CFC Annotator: A Cluster-Focused Combination Algorithm for Annotating Electronic Health Records by Referencing Interface Terminology

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Abstract: In this paper, we present a novel algorithm designed to address the challenge of annotating electronic health record (EHR) text using an interface terminology dataset. Annotated text datasets are essential for the continued development of Large Language Models (LLMs). However, creating these datasets is labor-intensive and time-consuming, highlighting the urgent need for automated annotation methods. Our proposed method, the Cluster-Focused Combination (CFC) Algorithm, which stores intermediate results to minimize annotation loss from terminology-based annotators, such as BioPortal's (*mgrep*), while achieving high coverage and significantly improving execution efficiency. We conduct a thorough evaluation of CFC on the benchmark dataset MIMIC-III, using the previously developed Cardiology Interface Terminology (CIT). Results show that CFC captured approximately 5,756 missed annotations from the baseline BioPortal (*mgrep*) while achieving a remarkable improvement in execution speed across different size of datasets. These findings demonstrate CFC's scalability and robustness in processing large datasets, offering an efficient solution for EHR text annotation. This work contributes to the preparation of large, high-quality training datasets for Natural Language Processing (NLP) tasks in biomedical domains.

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1 INTRODUCTION

The field of healthcare has undergone significant transformation in recent decades, driven by the widespread adoption and application of Electronic Health Records (EHRs) (Blumenthal, 2009). EHRs, which include diagnoses, radiology descriptions, discharge summaries, and lab reports, are particularly valuable as they contain up-to-date, patient-specific information. However, despite their importance in detailing individual patient conditions, much of this information is recorded as unstructured text, often using highly specialized clinical jargons.

To enable interoperability and enhance healthcare quality through post hoc research, it is critical to annotate these records with concepts from standardized terminologies. Without such annotations, the text remains ambiguous, vague, and unsuitable for automated processing. Annotated data not only enhances information sharing but also empowers downstream applications, including clinical decision support systems, disease surveillance, and the development of cutting-edge Large Language Models (LLMs). Numerous annotation tools have been developed

Numerous annotation tools have been developed to extract information from biomedical literature and EHRs. Some of the most widely used tools in the biomedical domain include MetaMap (Aronson & Lang, 2010), cTAKES, and the NCBO Annotator:

 MetaMap (Demner-Fushman et al., 2017): Developed by the National Library of Medicine (NLM), MetaMap maps concepts extracted from biomedical and clinical text to the Unified Medical Language System

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(UMLS) (Bodenreider, 2004).

- cTAKES: The Clinical Text Analysis and Knowledge Extraction System combines rule-based and machine learning techniques to extract and normalize clinical concepts, also for integration into UMLS (Savova & others, 2010).
- 3) NCBO Annotator: This tool provides broader functionality by mapping concepts from text to any ontology hosted on BioPortal (Jonquet et al., 2009)(Musen et al., 2008). Its back-end Linux application, Mgrep, efficiently matches text with billions of ontology terms.

Ontologies, such as SNOMED CT (Donnelly, 2006), play a vital role in capturing meaningful knowledge and representing relationships between medical concepts (M & Chacko, 2020). Ontologies and reference terminologies are designed to represent conceptual knowledge in medicine and are not optimized for application purposes (Rosenbloom & others, 2006). For practical applications, interface terminologies -- tailored specifically for specific application -- are typically employed (Rosenbloom & others, 2006)(Kanter et al., 2008)(Rosenbloom & others, 2008).

In our prior research (V. Keloth et al., 2020)(V. K. Keloth et al., 2023)(Dehkordi & others, 2023), we developed two interface terminologies for COVID-19 and cardiology, leveraging the NCBO Annotator during the iterative process of mining fine-grained concepts. While NCBO Annotator performed well in identifying seed phrases, they observed a key limitation: it struggled to accurately annotate the longest relevant text chunks. This occurred because the tool indiscriminately identified all potential concepts in the EHR text, leaving overlapping matches unresolved.

To address this issue, we propose a Cluster-Focused Combination (CFC) Annotation algorithm. This algorithm is designed to annotate EHR text using interface terminologies while maximizing annotation coverage. Optimized CFC incorporates a strategy storing intermediate results to address limitations in conventional annotation techniques, specifically those relying on the BioPortal (mgrep) terminologybased annotator (Noy & others, 2009). Traditional methods often fall short in maximizing coverage and can miss important concepts due to sequential matching constraints. In contrast, our approach optimizes coverage by dynamically clustering relevant terms and efficiently matching them to text, minimizing the risk of annotation loss while maintaining reference to interface terminologies.

