

The Computational Linguistics Summarization Pilot Task

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Abstract

The Computational Linguistics (CL) Summarization Pilot Task was created to encourage a community effort to address the research problem of summarizing research articles as “faceted summaries” in the domain of computational linguistics. In this pilot stage, a hand-annotated set of citing papers was provided for ten reference papers to help in automating the citation span and discourse facet identification problems. This paper details the corpus construction efforts by the organizers and the participating teams, who also participated in the task-based evaluation. The annotated development corpus used for this pilot task is publicly available at:

<https://github.com/WING-NUS/scisumm-corpus>

1 Introduction

The Computational Linguistics Summarization Pilot task provides resources aimed to encourage research on scientific document summarization. Specifically, the task considers summarization utilizing the set

of citation sentences (i.e., “citances”) that cite a reference article as a (community created) summary of a topic or paper (Nanba et al., 2011; Qazvinian and Radev, 2010). Citances for a reference paper are considered a synopsis of its key points and also its key contributions and importance within an academic community. An advantage of citances is that they embed meta-commentary and offer a contextual, interpretative layer to the cited text. A drawback of this approach is that citances usually do not consider the context of the target user (Jones, 2007; Teufel and Moens, 2002), verify the claim of the citation, nor provide context from the reference paper (in terms of the kind of information cited, or where it is in the referenced paper).

Existing scientific article summarization systems have approached the task by automatically generating related work sections for a target paper via a hierarchical topic tree (Hoang and Kan, 2010), generating model citation sentences (Mohammad et al., 2009) or implementing a literature review framework (Jaidka et al., 2013a). However, limited evaluation resources – i.e., human-created summaries – means that the efficacy of these approaches cannot be verified by others, hurting the replicability of works in this domain. The goals of the Computa-

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tional Linguistics (CL) shared task was to highlight the challenges and relevance of the scientific summarization, and provide evaluation resources for advancing the state-of-the-art.

The CL pilot task was constructed by sampling papers from the Association of Computational Linguistics’ (ACL) anthology (Bird et al., 2008). This task was run concurrently with the Text Analysis Conference 2014 (TAC ’14), with the approval and guidance by the TAC organizers, although not formally affiliated with it. It shares the same basic structure and guidelines as the formal TAC 2014 Biomedical Summarization (BiomedSumm) track. We released a training corpus of “topics” from CL research papers, each comprising a reference paper along with sample papers that cited the reference paper. Participants were then asked to participate in a task-based evaluation and include a system description and self-reported results as part of this report.

An ideal summary of a CL research paper would distill its overall contribution in the state of the art through a discussion of its goals, methods and results. Previous work (Mohammad et al., 2009; Abu-Jbara and Radev, 2011) in scientific document summarization have used citances from citing papers (hereafter, *CPs*) to create a multidocument summary of a reference paper (hereafter, *RP*). Their approach followed a three-part process: finding the relevant documents (*CPs*), then selecting sentences which justify the citation in the *RP*, and finally, generating the summary. In this task, we follow this method and have created a training corpus comprising human annotations for each of these sub-problems. Human annotators identified the citances in each of (up to) ten randomly sampled *CPs* for the *RP*.

Previous work also indicated that most citations clearly refer to one or more specific discourse facets of the *CP* (Jaidka et al., 2013b). Discourse facets indicate the type of information described in the reference span: e.g., “Aim” indicates that the citation concerns the aims of the reference paper. From our exploration of the computational linguistics domain, we observed that the discourse facets being cited were usually the aim of the paper, its methods and the results or implications of the work. We also applied these observations in annotating discourse facets in our training corpus.

2 Corpus Construction

A large and important body of scholarly communication in the domain of computational linguistics is publicly accessible and archived at the ACL Anthology¹. The texts from this archive are also under a Creative Commons license, which allows unfettered access to the works for any purpose, including downstream research on summarization.

