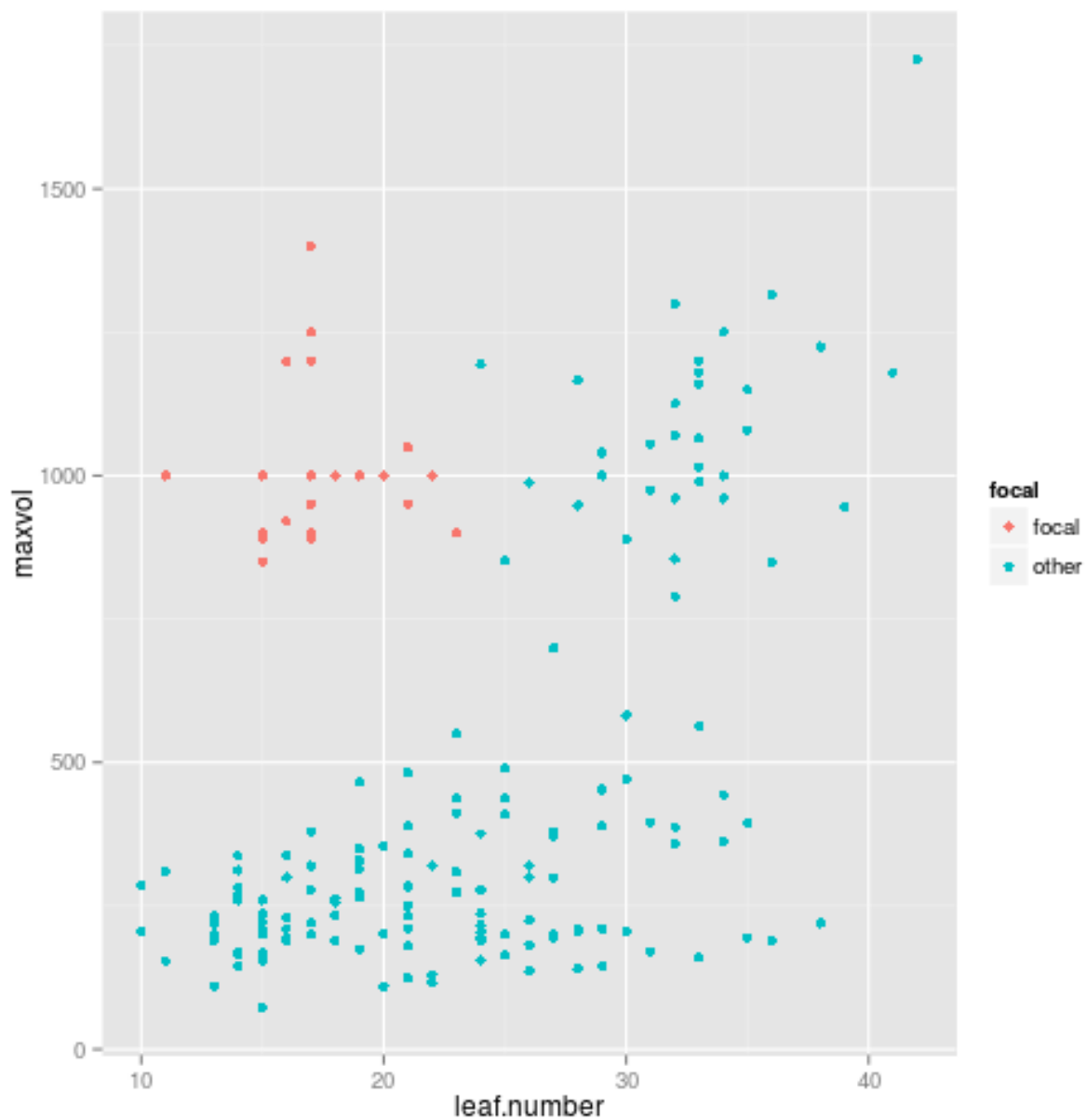


Data from macae

Here is a simple plot:

```
physical %>%  
  mutate(focal = ifelse(site == dataset, "focal", "other")) %>%  
  ggplot(aes(x = leaf.number, y = maxvol, colour = focal)) +  
  geom_point()
```

