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StatKeyEval

<u>A Statistical Framework for Dynamic Keyword Extraction, Evaluation, and Assessment Automation</u>

Aim:

To implement an API for the available research paper models to give the answers for the short answers.

Research Paper:

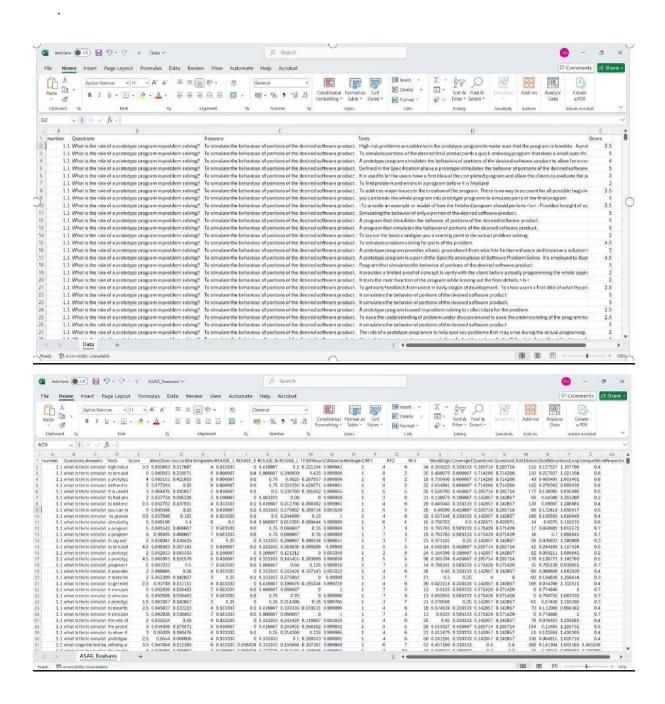
Title: Feature Engineering and Ensemble-Based Approach for Improving Automatic Short-Answer Grading Performance

Authors: Archana Sahu and Plaban Kumar Bhowmick.

Conference/Journal: Educational Data Mining Conference (2018)

Datasets:

- 1. UNT Dataset
- 2. SciEntsBank Dataset
- 3. Beetle Dataset



Statistical Function for Keyword Extraction

 $\operatorname{Relevance}(W) = \log \left(1 + \operatorname{FreqRatio}(W)\right)^{\alpha} \times \left[1 + \operatorname{InverseDistance}(W)\right]^{\beta} \times \operatorname{Specificity}(W)^{\gamma}$

Where:

• Relevance(W) is the final importance score for word W in the lexical extraction process

- FreqRatio(W) is the ratio of word W's frequency in relevant contexts to its frequency in irrelevant contexts plus a smoothing constant δ: frequency in relevant / (frequency in irrelevant + δ)
- InverseDistance(W) is the reciprocal of the average distance to other key terms plus 1: 1/(average distance to other key terms + 1)
- Specificity(W) is a measure of word uniqueness calculated as log(total corpus words / document frequency of W)
- α , β , and γ are tunable exponential parameters that control the influence of each component (typical values: α =0.5, β =0.7, γ =0.4)

Statistical Function for Keyword Mutation

ExpandMetric(
$$K_1, K_2$$
) = [SymbioticOverlap(K_1, K_2) × LogisticDecay($|K_1|, |K_2|$)] × $[1 + \log(1 + \text{SemanticDensity}(K_1 \cap K_2))]$

Where: • ExpandMetric (K_1, K_2) is the final expansion benefit score between keyword sets

K₁ and K₂

- SymbioticOverlap(K₁, K₂) is the quadratic overlap measure calculated as $|K_1 \cap K_2|^2 / (|K_1| \times |K_2|)$
- LogisticDecay($|K_1|$, $|K_2|$) is a balanced size similarity function calculated as $2/(1 + \exp(|abs(|K_1| |K_2|)|/\lambda))$, where λ is a scaling parameter
- SemanticDensity($K_1 \cap K_2$) is the sum of co-occurrence frequencies for all word pairs in the intersection
- $|K_1|$ and $|K_2|$ are the cardinalities (sizes) of the keyword sets
- log is the natural logarithm function
- exp is the exponential function

