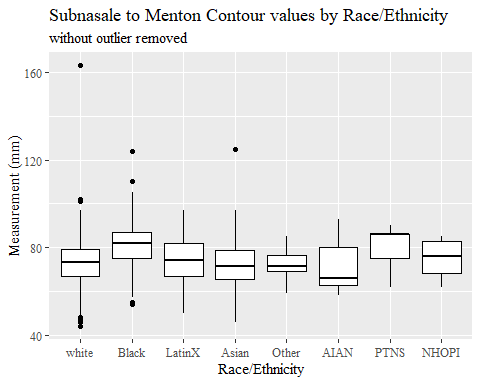
#%>% set\_header\_Cabels(values = list(SnasM\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, SnasM\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=SnasM\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Subnasale to Menton Contour values by Race/Ethnicity",  
 subtitle = "without outlier removed",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 236 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=SnasM\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Subnasale to Menton Contour values by Race/Ethnicity",  
 subtitle = "without outlier removed",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 236 rows containing non-finite values (stat\_boxplot).



#SnasM\_C gender sumstats  
SnasM\_Cgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(SnasM\_C, na.rm = TRUE),  
 max = max(SnasM\_C, na.rm = TRUE),  
 mean = mean(SnasM\_C, na.rm = TRUE),  
 mdn = median(SnasM\_C, na.rm = TRUE),  
 sd = sd(SnasM\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(SnasM\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(SnasM\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(SnasM\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(SnasM\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(SnasM\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(SnasM\_C)))  
  
SnasM\_Cgender\_sumstats <- SnasM\_Cgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(SnasM\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Subnasale to Menton Contour SumStats by Gender")

**Table** : Subnasale to Menton Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 46 | 125 | 73.03 | 73 | 10.04 | 0.31 | 55.0 | 67.0 | 73 | 80.00 | 88.00 | 23 |
| Male | 939 | 46 | 163 | 78.11 | 79 | 11.14 | 0.36 | 60.0 | 71.0 | 79 | 85.00 | 94.75 | 213 |
| Non-binary or Other | 5 | 44 | 81 | 67.20 | 75 | 16.07 | 7.19 | 46.6 | 57.0 | 75 | 79.00 | 80.60 | 0 |
| Prefer not to say | 1 | 90 | 90 | 90.00 | 90 |  |  | 90.0 | 90.0 | 90 | 90.00 | 90.00 | 0 |
|  | 8 | 64 | 87 | 75.50 | 75 | 8.38 | 2.96 | 64.7 | 70.5 | 75 | 80.75 | 86.65 | 0 |

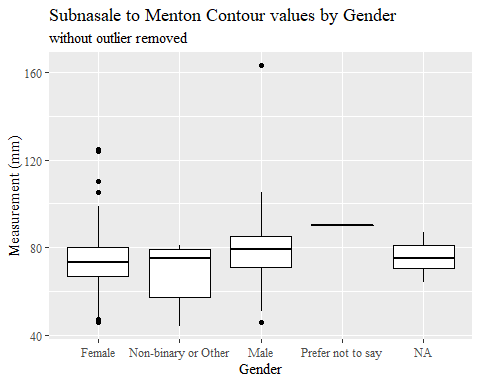
#%>% set\_header\_Cabels(values = list(SnasM\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(SnasM\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Subnasale to Menton Contour SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Subnasale to Menton Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 46 | 125 | 73.03 | 73 | 10.04 | 0.31 | 55.0 | 67.0 | 73 | 80.00 | 88.00 | 23 |
| Male | 939 | 46 | 163 | 78.11 | 79 | 11.14 | 0.36 | 60.0 | 71.0 | 79 | 85.00 | 94.75 | 213 |
| Non-binary or Other | 5 | 44 | 81 | 67.20 | 75 | 16.07 | 7.19 | 46.6 | 57.0 | 75 | 79.00 | 80.60 | 0 |
| Prefer not to say | 1 | 90 | 90 | 90.00 | 90 |  |  | 90.0 | 90.0 | 90 | 90.00 | 90.00 | 0 |
|  | 8 | 64 | 87 | 75.50 | 75 | 8.38 | 2.96 | 64.7 | 70.5 | 75 | 80.75 | 86.65 | 0 |

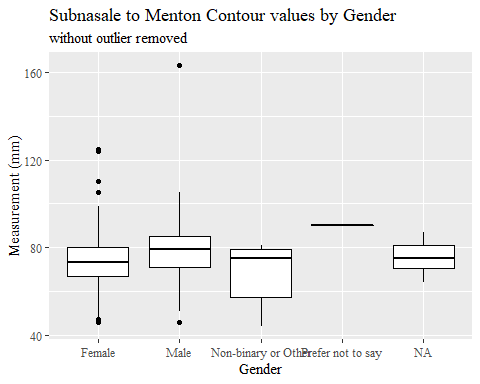
#%>% set\_header\_Cabels(values = list(SnasM\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, SnasM\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=SnasM\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Subnasale to Menton Contour values by Gender",  
 subtitle = "without outlier removed",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 236 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=SnasM\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Subnasale to Menton Contour values by Gender",  
 subtitle = "without outlier removed",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 236 rows containing non-finite values (stat\_boxplot).



#SnasM\_C age group sumstats  
SnasM\_Cage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(SnasM\_C, na.rm = TRUE),  
 max = max(SnasM\_C, na.rm = TRUE),  
 mean = mean(SnasM\_C, na.rm = TRUE),  
 sd = sd(SnasM\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(SnasM\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(SnasM\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(SnasM\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(SnasM\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(SnasM\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(SnasM\_C)))  
  
SnasM\_Cage\_sumstats <- SnasM\_Cage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(SnasM\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Subnasale to Menton Contour SumStats by Age Group")

**Table** : Subnasale to Menton Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 44 | 163 | 74.98 | 10.77 | 0.34 | 57 | 68 | 75.0 | 82 | 92.00 | 124 |
| 37-54 | 940 | 47 | 125 | 75.38 | 10.82 | 0.35 | 57 | 69 | 75.0 | 83 | 93.00 | 106 |
| 55-72 | 84 | 49 | 102 | 73.50 | 11.13 | 1.21 | 53 | 66 | 71.5 | 82 | 88.45 | 6 |
|  | 1 | 71 | 71 | 71.00 |  |  | 71 | 71 | 71.0 | 71 | 71.00 | 0 |

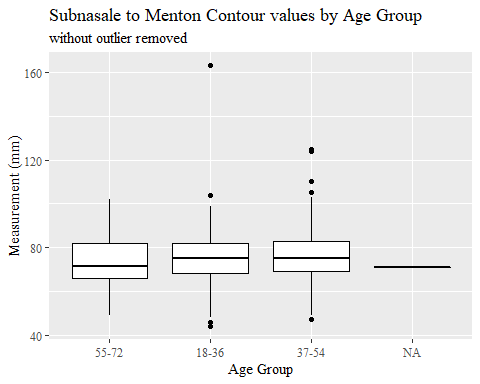
#%>% set\_header\_Cabels(values = list(SnasM\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(SnasM\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Subnasale to Menton Contour SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Subnasale to Menton Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 44 | 163 | 74.98 | 10.77 | 0.34 | 57 | 68 | 75.0 | 82 | 92.00 | 124 |
| 37-54 | 940 | 47 | 125 | 75.38 | 10.82 | 0.35 | 57 | 69 | 75.0 | 83 | 93.00 | 106 |
| 55-72 | 84 | 49 | 102 | 73.50 | 11.13 | 1.21 | 53 | 66 | 71.5 | 82 | 88.45 | 6 |
|  | 1 | 71 | 71 | 71.00 |  |  | 71 | 71 | 71.0 | 71 | 71.00 | 0 |

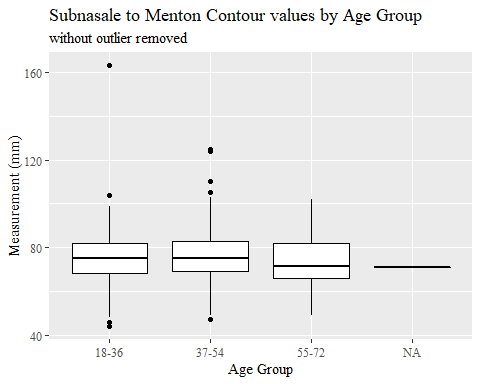
#%>% set\_header\_Cabels(values = list(SnasM\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, SnasM\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=SnasM\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Subnasale to Menton Contour values by Age Group",  
 subtitle = "without outlier removed",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 236 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=SnasM\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Subnasale to Menton Contour values by Age Group",  
 subtitle = "without outlier removed",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 236 rows containing non-finite values (stat\_boxplot).



SnasM\_C with outlier removed

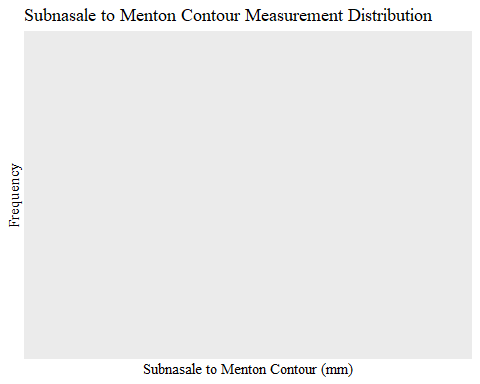
#REMOVING OUTLIER SnasM\_C  
  
max(headscan\_full$SnasM\_C, na.rm = TRUE)

## [1] 163

#CODE TO REMOVE OUTLIER, use later to remove from headscan\_full  
SnasM\_C\_no\_out <- headscan\_full %>% mutate(SnasM\_C = replace(SnasM\_C, SnasM\_C>16, NA))

#histogram of all SnasM\_C values  
ggplot(data=SnasM\_C\_no\_out, aes(x=SnasM\_C))+  
 geom\_bar(stat="bin", bins=20, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Subnasale to Menton Contour Measurement Distribution",  
 y="Frequency",  
 x="Subnasale to Menton Contour (mm)")

## Warning: Removed 2017 rows containing non-finite values (stat\_bin).



#SnasM\_C race/eth sumstats  
SnasM\_Crace\_sumstats1 <- SnasM\_C\_no\_out %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(SnasM\_C, na.rm = TRUE),  
 max = max(SnasM\_C, na.rm = TRUE),  
 mean = mean(SnasM\_C, na.rm = TRUE),  
 mdn = median(SnasM\_C, na.rm = TRUE),  
 sd = sd(SnasM\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(SnasM\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(SnasM\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(SnasM\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(SnasM\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(SnasM\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(SnasM\_C)))

## Warning in min(SnasM\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(SnasM\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(SnasM\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(SnasM\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(SnasM\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(SnasM\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(SnasM\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(SnasM\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf

## Warning in max(SnasM\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(SnasM\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(SnasM\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(SnasM\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(SnasM\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(SnasM\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(SnasM\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(SnasM\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf

SnasM\_Crace\_sumstats1 <- SnasM\_Crace\_sumstats1 %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(SnasM\_Crace\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Subnasale to Menton Contour SumStats by Race/Ethnicity")

**Table** : Subnasale to Menton Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1,240 |
| Black | 548 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 548 |
| LatinX | 100 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 100 |
| Asian | 91 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 91 |
| Other | 21 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 21 |
| AIAN | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |
| PTNS | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| NHOPI | 4 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 4 |

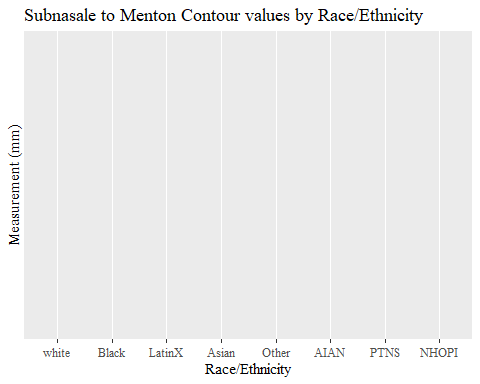
#%>% set\_header\_Cabels(values = list(SnasM\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(SnasM\_Crace\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Subnasale to Menton Contour SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Subnasale to Menton Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1,240 |
| Black | 548 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 548 |
| LatinX | 100 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 100 |
| Asian | 91 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 91 |
| Other | 21 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 21 |
| AIAN | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |
| PTNS | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| NHOPI | 4 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 4 |

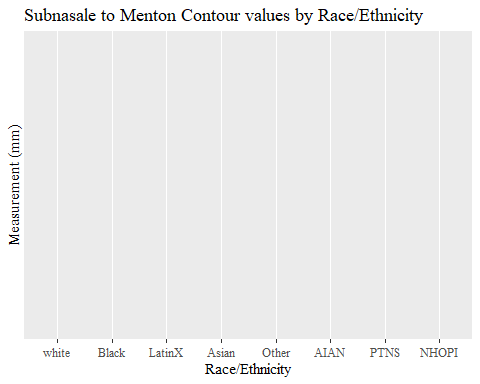
#%>% set\_header\_Cabels(values = list(SnasM\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
SnasM\_C\_no\_out %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, SnasM\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=SnasM\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Subnasale to Menton Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=SnasM\_C\_no\_out, aes(y=SnasM\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Subnasale to Menton Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#SnasM\_C gender sumstats  
SnasM\_Cgender\_sumstats1 <- SnasM\_C\_no\_out %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(SnasM\_C, na.rm = TRUE),  
 max = max(SnasM\_C, na.rm = TRUE),  
 mean = mean(SnasM\_C, na.rm = TRUE),  
 mdn = median(SnasM\_C, na.rm = TRUE),  
 sd = sd(SnasM\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(SnasM\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(SnasM\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(SnasM\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(SnasM\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(SnasM\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(SnasM\_C)))

## Warning in min(SnasM\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(SnasM\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(SnasM\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(SnasM\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(SnasM\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf

## Warning in max(SnasM\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(SnasM\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(SnasM\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(SnasM\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(SnasM\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf

SnasM\_Cgender\_sumstats1 <- SnasM\_Cgender\_sumstats1 %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(SnasM\_Cgender\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Subnasale to Menton Contour SumStats by Gender")

**Table** : Subnasale to Menton Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1,064 |
| Male | 939 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 939 |
| Non-binary or Other | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| Prefer not to say | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |
|  | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |

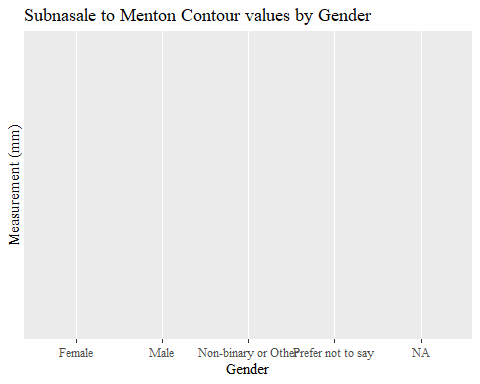
#%>% set\_header\_Cabels(values = list(SnasM\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(SnasM\_Cgender\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Subnasale to Menton Contour SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Subnasale to Menton Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1,064 |
| Male | 939 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 939 |
| Non-binary or Other | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| Prefer not to say | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |
|  | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |

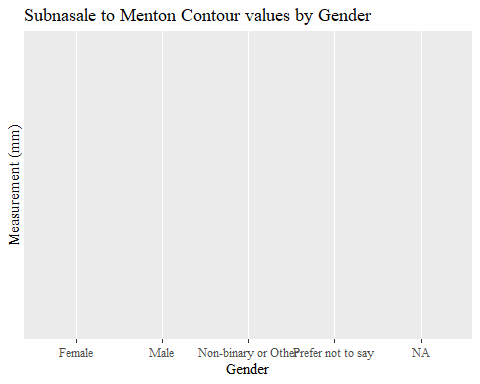
#%>% set\_header\_Cabels(values = list(SnasM\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
SnasM\_C\_no\_out %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, SnasM\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=SnasM\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Subnasale to Menton Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=SnasM\_C\_no\_out, aes(y=SnasM\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Subnasale to Menton Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#SnasM\_C age group sumstats  
SnasM\_Cage\_sumstats1 <- SnasM\_C\_no\_out %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(SnasM\_C, na.rm = TRUE),  
 max = max(SnasM\_C, na.rm = TRUE),  
 mean = mean(SnasM\_C, na.rm = TRUE),  
 sd = sd(SnasM\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(SnasM\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(SnasM\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(SnasM\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(SnasM\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(SnasM\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(SnasM\_C)))

## Warning in min(SnasM\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(SnasM\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(SnasM\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(SnasM\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf

## Warning in max(SnasM\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(SnasM\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(SnasM\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(SnasM\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf

SnasM\_Cage\_sumstats1 <- SnasM\_Cage\_sumstats1 %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(SnasM\_Cage\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Subnasale to Menton Contour SumStats by Age Group")

**Table** : Subnasale to Menton Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | Inf | -Inf |  |  |  |  |  |  |  |  | 992 |
| 37-54 | 940 | Inf | -Inf |  |  |  |  |  |  |  |  | 940 |
| 55-72 | 84 | Inf | -Inf |  |  |  |  |  |  |  |  | 84 |
|  | 1 | Inf | -Inf |  |  |  |  |  |  |  |  | 1 |

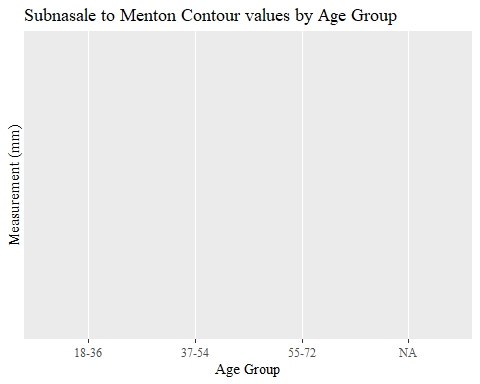
#%>% set\_header\_Cabels(values = list(SnasM\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(SnasM\_Cage\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Subnasale to Menton Contour SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Subnasale to Menton Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | Inf | -Inf |  |  |  |  |  |  |  |  | 992 |
| 37-54 | 940 | Inf | -Inf |  |  |  |  |  |  |  |  | 940 |
| 55-72 | 84 | Inf | -Inf |  |  |  |  |  |  |  |  | 84 |
|  | 1 | Inf | -Inf |  |  |  |  |  |  |  |  | 1 |

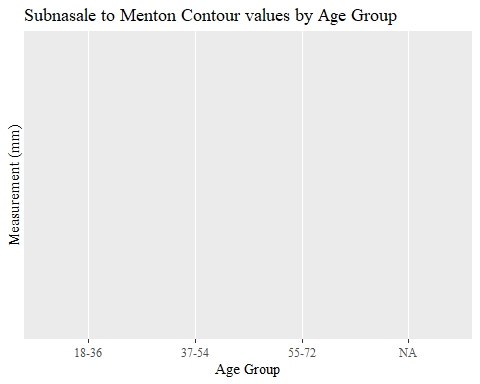
#%>% set\_header\_Cabels(values = list(SnasM\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
SnasM\_C\_no\_out %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, SnasM\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=SnasM\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Subnasale to Menton Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=SnasM\_C\_no\_out, aes(y=SnasM\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Subnasale to Menton Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

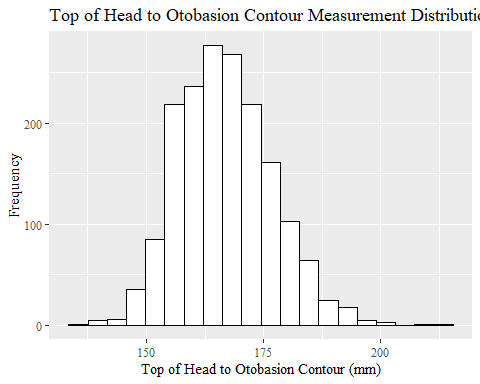
## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



TrHO\_C

#histogram of all TrHO\_C values  
ggplot(data=headscan\_full, aes(x=TrHO\_C))+  
 geom\_bar(stat="bin", bins=20, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Top of Head to Otobasion Contour Measurement Distribution",  
 y="Frequency",  
 x="Top of Head to Otobasion Contour (mm)")