As of the corpus construction date (18 September 2014), the live Anthology contained approximately 25K publications, exclusive of the third-party papers hosted (i.e., with metadata but without the actual PDF version of the paper) and extraneous files (i.e., front matter and full volumes). We randomly sampled research papers for use as *RPs* using the following procedure:

- We considered only papers published after and including 2006, leaving 13.8K publications. We randomized this list to remove any ordering effects;
- Starting from the top of the list, we used a combination of Google Web and Google Scholar searches to approximate the number of *CPs* for the *RP*. We retained any paper as an *RP* if it was reported to have over 10 citations;
- We vetted the citations to ensure that the citation spread was at least a window of three years, as previous work indicates that citations over different time periods (with respect to the publication date of the *RP*) exhibit different tendencies (Abu-Jbara et al., 2013);
- We then used the title search facility of the ACL Anthology Network² (AAN, February 2013 version), to locate the paper and, inspected all citing papers’ Anthology ID, title and year of publication. The citation count from Google / Google Scholar and AAN often differed substantially;

For every *RP*, we aimed to provide at least three *CPs* based on the following criteria (in order of priority):

1. Non-list citation (i.e., at least one citation in the body of the *CP* for the *RP* not of the form [*RP*,*a*,*b*,*c*]);

¹<http://aclweb.org/anthology/>

²<http://clair.eecs.umich.edu/aan/index.php>

2. The oldest and newest citations within AAN, and;
3. Citations from different years.

We included the oldest and newest citation regardless of criteria 1) and 3), and included a randomized sample of up to 8 additional citing paper IDs that met either criterion 1) and 3). The final list was divided among the annotator group, who are a subset of the authors of this paper, from the National University of Singapore and Nanyang Technological University, Singapore. Annotators re-used the resources created for BiomedSumm, which reduced the efforts required; however, a different set of discourse facets were used to best represent the content of computational linguistics research papers. The resultant corpus should be viewed as a development corpus only, such that later efforts by the community can enlarge it to a proper shared task with training, development and testing set divisions.

2.1 Corpus Preprocessing

The original source text for the papers in the CL-Summ corpus was not sentence-segmented, which made it difficult to compute evaluation metrics. Both the `clair_umich` system and the `TALN.UPF` system performed significant corpus pre-processing to create a sanitized, annotated dataset for their systems.

In the `clair_umich` system, for each RP, citing sentences were extracted from all its CPs. Each CP sentence was matched to the RP to create the final annotated dataset. Given a citing sentence, matching sentences from the RP were compared to the gold standard RP sentences to compute precision / recall. On an average, each CP sentence matched 1.28 RP sentences. The maximum number of matches was 7.

The UPF system performed the following sanitization process to overcome corpus encoding issues:

1. Automatic PDF-to-text conversion: Conversion of PDF versions of the paper into text, by means of Poppler³, a robust PDF-to-text converter;
2. Manual verification of output: Manual validation of the PDF-to-text conversion errors in order to get a clean textual version of each paper;

3. Sentence splitter and Sentence Sanitizer: Use of a rule-based sentence splitter and sanitizer to identify candidate sentences, and to remove incorrectly annotated sentences;
4. Mapping annotations to clean textual versions: Inspection of the textual contents of each of the annotation files, and manual mapping of the annotations to the clean textual version of each paper.

These resulted in two sanitized versions of the initial corpus that was shared as a part of this task. Both versions are shared, along with the original, in the official repository of the CL Corpus with the consent of the participants.

3 The CL-Summ Task

This shared task proposes to solve the same problems posed in the Biomedical Summarization Track of the 2014 Text Analysis Conference 2014⁴, but in the domain of Computational Linguistics. It poses the research problem of building a structured summary of a research paper – which incorporates facet information (such as Aims, Methods, Results and Implications) from the text of the paper, and “community summaries” from its citing papers.

We define the *CL-Summ Task* as follows:

Given: A topic, comprising of the PDF and extracted text of an reference paper (RP) and up to 10 citing papers (CPs). In each provided CP, the citations to the RP (or citances) have been identified and manually annotated. The information referenced in the RP is also annotated.

Output: Systems are required to perform the following tasks, where the numbering of the task corresponds to those used in the BiomedSumm task.

- Task 1A: Identify the text span in the RP which corresponds to the citances from the CP. These may be of the granularity of a full sentence or several sentences (up to five sentences), and may be contiguous or not. It may also be a sentence fragment.
- Task 1B: Identify the discourse facet for every cited text span from a predefined set of facets.

³<http://poppler.freedesktop.org/>

⁴<http://www.nist.gov/tac/2014/BiomedSumm/>

Discourse facets categorize the type of information described in the reference span. A maximum of three reference spans can be marked for every citance. In case these spans describe different different discourse facets, the most prevalent discourse facet is annotated.

