

Analysis of Cancer Cell Line Variants

Christiaan Klijn

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Test the plotting of a legend

```
plot(0, 0, xlim = c(0, 1000), ylim = c(0, 100), type = "n",  
     axes = F, xlab = "", ylab = "")  
rect(1, 20, 1000, 30, border = "black")  
labels <- c("Testing 111111111", "Testing 22222222", "Testing 3333333")  
legend("bottom", legend = labels, fill = rainbow(n = 3),  
       ncol = 3)
```

