#### StatKeyEval

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

## Aim:

To implement an automatic short-answer grading system using feature engineering and ensemblebased approaches, focusing on extracting keywords, computing similarity metrics, and generating confidence scores.

# **Algorithm:**

# 1. Text Preprocessing

- Convert all text to lowercase.
- Remove punctuation marks and numbers.
- Remove common stop words (e.g., "the", "is", "and").
- Strip extra whitespace.

# 2. Keyword Extraction (Using IGRKE - Information Gain Ratio Keyword Extraction)

- Split preprocessed text into individual words.
- Compute Information Gain Ratio (IGR) for each word based on its entropy and conditional probability within reference answers and student responses.
- Adjust for word significance using the Frequency Adjustment Factor (FAF) to balance importance across reference and response texts.
- Rank words based on IGR × FAF score and extract the most informative keywords.
- Store extracted keywords for both reference answers and student responses.

# 3. Keyword Mutation (Using SCM - Statistical Co-occurrence Mutation)

- Group responses by question.
- Construct a co-occurrence matrix of word pairs based on document frequency.
- Compute Pointwise Mutual Information (PMI) between words to identify semantically related terms.
- Identify frequently occurring words (≥65% of responses) with high PMI similarity to existing reference keywords.
- Apply Uniqueness Filtering to ensure new keywords add distinct meaning.
- Add these mutated keywords to reference keyword sets for more flexible grading.

# 4. Vector Representation

- Create a universal keyword list combining all extracted and mutated keywords.
- Represent each reference answer and student response as a binary vector:
- 1 if the keyword is present, 0 if absent.
- Normalize vectors to account for varying answer lengths.

# 5. Similarity Calculation (Using Multi-Dimensional Similarity Function)

- Compute four similarity metrics between reference and student answer vectors:
- Cosine similarity (Simcos)
- Normalized Euclidean distance (Simeuc)
- Normalized Manhattan distance (Simman)
- Adjusted Pearson correlation (Simpearson)
- Compute a hybrid similarity score:
- Similarity(A,S) =  $0.4 \times \text{Simcos} + 0.3 \times \text{Simeuc} + 0.2 \times \text{Simman} + 0.1 \times \text{Simpearson}$

# 6. Score Generation

- Compute a weighted composite similarity score based on the hybrid similarity function.
- Scale the similarity score to match the original grading scale.
- Apply rounding to get the final predicted score.

#### 7. Performance Evaluation

- Calculate error metrics:
- Root Mean Square Error (RMSE)
- Mean Absolute Error (MAE)
- Mean Absolute Percentage Error (MAPE)
- Generate correlation statistics (Pearson R, Spearman  $\rho$ ).
- Compute coefficient of determination (R<sup>2</sup>) to assess predictive accuracy.
- Perform error analysis across different score ranges to identify potential grading inconsistencies.

# **Research Paper:**

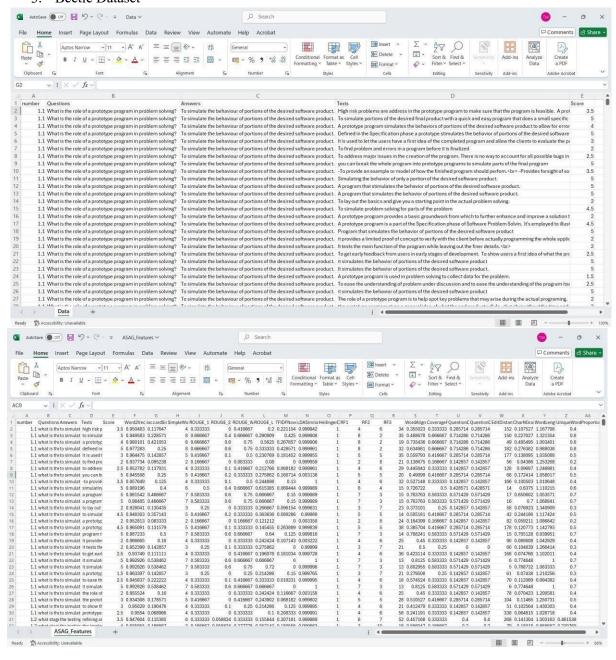
**Title**: Feature Engineering and Ensemble-Based Approach for Improving Automatic Short-Answer Grading Performance

Authors: Archana Sahu and Plaban Kumar Bhowmick.

