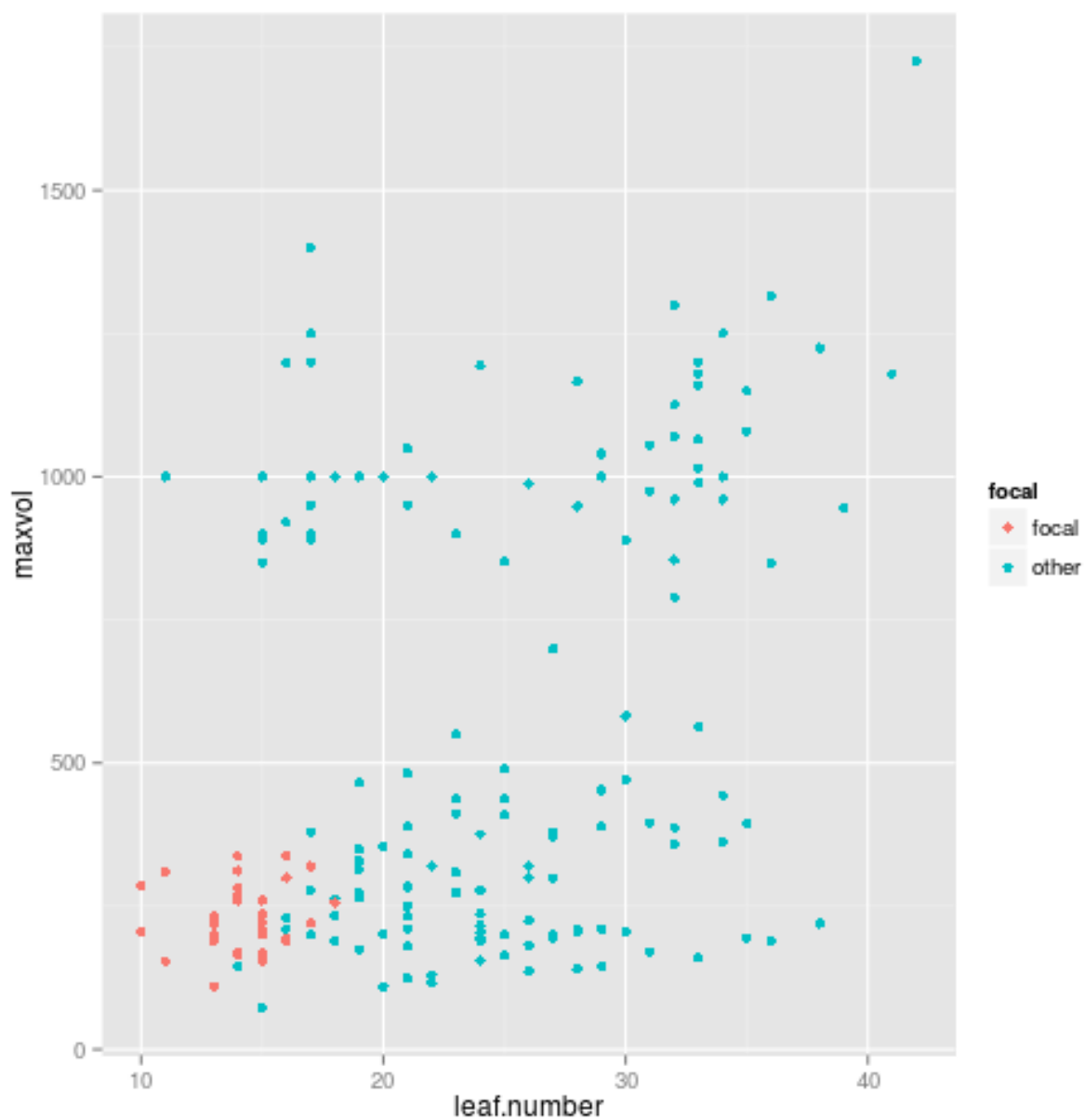


## Data from frenchguiana

Here is a simple plot:

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

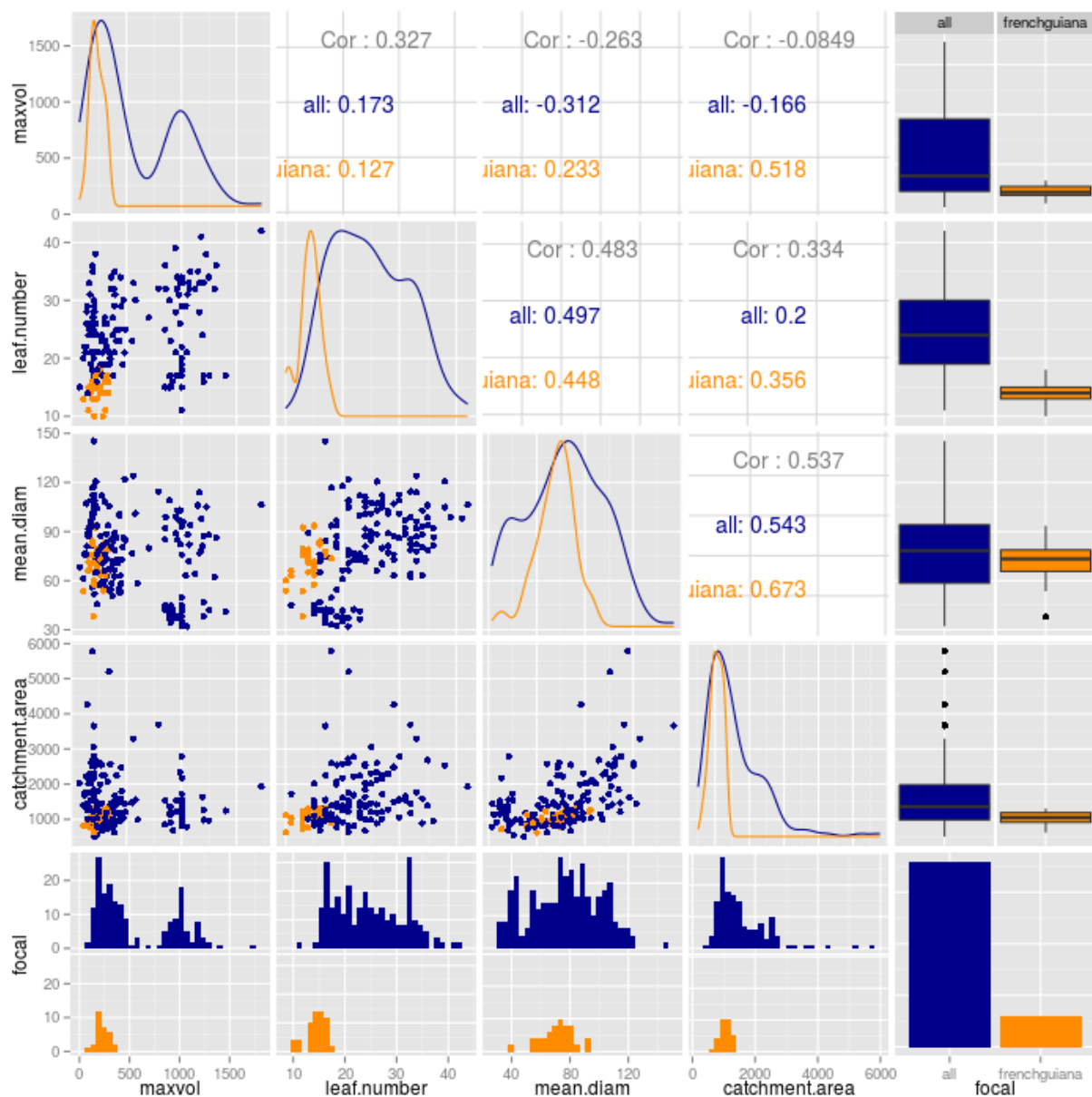
## Warning: Removed 30 rows containing missing values (geom\_point).