The annotation process is divided into two major steps:

- Phrase Matching: Mgrep (Dai, 2021) is used to generate a list of all matched phrases in the text based on a user-provided interface terminology. For this study, we applied the Cardiology Interface Terminology (CIT) (V. Keloth et al., 2020)(Dehkordi & others, 2023).
- 2) **Cluster-Focused Annotation:** Using the matched phrases list, the CFC annotation algorithm identifies an optimal combination of concepts for each sentence to ensure high-coverage annotations in the EHR text.

Our approach is generic and can be applied to any user-generated terminology, enabling robust and comprehensive annotation of EHRs for various applications with the optimal execution time.

2 BACKGROUNDS

2.1 MIMIC-III Dataset

MIMIC-III, short for Medical Information Mart for Intensive Care version 3, is an open-sourced, deidentified database that captures critical care data from patients admitted to the intensive care units at Beth Israel Deaconess Medical Canter in Boston, Massachusetts (Saeed et al., 2002). This extensive dataset includes a wide array of information such as vital signs, medications, laboratory results, procedural and diagnostic codes, as well as billing details.

2.2 Clinical Interface Terminology

Rosenbloom et al. (Rosenbloom & others, 2006)(Kanter et al., 2008)(Rosenbloom & others, 2008) describe that clinical interface terminologies aid healthcare practitioners by enabling rapid retrieval of patient information through a structured set of healthcare-related terms. These terminologies are specifically tailored for end-users, incorporating commonly used clinical phrases and colloquial expressions, in contrast to reference terminologies that aggregate clinical data based on standardized concepts.

In previous work (V. K. Keloth et al., 2023)(V. Keloth et al., 2020)(Dehkordi & others, 2023), we developed two pipelines for iteratively generating fine-granular clinical phrases from the Mimic-III cardiology notes and the Radiopeadia scan descriptions by using concatenation and anchoring

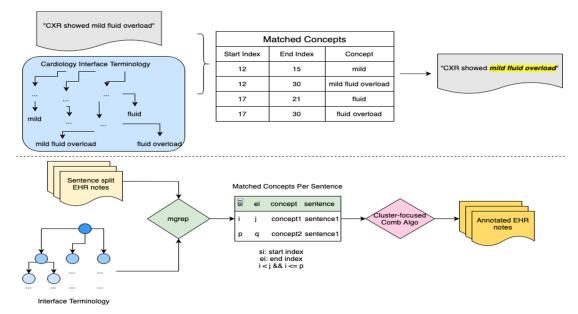


Figure 1: The entire workflow of the annotation pipeline.

operations. The generated concepts from these processes were used to construct two distinct interface terminologies: one for cardiology(V. Keloth et al., 2020)(Dehkordi & others, 2023) and one for Covid-19(V. K. Keloth et al., 2023).

2.3 NCBO BioPortal

The NCBO BioPortal is an open-source platform maintaining a continually growing collection of 1,158 biomedical ontologies. These ontologies are available in four different formats, including: 813 are in OWL (Web Ontology Language) format (Dean et al., 2004), 106 in OBO (Open Biomedical Ontologies) format, 30 in the UMLS's RRF format, and 122 in SKOS (Simple Knowledge Organization System) format (Miles & Bechhofer, 2009). BioPortal covers a broad range of topics ranging from Biological Process, to chemistry (Musen et al., 2008)(V. Keloth et al., 2020).

The BioPortal also provides varied of services for ontology users and allows them to upload their own developed terminologies. Once successfully submitting the terminology, the BioPortal generates and displays additional statistics about the uploaded terminology, including the number of classes and properties.

2.4 *Mgrep* Text Matching Tool

Mgrep is a command line tool that enables users to search through text files for lines that match a

specified regular expression (Dai, 2021). It is used as the back-end algorithm for BioPortal Annotator function. It functions similarly to the well-known Unix tool, grep, but extends its capabilities by supporting searches for patterns spanning multiple lines. While *mgrep* is not designed to replace grep, it is built to be compatible with it, meaning its options and behaviours closely align with those of grep. In our algorithm, we use *mgrep* to identify all relevant concepts from CIT that match phrases in the text, which are then used as candidates for our clusterfocused combination algorithm to annotate the EHR text.