## Warning: Removed 283 rows containing non-finite values (stat\_bin).



#TrHO\_C race/eth sumstats  
TrHO\_Crace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(TrHO\_C, na.rm = TRUE),  
 max = max(TrHO\_C, na.rm = TRUE),  
 mean = mean(TrHO\_C, na.rm = TRUE),  
 mdn = median(TrHO\_C, na.rm = TRUE),  
 sd = sd(TrHO\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrHO\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrHO\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrHO\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrHO\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrHO\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrHO\_C)))   
  
TrHO\_Crace\_sumstats <- TrHO\_Crace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(TrHO\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Top of Head to Otobasion Contour SumStats by Race/Ethnicity")

**Table** : Top of Head to Otobasion Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 135 | 213 | 166.19 | 166.0 | 9.71 | 0.28 | 152.00 | 159.00 | 166.0 | 172.00 | 183.00 | 117 |
| Black | 548 | 141 | 203 | 168.11 | 168.0 | 10.85 | 0.46 | 151.00 | 160.00 | 168.0 | 175.00 | 185.00 | 138 |
| LatinX | 100 | 150 | 196 | 168.34 | 167.0 | 10.11 | 1.01 | 155.00 | 161.00 | 167.0 | 174.00 | 185.60 | 11 |
| Asian | 91 | 151 | 188 | 169.96 | 169.0 | 7.90 | 0.83 | 156.05 | 165.00 | 169.0 | 175.75 | 182.90 | 9 |
| Other | 21 | 142 | 184 | 167.50 | 167.5 | 10.11 | 2.21 | 154.75 | 162.25 | 167.5 | 174.75 | 184.00 | 3 |
| AIAN | 8 | 153 | 182 | 167.00 | 167.0 | 12.15 | 4.30 | 153.60 | 156.00 | 167.0 | 177.50 | 180.80 | 1 |
| PTNS | 5 | 181 | 181 | 181.00 | 181.0 |  |  | 181.00 | 181.00 | 181.0 | 181.00 | 181.00 | 4 |
| NHOPI | 4 | 166 | 179 | 173.25 | 174.0 | 5.38 | 2.69 | 167.20 | 172.00 | 174.0 | 175.25 | 178.25 | 0 |

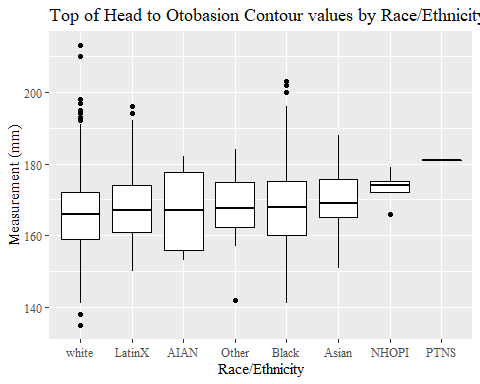
#%>% set\_header\_Cabels(values = list(TrHO\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrHO\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Top of Head to Otobasion Contour SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Top of Head to Otobasion Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 135 | 213 | 166.19 | 166.0 | 9.71 | 0.28 | 152.00 | 159.00 | 166.0 | 172.00 | 183.00 | 117 |
| Black | 548 | 141 | 203 | 168.11 | 168.0 | 10.85 | 0.46 | 151.00 | 160.00 | 168.0 | 175.00 | 185.00 | 138 |
| LatinX | 100 | 150 | 196 | 168.34 | 167.0 | 10.11 | 1.01 | 155.00 | 161.00 | 167.0 | 174.00 | 185.60 | 11 |
| Asian | 91 | 151 | 188 | 169.96 | 169.0 | 7.90 | 0.83 | 156.05 | 165.00 | 169.0 | 175.75 | 182.90 | 9 |
| Other | 21 | 142 | 184 | 167.50 | 167.5 | 10.11 | 2.21 | 154.75 | 162.25 | 167.5 | 174.75 | 184.00 | 3 |
| AIAN | 8 | 153 | 182 | 167.00 | 167.0 | 12.15 | 4.30 | 153.60 | 156.00 | 167.0 | 177.50 | 180.80 | 1 |
| PTNS | 5 | 181 | 181 | 181.00 | 181.0 |  |  | 181.00 | 181.00 | 181.0 | 181.00 | 181.00 | 4 |
| NHOPI | 4 | 166 | 179 | 173.25 | 174.0 | 5.38 | 2.69 | 167.20 | 172.00 | 174.0 | 175.25 | 178.25 | 0 |

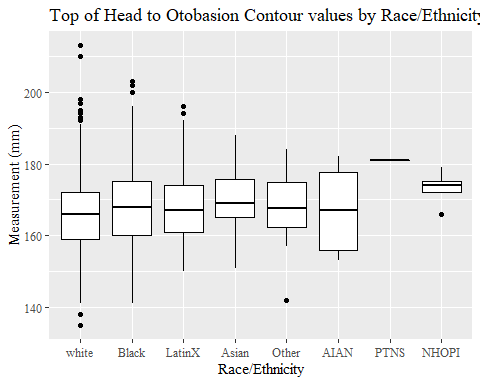
#%>% set\_header\_Cabels(values = list(TrHO\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, TrHO\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrHO\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Top of Head to Otobasion Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 283 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=TrHO\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Top of Head to Otobasion Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 283 rows containing non-finite values (stat\_boxplot).



#TrHO\_C gender sumstats  
TrHO\_Cgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(TrHO\_C, na.rm = TRUE),  
 max = max(TrHO\_C, na.rm = TRUE),  
 mean = mean(TrHO\_C, na.rm = TRUE),  
 mdn = median(TrHO\_C, na.rm = TRUE),  
 sd = sd(TrHO\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrHO\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrHO\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrHO\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrHO\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrHO\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrHO\_C)))

## Warning in min(TrHO\_C, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf

## Warning in max(TrHO\_C, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf

TrHO\_Cgender\_sumstats <- TrHO\_Cgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(TrHO\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Top of Head to Otobasion Contour SumStats by Gender")

**Table** : Top of Head to Otobasion Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 135 | 202 | 164.41 | 164 | 9.94 | 0.30 | 151.0 | 157 | 164 | 170 | 182.0 | 215 |
| Male | 939 | 141 | 213 | 169.45 | 169 | 9.43 | 0.31 | 154.0 | 163 | 169 | 175 | 185.0 | 64 |
| Non-binary or Other | 5 | 156 | 184 | 172.00 | 175 | 10.42 | 4.66 | 158.6 | 169 | 175 | 176 | 182.4 | 0 |
| Prefer not to say | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |
|  | 8 | 159 | 175 | 165.40 | 164 | 6.35 | 2.24 | 159.4 | 161 | 164 | 168 | 173.6 | 3 |

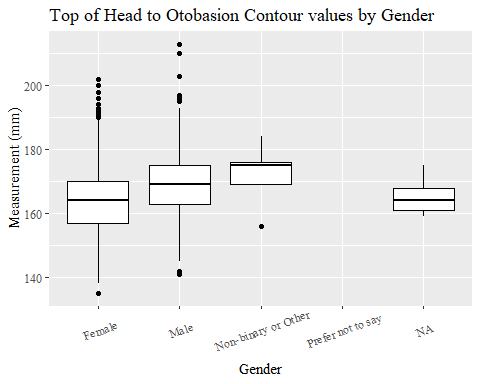
#%>% set\_header\_Cabels(values = list(TrHO\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrHO\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Top of Head to Otobasion Contour SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Top of Head to Otobasion Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 135 | 202 | 164.41 | 164 | 9.94 | 0.30 | 151.0 | 157 | 164 | 170 | 182.0 | 215 |
| Male | 939 | 141 | 213 | 169.45 | 169 | 9.43 | 0.31 | 154.0 | 163 | 169 | 175 | 185.0 | 64 |
| Non-binary or Other | 5 | 156 | 184 | 172.00 | 175 | 10.42 | 4.66 | 158.6 | 169 | 175 | 176 | 182.4 | 0 |
| Prefer not to say | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |
|  | 8 | 159 | 175 | 165.40 | 164 | 6.35 | 2.24 | 159.4 | 161 | 164 | 168 | 173.6 | 3 |

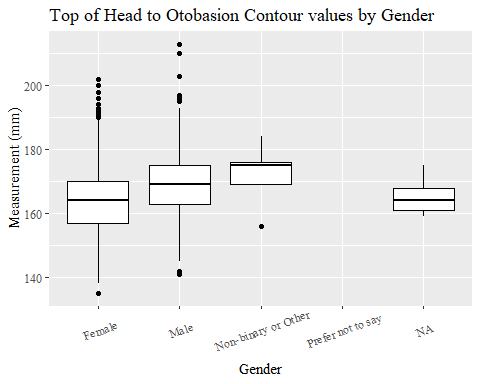
#%>% set\_header\_Cabels(values = list(TrHO\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, TrHO\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrHO\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Top of Head to Otobasion Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 283 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=TrHO\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Top of Head to Otobasion Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 283 rows containing non-finite values (stat\_boxplot).



#TrHO\_C age group sumstats  
TrHO\_Cage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(TrHO\_C, na.rm = TRUE),  
 max = max(TrHO\_C, na.rm = TRUE),  
 mean = mean(TrHO\_C, na.rm = TRUE),  
 mdn = median(TrHO\_C, na.rm = TRUE),  
 sd = sd(TrHO\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrHO\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrHO\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrHO\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrHO\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrHO\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrHO\_C)))  
  
TrHO\_Cage\_sumstats <- TrHO\_Cage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(TrHO\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Top of Head to Otobasion Contour SumStats by Age Group")

**Table** : Top of Head to Otobasion Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 135 | 213 | 168.04 | 167.0 | 10.13 | 0.32 | 153.00 | 161 | 167.0 | 175 | 184.1 | 153 |
| 37-54 | 940 | 138 | 210 | 165.84 | 166.0 | 9.68 | 0.32 | 151.00 | 159 | 166.0 | 172 | 182.0 | 122 |
| 55-72 | 84 | 147 | 192 | 167.43 | 166.5 | 10.83 | 1.18 | 150.75 | 161 | 166.5 | 174 | 186.0 | 8 |
|  | 1 | 169 | 169 | 169.00 | 169.0 |  |  | 169.00 | 169 | 169.0 | 169 | 169.0 | 0 |

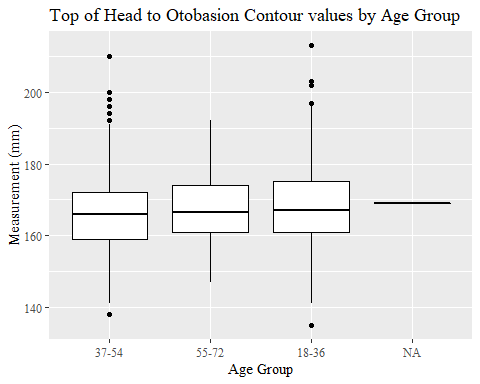
#%>% set\_header\_Cabels(values = list(TrHO\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrHO\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Top of Head to Otobasion Contour SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Top of Head to Otobasion Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 135 | 213 | 168.04 | 167.0 | 10.13 | 0.32 | 153.00 | 161 | 167.0 | 175 | 184.1 | 153 |
| 37-54 | 940 | 138 | 210 | 165.84 | 166.0 | 9.68 | 0.32 | 151.00 | 159 | 166.0 | 172 | 182.0 | 122 |
| 55-72 | 84 | 147 | 192 | 167.43 | 166.5 | 10.83 | 1.18 | 150.75 | 161 | 166.5 | 174 | 186.0 | 8 |
|  | 1 | 169 | 169 | 169.00 | 169.0 |  |  | 169.00 | 169 | 169.0 | 169 | 169.0 | 0 |

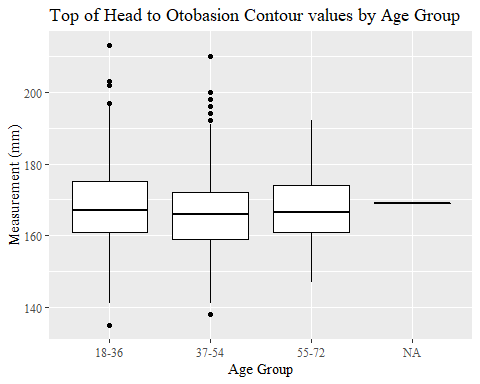
#%>% set\_header\_Cabels(values = list(TrHO\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, TrHO\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrHO\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Top of Head to Otobasion Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 283 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=TrHO\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Top of Head to Otobasion Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

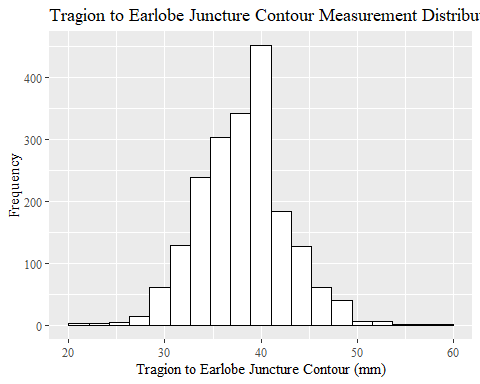
## Warning: Removed 283 rows containing non-finite values (stat\_boxplot).



TrEJ\_C

#histogram of all TrEJ\_C values  
ggplot(data=headscan\_full, aes(x=TrEJ\_C))+  
 geom\_bar(stat="bin", bins=20, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Tragion to Earlobe Juncture Contour Measurement Distribution",  
 y="Frequency",  
 x="Tragion to Earlobe Juncture Contour (mm)")

## Warning: Removed 33 rows containing non-finite values (stat\_bin).



#TrEJ\_C race/eth sumstats  
TrEJ\_Crace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(TrEJ\_C, na.rm = TRUE),  
 max = max(TrEJ\_C, na.rm = TRUE),  
 mean = mean(TrEJ\_C, na.rm = TRUE),  
 mdn = median(TrEJ\_C, na.rm = TRUE),  
 sd = sd(TrEJ\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrEJ\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrEJ\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrEJ\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrEJ\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrEJ\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrEJ\_C)))   
  
TrEJ\_Crace\_sumstats <- TrEJ\_Crace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(TrEJ\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Earlobe Juncture Contour SumStats by Race/Ethnicity")

**Table** : Tragion to Earlobe Juncture Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 20 | 60 | 38.40 | 38.0 | 4.58 | 0.13 | 32.0 | 35.00 | 38.0 | 41.00 | 46.00 | 17 |
| Black | 548 | 22 | 54 | 36.68 | 36.5 | 4.65 | 0.20 | 30.0 | 34.00 | 36.5 | 40.00 | 44.00 | 10 |
| LatinX | 100 | 28 | 52 | 39.63 | 39.0 | 4.47 | 0.45 | 33.9 | 37.00 | 39.0 | 43.00 | 48.00 | 1 |
| Asian | 91 | 28 | 54 | 39.44 | 39.0 | 4.42 | 0.46 | 32.8 | 37.00 | 39.0 | 42.00 | 48.00 | 2 |
| Other | 21 | 30 | 51 | 37.65 | 35.0 | 6.19 | 1.35 | 30.0 | 33.00 | 35.0 | 43.25 | 46.25 | 1 |
| AIAN | 8 | 34 | 44 | 38.50 | 39.0 | 4.14 | 1.46 | 34.0 | 34.75 | 39.0 | 41.00 | 43.25 | 2 |
| PTNS | 5 | 28 | 44 | 36.40 | 36.0 | 6.07 | 2.71 | 29.2 | 34.00 | 36.0 | 40.00 | 43.20 | 0 |
| NHOPI | 4 | 31 | 48 | 37.00 | 34.5 | 7.62 | 3.81 | 31.3 | 32.50 | 34.5 | 39.00 | 46.20 | 0 |

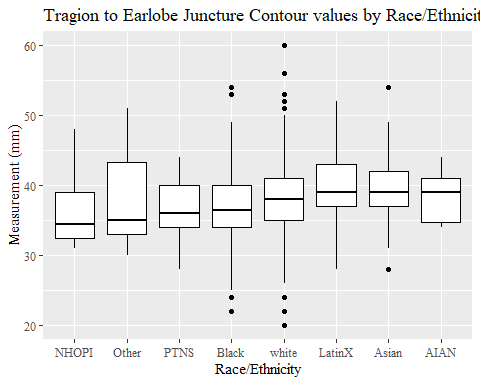
#%>% set\_header\_Cabels(values = list(TrEJ\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrEJ\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Earlobe Juncture Contour SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Earlobe Juncture Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 20 | 60 | 38.40 | 38.0 | 4.58 | 0.13 | 32.0 | 35.00 | 38.0 | 41.00 | 46.00 | 17 |
| Black | 548 | 22 | 54 | 36.68 | 36.5 | 4.65 | 0.20 | 30.0 | 34.00 | 36.5 | 40.00 | 44.00 | 10 |
| LatinX | 100 | 28 | 52 | 39.63 | 39.0 | 4.47 | 0.45 | 33.9 | 37.00 | 39.0 | 43.00 | 48.00 | 1 |
| Asian | 91 | 28 | 54 | 39.44 | 39.0 | 4.42 | 0.46 | 32.8 | 37.00 | 39.0 | 42.00 | 48.00 | 2 |
| Other | 21 | 30 | 51 | 37.65 | 35.0 | 6.19 | 1.35 | 30.0 | 33.00 | 35.0 | 43.25 | 46.25 | 1 |
| AIAN | 8 | 34 | 44 | 38.50 | 39.0 | 4.14 | 1.46 | 34.0 | 34.75 | 39.0 | 41.00 | 43.25 | 2 |
| PTNS | 5 | 28 | 44 | 36.40 | 36.0 | 6.07 | 2.71 | 29.2 | 34.00 | 36.0 | 40.00 | 43.20 | 0 |
| NHOPI | 4 | 31 | 48 | 37.00 | 34.5 | 7.62 | 3.81 | 31.3 | 32.50 | 34.5 | 39.00 | 46.20 | 0 |

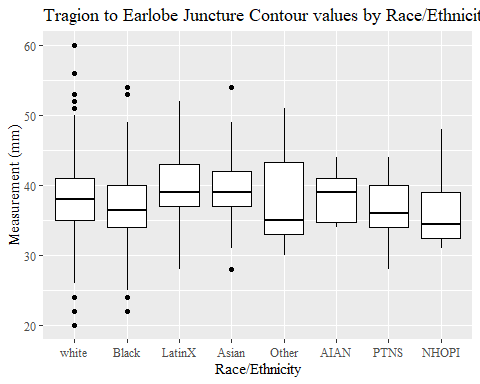
#%>% set\_header\_Cabels(values = list(TrEJ\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, TrEJ\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrEJ\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Earlobe Juncture Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 33 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=TrEJ\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Earlobe Juncture Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 33 rows containing non-finite values (stat\_boxplot).



#TrEJ\_C gender sumstats  
TrEJ\_Cgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(TrEJ\_C, na.rm = TRUE),  
 max = max(TrEJ\_C, na.rm = TRUE),  
 mean = mean(TrEJ\_C, na.rm = TRUE),  
 mdn = median(TrEJ\_C, na.rm = TRUE),  
 sd = sd(TrEJ\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrEJ\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrEJ\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrEJ\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrEJ\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrEJ\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrEJ\_C)))  
  
TrEJ\_Cgender\_sumstats <- TrEJ\_Cgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(TrEJ\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Earlobe Juncture Contour SumStats by Gender")

**Table** : Tragion to Earlobe Juncture Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 22 | 53 | 37.35 | 37.0 | 4.38 | 0.13 | 30.00 | 34.00 | 37.0 | 40 | 44.0 | 23 |
| Male | 939 | 20 | 60 | 38.80 | 39.0 | 4.91 | 0.16 | 31.00 | 35.00 | 39.0 | 42 | 47.0 | 10 |
| Non-binary or Other | 5 | 33 | 45 | 37.80 | 36.0 | 5.07 | 2.27 | 33.20 | 34.00 | 36.0 | 41 | 44.2 | 0 |
| Prefer not to say | 1 | 36 | 36 | 36.00 | 36.0 |  |  | 36.00 | 36.00 | 36.0 | 36 | 36.0 | 0 |
|  | 8 | 31 | 44 | 36.62 | 36.5 | 4.72 | 1.67 | 31.35 | 32.75 | 36.5 | 40 | 42.6 | 0 |

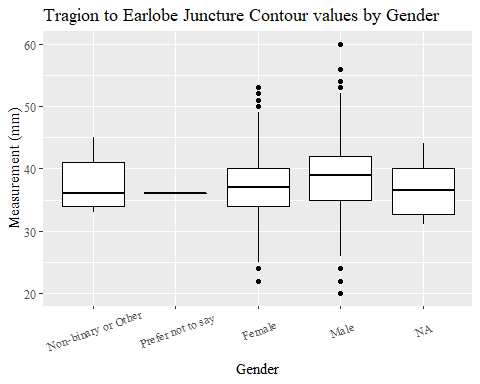
#%>% set\_header\_Cabels(values = list(TrEJ\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrEJ\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Earlobe Juncture Contour SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Earlobe Juncture Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 22 | 53 | 37.35 | 37.0 | 4.38 | 0.13 | 30.00 | 34.00 | 37.0 | 40 | 44.0 | 23 |
| Male | 939 | 20 | 60 | 38.80 | 39.0 | 4.91 | 0.16 | 31.00 | 35.00 | 39.0 | 42 | 47.0 | 10 |
| Non-binary or Other | 5 | 33 | 45 | 37.80 | 36.0 | 5.07 | 2.27 | 33.20 | 34.00 | 36.0 | 41 | 44.2 | 0 |
| Prefer not to say | 1 | 36 | 36 | 36.00 | 36.0 |  |  | 36.00 | 36.00 | 36.0 | 36 | 36.0 | 0 |
|  | 8 | 31 | 44 | 36.62 | 36.5 | 4.72 | 1.67 | 31.35 | 32.75 | 36.5 | 40 | 42.6 | 0 |

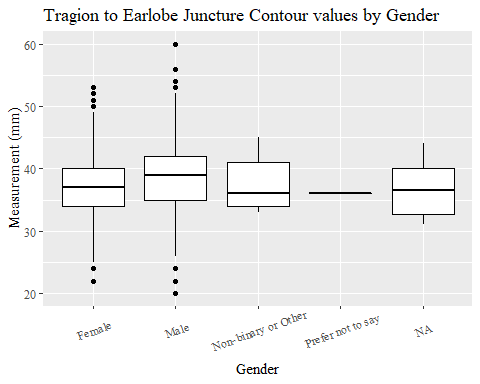
#%>% set\_header\_Cabels(values = list(TrEJ\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, TrEJ\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrEJ\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Earlobe Juncture Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 33 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=TrEJ\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Earlobe Juncture Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 33 rows containing non-finite values (stat\_boxplot).



#TrEJ\_C age group sumstats  
TrEJ\_Cage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(TrEJ\_C, na.rm = TRUE),  
 max = max(TrEJ\_C, na.rm = TRUE),  
 mean = mean(TrEJ\_C, na.rm = TRUE),  
 mdn = median(TrEJ\_C, na.rm = TRUE),  
 sd = sd(TrEJ\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrEJ\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrEJ\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrEJ\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrEJ\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrEJ\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrEJ\_C)))  
  
TrEJ\_Cage\_sumstats <- TrEJ\_Cage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(TrEJ\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Earlobe Juncture Contour SumStats by Age Group")

**Table** : Tragion to Earlobe Juncture Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 22 | 56 | 37.48 | 37 | 4.59 | 0.15 | 30 | 34 | 37 | 40 | 45 | 12 |
| 37-54 | 940 | 20 | 60 | 38.43 | 38 | 4.72 | 0.15 | 31 | 35 | 38 | 41 | 46 | 19 |
| 55-72 | 84 | 29 | 52 | 40.09 | 40 | 4.66 | 0.51 | 34 | 36 | 40 | 43 | 48 | 2 |
|  | 1 | 33 | 33 | 33.00 | 33 |  |  | 33 | 33 | 33 | 33 | 33 | 0 |

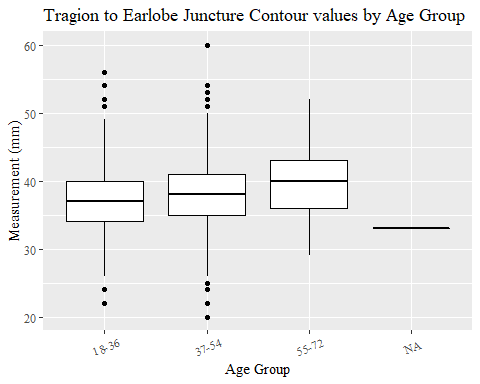
#%>% set\_header\_Cabels(values = list(TrEJ\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrEJ\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Earlobe Juncture Contour SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Earlobe Juncture Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 22 | 56 | 37.48 | 37 | 4.59 | 0.15 | 30 | 34 | 37 | 40 | 45 | 12 |
| 37-54 | 940 | 20 | 60 | 38.43 | 38 | 4.72 | 0.15 | 31 | 35 | 38 | 41 | 46 | 19 |
| 55-72 | 84 | 29 | 52 | 40.09 | 40 | 4.66 | 0.51 | 34 | 36 | 40 | 43 | 48 | 2 |
|  | 1 | 33 | 33 | 33.00 | 33 |  |  | 33 | 33 | 33 | 33 | 33 | 0 |

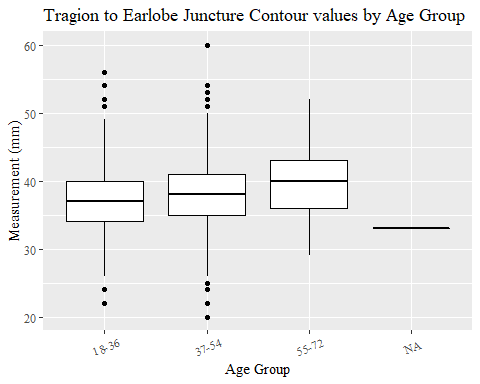
#%>% set\_header\_Cabels(values = list(TrEJ\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, TrEJ\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrEJ\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Earlobe Juncture Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 33 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=TrEJ\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Earlobe Juncture Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

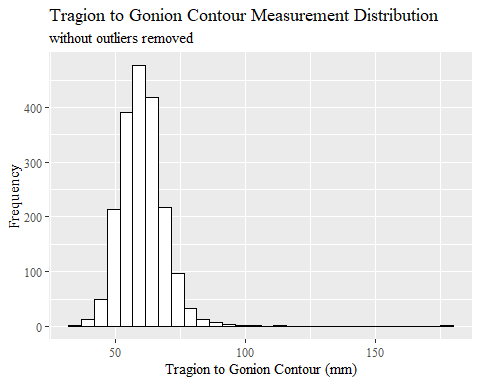
## Warning: Removed 33 rows containing non-finite values (stat\_boxplot).



TrGo\_C without outliers removed

#histogram of all TrGo\_C values  
ggplot(data=headscan\_full, aes(x=TrGo\_C))+  
 geom\_bar(stat="bin", bins=30, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Tragion to Gonion Contour Measurement Distribution",  
 subtitle = "without outliers removed",  
 y="Frequency",  
 x="Tragion to Gonion Contour (mm)")

## Warning: Removed 80 rows containing non-finite values (stat\_bin).



#TrGo\_C race/eth sumstats  
TrGo\_Crace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(TrGo\_C, na.rm = TRUE),  
 max = max(TrGo\_C, na.rm = TRUE),  
 mean = mean(TrGo\_C, na.rm = TRUE),  
 mdn = median(TrGo\_C, na.rm = TRUE),  
 sd = sd(TrGo\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrGo\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrGo\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrGo\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrGo\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrGo\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrGo\_C)))   
  
TrGo\_Crace\_sumstats <- TrGo\_Crace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(TrGo\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Gonion Contour SumStats by Race/Ethnicity")

**Table** : Tragion to Gonion Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 35 | 178 | 60.10 | 60.0 | 9.19 | 0.26 | 48.00 | 54.00 | 60.0 | 65.00 | 74.00 | 53 |
| Black | 548 | 35 | 96 | 59.77 | 59.0 | 8.08 | 0.35 | 48.00 | 54.00 | 59.0 | 64.50 | 73.00 | 17 |
| LatinX | 100 | 41 | 83 | 60.43 | 61.0 | 7.27 | 0.73 | 49.50 | 56.00 | 61.0 | 64.25 | 73.00 | 4 |
| Asian | 91 | 44 | 87 | 61.39 | 60.0 | 8.42 | 0.88 | 48.80 | 55.00 | 60.0 | 66.00 | 76.00 | 2 |
| Other | 21 | 43 | 76 | 57.11 | 56.5 | 8.32 | 1.82 | 45.55 | 51.75 | 56.5 | 61.75 | 67.50 | 3 |
| AIAN | 8 | 56 | 70 | 61.14 | 60.0 | 5.11 | 1.81 | 56.30 | 57.50 | 60.0 | 63.50 | 68.80 | 1 |
| PTNS | 5 | 58 | 81 | 69.60 | 70.0 | 9.02 | 4.03 | 59.20 | 64.00 | 70.0 | 75.00 | 79.80 | 0 |
| NHOPI | 4 | 49 | 71 | 62.00 | 64.0 | 9.27 | 4.64 | 51.25 | 60.25 | 64.0 | 65.75 | 69.95 | 0 |

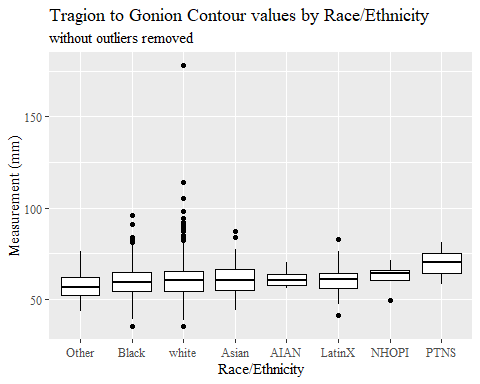
#%>% set\_header\_Cabels(values = list(TrGo\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrGo\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Gonion Contour SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Gonion Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 35 | 178 | 60.10 | 60.0 | 9.19 | 0.26 | 48.00 | 54.00 | 60.0 | 65.00 | 74.00 | 53 |
| Black | 548 | 35 | 96 | 59.77 | 59.0 | 8.08 | 0.35 | 48.00 | 54.00 | 59.0 | 64.50 | 73.00 | 17 |
| LatinX | 100 | 41 | 83 | 60.43 | 61.0 | 7.27 | 0.73 | 49.50 | 56.00 | 61.0 | 64.25 | 73.00 | 4 |
| Asian | 91 | 44 | 87 | 61.39 | 60.0 | 8.42 | 0.88 | 48.80 | 55.00 | 60.0 | 66.00 | 76.00 | 2 |
| Other | 21 | 43 | 76 | 57.11 | 56.5 | 8.32 | 1.82 | 45.55 | 51.75 | 56.5 | 61.75 | 67.50 | 3 |
| AIAN | 8 | 56 | 70 | 61.14 | 60.0 | 5.11 | 1.81 | 56.30 | 57.50 | 60.0 | 63.50 | 68.80 | 1 |
| PTNS | 5 | 58 | 81 | 69.60 | 70.0 | 9.02 | 4.03 | 59.20 | 64.00 | 70.0 | 75.00 | 79.80 | 0 |
| NHOPI | 4 | 49 | 71 | 62.00 | 64.0 | 9.27 | 4.64 | 51.25 | 60.25 | 64.0 | 65.75 | 69.95 | 0 |

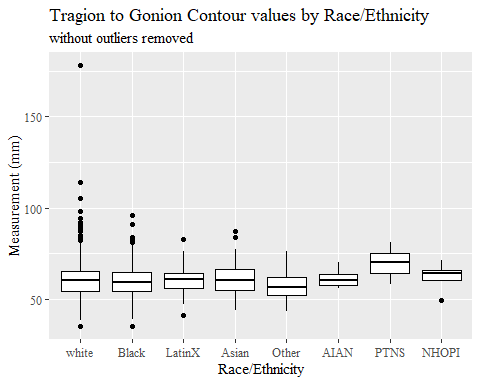
#%>% set\_header\_Cabels(values = list(TrGo\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, TrGo\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrGo\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Gonion Contour values by Race/Ethnicity",  
 subtitle = "without outliers removed",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 80 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=TrGo\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Gonion Contour values by Race/Ethnicity",  
 subtitle = "without outliers removed",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 80 rows containing non-finite values (stat\_boxplot).



#TrGo\_C gender sumstats  
TrGo\_Cgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(TrGo\_C, na.rm = TRUE),  
 max = max(TrGo\_C, na.rm = TRUE),  
 mean = mean(TrGo\_C, na.rm = TRUE),  
 mdn = median(TrGo\_C, na.rm = TRUE),  
 sd = sd(TrGo\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrGo\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrGo\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrGo\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrGo\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrGo\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrGo\_C)))  
  
TrGo\_Cgender\_sumstats <- TrGo\_Cgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(TrGo\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Gonion Contour SumStats by Gender")

**Table** : Tragion to Gonion Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 35 | 98 | 58.24 | 58 | 7.68 | 0.24 | 47.0 | 53 | 58 | 63 | 70.00 | 24 |
| Male | 939 | 35 | 178 | 62.32 | 62 | 9.45 | 0.31 | 49.0 | 56 | 62 | 67 | 76.00 | 56 |
| Non-binary or Other | 5 | 45 | 59 | 53.80 | 55 | 5.40 | 2.42 | 46.6 | 53 | 55 | 57 | 58.60 | 0 |
| Prefer not to say | 1 | 58 | 58 | 58.00 | 58 |  |  | 58.0 | 58 | 58 | 58 | 58.00 | 0 |
|  | 8 | 52 | 73 | 58.38 | 55 | 8.05 | 2.85 | 52.0 | 52 | 55 | 62 | 71.25 | 0 |

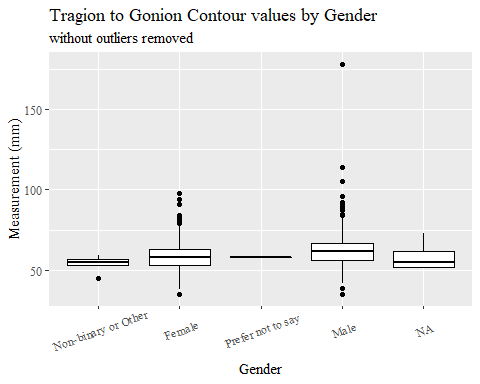
#%>% set\_header\_Cabels(values = list(TrGo\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrGo\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Gonion Contour SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Gonion Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 35 | 98 | 58.24 | 58 | 7.68 | 0.24 | 47.0 | 53 | 58 | 63 | 70.00 | 24 |
| Male | 939 | 35 | 178 | 62.32 | 62 | 9.45 | 0.31 | 49.0 | 56 | 62 | 67 | 76.00 | 56 |
| Non-binary or Other | 5 | 45 | 59 | 53.80 | 55 | 5.40 | 2.42 | 46.6 | 53 | 55 | 57 | 58.60 | 0 |
| Prefer not to say | 1 | 58 | 58 | 58.00 | 58 |  |  | 58.0 | 58 | 58 | 58 | 58.00 | 0 |
|  | 8 | 52 | 73 | 58.38 | 55 | 8.05 | 2.85 | 52.0 | 52 | 55 | 62 | 71.25 | 0 |

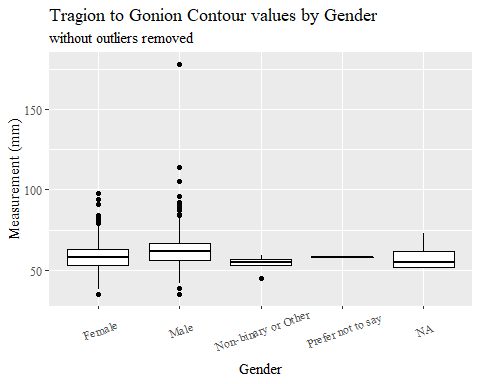
#%>% set\_header\_Cabels(values = list(TrGo\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, TrGo\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrGo\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Gonion Contour values by Gender",  
 subtitle = "without outliers removed",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 80 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=TrGo\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Gonion Contour values by Gender",  
 subtitle = "without outliers removed",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 80 rows containing non-finite values (stat\_boxplot).



#TrGo\_C age group sumstats  
TrGo\_Cage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(TrGo\_C, na.rm = TRUE),  
 max = max(TrGo\_C, na.rm = TRUE),  
 mean = mean(TrGo\_C, na.rm = TRUE),  
 mdn = median(TrGo\_C, na.rm = TRUE),  
 sd = sd(TrGo\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrGo\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrGo\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrGo\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrGo\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrGo\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrGo\_C)))  
  
TrGo\_Cage\_sumstats <- TrGo\_Cage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(TrGo\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Gonion Contour SumStats by Age Group")

**Table** : Tragion to Gonion Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 38 | 92 | 60.07 | 60 | 8.10 | 0.26 | 48 | 54.00 | 60 | 65.00 | 74.00 | 33 |
| 37-54 | 940 | 35 | 114 | 59.94 | 60 | 8.62 | 0.28 | 47 | 54.00 | 60 | 65.00 | 73.30 | 45 |
| 55-72 | 84 | 41 | 178 | 62.06 | 61 | 15.44 | 1.69 | 48 | 55.25 | 61 | 65.75 | 75.95 | 2 |
|  | 1 | 62 | 62 | 62.00 | 62 |  |  | 62 | 62.00 | 62 | 62.00 | 62.00 | 0 |

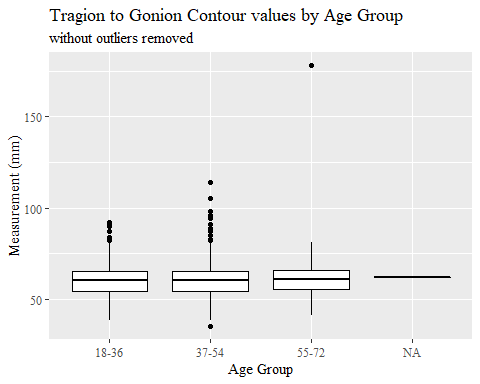
#%>% set\_header\_Cabels(values = list(TrGo\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrGo\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Gonion Contour SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Gonion Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 38 | 92 | 60.07 | 60 | 8.10 | 0.26 | 48 | 54.00 | 60 | 65.00 | 74.00 | 33 |
| 37-54 | 940 | 35 | 114 | 59.94 | 60 | 8.62 | 0.28 | 47 | 54.00 | 60 | 65.00 | 73.30 | 45 |
| 55-72 | 84 | 41 | 178 | 62.06 | 61 | 15.44 | 1.69 | 48 | 55.25 | 61 | 65.75 | 75.95 | 2 |
|  | 1 | 62 | 62 | 62.00 | 62 |  |  | 62 | 62.00 | 62 | 62.00 | 62.00 | 0 |

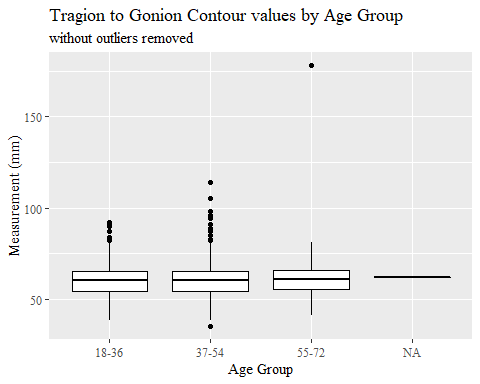
#%>% set\_header\_Cabels(values = list(TrGo\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, TrGo\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrGo\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Gonion Contour values by Age Group",  
 subtitle = "without outliers removed",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 80 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=TrGo\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Gonion Contour values by Age Group",  
 subtitle = "without outliers removed",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 80 rows containing non-finite values (stat\_boxplot).



