Performance of linking graduates

Flavio & Christoph & Mona

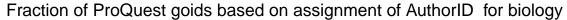
23 July, 2023

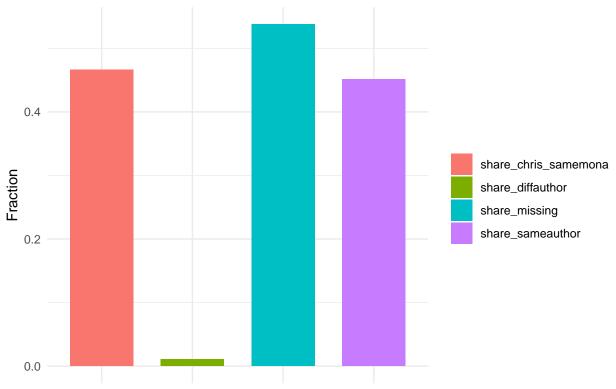
Contents

```
# Function to process the data for a specific field
process_data <- function(field) {</pre>
 # Read the data for the specified field
 if (field %in% c("history")) {
 links_graduates_mona <- read.csv(paste0(datapath, "links_graduates_", field, "_mona_degree0_19852015.c
   filter(link_score>0.7) %>%
   rename(authorid_mona = AuthorId) %>%
   rename(linkscore_mona=link_score)
 } else {
   links_graduates_mona <- read.csv(paste0(datapath, "/links_graduates_", field, "_mona_degree0_1985201
   filter(link_score>0.7) %>%
   rename(authorid_mona = grantid_authorposition) %>%
   rename(goid = AuthorId) %>%
   rename(linkscore_mona=link_score)
}
 if (field %in% c("biology", "computer science", "economics", "engineering", "environmental science", "
 links_graduates_christoph <- read.csv(paste0(datapath,"links_graduates_", field, "_christoph_fielddeg</pre>
   filter(link_score>0.7) %>%
   rename(authorid_christoph = AuthorId) %>%
   rename(linkscore_christoph=link_score)
 links_graduates_christoph <- read.csv(paste0(datapath, "links_graduates_", field, "_christoph_degree0_</pre>
   filter(link_score>0.7) %>%
   rename(authorid_christoph = AuthorId) %>%
   rename(linkscore_christoph=link_score)
 }
links_graduates_mona <- collect(links_graduates_mona)</pre>
links_graduates_christoph <- collect(links_graduates_christoph)</pre>
# Performs the full join: bothlink=1 if same authorID assigned in both, 0 if different authorID assigne
# Then calculates the share of links found by Christoph also found by Mona (number links found by both
```

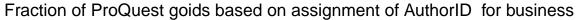
```
links_graduates <- links_graduates_mona %>%
  full_join(links_graduates_christoph, by = c("goid")) %>%
  mutate(
   field = field,
   monalink = ifelse(!is.na(authorid_mona), 1, 0),
    chrislink = ifelse(!is.na(authorid_christoph), 1, 0),
   bothlink = ifelse(is.na(authorid_christoph) | is.na(authorid_mona),
                      ifelse(authorid_christoph == authorid_mona, 1, 0)),
    share_bothlink = sum(bothlink == 1 & chrislink == 1, na.rm = TRUE) / sum(chrislink == 1, na.rm = T
  )
# Look closer at link differences:
# share of ProQuest goids assigned to same AuthorId (share_sameauthor), distinct AuthorId (share_diffau
links_graduates <- links_graduates %>%
mutate(
    share_sameauthor = sum(bothlink == 1, na.rm = TRUE) / n_distinct(goid),
   share_diffauthor = sum(bothlink == 0 & !is.na(bothlink), na.rm = TRUE) / n_distinct(goid),
   share_missing = sum(is.na(bothlink)) / n_distinct(goid),
   share_missing_mona = sum(is.na(bothlink) & monalink == 0) / n_distinct(goid),
   share_missing_chris = sum(is.na(bothlink) & chrislink == 0) / n_distinct(goid),
   share_chris_samemona = sum(chrislink & bothlink==1, na.rm=TRUE) / sum(chrislink, na.rm=TRUE)
# Create table with the shares by field
# Problem: not shown in pdf, ugly table here
shares_table <- links_graduates %>%
  select(field, share_bothlink, share_sameauthor, share_diffauthor, share_missing, share_chris_samemona
  group_by(field) %>%
  summarize(
   share_bothlink = mean(share_bothlink, na.rm = TRUE),
   share_sameauthor = mean(share_sameauthor, na.rm = TRUE),
   share_diffauthor = mean(share_diffauthor, na.rm = TRUE),
   share_missing = mean(share_missing, na.rm = TRUE),
   share_chris_samemona = mean(share_chris_samemona, na.rm = TRUE)
  )
# Select the shares for the bar chart: total number of goids as base
shares_data <- links_graduates %>%
  summarise(
   share_sameauthor = mean(share_sameauthor, na.rm = TRUE),
   share_diffauthor = mean(share_diffauthor, na.rm = TRUE),
   share_missing = mean(share_missing, na.rm = TRUE),
    share_chris_samemona = mean(share_chris_samemona, na.rm = TRUE)
  ) %>%
  gather(variable, value)
# Create the bar chart
bar_chart <- ggplot(shares_data, aes(x = variable, y = value, fill = variable)) +</pre>
```

