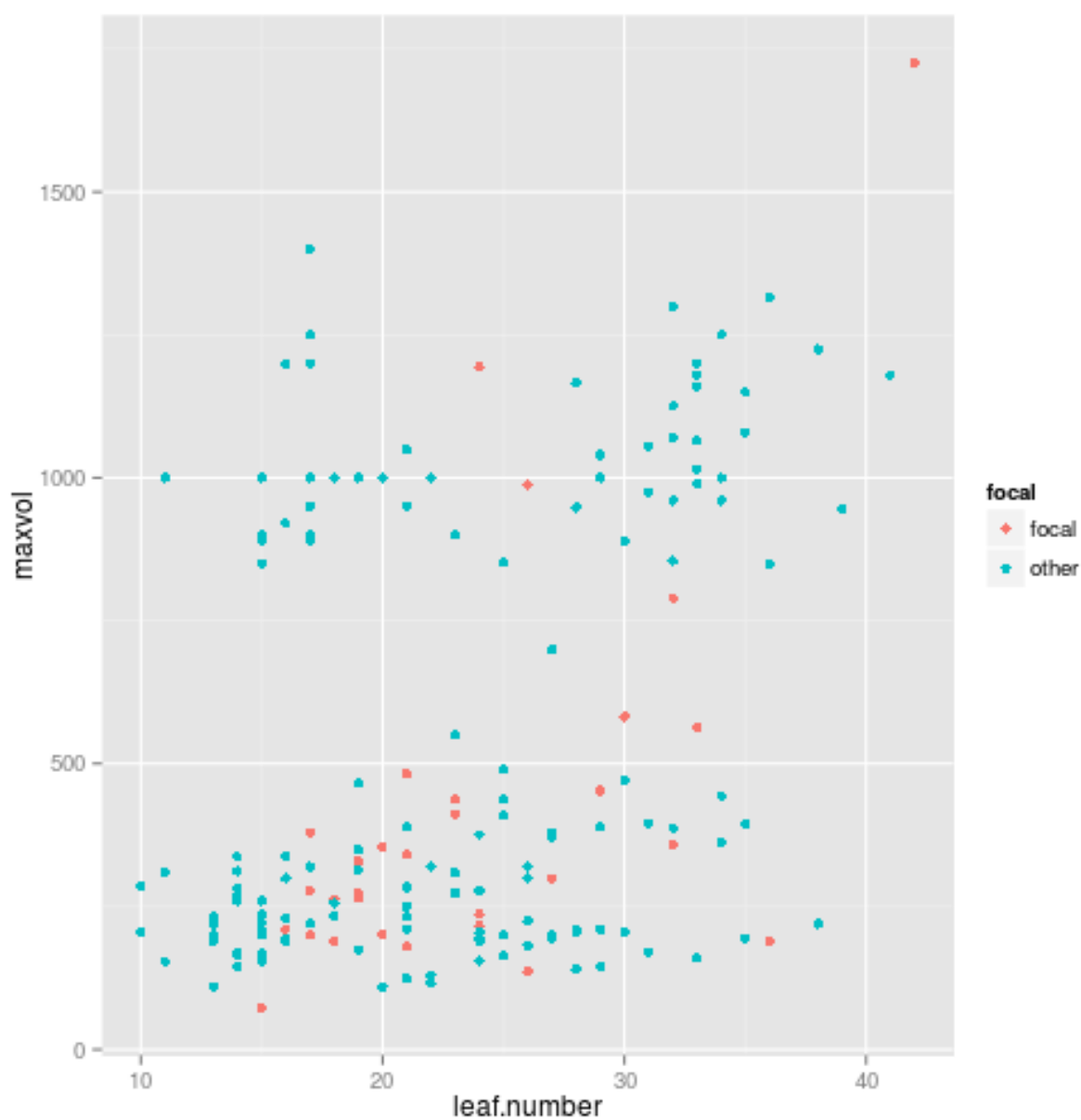


Data from colombia

Here is a simple plot:

