Feature selection

```
# library(extrafont)
# extrafont::loadfonts(quiet = TRUE)
set.seed(42)
library(igraph)
##
## Attaching package: 'igraph'
## The following objects are masked from 'package:stats':
##
##
       decompose, spectrum
## The following object is masked from 'package:base':
##
##
       union
library(tidyverse)
## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplyr
            1.1.4
                         v readr
                                      2.1.5
## v forcats 1.0.0
                         v stringr
                                      1.5.1
## v ggplot2 3.5.1
                        v tibble
                                     3.2.1
## v lubridate 1.9.3
                          v tidyr
                                      1.3.1
## v purrr
               1.0.2
## -- Conflicts ------ tidyverse_conflicts() --
## x lubridate::%--%()
                            masks igraph::%--%()
                        masks igraph::compose()
masks igraph::crossing()
masks stats::filter()
## x dplyr::as_data_frame() masks tibble::as_data_frame(), igraph::as_data_frame()
## x purrr::compose()
## x tidyr::crossing()
## x dplyr::filter()
## x dplyr::lag()
                            masks stats::lag()
## x apryr::rag()
## x purrr::simplify()
                            masks igraph::simplify()
## i Use the conflicted package (<a href="http://conflicted.r-lib.org/">http://conflicted.r-lib.org/</a>) to force all conflicts to become error
```

Load and tidy data

```
data <- read_csv("../measurements/measurements.csv")

## Rows: 754 Columns: 96

## -- Column specification ------

## Delimiter: ","

## chr (9): fpath, KUK_ID, class, FileName, FolderPath, subcorpus, DocumentTit...

## dbl (85): RuleAbstractNouns, RuleAmbiguousRegards, RuleAnaphoricReferences, ...

## 1gl (2): ClarityPursuit, SyllogismBased

##

## i Use `spec()` to retrieve the full column specification for this data.</pre>
```

```
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.
data clean <- data %>%
  select(!c(
   fpath,
    # KUK_ID,
    # FileName,
   FolderPath,
    # subcorpus,
   DocumentTitle,
   ClarityPursuit,
   Readability,
   SyllogismBased,
   SourceDB
  )) %>%
  # replace -1s in variation coefficients with NAs
  mutate(across(c(
    `RuleDoubleAdpos.max_allowable_distance.v`,
    `RuleTooManyNegations.max_negation_frac.v`,
    `RuleTooManyNegations.max_allowable_negations.v`,
    `RuleTooManyNominalConstructions.max_noun_frac.v`,
    `RuleTooManyNominalConstructions.max allowable nouns.v`,
    `RuleCaseRepetition.max_repetition_count.v`,
    `RuleCaseRepetition.max repetition frac.v`,
    `RulePredSubjDistance.max_distance.v`,
    `RulePredObjDistance.max distance.v`,
    `RuleInfVerbDistance.max distance.v`,
    `RuleMultiPartVerbs.max distance.v`,
    `RuleLongSentences.max_length.v`,
    `RulePredAtClauseBeginning.max_order.v`,
    `mattr.v`,
    `maentropy.v`
  ), ~ na_if(.x, -1))) %>%
  # replace NAs with Os
  replace_na(list(
   RuleGPcoordovs = 0,
    RuleGPdeverbaddr = 0,
   RuleGPpatinstr = 0,
   RuleGPdeverbsubj = 0,
   RuleGPadjective = 0,
   RuleGPpatbenperson = 0,
   RuleGPwordorder = 0,
   RuleDoubleAdpos = 0,
   RuleDoubleAdpos.max allowable distance = 0,
   RuleDoubleAdpos.max allowable distance.v = 0,
   RuleAmbiguousRegards = 0,
   RuleReflexivePassWithAnimSubj = 0,
   RuleTooManyNegations = 0,
   RuleTooManyNegations.max_negation_frac = 0,