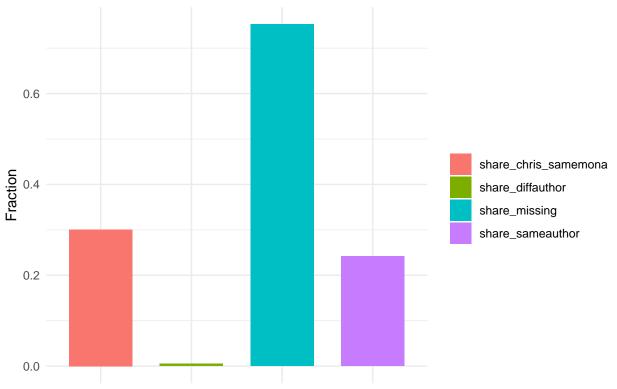
```
geom_bar(stat = "identity", width = 0.7) +
  theme_minimal() +
  labs(
    x = NULL,
    y = "Fraction",
   title = paste("Fraction of ProQuest goids based on assignment of AuthorID for", field),
   fill= NULL
  scale_x_discrete(labels = c("Mona same author ID|Chris linked", "different author ID", " author ID mis
# Print the bar chart
print(bar_chart)
# Print the summary table
shares_table %>%
 kable(format = "html",
        align = c("1", "c", "c", "c", "c", "c"),
        digits = 2,
        caption = "Share Statistics by Field",
        booktabs = TRUE)
}
# Fields to process
# missing fields: ""art, "chemistry", "geography", "history", "mathematics", "medicine", "sociology"
fields_to_process <- c("biology", "business", "computer science", "economics", "engineering", "environm
# Loop through the fields
for (field in fields_to_process) {
 table <- process_data(field)</pre>
  print(table)
  cat("\n\n")
}
```



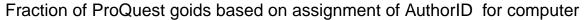


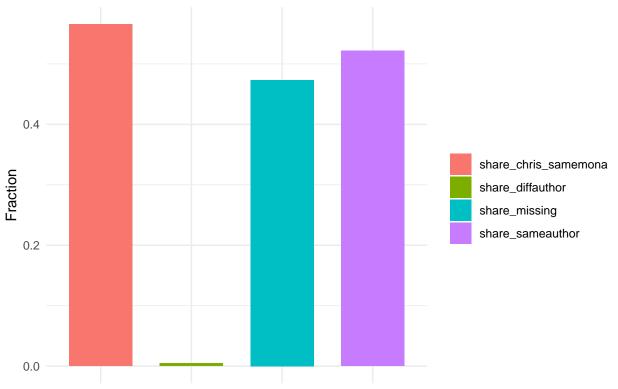
```
## <caption>Share Statistics by Field</caption>
##
 <thead>
##
  field 
##
##
   share_bothlink 
##
   share_sameauthor 
   share_diffauthor 
##
##
   share_missing 
   share_chris_samemona 
##
##
 </thead>
##
## 
##
  biology 
##
   0.47 
##
##
   0.45 
   0.01 
##
   0.54 
##
   0.47 
##
 ##
## 
##
```