Statistical Functions for Score calculation

AdjustedMetric(R) =
$$M \times \left[1 - e^{-R/\tau}\right] \times \left[1 + \beta \times \sin\left(\frac{\pi \times R}{2M}\right)\right]$$

Where:

- AdjustedMetric(R) is the final calibrated value after quantization to the nearest 0.5
- R is the original raw measurement
- M is the maximum threshold value
- τ is a scaling coefficient governing the saturation rate
- β is a modulation parameter controlling oscillation intensity
- e is the base of the natural logarithm

• sin is the sine function

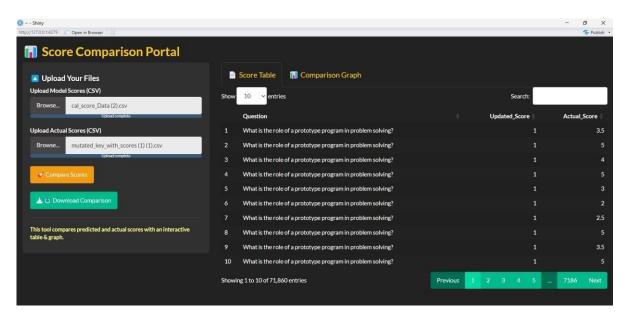
Code for Performance Graph:

Code:

```
# Load necessary libraries if (!require("shiny"))
install.packages("shiny") if (!require("ggplot2"))
install.packages("ggplot2") if (!require("DT"))
install.packages("DT") if (!require("readr"))
install.packages("readr") if (!require("dplyr"))
install.packages("dplyr") if (!require("Metrics"))
install.packages("Metrics")
library(shiny)
library(ggplot2)
library(DT) library(readr)
library(dplyr)
library(Metrics)
# Define UI ui
<- fluidPage(
 titlePanel("Score Comparison Portal"),
sidebarLayout(
                  sidebarPanel(
   fileInput("file model", "Upload Model Scores (CSV)", accept = c(".csv")),
fileInput("file existing", "Upload Actual Scores (CSV)", accept = c(".csv")),
actionButton("compare btn", "Compare Scores")
  ),
  mainPanel(
                  tabsetPanel(
                                    tabPanel("Score
Table", DTOutput("score table")),
     tabPanel("Comparison Graph", plotOutput("comparison plot"))
   )
  )
 )
# Define Server logic
server <- function(input, output, session)</pre>
 { results <- reactiveValues(data = NULL)
 observeEvent(input$compare btn,
{ req(input$file model, input$file existing)
  model data <- read csv(input$file model$datapath) %>% select(Questions, Calculated Score)
existing data <- read csv(input$file existing$datapath) %>% select(Questions, Score)
```

```
colnames(model data) <- c("Question", "Updated Score")
colnames(existing data) <- c("Question", "Actual Score")
  comparison data <- model data %>%
   inner_join(existing_data, by = "Question")
  results$data <- comparison data
 })
 # Render Data Table
output$score table <-
renderDT({ req(results$data)
  datatable(results$data, options = list(pageLength = 10))
 })
 # Render Comparison Graph output$comparison plot <-
renderPlot({ req(results$data) ggplot(results$data, aes(x = Updated Score, y =
Actual Score))
     geom point(color = "blue", size = 3) +
                                              geom smooth(method = "lm", se =
FALSE, color = "red", linetype = "dashed") +
                                              labs(title = "Comparison: Updated
Score vs. Actual Score",
                              x = "Updated Score (Calculated)", y = "Actual
Score")+
             theme_minimal()
 })
# Run the app
shinyApp(ui = ui, server = server)
```

Output:





INPUT FILE LINK:

https://drive.google.com/file/d/108GJwpX2duVMxNAOZK6cKh0kc3n6wees/view?usp=sharing

https://drive.google.com/file/d/10r2z1S11ffp wGnfYmyEvKfw1LFH717s/view?usp=sharing

OUTPUT FILE LINK:

https://drive.google.com/file/d/1m7MykvAg7lEBvyVSePob4iA-XNBGhO6Q/view?usp=sharing

CODE FOR API: try to build upon with this code: if (!require("shiny")) install.packages("shiny") if

