Code for Variety and Mainstays of the R Developer Community

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- Text Mining
 - 1 Data Cleaning
 - 1.1 Length of the descriptions
 - 1.2 Convert upper cases, delete web links and doi
 - 1.3 Lemmatization
 - 1.4 Delete numbers and symbols
 - 1.5 remove common stopwords
 - 2 Frequency Analysis
 - 2.1 Word Frequency
 - 2.2 Phrase Frequency
 - o 3 Topic Modeling
 - 3.1 Determine the number of topics
 - **3.2 Topic 20**
- Network Analysis
 - o 4 Package dependency network and author collaboration network
 - 4.1 Build network and set edges
 - 4.2 Measures of Influence
 - 4.3 Sensitivity Analysis
 - 4.4 Select Important Packages and Authors
 - 4.5 Visualization
 - 4.6 Correlation among importance indexes and downloads
 - 5 Bipartite Network
 - 5.1 Build bipartite network and set edges (weights)
 - 5.2 Visulization
 - 5.3 Centrality

Here we provided the code and data for the paper titled "Variety and Mainstays of the R Developer Community".

```
## package and function perparation
library(dplyr)
library(tidytext)
library(janeaustenr)
library(ggplot2)
library(scales)
library(textdata)
library(tidyr)
library(igraph)
library(ggraph)
library(tm)
library(stringr)
library (wordcloud)
library(topicmodels)
library(wordcloud2)
library(spacyr) # for lemmatizing verbs / aux by importing python pa
        ckages
library(SemNetCleaner) # for lemmatizing nouns
# function for save data
write.csv.utf8.BOM <- function(df, filename) {</pre>
    con <- file(filename, "w")</pre>
    tryCatch({
        for (i in 1:ncol(df))
            df[,i] = iconv(df[,i], to = "UTF-8")
        writeChar(iconv("\ufeff", to = "UTF-8"), con, eos = NULL)
        write.csv(df, file = con, row.names=FALSE)
    }, finally = {close(con)})
# function for spliting the name and email of maintainer
split mt <- function(maintainer) {</pre>
  mt2name email = strsplit(maintainer, '<')[[1]]</pre>
  name = mt2name email[1]
  email = strsplit(mt2name email[2],'>')[[1]]
  return(mt name email = list(name = name, email = email))
```

Extracted the data from CRAN on 2022-11-27.

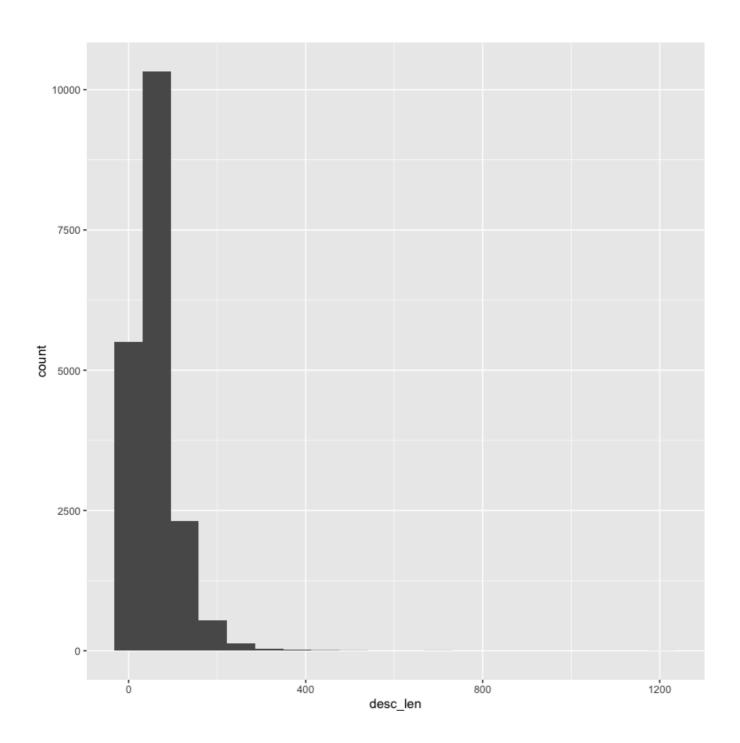
Load the extracted data which can be obtained from https://github.com/zhanglj37/R_Developer_Community.

```
load("pkg_221127.RData")
```

Text Mining

1 Data Cleaning

1.1 Length of the descriptions



```
summary(desc_len)
```

```
## Min. 1st Qu. Median Mean 3rd Qu. Max.
## 1.00 28.00 49.00 60.15 78.00 1207.00
```

1.2 Convert upper cases, delete web links and doi

```
desc[,"Description"] = tolower(desc[,"Description"])
# delete links
```

1.3 Lemmatization

```
desc sentence = desc[, "Description"]
names(desc sentence) = desc[,"Package"]
# lemmatize verbs (applied, applies --> apply) and aux (was --> be)
         using spacyr spacy parse()
# lemmatize NOUNs using SemNetCleaner singularize()
# https://spacy.io/usage
desc token = spacy parse(desc sentence)
desc token2 = desc token
for (i in 1:nrow(desc token2)){
    if (desc_token2[i,"pos"] == "VERB" || desc_token2[i,"pos"] == "AUX" )
        desc token2[i,"lemma"] = desc token[i,"lemma"]
    }else if(desc token2[i,"pos"]=="NOUN") {
        desc token2[i,"lemma"] = singularize(as.character(desc token
        2[i, "token"]))
    }else if(desc token2[i,"pos"]=="PROPN") {
        desc token2[i,"lemma"] = singularize(as.character(desc token
        2[i,"token"]))
    }else if(desc token2[i,"pos"]=="PRON") {
        desc token2[i,"lemma"] = singularize(as.character(desc token
        2[i, "token"]))
    }else{
        desc token2[i,"lemma"] = desc token[i,"token"]
}
colnames(desc token2) = c("doc id", "sentence id", "token id", "toke
        n", "word", "pos", "entity")
```

1.4 Delete numbers and symbols

```
delete pos = c("PUNCT", "NUM", "SYM")
pos loc1 = which(desc token2[,"pos"] == delete pos[1])
pos loc2 = which(desc token2[,"pos"] == delete pos[2])
pos loc3 = which(desc token2[,"pos"]==delete pos[3])
loc token = c(1:nrow(desc token2))
# detect numbers # numbers that are not detected by spacy sparse
numbers only <- function(x) !grepl("\\D", x)</pre>
loc number = loc token[numbers only(desc token2[,"word"])]
# detect space
space only <- function(x) !grepl("\\S", x)</pre>
loc space = loc token[space only(desc token2[,"word"])]
# detect symbols # symbols that are not detected by spacy sparse
no str num <- function(x) !grepl("\\w", x)</pre>
loc no str num = loc token[no str num(desc token2[,"word"])]
one char1 <- function(x) grepl("^\\w\\\\", x)</pre>
one char2 <- function(x) grepl("^{\w}, x)
loc onechar1 = loc token[one_char1(desc_token2[,"word"])]
loc onechar2 = loc token[one char2(desc token2[,"word"])]
pos delete = c(pos loc1, pos loc2, pos loc3, loc number, loc space,
         loc no str num, loc onechar1, loc onechar2)
token clean = desc token2[-pos delete,]
token clean = tibble(token clean)
```

1.5 remove common stopwords

```
## Joining, by = "word"
```

```
# TF-IDF value
#token_count = token_nostop %>%
# count(doc_id, word)

#total_words = token_count %>%
# group_by(doc_id) %>%
# summarize(total = sum(n))

#token_count2 <- left_join(token_count, total_words)

#token_tf_idf <- token_count2 %>%
# bind_tf_idf(word, doc_id, n)

#temp = token_tf_idf %>%
# select(-total) %>%
# arrange(desc(tf_idf))
```

2 Frequency Analysis

2.1 Word Frequency

```
word_totals <- token_nostop %>%
  count(word, sort = TRUE)
```

```
produce change expression
                                                                                                                                                                                        dynamic
                                                                                count<sub>normal</sub> clustering
                                                                                                                                                                                         visualize
                                            observation calculation functionality graphical
                                                   association estimates input population search
                     combine common component correlation
                              space learning apply object continuous study add likelihood generalize conditional interface inference simulate
                             variance develop compute values result standard typ optimization
      write build matrix multiple parameter power spatial information estimate estimate records access error designplot algorithm define format linear roject gene process time define format linear multiple parameter estimate estimate roject gene process time define format linear roject gene process time define format linear multiple parameter estimate estimation multiple parameter estimate reconstruction access access error designplot algorithm text properties and properties application in the properties of the parameter estimate estimation access access error designplot algorithm text properties application in the properties of the parameter estimate estimation access access error designplot algorithm text properties application in the properties application in the properties and properties application in the properties and properties application in the properties access access access access access access access access access time access and properties application in the properties access acce
                         gaussian packages
project control variables tool mo
 miss obtain : estimation
       performance event interval modeling individual genetic testing decrease series map out apus matric software factor probability multivariate prediction
                                                                                                                                                                                        versionform easyspecie
                                                                       testing density multivariate prediction robust estimator structure binary shiny glog
                                                                 distance
                                                                        classification specific select parametric
                                                                                        coefficient identify confidence
                                                                                        download computationequation
```