TrGo\_C with outliers removed

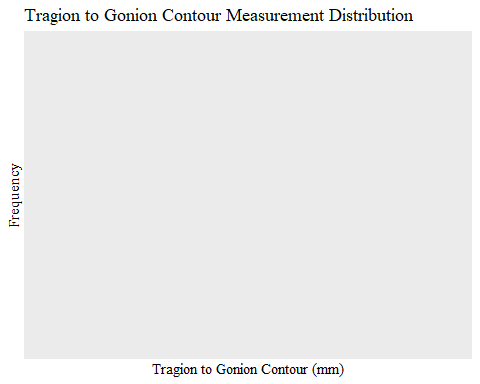
max(headscan\_full$TrGo\_C, na.rm = TRUE)

## [1] 178

#CODE TO REMOVE OUTLIER, use later to remove from headscan\_full  
TrGo\_C\_no\_out <- headscan\_full %>% mutate(TrGo\_C = replace(TrGo\_C, TrGo\_C>17, NA))

#histogram of all TrGo\_C values  
ggplot(data=TrGo\_C\_no\_out, aes(x=TrGo\_C))+  
 geom\_bar(stat="bin", bins=25, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Tragion to Gonion Contour Measurement Distribution",  
 y="Frequency",  
 x="Tragion to Gonion Contour (mm)")

## Warning: Removed 2017 rows containing non-finite values (stat\_bin).



#TrGo\_C race/eth sumstats  
TrGo\_Crace\_sumstats1 <- TrGo\_C\_no\_out %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(TrGo\_C, na.rm = TRUE),  
 max = max(TrGo\_C, na.rm = TRUE),  
 mean = mean(TrGo\_C, na.rm = TRUE),  
 mdn = median(TrGo\_C, na.rm = TRUE),  
 sd = sd(TrGo\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrGo\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrGo\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrGo\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrGo\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrGo\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrGo\_C)))

## Warning in min(TrGo\_C, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf  
  
## Warning in min(TrGo\_C, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf  
  
## Warning in min(TrGo\_C, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf  
  
## Warning in min(TrGo\_C, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf  
  
## Warning in min(TrGo\_C, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf  
  
## Warning in min(TrGo\_C, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf  
  
## Warning in min(TrGo\_C, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf  
  
## Warning in min(TrGo\_C, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf

## Warning in max(TrGo\_C, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf  
  
## Warning in max(TrGo\_C, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf  
  
## Warning in max(TrGo\_C, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf  
  
## Warning in max(TrGo\_C, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf  
  
## Warning in max(TrGo\_C, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf  
  
## Warning in max(TrGo\_C, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf  
  
## Warning in max(TrGo\_C, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf  
  
## Warning in max(TrGo\_C, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf

TrGo\_Crace\_sumstats1 <- TrGo\_Crace\_sumstats1 %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(TrGo\_Crace\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Gonion Contour SumStats by Race/Ethnicity")

**Table** : Tragion to Gonion Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1,240 |
| Black | 548 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 548 |
| LatinX | 100 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 100 |
| Asian | 91 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 91 |
| Other | 21 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 21 |
| AIAN | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |
| PTNS | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| NHOPI | 4 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 4 |

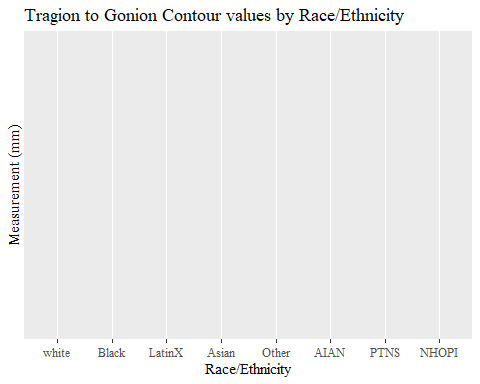
#%>% set\_header\_Cabels(values = list(TrGo\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrGo\_Crace\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Gonion Contour SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Gonion Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1,240 |
| Black | 548 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 548 |
| LatinX | 100 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 100 |
| Asian | 91 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 91 |
| Other | 21 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 21 |
| AIAN | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |
| PTNS | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| NHOPI | 4 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 4 |

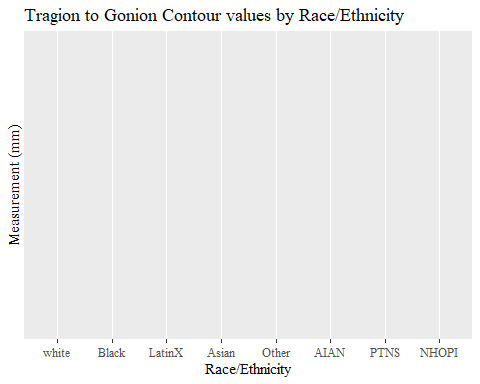
#%>% set\_header\_Cabels(values = list(TrGo\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
TrGo\_C\_no\_out %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, TrGo\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrGo\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Gonion Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=TrGo\_C\_no\_out, aes(y=TrGo\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Gonion Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#TrGo\_C gender sumstats  
TrGo\_Cgender\_sumstats1 <- TrGo\_C\_no\_out %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(TrGo\_C, na.rm = TRUE),  
 max = max(TrGo\_C, na.rm = TRUE),  
 mean = mean(TrGo\_C, na.rm = TRUE),  
 mdn = median(TrGo\_C, na.rm = TRUE),  
 sd = sd(TrGo\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrGo\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrGo\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrGo\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrGo\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrGo\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrGo\_C)))

## Warning in min(TrGo\_C, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf  
  
## Warning in min(TrGo\_C, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf  
  
## Warning in min(TrGo\_C, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf  
  
## Warning in min(TrGo\_C, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf  
  
## Warning in min(TrGo\_C, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf

## Warning in max(TrGo\_C, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf  
  
## Warning in max(TrGo\_C, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf  
  
## Warning in max(TrGo\_C, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf  
  
## Warning in max(TrGo\_C, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf  
  
## Warning in max(TrGo\_C, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf

TrGo\_Cgender\_sumstats1 <- TrGo\_Cgender\_sumstats1 %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(TrGo\_Cgender\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Gonion Contour SumStats by Gender")

**Table** : Tragion to Gonion Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1,064 |
| Male | 939 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 939 |
| Non-binary or Other | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| Prefer not to say | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |
|  | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |

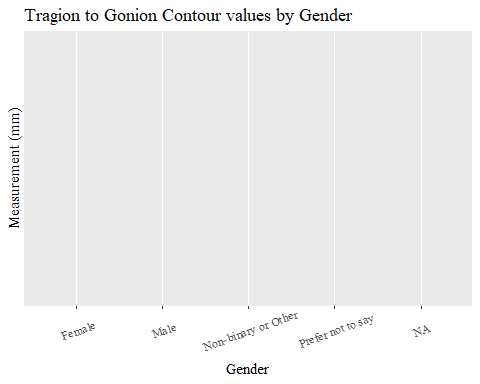
#%>% set\_header\_Cabels(values = list(TrGo\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrGo\_Cgender\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Gonion Contour SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Gonion Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1,064 |
| Male | 939 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 939 |
| Non-binary or Other | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| Prefer not to say | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |
|  | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |

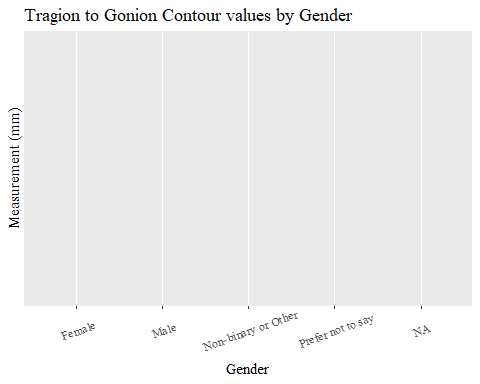
#%>% set\_header\_Cabels(values = list(TrGo\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
TrGo\_C\_no\_out %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, TrGo\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrGo\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Gonion Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=TrGo\_C\_no\_out, aes(y=TrGo\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Gonion Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#TrGo\_C age group sumstats  
TrGo\_Cage\_sumstats1 <- TrGo\_C\_no\_out %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(TrGo\_C, na.rm = TRUE),  
 max = max(TrGo\_C, na.rm = TRUE),  
 mean = mean(TrGo\_C, na.rm = TRUE),  
 mdn = median(TrGo\_C, na.rm = TRUE),  
 sd = sd(TrGo\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrGo\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrGo\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrGo\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrGo\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrGo\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrGo\_C)))

## Warning in min(TrGo\_C, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf  
  
## Warning in min(TrGo\_C, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf  
  
## Warning in min(TrGo\_C, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf  
  
## Warning in min(TrGo\_C, na.rm = TRUE): no non-missing arguments to min; returning  
## Inf

## Warning in max(TrGo\_C, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf  
  
## Warning in max(TrGo\_C, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf  
  
## Warning in max(TrGo\_C, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf  
  
## Warning in max(TrGo\_C, na.rm = TRUE): no non-missing arguments to max; returning  
## -Inf

TrGo\_Cage\_sumstats1 <- TrGo\_Cage\_sumstats1 %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(TrGo\_Cage\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Gonion Contour SumStats by Age Group")

**Table** : Tragion to Gonion Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 992 |
| 37-54 | 940 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 940 |
| 55-72 | 84 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 84 |
|  | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |

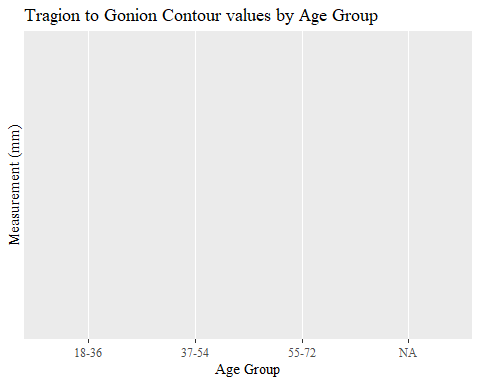
#%>% set\_header\_Cabels(values = list(TrGo\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrGo\_Cage\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Gonion Contour SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Gonion Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 992 |
| 37-54 | 940 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 940 |
| 55-72 | 84 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 84 |
|  | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |

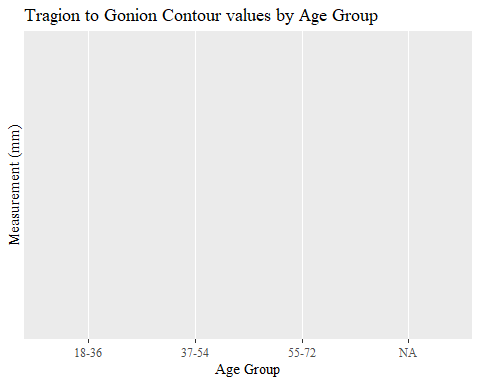
#%>% set\_header\_Cabels(values = list(TrGo\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
TrGo\_C\_no\_out %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, TrGo\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrGo\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Gonion Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=TrGo\_C\_no\_out, aes(y=TrGo\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Gonion Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

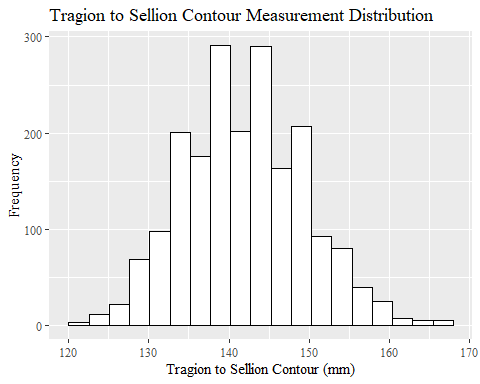
## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



TrSel\_C

#histogram of all TrSel\_C values  
ggplot(data=headscan\_full, aes(x=TrSel\_C))+  
 geom\_bar(stat="bin", bins=20, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Tragion to Sellion Contour Measurement Distribution",  
 y="Frequency",  
 x="Tragion to Sellion Contour (mm)")

## Warning: Removed 31 rows containing non-finite values (stat\_bin).



#TrSel\_C race/eth sumstats  
TrSel\_Crace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(TrSel\_C, na.rm = TRUE),  
 max = max(TrSel\_C, na.rm = TRUE),  
 mean = mean(TrSel\_C, na.rm = TRUE),  
 mdn = median(TrSel\_C, na.rm = TRUE),  
 sd = sd(TrSel\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrSel\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrSel\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrSel\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrSel\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrSel\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrSel\_C)))   
  
TrSel\_Crace\_sumstats <- TrSel\_Crace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(TrSel\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Sellion Contour SumStats by Race/Ethnicity")

**Table** : Tragion to Sellion Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 120 | 168 | 140.81 | 141.0 | 7.25 | 0.21 | 130.00 | 136.00 | 141.0 | 146 | 153.00 | 18 |
| Black | 548 | 124 | 168 | 144.77 | 145.0 | 7.28 | 0.31 | 133.00 | 140.00 | 145.0 | 149 | 157.00 | 7 |
| LatinX | 100 | 125 | 162 | 141.15 | 141.0 | 7.32 | 0.73 | 129.00 | 137.00 | 141.0 | 146 | 151.10 | 1 |
| Asian | 91 | 122 | 165 | 141.56 | 140.0 | 7.80 | 0.82 | 132.00 | 136.00 | 140.0 | 147 | 155.30 | 3 |
| Other | 21 | 129 | 162 | 142.40 | 143.0 | 7.68 | 1.68 | 130.90 | 138.25 | 143.0 | 145 | 151.55 | 1 |
| AIAN | 8 | 133 | 145 | 137.29 | 136.0 | 4.15 | 1.47 | 133.30 | 134.50 | 136.0 | 139 | 143.50 | 1 |
| PTNS | 5 | 135 | 150 | 143.00 | 145.0 | 5.96 | 2.66 | 135.80 | 139.00 | 145.0 | 146 | 149.20 | 0 |
| NHOPI | 4 | 142 | 148 | 145.75 | 146.5 | 2.87 | 1.44 | 142.45 | 144.25 | 146.5 | 148 | 148.00 | 0 |

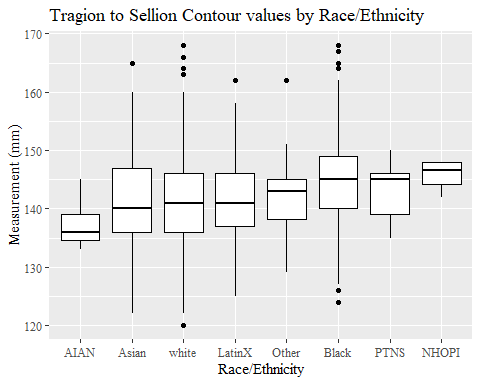
#%>% set\_header\_Cabels(values = list(TrSel\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrSel\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Sellion Contour SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Sellion Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 120 | 168 | 140.81 | 141.0 | 7.25 | 0.21 | 130.00 | 136.00 | 141.0 | 146 | 153.00 | 18 |
| Black | 548 | 124 | 168 | 144.77 | 145.0 | 7.28 | 0.31 | 133.00 | 140.00 | 145.0 | 149 | 157.00 | 7 |
| LatinX | 100 | 125 | 162 | 141.15 | 141.0 | 7.32 | 0.73 | 129.00 | 137.00 | 141.0 | 146 | 151.10 | 1 |
| Asian | 91 | 122 | 165 | 141.56 | 140.0 | 7.80 | 0.82 | 132.00 | 136.00 | 140.0 | 147 | 155.30 | 3 |
| Other | 21 | 129 | 162 | 142.40 | 143.0 | 7.68 | 1.68 | 130.90 | 138.25 | 143.0 | 145 | 151.55 | 1 |
| AIAN | 8 | 133 | 145 | 137.29 | 136.0 | 4.15 | 1.47 | 133.30 | 134.50 | 136.0 | 139 | 143.50 | 1 |
| PTNS | 5 | 135 | 150 | 143.00 | 145.0 | 5.96 | 2.66 | 135.80 | 139.00 | 145.0 | 146 | 149.20 | 0 |
| NHOPI | 4 | 142 | 148 | 145.75 | 146.5 | 2.87 | 1.44 | 142.45 | 144.25 | 146.5 | 148 | 148.00 | 0 |

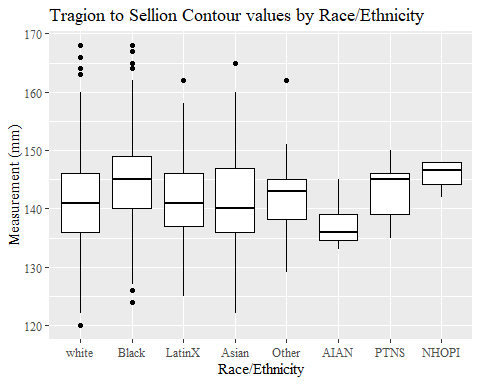
#%>% set\_header\_Cabels(values = list(TrSel\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, TrSel\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrSel\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Sellion Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 31 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=TrSel\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Sellion Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 31 rows containing non-finite values (stat\_boxplot).



#TrSel\_C gender sumstats  
TrSel\_Cgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(TrSel\_C, na.rm = TRUE),  
 max = max(TrSel\_C, na.rm = TRUE),  
 mean = mean(TrSel\_C, na.rm = TRUE),  
 mdn = median(TrSel\_C, na.rm = TRUE),  
 sd = sd(TrSel\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrSel\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrSel\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrSel\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrSel\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrSel\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrSel\_C)))  
  
TrSel\_Cgender\_sumstats <- TrSel\_Cgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(TrSel\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Sellion Contour SumStats by Gender")

**Table** : Tragion to Sellion Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 120 | 168 | 138.62 | 138 | 6.66 | 0.20 | 129.00 | 134 | 138 | 143.00 | 150.00 | 22 |
| Male | 939 | 125 | 168 | 145.72 | 146 | 6.55 | 0.21 | 135.00 | 141 | 146 | 150.00 | 156.55 | 9 |
| Non-binary or Other | 5 | 130 | 144 | 139.00 | 140 | 5.29 | 2.37 | 132.00 | 140 | 140 | 141.00 | 143.40 | 0 |
| Prefer not to say | 1 | 146 | 146 | 146.00 | 146 |  |  | 146.00 | 146 | 146 | 146.00 | 146.00 | 0 |
|  | 8 | 135 | 145 | 140.38 | 141 | 3.38 | 1.19 | 135.35 | 139 | 141 | 142.25 | 144.30 | 0 |

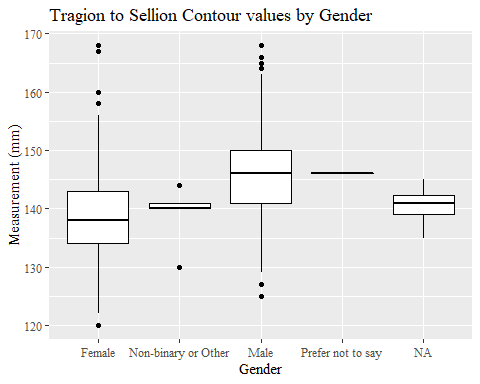
#%>% set\_header\_Cabels(values = list(TrSel\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrSel\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Sellion Contour SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Sellion Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 120 | 168 | 138.62 | 138 | 6.66 | 0.20 | 129.00 | 134 | 138 | 143.00 | 150.00 | 22 |
| Male | 939 | 125 | 168 | 145.72 | 146 | 6.55 | 0.21 | 135.00 | 141 | 146 | 150.00 | 156.55 | 9 |
| Non-binary or Other | 5 | 130 | 144 | 139.00 | 140 | 5.29 | 2.37 | 132.00 | 140 | 140 | 141.00 | 143.40 | 0 |
| Prefer not to say | 1 | 146 | 146 | 146.00 | 146 |  |  | 146.00 | 146 | 146 | 146.00 | 146.00 | 0 |
|  | 8 | 135 | 145 | 140.38 | 141 | 3.38 | 1.19 | 135.35 | 139 | 141 | 142.25 | 144.30 | 0 |

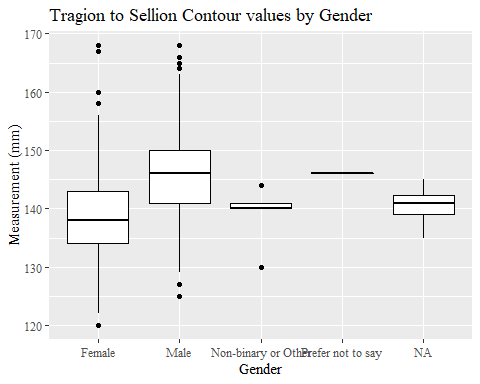
#%>% set\_header\_Cabels(values = list(TrSel\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, TrSel\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrSel\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Sellion Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 31 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=TrSel\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Sellion Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 31 rows containing non-finite values (stat\_boxplot).