   RuleTooManyNegations.max_negation_frac.v = 0,
   RuleTooManyNegations.max_allowable_negations = 0,
   RuleTooManyNegations.max_allowable_negations.v = 0,
   RuleTooManyNominalConstructions.max_noun_frac.v = 0,
    RuleTooManyNominalConstructions.max_allowable_nouns.v = 0,
```

```
RuleFunctionWordRepetition = 0,
 RuleCaseRepetition.max_repetition_count.v = 0,
 RuleCaseRepetition.max repetition frac.v = 0,
 RuleWeakMeaningWords = 0,
  RuleAbstractNouns = 0,
 RuleRelativisticExpressions = 0,
 RuleConfirmationExpressions = 0,
 RuleRedundantExpressions = 0,
 RuleTooLongExpressions = 0,
 RuleAnaphoricReferences = 0,
 RuleLiteraryStyle = 0,
 RulePassive = 0,
 RulePredSubjDistance = 0,
  RulePredSubjDistance.max_distance = 0,
 RulePredSubjDistance.max_distance.v = 0,
  RulePredObjDistance = 0,
 RulePredObjDistance.max_distance = 0,
 RulePredObjDistance.max_distance.v = 0,
 RuleInfVerbDistance = 0,
 RuleInfVerbDistance.max_distance = 0,
 RuleInfVerbDistance.max distance.v = 0,
 RuleMultiPartVerbs = 0,
 RuleMultiPartVerbs.max_distance = 0,
 RuleMultiPartVerbs.max_distance.v = 0,
 RuleLongSentences.max_length.v = 0,
 RulePredAtClauseBeginning.max_order.v = 0,
 RuleVerbalNouns = 0,
 RuleDoubleComparison = 0,
 RuleWrongValencyCase = 0,
 RuleWrongVerbonominalCase = 0,
 RuleIncompleteConjunction = 0
)) %>%
# norm data expected to correlate with text length
mutate(across(c(
 RuleGPcoordovs,
 RuleGPdeverbaddr,
 RuleGPpatinstr,
 RuleGPdeverbsubj,
 RuleGPadjective,
 RuleGPpatbenperson,
 RuleGPwordorder,
 RuleDoubleAdpos,
 RuleAmbiguousRegards,
 RuleFunctionWordRepetition,
 RuleWeakMeaningWords,
 RuleAbstractNouns,
 RuleRelativisticExpressions,
 RuleConfirmationExpressions,
 RuleRedundantExpressions,
  RuleTooLongExpressions,
 RuleAnaphoricReferences,
 RuleLiteraryStyle,
 RulePassive,
```

```
RuleVerbalNouns,
   RuleDoubleComparison,
   RuleWrongValencyCase,
   RuleWrongVerbonominalCase,
   RuleIncompleteConjunction,
   num_hapax,
   RuleReflexivePassWithAnimSubj,
   RuleTooManyNominalConstructions,
   RulePredSubjDistance,
   RuleMultiPartVerbs,
   RulePredAtClauseBeginning
  ), ~ .x / word_count)) %>%
  mutate(across(c(
    RuleTooFewVerbs,
   RuleTooManyNegations,
   RuleCaseRepetition,
   RuleLongSentences,
   RulePredObjDistance,
   RuleInfVerbDistance
  ), ~ .x / sent_count)) %>%
  # remove variables identified as "u counts"
  select(!c(
   RuleTooFewVerbs,
   RuleTooManyNegations,
   RuleTooManyNominalConstructions,
   RuleCaseRepetition,
   RuleLongSentences,
   RulePredAtClauseBeginning,
    sent_count,
   word_count,
    syllab_count,
    char_count
  )) %>%
  # remove variables identified as unreliable
  select(!c(
   RuleAmbiguousRegards,
   RuleFunctionWordRepetition,
   RuleDoubleComparison,
   RuleWrongValencyCase,
   RuleWrongVerbonominalCase
  )) %>%
  # remove further variables belonging to the 'acceptability' category
  select(!RuleIncompleteConjunction) %>%
  mutate(across(c(class), ~ as.factor(.x)))
# no NAs should be present now
data_clean[!complete.cases(data_clean), ]
## # A tibble: 0 x 73
## # i 73 variables: KUK_ID <chr>, class <fct>, FileName <chr>, subcorpus <chr>,
       RuleAbstractNouns <dbl>, RuleAnaphoricReferences <dbl>,
## #
       RuleCaseRepetition.max_repetition_count <dbl>,
## #
       RuleCaseRepetition.max_repetition_count.v <dbl>,
       RuleCaseRepetition.max_repetition_frac <dbl>,
## #
```