2.2 Phrase Frequency

Get Phrases and Frequency

save the frequency results and select meaningful phrases manully

```
#### 2-gram
desc lemma.2gram <- unnest tokens(desc, word, Description, token =</pre>
        "ngrams", n = 2)
desc lemma.2gram %>% filter(word=="p value" | word == "p values") %
        >% count(word, sort = T)
##
       word n
## 1 p values 233
## 2 p value 161
desc lemma.2count <- desc lemma.2gram %>% separate(word, c("word1",
        "word2"), sep = " ") %>%
    filter(!word1 %in% stop words$word & !word2 %in% stop words$wor
        d) %>% # remove phrase consist with all stop words
    unite (word, word1, word2, sep = " ") %>%
    count(word, sort = TRUE)
head (desc lemma.2count)
##
                   word n
## 1
           time series 1018
             data sets 523
## 2
## 3 maximum likelihood 410
## 4 regression models 395
## 5 data analysis 393
## 6
           monte carlo 369
#write.csv(desc lemma.2count, 'freq/2word freq.csv', row.names=F)
#### 3-gram
desc lemma.3gram <- unnest tokens(desc, word, Description, token =</pre>
        "ngrams", n = 3)
desc lemma.3count <- desc lemma.3gram %>% separate(word, c("word1",
        "word2", "word3"), sep = " ") %>%
    filter(!word1 %in% stop words$word & !word2 %in% stop words$word
        & !word3 %in% stop words$word) %>% # remove phrase consist
         with all stop words
    unite(word, word1, word2, word3, sep = " ") %>%
    count(word, sort = TRUE)
```

head(desc lemma.3count)

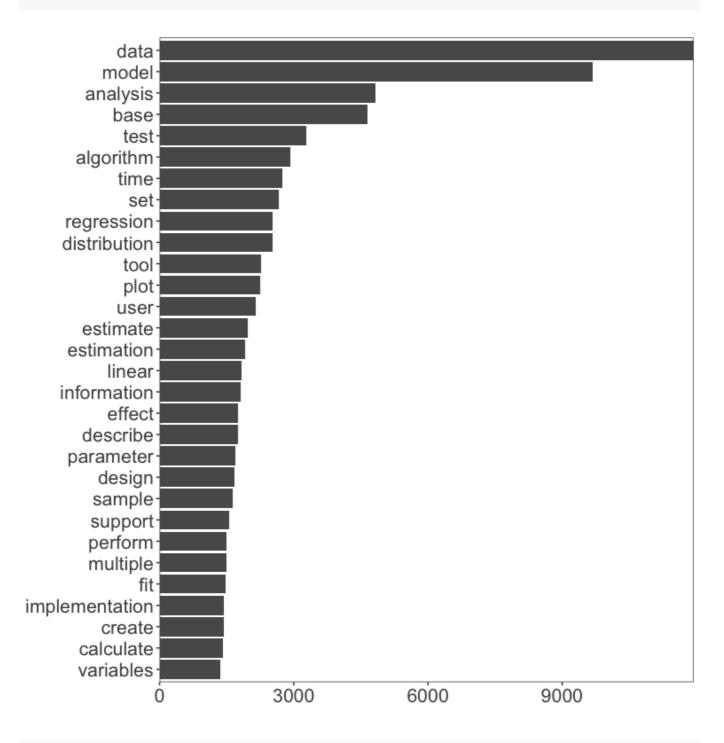
Frequency Figures

```
word1_plot = word_totals %>%
   top_n(30) %>%
   mutate(word = reorder(word, n)) %>%
   ggplot(aes(word, n)) +
   theme(axis.title.x = element_text(face = 'bold'), axis.title.y =
        element_text(face = 'bold'), axis.title = element_text(face
        = 'bold'), text=element_text(size=20)) +
   geom_col(show.legend = FALSE) +
   coord_flip() +
```

```
scale_y_continuous(expand = c(0,0)) +
labs(x = NULL, y = NULL) +
theme(panel.grid=element_blank())
```

```
## Selecting by n
```

```
word1 plot
```

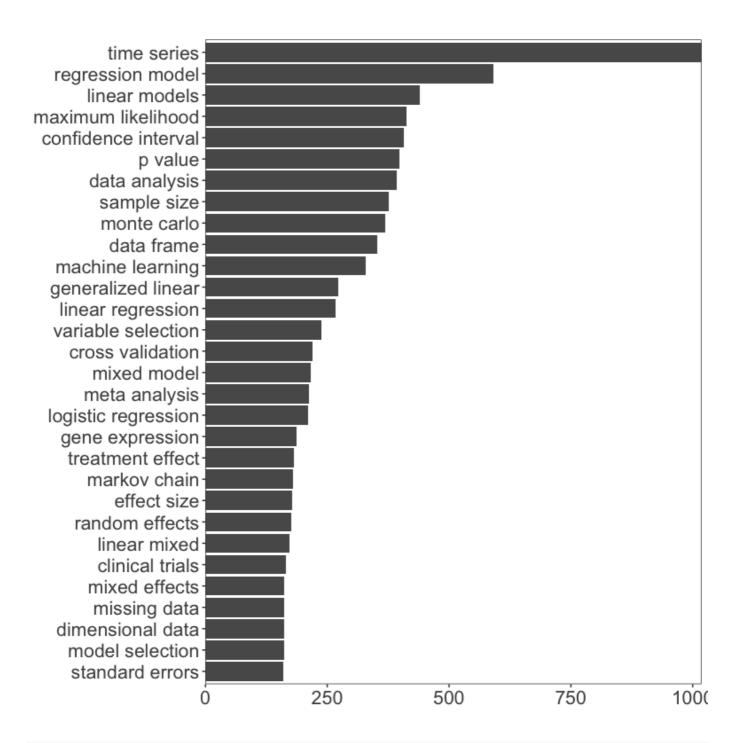


```
word2_plot = word2 %>%
  top_n(30) %>%
  mutate(term = reorder(term, frequency)) %>%
```

```
ggplot(aes(term, frequency)) +
theme(axis.title.x = element_text(face = 'bold'), axis.title.y =
        element_text(face = 'bold'), axis.title = element_text(face
        = 'bold'), text=element_text(size=20)) +
geom_col(show.legend = FALSE) +
coord_flip() +
scale_y_continuous(expand = c(0,0)) +
labs(x = NULL, y = NULL) +
theme(panel.grid=element_blank())
```

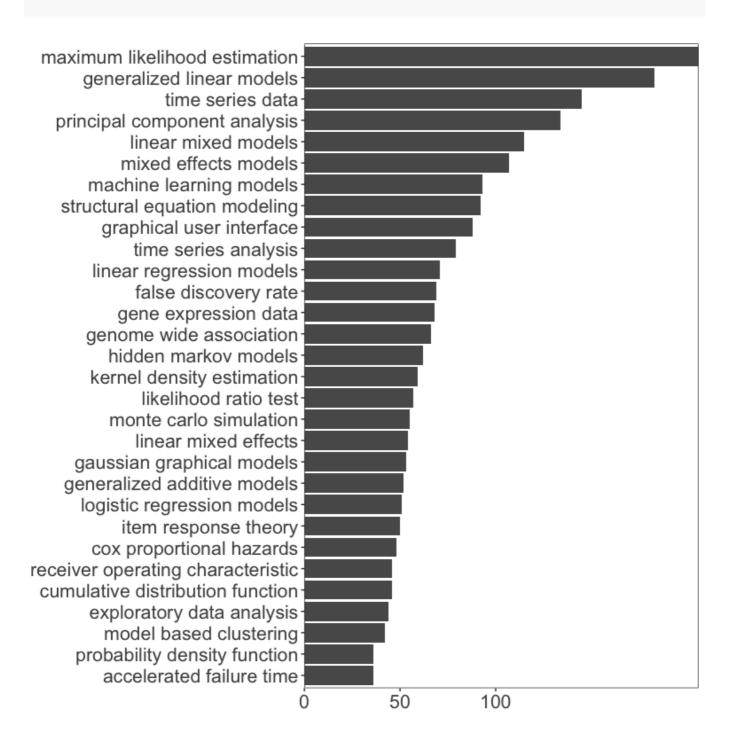
```
## Selecting by frequency
```

```
word2 plot
```



```
word3_plot = word3 %>%
  top_n(30) %>%
  mutate(term = reorder(term, frequency)) %>%
  ggplot(aes(term, frequency)) +
  theme(axis.title.x = element_text(face = 'bold'), axis.title.y =
        element_text(face = 'bold'), axis.title = element_text(face
        = 'bold'), text=element_text(size=20)) +
  geom_col(show.legend = FALSE) +
  coord_flip() +
  scale_y_continuous(expand = c(0,0),breaks = c(0, 50,100)) +
  labs(x = NULL, y = NULL) +
  theme(panel.grid=element_blank())
```

word3 plot



```
ggsave("freq/word1.pdf", word1_plot, dpi = 400,width=5, height = 7)
ggsave("freq/word2.pdf", word2_plot, dpi = 400,width=5, height = 7)
ggsave("freq/word3.pdf", word3_plot, dpi = 400,width=5, height = 7)
```