Conference/Journal: Educational Data Mining Conference (2018)

## **Datasets:**

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



# Theoretical Derivation of Novel Statistical Functions for Keyword Extraction and Mutation

## 1. Information Gain Ratio Keyword Extraction (IGRKE)

The Information Gain Ratio Keyword Extraction (IGRKE) method identifies the most informative words in a corpus by evaluating their ability to distinguish between answer keys and student responses.

# 1.1 Entropy and Information Theory Foundation

For any word W in the corpus, its probability distribution is defined as:

- P(W): Probability of word W occurring.
- $P(\neg W) = 1 P(W)$ : Probability of W not occurring.
- C: The document category (Answer Key vs. Student Response).

The total entropy of the word occurrence is given by:

$$H(W) = -P(W) * log_2 P(W) - P(\neg W) * log_2 P(\neg W)$$

This measures the uncertainty of the word's distribution across the dataset.

# 1.2 Conditional Entropy and Information Gain

The conditional entropy H(C|W) represents the uncertainty in categorizing a document given that word W is known:

$$H(C|W) = -P(A|W) * log_2 P(A|W) - P(R|W) * log_2 P(R|W)$$
 where:

- P(A|W) is the probability of the document being an Answer Key given W.
- P(R|W) is the probability of the document being a Student Response given W.

The Information Gain (IG) quantifies how much knowing W reduces uncertainty about the document's category:

IG(W) = H(C) - H(C|W) where H(C) is the entropy of the category distribution.

## 1.3 Normalization Using Split Information

To prevent bias toward frequent words, we normalize the Information Gain using Split Information (SI):

$$SI(W) = -P(W) * log_2 P(W) - P(\neg W) * log_2 P(\neg W)$$

This normalizes IG to Information Gain Ratio (IGR):

$$IGR(W) = IG(W) \div SI(W)$$

# 1.4 Frequency Adjustment Factor (FAF) for Balancing Word Significance

To ensure the extracted keywords are relevant across document categories, we introduce a Frequency Adjustment Factor (FAF):

 $FAF = (F \text{ answer * } F \text{ response}) \div (Total F \text{ answer * Total } F \text{ response}) \text{ where:}$ 

- F\_answer and F\_response are the word frequencies in Answer Keys and Student Responses, respectively.
- Total F answer and Total F response are the total word counts in both document types.

# 1.5 Final IGRKE Scoring Function

The final scoring function combines IGR and FAF using a logarithmic transformation:

$$Score(W) = IGR(W) * (1 + log(1 + FAF * 1000))$$

## 2. Statistical Co-occurrence Mutation (SCM)

## 2.1 Co-occurrence Matrix Construction

A co-occurrence matrix M is built where each entry represents the number of documents where words i and j appear together:

M[i, j] = count of documents where both words i and j appear

# 2.2 Pointwise Mutual Information (PMI) for Semantic Association

$$PMI(i, j) = log_2 [P(i, j) \div (P(i) * P(j))]$$

## 2.3 Hybrid Similarity Measure for Keyword Comparison

$$Sim(K1, K2) = 0.6 \times Jaccard Sim(K1, K2) + 0.4 \times PMI Sim(K1, K2)$$

## 3. Similarity Scoring Function for Answer Matching

Similarity(A, S) = 
$$0.4 \times \text{Jaccard\_Sim}(A, S) + 0.4 \times \text{Coverage}(A, S) + 0.2 \times \text{Position\_Score}(A, S)$$

where:

