Intro to RQDA

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Steps in Analysis of Qualitative Data

- 1. Preparation
- 2. Coding
- 3. Data Attributes
- 4. Codes abstraction
- 5. Code plotting and sharing
- 6. Theory building

1. Preparation

RQDA – based analysis

- #1. Install R, RSQLite, GTK, RQDA
- #2. Transform textual data into individual ASCII or .txt files
- #3. Launch RQDA
- #4. Create a new project
- #5. Upload all .txt files into RQDA

Advantages and Disadvantages

- All data files are transportable online via emails,
- Dropbox, or using a device (e.g. USB stick)
- Ease of logistics handling when moving offices,
- involving multiple researchers
- Not all researchers are familiar with R

Let's start with: What is R

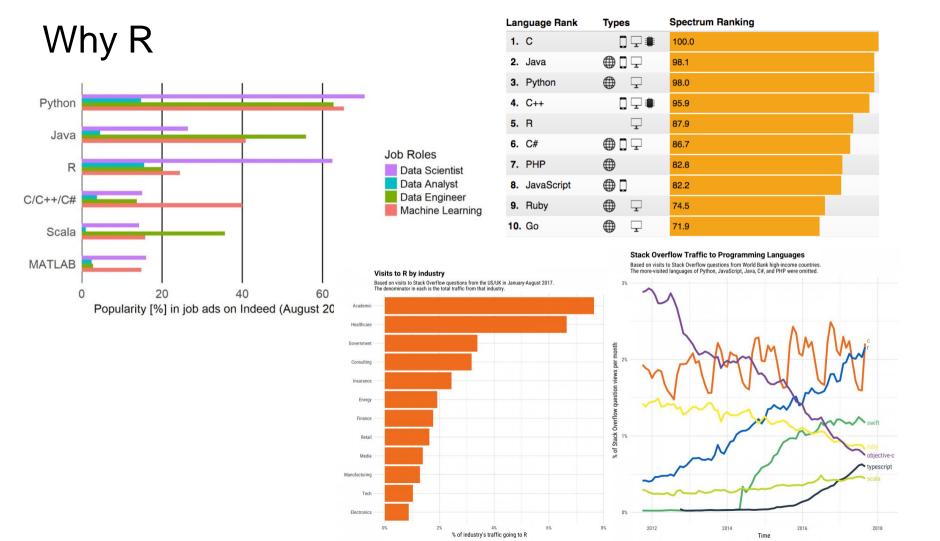


What is R Studio



- 1) A company
- 2) A software program that makes working in R easier

IDE: Integrated Development Environment



Why R

Full programming language: customizable, extensible

Open source: contributed packages

Designed for data analysis

Why R

Support for reproducible analysis

Widely used

R Community

R Community



RStudio Community https://community.rstudio.com/

R Bloggers https://www.r-bloggers.com/

R User Group Nepal https://www.r-nepal.org/



Things to Know About R





You can't "learn R"



You learn the basics

and learn how to learn more

It's expected that you'll install and use additional packages

There are multiple ways to do most things Some ways are better than others (efficient)

You can, and often will, have more than one dataset open in R at the same time

Run R in multiple ways:

Interactively: R console, RStudio, R Notebooks, Jupyter Notebooks

Batch jobs

Recently it got on cloud as well

R packages



Installing Packages

RCRAN network

install.packages("tidyverse") #install the packages library(tidyverse) #load the packages

Bioconductor

if (!requireNamespace("BiocManager"))
 install.packages("BiocManager")