#TrSel\_C age group sumstats  
TrSel\_Cage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(TrSel\_C, na.rm = TRUE),  
 max = max(TrSel\_C, na.rm = TRUE),  
 mean = mean(TrSel\_C, na.rm = TRUE),  
 mdn = median(TrSel\_C, na.rm = TRUE),  
 sd = sd(TrSel\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrSel\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrSel\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrSel\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrSel\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrSel\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrSel\_C)))  
  
TrSel\_Cage\_sumstats <- TrSel\_Cage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(TrSel\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Sellion Contour SumStats by Age Group")

**Table** : Tragion to Sellion Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 122 | 168 | 141.68 | 142 | 7.54 | 0.24 | 130 | 137 | 142 | 147 | 155.0 | 13 |
| 37-54 | 940 | 120 | 168 | 142.04 | 142 | 7.44 | 0.24 | 131 | 136 | 142 | 147 | 155.0 | 17 |
| 55-72 | 84 | 127 | 160 | 144.22 | 146 | 6.87 | 0.75 | 134 | 138 | 146 | 149 | 154.8 | 1 |
|  | 1 | 145 | 145 | 145.00 | 145 |  |  | 145 | 145 | 145 | 145 | 145.0 | 0 |

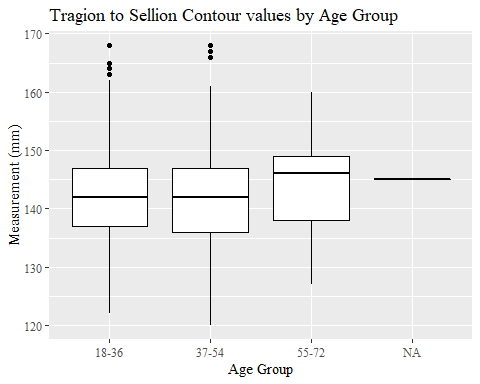
#%>% set\_header\_Cabels(values = list(TrSel\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrSel\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Sellion Contour SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Sellion Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 122 | 168 | 141.68 | 142 | 7.54 | 0.24 | 130 | 137 | 142 | 147 | 155.0 | 13 |
| 37-54 | 940 | 120 | 168 | 142.04 | 142 | 7.44 | 0.24 | 131 | 136 | 142 | 147 | 155.0 | 17 |
| 55-72 | 84 | 127 | 160 | 144.22 | 146 | 6.87 | 0.75 | 134 | 138 | 146 | 149 | 154.8 | 1 |
|  | 1 | 145 | 145 | 145.00 | 145 |  |  | 145 | 145 | 145 | 145 | 145.0 | 0 |

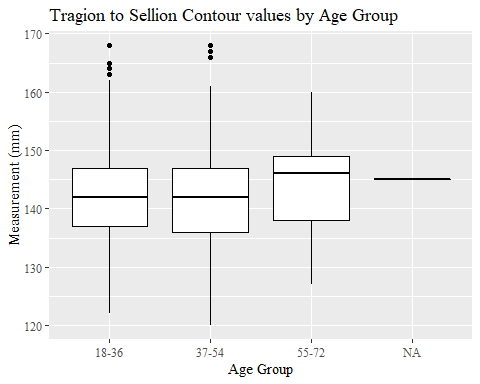
#%>% set\_header\_Cabels(values = list(TrSel\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, TrSel\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrSel\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Sellion Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 31 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=TrSel\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Sellion Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

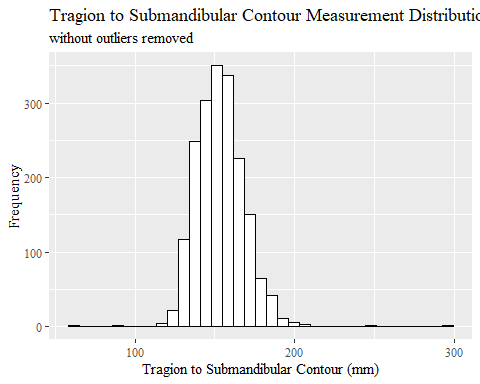
## Warning: Removed 31 rows containing non-finite values (stat\_boxplot).



TrSman\_C without outlier removed

#histogram of all TrSman\_C values  
ggplot(data=headscan\_full, aes(x=TrSman\_C))+  
 geom\_bar(stat="bin", bins=35, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Tragion to Submandibular Contour Measurement Distribution",  
 subtitle = "without outliers removed",  
 y="Frequency",  
 x="Tragion to Submandibular Contour (mm)")

## Warning: Removed 132 rows containing non-finite values (stat\_bin).



#TrSman\_C race/eth sumstats  
TrSman\_Crace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(TrSman\_C, na.rm = TRUE),  
 max = max(TrSman\_C, na.rm = TRUE),  
 mean = mean(TrSman\_C, na.rm = TRUE),  
 mdn = median(TrSman\_C, na.rm = TRUE),  
 sd = sd(TrSman\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrSman\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrSman\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrSman\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrSman\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrSman\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrSman\_C)))   
  
TrSman\_Crace\_sumstats <- TrSman\_Crace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(TrSman\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Submandibular Contour SumStats by Race/Ethnicity")

**Table** : Tragion to Submandibular Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 64 | 298 | 152.31 | 151.0 | 15.58 | 0.44 | 131.0 | 142.0 | 151.0 | 161.00 | 178.65 | 72 |
| Black | 548 | 122 | 201 | 156.78 | 156.0 | 12.65 | 0.54 | 138.0 | 148.0 | 156.0 | 165.50 | 178.90 | 45 |
| LatinX | 100 | 126 | 195 | 153.07 | 150.5 | 14.70 | 1.47 | 132.0 | 142.0 | 150.5 | 162.25 | 176.75 | 4 |
| Asian | 91 | 122 | 196 | 150.31 | 150.0 | 13.67 | 1.43 | 130.0 | 140.5 | 150.0 | 158.00 | 173.10 | 4 |
| Other | 21 | 132 | 165 | 150.00 | 150.5 | 9.91 | 2.16 | 133.5 | 146.0 | 150.5 | 156.75 | 162.00 | 5 |
| AIAN | 8 | 130 | 169 | 144.43 | 146.0 | 13.88 | 4.91 | 130.9 | 133.0 | 146.0 | 150.00 | 164.20 | 1 |
| PTNS | 5 | 139 | 181 | 155.80 | 154.0 | 15.83 | 7.08 | 140.6 | 147.0 | 154.0 | 158.00 | 176.40 | 0 |
| NHOPI | 4 | 143 | 151 | 147.67 | 149.0 | 4.16 | 2.08 | 143.6 | 146.0 | 149.0 | 150.00 | 150.80 | 1 |

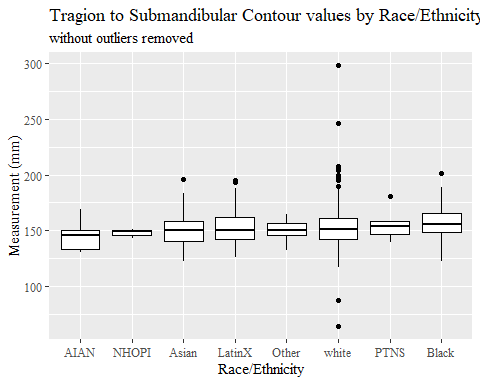
#%>% set\_header\_Cabels(values = list(TrSman\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrSman\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Submandibular Contour SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Submandibular Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 64 | 298 | 152.31 | 151.0 | 15.58 | 0.44 | 131.0 | 142.0 | 151.0 | 161.00 | 178.65 | 72 |
| Black | 548 | 122 | 201 | 156.78 | 156.0 | 12.65 | 0.54 | 138.0 | 148.0 | 156.0 | 165.50 | 178.90 | 45 |
| LatinX | 100 | 126 | 195 | 153.07 | 150.5 | 14.70 | 1.47 | 132.0 | 142.0 | 150.5 | 162.25 | 176.75 | 4 |
| Asian | 91 | 122 | 196 | 150.31 | 150.0 | 13.67 | 1.43 | 130.0 | 140.5 | 150.0 | 158.00 | 173.10 | 4 |
| Other | 21 | 132 | 165 | 150.00 | 150.5 | 9.91 | 2.16 | 133.5 | 146.0 | 150.5 | 156.75 | 162.00 | 5 |
| AIAN | 8 | 130 | 169 | 144.43 | 146.0 | 13.88 | 4.91 | 130.9 | 133.0 | 146.0 | 150.00 | 164.20 | 1 |
| PTNS | 5 | 139 | 181 | 155.80 | 154.0 | 15.83 | 7.08 | 140.6 | 147.0 | 154.0 | 158.00 | 176.40 | 0 |
| NHOPI | 4 | 143 | 151 | 147.67 | 149.0 | 4.16 | 2.08 | 143.6 | 146.0 | 149.0 | 150.00 | 150.80 | 1 |

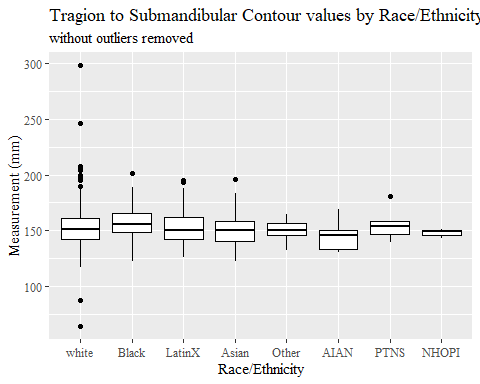
#%>% set\_header\_Cabels(values = list(TrSman\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, TrSman\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrSman\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Submandibular Contour values by Race/Ethnicity",  
 subtitle = "without outliers removed",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 132 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=TrSman\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Submandibular Contour values by Race/Ethnicity",  
 subtitle = "without outliers removed",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 132 rows containing non-finite values (stat\_boxplot).



#TrSman\_C gender sumstats  
TrSman\_Cgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(TrSman\_C, na.rm = TRUE),  
 max = max(TrSman\_C, na.rm = TRUE),  
 mean = mean(TrSman\_C, na.rm = TRUE),  
 mdn = median(TrSman\_C, na.rm = TRUE),  
 sd = sd(TrSman\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrSman\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrSman\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrSman\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrSman\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrSman\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrSman\_C)))  
  
TrSman\_Cgender\_sumstats <- TrSman\_Cgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(TrSman\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Submandibular Contour SumStats by Gender")

**Table** : Tragion to Submandibular Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 87 | 195 | 146.83 | 145 | 11.87 | 0.36 | 130.00 | 139.00 | 145 | 154.00 | 167.45 | 32 |
| Male | 939 | 64 | 298 | 161.52 | 160 | 14.00 | 0.46 | 142.90 | 153.00 | 160 | 169.00 | 184.00 | 100 |
| Non-binary or Other | 5 | 119 | 161 | 143.20 | 147 | 17.21 | 7.70 | 121.80 | 133.00 | 147 | 156.00 | 160.00 | 0 |
| Prefer not to say | 1 | 181 | 181 | 181.00 | 181 |  |  | 181.00 | 181.00 | 181 | 181.00 | 181.00 | 0 |
|  | 8 | 143 | 175 | 152.25 | 149 | 10.19 | 3.60 | 144.05 | 146.75 | 149 | 152.75 | 169.05 | 0 |

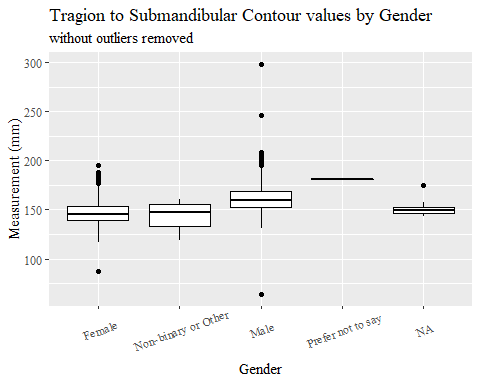
#%>% set\_header\_Cabels(values = list(TrSman\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrSman\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Submandibular Contour SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Submandibular Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 87 | 195 | 146.83 | 145 | 11.87 | 0.36 | 130.00 | 139.00 | 145 | 154.00 | 167.45 | 32 |
| Male | 939 | 64 | 298 | 161.52 | 160 | 14.00 | 0.46 | 142.90 | 153.00 | 160 | 169.00 | 184.00 | 100 |
| Non-binary or Other | 5 | 119 | 161 | 143.20 | 147 | 17.21 | 7.70 | 121.80 | 133.00 | 147 | 156.00 | 160.00 | 0 |
| Prefer not to say | 1 | 181 | 181 | 181.00 | 181 |  |  | 181.00 | 181.00 | 181 | 181.00 | 181.00 | 0 |
|  | 8 | 143 | 175 | 152.25 | 149 | 10.19 | 3.60 | 144.05 | 146.75 | 149 | 152.75 | 169.05 | 0 |

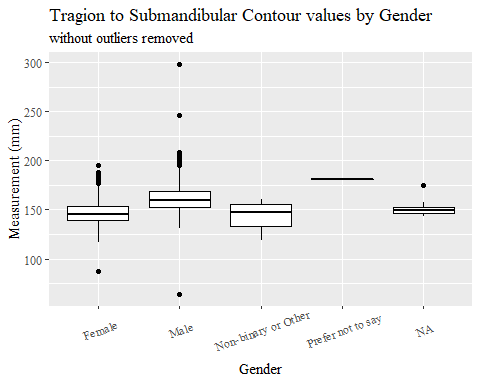
#%>% set\_header\_Cabels(values = list(TrSman\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, TrSman\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrSman\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Submandibular Contour values by Gender",  
 subtitle = "without outliers removed",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 132 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=TrSman\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Submandibular Contour values by Gender",  
 subtitle = "without outliers removed",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 132 rows containing non-finite values (stat\_boxplot).



#TrSman\_C age group sumstats  
TrSman\_Cage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(TrSman\_C, na.rm = TRUE),  
 max = max(TrSman\_C, na.rm = TRUE),  
 mean = mean(TrSman\_C, na.rm = TRUE),  
 mdn = median(TrSman\_C, na.rm = TRUE),  
 sd = sd(TrSman\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrSman\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrSman\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrSman\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrSman\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrSman\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrSman\_C)))  
  
TrSman\_Cage\_sumstats <- TrSman\_Cage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(TrSman\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Submandibular Contour SumStats by Age Group")

**Table** : Tragion to Submandibular Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 119 | 205 | 150.23 | 149 | 13.47 | 0.43 | 131.00 | 140.00 | 149 | 159.00 | 174.00 | 67 |
| 37-54 | 940 | 87 | 298 | 156.31 | 155 | 15.29 | 0.50 | 136.00 | 146.00 | 155 | 165.00 | 182.00 | 63 |
| 55-72 | 84 | 64 | 189 | 157.98 | 159 | 16.36 | 1.79 | 137.05 | 149.25 | 159 | 167.75 | 176.95 | 2 |
|  | 1 | 158 | 158 | 158.00 | 158 |  |  | 158.00 | 158.00 | 158 | 158.00 | 158.00 | 0 |

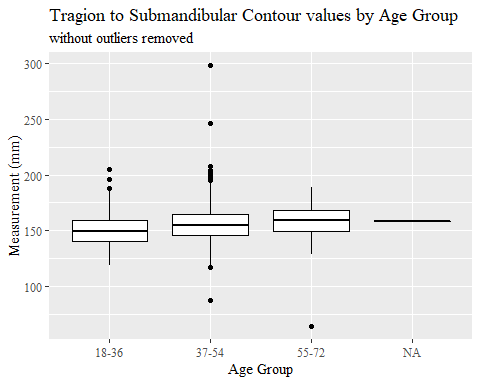
#%>% set\_header\_Cabels(values = list(TrSman\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrSman\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Submandibular Contour SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Submandibular Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 119 | 205 | 150.23 | 149 | 13.47 | 0.43 | 131.00 | 140.00 | 149 | 159.00 | 174.00 | 67 |
| 37-54 | 940 | 87 | 298 | 156.31 | 155 | 15.29 | 0.50 | 136.00 | 146.00 | 155 | 165.00 | 182.00 | 63 |
| 55-72 | 84 | 64 | 189 | 157.98 | 159 | 16.36 | 1.79 | 137.05 | 149.25 | 159 | 167.75 | 176.95 | 2 |
|  | 1 | 158 | 158 | 158.00 | 158 |  |  | 158.00 | 158.00 | 158 | 158.00 | 158.00 | 0 |

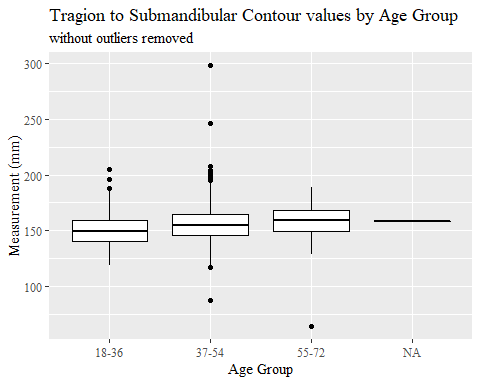
#%>% set\_header\_Cabels(values = list(TrSman\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, TrSman\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrSman\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Submandibular Contour values by Age Group",  
 subtitle = "without outliers removed",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 132 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=TrSman\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Submandibular Contour values by Age Group",  
 subtitle = "without outliers removed",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 132 rows containing non-finite values (stat\_boxplot).



TrSman\_C with outlier removed

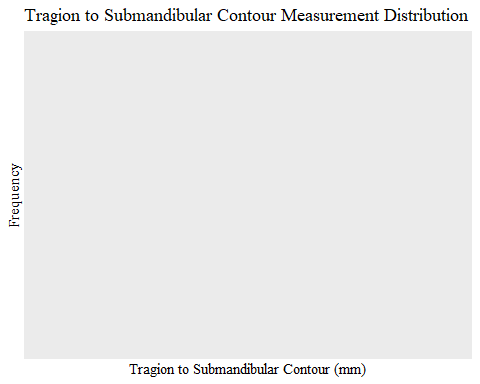
max(headscan\_full$TrSman\_C, na.rm = TRUE)

## [1] 298

#CODE TO REMOVE OUTLIER, use later to remove from headscan\_full  
TrSman\_C\_no\_out <- headscan\_full %>% mutate(TrSman\_C = replace(TrSman\_C, TrSman\_C>29, NA))

#histogram of all TrSman\_C values  
ggplot(data=TrSman\_C\_no\_out, aes(x=TrSman\_C))+  
 geom\_bar(stat="bin", bins=25, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Tragion to Submandibular Contour Measurement Distribution",  
 y="Frequency",  
 x="Tragion to Submandibular Contour (mm)")

## Warning: Removed 2017 rows containing non-finite values (stat\_bin).



#TrSman\_C race/eth sumstats  
TrSman\_Crace\_sumstats1 <- TrSman\_C\_no\_out %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(TrSman\_C, na.rm = TRUE),  
 max = max(TrSman\_C, na.rm = TRUE),  
 mean = mean(TrSman\_C, na.rm = TRUE),  
 mdn = median(TrSman\_C, na.rm = TRUE),  
 sd = sd(TrSman\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrSman\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrSman\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrSman\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrSman\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrSman\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrSman\_C)))

## Warning in min(TrSman\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSman\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSman\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSman\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSman\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSman\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSman\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSman\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf

## Warning in max(TrSman\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSman\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSman\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSman\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSman\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSman\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSman\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSman\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf

TrSman\_Crace\_sumstats1 <- TrSman\_Crace\_sumstats1 %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(TrSman\_Crace\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Submandibular Contour SumStats by Race/Ethnicity")

**Table** : Tragion to Submandibular Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1,240 |
| Black | 548 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 548 |
| LatinX | 100 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 100 |
| Asian | 91 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 91 |
| Other | 21 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 21 |
| AIAN | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |
| PTNS | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| NHOPI | 4 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 4 |

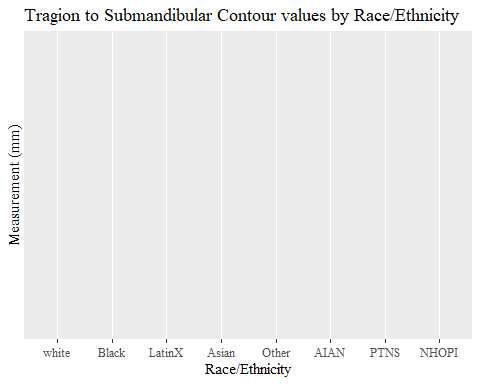
#%>% set\_header\_Cabels(values = list(TrSman\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrSman\_Crace\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Submandibular Contour SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Submandibular Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1,240 |
| Black | 548 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 548 |
| LatinX | 100 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 100 |
| Asian | 91 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 91 |
| Other | 21 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 21 |
| AIAN | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |
| PTNS | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| NHOPI | 4 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 4 |