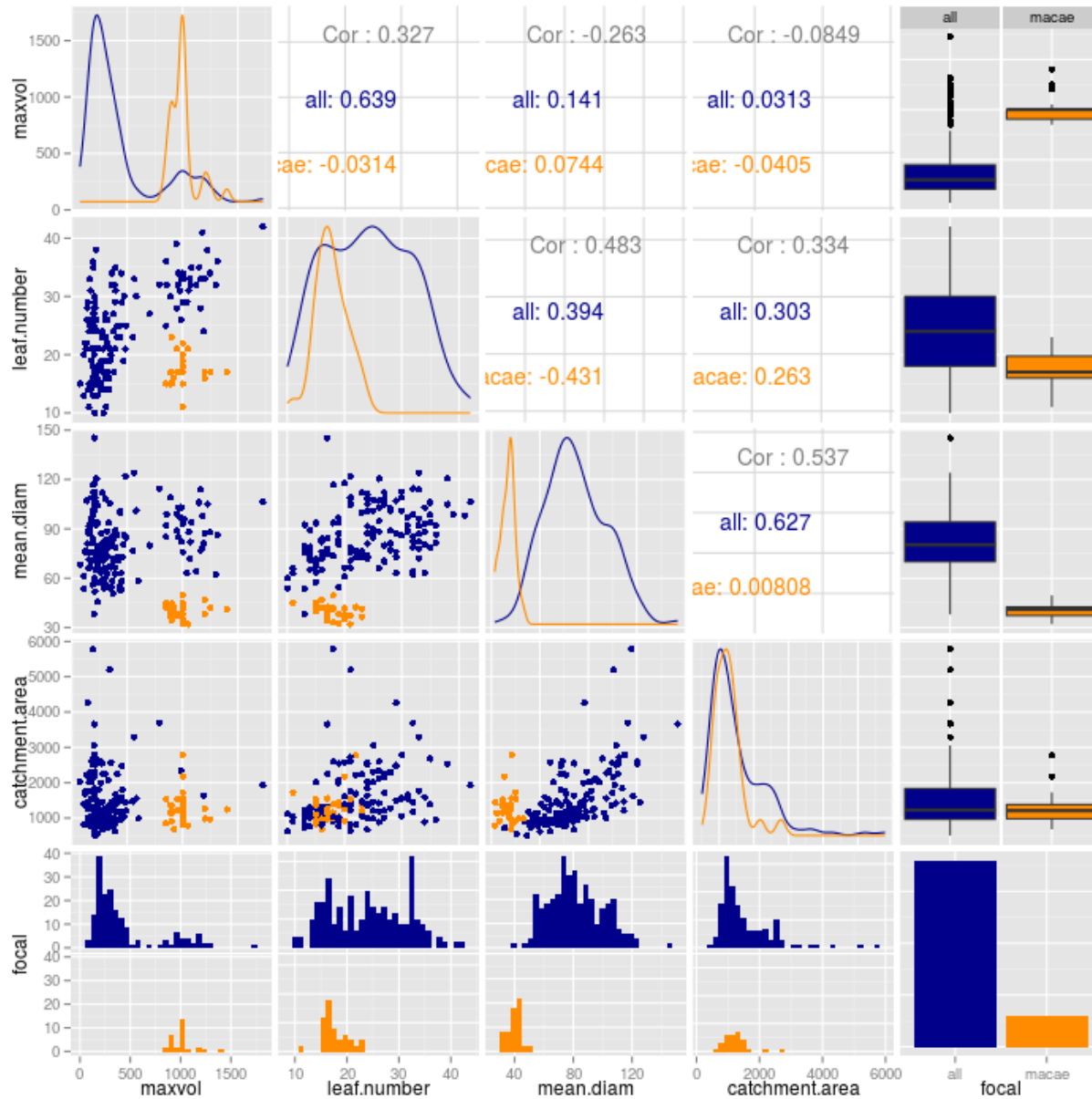
Warning: Removed 30 rows containing missing values (geom_point).



Size variables

```
sizepairs <- physical %>%
  mutate(focal = ifelse(site == dataset, dataset, "all")) %>%
  select(maxvol:catchment.area, focal) %>%
  ggpairs(colour = "focal")

add_theme_to_ggpairs(sizepairs)
```



Chemical variables – initial

```
sizepairs <- physical %>%
  mutate(focal = ifelse(site == dataset, dataset, "all")) %>%
  select(turbidity.initial:chlorophyll.initial, focal) %>%
  ggpairs(colour = "focal")

add_theme_to_ggpairs(sizepairs)
```