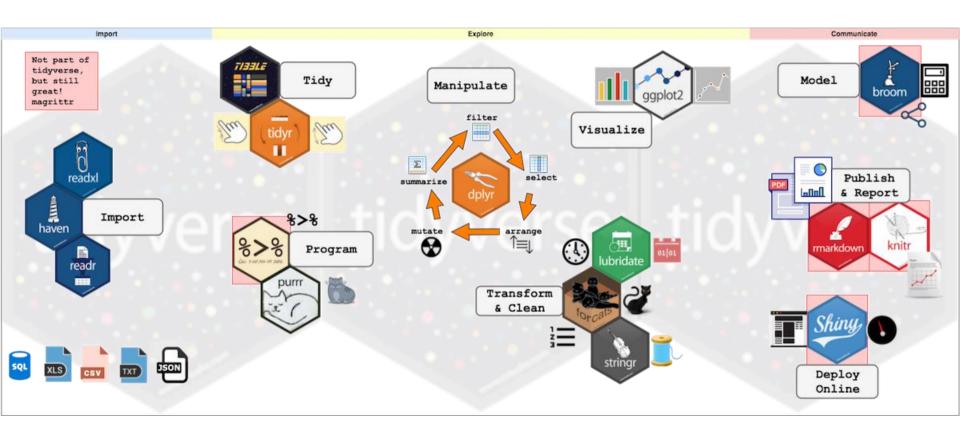
BiocManager::install()

BiocManager::install(c("GenomicFeatures", "AnnotationDbi"))

Github

library(devtools)

devtools::install_github("strengejacke/sjPlot")



Install and Download

https://github.com/prabinrs/intro_RQDA

2. Coding

- One or more coder performs inductive data analysis together, by highlighting the text and clicking "Mark" ("Unmark") to do (undo) the coding
- Other analyst(s) critiques the coding results; reconcile differences in interpretation

- Increased reliability and accuracy of data interpretation
- Ease of capturing new, unexpected insights through re-coding of data
- Possible mechanical errors for inexperienced users
- Possible loss of reflexivity or decontextualizing data

Code and Categories

Inductive coding or Open Coding (Strauss and Corbin, 1990; Gioia et al. 2013)

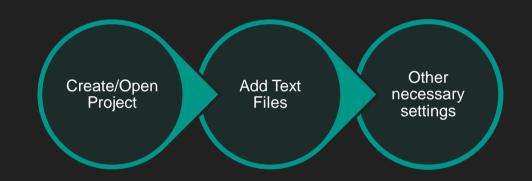
 Code a textual unit that is close to (i.e basic or lowest level) the data without being predicated on any theory.

Eg "they used my debit card to charge amounts I never authorized": code "Monetary Loss"

Deductive Coding: (Fereday and Muir-Cochrane, 2006; Bazeley and Jackson, 2013)

- Categories are coded based on established concept, variables or theories

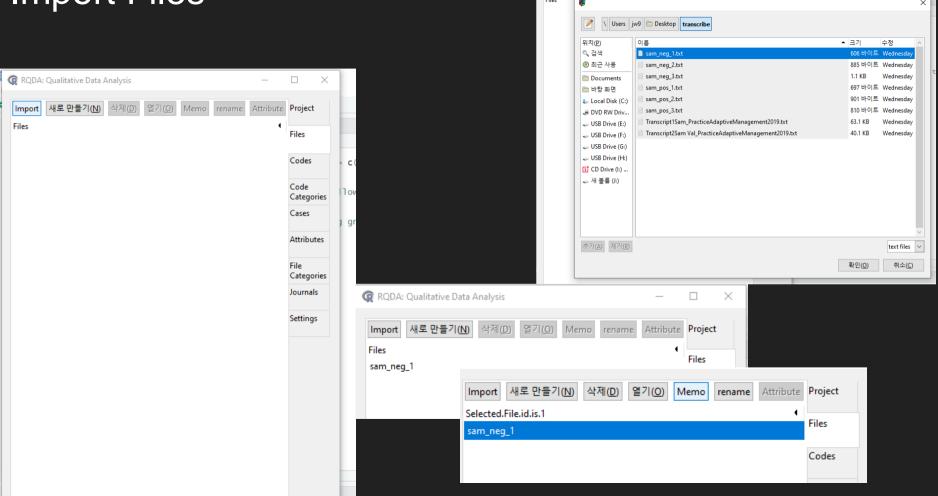
Eg "the customer service department doesn't work well with the technical support and that's why they took six months to solve my technical problems; Code "Market Orientatin" (based on Jaworki and Kohli, 1993)