3 METHOD

A user constructed interface terminology contains plenty of phrases from the user provided text dataset: some of them are high granular phrases, and some of them are relatively short in the length. Annotating the text with this set of phrases from the interface terminology with our method can be illustrated in the Figure 1.

Initially, we use *mgrep* to align text segments with the Cardiology Interface Terminology (CIT) (V. Keloth et al., 2020). *Mgrep* generates all matched phrases including cases of overlapping concepts of CIT.

Within the sentence, these matched phrases can be classified into 5 different conditions of positioning which will be discussed in section 3.1. Subsequently, we apply a cluster-focused algorithm to determine the optimal combination of concepts in each sentence and highlight each concept in the text.

3.1 Five Conditions

Consider a sentence S of n characters S [0, n-1], which is annotated by a terminology T. Suppose that m occurrences of concepts of T were identified in the sentence S. Let $C_1, C_2, ..., C_m$ be the concepts identified in S and stored in Dictionary D. Let L_i and Ri be the respective starting index and ending index of C_i in S, where $L_i \leq R_i$ for i = 1, m. The concepts C_i in S are ordered by their Li: $(L_1 \leq L_2 ... \leq L_m)$. Example 1: Sentence S: "*CXR showed mild fluid overload*."

	L(start)	R(end)	Concept
C1	12	15	mild
C ₂	12	30	mild fluid overload
C ₃	17	21	fluid
C ₄	17	30	fluid overload

- 1. Two concepts C_j and C_k (j < k) in S are called disjoint, if $R_j + 1 < L_k$. (Plus 1 is necessary, since it must have a space between C_j and C_k), for examples:
 - a. Concept C_1 "mild" and C_4 "fluid overload" are disjoint.
 - b. Concept C₁ "mild" and C₃ "fluid" are disjoint
- Two concepts C_j and C_k (j < k) in S are called leftadjusted, if L_j = L_k and R_j < R_k. for example:
 - a. Concept C₃ "fluid" and C₄ "fluid overload" are left-adjusted
- 3. Two concepts C_j and C_k (j < k) in S are called right-adjusted, if $L_j < L_k$ and $R_j = R_k$. See examples:
 - a. Concept C₃ "mild fluid overload" and C₄ "fluid overload" are right adjusted
- Two concepts C_j and C_k (j < k) in S are called overlap, if L_j < L_k < R_j < R_k.
 - For example, supposing the two concepts are "mild fluid overload" and "fluid overload XXX" (XXX refers to a word), then "fluid overload" is the overlapping.
- Two concepts C_j and C_k (j<k) in S satisfy that C_j contains C_k, if L_j < L_k and R_j > R_k. See examples:
 - a. Concept C₂ "mild fluid overload" contains C₃ "fluid".

By annotation of S using a terminology T, we refer to a subset of Dictionary D, such that all C_i in

the subset are mutually disjoint. When we consider an option of annotation for sentence by the terminology T, we have two purposes. The first purpose is to find a set of concepts for annotating such that the total number of words in those concepts is maximized. This purpose will lead to a high value of the annotation coverage for a given text. The other purpose is capturing the semantics of the sentence as intended by its author. Towards this goal, the CIT includes high granularity concepts, which better capture the semantic in the sentence. These concepts are typically longer and contribute to higher breadth of the annotation. The ultimate purpose is to serve customers of the annotation which are trying to comprehend the content of the text from its annotated parts. For this purpose, our annotation is not only matching the text to the existing terminology and listing all the matched concepts which are possibly redundant or overlapped, but we also output the text with highlighted annotations in the format of html. It is more user friendly.

Consider the case that all K concepts of T found in S are mutually disjoint. In this case, all of them are selected for the annotation, and the coverage of the sentence is optimal. However, in most cases, we encounter pairs of concepts which are left-adjusted, right-adjusted, overlapping, or in containment relation.

3.2 Basic Cluster-Focused Combination (CFC) Algorithm

Considering the sentence: "The patient was taken on to the operating room for a redo mitral valve surgery with mitral valve replacement". The phrases of the sentences which are matched to the concepts in CIT are listed in Table 2. We observe that the concepts "patient" "operating room" and "redo" where isolated concepts which did not interact with any other concepts of the dictionary. Hence, those disjoint concepts of the dictionary are selected for the annotation because they will appear in any annotation of the sentence using D. The challenge for selecting concepts for the annotation appears when we have a set of concepts with overlap. We see the clusters for five concepts C_4 to C_8 and three concepts C_9 to C_{11} in Table 2. The challenge is how to pick a combination of concepts for such clusters which optimizes in coverage and breadth.