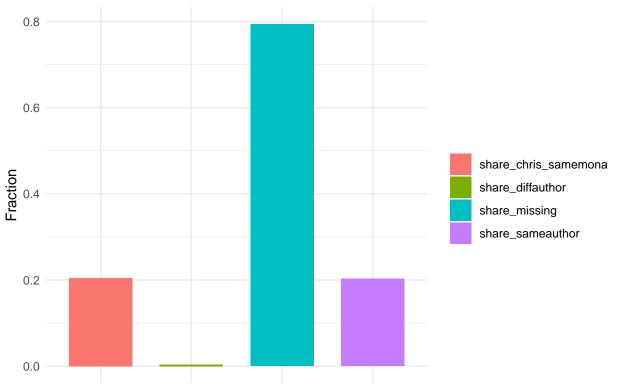
```
## <caption>Share Statistics by Field</caption>
##
 <thead>
##
  field 
##
##
   share_bothlink 
##
   share_sameauthor 
   share_diffauthor 
##
##
   share_missing 
   share_chris_samemona 
##
##
 ##
 </thead>
## 
##
  business 
##
##
   0.3 
##
   0.24 
   0.01 
##
   0.75 
##
   0.3 
##
 ##
## 
##
```





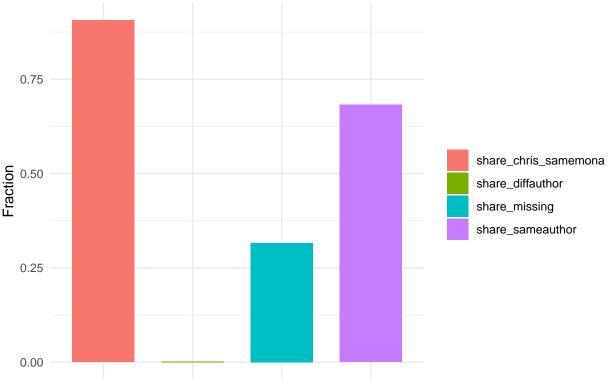
```
## <caption>Share Statistics by Field</caption>
##
 <thead>
##
  field 
##
##
   share_bothlink 
##
   share_sameauthor 
   share_diffauthor 
##
##
   share_missing 
   share_chris_samemona 
##
##
 </thead>
##
## 
##
  computer science 
##
   0.57 
##
##
   0.52 
   0 
##
   0.47 
##
   0.57 
##
 ##
## 
##
```

Fraction of ProQuest goids based on assignment of AuthorID for economic

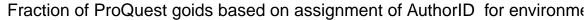


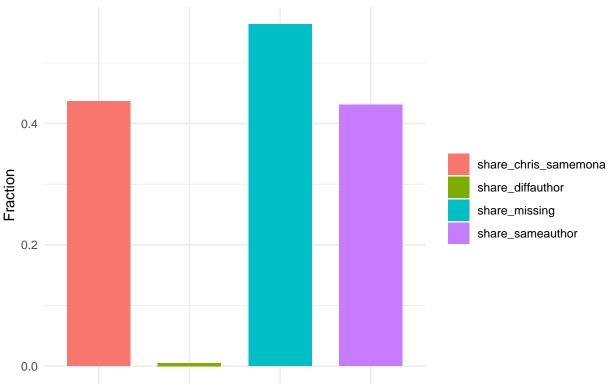
```
## <caption>Share Statistics by Field</caption>
##
 <thead>
##
  field 
##
##
   share_bothlink 
##
   share_sameauthor 
   share_diffauthor 
##
##
   share_missing 
   share_chris_samemona 
##
##
 </thead>
##
## 
##
  economics 
##
##
   0.2 
##
   0.2 
   0 
##
   0.79 
##
   0.2 
##
 ##
## 
##
```

Fraction of ProQuest goids based on assignment of AuthorID for engineer

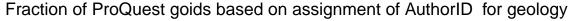


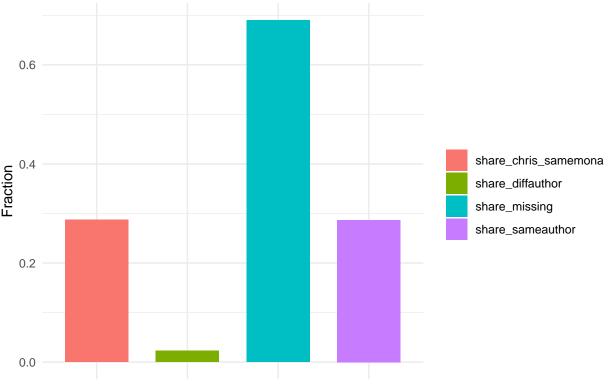
```
## <caption>Share Statistics by Field</caption>
##
 <thead>
##
  field 
##
##
   share_bothlink 
##
   share_sameauthor 
   share_diffauthor 
##
##
   share_missing 
   share_chris_samemona 
##
##
 </thead>
##
## 
 ##
##
   engineering 
##
   0.91 
##
   0.68 
   0 
##
   0.32 
##
   0.91 
##
 ##
## 
##
```