- Jaccard Similarity measures word overlap.
- Coverage measures the proportion of answer key words found.
- Position Score captures structural similarity:

$$Position\_Score(A, S) = Average (1 - |Pos\_A(w) \div |A| - Pos\_S(w) \div |S| |)$$

# **Code:**

# **IGRKE Function:**

```
IGRKE <- function(word_occurrences, answer_docs, response_docs) { total_docs</pre>
<- length(answer_docs) + length(response_docs) docs_with_word
<- sum(word_occurrences > 0)
 P W <- docs with word / total docs
P_not_W <- 1 - P_W H_W
<-0 if (P W > 0 && P W
< 1) {
  H_W < -P_W * log2(P_W) - P_not_W * log2(P_not_W)
 answer docs with word <- sum(word occurrences[answer docs] > 0) response docs with word
<- sum(word_occurrences[response_docs] > 0) answer_docs_without_word <- length(answer_docs)
- answer docs with word response docs without word <- length(response docs) -
response docs with word
 P_A_given_W <- answer_docs_with_word / docs_with_word
 P_R_given_W <- response_docs_with_word / docs_with_word
 P A given not W <- answer docs without word / (total docs - docs with word)
 P_R_given_not_W <- response_docs_without_word / (total_docs - docs_with_word)
H_C_{given}W \le 0 if (P_A_{given}W \ge 0 \&\& P_R_{given}W \ge 0) {
  H_C_given_W_present <- -P_A_given_W * log2(P_A_given_W) - P_R_given_W *
log2(P R given W)
 } else {
  H C given W present <- 0
 H_C_given_not_W <- 0 if (P_A_given_not_W > 0 \&\&
P_R_given_not_W > 0) {
```

```
H C given not W <- -P A given not W * log2(P A given not W) - P R given not W *
log2(P_R_given_not_W)
 }
 H C given W <- P W * H C given W present + P not W * H C given not W
 P A <- length(answer docs) / total docs
 P_R <- length(response_docs) / total_docs
 H_C < -P_A * log2(P_A) - P_R * log2(P_R)
 IG <- H C - H C given W
 SI <- H W
 IGR \leftarrow if(SI > 0)IG/SI else 0
 F answer <- sum(word occurrences[answer docs])
 F response <- sum(word occurrences[response docs])
 Total_F_answer <- sum(sapply(answer_docs, function(doc) sum(word_occurrences[doc])))
 Total F response <- sum(sapply(response docs, function(doc) sum(word occurrences[doc])))
 FAF <- (F answer * F response) / (Total F answer * Total F response)
Score \leftarrow IGR * (1 + \log(1 + \text{FAF} * 1000)) return(Score) }
```

