test\_word

descr(iris) %>% print() %>% knit\_print()

## Warning: The `validate` argument of `as\_tibble()` is deprecated as of tibble 2.0.0.  
## Please use the `.name\_repair` argument instead.

| Variables | Total | p |
| --- | --- | --- |
|  | (N=150) |  |
| **Sepal.Length** |  |  |
| N | 150 | <0.001tt1 |
| mean | 5.8 |  |
| sd | 0.83 |  |
| median | 5.8 |  |
| Q1 - Q3 | 5.1 -- 6.4 |  |
| min - max | 4.3 -- 7.9 |  |
| **Sepal.Width** |  |  |
| N | 150 | <0.001tt1 |
| mean | 3.1 |  |
| sd | 0.44 |  |
| median | 3 |  |
| Q1 - Q3 | 2.8 -- 3.3 |  |
| min - max | 2 -- 4.4 |  |
| **Petal.Length** |  |  |
| N | 150 | <0.001tt1 |
| mean | 3.8 |  |
| sd | 1.8 |  |
| median | 4.3 |  |
| Q1 - Q3 | 1.6 -- 5.1 |  |
| min - max | 1 -- 6.9 |  |
| **Petal.Width** |  |  |
| N | 150 | <0.001tt1 |
| mean | 1.2 |  |
| sd | 0.76 |  |
| median | 1.3 |  |
| Q1 - Q3 | 0.3 -- 1.8 |  |
| min - max | 0.1 -- 2.5 |  |
| **Species** |  |  |
| setosa | 50 (33%) | >0.999chi1 |
| versicolor | 50 (33%) |  |
| virginica | 50 (33%) |  |
| tt1Students one-sample t-test | | |
| chi1Chi-squared goodness-of-fit test | | |

descr(  
 iris,  
 "Species",  
 group\_labels = list(setosa = "My custom group label"),  
 var\_options = list(Sepal.Length = list(label = "My custom variable label"))  
) %>% print() %>% knit\_print()

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## Warning: The `value` argument of ``names<-`()` must be a character vector as of  
## tibble 3.0.0.

| Variables | My custom group label | versicolor | virginica | Total | p |
| --- | --- | --- | --- | --- | --- |
|  | (N=50) | (N=50) | (N=50) | (N=150) |  |
| My custom variable label |  |  |  |  |  |
| N | 150 | 150 | 150 | 150 | <0.001F |
| mean | 5.8 | 5.8 | 5.8 | 5.8 |  |
| sd | 0.83 | 0.83 | 0.83 | 0.83 |  |
| median | 5.8 | 5.8 | 5.8 | 5.8 |  |
| Q1 - Q3 | 5.1 -- 6.4 | 5.1 -- 6.4 | 5.1 -- 6.4 | 5.1 -- 6.4 |  |
| min - max | 4.3 -- 7.9 | 4.3 -- 7.9 | 4.3 -- 7.9 | 4.3 -- 7.9 |  |
| **Sepal.Width** |  |  |  |  |  |
| N | 150 | 150 | 150 | 150 | <0.001F |
| mean | 3.1 | 3.1 | 3.1 | 3.1 |  |
| sd | 0.44 | 0.44 | 0.44 | 0.44 |  |
| median | 3 | 3 | 3 | 3 |  |
| Q1 - Q3 | 2.8 -- 3.3 | 2.8 -- 3.3 | 2.8 -- 3.3 | 2.8 -- 3.3 |  |
| min - max | 2 -- 4.4 | 2 -- 4.4 | 2 -- 4.4 | 2 -- 4.4 |  |
| **Petal.Length** |  |  |  |  |  |
| N | 150 | 150 | 150 | 150 | <0.001F |
| mean | 3.8 | 3.8 | 3.8 | 3.8 |  |
| sd | 1.8 | 1.8 | 1.8 | 1.8 |  |
| median | 4.3 | 4.3 | 4.3 | 4.3 |  |
| Q1 - Q3 | 1.6 -- 5.1 | 1.6 -- 5.1 | 1.6 -- 5.1 | 1.6 -- 5.1 |  |
| min - max | 1 -- 6.9 | 1 -- 6.9 | 1 -- 6.9 | 1 -- 6.9 |  |
| **Petal.Width** |  |  |  |  |  |
| N | 150 | 150 | 150 | 150 | <0.001F |
| mean | 1.2 | 1.2 | 1.2 | 1.2 |  |
| sd | 0.76 | 0.76 | 0.76 | 0.76 |  |
| median | 1.3 | 1.3 | 1.3 | 1.3 |  |
| Q1 - Q3 | 0.3 -- 1.8 | 0.3 -- 1.8 | 0.3 -- 1.8 | 0.3 -- 1.8 |  |
| min - max | 0.1 -- 2.5 | 0.1 -- 2.5 | 0.1 -- 2.5 | 0.1 -- 2.5 |  |
| FF-test (ANOVA) | | | | | |

descr(iris) %>% print(silent=T) %>% .$ft %>% knit\_print()

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| Variables | Total | p |
| --- | --- | --- |
|  | (N=150) |  |
| **Sepal.Length** |  |  |
| N | 150 | <0.001tt1 |
| mean | 5.8 |  |
| sd | 0.83 |  |
| median | 5.8 |  |
| Q1 - Q3 | 5.1 -- 6.4 |  |
| min - max | 4.3 -- 7.9 |  |
| **Sepal.Width** |  |  |
| N | 150 | <0.001tt1 |
| mean | 3.1 |  |
| sd | 0.44 |  |
| median | 3 |  |
| Q1 - Q3 | 2.8 -- 3.3 |  |
| min - max | 2 -- 4.4 |  |
| **Petal.Length** |  |  |
| N | 150 | <0.001tt1 |
| mean | 3.8 |  |
| sd | 1.8 |  |
| median | 4.3 |  |
| Q1 - Q3 | 1.6 -- 5.1 |  |
| min - max | 1 -- 6.9 |  |
| **Petal.Width** |  |  |
| N | 150 | <0.001tt1 |
| mean | 1.2 |  |
| sd | 0.76 |  |
| median | 1.3 |  |
| Q1 - Q3 | 0.3 -- 1.8 |  |
| min - max | 0.1 -- 2.5 |  |
| **Species** |  |  |
| setosa | 50 (33%) | >0.999chi1 |
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| virginica | 50 (33%) |  |
| tt1Students one-sample t-test | | |
| chi1Chi-squared goodness-of-fit test | | |