New Project



Import Files



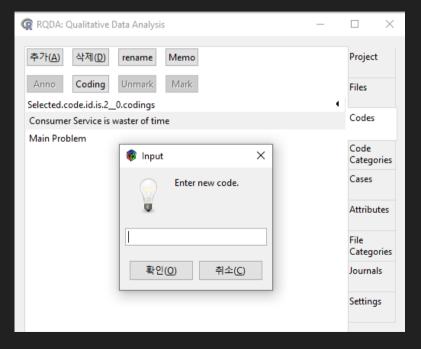
RQDA: Qualitative Data Analysis

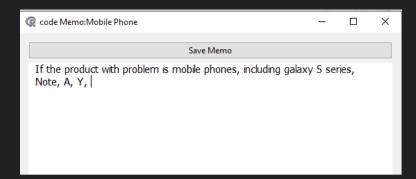
새로 만들기(N)

- 🗆 ×

삭제(D) 열기(O) Memo rename Attribute Project

Codes





3. Data Attributes

- Click "Attributes" and record key data attributes (e.g. data sources, informants' demographics, dependent variables)
- Click "Memo" to write memos for each case

- Allow easy capture of meta information of the data and attributes of informants
- Memos can be copied-and-pasted to a Word processor to aid data analysis

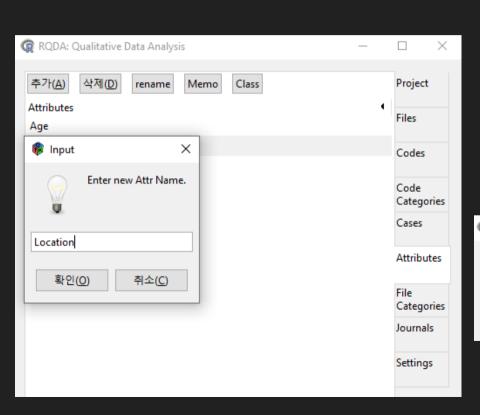
- that coders add data attributes or memos to enrich understanding
- enables users to add variables to individual files to show a subset of files and to perform statistical analysis on attributes.

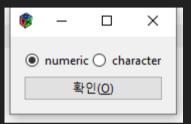
Eg:

Attributes: Type of File,

Memo: Interview with p07 was conducted in presence of her colleagues and seniors.

