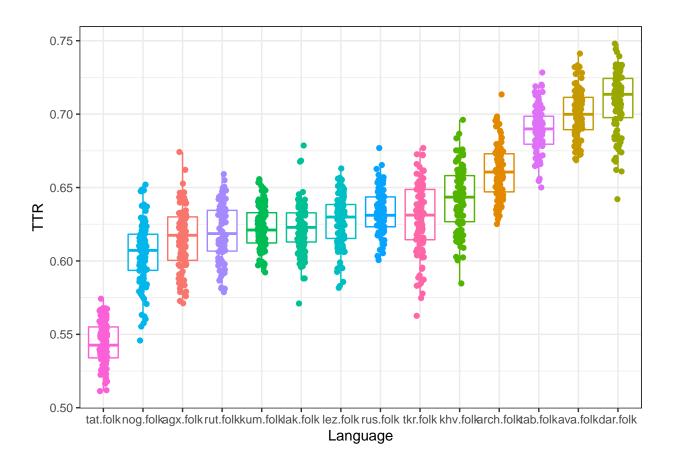
TTR across languages of Daghestan

100 TTRs per language

```
df1 <- read.csv("TTR_100_datapoints_per_language.txt",</pre>
                header = TRUE, sep = ",")
df1_long <- gather(df1)</pre>
head(df1_long)
##
          key
                  value
## 1 agx.folk 0.5849057
## 2 agx.folk 0.6420846
## 3 agx.folk 0.6250000
## 4 agx.folk 0.6620553
## 5 agx.folk 0.6105675
## 6 agx.folk 0.5928144
df1_long %>%
  ggplot(aes(reorder(key, value, FUN = median), value, color = key))+
  geom_boxplot(outlier.shape = NA)+
  geom_jitter(width = 0.1)+
  labs(x = "Language",
      y = "TTR")+
  theme_bw()+
  theme(legend.position = "none")
```



Pairwise t-test

Holm-Bonferroni method

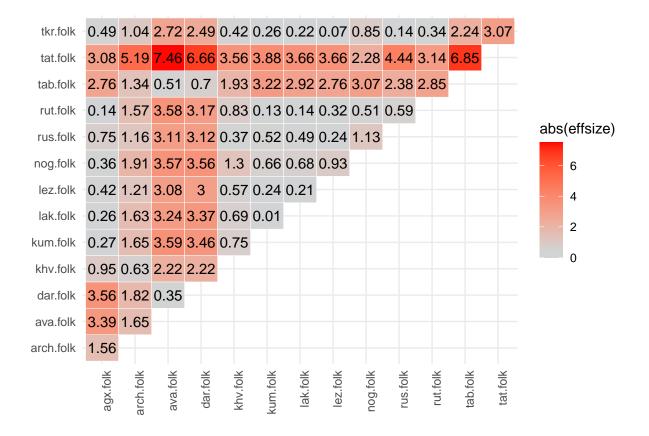
```
##
##
   Pairwise comparisons using paired t tests
##
## data: df1_long$value and df1_long$key
##
##
            agx.folk arch.folk ava.folk dar.folk khv.folk kum.folk lak.folk
## arch.folk < 2e-16 -
## ava.folk < 2e-16 < 2e-16
## dar.folk < 2e-16 < 2e-16
                               0.0115
## khv.folk 6.2e-14 2.0e-07
                               < 2e-16 < 2e-16
## kum.folk 0.0929
                     < 2e-16
                               < 2e-16
                                       < 2e-16 8.1e-10
## lak.folk 0.1310
                     < 2e-16
                               < 2e-16
                                       < 2e-16 1.6e-08
                                                        1.0000
                                                                  0.2973
## lez.folk 0.0011
                     < 2e-16
                               < 2e-16
                                        < 2e-16
                                                3.8e-06
                                                         0.1695
## nog.folk 0.0097
                     < 2e-16
                               < 2e-16
                                       < 2e-16 < 2e-16
                                                        5.9e-08
                                                                  2.9e-08
                                       < 2e-16 0.0060
## rus.folk 1.1e-09 < 2e-16
                               < 2e-16
                                                         2.2e-05
                                                                  8.0e-05
## rut.folk 1.0000
                     < 2e-16
                               < 2e-16 < 2e-16 1.6e-11 1.0000
                                                                  1.0000
```

```
## tab.folk < 2e-16 < 2e-16 4.1e-05 8.2e-09 < 2e-16 < 2e-16 < 2e-16
## tat.folk < 2e-16 
## tkr.folk 9.0e-05 7.1e-16 < 2e-16 < 2e-16 0.0013 0.1310
           lez.folk nog.folk rus.folk rut.folk tab.folk tat.folk
## arch.folk -
## ava.folk -
## dar.folk -
## khv.folk -
## kum.folk -
## lak.folk -
## lez.folk -
## nog.folk 1.6e-13 -
## rus.folk 0.1695
                   < 2e-16 -
## rut.folk 0.0283
                    3.9e-05 1.6e-06 -
## tab.folk < 2e-16 < 2e-16 < 2e-16 < 2e-16 -
## tat.folk < 2e-16 < 2e-16 < 2e-16 < 2e-16 < 2e-16 -
## tkr.folk 1.0000 6.7e-12 1.0000 0.0157 < 2e-16 < 2e-16
##
## P value adjustment method: holm
```