3 Topic Modeling

3.1 Determine the number of topics

Since the cross-validation is very time-consuming, here we just show the code but didn't run it.

```
# data perparation
library(topicmodels)
desc_dtm_select = token_nostop[,c("doc_id","word")]
desc_dtm_select2 = desc_dtm_select %>%
    group_by(doc_id) %>%
    count(word) %>% ungroup()

desc_dtm = desc_dtm_select2 %>% cast_dtm(doc_id, word, n)
tm::inspect(desc_dtm)
```

```
## <<DocumentTermMatrix (documents: 18898, terms: 42268)>>
## Non-/sparse entries: 437158/798343506
## Sparsity
           : 100%
## Maximal term length: 65
## Weighting : term frequency (tf)
## Sample
##
            Terms
## Docs algorithm analysis base data distribution model regre
## arthistory
             0
                          0 0 1
  ff
                           0
                              0
                                             0
##
                   0
                                  6
                                                  0
                              0 10
  frailtypack
                   0
                           1
                                             2
                                                 26
##
##
  KoulMde
                   0
                          4
                              3
                                  1
                                             0
                                                 8
                   0
                          1
                              2
                                             0
                                                  9
##
   lactcurves
                                  1
                              2 14
                          2
## MHCtools
                                             0
                                                  2
  mMARCH.AC
                              0 13
                                             0
                                                  0
                   1
                          6
                              1
                                  9
                                                  6
##
   RJafroc
                                             0
  rSHAPE
                   0
                              0
##
                          1
                                  0
                                             0
                                                  2
                          2 0 6
##
   spatstat
                                             0
                                                 12
##
            Terms
## Docs
       test time
##
  arthistory 0 0
               0
  frailtypack 0
##
                   3
   KoulMde
##
```

```
##
    lactcurves
##
    MHCtools
                    0
                         \cap
    mMARCH.AC
##
                   0
                         1
##
    RJafroc
                   0
                         0
##
    rSHAPE
    spatstat
                         1
##
```

```
k_max = 30
k.topics <- 2:k_max
doc_num = nrow(cran)

folding4 <- rep(1:4, each = round(doc_num/5,0))
folding = c(folding4, rep(5, (doc_num-length(folding4))))</pre>
```

```
## parallel computation
runonce paral <- function(sed) {</pre>
   res <- NULL
    for (k in k.topics) {
        for (fold in 1:5) {
            testing.dtm <- which(folding == fold)</pre>
            training.dtm <- which(folding != fold)</pre>
            training.model <- LDA(desc dtm[training.dtm, ], k = k, c</pre>
        ontrol = list(seed = sed*100))
            test.model <- LDA(desc dtm[testing.dtm, ], model = train</pre>
        ing.model, control = list(estimate.beta = FALSE))
            prep = perplexity(test.model)
            cat(paste(sed, k, fold, prep, "\n", sep="\t"),append=T)
            res <- rbind(res, c(sed, k, fold, prep))</pre>
    return (res)
}
library(doParallel)
ncores <- 32
#switch between %do% (serial) and %dopar% (parallel)
`%is par%` <- `%do%`
}else{ #parallel
    `%is par%` <- `%dopar%`
```

```
cl <- makeCluster(ncores) #, outfile="res.txt")
    registerDoParallel(cores = ncores)
}

foreach(sed = 1:100, .packages = c("topicmodels") ) %is_par%{
    sink("res.txt", append=TRUE) # divert the output to the log file
    res = runonce_paral(sed)
}

if(ncores > 1) stopCluster(cl)
```

Import the results of cross-validation

```
res = read.table("lda/res.txt", sep="\t")

colnames(res) = c('sed', 'topics', 'fold', 'perplexity')

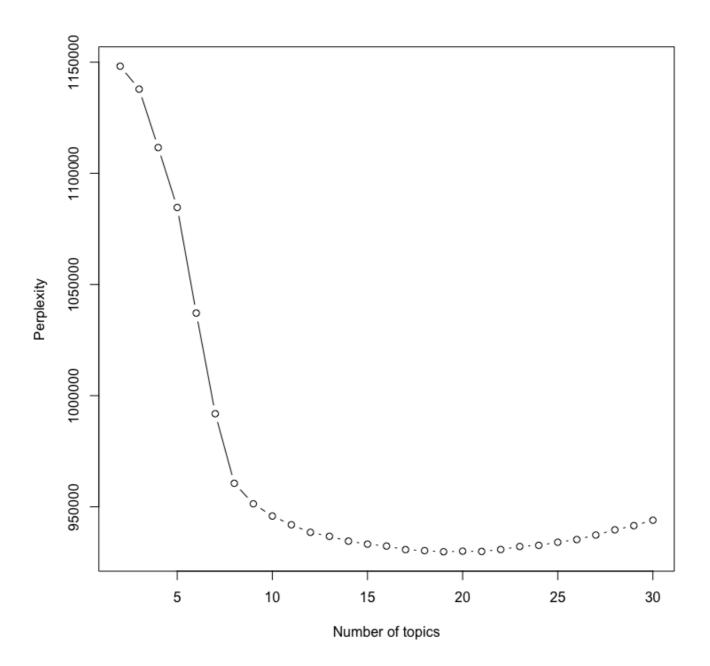
total.perp <- NULL

for(sedi in 1:100){
    for(ki in 2:k_max){
        loc = which(res[,'sed']==sedi & res[,'topics']==ki)
            total.perp = rbind(total.perp, c(sedi, ki, mean(res[loc,'per plexity'])))
    }
}

# round(total.perp)

total.perp.all = tapply(total.perp[, 3], total.perp[, 2], mean)
sort(total.perp.all)</pre>
```

```
##
         19
                   21
                             20
                                      18
                                                 17
                                                           22
   929789.7 929932.2
                      930054.1 930299.9
                                           930744.0
                                                     930788.1
##
                                                               932151
         24
                             25
                                       14
                                                 26
##
   932677.5
             933219.6
                      934061.0
                                 934571.9
                                          935270.0
                                                     936731.0
##
                                                              937306
         28
                   29
                                       30
##
                             11
                                                 10
##
   939651.4 941527.3
                      941917.2 943974.6
                                          945859.9
                                                     951360.7
                                                              960580
                              4
                                        3
##
## 1037146.3 1084631.1 1111598.1 1137845.6 1148172.4
```



```
## 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 2 ## 0 0 0 0 0 0 0 1 0 3 2 2 1 2 3 1 4 3 2 1 2 ## 28 29 30 ## 0 0 0
```

3.2 Topic 20

```
desc 1da20 \leftarrow LDA(desc dtm, k = 20, control = list(seed = 1000))
desc topics <- tidy(desc lda20, matrix = "beta")</pre>
desc topics
## # A tibble: 845,360 \times 3
    topic term beta
    <int> <chr> <dbl>
##
##
   1 1 analyze 1.05e- 3
##
         2 analyze 1.57e- 3
        3 analyze 1.42e-10
##
##
        4 analyze 8.08e-12
        5 analyze 6.57e-30
##
  6 6 analyze 2.62e- 3
##
   7
##
        7 analyze 4.81e- 3
        8 analyze 6.18e- 4
## 8
## 9
        9 analyze 2.02e- 4
## 10 10 analyze 1.87e-12
## # ... with 845,350 more rows
```