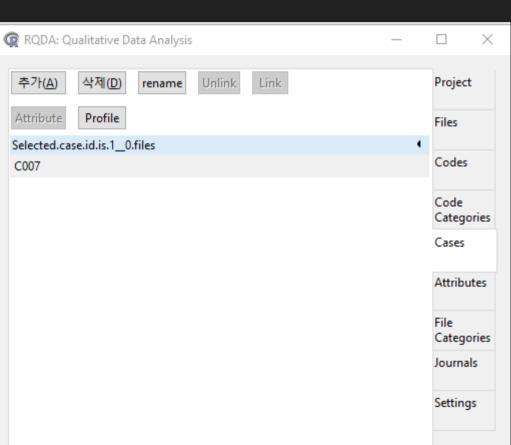
Attributes

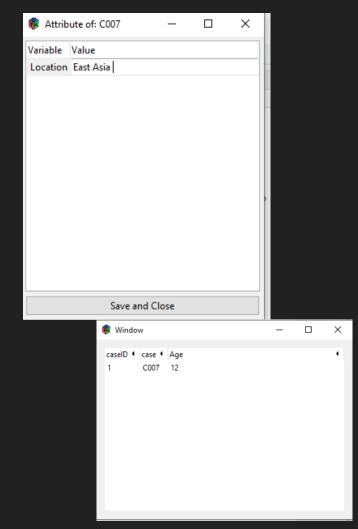




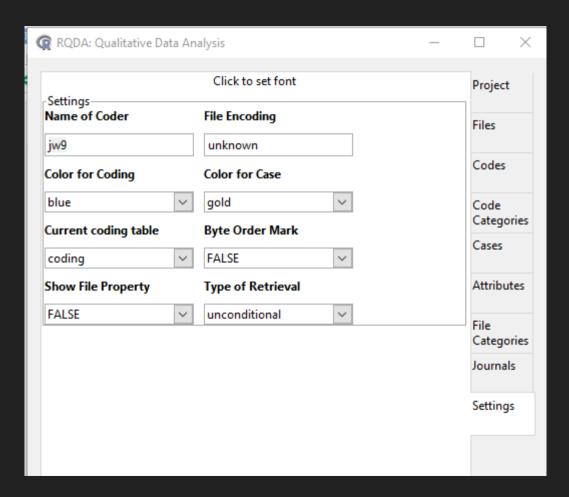


Cases

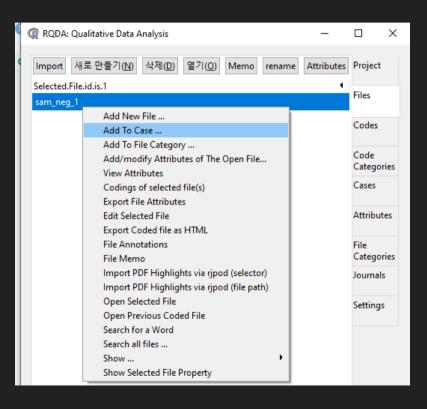


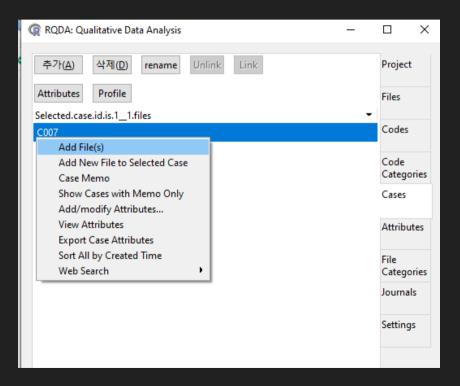


Settings

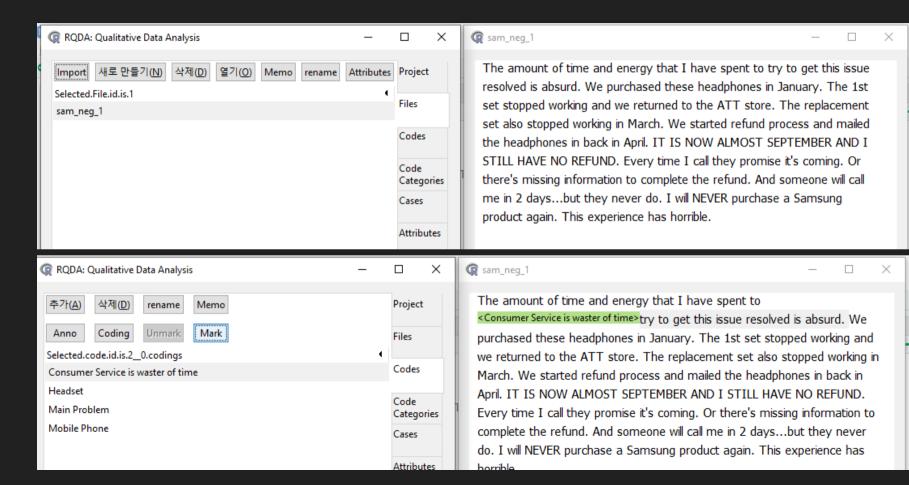


Add File to case





Coding



4. Codes Abstraction

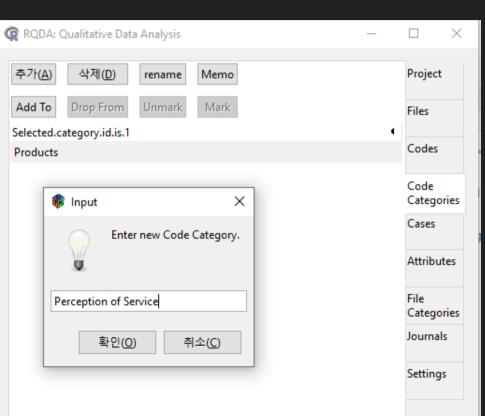
 Use "Add To" button to assign all first order codes to second-order code categories
 To create third-order code categories, copy and paste all codes and re-arrange them using color font with Excel
 Based on Constructivist theory-building Process

