## **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

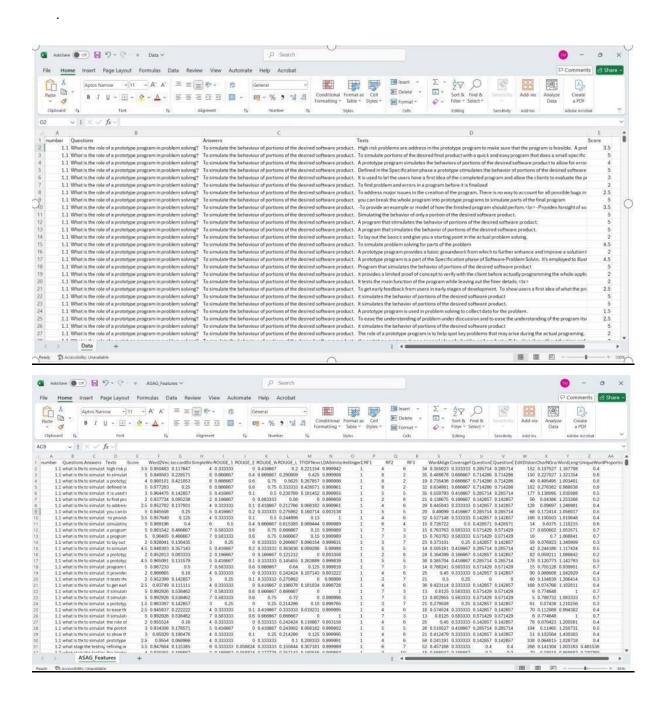
Grading Performance

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**

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

## 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)
- $\alpha$ ,  $\beta$ , and  $\gamma$  are tunable exponential parameters that control the influence of each component (typical values:  $\alpha$ =0.5,  $\beta$ =0.7,  $\gamma$ =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

K<sub>1</sub> and K<sub>2</sub>

- SymbioticOverlap(K<sub>1</sub>, K<sub>2</sub>) 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  $\lambda$  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
- $\tau$  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

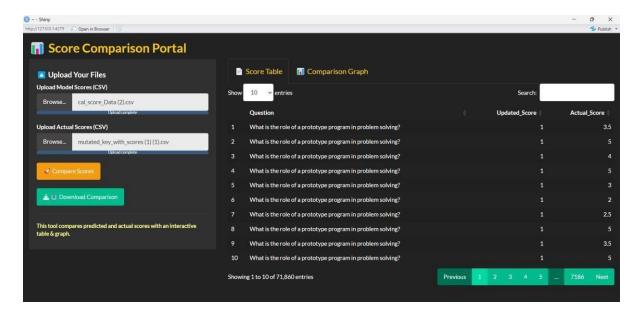
# **Code for Performance Graph:**

#### **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

<u>CODE FOR API:</u> 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) { 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 length(answer keywords) == 0 \parallel 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(</pre>
                                                 word
              score = freq,
                                 uniqueness = 1 - freq
= word,
    )
  }
 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_dataText_Keywords, function(x) { if (is.na(x) || x = "") return(character(0))
unlist(strsplit(x, ", "))
 })
 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))) {
<- 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)
}) %>%
                      return(result)
           ungroup()
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 <- reactive Values(
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$student answer) score result
<- calculate score(model kw, student kw)
  results$model keywords <- model kw results$student keywords
<- student kw results$score
<- score result
  output$score output <- renderPrint({ cat("WPCS Score: ", results$score$wpcs score,
                    cat("Similarity: ", round(results$score$similarity * 100, 2), "%\n", sep
"/5.0\n", sep = "")
        cat("Matching: ", length(intersect(results$model keywords, results$student keywords)),
     " out of ", length(union(results$model keywords, results$student keywords)), "\n", sep="")
  })
  output$keywords_output <- renderPrint({
                                             cat("Model Answer
                  cat(paste(results$model keywords, collapse = ",
Keywords:\n")
"), "\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$batch data,
options = list(pageLength = 5))
    })
  }, error = function(e) {       showNotification("Error reading file",
type = "error")
  })
 })
 observeEvent(input$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$batch stats <- renderPrint({</pre>
("WPCS Score" %in% colnames(processed data)) {
                                                    cat("Records:
", nrow(processed data), "\n")
    if ("Score" %in% colnames(processed data)) { cat("Average Score: ",
round(mean(processed_data$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")
    }
             cat("Data loaded")
   } else {
}
  })
 })
 output$download results <- downloadHandler(
                                                 filename
= function() {
   "results.csv"
  },
       content = function(file) {
write csv(results$processed data, file)
  }
```

}

shinyApp(ui = ui, server = server)RESULT:

