# Measuring Typological and Goegraphical Distances in R

## Yingqi Jing

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## 1 R programming language

### 1.1 Conditional statement (ifelse & case\_when)

```
pi = 3.14
if(is.numeric(pi)){
  print("You have a numeric pi :)")
  print("You do not have a numeric pi :(")
[1] "You have a numeric pi :)"
mylogic = TRUE
if(is.logical(mylogic)){
  print("You have a logical variable :)")
}else{
  print("You do not have a logical variable :(")
[1] "You have a logical variable :)"
lang = "Estonian"
if(is.character(lang) & lang == "Estonian"){
  print("It is a character variable and it is Estonian!")
}else{
  print("It is not Estonian!")
[1] "It is a character variable and it is Estonian!"
lang = "Estonian"
if(is.character(lang) | lang == "Estonian"){
  print("It is a character or it is Estonian!")
}else{
  print("It is not Estonian!")
```

[1] "It is a character or it is Estonian!"

```
lang_df = data.frame(id = 1:5,
          lang = c("Estonian", "Finnish", "English",
                   "Chinese", "Esperanto"))
lang_df
  id
          lang
1 1 Estonian
2 2
      Finnish
3 3
      English
4 4
       Chinese
5 5 Esperanto
lang_df %>%
  mutate(family = ifelse(lang == "Estonian", "Estonian", "not Estonian"))
  id
          lang
                     family
1 1 Estonian
                   Estonian
2 2
      Finnish not Estonian
3 3
      English not Estonian
       Chinese not Estonian
4
  4
5 5 Esperanto not Estonian
We can generalize the ifelse into a case of multiple conditions, so that you have not only two options but
multiple options.
lang_df %>%
  mutate(family = case_when(lang %in% c("Estonian", "Finnish") ~ "Uralic",
                            lang %in% c("English") ~ "IE",
                            lang %in% c("Chinese") ~ "Sino-Tibetan",
                            TRUE ~ "unknown"))
```

## 1.2 For loop (for and rowwise)

```
names = c("Yingqi", "John", "Emily", "David")
for(name in names){
  print(nchar(name))
Γ17 6
```

[1] 4

[1] 5

[1] 5

Dplyr provides a convenient rowwise operation to work on each row of a data frame. This can be viewed as an alternative to for loop in a data.frame. Note: rowwise groups your data by row (class: rowwise\_df), and it is best to ungroup immediately.<sup>1</sup>

To facilitate rowwise operation, you may need to use c\_across to select specific columns. For example, if you want to calculate the number of characters of language names in a data.frame.

```
mydf = data.frame(id = 1:4,
           names = c("Yingqi", "John", "Emily", "David"))
name_length = mydf %>%
 rowwise() %>%
  mutate(nchar_name = nchar(c_across(names))) %>%
  ungroup
name_length
```

```
# A tibble: 4 x 3
     id names nchar_name
  <int> <chr>
                    <int>
     1 Yingqi
1
2
     2 John
                        4
3
      3 Emily
                        5
      4 David
                        5
```

<sup>&</sup>lt;sup>1</sup>https://medium.com/p/da3638b5f46c

## 1.3 Function

```
myaverage = function(numvec){
   average = sum(numvec)/length(numvec)
   return(average)
}

myvec = c(2, 3)
myaverage(myvec)

[1] 2.5

name_length %>%
   pull(nchar_name) %>%
   myaverage()
```

[1] 5

## 2 Typological similarities between Uralic languages

#### 2.1 Function

```
typological_sim = function(data = ut_final, x, y){
   subdata = data %>%
   dplyr::select(-matches("subfamily|area", ignore.case = T)) %>%
   column_to_rownames(var = "Name")
   sim = sum(abs(as.vector(subdata[x, ]) == as.vector(subdata[y, ])))
   sim_p = sim/ncol(subdata)
   return(sim_p)
}
```

#### 2.2 Data preprocessing

```
uratyp_df = read.csv("../data/values.csv")
lang_df = read.csv("../data/languages.csv")
ut_data = uratyp_df %>%
   inner_join(., lang_df, by = c("Language_ID" = "ID")) %>%
   dplyr::select(Name, Parameter_ID, Value, Subfamily) %>%
   filter(grepl("UT", Parameter_ID))
```

(1) Convert all data into binary (0, 1)

```
ut_wide = ut_data %>%
mutate(Value = case_when(
   Value == "0" ~ OL,
   Value == "1" ~ 1L,
   TRUE ~ NA_integer_ # convert all "?" into NA
)) %>%
pivot_wider(., names_from = Parameter_ID, values_from = Value)
```

(2) Remove all columns with missing values

```
ut_wide = ut_wide %%
select_if(function(x) !any(is.na(x)))
# alternatively, select_if(~ !any(is.na(.x)))
# select_if(~ sum(is.na(.x)) == 0)
# select(where(~ sum(is.na(.x)) == 0))
```

(3) Remove all constant columns

```
ut_final = ut_wide %>%
  remove_constant(.)
  # select_if(~ length(unique(.x)) > 1)
head(ut_final)
```

# A tibble: 6 x 143 Subfamily UT001 UT002 UT003 UT004 UT005 UT006 UT007 UT008 UT009 UT010 Name <chr>> 1 Central~ Finnic 2 Courlan~ Finnic 3 East\_Ma~ Ugric 4 Erzya Mordvin 5 Estonian Finnic 6 Finnish Finnic 

```
# ... with 131 more variables: UT011 <int>, UT012 <int>, UT013 <int>,
   UT015 <int>, UT016 <int>, UT017 <int>, UT020 <int>, UT021 <int>,
   UT022 <int>, UT023 <int>, UT024 <int>, UT025 <int>, UT026 <int>,
   UT028 <int>, UT029 <int>, UT031 <int>, UT032 <int>, UT034 <int>,
   UT035 <int>, UT036 <int>, UT037 <int>, UT038 <int>, UT039 <int>,
   UT042 <int>, UT043 <int>, UT044 <int>, UT045 <int>, UT047 <int>,
   UT048 <int>, UT049 <int>, UT050 <int>, UT051 <int>, UT052 <int>, ...
 (4) reshape the data into long format
lang sorted = ut final %>%
  arrange(Subfamily) %>%
  pull(Name)
ut_final_long = ut_final %>%
  pivot_longer(., names_to = "feature", values_to = "value", -c("Name", "Subfamily")) %>%
  mutate(value = factor(value),
         Name = factor(Name, levels = lang_sorted))
head(ut_final_long)
# A tibble: 6 x 4
  Name
               Subfamily feature value
  <fct>
               <chr>
                         <chr>
                                 <fct>
1 Central_Veps Finnic
                         UT001
2 Central_Veps Finnic
                         UT002
                                 1
3 Central_Veps Finnic
                         UT003
4 Central_Veps Finnic
                         UT004
                                 1
5 Central_Veps Finnic
                         UT005
                                 1
6 Central_Veps Finnic
                         UT006
                                 0
 (5) Visualize data via heatmap
ut_final_long %>%
  ggplot(., aes(x = feature, y = Name, fill = value)) +
  geom_tile() +
  scale_fill_manual(values = alpha(c("blue", "red"), 0.65)) +
  theme(axis.text.x = element_text(angle = 90, size = 6, hjust = 0),
        axis.ticks = element blank()) +
  labs(x = NULL, y = NULL)
```