	L	R	# of words	Concept
C_1	5	11	1	patient
C2	33	46	2	operating room
C3	54	57	1	redo
C4	59	70	2	mitral valve
C5	59	78	3	mitral valve surgery
C6	66	70	1	valve
C7	66	78	2	valve surgery
C_8	71	78	1	surgery
C9	85	96	2	mitral valve
C10	92	96	1	valve
C ₁₁	85	107	3	mitral valve replacement

Table 2: Dictionary D associated to Example 1.

Note*: L refers to start index, and R refers to end index. The reason for not using S to represent start index is to distinguish with the abbreviation of Sentence S.

In Example 2: "*CXR showed mild fluid overload*", we have the matched concepts as in Table 3.

Table 3: Matched concepts in Example 1.

	L	R	Concept
C1	12	15	mild
C3	12	30	mild fluid overload
C4	17	21	fluid
C5	17	30	fluid overload

There are two combinations of concepts sharing the maximum annotated words: $COMB_1 = C_1 + C_5$ and $COMB_2 = C_3$.

Since COMB₁ has two concepts and COMB₂ has one concept, we select COMB₂ for annotation in S, because there is COMB₂ contains only one concept. When a tie between two combinations on the number of words, the one contains less concepts is preferable to increase the breadth. As a result, the sentence S is annotated as: "*CXR showed mild fluid overload*".

Before generating the COMB of the whole sentence, it is necessary to recap the definition of "cluster" in the algorithm. "Cluster" of a sentence. For this, we define a graph where each annotated concept is a node. Two nodes are connected by an edge if there is an overlap between the phrases of the concepts represented by the nodes. That is the concepts are leftadjusted, right-adjusted or overlapping. A maximal connected subgraph of the graph representing the sentence is called a "cluster". The purpose for finding clusters of a sentence is: when a sentence contains a cluster, a decision will be made on which concept(s) should be selected in the cluster for annotating the sentence. That is, which concept(s) should be collected and kept in the COMB set.

3.3 Optimized Cluster-Focused Combination (CFC) Algorithm

The algorithm for finding an annotation of a cluster of k concepts with maximum coverage and breadth scans the concepts of the cluster from left to right, considering for each concept the solutions with and without this concept. The complexity is $O(N^k)$. In practices, we observed that for only 150 notes, it takes more than half hour. Due to the issue of high timeconsuming, we optimize the algorithm by importing a memorization mechanism.

3.3.1 Workflow of Annotation Process by Using Optimized CFC

Like in the basic version of the Cluster-focused Combination Algorithm, concepts are first sorted by their start index. If two or more concepts share the same start index, they are then sorted by their end index in ascending order. A strategy is then applied to determine whether each matched concept should be selected or omitted from the solution. Based on the current concept selection, there are five possible scenarios:

- If the Current Concept is selected, and when the Next Concept overlaps with the Current Concept, then, keep the Current Concept and continue checking the subsequent concepts to find one that doesn't overlap.
- 2) If the Current Concept is selected, and when the Next Concept does NOT overlap with the Current Concept, then, select the Next Concept and add to the Set.
- 3) If the Current Concept is selected, when the Next Concept does NOT overlap with the Current Concept, do not select the Next Concept.
- 4) If the Current Concept is not selected, select the Next Concept and add to the Set.
- 5) If the Current Concept is not selected, do not select the Next Concept either.

In this project, we divide the problem into smaller subproblems by considering subsets of concepts starting from each concept. For each concept, we maintain two scores: the maximum number of words covered when (i) selecting the concept and (ii) omitting it. Finally, after calculating the scores for all concepts in reverse order, we determine the result by checking the score for the first concept, which represents the overall score for the complete solution.

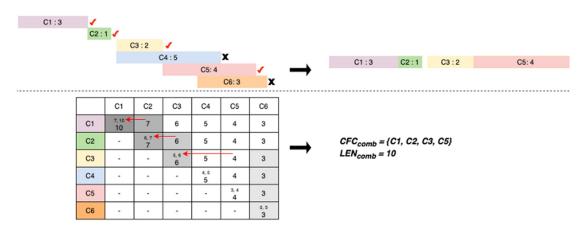


Figure 2: Example of the workflow by optimized CFC.