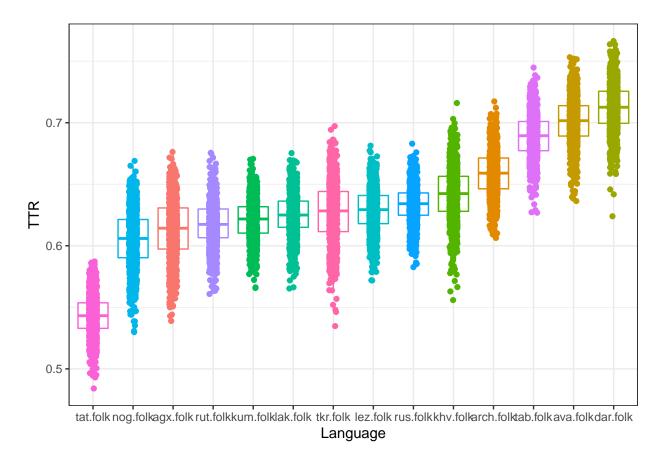
Effect sizes

```
library("rstatix")
##
## Attaching package: 'rstatix'
## The following object is masked from 'package:stats':
##
##
      filter
effect.sizes <- cohens_d(df1_long, value ~ key, paired = T)
print(effect.sizes)
## # A tibble: 91 x 7
           group1 group2
                             effsize
                                      n1
                                            n2 magnitude
     .у.
## * <chr> <chr>
                   <chr>
                             <dbl> <int> <int> <ord>
## 1 value agx.folk arch.folk -1.56 100 100 large
## 2 value agx.folk ava.folk -3.39
                                      100
                                           100 large
## 3 value agx.folk dar.folk -3.56
                                     100
                                          100 large
## 4 value agx.folk khv.folk -0.946 100
                                          100 large
## 5 value agx.folk kum.folk -0.275
                                     100
                                          100 small
## 6 value agx.folk lak.folk -0.257
                                      100
                                          100 small
                                     100
                                          100 small
## 7 value agx.folk lez.folk -0.421
## 8 value agx.folk nog.folk
                            0.356 100 100 small
## 9 value agx.folk rus.folk -0.747
                                      100 100 moderate
## 10 value agx.folk rut.folk
                                           100 negligible
                             -0.136 100
## # ... with 81 more rows
```

Effect size heatmap



1000 TTRs per language



Pairwise t-test

Holm-Bonferroni method

##

```
## Pairwise comparisons using paired t tests
##
## data: df2_long$value and df2_long$key
##
                                agx.folk arch.folk ava.folk dar.folk khv.folk kum.folk lak.folk
## arch.folk < 2e-16 -
## ava.folk < 2e-16 < 2e-16 -
## dar.folk < 2e-16 < 2e-16 < 2e-16 -
## khv.folk < 2e-16 < 2e-16 < 2e-16 < 2e-16 -
## kum.folk 6.2e-15 < 2e-16 < 2e-16 < 2e-16 < 2e-16 -
## lak.folk < 2e-16 < 2e-16 < 2e-16 < 2e-16 < 2e-16 5.0e-07
## lez.folk < 2e-16 < 3.3e-07 
## nog.folk 6.1e-15 < 2e-16 < 2e-16 < 2e-16 < 2e-16 < 2e-16 < 2e-16
## rus.folk < 2e-16 < 2e-16 < 2e-16 < 2e-16 < 2e-16 < 2e-16 < 2e-16
## rut.folk 3.3e-05 < 2e-16 < 2e-16 < 2e-16 < 2e-16 4.9e-05 < 2e-16
## tab.folk < 2e-16 < 2e-16 < 2e-16 < 2e-16 < 2e-16 < 2e-16
## tat.folk < 2e-16 
## tkr.folk < 2e-16 < 2e-16 < 2e-16 < 2e-16 < 2e-16 1.0e-11 0.0067
                               lez.folk nog.folk rus.folk rut.folk tab.folk tat.folk
## arch.folk -
## ava.folk -
## dar.folk -
## khv.folk -
## kum.folk -
## lak.folk -
## lez.folk -
## nog.folk < 2e-16 -
## rus.folk 3.6e-08 < 2e-16 -
## rut.folk < 2e-16 < 2e-16 < 2e-16 -
## tab.folk < 2e-16 < 2e-16 < 2e-16 < 2e-16 -
## tat.folk < 2e-16 < 2e-16 < 2e-16 < 2e-16 < 2e-16 < 2e-16 -
## tkr.folk 0.1477 < 2e-16 2.1e-09 < 2e-16 < 2e-16 < 2e-16
## P value adjustment method: holm
```

