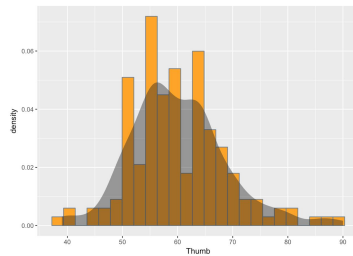


Basics <pre>print("hello") mynumber <- 5 # assigns value to object myvector <- c(1,2,3) # combines elements into vector myvector[1] # first element in vector Fingers\$Sex # variable in data frame</pre>	Tables <pre>tally(myvector) tally(~ Condition, data = MindsetMatters) tally(~ Thumb > 65, data = Fingers) tally(Thumb ~ Sex, data = Fingers, margins = TRUE, format = "proportion")</pre>	Fitting and Evaluating Models <pre>Empty.model <- lm(Thumb ~ NULL, data = Fingers) Sex.model <- lm(Thumb ~ Sex, data = Fingers) Height.fun<- makeFun(Height.model) predict(Empty.model) resid(Empty.model) anova(Empty.model) supernova(Sex.model)</pre>	Data <pre>str(MindsetMatters) head(MindsetMatters) tail(MindsetMatters) sort(myvector) arrange(Fingers, Thumb) select(Fingers, Sex, RaceEthnic, Thumb) # selects variables filter(Fingers, SLast != "NA") # selects cases head(select(Fingers, Thumb))</pre>
Operators <pre>sum(1,2,100) +, -, *, / >, <, >=, <=, ==, != Fingers\$RingLonger <- Fingers\$Ring > Fingers\$Index # results in TRUE or FALSE abs(Fingers\$Residual) Fingers\$Residual^2 sqrt(157)</pre>	Simple Statistics <pre>mean(Fingers\$Thumb) var(Fingers\$Thumb) sd(Fingers\$Thumb) favstats(~ Wt, data = MindsetMatters) cohensD(Thumb ~ Sex, data = Fingers) cor(Thumb ~ Height, data = Fingers)</pre> Probability Distributions <pre>xpnorm(65.1, Thumb.stats\$mean, Thumb.stats\$sd) zscore(Fingers\$Thumb) qt(.975, df = 999) # returns t at this probability qf(.95, df1 = 1, df2 = 100) # returns F at this probability confint(Empty.model) # CI using t distribution</pre>	Simulation & Resampling <pre>sample(Fingers\$Thumb, 10) # sample without replacement resample(Fingers\$Thumb, 157) # sample with replacement do(3) * resample(Fingers\$Thumb,10) shuffle(Servers\$RandomGroups1) # mixes up values in a variable simThumb <- rnorm(10000, Thumb.stats\$mean, Thumb.stats\$sd) # simulates sampling from a normal distribution simPop <- data.frame(simThumb) simSDoM <- do(10000) * mean(rnorm(157, Thumb.stats\$mean, Thumb.stats\$sd)) # simulates sampling distribution of means bootSDoM <- do(10000) * mean(resample(Fingers\$Thumb,157)) # bootstraps sampling distribution of means</pre>	<pre>as.factor(Fingers\$Sex) as.numeric(Fingers\$Interest) factor(Fingers\$Sex, levels = c(1,2), labels = c("female", "male")) recode(Fingers\$Job, "0" = 0, "1" = 50, "2" = 100) ntile(Fingers\$Height, 2) # creates equal sized groups aggregate(Happiness ~ Region, data = HappyPlanetIndex, FUN = mean)</pre>

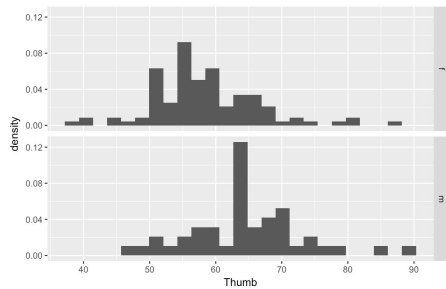
Plots

```
gf_histogram(~ Thumb, data = Fingers, color =
"red", fill = "gray", bins = 10, binwidth = 4)
```

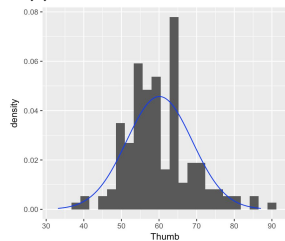
```
%>% gf_labs(title= "Student Thumb Lengths",
x= "Thumb Length (mm)")
# changes labels
%>% gf_density()
# adds density curve to a histogram
```



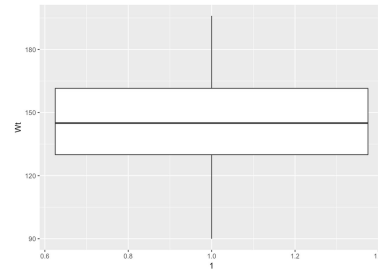
```
gf_histogram(..density..~ Thumb, data =
Fingers)
%>% gf_facet_grid(Sex ~ .)
```



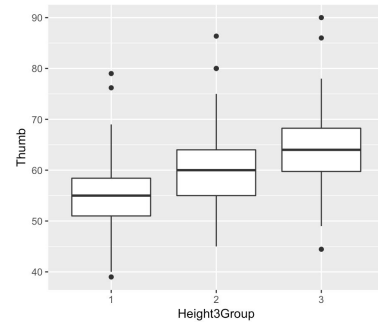
```
%>% gf_dist("norm", color = "blue", params =
list(Thumb.stats$mean,
Thumb.stats$sd))
```



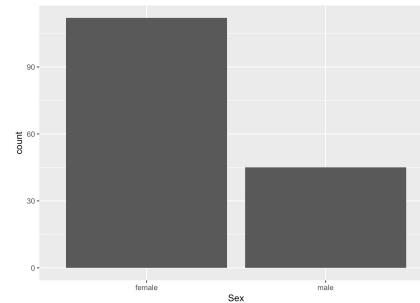
```
gf_boxplot(Wt ~ 1, data = MindsetMatters)
```



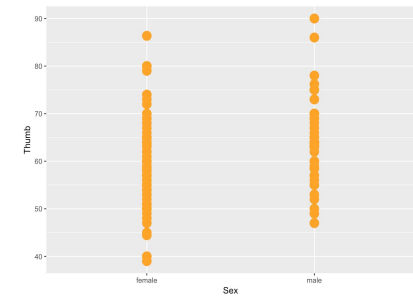
```
gf_boxplot(Thumb ~ Height3Group, data =
Fingers)
```



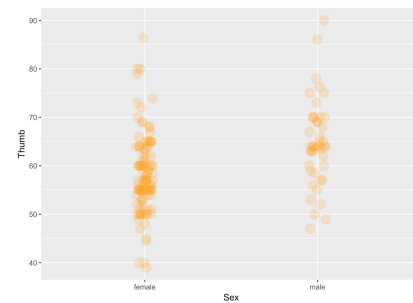
```
gf_bar(~ Sex, data = Fingers)
# creates bar graph
```



```
gf_point(Thumb ~ Sex, data = Fingers, color =
"orange", size = 5)
```



```
gf_jitter(Thumb ~ Sex, data = Fingers, color =
"orange", size = 5, alpha = .5)
```



```
gf_point(Thumb ~ Height, data = Fingers, size
= 4) %>%
gf_lm(color = "orange")
# adds a regression line
```