Figure 2 illustrates an example with six matched concepts within a single sentence. In Figure 2, this process is illustrated using a backwards dynamic selection approach. For each concept, we represent a pair as (a, b), where *a* and *b* denote the scores when omitting and selecting the concept, respectively. Starting with the last concept, C6, we assume only this concept is available. If C6 is selected, the Current Set becomes {{C6} Len:3}, with a length of 3 words; if C6 is omitted, the set is empty, represented as {{} Len:0} with a length of 0 words. Thus, for C6, we store (0, 3), where 0 and 3 are the scores for omitting and selecting C6, respectively.

Next, we consider C5, which overlaps with C6. If we add C5 to the Current Set, we cannot select C6, resulting in $\{\{C5\} Len:4\}$. By omitting C5, we retain the option to select C6, which produces the set $\{\{C6\} Len:3\}$. Since, for each concept, we store the maximum score between selecting and omitting it, we record (3, 4) for C5.

Moving to C4, which overlaps with both C5 and C6, we find that selecting C4 requires omitting both C5 and C6, resulting in a score of 5. By omitting C4, we retain the highest score starting from C5, which is 4. Thus, we store (5, 4) as the scores for C4, corresponding to the sets {{C4} Len:5} and {{C5} Len:4}, respectively.

Continuing this process, if we select C3, we cannot select C4, so we choose the best result from C5, forming the set { $\{C3, C5\}$ Len:6} (calculated as { $\{C3, C5\}$ Len:2+4}). C2 and C1 can be added without any conflict, as they do not overlap with other concepts, resulting in the set { $\{C1, C2, C3, C5\}$ Len:10}, calculated as { $\{C1, C2, C3, C5\}$ Len:3+1+6}.

Finally, the optimal score is achieved with C1, which represents the best possible combination of

concepts. Thus, the best concept combination for the annotation text is $\{C1, C2, C3, C5\}$ with L=10.

3.3.2 Pseudo Code of Optimized CFC

In Algorithm 1, we present the pseudo-code for the optimized (CFC) algorithm. This optimization merges two tasks: "finding the maximum number of words annotated" and "determining the best combination of concepts"--- into a single function: *find_decide_best_comb_dp*. This function is organized into five key sections that guide the decision-making process for selecting the best concept combination at each stage.

The first two sections address the initialization of the case base and previously traversed cases, laying the groundwork for recursive exploration. Section 3 handles the scenario where the current concept is "not" included in the solution. In this case, the algorithm simply copies the previous state of results to a list, not take curr concept, for storage and continues the recursion without modifying the solution. If we decide to include the current concept in the combination, Section 4 initiates two primary operations. First, it locates a a next qualified concept that does not overlap with the current concept; this non-overlapping condition ensures that we maintain a conflict-free solution as we proceed. Next, the current concept's index is added to next decision concept list, which accumulates the indices of concepts chosen in the optimal combination so far. At this point, we update the *take curr concept* list by:

1. Updating the "Maximum Number of Words Annotated" by adding the word count of the current concept.

Data:	list of matched concepts,
	number of concepts,
	current index of concept
	t: the optimal combination of concepts for annotating the sentence
1:	Function: find_decide_best_comb_dp (in: dict D, num_concept N, curr_index i, out: curr_comb) // *note: dict D is a list of tuples, not python dictionary, but function as a dictionary
2:	/*
	Section 1: Base case
	*/
3:	if <i>i</i> reaches to the N
4:	return {0, 0, []}
5:	/*
	Section 2: Already considered case */
6:	if Concept C at i-th index of curr_comb is not None
7:	return curr_comb[i]
8:	/*
	Section 3: Don't take the current concept */
9:	next_decisions = find_decide_best_comb_dp(D, N, i+1, curr_comb)
10:	assign next_decisions[0]: maximum number of words annotated,
	next_decisions[1]:minimum number of concepts used,
	next_decisions[2] curr best concept list to not take curr concept list
11:	/*
	Section 4: Take the current concept
12:	assign next_concept_index j with the index of $i + 1$
13:	while j is not reach to N and start index s of D[j]is less or equal than end index e of D[i]
14:	update j by increasing 1
15:	next decisions = find decide best comb dp(D, N, j, curr comb)
16:	append index i into next decisions concepts list
17:	if next_decisions[2]: curr best concept list is Empty
18:	next_decisions_concepts_list extend next_decisions[2]
19:	take_curr_concept = D[i][2] + next_decisions[0], 1 + next_decisions[1], next_decisions_concepts_list
20:	/*
	Section 5: Decide the best combination by comparing the take_curr_concept and not_take_curr_concept */
21:	if maximum number of words annotated of take_curr_concept greater than not_take_curr_concept
	or if maximum number of words annotated are the same , and minimum number of concepts used of take_curr_concept smaller than not_take_curr_concept
22:	add take_curr_concept to curr_comb[i]
23:	else
24:	<pre>add not_take_curr_concept to curr_comb[i]</pre>
25:	return curr_comb[i]

Algorithm 1: Find the optimal combination of concepts for annotating the sentence.

