Iris Demo

```
library(tidyverse)

df <- iris %>%
    filter(Species == params$species)

Report for the species: versicolor

df %>%
    filter(Species == params$species) %>%
    ggplot(aes(Sepal.Width))+
    geom_histogram() +
    ggtitle(str_c("Distribution of Sepal.Length for ", params$species))

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

Distribution of Sepal.Length for versicolor