Relationships between topic and packages

```
desc_alpha <- tidy(desc_lda20, matrix = "gamma")

desc_alpha[which(desc_alpha$document == 'Rcpp'),]</pre>
```

```
## # A tibble: 20 × 3
## document topic gamma
## <chr> <int> <dbl>
```

```
## 1 Rcpp
                1 0.0752
                 2 0.00191
## 2 Rcpp
## 3 Rcpp
                3 0.00191
               4 0.00191
## 4 Rcpp
## 5 Rcpp
                5 0.00191
## 6 Rcpp
              6 0.00191
## 7 Rcpp
                7 0.00191
              8 0.00191
## 8 Rcpp
## 9 Rcpp
                9 0.00191
              10 0.00191
## 10 Rcpp
## 11 Rcpp
               11 0.00191
              12 0.122
13 0.00191
## 12 Rcpp
## 13 Rcpp
              14 0.00191
## 14 Rcpp
## 15 Rcpp
               15 0.00191
## 16 Rcpp
               16 0.771
              17 0.00191
## 17 Rcpp
## 18 Rcpp
               18 0.00191
              19 0.00191
## 19 Rcpp
               20 0.00191
## 20 Rcpp
```

Network Analysis

4 Package dependency network and author collaboration network

4.1 Build network and set edges

```
library(cranly)
library(influential)
```

```
##
## Attaching package: 'influential'
```

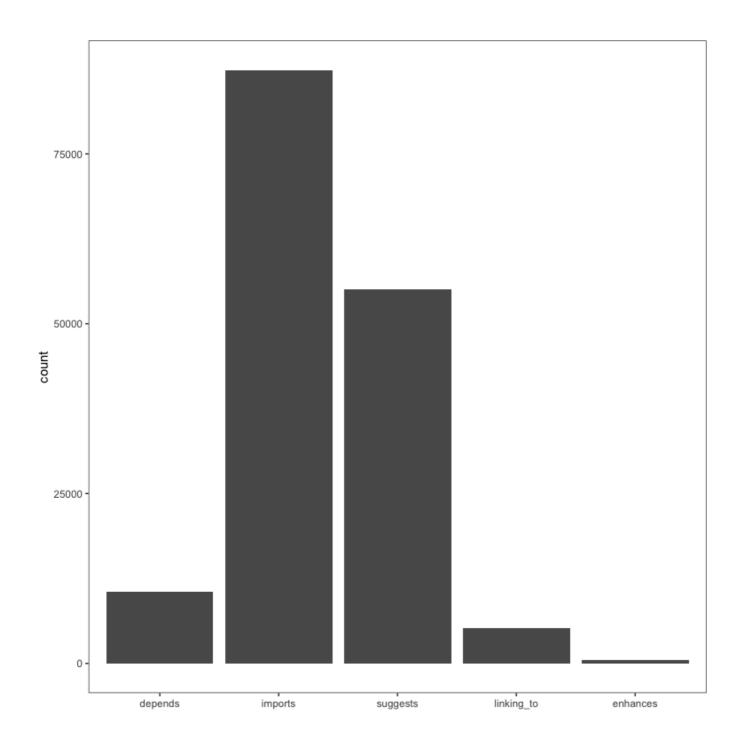
```
## The following objects are masked from 'package:igraph':
##
## betweenness, graph_from_data_frame
```

```
library(igraph)
load("pkg 221127.RData") # load cran
cran ly <- clean CRAN db(cran)</pre>
pkg net <- build network(cran ly, perspective = "package")</pre>
aut net <- build network(cran ly, perspective = "author")</pre>
pkg graph = as.igraph(pkg net,reverse=TRUE) # reverse the direction
aut graph = as.igraph(aut net)
####### flag weights
######## pkg graph: weight = relationships
pkg net type = E(pkg graph)$type
pkg net type num = pkg net type
pkg type = c("depends", "imports", "suggests", "linking to", "enhanc
        es")
pkg type num = c(5:1)
for (typei in 1:length(pkg type)){
    loc = which(pkg net type==pkg type[typei])
    pkg net type num[loc] = pkg type num[typei]
pkg graph w = pkg graph %>% set edge attr("weight", value = pkg net
        type num)
####### special cases: same anthor with different names (Gábor Csá
        rdi and Kirill Müller)
aut adj matrix = as adjacency matrix(aut graph)
loc1 = which(colnames(aut adj matrix) == "Gabor Csardi")
loc2 = which(colnames(aut adj matrix) == "Gábor Csárdi")
loc11 = which(aut adj matrix[loc1,]>0)
loc21 = which(aut adj matrix[loc2,]>0)
aut adj matrix[loc2,loc11] = aut adj matrix[loc2,loc11] + aut adj matrix
        [loc1, loc11]
aut adj matrix[loc11,loc2]=aut adj matrix[loc11,loc2]+aut adj matrix
        [loc11, loc1]
aut adj matrix = aut adj matrix[-loc1, -loc1]
loc1 = which(colnames(aut adj matrix) == "Kirill Muller")
loc2 = which(colnames(aut adj matrix) == "Kirill Müller")
loc11 = which(aut adj matrix[loc1,]>0)
```

```
loc21 = which(aut adj matrix[loc2,]>0)
aut adj matrix[loc2,loc11] = aut adj matrix[loc2,loc11] + aut adj matrix
        [loc1, loc11]
aut adj matrix[loc11,loc2]=aut adj matrix[loc11,loc2]+aut adj matrix
        [loc11, loc1]
aut adj matrix = aut adj matrix[-loc1, -loc1]
######## aut graph: weight = number of coauthored pkgs (collaborati
        on intensity)
aut graph w = graph from adjacency matrix(aut adj matrix, mode="undir
        ected", weighted=TRUE, diag=FALSE)
aut graph nw = graph from adjacency matrix(aut adj matrix, mode="undi
        rected", weighted=NULL, diag=FALSE)
####### frequency figure
pkg net type = factor(pkg_net_type, levels = pkg_type)
table(pkg_net type)
## pkg net type
## depends
                imports suggests linking to enhances
        10523
                              54991
                                          5153
                                                       558
##
                   87328
ggplot(as.data.frame(pkg net type), aes(pkg net type)) +
    geom bar() +
```

labs(x = NULL) +

theme(panel.grid=element blank())



4.2 Measures of Influence

```
autw eigen = eigen centrality(aut graph w, weights = E(aut graph w)$
        weight)
## sort
pkg rank dg = sort(pkgw dg, decreasing = TRUE)
aut rank dg = sort(autw dg, decreasing = TRUE)
pkg rank betw = sort(pkgw betw, decreasing = TRUE)
aut rank betw = sort(autw betw, decreasing = TRUE)
pkg rank eigen = sort(pkgw eigen[["vector"]], decreasing = TRUE)
aut rank eigen = sort(autw eigen[["vector"]], decreasing = TRUE)
pkg rank pgRk = sort(pkgw pgRk[["vector"]], decreasing = TRUE)
aut rank pgRk = sort(autw pgRk[["vector"]], decreasing = TRUE)
## organize
table pkg = array(NA, dim=c(50,4))
colnames(table pkg) = c("In-degree", "Betweenness",
 "Eigenvector", "PageRank")
table pkg[,"In-degree"] = names(pkg rank dg[1:50])
table pkg[,"Betweenness"] = names(pkg rank betw[1:50])
table pkg[,"PageRank"] = names(pkg rank pgRk[1:50])
table pkg[,"Eigenvector"] = names(pkg rank eigen[1:50])
write.csv(table pkg, "network/50pkg w 54321.csv")
table aut = array(NA, dim=c(50,4))
colnames(table aut) = c("In-degree", "Betweenness",
 "Eigenvector", "PageRank")
table aut[,"In-degree"] = names(aut rank dg[1:50])
table aut[,"Betweenness"] = names(aut rank betw[1:50])
table aut[,"PageRank"] = names(aut rank pgRk[1:50])
table aut[,"Eigenvector"] = names(aut rank eigen[1:50])
write.csv.utf8.BOM(table aut, "network/50aut w.csv")
```

table pkg

```
##
         In-degree
                          Betweenness
                                              Eigenvector
                                                               PageRank
##
    [1,] "knitr"
                          "ggplot2"
                                              "knitr"
                                                               "testthat"
    [2,] "testthat"
                          "knitr"
                                              "rmarkdown"
                                                               "utils"
##
    [3,] "rmarkdown"
                          "broom"
                                              "testthat"
                                                               "methods"
##
   [4,] "stats"
                                              "stats"
                                                               "stats"
##
                          "dplyr"
##
   [5,] "Rcpp"
                          "emmeans"
                                              "ggplot2"
                                                               "knitr"
##
   [6,] "ggplot2"
                          "testthat"
                                              "dplyr"
                                                               "patchSynct
                                                               "rmarkdown"
##
   [7,] "methods"
                          "shiny"
                                              "methods"
   [8,] "dplyr"
                          "stats"
##
                                              "utils"
                                                               "tools"
   [9,] "utils"
                          "rmarkdown"
                                              "Rcpp"
                                                               "covr"
##
## [10,] "graphics"
                          "survival"
                                              "rlang"
                                                               "stringr"
```