(!require("tm")) install.packages("tm") if

(!require("dplyr")) install.packages("dplyr") if

(!require("readr")) install.packages("readr") if

(!require("DT")) install.packages("DT")

library(shiny) library(tm)

library(dplyr)

library(readr) library(DT)

Basic functions extract keywords <- function(text) { text

<- tolower(text) words <- unlist(strsplit(gsub("[[:punct:]]",

```
"", text), "\\s+")) words <- words[words != ""] stopwords <- c("the", "and",
"a", "an", "in", "on", "at", "to", "for", "of", "with",
"is", "are", "was", "were", "be", "been", "being", "have", "has", "had",
          "do", "does", "did", "can", "could", "will", "would", "should", "may",
          "might", "must", "shall", "this", "that", "these", "those", "it", "its",
          "they", "them", "their", "we", "us", "our", "i", "me", "my", "you", "your")
words <- words[!words %in% stopwords] words <- words[nchar(words) > 1]
words <- words[!grepl("^\\d+$", words)] return(unique(words))
}
calculate score <- function(answer keywords, student keywords)</pre>
{ jaccard sim <- length(intersect(answer keywords, student keywords)) /
length(union(answer keywords, student keywords)) raw score <- jaccard sim
* 5 wpcs score <- min(raw score * 1.05, 5) wpcs score
<-round(wpcs score * 2) / 2
return(list( raw score = raw score,
wpcs score = wpcs score,
similarity = jaccard sim
 ))
}
SCM <- function(corpus, answer keywords, student keywords, threshold = 0.3) { if (length(corpus)
== 0 \parallel \text{length(answer keywords)} == 0 \parallel \text{length(student keywords)} == 0)
{ return(list(mutation candidates = list(), similarity score = 0))
 corpus <- lapply(corpus, function(x) if(length(x) == 0) c("") else x)
candidates <- setdiff(student keywords, answer keywords) if
(length(candidates) == 0) { return(list(mutation_candidates = list(),
similarity score = 0)
```

```
}
 candidate_freq <- sapply(candidates, function(word)</pre>
{ sum(sapply(corpus, function(doc) word %in% doc))
 })
 candidate rel freq <- candidate freq / length(corpus) mutation candidates
<- list() for (i in
1:length(candidates)) {
                          word <- candidates[i]
                                                   freq
<- candidate_rel_freq[i]
                          if (freq >= threshold)
      mutation_candidates[[word]] <- list(
                                                word
= word,
              score = freq,
                                uniqueness = 1 - freq
   )
  }
 if (length(mutation candidates) > 0) {
                                          sorted candidates
<-
                                 mutation candidates[order(
sapply(mutation candidates,
                                  function(x)
                                                   x$score),
decreasing = TRUE
  )]
 } else {
            sorted candidates
<- list()
 }
 jaccard sim <- length(intersect(answer keywords, student keywords))/
length(union(answer keywords, student keywords))
return(list(
              mutation candidates =
sorted candidates,
                      similarity score = jaccard sim
 ))
}
```

```
update keywords <- function(question data) { answer keywords <-
unlist(strsplit(question data$Answer Keywords[1], ", ")) all text keywords <-
lapply(question data$Text Keywords, function(x) { if (is.na(x) || x == "")
                     unlist(strsplit(x, ", "))
return(character(0))
 })
 all student keywords <- unique(unlist(all text keywords)) threshold <- 0.65 scm result
<- SCM(all text keywords, answer keywords, all student keywords, threshold)
mutation candidates <- scm result$mutation candidates new keywords <-
names(mutation candidates)
                             return(list( new keywords = paste(new keywords,
collapse = ", "), similarity score = scm result$similarity score
))
}
process_batch <- function(data) { if ("Score" %in% colnames(data) &&
!"WPCS Score" %in% colnames(data)) {
                                         data <- data %>%
                                         mutate( Score =
    as.numeric(Score),
    WPCS Score = pmin(Score * 1.05, 5),
    WPCS Score = round(WPCS Score *2) / 2
   )
 if (all(c("Questions", "Answer Keywords", "Text Keywords") %in% colnames(data)))
{ result <- data %>% group by(Questions) %>%