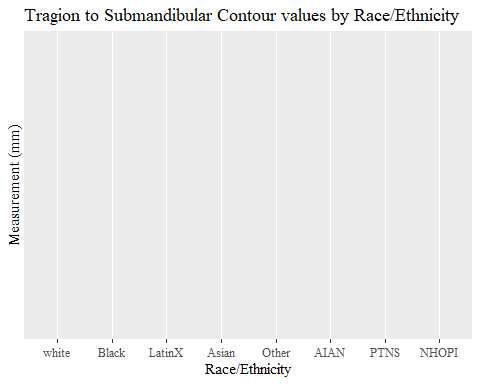
#%>% set\_header\_Cabels(values = list(TrSman\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
TrSman\_C\_no\_out %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, TrSman\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrSman\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Submandibular Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=TrSman\_C\_no\_out, aes(y=TrSman\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Submandibular Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#TrSman\_C gender sumstats  
TrSman\_Cgender\_sumstats1 <- TrSman\_C\_no\_out %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(TrSman\_C, na.rm = TRUE),  
 max = max(TrSman\_C, na.rm = TRUE),  
 mean = mean(TrSman\_C, na.rm = TRUE),  
 mdn = median(TrSman\_C, na.rm = TRUE),  
 sd = sd(TrSman\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrSman\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrSman\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrSman\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrSman\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrSman\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrSman\_C)))

## Warning in min(TrSman\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSman\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSman\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSman\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSman\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf

## Warning in max(TrSman\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSman\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSman\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSman\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSman\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf

TrSman\_Cgender\_sumstats1 <- TrSman\_Cgender\_sumstats1 %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(TrSman\_Cgender\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Submandibular Contour SumStats by Gender")

**Table** : Tragion to Submandibular Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1,064 |
| Male | 939 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 939 |
| Non-binary or Other | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| Prefer not to say | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |
|  | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |

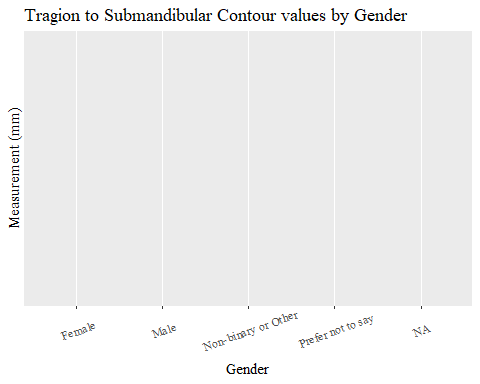
#%>% set\_header\_Cabels(values = list(TrSman\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrSman\_Cgender\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Submandibular Contour SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Submandibular Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1,064 |
| Male | 939 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 939 |
| Non-binary or Other | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| Prefer not to say | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |
|  | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |

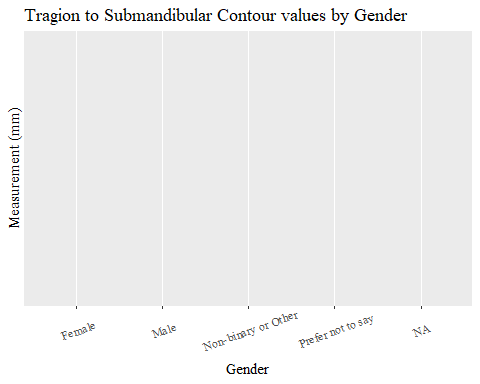
#%>% set\_header\_Cabels(values = list(TrSman\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
TrSman\_C\_no\_out %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, TrSman\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrSman\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Submandibular Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=TrSman\_C\_no\_out, aes(y=TrSman\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Submandibular Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#TrSman\_C age group sumstats  
TrSman\_Cage\_sumstats1 <- TrSman\_C\_no\_out %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(TrSman\_C, na.rm = TRUE),  
 max = max(TrSman\_C, na.rm = TRUE),  
 mean = mean(TrSman\_C, na.rm = TRUE),  
 mdn = median(TrSman\_C, na.rm = TRUE),  
 sd = sd(TrSman\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrSman\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrSman\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrSman\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrSman\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrSman\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrSman\_C)))

## Warning in min(TrSman\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSman\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSman\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSman\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf

## Warning in max(TrSman\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSman\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSman\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSman\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf

TrSman\_Cage\_sumstats1 <- TrSman\_Cage\_sumstats1 %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(TrSman\_Cage\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Submandibular Contour SumStats by Age Group")

**Table** : Tragion to Submandibular Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 992 |
| 37-54 | 940 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 940 |
| 55-72 | 84 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 84 |
|  | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |

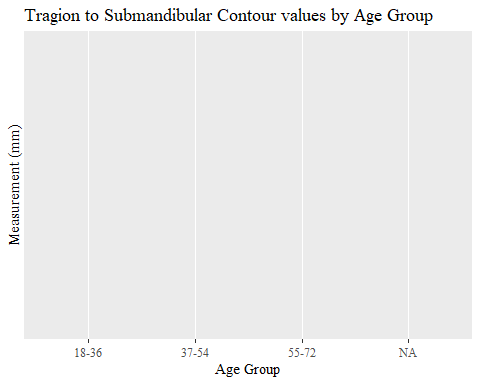
#%>% set\_header\_Cabels(values = list(TrSman\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrSman\_Cage\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Submandibular Contour SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Submandibular Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 992 |
| 37-54 | 940 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 940 |
| 55-72 | 84 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 84 |
|  | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |

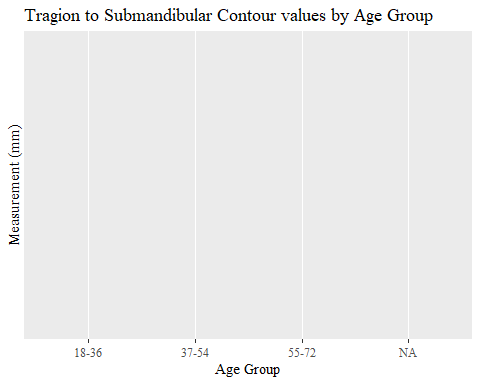
#%>% set\_header\_Cabels(values = list(TrSman\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
TrSman\_C\_no\_out %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, TrSman\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrSman\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Submandibular Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=TrSman\_C\_no\_out, aes(y=TrSman\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Submandibular Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

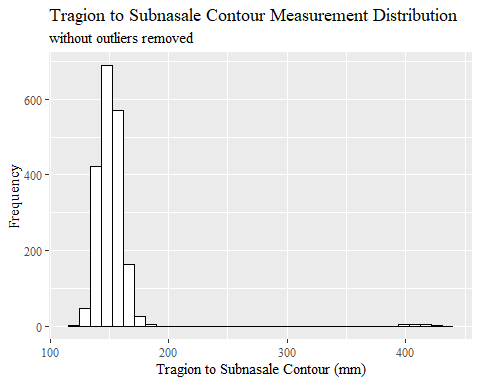
## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



TrSnas\_C without outliers removed

#histogram of all TrSnas\_C values  
ggplot(data=headscan\_full, aes(x=TrSnas\_C))+  
 geom\_bar(stat="bin", bins=35, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Tragion to Subnasale Contour Measurement Distribution",  
 subtitle = "without outliers removed",  
 y="Frequency",  
 x="Tragion to Subnasale Contour (mm)")

## Warning: Removed 71 rows containing non-finite values (stat\_bin).



#TrSnas\_C race/eth sumstats  
TrSnas\_Crace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(TrSnas\_C, na.rm = TRUE),  
 max = max(TrSnas\_C, na.rm = TRUE),  
 mean = mean(TrSnas\_C, na.rm = TRUE),  
 mdn = median(TrSnas\_C, na.rm = TRUE),  
 sd = sd(TrSnas\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrSnas\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrSnas\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrSnas\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrSnas\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrSnas\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrSnas\_C)))   
  
TrSnas\_Crace\_sumstats <- TrSnas\_Crace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(TrSnas\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Subnasale Contour SumStats by Race/Ethnicity")

**Table** : Tragion to Subnasale Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 122 | 437 | 152.64 | 148.0 | 34.19 | 0.97 | 135.10 | 142.00 | 148.0 | 155.00 | 166.00 | 57 |
| Black | 548 | 131 | 181 | 154.98 | 155.0 | 8.34 | 0.36 | 142.00 | 149.00 | 155.0 | 160.00 | 169.00 | 8 |
| LatinX | 100 | 131 | 169 | 149.37 | 148.0 | 8.21 | 0.82 | 136.85 | 144.00 | 148.0 | 156.75 | 161.15 | 2 |
| Asian | 91 | 129 | 172 | 148.49 | 147.0 | 9.64 | 1.01 | 135.40 | 142.00 | 147.0 | 154.00 | 166.60 | 2 |
| Other | 21 | 135 | 172 | 151.55 | 150.0 | 8.83 | 1.93 | 140.70 | 145.75 | 150.0 | 158.50 | 163.45 | 1 |
| AIAN | 8 | 138 | 160 | 146.00 | 144.0 | 6.90 | 2.44 | 139.50 | 143.00 | 144.0 | 147.00 | 156.40 | 1 |
| PTNS | 5 | 142 | 162 | 152.80 | 152.0 | 7.69 | 3.44 | 143.60 | 150.00 | 152.0 | 158.00 | 161.20 | 0 |
| NHOPI | 4 | 150 | 155 | 152.00 | 151.5 | 2.45 | 1.22 | 150.00 | 150.00 | 151.5 | 153.50 | 154.70 | 0 |

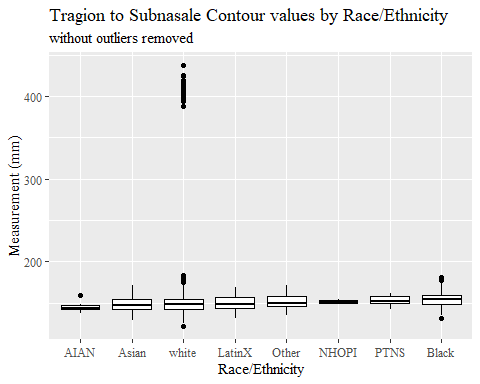
#%>% set\_header\_Cabels(values = list(TrSnas\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrSnas\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Subnasale Contour SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Subnasale Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 122 | 437 | 152.64 | 148.0 | 34.19 | 0.97 | 135.10 | 142.00 | 148.0 | 155.00 | 166.00 | 57 |
| Black | 548 | 131 | 181 | 154.98 | 155.0 | 8.34 | 0.36 | 142.00 | 149.00 | 155.0 | 160.00 | 169.00 | 8 |
| LatinX | 100 | 131 | 169 | 149.37 | 148.0 | 8.21 | 0.82 | 136.85 | 144.00 | 148.0 | 156.75 | 161.15 | 2 |
| Asian | 91 | 129 | 172 | 148.49 | 147.0 | 9.64 | 1.01 | 135.40 | 142.00 | 147.0 | 154.00 | 166.60 | 2 |
| Other | 21 | 135 | 172 | 151.55 | 150.0 | 8.83 | 1.93 | 140.70 | 145.75 | 150.0 | 158.50 | 163.45 | 1 |
| AIAN | 8 | 138 | 160 | 146.00 | 144.0 | 6.90 | 2.44 | 139.50 | 143.00 | 144.0 | 147.00 | 156.40 | 1 |
| PTNS | 5 | 142 | 162 | 152.80 | 152.0 | 7.69 | 3.44 | 143.60 | 150.00 | 152.0 | 158.00 | 161.20 | 0 |
| NHOPI | 4 | 150 | 155 | 152.00 | 151.5 | 2.45 | 1.22 | 150.00 | 150.00 | 151.5 | 153.50 | 154.70 | 0 |

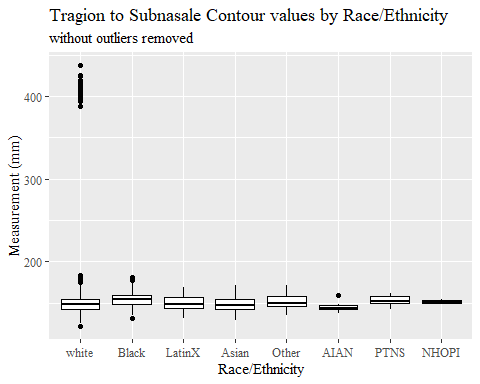
#%>% set\_header\_Cabels(values = list(TrSnas\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, TrSnas\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrSnas\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Subnasale Contour values by Race/Ethnicity",  
 subtitle = "without outliers removed",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 71 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=TrSnas\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Subnasale Contour values by Race/Ethnicity",  
 subtitle = "without outliers removed",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 71 rows containing non-finite values (stat\_boxplot).



#TrSnas\_C gender sumstats  
TrSnas\_Cgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(TrSnas\_C, na.rm = TRUE),  
 max = max(TrSnas\_C, na.rm = TRUE),  
 mean = mean(TrSnas\_C, na.rm = TRUE),  
 mdn = median(TrSnas\_C, na.rm = TRUE),  
 sd = sd(TrSnas\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrSnas\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrSnas\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrSnas\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrSnas\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrSnas\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrSnas\_C)))  
  
TrSnas\_Cgender\_sumstats <- TrSnas\_Cgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(TrSnas\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Subnasale Contour SumStats by Gender")

**Table** : Tragion to Subnasale Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 122 | 420 | 149.03 | 145 | 27.97 | 0.86 | 135.0 | 140 | 145 | 151.00 | 162.00 | 36 |
| Male | 939 | 127 | 437 | 157.38 | 155 | 25.86 | 0.84 | 142.0 | 149 | 155 | 161.00 | 170.00 | 35 |
| Non-binary or Other | 5 | 136 | 151 | 146.00 | 147 | 5.96 | 2.66 | 138.0 | 146 | 147 | 150.00 | 150.80 | 0 |
| Prefer not to say | 1 | 152 | 152 | 152.00 | 152 |  |  | 152.0 | 152 | 152 | 152.00 | 152.00 | 0 |
|  | 8 | 141 | 157 | 148.12 | 149 | 5.46 | 1.93 | 141.7 | 143 | 149 | 151.25 | 155.25 | 0 |

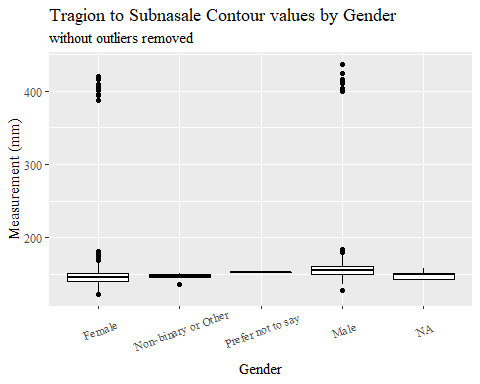
#%>% set\_header\_Cabels(values = list(TrSnas\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrSnas\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Subnasale Contour SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Subnasale Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 122 | 420 | 149.03 | 145 | 27.97 | 0.86 | 135.0 | 140 | 145 | 151.00 | 162.00 | 36 |
| Male | 939 | 127 | 437 | 157.38 | 155 | 25.86 | 0.84 | 142.0 | 149 | 155 | 161.00 | 170.00 | 35 |
| Non-binary or Other | 5 | 136 | 151 | 146.00 | 147 | 5.96 | 2.66 | 138.0 | 146 | 147 | 150.00 | 150.80 | 0 |
| Prefer not to say | 1 | 152 | 152 | 152.00 | 152 |  |  | 152.0 | 152 | 152 | 152.00 | 152.00 | 0 |
|  | 8 | 141 | 157 | 148.12 | 149 | 5.46 | 1.93 | 141.7 | 143 | 149 | 151.25 | 155.25 | 0 |

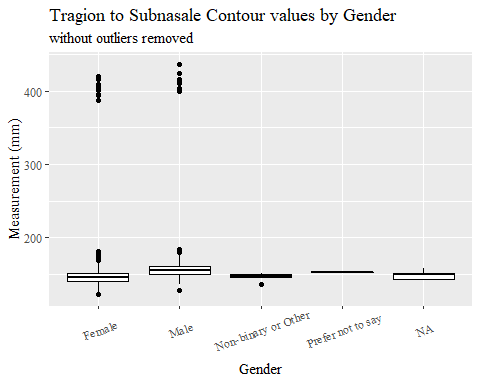
#%>% set\_header\_Cabels(values = list(TrSnas\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, TrSnas\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrSnas\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Subnasale Contour values by Gender",  
 subtitle = "without outliers removed",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 71 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=TrSnas\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Subnasale Contour values by Gender",  
 subtitle = "without outliers removed",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 71 rows containing non-finite values (stat\_boxplot).



#TrSnas\_C age group sumstats  
TrSnas\_Cage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(TrSnas\_C, na.rm = TRUE),  
 max = max(TrSnas\_C, na.rm = TRUE),  
 mean = mean(TrSnas\_C, na.rm = TRUE),  
 mdn = median(TrSnas\_C, na.rm = TRUE),  
 sd = sd(TrSnas\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrSnas\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrSnas\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrSnas\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrSnas\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrSnas\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrSnas\_C)))  
  
TrSnas\_Cage\_sumstats <- TrSnas\_Cage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(TrSnas\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Subnasale Contour SumStats by Age Group")

**Table** : Tragion to Subnasale Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 129 | 437 | 152.23 | 149 | 27.36 | 0.87 | 136 | 143 | 149 | 156 | 167 | 31 |
| 37-54 | 940 | 122 | 411 | 153.21 | 150 | 26.68 | 0.87 | 136 | 144 | 150 | 157 | 168 | 37 |
| 55-72 | 84 | 130 | 425 | 157.56 | 155 | 31.33 | 3.42 | 142 | 147 | 155 | 160 | 170 | 3 |
|  | 1 | 147 | 147 | 147.00 | 147 |  |  | 147 | 147 | 147 | 147 | 147 | 0 |

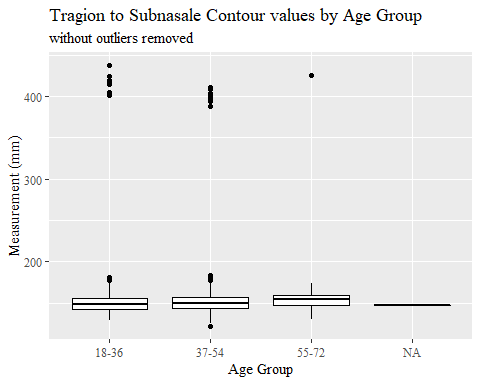
#%>% set\_header\_Cabels(values = list(TrSnas\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrSnas\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Subnasale Contour SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Subnasale Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 129 | 437 | 152.23 | 149 | 27.36 | 0.87 | 136 | 143 | 149 | 156 | 167 | 31 |
| 37-54 | 940 | 122 | 411 | 153.21 | 150 | 26.68 | 0.87 | 136 | 144 | 150 | 157 | 168 | 37 |
| 55-72 | 84 | 130 | 425 | 157.56 | 155 | 31.33 | 3.42 | 142 | 147 | 155 | 160 | 170 | 3 |
|  | 1 | 147 | 147 | 147.00 | 147 |  |  | 147 | 147 | 147 | 147 | 147 | 0 |

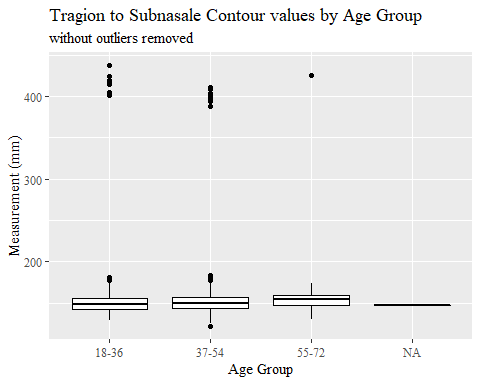
#%>% set\_header\_Cabels(values = list(TrSnas\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, TrSnas\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrSnas\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Subnasale Contour values by Age Group",  
 subtitle = "without outliers removed",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 71 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=TrSnas\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Subnasale Contour values by Age Group",  
 subtitle = "without outliers removed",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 71 rows containing non-finite values (stat\_boxplot).