Effect sizes

```
effect.sizes <- cohens_d(df2_long, value ~ key, paired = T)
head(effect.sizes)</pre>
```

```
## # A tibble: 6 x 7
    .у.
          group1 group2
                            effsize
                                      n1
                                            n2 magnitude
    <chr> <chr>
                             <dbl> <int> <int> <ord>
                  <chr>
## 1 value agx.folk arch.folk -1.51
                                     1000 1000 large
## 2 value agx.folk ava.folk -2.93
                                     1000 1000 large
                                     1000 1000 large
## 3 value agx.folk dar.folk -3.21
## 4 value agx.folk khv.folk -0.878 1000 1000 large
## 5 value agx.folk kum.folk -0.260
                                     1000
                                          1000 small
## 6 value agx.folk lak.folk
                            -0.405 1000 1000 small
```

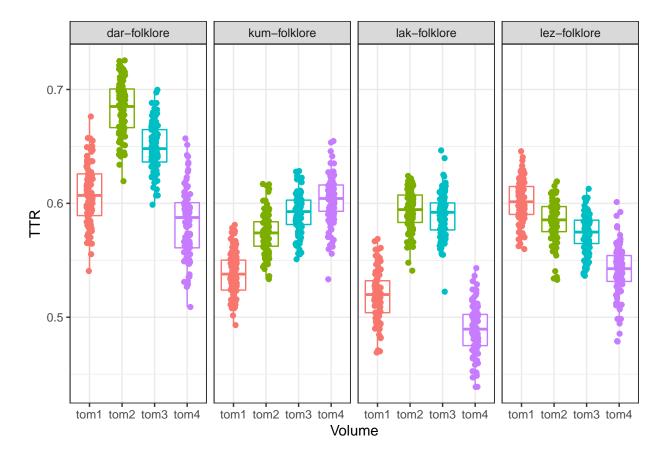
Effect size heatmap



Volume comparison

```
## 3 tom1 0.5345850 kum-folklore
## 4 tom1 0.5811209 kum-folklore
## 5 tom1 0.5374016 kum-folklore
## 6 tom1 0.5351137 kum-folklore

df3 %>%
```



Pairwise t-tests

Holm-Bonferroni method

```
##
## Pairwise comparisons using paired t tests
## data: filter(df3, Language == "dar-folklore")$TTR and filter(df3, Language == "dar-folklore")$Volum
##
        tom1
##
                tom2
                        tom3
## tom2 < 2e-16 -
## tom3 < 2e-16 < 2e-16 -
## tom4 6.1e-08 < 2e-16 < 2e-16
## P value adjustment method: holm
p.values <- pairwise.t.test(filter(df3, Language == "kum-folklore")$TTR,
                              filter(df3, Language == "kum-folklore")$Volume,
                              paired = T, p.adjust.method = "holm")
p.values
##
## Pairwise comparisons using paired t tests
## data: filter(df3, Language == "kum-folklore")$TTR and filter(df3, Language == "kum-folklore")$Volum
##
##
        tom1
                tom2
                        tom3
## tom2 < 2e-16 -
## tom3 < 2e-16 1.6e-12 -
## tom4 < 2e-16 < 2e-16 2.5e-05
## P value adjustment method: holm
p.values <- pairwise.t.test(filter(df3, Language == "lak-folklore")$TTR,
                              filter(df3, Language == "lak-folklore")$Volume,
                              paired = T, p.adjust.method = "holm")
p.values
## Pairwise comparisons using paired t tests
## data: filter(df3, Language == "lak-folklore")$TTR and filter(df3, Language == "lak-folklore")$Volum
##
##
        tom1
               tom2
                      tom3
## tom2 <2e-16 -
## tom3 <2e-16 0.14
## tom4 <2e-16 <2e-16 <2e-16
## P value adjustment method: holm
p.values <- pairwise.t.test(filter(df3, Language == "lez-folklore")$TTR,
                              filter(df3, Language == "lak-folklore")$Volume,
                              paired = T, p.adjust.method = "holm")
p.values
```

##

```
## Pairwise comparisons using paired t tests
##

## data: filter(df3, Language == "lez-folklore")$TTR and filter(df3, Language == "lak-folklore")$Volum
##

## tom1 tom2 tom3
## tom2 9.6e-09 - -
## tom3 < 2e-16 2.5e-05 -
## tom4 < 2e-16 < 2e-16 < 2e-16
##

## P value adjustment method: holm</pre>
```