TP bourdon
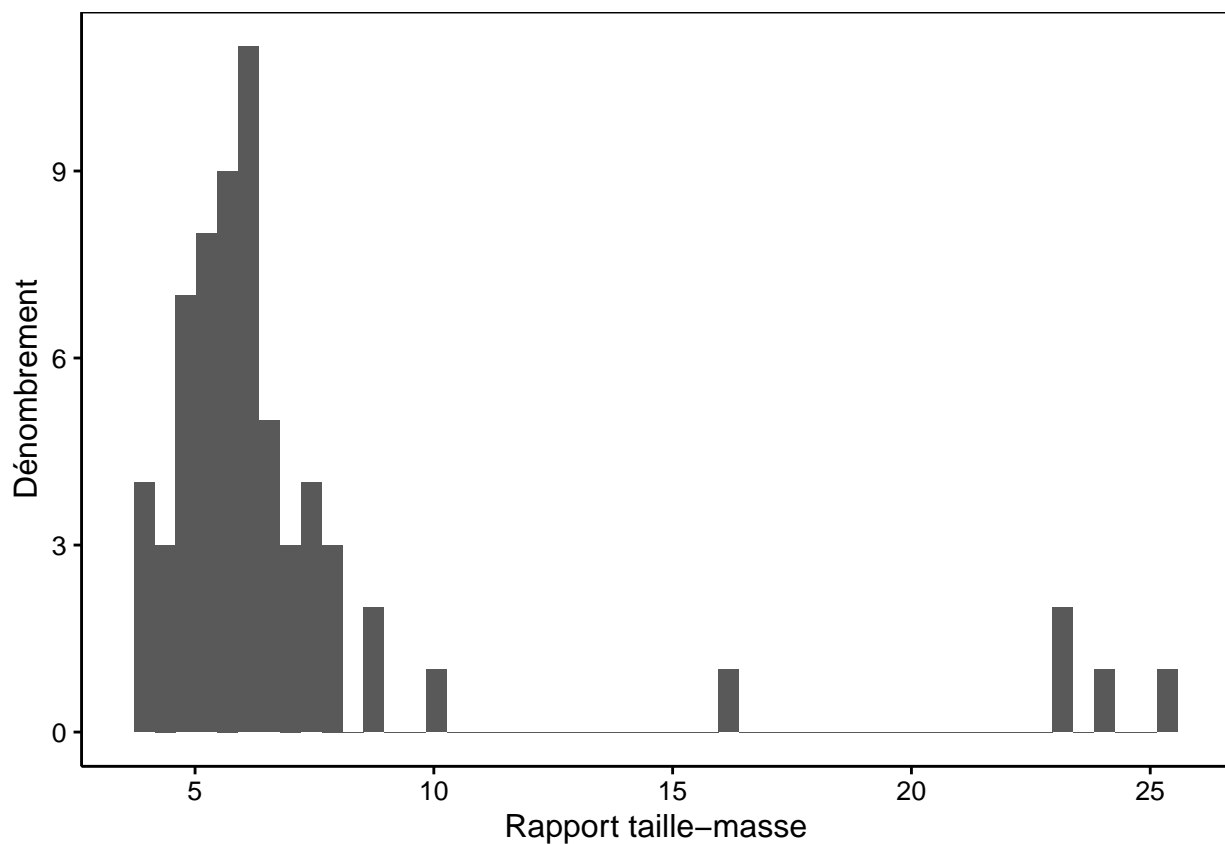
```
colonie <- read("bourdonR.xlsx")

## New names:
## * `` -> ...1
colonie %>.%
  filter(., Type == "larve") %>.%
  select(., -ITD) %>.%
  mutate(., t_m = taille/masse) -> colonie_larve

chart(data = colonie_larve, ~t_m)+
  geom_histogram(bins = 50) +
  ylab("Dénombrement") +
  xlab("Rapport taille-masse")

## Warning: Removed 6 rows containing non-finite values (stat_bin).
```



```
colonie %>.%
  filter(., Type == "ouvrière") %>.%
  select(., -taille) %>.%
```

```
  mutate(., t_m = ITD/masse) -> colonie_ouvrière
```

```
chart(data = colonie_ouvrière, ~t_m)+
  geom_histogram(bins = 50) +
  ylab("Dénombrement") +
  xlab("Rapport taille-masse")
```