##	[11,]	"MASS"	"bayestestR"	"magrittr"	"graphics"
##	[12,]	"magrittr"	"sf"	"tibble"	"mapmisc"
##	[13,]	"covr"	"gap"	"tidyr"	"Rcpp"
##	[14,]	"rlang"	"MASS"	"graphics"	"tis"
##	[15,]	"tibble"	"targets"	"covr"	"grDevices"
##	[16,]	"tidyr"	"caret"	"stringr"	"rlang"
##	[17,]	"stringr"	"multcomp"	"purrr"	"MASS"
##	[18,]	"grDevices"	"Hmisc"	"MASS"	"ggplot2"
##	[19,]	"purrr"	"texreg"	"grDevices"	"magrittr"
##	[20,]	"parallel"	"insight"	"data.table"	"withr"
##	[21,]	"Matrix"	"parameters"	"jsonlite"	"jsonlite"
##	[22,]	"data.table"	"enrichwith"	"Matrix"	"parallel"
##	[23,]	"jsonlite"	"cops"	"shiny"	"htmltools"
##	[24,]	"RcppArmadillo"	"mlr"	"parallel"	"scales"
##	[25,]	"shiny"	"Z00"	"httr"	"fastcluste
##	[26,]	"httr"	"earth"	"glue"	"enrichwith
##	[27,]	"mvtnorm"	"tibble"	"scales"	"cops"
##	[28,]	"survival"	"AER"	"sf"	"QuasiSeq"
##	[29,]	"foreach"	"lme4"	"readr"	"tibble"
##	[30,]	"scales"	"jsonlite"	"lubridate"	"dplyr"
##	[31,]	"plyr"	"robustbase"	"igraph"	"lattice"
##	[32,]	"igraph"	"plotmo"	"reshape2"	"plyr"
##	[33,]	"reshape2"	"quantreg"	"withr"	"R6"
##	[34,]	"lubridate"	"effects"	"gridExtra"	"vctrs"
##	[35,]	"doParallel"	"rgl"	"foreach"	"reshape"
##	[36,]	"grid"	"nnet"	"cli"	"glue"
##	[37,]	"sp"	"nlme"	"tidyselect"	"digest"
##	[38,]	"gridExtra"	"sp"	"plyr"	"tinytest"
##	[39,]	"readr"	"doParallel"	"xml2"	"RUnit"
##	[40,]	"spelling"	"surveillance"	"sp"	"httr"
##	[41,]	"lattice"	"mice"	"plotly"	"Matrix"
##	[42,]	"glue"	"Matrix"	"grid"	"xml2"
##	[43,]	"RColorBrewer"	"data.tree"	"survival"	"shiny"
##	[44,]	"sf"	"car"	"lme4"	"curl"
##	[45,]	"xml2"	"spelling"	"vctrs"	"rstudioapi
##	[46,]	"raster"	"metafor"	"lifecycle"	"cli"
##	[47,]	"ZOO"	"marginaleffects"	"broom"	"yaml"
##	[48,]	"markdown"	"gtools"	"mvtnorm"	"survival"
##	[49,]	"R6"	"partykit"	"RcppArmadillo"	"markdown"
##	[50,]	"glmnet"	"hunspell"	"doParallel"	"htmlwidget

table_aut

##	In-degree	Betweenness	Eigenvector
##	[1,] "Hadley Wickham"	"R Core"	"RStudio"
0.0			

##	[2,]	"RStudio"	"Hadley Wickham"	"Hadley Wickham"
##	[3,]	"Dirk Eddelbuettel"	"Dirk Eddelbuettel"	"Jim Hester"
##	[4,]	"Ben Bolker"	"Martin Maechler"	"JJ Allaire"
##	[5 ,]	"R Core"	"RStudio"	"Winston Chang"
##	[6,]	"Martin Maechler"	"Ben Bolker"	"Yihui Xie"
##	[7,]	"Yihui Xie"	"Kurt Hornik"	"Joe Cheng"
##	[8,]	"Brian Ripley"	"Brian Ripley"	"Max Kuhn"
##	[9,]	"Michael Friendly"	"Roger Bivand"	"Gábor Csárdi"
##	[10,]	"Jim Hester"	"Achim Zeileis"	"Lionel Henry"
##	[11,]	"Roger Bivand"	"Scott Chamberlain"	"Kirill Müller"
##	[12,]	"JJ Allaire"	"Jeroen Ooms"	"Kevin Ushey"
##	[13,]	"Henrik Bengtsson"	"Yihui Xie"	"Christophe Dervi
##	[14,]	"Bill Venables"	"Zhian N Kamvar"	"Barret Schloerke
##	[15,]	"Kevin Ushey"	"Michael Friendly"	"Jennifer Bryan"
##	[16,]	"Achim Zeileis"	"John Muschelli"	"Jeroen Ooms"
##	[17,]	"Kurt Hornik"	"Rob Hyndman"	"Carson Sievert"
##	[18,]	"Max Kuhn"	"Tyler Rinker"	"Javier Luraschi"
##	[19,]	"Romain Francois"	"Bill Denney"	"Romain Francois"
##	[20,]	"Duncan Murdoch"	"John Wiseman"	"Daniel Falbel"
##	[21,]	"Zhian N Kamvar"	"Thomas Lumley"	"PBC"
##	[22,]	"David Robinson"	"Jim Hester"	"R Core"
##	[23,]	"Hao Zhu"	"Sahir Bhatnagar"	"Henrik Bengtsson
##	[24,]	"Michal Bojanowski"	"Henrik Bengtsson"	"Ben Bolker"
##	[25,]	"Joe Cheng"	"Toby Hocking"	"David Robinson"
##	[26,]	"Vilmantas Gegzna"	"Carl Boettiger"	"Michal Bojanowsk
##	[27,]	"Christophe Dervieux"	"Kevin Ushey"	"Thomas Lin Peder
##	[28,]	"Bill Denney"	"Kirill Müller"	"Davis Vaughan"
##	[29,]	"Adrian Baddeley"	"Bob Rudis"	"JooYoung Seo"
##	[30,]	"Jeroen Ooms"	"Cleve Moler"	"Atsushi Yasumoto
##	[31,]	"Hong Ooi"	"JJ Allaire"	"Yixuan Qiu"
##	[32,]	"Noam Ross"	"Noam Ross"	"Joseph Larmarang
##	[33,]	"David Hugh-Jones"	"Bill Venables"	"Dirk Eddelbuette
##	[34,]	"Joseph Larmarange"	"Max Kuhn"	"Malcolm Barrett"
##	[35,]	"John Muschelli"	"Ben Goodrich"	"Noam Ross"
##	[36,]	"Jonah Gabry"	"Jonah Gabry"	"Jeff Allen"
11 11				W1 W
		"Malcolm Barrett"	"Christophe Dutang"	"Hao Zhu"
		"Greg Snow"	"Torsten Hothorn"	"Michael Friendly
##		"Jim Lemon"	"Winston Chang"	"Jenny Bryan"
##		"Alessandro Gasparini"		"Google Inc"
##		"Eduard Szoecs"	"Apache Foundation"	"David Hugh-Jones
##		"Tal Galili"	"Arni Magnusson"	"jQuery Foundatio
##		"Jenny Bryan"	"Romain Francois"	"Roger Bivand"
##		"Torsten Hothorn"	"Joe Cheng"	"Hiroaki Yutani"
##		"Matthieu Stigler"	"Duncan Murdoch"	"Martin Maechler"
##		"Garrick Aden-Buie"	"Steven G Johnson"	"Alex Hayes"
##		"Dieter Menne"	"Yuan Tang"	"Mango Solutions"
##	[48,]	"Frank E Harrell Jr"	"Jacob Bien"	"Richard Iannone"

```
## [49,] "Rolf Turner" "Lampros Mouselimis" "Simon Couch"
 ## [50,] "Michael Chirico" "Yi Yang"
                                                      "Frederik Aust"
 ##
         PageRank
 ##
    [1,] "RStudio"
     [2,] "Hadley Wickham"
 ##
    [3,] "R Core"
 ##
 ##
     [4,] "Martin Maechler"
     [5,] "Ben Bolker"
 ##
 ##
     [6,] "Dirk Eddelbuettel"
     [7,] "Yihui Xie"
 ##
 ##
    [8,] "Achim Zeileis"
 ##
    [9,] "JJ Allaire"
 ## [10,] "Kurt Hornik"
 ## [11,] "Brian Ripley"
 ## [12,] "Roger Bivand"
 ## [13,] "Jim Hester"
 ## [14,] "Jeroen Ooms"
 ## [15,] "Scott Chamberlain"
 ## [16,] "Michael Friendly"
 ## [17,] "Zhian N Kamvar"
 ## [18,] "Winston Chang"
 ## [19,] "Joe Cheng"
 ## [20,] "Henrik Bengtsson"
 ## [21,] "Bob Rudis"
 ## [22,] "Kevin Ushey"
 ## [23,] "Duncan Murdoch"
 ## [24,] "Max Kuhn"
 ## [25,] "Bill Venables"
 ## [26,] "John Muschelli"
 ## [27,] "Gábor Csárdi"
 ## [28,] "Rob Hyndman"
 ## [29,] "Torsten Hothorn"
 ## [30,] "Kirill Müller"
 ## [31,] "Noam Ross"
 ## [32,] "Romain Francois"
 ## [33,] "Christophe Dervieux"
 ## [34,] "Carl Boettiger"
 ## [35,] "David Robinson"
 ## [36,] "Jonah Gabry"
 ## [37,] "Maëlle Salmon"
 ## [38,] "Michael Sumner"
 ## [39,] "Barret Schloerke"
 ## [40,] "Stéphane Laurent"
 ## [41,] "Bill Denney"
 ## [42,] "Edzer Pebesma"
 ## [43,] "Hao Zhu"
 ## [44,] "Indrajeet Patil"
```

```
## [45,] "Michel Lang"
## [46,] "Michal Bojanowski"
## [47,] "John Fox"
## [48,] "Google Inc"
## [49,] "Arni Magnusson"
## [50,] "Uwe Ligges"
```