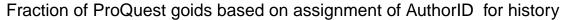


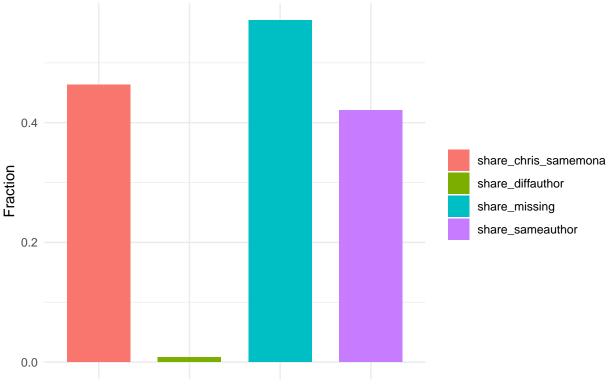
```
## <caption>Share Statistics by Field</caption>
##
 <thead>
##
  field 
##
##
   share_bothlink 
##
   share_sameauthor 
   share_diffauthor 
##
##
   share_missing 
   share_chris_samemona 
##
##
 </thead>
##
## 
 ##
   environmental science 
##
##
   0.44 
##
   0.43 
   0.01 
##
   0.56 
##
   0.44 
##
 ##
## 
##
```



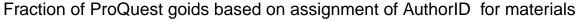


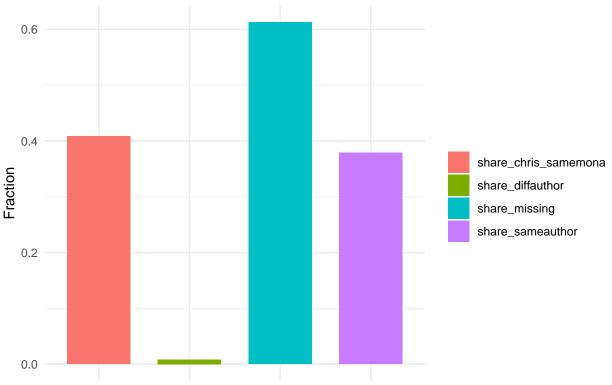
```
## <caption>Share Statistics by Field</caption>
##
 <thead>
##
  field 
##
##
   share_bothlink 
##
   share_sameauthor 
   share_diffauthor 
##
##
   share_missing 
   share_chris_samemona 
##
##
 </thead>
##
## 
 ##
   geology 
##
   0.29 
##
##
   0.29 
   0.02 
##
   0.69 
##
   0.29 
##
 ##
## 
##
```



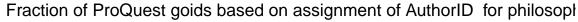


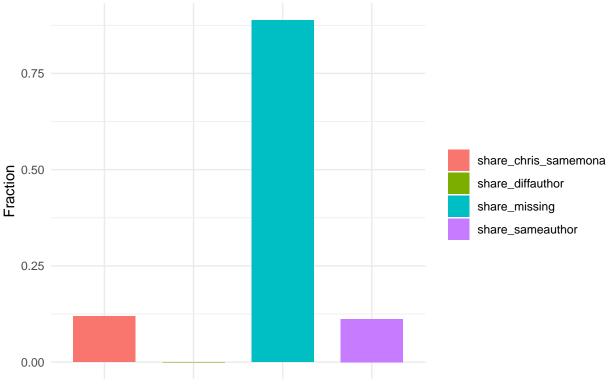
```
## <caption>Share Statistics by Field</caption>
##
 <thead>
##
  field 
##
##
   share_bothlink 
##
   share_sameauthor 
   share_diffauthor 
##
##
   share_missing 
   share_chris_samemona 
##
##
 ##
 </thead>
## 
 ##
   history 
##
   0.46 
##
##
   0.42 
   0.01 
##
   0.57 
##
   0.46 
##
 ##
## 
##
```



