

# Iris Demo

```
library(tidyverse)

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

Report for the species: setosa

```
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 setosa