- 2. Increasing the "Minimum Number of Concepts" by one, as we add the current concept to the optimal set.
- 3. Updating the best current concepts list to reflect the newly added concept.

With *not_take_curr_concept* and *take_curr_concept* lists now fully populated, we have enough information to decide whether to include the current concept.

Finally, Section 5 performs a comparison between not take_curr_concept and take_curr_concept lists

based on two prioritized criteria. First, it evaluates the "number of words annotated" -- the decision that annotates the maximum words is preferred. If there is a tie in this criterion, it then considers the "number of concepts", preferring the option that includes fewer concepts for efficiency. The "winning" combination at this stage is stored in *curr_comb*, representing the best possible combination up to the current point in recursion. at this stage is stored in *curr_comb*, representing the best possible combination up to the current point in recursion.

4 **RESULTS**

We tested our algorithm on a corpus of de-identified Cardiology Information notes from the Mimic-III dataset. The objective is to annotate relevant terms within the notes by referencing the CIT (V. Keloth et al., 2020)(Dehkordi & others, 2023). For the evaluation, we employed two metrics: coverage and breadth. Coverage measures the percentage of concepts from the CIT captured within the EHR notes, reflecting the extent to which the interface terminology comprehensively annotates the text. Breadth, on the other hand, indicates the average number of words per annotated concept, representing the specificity or granularity of concepts within the CIT. High coverage implies extensive text annotation by the CIT, signifying its thoroughness, while breadth highlights the conceptual detail included in the terminology. The equations are as follows:

$$Coverage = \frac{\sum number of words in annotated concept C}{\sum number of words in each note} * 100$$

$Breadth = \frac{Number of words in all annotated concepts}{Number of annotated concepts}$

We selected the BioPortal Annotator and a baseline Cluster-focused-Combination annotation algorithm for comparison. Given that our annotation approach relies on precise matching with the concepts within a specialized domain-specific interface terminology using large language models is not suitable for this study. This focus on domain-specific, deterministic concept matching ensures that the annotations remain consistent with the defined terminological scope of CIT, which is essential for accurate representation and evaluation in this context.

4.1 Coverage and Breadth

The experiments were conducted on two datasets: a small-scale text dataset annotated with an earlier version of the Cardiology Interface Terminology (CIT) and a large-scale dataset annotated with the final version. For the smaller dataset, we randomly selected 150 cases from MIMIC-III Event Notes and annotated them using CIT_V_{3.2}. For the larger dataset, comprising 500 notes, we applied the final version, CIT_V_{5.2}, to evaluate the capacity of the annotators in handling a larger volume of data.

These phrases are constructed by concatenating and anchoring CIT_V_{3.2} concepts. Consequently, annotating the 500-note dataset with CIT_V_{5.2} introduces a higher number of potential matched phrases and more complex overlapping cases when identifying concepts for annotating in the text. This second dataset, therefore, presents a greater challenge for the annotation process, testing the robustness and accuracy of the annotators under more intricate conditions.

Table 4: The Comparison of # of Annotated Concepts, Coverage and Breadth among Two Annotators on 150 Notes.

	# of Annotated Concepts	# Annotated Words	Coverage	Breadth
BioPortal (mgrep)	8518	17718	44.85%	2.08
Basic/Optimized CFC	9022	18338	46.42%	2.03

Table 4 shows that 504 concepts were missed by the BioPortal (*mgrep*) annotator but were successfully identified by the Cluster-Focused Combination (CFC) algorithm. Although the dataset is relatively small, the increase in coverage between BioPortal (*mgrep*) and the CFC algorithm is modest. The slightly lower breadth in CFC annotations reflects two key points:

- 1) The generated interface terminology includes highly granular concepts.
- The BioPortal annotator prioritizes the longest matching concept per sentence, often omitting shorter, overlapping concepts. This approach sacrifices some information by reducing the text coverage.

This second limitation motivated the development of the CFC algorithm, designed to address these gaps by capturing both long and short overlapping concepts. Additionally, the optimized CFC algorithm shares core functionalities with the basic CFC version, resulting in identical outcomes for the number of annotations, coverage, and breadth.