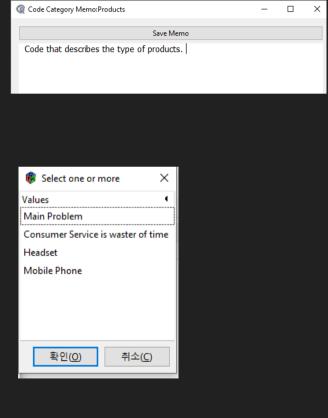
- A highly systematic way of aggregating codes to a higher level of meaning
- Challenging to aggregate too many firstorder codes (thousands of codes) to second-order codes

Abstraction to Second Level categories i.e after coding completion, first level codes can be aggreagated into higher-level **code categories**

Eg. "they used my debit card to charge amounts that I never authorize" and "this faulty laptop means I couldn't work for four days" can be aggregated to a higher-level concept called "monetary loss"

Code Categories





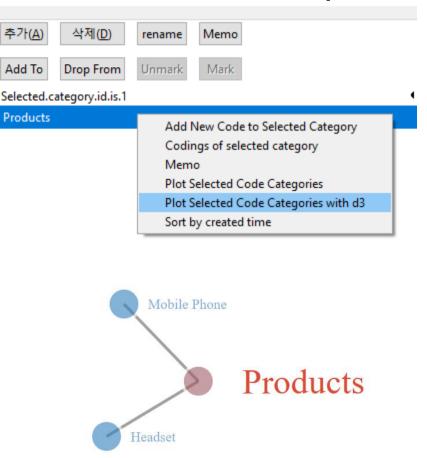
5. Codes Plotting and Sharing

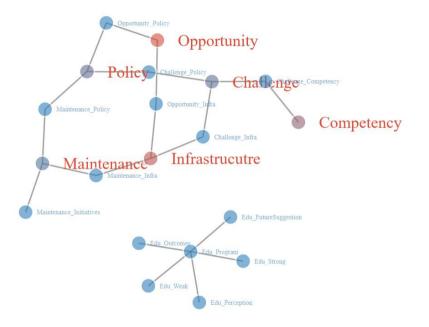
 Select all second-order using cursor, right click and press "Plot selected Code categories".

- Produce images of networks of codes
- Can re-arrange network codes using various network plot models (e.g

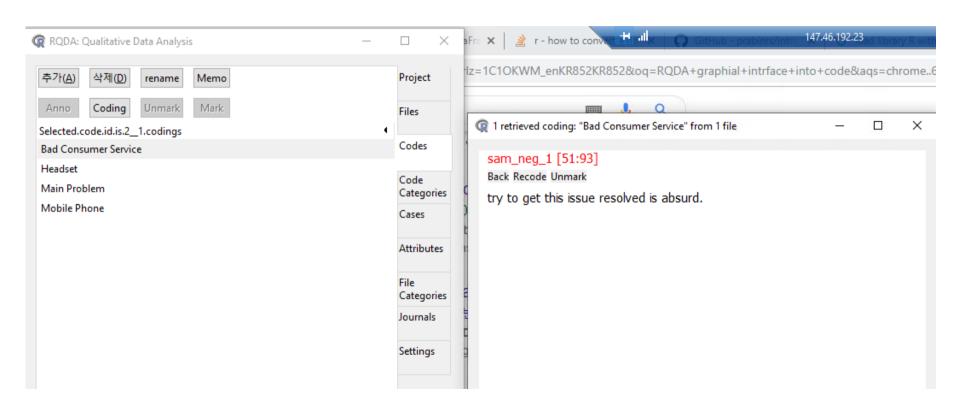
Allows User to visualize the coding results, enhance the data interpretation and analysis process. Transition from an inductive to abductive theorizing process- moving back and forth between the emergent findings and literature/theories to find new concepts within the data.