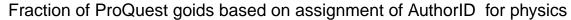


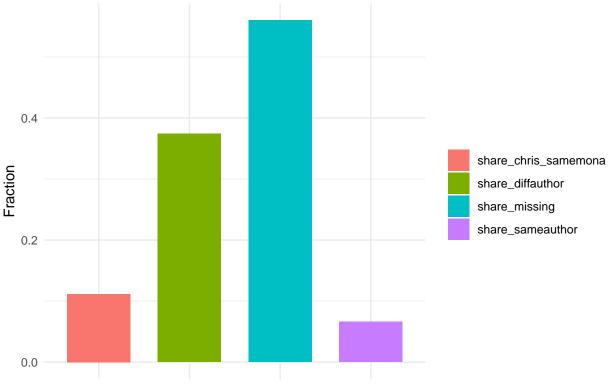
```
## <caption>Share Statistics by Field</caption>
##
 <thead>
##
  field 
##
##
   share_bothlink 
##
   share_sameauthor 
   share_diffauthor 
##
##
   share_missing 
   share_chris_samemona 
##
##
 </thead>
##
## 
 ##
   materials science 
##
   0.41 
##
##
   0.38 
   0.01 
##
   0.61 
##
   0.41 
##
 ##
## 
##
```



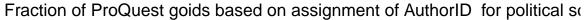


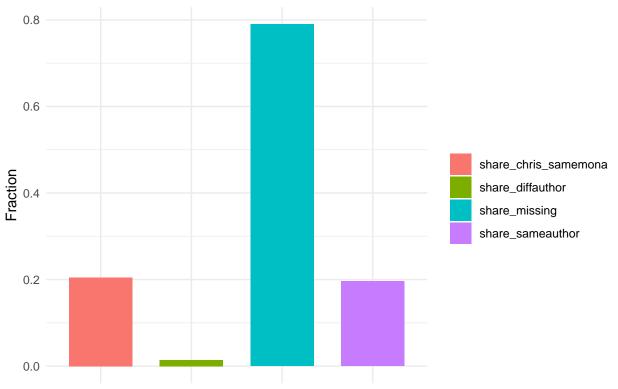
```
## <caption>Share Statistics by Field</caption>
##
 <thead>
##
  field 
##
##
   share_bothlink 
##
   share_sameauthor 
   share_diffauthor 
##
##
   share_missing 
   share_chris_samemona 
##
##
 </thead>
##
## 
 ##
   philosophy 
##
   0.12 
##
##
   0.11 
   0 
##
   0.89 
##
   0.12 
##
 ##
## 
##
```



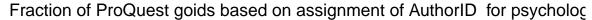


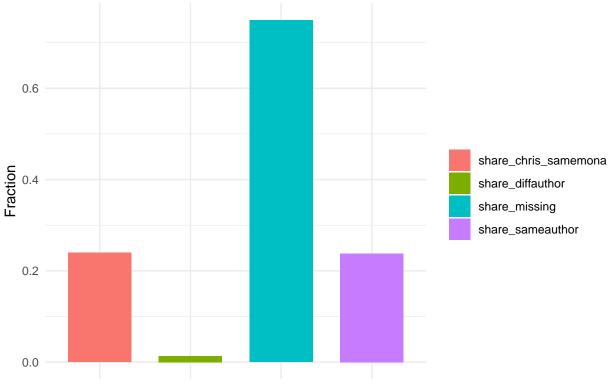
```
## <caption>Share Statistics by Field</caption>
##
 <thead>
##
  field 
##
##
   share_bothlink 
##
   share_sameauthor 
   share_diffauthor 
##
##
   share_missing 
   share_chris_samemona 
##
##
 ##
 </thead>
## 
 ##
##
   physics 
   0.11 
##
##
   0.07 
   0.37 
##
   0.56 
##
   0.11 
##
 ##
## 
##
```





```
## <caption>Share Statistics by Field</caption>
##
 <thead>
##
  field 
##
##
   share_bothlink 
##
   share_sameauthor 
   share_diffauthor 
##
##
   share_missing 
   share_chris_samemona 
##
##
 </thead>
##
## 
 ##
   political science 
##
   0.2 
##
##
   0.2 
   0.01 
##
   0.79 
##
   0.2 
##
 ##
## 
##
```





```
## <caption>Share Statistics by Field</caption>
##
 <thead>
##
 ##
   field 
##
   share_bothlink 
##
   share_sameauthor 
   share_diffauthor 
##
   share_missing 
##
   share_chris_samemona 
##
##
 </thead>
##
## 
 ##
##
   psychology 
##
   0.24 
##
   0.24 
   0.01 
##
##
   0.75 
   0.24 
##
 ##
## 
##
```