Table 5: The Comparison of # of Annotated Concepts, Coverage and Breadth among Two Annotators on 500 Notes.

	Annotated Concepts	Annotated Words	Coverage	Breadth
BioPortal (mgrep)	50087	99977	69.13%	2.02
Optimized CFC	55843	103287	71.42%	1.85

Table 5 compares the performance of two annotation methods on the larger 500-note dataset. Here, the CFC algorithm captures 5,756 concepts missed by BioPortal, a significant increase compared to the 504 missed concepts in the 150-note dataset. While the dataset size grew by 3.3 times, the number of missed annotations increased approximately 11 time, highlighting the annotation or information loss issue faced by BioPortal.

For the larger dataset, we did not include

experiments with the basic CFC algorithm due to its high time complexity, O (2^k) per sentence, where N=2 represents the number of matched candidate concepts, and *k* is the size of the concept combination. Detailed runtime analysis can be found in section 4.2.

4.2 Runtime Analysis

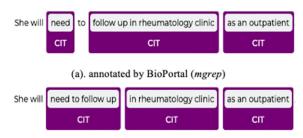
The optimized CFC algorithm significantly enhances the time efficiency of the basic CFC algorithm by employing memorization techniques. Instead of processing every possible combination of concepts, the optimized CFC algorithm evaluates scores for all subsequence in the sorted list of concepts. Therefore, the theoretical time complexity is improved from $O(2^k)$ to $O(k^2)$. As shown in Table 3, the runtime of the basic CFC on the 150-note dataset is 2,952 seconds and for BioPortal with 500-notes which took 5564 seconds, meaning that is not efficient for processing larger datasets. In contrast, the optimized CFC drastically reduces execution time from hours to milliseconds. Annotating the 500-note dataset with the optimized CFC takes less than one second, demonstrating its scalability and feasibility for processing large biomedical datasets.

Table 6: The Comparison of Execution Runtime (seconds)among Three Annotators on Two Datasets.

	150-dataset	500-dataset
BioPortal (mgrep)	169.68	5564.46
Basic CFC	2952.66	
Optimized CFC	0.02	0.71

4.3 Case Study

Figure 6 presents an example for sentence "She will need to follow up in rheumatology clinic as an outpatient.", annotated using both the BioPortal (mgrep) and CFC algorithms. In Figure 3(a), the BioPortal (mgrep) identifies and annotates the longest matched phrase, "follow up in rheumatology clinic". The remaining two segments of the sentence are non-overlapping annotated with concepts, specifically "need" and "as outpatient". an Altogether, this approach annotates 9 words, covering 75% of the sentence. By contrast, the CFC algorithm identifies three additional concepts within the sentence: "need to follow up", "in rheumatology clinic", and "as an outpatient". Although CFC does not select the longest matched concept, "follow up in rheumatology clinic", it annotates a total of 10 words -- one more than BioPortal (mgrep).



(b). annotated by optimized CFC algorithm

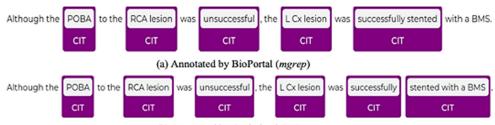
Figure 3: Annotation for example one by BioPortal (*mgrep*) and optimized CFC annotator.

In another instance, important information is missed by the BioPortal (*mgrep*) annotator. Figure 4 illustrates a comparison between BioPortal (*mgrep*) and CFC annotations for Example Two, the sentence: "Although the POBA to the RCA lesion was unsuccessful, the L Cx lesion was successfully stented with a BMS." In Figure 4, the BioPortal (*mgrep*) misses the important information "BMS" in its annotation, as it selects the concept "successfully stented" over the shorter "successfully". This choice results in the omission of the important detail "BMS". In contrast, the CFC algorithm captures all relevant information, providing a more comprehensive and accurate annotation.

A similar issue arises in Figure 5 with the sentence: "Had asymptomatic run of NSVT with stable vital signs." In Figure 5(a), the BioPortal fails to identify the term "NSVT". Instead, it selects "asymptomatic run" as the annotated concept due to its longer phrase length compared to "asymptomatic". However, "NSVT" appears within another concept, "run of NSVT", which overlaps with "asymptomatic run". This overlap leads BioPortal to skip annotating "run of NSVT", resulting in the omission of "NSVT" from the annotation.