Visualize the concepts

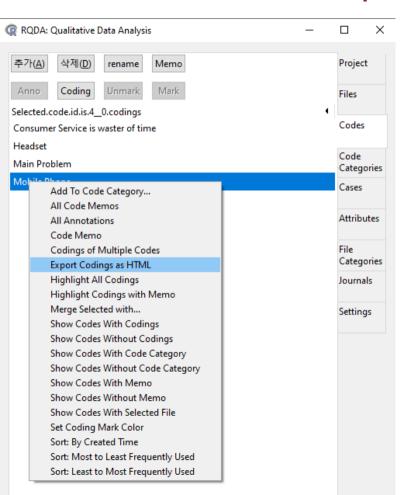


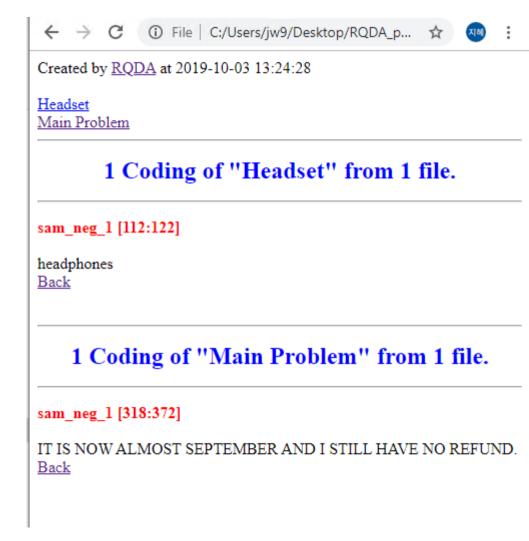


Review codes and Export



Review codes and Export





Complex Queries

- RQDA uses SQLite for the data management,
- SQL Queries can be used from R (not RQDA GUI)
- Build in Functions

```
#Load Data from RODA
openProject("leaders.rgda")
interviews <- RODAQuery("SELECT file FROM source")
interviews file <- apply(interviews, 1, function(x) qsub("...", "...", x))
interviews file <- apply(interviews, 1, function(x) qsub("'", "", x))
interviews <- Corpus(VectorSource(interviews$file))
interviews <- tm_map(interviews, stripWhitespace)</pre>
interviews <- tm_map(interviews, content_transformer(tolower))</pre>
interviews <- tm_map(interviews, removeWords, stopwords("english"))</pre>
interviews <- tm_map(interviews, removePunctuation)</pre>
interviews <- tm_map(interviews, removeNumbers)
interviews <- tm_map(interviews, removeWords, c("L01","L02","L03","L04" ))</pre>
interviews <- tm_map(interviews, removewords, c("prabin", "raj", "shakya","jk", "like", "jieun", "know"))
interviews <- tm_map(interviews, removeWords, stopwords(kind = "en"))
interviews <- tm_map(interviews, removeWords, c("prs", "will", "can", "quite",
                                                     "day", "yes", "even", "give",
"done", "also", "actually", "yeah", "things", "thing",
"see", "much", "just", "bit", "maybe",
                                                     "got", "kind", "tow", "example"))
#wordcloud
set.seed(1985)
wordcloud(interviews, min.freq = 10, max.words =50, rot.per=0.35,
          colors = brewer.pal(8, "Blues")[-1:-5])
#Topic analysis
leader_dtm<-DocumentTermMatrix(interviews)</pre>
leader_dtm<-removeSparseTerms(leader_dtm,0.99)</pre>
leaderTerms<-LDA(leader_dtm, k=5)</pre>
terms(leaderTerms.10)
```

Complex Queries

```
help(RQDA)
help(RQDATables")
help("RQDAQuery")
```

```
# Few useful inbuild functions
filesByCodes() # files contain codes with frequency
getCaseIds(fid) # Retrieve the number of cases and the case name
getCaseNames(caseID) # Return the names of the IDs
getCaseIds(fid) #Return the case Name or IDs
summaryCodings() # return coding frequencies |
getCodingTable() # get details on the codes
getCodingsByOne() # serach for specific code in the data.
codingBySearch() # finding speciific text in file and apply specific code.
```

```
filesByCodes() # files contain codes with frequency
fid filename codedBy.Bad Consumer Service codedBy.Headset codedBy.Main Problem
1 sam_neg_1 1 1 1
```

Codes are available in gitub, please open Rproj file and play around.

6. Theory Building

 Identify the relationships among themes / concepts / variables that emerged to create a process mode. This is manual process that can only be done using human interpreter.

iteratively re-categorized all first- and second-level code categories in a "data structure" to refine codes to the best possible higher-level categories that describe and explain the empirical reality and the literature. Should give clear ideas on who's and why's