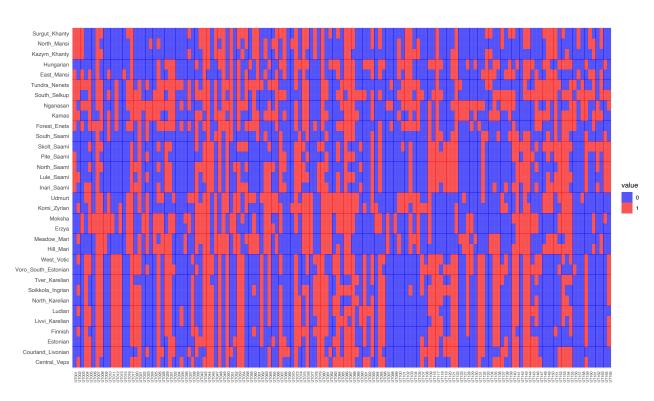
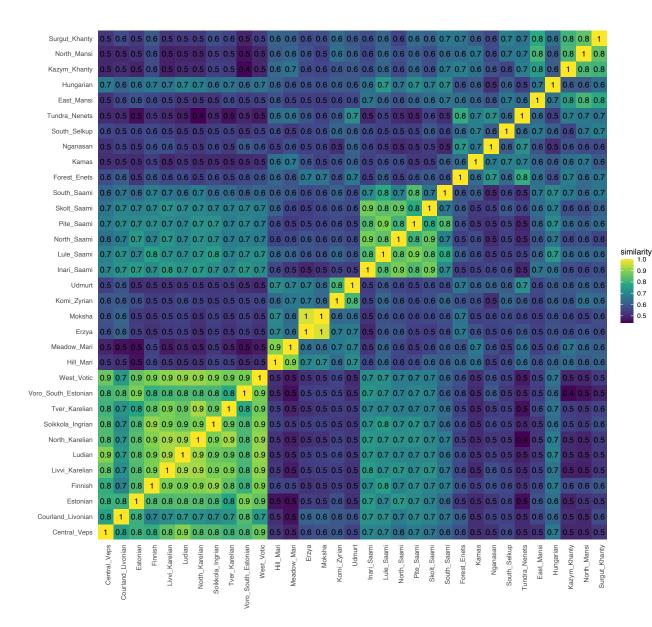


Figure 1: Overview of typological data in UT database

### 2.3 Calculating typological similarities

```
lgs = ut_final$Name
lgs_sim = expand.grid(lgs, lgs, stringsAsFactors = F) %>%
 rowwise() %>%
  mutate(similarity = typological_sim(ut_final, Var1, Var2))
lgs_sim_sorted = lgs_sim %>%
  mutate(Var1 = factor(Var1, levels = lang_sorted),
        Var2 = factor(Var2, levels = lang_sorted))
ggplot(lgs_sim_sorted, aes(Var1, Var2, fill = similarity)) +
  geom tile() +
  geom_text(data = lgs_sim_sorted,
            mapping = aes(Var1, Var2,
                          label = round(similarity, digit = 1))) +
  scale_fill_continuous(type = "viridis") +
  labs(x = NULL, y = NULL) +
  theme(plot.title = element_text(hjust = 0.5, face = "bold"),
        axis.text.y = element_text(size = 9),
       axis.text.x = element_text(angle = 90, size = 9, hjust = 1),
       axis.ticks = element_blank(),
       panel.grid.major = element_blank(),
        panel.grid.minor = element_blank(),
       panel.background = element_blank()) +
  coord fixed()
```



#### 2.4 Exercises

(1) Pls add the information of typological areas (parameters.csv) to UT dataset. Note: you can use the preprocessed UT data ut\_final\_long.

```
feat_areas = read.csv("../Data/parameters.csv")
feat_areas = feat_areas %>%
  dplyr::select(ID, Area)
ut_areas = ut_final_long %>%
  inner_join(., feat_areas, by = c("feature" = "ID"))
```

(2) Pls calculate the typological similarities between languages across typological areas (phonology, morphology and syntax) in UT dataset, and plot them as heatmaps separately. Note: you can remove the lexicon features, and use facet\_wrap function to create subpanels.

```
ut_area_sim = ut_areas %>%
  filter(Area %in% c("Phonology", "Morphology", "Syntax")) %>%
  split(.$Area) %>%
  map_dfr(., ~{subdata = .x %>% spread(., key = feature, value = value)
               expand.grid(lgs, lgs, stringsAsFactors = F) %>%
                 rowwise() %>%
                 mutate(similarity = typological_sim(subdata, Var1, Var2)) %>%
                 mutate(Area = subdata$Area[1])})
ut_area_sim_sorted = ut_area_sim %>%
  mutate(Var1 = factor(Var1, levels = lang_sorted),
         Var2 = factor(Var2, levels = lang_sorted),
         Area = factor(Area, levels = c("Phonology", "Morphology", "Syntax")))
ut area sim sorted %>%
  ggplot(., aes(Var1, Var2, fill = similarity)) +
  geom_tile() +
  # geom_text(data = lgs_sim_sorted,
             mapping = aes(Var1, Var2,
                            label = round(similarity, digit = 1))) +
  scale fill continuous(type = "viridis") +
  facet_wrap(~Area) +
  # scale_x_discrete(position = "bottom") +
  labs(x = NULL, y = NULL) +
  theme(plot.title = element_text(hjust = 0.5, face = "bold"),
        axis.text.y = element_text(size = 9),
        axis.text.x = element_text(angle = 90, size = 9, hjust = 1),
        axis.ticks = element_blank(),
        panel.grid.major = element_blank(),
        panel.grid.minor = element_blank(),
        panel.background = element_blank()) +
  coord fixed()
```