4.3 Sensitivity Analysis

weight: 321

no weight

```
pkgw_1_betw = betweenness(pkg_graph)
pkgw_1_dg = degree(pkg_graph, mode = c("in"))
pkgw_1_pgRk = page_rank(pkg_graph)
pkgw_1_eigen = eigen_centrality(pkg_graph)
```

```
## 1 1
## 1 1
## 0.9662078 0.9805951
## 0.9926748 0.9941666
```

4.4 Select Important Packages and Authors

Packages

```
imp pkg = NULL
for(i in 1:nrow(table pkg)){
  for(j in 1:ncol(table pkg)) {
    pkg temp = as.character(table pkg[i,j])
    loc = which(table pkg==pkg temp)
    if (length(loc)>1) {
      if (length(which(imp pkg==pkg temp))==0) {
        imp pkg = c(imp pkg, pkg temp)
      }
   }
}
imp pkg2 = array(NA, dim=c(length(imp pkg), 3))
imp pkg2[,1] = imp pkg
colnames(imp pkg2) = c('pkg','title','mt')
for(i in 1:length(imp pkg)){
  imp pkg2[i,'pkg'] = imp pkg[i]
  loc = which(cran[,'Package'] == imp pkg[i])
  temp = try(cran[loc,'Title'])
  if (length (temp) > 0) {
    imp pkg2[i,'title'] = temp
  temp = try(cran[loc, 'Maintainer'])
```

```
if(length(temp)>0) {
    imp_pkg2[i,'mt'] = split_mt(temp) $name
  }
}
write.csv.utf8.BOM(imp_pkg2,'network/imp_pkg2_54321.csv')
imp_pkg2
```

```
##
   pkg
   [1,] "knitr"
##
   [2,] "ggplot2"
##
   [3,] "testthat"
##
##
   [4,] "rmarkdown"
   [5,] "utils"
##
  [6,] "broom"
##
##
  [7,] "methods"
## [8,] "stats"
## [9,] "dplyr"
## [10,] "Rcpp"
## [11,] "shiny"
## [12,] "covr"
## [13,] "graphics"
## [14,] "survival"
## [15,] "rlang"
## [16,] "stringr"
## [17,] "MASS"
## [18,] "magrittr"
## [19,] "sf"
## [20,] "tibble"
## [21,] "tidyr"
## [22,] "grDevices"
## [23,] "purrr"
## [24,] "parallel"
## [25,] "data.table"
## [26,] "withr"
## [27,] "Matrix"
## [28,] "jsonlite"
## [29,] "enrichwith"
## [30,] "cops"
## [31,] "RcppArmadillo"
## [32,] "scales"
## [33,] "zoo"
## [34,] "httr"
## [35,] "glue"
## [36,] "mvtnorm"
## [37,] "foreach"
```

```
## [38,] "lme4"
 ## [39,] "readr"
 ## [40,] "lubridate"
 ## [41,] "plyr"
 ## [42,] "igraph"
 ## [43,] "lattice"
 ## [44,] "reshape2"
 ## [45,] "R6"
 ## [46,] "gridExtra"
 ## [47,] "vctrs"
 ## [48,] "doParallel"
 ## [49,] "grid"
 ## [50,] "cli"
 ## [51,] "sp"
 ## [52,] "xml2"
 ## [53,] "spelling"
 ## [54,] "markdown"
 ##
          title
 ##
    [1,] "A General-Purpose Package for Dynamic Report Generation in R
     [2,] "Create Elegant Data Visualisations Using the Grammar of Grap
     [3,] "Unit Testing for R"
 ##
     [4,] "Dynamic Documents for R"
 ##
     [5,] NA
 ##
     [6,] "Convert Statistical Objects into Tidy Tibbles"
 ##
 ##
    [7,] NA
 ##
     [8,] NA
 ##
    [9,] "A Grammar of Data Manipulation"
 ## [10,] "Seamless R and C++ Integration"
 ## [11,] "Web Application Framework for R"
 ## [12,] "Test Coverage for Packages"
 ## [13,] NA
 ## [14,] "Survival Analysis"
 ## [15,] "Functions for Base Types and Core R and 'Tidyverse' Features
 ## [16,] "Simple, Consistent Wrappers for Common String Operations"
 ## [17,] "Support Functions and Datasets for Venables and Ripley's MAS
 ## [18,] "A Forward-Pipe Operator for R"
 ## [19,] "Simple Features for R"
 ## [20,] "Simple Data Frames"
 ## [21,] "Tidy Messy Data"
 ## [22,] NA
 ## [23,] "Functional Programming Tools"
 ## [24,] NA
 ## [25,] "Extension of `data.frame`"
 ## [26,] "Run Code 'With' Temporarily Modified Global State"
 ## [27,] "Sparse and Dense Matrix Classes and Methods"
 ## [28,] "A Simple and Robust JSON Parser and Generator for R"
 ## [29,] "Methods to Enrich R Objects with Extra Components"
```

```
## [30,] "Cluster Optimized Proximity Scaling"
 ## [31,] "'Rcpp' Integration for the 'Armadillo' Templated Linear Alge
 ## [32,] "Scale Functions for Visualization"
 ## [33,] "S3 Infrastructure for Regular and Irregular Time Series (Z's
 ## [34,] "Tools for Working with URLs and HTTP"
 ## [35,] "Interpreted String Literals"
 ## [36,] "Multivariate Normal and t Distributions"
 ## [37,] "Provides Foreach Looping Construct"
 ## [38,] "Linear Mixed-Effects Models using 'Eigen' and S4"
 ## [39,] "Read Rectangular Text Data"
 ## [40,] "Make Dealing with Dates a Little Easier"
 ## [41,] "Tools for Splitting, Applying and Combining Data"
 ## [42,] "Network Analysis and Visualization"
 ## [43,] "Trellis Graphics for R"
 ## [44,] "Flexibly Reshape Data: A Reboot of the Reshape Package"
 ## [45,] "Encapsulated Classes with Reference Semantics"
 ## [46,] "Miscellaneous Functions for \"Grid\" Graphics"
 ## [47,] "Vector Helpers"
 ## [48,] "Foreach Parallel Adaptor for the 'parallel' Package"
 ## [49,] NA
 ## [50,] "Helpers for Developing Command Line Interfaces"
 ## [51,] "Classes and Methods for Spatial Data"
 ## [52,] "Parse XML"
 ## [53,] "Tools for Spell Checking in R"
 ## [54,] "Render Markdown with 'commonmark'"
 ##
         mt
    [1,] "Yihui Xie "
 ##
 ##
    [2,] "Thomas Lin Pedersen "
 ##
    [3,] "Hadley Wickham"
    [4,] "Yihui Xie "
 ##
 ##
    [5,] NA
    [6,] "Simon Couch "
 ##
 ##
    [7,] NA
    [8,] NA
 ##
 ##
    [9,] "Hadley Wickham "
 ## [10,] "Dirk Eddelbuettel "
 ## [11,] "Winston Chang "
 ## [12,] "Jim Hester "
 ## [13,] NA
 ## [14,] "Terry M Therneau "
 ## [15,] "Lionel Henry "
 ## [16,] "Hadley Wickham"
 ## [17,] "Brian Ripley "
 ## [18,] "Lionel Henry "
 ## [19,] "Edzer Pebesma "
 ## [20,] "Kirill Müller "
 ## [21,] "Hadley Wickham"
```

```
## [22,] NA
 ## [23,] "Lionel Henry "
 ## [24,] NA
 ## [25,] "Matt Dowle"
 ## [26,] "Lionel Henry "
 ## [27,] "Martin Maechler"
 ## [28,] "Jeroen Ooms "
 ## [29,] "Ioannis Kosmidis "
 ## [30,] "Thomas Rusch "
 ## [31,] "Dirk Eddelbuettel "
 ## [32,] "Hadley Wickham "
 ## [33,] "Achim Zeileis "
 ## [34,] "Hadley Wickham"
 ## [35,] "Jennifer Bryan "
 ## [36,] "Torsten Hothorn"
 ## [37,] "Folashade Daniel "
 ## [38,] "Ben Bolker"
 ## [39,] "Jennifer Bryan "
 ## [40,] "Vitalie Spinu "
 ## [41,] "Hadley Wickham "
 ## [42,] "Tamás Nepusz "
 ## [43,] "Deepayan Sarkar"
 ## [44,] "Hadley Wickham "
 ## [45,] "Winston Chang"
 ## [46,] "Baptiste Auguie "
 ## [47,] "Lionel Henry "
 ## [48,] "Folashade Daniel "
 ## [49,] NA
 ## [50,] "Gábor Csárdi "
 ## [51,] "Edzer Pebesma "
 ## [52,] "Hadley Wickham "
 ## [53,] "Jeroen Ooms "
 ## [54,] "Yihui Xie "
```