 TrSnas\_C with outliers removed

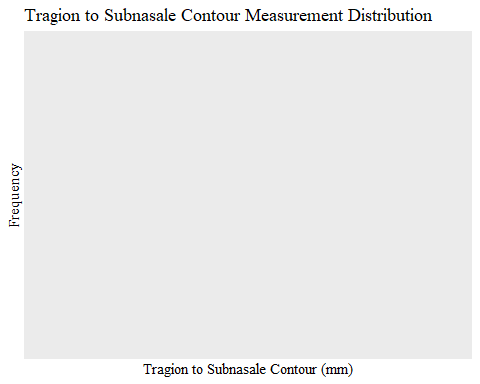
max(headscan\_full$TrSnas\_C, na.rm = TRUE)

## [1] 437

#CODE TO REMOVE OUTLIER, use later to remove from headscan\_full  
TrSnas\_C\_no\_out <- headscan\_full %>% mutate(TrSnas\_C = replace(TrSnas\_C, TrSnas\_C>30, NA))

#histogram of all TrSnas\_C values  
ggplot(data=TrSnas\_C\_no\_out, aes(x=TrSnas\_C))+  
 geom\_bar(stat="bin", bins=20, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Tragion to Subnasale Contour Measurement Distribution",  
 y="Frequency",  
 x="Tragion to Subnasale Contour (mm)")

## Warning: Removed 2017 rows containing non-finite values (stat\_bin).



#TrSnas\_C race/eth sumstats  
TrSnas\_Crace\_sumstats1 <- TrSnas\_C\_no\_out %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(TrSnas\_C, na.rm = TRUE),  
 max = max(TrSnas\_C, na.rm = TRUE),  
 mean = mean(TrSnas\_C, na.rm = TRUE),  
 mdn = median(TrSnas\_C, na.rm = TRUE),  
 sd = sd(TrSnas\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrSnas\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrSnas\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrSnas\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrSnas\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrSnas\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrSnas\_C)))

## Warning in min(TrSnas\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSnas\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSnas\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSnas\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSnas\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSnas\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSnas\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSnas\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf

## Warning in max(TrSnas\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSnas\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSnas\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSnas\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSnas\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSnas\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSnas\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSnas\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf

TrSnas\_Crace\_sumstats1 <- TrSnas\_Crace\_sumstats1 %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(TrSnas\_Crace\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Subnasale Contour SumStats by Race/Ethnicity")

**Table** : Tragion to Subnasale Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1,240 |
| Black | 548 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 548 |
| LatinX | 100 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 100 |
| Asian | 91 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 91 |
| Other | 21 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 21 |
| AIAN | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |
| PTNS | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| NHOPI | 4 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 4 |

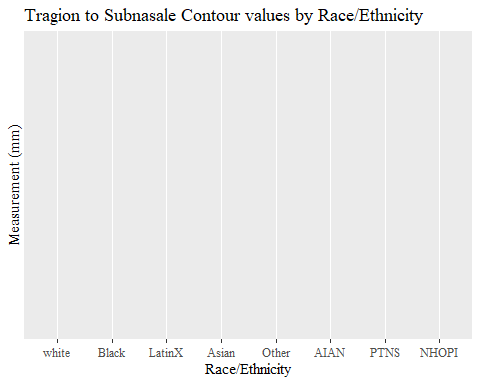
#%>% set\_header\_Cabels(values = list(TrSnas\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrSnas\_Crace\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Subnasale Contour SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Subnasale Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1,240 |
| Black | 548 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 548 |
| LatinX | 100 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 100 |
| Asian | 91 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 91 |
| Other | 21 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 21 |
| AIAN | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |
| PTNS | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| NHOPI | 4 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 4 |

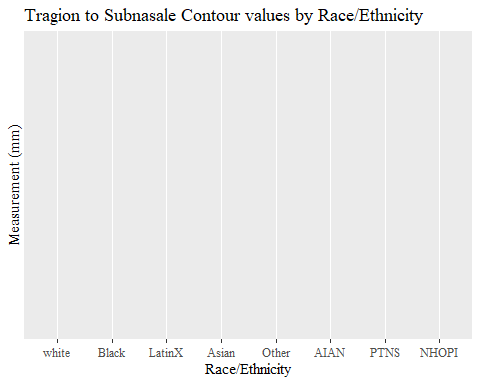
#%>% set\_header\_Cabels(values = list(TrSnas\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
TrSnas\_C\_no\_out %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, TrSnas\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrSnas\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Subnasale Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=TrSnas\_C\_no\_out, aes(y=TrSnas\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Subnasale Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#TrSnas\_C gender sumstats  
TrSnas\_Cgender\_sumstats1 <- TrSnas\_C\_no\_out %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(TrSnas\_C, na.rm = TRUE),  
 max = max(TrSnas\_C, na.rm = TRUE),  
 mean = mean(TrSnas\_C, na.rm = TRUE),  
 mdn = median(TrSnas\_C, na.rm = TRUE),  
 sd = sd(TrSnas\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrSnas\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrSnas\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrSnas\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrSnas\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrSnas\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrSnas\_C)))

## Warning in min(TrSnas\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSnas\_C, na.rm = TRUE): no non-missing arguments to min;  
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## returning Inf  
  
## Warning in min(TrSnas\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSnas\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf

## Warning in max(TrSnas\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSnas\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSnas\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSnas\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSnas\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf

TrSnas\_Cgender\_sumstats1 <- TrSnas\_Cgender\_sumstats1 %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(TrSnas\_Cgender\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Subnasale Contour SumStats by Gender")

**Table** : Tragion to Subnasale Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1,064 |
| Male | 939 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 939 |
| Non-binary or Other | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| Prefer not to say | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |
|  | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |

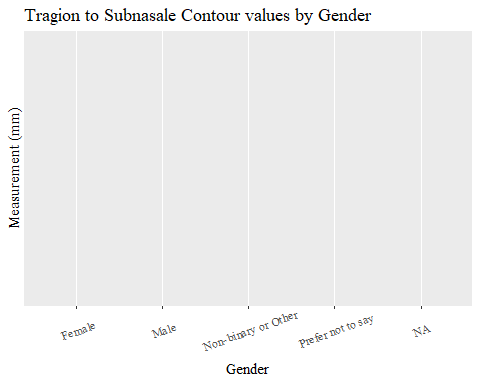
#%>% set\_header\_Cabels(values = list(TrSnas\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrSnas\_Cgender\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Subnasale Contour SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Subnasale Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1,064 |
| Male | 939 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 939 |
| Non-binary or Other | 5 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 5 |
| Prefer not to say | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |
|  | 8 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 8 |

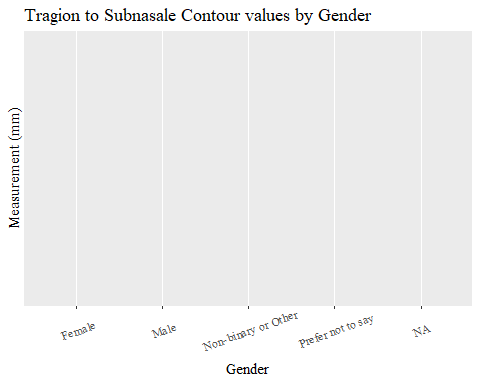
#%>% set\_header\_Cabels(values = list(TrSnas\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
TrSnas\_C\_no\_out %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, TrSnas\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrSnas\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Subnasale Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=TrSnas\_C\_no\_out, aes(y=TrSnas\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Subnasale Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#TrSnas\_C age group sumstats  
TrSnas\_Cage\_sumstats1 <- TrSnas\_C\_no\_out %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(TrSnas\_C, na.rm = TRUE),  
 max = max(TrSnas\_C, na.rm = TRUE),  
 mean = mean(TrSnas\_C, na.rm = TRUE),  
 mdn = median(TrSnas\_C, na.rm = TRUE),  
 sd = sd(TrSnas\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrSnas\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrSnas\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrSnas\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrSnas\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrSnas\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrSnas\_C)))

## Warning in min(TrSnas\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSnas\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSnas\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf  
  
## Warning in min(TrSnas\_C, na.rm = TRUE): no non-missing arguments to min;  
## returning Inf

## Warning in max(TrSnas\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSnas\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSnas\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf  
  
## Warning in max(TrSnas\_C, na.rm = TRUE): no non-missing arguments to max;  
## returning -Inf

TrSnas\_Cage\_sumstats1 <- TrSnas\_Cage\_sumstats1 %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(TrSnas\_Cage\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Subnasale Contour SumStats by Age Group")

**Table** : Tragion to Subnasale Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 992 |
| 37-54 | 940 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 940 |
| 55-72 | 84 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 84 |
|  | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |

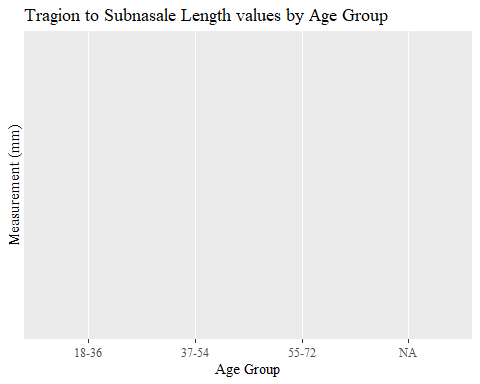
#%>% set\_header\_Cabels(values = list(TrSnas\_C = "Subnasale/SubnasaleCont"))  
  
#Autofit Width Table TNR  
flextable(TrSnas\_Cage\_sumstats1) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Subnasale Contour SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Subnasale Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 992 |
| 37-54 | 940 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 940 |
| 55-72 | 84 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 84 |
|  | 1 | Inf | -Inf |  |  |  |  |  |  |  |  |  | 1 |

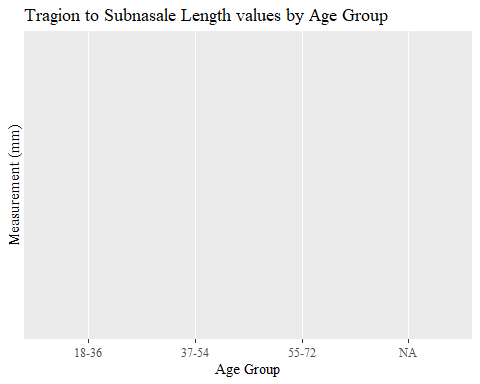
#%>% set\_header\_Cabels(values = list(TrSnas\_C = "Subnasale/SubnasaleCont"))  
  
#boxplot reorderd by median  
TrSnas\_C\_no\_out %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, TrSnas\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrSnas\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Subnasale Length values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=TrSnas\_C\_no\_out, aes(y=TrSnas\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Subnasale Length values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

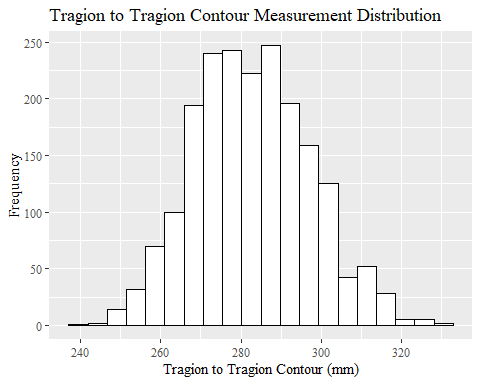
## Warning: Removed 2017 rows containing non-finite values (stat\_boxplot).



TrTr\_C

#histogram of all TrTr\_C values  
ggplot(data=headscan\_full, aes(x=TrTr\_C ))+  
 geom\_bar(stat="bin", bins=20, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Tragion to Tragion Contour Measurement Distribution",  
 y="Frequency",  
 x="Tragion to Tragion Contour (mm)")

## Warning: Removed 38 rows containing non-finite values (stat\_bin).



#TrTr\_C race/eth sumstats  
TrTr\_Crace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(TrTr\_C, na.rm = TRUE),  
 max = max(TrTr\_C, na.rm = TRUE),  
 mean = mean(TrTr\_C, na.rm = TRUE),  
 mdn = median(TrTr\_C, na.rm = TRUE),  
 sd = sd(TrTr\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrTr\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrTr\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrTr\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrTr\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrTr\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrTr\_C)))   
  
TrTr\_Crace\_sumstats <- TrTr\_Crace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(TrTr\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Tragion Contour SumStats by Race/Ethnicity")

**Table** : Tragion to Tragion Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 241 | 332 | 280.65 | 280.0 | 13.80 | 0.39 | 260.00 | 271.00 | 280.0 | 290.00 | 304.00 | 23 |
| Black | 548 | 248 | 329 | 288.06 | 288.0 | 14.15 | 0.60 | 265.00 | 278.00 | 288.0 | 297.00 | 312.00 | 9 |
| LatinX | 100 | 250 | 317 | 280.62 | 279.0 | 14.11 | 1.41 | 257.90 | 272.50 | 279.0 | 290.00 | 302.50 | 1 |
| Asian | 91 | 243 | 325 | 281.16 | 278.0 | 15.20 | 1.59 | 261.00 | 271.00 | 278.0 | 291.25 | 309.00 | 3 |
| Other | 21 | 253 | 314 | 282.80 | 283.5 | 14.59 | 3.18 | 264.40 | 275.50 | 283.5 | 289.75 | 303.55 | 1 |
| AIAN | 8 | 265 | 296 | 273.00 | 267.0 | 11.21 | 3.96 | 265.30 | 266.50 | 267.0 | 275.00 | 290.90 | 1 |
| PTNS | 5 | 269 | 298 | 283.80 | 285.0 | 12.68 | 5.67 | 269.80 | 273.00 | 285.0 | 294.00 | 297.20 | 0 |
| NHOPI | 4 | 283 | 297 | 289.25 | 288.5 | 6.85 | 3.42 | 283.15 | 283.75 | 288.5 | 294.00 | 296.40 | 0 |

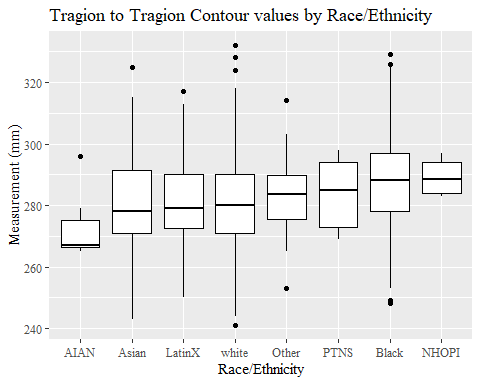
#%>% set\_header\_Cabels(values = list(TrTr\_C = "Tragion/TragionCont"))  
  
#Autofit Width Table TNR  
flextable(TrTr\_Crace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Tragion Contour SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Tragion Contour SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 241 | 332 | 280.65 | 280.0 | 13.80 | 0.39 | 260.00 | 271.00 | 280.0 | 290.00 | 304.00 | 23 |
| Black | 548 | 248 | 329 | 288.06 | 288.0 | 14.15 | 0.60 | 265.00 | 278.00 | 288.0 | 297.00 | 312.00 | 9 |
| LatinX | 100 | 250 | 317 | 280.62 | 279.0 | 14.11 | 1.41 | 257.90 | 272.50 | 279.0 | 290.00 | 302.50 | 1 |
| Asian | 91 | 243 | 325 | 281.16 | 278.0 | 15.20 | 1.59 | 261.00 | 271.00 | 278.0 | 291.25 | 309.00 | 3 |
| Other | 21 | 253 | 314 | 282.80 | 283.5 | 14.59 | 3.18 | 264.40 | 275.50 | 283.5 | 289.75 | 303.55 | 1 |
| AIAN | 8 | 265 | 296 | 273.00 | 267.0 | 11.21 | 3.96 | 265.30 | 266.50 | 267.0 | 275.00 | 290.90 | 1 |
| PTNS | 5 | 269 | 298 | 283.80 | 285.0 | 12.68 | 5.67 | 269.80 | 273.00 | 285.0 | 294.00 | 297.20 | 0 |
| NHOPI | 4 | 283 | 297 | 289.25 | 288.5 | 6.85 | 3.42 | 283.15 | 283.75 | 288.5 | 294.00 | 296.40 | 0 |

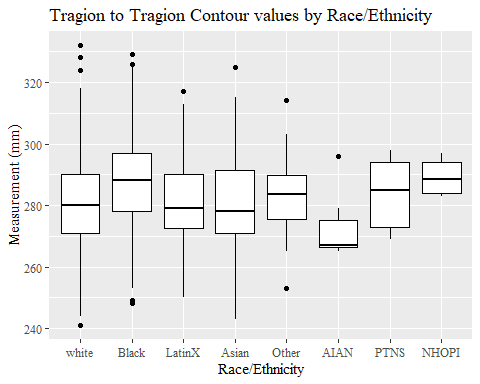
#%>% set\_header\_Cabels(values = list(TrTr\_C = "Tragion/TragionCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, TrTr\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrTr\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Tragion Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 38 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=TrTr\_C, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Tragion Contour values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 38 rows containing non-finite values (stat\_boxplot).



#TrTr\_C gender sumstats  
TrTr\_Cgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(TrTr\_C, na.rm = TRUE),  
 max = max(TrTr\_C, na.rm = TRUE),  
 mean = mean(TrTr\_C, na.rm = TRUE),  
 mdn = median(TrTr\_C, na.rm = TRUE),  
 sd = sd(TrTr\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrTr\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrTr\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrTr\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrTr\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrTr\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrTr\_C)))  
  
TrTr\_Cgender\_sumstats <- TrTr\_Cgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(TrTr\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Tragion Contour SumStats by Gender")

**Table** : Tragion to Tragion Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 241 | 329 | 275.78 | 275.0 | 12.33 | 0.38 | 257.0 | 267.0 | 275.0 | 283 | 297.15 | 26 |
| Male | 939 | 254 | 332 | 290.53 | 290.0 | 12.33 | 0.40 | 271.0 | 282.0 | 290.0 | 299 | 312.00 | 12 |
| Non-binary or Other | 5 | 260 | 288 | 277.20 | 278.0 | 10.45 | 4.67 | 263.6 | 278.0 | 278.0 | 282 | 286.80 | 0 |
| Prefer not to say | 1 | 294 | 294 | 294.00 | 294.0 |  |  | 294.0 | 294.0 | 294.0 | 294 | 294.00 | 0 |
|  | 8 | 270 | 302 | 279.12 | 277.5 | 10.52 | 3.72 | 270.0 | 271.5 | 277.5 | 281 | 295.70 | 0 |

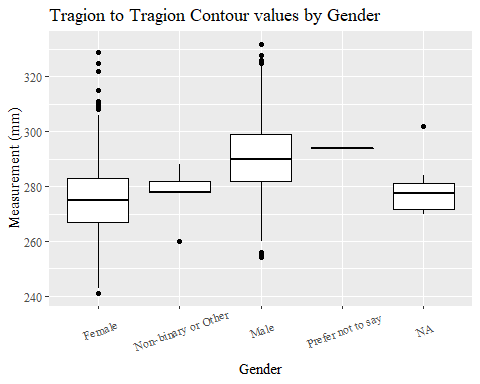
#%>% set\_header\_Cabels(values = list(TrTr\_C = "Tragion/TragionCont"))  
  
#Autofit Width Table TNR  
flextable(TrTr\_Cgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Tragion Contour SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Tragion Contour SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 241 | 329 | 275.78 | 275.0 | 12.33 | 0.38 | 257.0 | 267.0 | 275.0 | 283 | 297.15 | 26 |
| Male | 939 | 254 | 332 | 290.53 | 290.0 | 12.33 | 0.40 | 271.0 | 282.0 | 290.0 | 299 | 312.00 | 12 |
| Non-binary or Other | 5 | 260 | 288 | 277.20 | 278.0 | 10.45 | 4.67 | 263.6 | 278.0 | 278.0 | 282 | 286.80 | 0 |
| Prefer not to say | 1 | 294 | 294 | 294.00 | 294.0 |  |  | 294.0 | 294.0 | 294.0 | 294 | 294.00 | 0 |
|  | 8 | 270 | 302 | 279.12 | 277.5 | 10.52 | 3.72 | 270.0 | 271.5 | 277.5 | 281 | 295.70 | 0 |

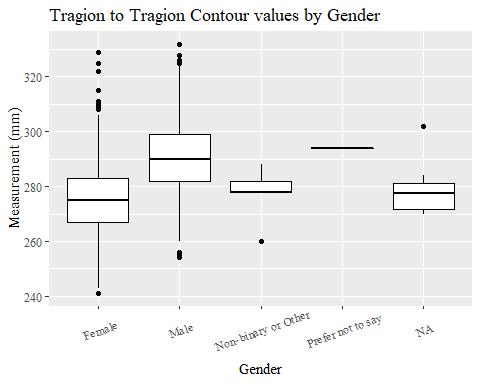
#%>% set\_header\_Cabels(values = list(TrTr\_C = "Tragion/TragionCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, TrTr\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrTr\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Tragion Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 38 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=TrTr\_C, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Tragion Contour values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 38 rows containing non-finite values (stat\_boxplot).