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

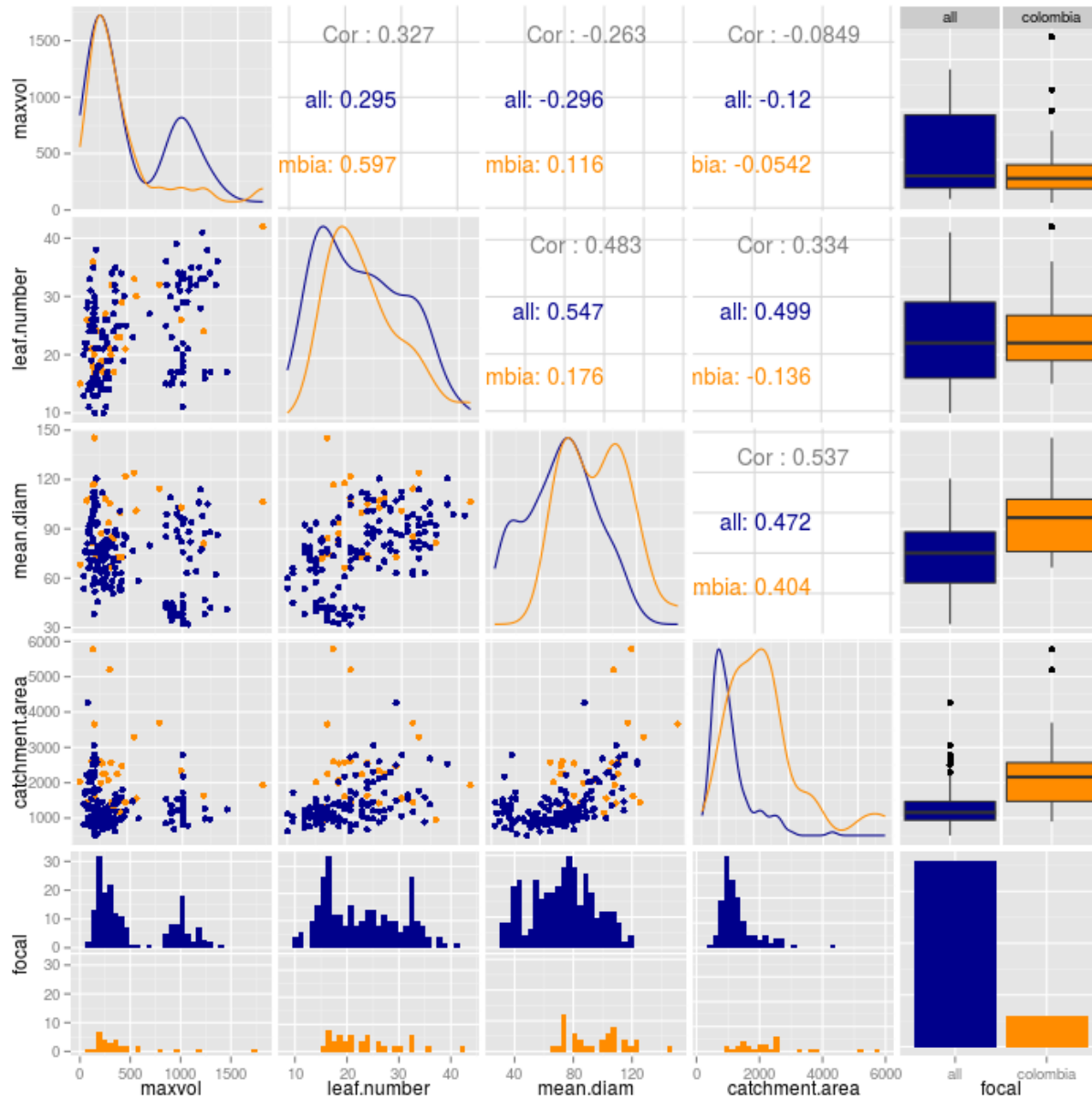
Warning: Removed 30 rows containing missing values (geom_point).