5 DISCUSSIONS

Our Cluster-Focused Combination (CFC) algorithm annotates electronic health record (EHR) texts by leveraging the interface terminology with the highest coverage, while maintaining optimal time complexity. In a comprehensive evaluation, we tested the CFC algorithm on two datasets of varying sizes, randomly selected from the MIMIC-III database, using two versions of Cardiology Interface Terminology (CIT). Compared to the traditional



(b) Annotated by Optimized CFC

Figure 4: Annotation for Example Two by the BioPortal (mgrep) and optimized CFC annotator.

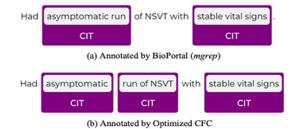


Figure 5: Annotation for Example Three by the BioPortal (*mgrep*) and optimized CFC annotator.

BioPortal (*mgrep*) longest-chunk annotator, our algorithm significantly improves the number of annotated concepts, the number of annotated words, and overall annotation coverage. Furthermore, the CFC algorithm offers a substantial reduction in execution time, making it well-suited for large-scale datasets and enhancing its scalability and efficiency in high-volume clinical settings. In Figure 6, we present the chart of the execution time on different number of matched concepts per sentence. We randomly re-generate 150-notes, 500-notes and 1000-notes for the execution time study.

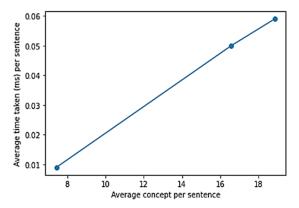


Figure 6: The execution time on average number of matched concepts per sentence of three datasets: 150-notes, 500-notes and 1000-notes.

The average number of matched concepts per sentence for three datasets are: 7.4, 18.9, 16.7

respectively. It is observed that the execution time is increasing in a linear-like curve. For a sentence with 18 matched concepts, it only takes about 0.06s by our optimized CFC algorithm. In addition, the tested 1000-notes dataset is processed only in 1.87s which again demonstrate the power of the algorithm on annotating enormous scale of dataset.

Despite these advances, the CFC annotator has certain limitations. It cannot recognize unstructured phrases that lack direct mappings to the semantics in the provided terminology. This reliance on the quality of the interface terminology means that if the terminology lacks high granularity concepts or fails to capture specific information, the CFC algorithm may miss annotating relevant phrases. Therefore, the success of CFC is closely tied to the comprehensiveness and detail of the terminology provided. As such, careful curation and frequent updates to the interface terminology are essential for ensuring that annotations remain accurate and exhaustive in real-world applications.

This performance improvement indicates great potential for automated annotation on a massive scale, alleviating the need for labour-intensive manual annotation. The optimized CFC can thereby facilitate the creation of high-quality, large-scale annotated datasets, supporting future training efforts for Large Language Models and other data-intensive AI applications.

The optimized CFC annotator's flexibility also makes it adaptable for use with other types of terminologies or ontologies beyond standard interface terminology. Users can supply their own terminology and datasets, allowing for customizable annotations across a wide array of domains. However, it is important to note that the broader and more detailed the terminology, the higher the expected annotation performance. In future work, we aim to develop related software based on the optimized CFC algorithm to provide more interactive annotation services and plan to test it on larger-scale datasets across diverse domains to further assess its versatility and robustness.

6 CONCLUSIONS

In conclusion, our research introduces a novel algorithm, the cluster-focused combination algorithm, designed to overcome the challenges associated with annotating Electronic Health Record (EHR) text using interface terminology. This algorithm addresses critical issues in text annotation by utilizing a dynamic programming approach, effectively balancing the need for high annotation coverage and breadth while mitigating common pitfalls of previous methods. Our extensive evaluation on benchmark datasets, such as Mimic III, reveals an improvement in annotation coverage and captured 5756 missed annotated concepts by the traditional BioPortal Annotator. Additionally, cluster-focused combination the algorithm demonstrates a notable reduction in execution time by an average of about 8000 times, enhancing its scalability for large datasets.

These findings make the optimized CFC a highly effective tool for real-world text annotation tasks that rely on interface terminology. By providing a more efficient and comprehensive solution, this work not only advances the capabilities in EHR text annotation but also contributes to the broader field of Natural Language Processing. This is particularly significant for the development of Large Language Models, which depend on vast, well-annotated datasets. Our algorithm paves the way for future innovations in dataset preparation, promising to streamline and accelerate the annotation process for large-scale NLP applications.

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