```
## # RuleCaseRepetition.max_repetition_frac.v <dbl>,
## # RuleConfirmationExpressions <dbl>, RuleDoubleAdpos <dbl>, ...

data_clean_scaled <- data_clean %>%
    mutate(across(class, ~ .x == "good")) %>%
    mutate(across(5:length(names(data_clean)), ~ scale(.x)))
```

Important features identification

```
data_clean_good <- data_clean_scaled %>% filter(class == "good")
data_clean_bad <- data_clean_scaled %>% filter(class == "bad")
feature_importances <- tibble(</pre>
  feat_name = character(), p_value = numeric()
for (i in 5:73) {
  fname <- names(data_clean)[i]</pre>
  formula_single <- reformulate(fname, "class")</pre>
  # print(formula_single)
  glm_model <- glm(formula_single, data_clean, family = "binomial")</pre>
  glm_coefficients <- summary(glm_model)$coefficients</pre>
  row_index <- which(rownames(glm_coefficients) == fname)</pre>
  p_value <- glm_coefficients[row_index, 4]</pre>
 feature_importances <- feature_importances %>%
    add_row(feat_name = fname, p_value = p_value)
}
feature_importances
## # A tibble: 69 x 2
      feat name
##
                                                      p value
                                                        <dbl>
##
      <chr>>
## 1 RuleAbstractNouns
                                                  0.00187
## 2 RuleAnaphoricReferences
                                                  0.660
## 3 RuleCaseRepetition.max_repetition_count
                                                  0.0722
## 4 RuleCaseRepetition.max_repetition_count.v 0.00479
## 5 RuleCaseRepetition.max_repetition_frac
                                                  0.00000740
## 6 RuleCaseRepetition.max_repetition_frac.v
                                                 0.000000472
## 7 RuleConfirmationExpressions
                                                  0.0985
## 8 RuleDoubleAdpos
                                                  0.312
## 9 RuleDoubleAdpos.max_allowable_distance
                                                  0.000154
## 10 RuleDoubleAdpos.max allowable distance.v 0.00000356
## # i 59 more rows
selected_features <- feature_importances %>%
  filter(p_value <= 0.05) %>%
  pull(feat_name)
```

Correlations

```
data_pure <- data_clean %>% select(any_of(selected_features))
cor_matrix <- cor(data_pure)</pre>
cor_tibble_long <- cor_matrix %>%
  as_tibble() %>%
 mutate(feat1 = selected_features) %>%
 pivot_longer(!feat1, names_to = "feat2", values_to = "cor") %>%
  mutate(abs_cor = abs(cor))
cor_matrix_upper <- cor_matrix</pre>
cor_matrix_upper[lower.tri(cor_matrix_upper)] <- 0</pre>
cor_tibble_long_upper <- cor_matrix_upper %>%
 as_tibble() %>%
 mutate(feat1 = selected_features) %>%
 pivot_longer(!feat1, names_to = "feat2", values_to = "cor") %>%
 mutate(abs_cor = abs(cor)) %>%
 filter(feat1 != feat2)
cor_tibble_long_upper_considerable <- cor_tibble_long_upper %>%
  filter(abs_cor >= 0.3)
max_correlations <- cor_tibble_long %>%
  filter(feat1 != feat2) %>%
  group_by(feat1) %>%
  summarize(maxcor = max(abs cor)) %>%
 ungroup()
```

Visualisation

```
library(paletteer)
my_colors <- paletteer::paletteer_d("ggthemes::Classic_10_Medium")</pre>
network <- graph_from_data_frame(</pre>
  cor_tibble_long_upper_considerable,
  directed = FALSE
E(network)$weight <- cor_tibble_long_upper_considerable$abs_cor</pre>
network_communities <- cluster_optimal(network)</pre>
network_membership <- membership(network_communities)</pre>
plot(
  network,
  layout = layout.fruchterman.reingold,
  vertex.color = map(
    network_communities$membership,
    function(x) my_colors[x]
  ) %>% unlist(use.names = FALSE),
  vertex.size = 6,
```

```
# vertex.frame.color = "#000000000",
# vertex.label.family = "Public Sans",
vertex.label.color = "black",
vertex.label.cex = 0.5
)
```