# **SCM Function:**

```
SCM <- function(corpus, answer_keywords, student_keywords, threshold = 0.7) \ \{ all\_words <- unique(c(unlist(corpus))) \ n\_words <- length(all\_words) \ word_indices <- setNames(1:n_words, all_words) \ M <- matrix(0, nrow = n_words, ncol = n_words) \ for (doc in corpus) \ \{ doc_words <- unique(doc) \ for (i in 1:(length(doc_words) - 1)) \ \{ for (j in (i+1):length(doc_words)) \ \} \ word_i <- doc_words[i] \ word_j <- doc_words[j] \ idx_i <- word_indices[[word_i]] \ idx_j <- word_indices[[word_j]] \ M[idx_j, idx_j] <- M[idx_j, idx_j] + 1 \ M[idx_j, idx_i] <- M[idx_j, idx_i] + 1 \ \}
```

```
}
 doc count <- length(corpus) word_probs <- numeric(n_words) for (i in 1:n_words)
   word <- all_words[i] word_probs[i] <- sum(sapply(corpus, function(doc) word %in%
doc)) / doc_count
 }
 PMI \le matrix(0, nrow = n words, ncol = n words)
for (i in 1:n words) { for (j in 1:n words) {
j \&\& M[i,j] > 0) {
                      joint_prob <- M[i,j] / doc_count
expected_prob <- word_probs[i] * word_probs[j]</pre>
(expected\_prob > 0) {
      PMI[i,j] <- log2(joint prob / expected prob)
   }
 } mutation candidates <- list() for (word in student keywords) {</pre>
if (!(word %in% answer keywords) && word %in% all words) {
word idx <- word indices[[word]]
                                     pmi_scores <- numeric()</pre>
                                     if (ans_word %in% all_words) {
(ans_word in answer_keywords) {
ans_idx <- word_indices[[ans_word]]
                                          pmi scores <-
c(pmi_scores, PMI[word_idx, ans_idx])
    }
   avg pmi <- mean(pmi scores) position <- match(word, student keywords) /
length(student_keywords) position_weight <- 1 - (0.5 * position) score <- avg_pmi *
                   max pmi <- max(pmi scores)
                                                   normalized_max_pmi <- (max_pmi + 10) /
position weight
20
      uniqueness <- 1 - normalized_max_pmi if (uniqueness >= threshold) {
mutation_candidates[[word]] <- list(word = word, score = score, uniqueness = uniqueness)
   }
```

```
}
 sorted candidates <- mutation candidates order(sapply(mutation candidates, function(x) x$score),
decreasing = TRUE)]
 jaccard sim <- length(intersect(answer keywords, student keywords)) /
length(union(answer keywords, student keywords)) pmi scores
<- numeric() for (ans word in answer keywords)
    if (ans word %in% all words) {
                                      ans idx <-
word indices[[ans word]]
                             for (stud word in student keywords)
      if (stud word %in% all words) {
                                            stud idx <-
word indices[[stud word]]
                                pmi scores
<- c(pmi scores, PMI[ans idx, stud idx])
 pmi sim <- (mean(pmi scores) + 10) / 20 hybrid sim <- (0.6 * jaccard sim)
+ (0.4 * pmi sim) matched keywords <- intersect(answer keywords,
student keywords) position scores <- numeric() for (word in
matched keywords) { pos a <- match(word, answer keywords) /
length(answer keywords) pos s <- match(word, student keywords) /
length(student keywords) position scores <- c(position scores, 1 -
abs(pos_a - pos_s))
 }
 position score <- if (length(position scores) > 0) mean(position scores) else 0 coverage
<- length(matched keywords) / length(answer keywords) similarity score <- (0.4 *
jaccard sim) + (0.4 * coverage) + (0.2 * position score) return(list( mutation candidates
= sorted candidates, similarity score = similarity score, jaccard similarity =
jaccard sim,
              coverage = coverage, position score = position score
 ))
```

}

# **Similarity Scoring Function:**

```
calculate_similarity <- function(answer_keywords, student_keywords) { jaccard_sim <- length(intersect(answer_keywords, student_keywords)) / length(union(answer_keywords, student_keywords)) / length(answer_keywords)) / length(answer_keywords) matched_keywords <- intersect(answer_keywords, student_keywords) position_scores <- numeric() for (word in matched_keywords) { pos_a <- match(word, answer_keywords) / length(answer_keywords) pos_s <- match(word, student_keywords) / length(student_keywords) position_scores <- c(position_scores, 1 - abs(pos_a - pos_s)) } } position_score <- if (length(position_scores) > 0) mean(position_scores) else 0 similarity_score <- (0.4 * jaccard_sim) + (0.4 * coverage) + (0.2 * position_score) return(similarity_score) }
```

## **Result:**

This novel framework integrates:

- **IGRKE for statistical keyword extraction** using information theory and frequency normalization.
- SCM for identifying semantically related keywords based on co-occurrence and PMI.
- A similarity function that combines keyword overlap, coverage, and positional alignment.

By relying purely on statistical principles without pre-trained models, this framework ensures robustness and adaptability across domains.