Authors

```
## important authors (identified by at least two indexes)
imp_aut = NULL
for(i in 1:nrow(table_aut)) {
   for(j in 1:ncol(table_aut)) {
     aut_temp = as.character(table_aut[i,j])
     loc = which(table_aut==aut_temp)
     if(length(loc)>1) {
        if(length(which(imp_aut==aut_temp))==0) {
            imp_aut = c(imp_aut, aut_temp)
        }
}
```

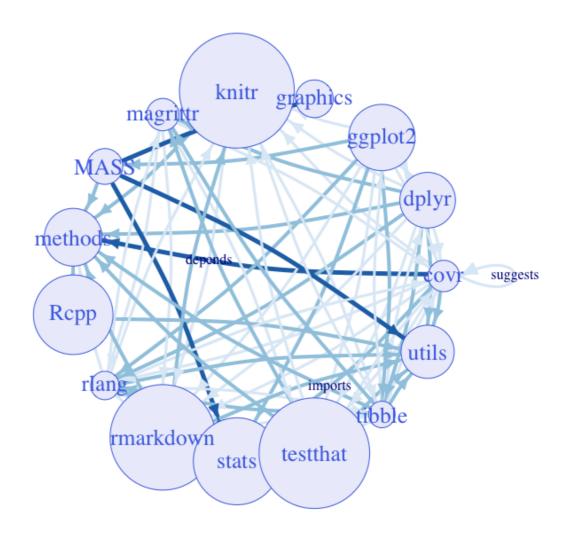
```
}
}
imp_aut
```

```
"R Core"
                                                     "RStudio"
   [1] "Hadley Wickham"
   [4] "Dirk Eddelbuettel"
##
                               "Jim Hester"
                                                     "Ben Bolker"
## [7] "Martin Maechler"
                               "JJ Allaire"
                                                     "Winston Chang"
## [10] "Yihui Xie"
                               "Kurt Hornik"
                                                     "Joe Cheng"
                                                     "Achim Zeileis"
## [13] "Brian Ripley"
                               "Max Kuhn"
## [16] "Michael Friendly"
                               "Roger Bivand"
                                                     "Gábor Csárdi"
## [19] "Scott Chamberlain"
                               "Kirill Müller"
                                                     "Jeroen Ooms"
## [22] "Kevin Ushey"
                               "Henrik Bengtsson"
                                                     "Christophe Dervie
## [25] "Bill Venables"
                               "Zhian N Kamvar"
                                                     "Barret Schloerke"
## [28] "John Muschelli"
                               "Rob Hyndman"
                                                     "Romain Francois"
## [31] "Bill Denney"
                               "Duncan Murdoch"
                                                     "Bob Rudis"
## [34] "David Robinson"
                               "Hao Zhu"
                                                     "Michal Bojanowski
## [37] "Carl Boettiger"
                               "Torsten Hothorn"
                                                     "Noam Ross"
## [40] "Joseph Larmarange"
                               "David Hugh-Jones"
                                                     "Malcolm Barrett"
## [43] "Jonah Gabry"
                               "Jenny Bryan"
                                                     "Google Inc"
## [46] "Arni Magnusson"
```

4.5 Visualization

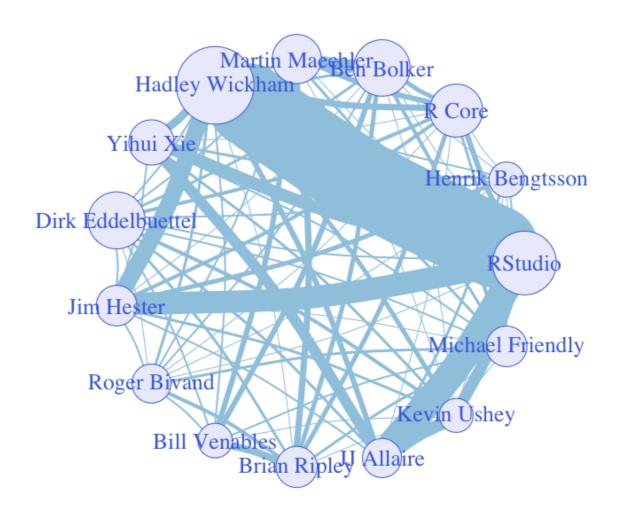
package sub-graph

```
loc = which(pkg net type==pkg type[typei])
    pkg net type color[loc] = pkg type color[typei]
pkg net type label = rep(NA, length(pkg net type))
for(typei in 1:length(pkg type)){ # select five labels as example
    loc = which(pkg net type==pkg type[typei])[1]
    pkg net type label[loc] = pkg type[typei]
}
pkg sub graph = pkg sub graph %>%
    set edge attr("color", value = pkg net type color) %>%
    set edge attr("label", value = pkg net type label)
plot.igraph(pkg sub graph,
    edge.color = E(pkg sub graph)$color,
    edge.width = as.numeric(E(pkg sub graph)$weight),
    edge.label = E(pkg sub graph)$label,
    edge.arrow.size = 1,
    edge.curved = .1,
    vertex.color = colors[5],
    vertex.frame.color=colors[1],
    vertex.label.color=colors[1],
    layout=layout.circle,
    vertex.size = (V(pkg sub graph) $indegree) / 120,
    vertex.label.cex = 1.6)
```



author sub-graph

```
edge.arrow.size = 1,
edge.curved = .1,
vertex.color = colors[5],
vertex.frame.color=colors[1],
vertex.label.color=colors[1],
layout=layout.circle,
vertex.size = (V(aut_sub_graph)$degree)/20,
vertex.label.cex = 1.6)
```



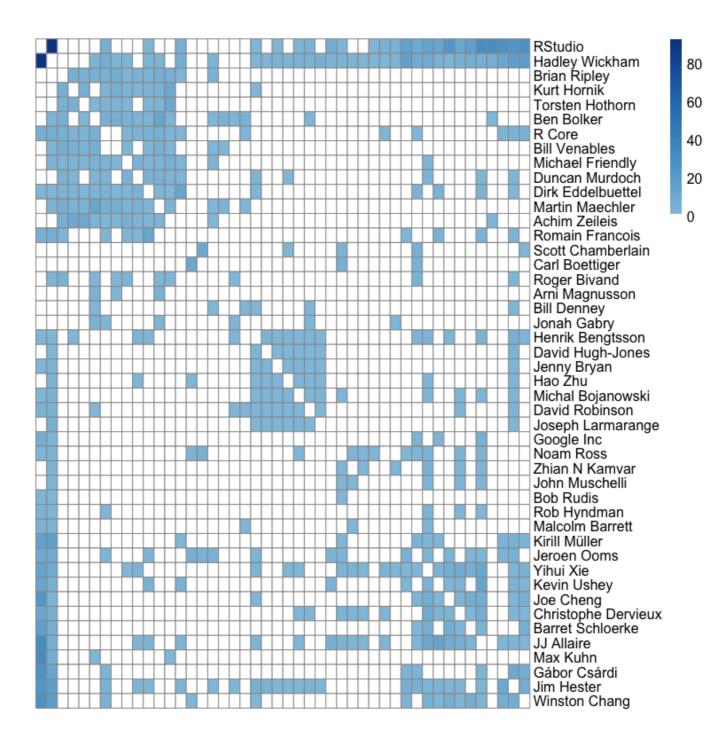
heatmap of author collaboration

```
library(corrplot)
```

```
library(pheatmap)
loc=NULL

for(i in 1:length(imp_aut)) {
   loc = c(loc, which(colnames(aut_adj_matrix)==imp_aut[i]))
}
aut_corsub_matrix = as.matrix(aut_adj_matrix[loc,loc])

pheatmap(aut_corsub_matrix,col=c("white",colorRampPalette(brewer.pal (8, "Blues"))(170)[80:170]),
   angle_col="45",
   show_colnames = F,
   fontsize=12,
   treeheight_row = 0, treeheight_col = 0)
```