## Size variables

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

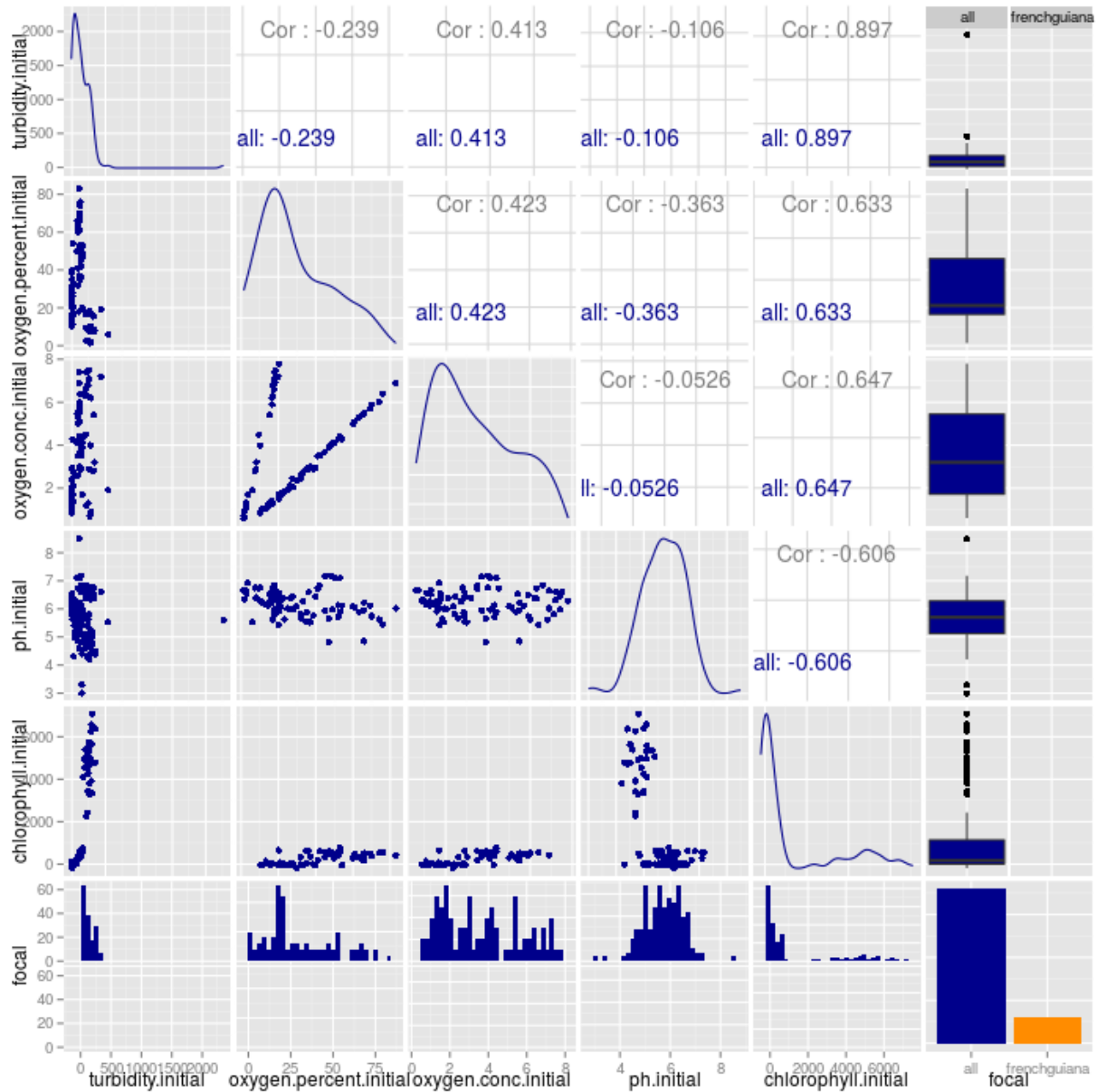
add_theme_to_ggpairs(sizepairs)
```



## 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)
```



```

set.seed(4812)
physical %>%
  filter(site == dataset) %>%
  select(site_brom.id, turbidity.initial:chlorophyll.initial) %>%
  sample_n(3) %>%
  mutate(`*CONFIRMED*` = "    ") %>%
  as.data.frame %>%
  kable

```

site_brom.id	turbidity.initial	oxygen.percent.initial	oxygen.conc.initial	ph.initial	chlorophyll.initial	<i>CONFIRMED</i>
frenchguiana_35	NA	NA	NA	NA	NA	
frenchguiana_33	NA	NA	NA	NA	NA	
frenchguiana_79	NA	NA	NA	NA	NA	

**Chemical variables to check**