- many missings between Christoph's and Mona's links
- share of Mona's links compared to Christoph's links low (share_bothlink) but mostly due to missings

- and different author assignment
- in most fields, goids linked to the same authors, only few that were linked to different ones
- exception in physics, most authors linked differently, why? (no obvious mistakes when renaming variables and joining the datasets)

Now do the same just for physics but for christoph, mona and flavio

```
process_data <- function(field) {</pre>
  # Reads data for specified field
  # Read the data for Mona
  links_graduates_mona <- read.csv(paste0(datapath, "links_graduates_", field, "_mona_degree0_19852015.c
    filter(link_score>0.7) %>%
   rename(authorid_mona = grantid_authorposition) %>%
   rename(goid = AuthorId) %>%
   rename(linkscore_mona=link_score)
  # Read the data for Flavio
  links_graduates_flavio <- read.csv(paste0(datapath, "links_graduates_", field, "_flavio_degree0_198520
   filter(link_score>0.7) %>%
    rename(authorid_flavio = AuthorId) %>%
   rename(linkscore_flavio=link_score)
  # Read the data for Christoph
  links_graduates_christoph <- read.csv(paste0(datapath, "links_graduates_", field, "_christoph_degree0_</pre>
   filter(link_score>0.7) %>%
   rename(authorid_christoph = AuthorId) %>%
   rename(linkscore_christoph=link_score)
  links_graduates_mona <- collect(links_graduates_mona)</pre>
  links_graduates_flavio <- collect(links_graduates_flavio)</pre>
  links_graduates_christoph <- collect(links_graduates_christoph)</pre>
  # Join the data
  links_graduates <- links_graduates_mona %>%
    full_join(links_graduates_flavio, by = c("goid")) %>%
   full_join(links_graduates_christoph, by = c("goid")) %>%
   mutate(
      field = field,
      mona_same_as_chris = ifelse(authorid_mona == authorid_christoph, 1, 0),
      flavio_same_as_chris = ifelse(authorid_flavio == authorid_christoph, 1, 0),
      mona_diff_from_chris = ifelse(authorid_mona != authorid_christoph & !is.na(authorid_mona) & !is.n
      flavio_diff_from_chris = ifelse(authorid_flavio != authorid_christoph & !is.na(authorid_flavio) &
   ) %>%
   replace_na(list(mona_same_as_chris = 0, flavio_same_as_chris = 0, mona_diff_from_chris = 0, flavio_
   mutate(
      # Calculate shares
      share_mona_same_as_chris = mean(mona_same_as_chris, na.rm = TRUE),
      share_flavio_same_as_chris = mean(flavio_same_as_chris, na.rm = TRUE),
      share_mona_diff_from_chris = mean(mona_diff_from_chris, na.rm = TRUE),
      share_flavio_diff_from_chris = mean(flavio_diff_from_chris, na.rm = TRUE)
```

```
# Create table with the shares by field
  shares_table <- links_graduates %>%
    select(field, share_mona_same_as_chris, share_flavio_same_as_chris, share_mona_diff_from_chris, sha
    group_by(field) %>%
    summarize(
      share_mona_same_as_chris = mean(share_mona_same_as_chris, na.rm = TRUE),
      share_flavio_same_as_chris = mean(share_flavio_same_as_chris, na.rm = TRUE),
      share_mona_diff_from_chris = mean(share_mona_diff_from_chris, na.rm = TRUE),
      share_flavio_diff_from_chris = mean(share_flavio_diff_from_chris, na.rm = TRUE)
    )
  # Return the table
  shares_table %>%
    kable(format = "latex",
          align = c("1", "c", "c", "c", "c"),
          digits = 2,
          caption = "Share Statistics by Field",
          booktabs = TRUE)
}
# Specify the field to process
field_to_process <- "physics"</pre>
# Process the data
table <- process_data(field_to_process)</pre>
# Print the table
print(table)
## \begin{table}
##
## \caption{\label{tab:unnamed-chunk-5}Share Statistics by Field}
## \centering
## \begin{tabular}[t]{lcccc}
## \toprule
## field & share\_mona\_same\_as\_chris & share\_flavio\_same\_as\_chris & share\_mona\_diff\_from\_chr
## \midrule
## physics & 0.06 & 0.5 & 0.36 & 0.03\\
## \bottomrule
## \end{tabular}
## \end{table}
```