#TrTr\_C age group sumstats  
TrTr\_Cage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(TrTr\_C, na.rm = TRUE),  
 max = max(TrTr\_C, na.rm = TRUE),  
 mean = mean(TrTr\_C, na.rm = TRUE),  
 mdn = median(TrTr\_C, na.rm = TRUE),  
 sd = sd(TrTr\_C, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrTr\_C, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrTr\_C, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrTr\_C, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrTr\_C, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrTr\_C, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrTr\_C)))  
  
TrTr\_Cage\_sumstats <- TrTr\_Cage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(TrTr\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Tragion Contour SumStats by Age Group")

**Table** : Tragion to Tragion Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 241 | 326 | 282.23 | 282 | 14.38 | 0.46 | 260.0 | 272 | 282 | 292.0 | 307.0 | 18 |
| 37-54 | 940 | 244 | 332 | 282.78 | 282 | 14.33 | 0.47 | 261.0 | 272 | 282 | 293.0 | 308.0 | 19 |
| 55-72 | 84 | 255 | 317 | 287.63 | 288 | 13.26 | 1.45 | 266.1 | 278 | 288 | 297.5 | 308.9 | 1 |
|  | 1 | 283 | 283 | 283.00 | 283 |  |  | 283.0 | 283 | 283 | 283.0 | 283.0 | 0 |

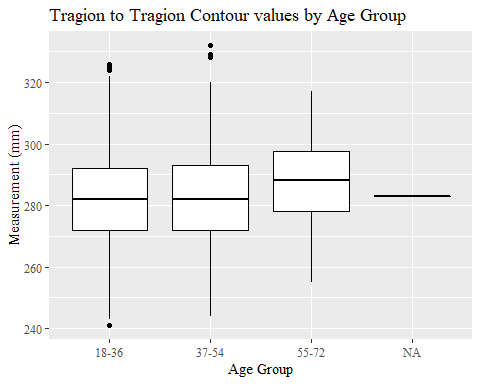
#%>% set\_header\_Cabels(values = list(TrTr\_C = "Tragion/TragionCont"))  
  
#Autofit Width Table TNR  
flextable(TrTr\_Cage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Tragion Contour SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Tragion Contour SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 241 | 326 | 282.23 | 282 | 14.38 | 0.46 | 260.0 | 272 | 282 | 292.0 | 307.0 | 18 |
| 37-54 | 940 | 244 | 332 | 282.78 | 282 | 14.33 | 0.47 | 261.0 | 272 | 282 | 293.0 | 308.0 | 19 |
| 55-72 | 84 | 255 | 317 | 287.63 | 288 | 13.26 | 1.45 | 266.1 | 278 | 288 | 297.5 | 308.9 | 1 |
|  | 1 | 283 | 283 | 283.00 | 283 |  |  | 283.0 | 283 | 283 | 283.0 | 283.0 | 0 |

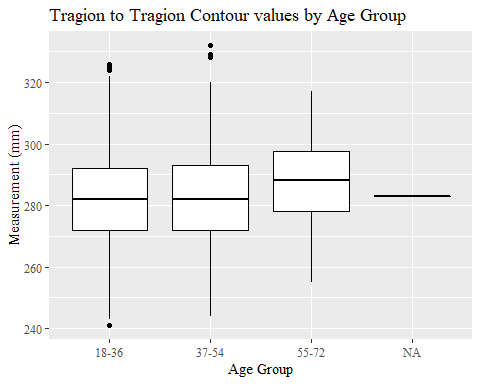
#%>% set\_header\_Cabels(values = list(TrTr\_C = "Tragion/TragionCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, TrTr\_C, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrTr\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Tragion Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 38 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=TrTr\_C, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Tragion Contour values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

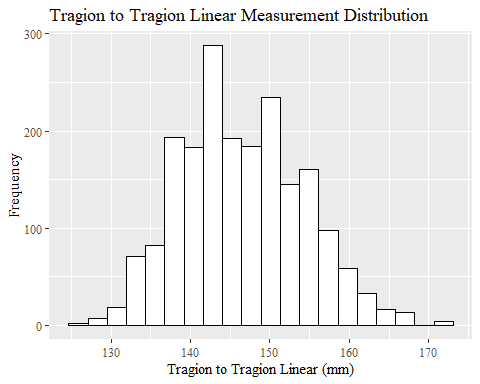
## Warning: Removed 38 rows containing non-finite values (stat\_boxplot).



TrTr\_L

#histogram of all TrTr\_L values  
ggplot(data=headscan\_full, aes(x=TrTr\_L))+  
 geom\_bar(stat="bin", bins=20, color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 labs(title="Tragion to Tragion Linear Measurement Distribution",  
 y="Frequency",  
 x="Tragion to Tragion Linear (mm)")

## Warning: Removed 34 rows containing non-finite values (stat\_bin).



#TrTr\_L race/eth sumstats  
TrTr\_Lrace\_sumstats <- headscan\_full %>%   
 group\_by(race\_eth) %>%   
 summarise(n = n(),  
 min = min(TrTr\_L, na.rm = TRUE),  
 max = max(TrTr\_L, na.rm = TRUE),  
 mean = mean(TrTr\_L, na.rm = TRUE),  
 mdn = median(TrTr\_L, na.rm = TRUE),  
 sd = sd(TrTr\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrTr\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrTr\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrTr\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrTr\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrTr\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrTr\_L)))   
  
TrTr\_Lrace\_sumstats <- TrTr\_Lrace\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(TrTr\_Lrace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Tragion Linear SumStats by Race/Ethnicity")

**Table** : Tragion to Tragion Linear SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 127 | 173 | 145.50 | 145.0 | 7.57 | 0.22 | 134.0 | 140.0 | 145.0 | 151.00 | 158.00 | 20 |
| Black | 548 | 130 | 172 | 148.47 | 148.0 | 7.57 | 0.32 | 137.0 | 143.0 | 148.0 | 154.00 | 161.10 | 9 |
| LatinX | 100 | 132 | 171 | 146.90 | 147.0 | 7.54 | 0.75 | 135.0 | 141.5 | 147.0 | 151.00 | 158.10 | 1 |
| Asian | 91 | 129 | 167 | 147.56 | 148.0 | 8.67 | 0.91 | 135.4 | 140.0 | 148.0 | 153.00 | 161.00 | 2 |
| Other | 21 | 138 | 159 | 147.45 | 146.0 | 6.56 | 1.43 | 139.9 | 142.0 | 146.0 | 151.75 | 158.05 | 1 |
| AIAN | 8 | 132 | 157 | 141.86 | 140.0 | 7.99 | 2.82 | 133.8 | 138.5 | 140.0 | 143.50 | 154.00 | 1 |
| PTNS | 5 | 137 | 156 | 147.80 | 153.0 | 8.64 | 3.87 | 137.6 | 140.0 | 153.0 | 153.00 | 155.40 | 0 |
| NHOPI | 4 | 153 | 160 | 156.50 | 156.5 | 3.11 | 1.55 | 153.3 | 154.5 | 156.5 | 158.50 | 159.70 | 0 |

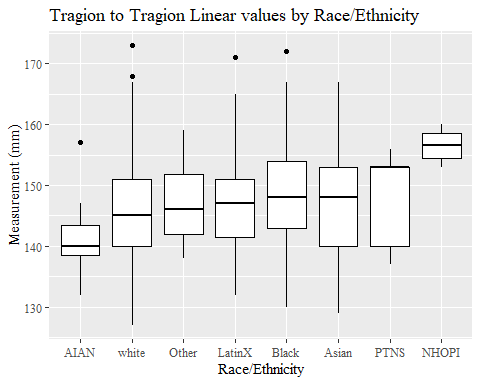
#%>% set\_header\_Labels(values = list(TrTr\_L = "Tragion/TragionCont"))  
  
#Autofit Width Table TNR  
flextable(TrTr\_Lrace\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Tragion Linear SumStats by Race/Ethnicity") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Tragion Linear SumStats by Race/Ethnicity

| **race\_eth** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| white | 1,240 | 127 | 173 | 145.50 | 145.0 | 7.57 | 0.22 | 134.0 | 140.0 | 145.0 | 151.00 | 158.00 | 20 |
| Black | 548 | 130 | 172 | 148.47 | 148.0 | 7.57 | 0.32 | 137.0 | 143.0 | 148.0 | 154.00 | 161.10 | 9 |
| LatinX | 100 | 132 | 171 | 146.90 | 147.0 | 7.54 | 0.75 | 135.0 | 141.5 | 147.0 | 151.00 | 158.10 | 1 |
| Asian | 91 | 129 | 167 | 147.56 | 148.0 | 8.67 | 0.91 | 135.4 | 140.0 | 148.0 | 153.00 | 161.00 | 2 |
| Other | 21 | 138 | 159 | 147.45 | 146.0 | 6.56 | 1.43 | 139.9 | 142.0 | 146.0 | 151.75 | 158.05 | 1 |
| AIAN | 8 | 132 | 157 | 141.86 | 140.0 | 7.99 | 2.82 | 133.8 | 138.5 | 140.0 | 143.50 | 154.00 | 1 |
| PTNS | 5 | 137 | 156 | 147.80 | 153.0 | 8.64 | 3.87 | 137.6 | 140.0 | 153.0 | 153.00 | 155.40 | 0 |
| NHOPI | 4 | 153 | 160 | 156.50 | 156.5 | 3.11 | 1.55 | 153.3 | 154.5 | 156.5 | 158.50 | 159.70 | 0 |

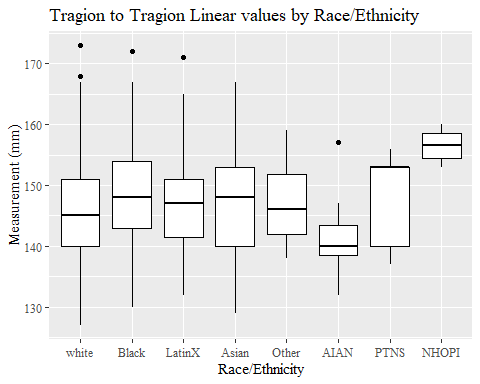
#%>% set\_header\_Labels(values = list(TrTr\_L = "Tragion/TragionCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(race\_eth= fct\_reorder(race\_eth, TrTr\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrTr\_L, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Tragion Linear values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 34 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=TrTr\_L, x=race\_eth))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Tragion Linear values by Race/Ethnicity",  
 y="Measurement (mm)",  
 x="Race/Ethnicity")

## Warning: Removed 34 rows containing non-finite values (stat\_boxplot).



#TrTr\_L gender sumstats  
TrTr\_Lgender\_sumstats <- headscan\_full %>%   
 group\_by(gender) %>%   
 summarise(n = n(),  
 min = min(TrTr\_L, na.rm = TRUE),  
 max = max(TrTr\_L, na.rm = TRUE),  
 mean = mean(TrTr\_L, na.rm = TRUE),  
 mdn = median(TrTr\_L, na.rm = TRUE),  
 sd = sd(TrTr\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrTr\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrTr\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrTr\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrTr\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrTr\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrTr\_L)))  
  
TrTr\_Lgender\_sumstats <- TrTr\_Lgender\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
#Size 12 Table TNR  
flextable(TrTr\_Lgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Tragion Linear SumStats by Gender")

**Table** : Tragion to Tragion Linear SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 127 | 165 | 141.88 | 142 | 5.82 | 0.18 | 133.0 | 138.00 | 142 | 145.00 | 152.0 | 23 |
| Male | 939 | 132 | 173 | 151.72 | 152 | 6.18 | 0.20 | 142.0 | 147.00 | 152 | 156.00 | 162.0 | 11 |
| Non-binary or Other | 5 | 139 | 149 | 143.20 | 141 | 4.15 | 1.85 | 139.4 | 141.00 | 141 | 146.00 | 148.4 | 0 |
| Prefer not to say | 1 | 153 | 153 | 153.00 | 153 |  |  | 153.0 | 153.00 | 153 | 153.00 | 153.0 | 0 |
|  | 8 | 138 | 150 | 144.50 | 144 | 3.82 | 1.35 | 139.4 | 142.75 | 144 | 147.25 | 149.3 | 0 |

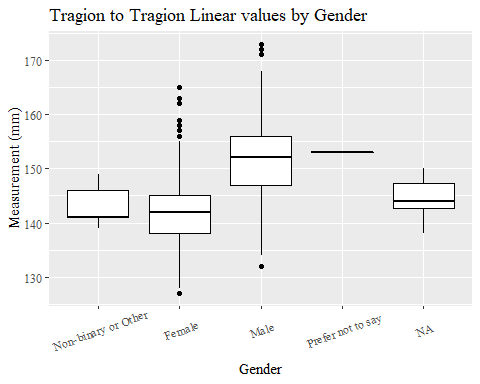
#%>% set\_header\_Labels(values = list(TrTr\_L = "Tragion/TragionCont"))  
  
#Autofit Width Table TNR  
flextable(TrTr\_Lgender\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Tragion Linear SumStats by Gender") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Tragion Linear SumStats by Gender

| **gender** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| Female | 1,064 | 127 | 165 | 141.88 | 142 | 5.82 | 0.18 | 133.0 | 138.00 | 142 | 145.00 | 152.0 | 23 |
| Male | 939 | 132 | 173 | 151.72 | 152 | 6.18 | 0.20 | 142.0 | 147.00 | 152 | 156.00 | 162.0 | 11 |
| Non-binary or Other | 5 | 139 | 149 | 143.20 | 141 | 4.15 | 1.85 | 139.4 | 141.00 | 141 | 146.00 | 148.4 | 0 |
| Prefer not to say | 1 | 153 | 153 | 153.00 | 153 |  |  | 153.0 | 153.00 | 153 | 153.00 | 153.0 | 0 |
|  | 8 | 138 | 150 | 144.50 | 144 | 3.82 | 1.35 | 139.4 | 142.75 | 144 | 147.25 | 149.3 | 0 |

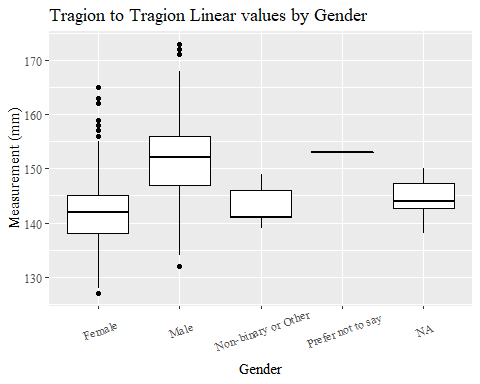
#%>% set\_header\_Labels(values = list(TrTr\_L = "Tragion/TragionCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(gender= fct\_reorder(gender, TrTr\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrTr\_L, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Tragion Linear values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 34 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=TrTr\_L, x=gender))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Tragion Linear values by Gender",  
 y="Measurement (mm)",  
 x="Gender")

## Warning: Removed 34 rows containing non-finite values (stat\_boxplot).



#TrTr\_L age group sumstats  
TrTr\_Lage\_sumstats <- headscan\_full %>%   
 group\_by(age\_group) %>%   
 summarise(n = n(),  
 min = min(TrTr\_L, na.rm = TRUE),  
 max = max(TrTr\_L, na.rm = TRUE),  
 mean = mean(TrTr\_L, na.rm = TRUE),  
 mdn = median(TrTr\_L, na.rm = TRUE),  
 sd = sd(TrTr\_L, na.rm = TRUE),  
 se = sd/sqrt(n),  
 percent5th = quantile(TrTr\_L, 0.05, na.rm=TRUE),  
 percent25th = quantile(TrTr\_L, 0.25, na.rm=TRUE),  
 percent50th = quantile(TrTr\_L, 0.50, na.rm=TRUE),  
 percent75th = quantile(TrTr\_L, 0.75, na.rm=TRUE),  
 percent95th = quantile(TrTr\_L, 0.95, na.rm=TRUE),  
 na = sum(is.na(TrTr\_L)))  
  
TrTr\_Lage\_sumstats <- TrTr\_Lage\_sumstats %>%   
 mutate(across(where(is.numeric), round, 2))  
  
  
#Size 12 Table TNR  
flextable(TrTr\_Lage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Tragion Linear SumStats by Age Group")

**Table** : Tragion to Tragion Linear SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 127 | 167 | 146.36 | 146 | 7.68 | 0.24 | 134.0 | 141 | 146 | 152 | 160.00 | 15 |
| 37-54 | 940 | 127 | 173 | 146.42 | 146 | 7.69 | 0.25 | 135.0 | 141 | 146 | 152 | 159.95 | 18 |
| 55-72 | 84 | 132 | 172 | 149.19 | 149 | 8.49 | 0.93 | 136.1 | 143 | 149 | 156 | 161.90 | 1 |
|  | 1 | 147 | 147 | 147.00 | 147 |  |  | 147.0 | 147 | 147 | 147 | 147.00 | 0 |

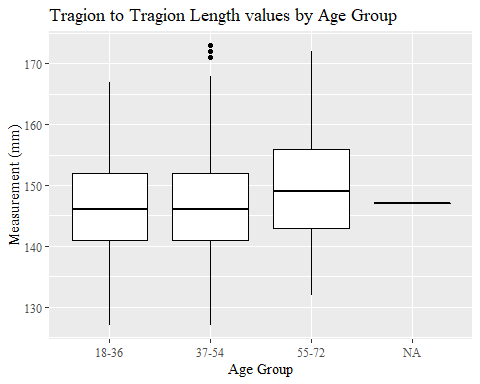
#%>% set\_header\_Labels(values = list(TrTr\_L = "Tragion/TragionCont"))  
  
#Autofit Width Table TNR  
flextable(TrTr\_Lage\_sumstats) %>%  
 my\_ft\_theme()%>%   
 bold(part = "header") %>%   
 set\_caption("Tragion to Tragion Linear SumStats by Age Group") %>%   
 fit\_to\_width(7.5)

**Table** : Tragion to Tragion Linear SumStats by Age Group

| **age\_group** | **n** | **min** | **max** | **mean** | **mdn** | **sd** | **se** | **percent5th** | **percent25th** | **percent50th** | **percent75th** | **percent95th** | **na** |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| 18-36 | 992 | 127 | 167 | 146.36 | 146 | 7.68 | 0.24 | 134.0 | 141 | 146 | 152 | 160.00 | 15 |
| 37-54 | 940 | 127 | 173 | 146.42 | 146 | 7.69 | 0.25 | 135.0 | 141 | 146 | 152 | 159.95 | 18 |
| 55-72 | 84 | 132 | 172 | 149.19 | 149 | 8.49 | 0.93 | 136.1 | 143 | 149 | 156 | 161.90 | 1 |
|  | 1 | 147 | 147 | 147.00 | 147 |  |  | 147.0 | 147 | 147 | 147 | 147.00 | 0 |

#%>% set\_header\_Labels(values = list(TrTr\_L = "Tragion/TragionCont"))  
  
#boxplot reorderd by median  
headscan\_full %>%   
 ungroup() %>%   
 mutate(age\_group= fct\_reorder(age\_group, TrTr\_L, FUN=median, na.rm=TRUE)) %>%   
 ggplot(aes(y=TrTr\_L, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Tragion Length values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 34 rows containing non-finite values (stat\_boxplot).



#boxplot not reordered by median  
ggplot(data=headscan\_full, aes(y=TrTr\_L, x=age\_group))+  
 geom\_boxplot(color= "black", fill = "white")+  
 theme(text=element\_text(family= "Times New Roman"))+  
 #theme(axis.text.x = element\_text(angle = 20, vjust=0.7))+  
 labs(title="Tragion to Tragion Length values by Age Group",  
 y="Measurement (mm)",  
 x="Age Group")

## Warning: Removed 34 rows containing non-finite values (stat\_boxplot).