Size variables

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

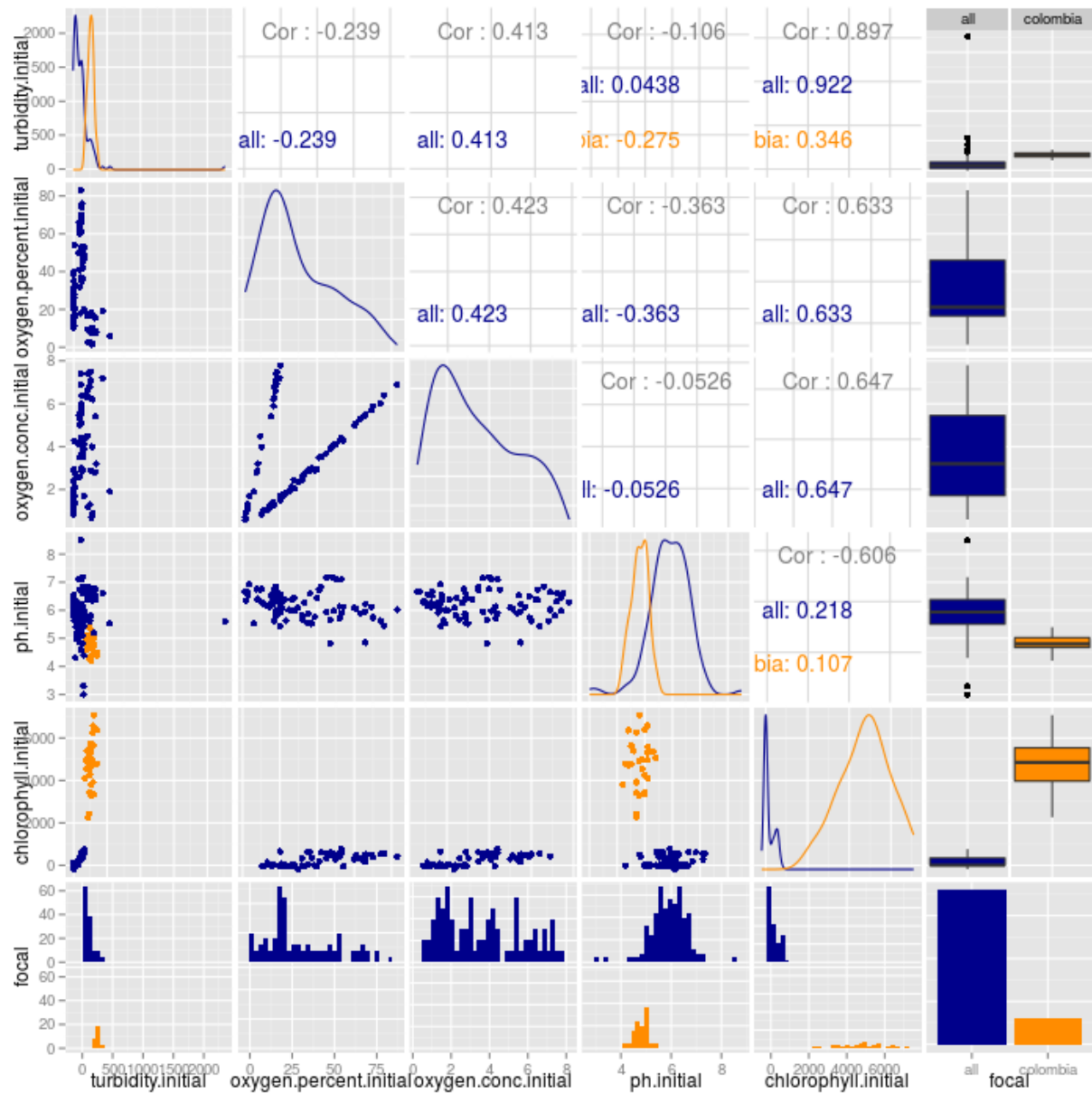
add_theme_to_ggpairs(sizepairs)
```



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>
colombia_11	302.7	NA	NA	4.8	7085	
colombia_9	298.5	NA	NA	5.0	6408	
colombia_10	262.2	NA	NA	4.2	3811	

Chemical variables to check