                                                   group modify(~{
                                                                        mutation result
<- update keywords(.x)
    .x$New Keywords <- mutation result$new keywords
    .x$Similarity Score <- mutation result$similarity score
    .x$Combined Keywords <- ifelse(.x$New Keywords!="",
paste(.x$Answer Keywords, .x$New Keywords, sep = ", "),
```

```
.x$Answer_Keywords)
return(.x)
            }) %>%
                        ungroup()
                                    return(result)
 }
return(data)
}
# Ultra simple UI ui <-
fluidPage(titlePanel("Keyword Analysis"),
#Single Analysis Tab h3("Single
Analysis"),
textInput("question", "Question"), textAreaInput("model answer", "Model
Answer"), textInput("manual keywords", "Model Answer Keywords (comma-
separated)"), textAreaInput("student_answer", "Student Answer"), actionButton("analyze_btn",
"Analyze"),
hr(),
h4("Results:"), verbatimTextOutput("score_output"),
verbatimTextOutput("keywords output"), hr(),
#Batch Processing Tab h3("Batch
Processing"),
 fileInput("file upload", "Upload CSV File"), checkboxInput("header",
"File has header", TRUE), actionButton("process btn", "Process"),
downloadButton("download results", "Download"),
```

```
hr(),
 DTOutput("results_table"), verbatimTextOutput("batch_stats")
)
# Server logic server <- function(input,
output, session) { results <--
reactiveValues( model_keywords = NULL,
student_keywords = NULL, score =
NULL, batch data = NULL,
processed data = NULL
 )
 observeEvent(input$analyze btn,
{ req(input$model answer,
input$student answer)
  if (input$manual keywords != "") {
                                   model kw <-
unlist(strsplit(input$manual keywords, ", "))
                                          model kw <- trimws(model kw)
  } else {
   model kw <- extract keywords(input$model answer)</pre>
  student_kw <- extract_keywords(input$student_answer) score_result
<- calculate score(model kw, student kw)
  <- student_kw
              results$score
<- score result
```

```
output$score output <- renderPrint({</pre>
                                           cat("WPCS Score: ", results$score$wpcs score,
"/5.0\n", sep = "")
                     cat("Similarity: ", round(results\score\similarity * 100, 2), "\n", sep
        cat("Matching: ", length(intersect(results$model keywords,
results\student_keywords)),
      "out of", length(union(results$model keywords, results$student keywords)), "\n", sep="")
  })
  output$keywords output <- renderPrint({</pre>
                                                cat("Model
Answer Keywords:\n")
                           cat(paste(results$model keywords,
collapse = ", "), "\n\n")
                           cat("Student Answer Keywords:\n")
                  cat(paste(results$student keywords,
collapse = ", "))
  })
 })
 observeEvent(input$file_upload,
{ req(input$file upload)
                 batch data <- read csv(input$file upload$datapath, col names = input$header)
  tryCatch({
results$batch data <- batch data
   output$results table <-
renderDT({ datatable(results$batch data, options =
list(pageLength = 5))
   })
                             showNotification("Error reading
  }, error = function(e) {
file", type = "error")
  })
```

```
})
 observeEvent(input$process_btn,
{ req(results$batch data)
  processed data <- process batch(results$batch data)
                                                       results$processed data
<- processed data
  output$results table <-
renderDT({ datatable(results$processed_data, options =
list(pageLength = 5))
  })
  output$batch stats <- renderPrint({
                                       if
("WPCS_Score" %in% colnames(processed_data)) { cat("Records:
", nrow(processed data), "\n")
    if ("Score" %in% colnames(processed data)) { cat("Average Score: ",
round(mean(processed_data\$Score, na.rm = TRUE), 2), "\n") cat("Average WPCS: ",
round(mean(processed data$WPCS Score, na.rm = TRUE), 2), "\n")
    }
    if ("Similarity Score" %in% colnames(processed data)) {
     cat("Average Similarity: ", round(mean(processed_data$Similarity_Score, na.rm = TRUE) *
100, 2), "%\n")
    }
   } else {
               cat("Data
loaded")
}
  })
```

shinyApp(ui = ui, server = server)<u>RESULT:</u>