4.6 Correlation among importance indexes and downloads

```
library(cranlogs)
pkg_down_year = array(NA, dim = c(nrow(cran),2))
colnames(pkg_down_year) = c("pkg", "downloads")
pkg_down_year[,"pkg"] = cran[,"Package"]

for(pkgi in 1:nrow(pkg_down_year)){ #6402
    cat(paste0(pkgi,"-",pkg_down_year[pkgi, "pkg"],"\n"))
```

```
pkg down year = read.csv('pkg down year.csv')
pkg down year rank = as.numeric(pkg down year[,2])
names(pkg down year rank) = pkg down year[,1]
pkg all index = cbind(pkgw dg, pkgw betw, pkgw eigen[["vector"]], pk
        gw_pgRk[["vector"]])
colnames(pkg all index) = c('dg','bet','eigen','pg')
pkg down rank year n = pkgw dg
no down = NULL
for(i in 1:length(pkg down rank year n)){
  loc = which(names(pkg down year rank) == names(pkgw dg)[i])
  if (length (loc) !=0) {
   pkg down rank year n[i] = pkg down year rank[loc]
    no down = c(no down, names(pkgw dg)[i])
}
for (i in 1:4) {
    cat(cor(pkg down rank year n, pkg all index[,i]))
    cat('\n')
```

```
## 0.4217525
## 0.4065768
## 0.4335971
## 0.3489176
```

5 Bipartite Network

5.1 Build bipartite network and set edges (weights)

```
basepkg = c("base", "boot", "class", "cluster", "codetools", "compil
        er", "datasets",
    "foreign", "graphics", "grDevices", "grid", "KernSmooth", "latti
        ce", "MASS",
    "Matrix", "methods", "mgcv", "nlme", "nnet", "parallel", "rpart"
        , "spatial",
    "splines", "stats", "stats4", "survival", "tcltk", "tools", "uti
        ls")
bi matrix w = array(0, dim = c(length(V(aut graph) name), length(V(p)
        kg graph w) $name)))
colnames(bi matrix w) = V(pkg graph w)$name
rownames(bi matrix w) = V(aut graph)$name
for (auti in 1:nrow(bi matrix w)){
    aut pkg = V(aut graph) $package[[auti]]
    aut pkg num = length(aut pkg)
    for (aut pkgi in 1:aut pkg num) {
        pkgi = which(colnames(bi matrix w) == aut pkg[aut pkgi])
        #if(aut pkg[aut pkgi] == colnames(bi matrix w)[pkgi]){
        bi matrix w[auti, pkgi] = 1
        if(V(aut graph) $name[auti] == V(pkg graph w) $maintainer[pkg
           bi matrix w[auti, pkgi] = 3
   }
}
# base r packages
locRcore = which(rownames(bi matrix w) == "R Core")
for (pkgi in 1:length(basepkg)) {
    locpkg = which(colnames(bi matrix w) == basepkg[pkgi])
    if(is.na(V(pkg graph w) $maintainer[locpkg])) {
            bi matrix w[locRcore, locpkg] = 3
bi matrix w0 = bi matrix w
loc1 = which(rownames(bi matrix w) == "Kirill Muller")
loc2 = which(rownames(bi matrix w) == "Kirill Müller")
loc11 = which(bi matrix w[loc1,]>0)
loc21 = which(bi matrix w[loc2,]>0)
bi matrix w[loc2,loc11]=bi matrix w[loc2,loc11]+bi matrix w[loc1,loc
bi matrix w = bi matrix w[-loc1, ]
loc1 = which(rownames(bi matrix w) == "Gabor Csardi")
```

```
loc2 = which(rownames(bi_matrix_w) == "Gábor Csárdi")
loc11 = which(bi_matrix_w[loc1,]>0)
loc21 = which(bi_matrix_w[loc2,]>0)
bi_matrix_w[loc2,loc11] = bi_matrix_w[loc2,loc11] + bi_matrix_w[loc1,loc_1]
bi_matrix_w = bi_matrix_w[-loc1,]
bi_graph_w = graph.incidence(bi_matrix_w, weighted = T) #graph_from_incidence_matrix
```

5.2 Visulization

```
loc sub1 = loc sub2 = NULL
for (i in 1:15) {
    loc sub1 = c(loc sub1, which(V(aut graph)$name==names(aut rank d
        q)[i]))
    loc sub2 = c(loc sub2, which(V(pkg graph w) $name==names(pkg rank
        dg)[i]))
bi matrix sub = bi matrix w[loc sub1, loc sub2]
bi graph w sub = graph.incidence(bi matrix sub, weighted = T) #subg
        raph(bi graph w,c(names(pkg rank dg)[1:15], names(aut rank d
        g) [1:15]))
# all influential pkg and author
loc sub1 = loc sub2 = NULL
for (i in 1:nrow(imp aut2)){
    loc sub1 = c(loc sub1, which(V(aut graph)$name==imp aut2[i,1]))
for (i in 1:nrow(imp pkq2)) {
    loc sub2 = c(loc sub2, which(V(pkg graph w)$name==imp pkg2[i,1
        ]))
bi matrix sub = bi matrix w[loc sub1, loc sub2]
bi graph w sub = graph.incidence(bi matrix sub, weighted = T)
V(bi graph w sub)$color <- V(bi_graph_w_sub)$type</pre>
V(bi graph w sub) $color=gsub("FALSE", "#FAEBD7", V(bi graph w sub) $col
V(bi graph w sub) $color=gsub("TRUE", colors[5], V(bi graph w sub) $colo
        r)
```

5.3 Centrality

```
library(bipartite)
library(birankr)
```

```
#bi_matrix_w
bi_adj = as_adjacency_matrix(bi_graph_w) #as_edgelist
bi_edge = as_edgelist(bi_graph_w)
bi_e_weight = E(bi_graph_w)$weight

bi_edge_dt_w = data.table(cbind(bi_edge,bi_e_weight))

bi_cohits_w_0 = br_cohits(bi_edge_dt_w, weight_name = "bi_e_weight")
bi_cohits_w = bi_cohits_w_0[,'rank']
names(bi_cohits_w) = bi_cohits_w_0[,1]
head(sort(bi_cohits_w,T))
```

```
## Dirk Eddelbuettel Hadley Wickham RStudio Stéphane Lau
## 0.001799382 0.001761922 0.001732088 0.00157
## Gábor Csárdi Scott Chamberlain
## 0.001487436 0.001469630
```

```
bi_birank_w_0 = br_birank(bi_edge_dt_w, weight_name = "bi_e_weight")
bi_birank_w = bi_birank_w_0[,'rank']
names(bi_birank_w) = bi_birank_w_0[,1]
head(sort(bi_birank_w,T))
```

```
## Dirk Eddelbuettel Hadley Wickham RStudio Stéphane Lau 0.0002311202 0.0002293970 0.0002287313 0.000214  
## Scott Chamberlain Gábor Csárdi  
## 0.0002120412 0.0002093672
```

Relationship between cohits and birank

```
cor(bi_cohits_w, bi_birank_w)

## [1] 0.8596433

bi_cohits_w_50 = sort(bi_cohits_w,T)[1:50]
bi_birank_w_50 = sort(bi_birank_w,T)[1:50]
```

Relationships between birank and other indexes

```
## dgree 0.178544

## betw 0.3635495

## eigen 0.2471052

## pgRk 0.4269204
```

Visualization

```
pkg_download = sort(pkg_down_year_rank, decreasing = TRUE)
downloads_sorted = cbind(names(pkg_download), as.numeric(pkg_download))
```