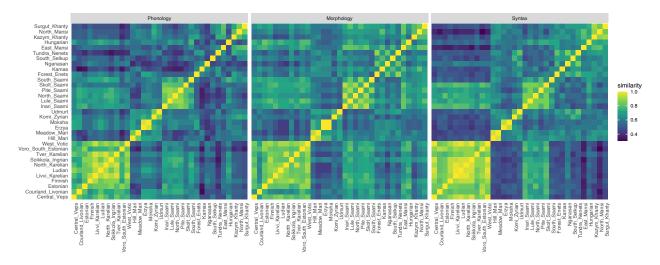


Figure 2: Typological similarities between Uralic languages across different areas in UT data

## 3 Geographical distances between languages

#### 3.1 Function

```
geographical_dist = function(data = lang_geo, x = lang1, y = lang2){
  lang1_location = data[x, ]
  lang2_location = data[y, ]
  return(distHaversine(lang1_location, lang2_location)/1000)
}
```

### 3.2 Measuring geographical distances

```
lang_geo = lang_df %>%
  dplyr::select(Name, Longitude, Latitude) %>%
  column_to_rownames(var = "Name")
geo_dist = expand.grid(lgs, lgs, stringsAsFactors = F) %>%
 rowwise() %>%
  mutate(distance = geographical_dist(data = lang_geo,
                                      x = Var1,
                                      y = Var2)) %>%
  ungroup %>%
  mutate(dist_scaled = distance/max(distance))
geo_dist_sorted = geo_dist %>%
  mutate(Var1 = factor(Var1, levels = lang_sorted),
         Var2 = factor(Var2, levels = lang_sorted))
geo_dist_sorted %>%
  ggplot(., aes(Var1, Var2, fill = distance)) +
  geom_tile() +
  geom_text(data = geo_dist_sorted,
            mapping = aes(Var1, Var2,
                          label = round(dist_scaled, digit = 1))) +
  scale_fill_continuous(type = "viridis", direction = -1) +
  labs(x = NULL, y = NULL) +
  theme(plot.title = element_text(hjust = 0.5, face = "bold"),
        axis.text.y = element_text(size = 9),
        axis.text.x = element text(angle = 90, size = 9, hjust = 1),
       axis.ticks = element_blank(),
        panel.grid.major = element_blank(),
        panel.grid.minor = element_blank(),
       panel.background = element_blank()) +
  coord_fixed()
```

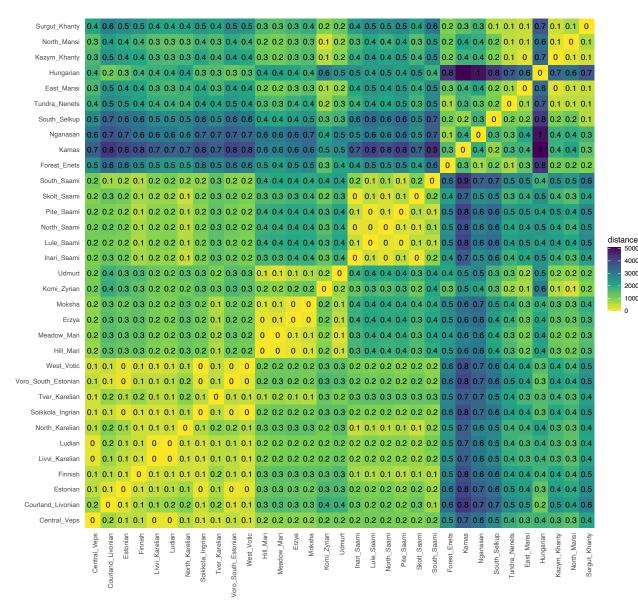


Figure 3: Geographical distances between Uralic languages