Share A Graph

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Consumer Goods Data Import

```
Consumer Goods Dataset
```

```
ConsumerGood <- read_csv("~/CSVs/ConsumerGood.csv")</pre>
## Rows: 108 Columns: 4
## -- Column specification -------
## Delimiter: ","
## dbl (4): rownames, distribution, share, price
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.
head(ConsumerGood)
## # A tibble: 6 x 4
##
   rownames distribution share price
##
      <dbl>
                  <dbl> <dbl> <dbl>
                   0.905 2.74 106.
## 1
        1
## 2
         2
                         3.01 106.
                   0.9
                   0.988 2.20 107.
         3
## 3
                        2.67 106.
## 4
          4
                   0.96
## 5
         5
                   0.954 2.87 106.
## 6
          6
                   0.988 2.77 106.
```

Share and distribution of consumer goods

```
# Modify Dataset to include Ypred
ConsumerGood = ConsumerGood %>% mutate(Ypred=-0.8896+3.547*distribution)

# Linear Model
dist_share_model <- lm(share ~ distribution, data = ConsumerGood)
summary(dist_share_model)

##
## Call:
## Im(formula = share ~ distribution, data = ConsumerGood)
##
## Residuals:
## Min    1Q Median    3Q Max
## -1.03024 -0.35559 -0.04637    0.30933    1.69116
##</pre>
```

```
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
##
                -0.8896
                           0.1718 -5.179 1.07e-06 ***
## (Intercept)
## distribution
                 3.5470
                            0.2250 15.764 < 2e-16 ***
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.5724 on 106 degrees of freedom
## Multiple R-squared: 0.701, Adjusted R-squared: 0.6982
## F-statistic: 248.5 on 1 and 106 DF, p-value: < 2.2e-16
# scatter plot
gf_point(
 share ~ distribution,
 data = ConsumerGood,
 # sets color based on residual
 color = ifelse(residuals(dist_share_model) == 0, "green",
  # if point not on line set as red or blue
  ifelse(residuals(dist_share_model) > 0, "red", "blue")
 )) %>%
  # plots line of best fit
 gf_lm(color="gray50",alpha=0.5) %>%
  # plots dotted residual lines from point to linear model
 gf_linerange(Ypred+share~distribution, linetype="dotted", color="gray30")
```