Evaluation: Evaluate Task 1A performance by using the ROUGE (Lin, 2004) score to compare the overlap of text spans in the system output versus the gold standard created by human annotators.

An additional task in BioMedSumm, which was not advertised with this shared task, was:

Task 2: Generate a faceted summary of up to 250 words, of the reference paper, using itself and the citing papers.

Nine teams expressed an interest in participating in the shared task, and three eventually submitted runs, system descriptions and self-assessed results. These three systems — **clair_umich** from University of Michigan; Ann Arbor, USA, **MQ**, from Macquarie University, Australia; and **Taln.UPF**, from Universitat Pompeu Fabra, Spain — are described in the following sections.

4 The clair_umich System — Comparing Overlap of Word Synsets

4.1 Baseline System

The team first created a baseline system based on the basic information retrieval measure: *term frequency* \times *inverse document frequency* (TF.IDF) cosine similarity. Note that for any citing sentence, the system computed the TF.IDF cosine similarity with all the sentences in the RP, thus the IDF values differed across each of the 10 RPs.

4.2 Supervised System

The supervised system used knowledge-based features derived from WordNet, syntactic dependency based features, and distributional features in addition to the simple lexical features like cosine similarity. These features are described below.

1. **Lexical Features:** Two lexical features were used – TF.IDF and the LCS (Longest Common Subsequence) between the citing sentence (C) and reference sentence S , which is computed as:

$$\frac{|LCS|}{\min(|C|, |S|)}$$

2. **Knowledge Based Features:** Six WordNet-based similarity measures were combined to obtain six sentence similarity features (Banea et al., 2012): path similarity, WUP similarity (Wu and Palmer, 1994), LCH similarity (Leacock and Chodorow, 1998), Resnik similarity (Resnik, 1995), Jiang-Conrath similarity (Jiang and Conrath, 1997), and Lin similarity (Lin, 1998). Using these measures, they computed the similarities between the citing and reference sentences by creating a set of senses for each of the words in each sentence:

$$sim_{wn}(C, R) = \frac{(\omega + \sum_{i=1}^{|\phi|} \phi_i) * (2|C||R|)}{|C| + |R|}$$

Here ω is the number of shared senses between C and R . The list ϕ contains the similarities of non-shared words in the shorter text, ϕ_i is the highest similarity score of the i^{th} word among all the words of the lower text (Zhu and Lan, 2013).

3. **Syntactic Features:** Given a candidate sentence pair, two syntactic dependencies were considered equal if they had the same dependency type, governing lemma, and dependent lemma (Zhu and Lan, 2013). The Stanford parser was used to obtain dependency parses of all the citing sentences and reference sentences. Then, if R_c and R_r are the set of all dependency relations in C and R , the dependency overlap score was computed using the formula:

$$sim_{dep}(C, R) = \frac{2 * |R_c \cap R_r| * |R_c||R_r|}{|R_c| + |R_r|}$$

5 The MQ System — Finding the Best Fit to a Citance

Given the text of a citance, the MQ system ranked the sentences of the reference paper according to its similarity to the citance. Every sentence and its citance was modeled as a vector and compared using cosine similarity.

Baseline – Using TF.IDF For the baseline system, the TF.IDF of all lowercased words was

$$\text{MMR} = \arg \max_{D_i \in R \setminus S} \left[\lambda(\text{sim}(D_i, Q)) - (1 - \lambda) \max_{D_j \in S} \text{sim}(D_i, D_j) \right]$$

Where:

- Q is the citance text.
- R is the set of sentences in the document.
- S is the set of sentences that haven been chosen in the summary so far.

Figure 1: Maximal Marginal Relevance (MMR) algorithm, as used in the MQ system.

used, without removing stop words (similar to the `clair_umich` team). Separate TF.IDF statistics were computed for each reference paper, using the set of sentences in RP and the citance text of all citing papers (CPs).

Adding texts of the same topic: Since the amount of text used to compute the TF.IDF was relatively little, it was presumed that citing papers are of the same topic. Accordingly the complete text of all citing papers was added in calculations for the IDF component.

Adding context: To extend the information on each sentence in the RP, the text from the RPs was added within a context window of 20 neighboring sentences to the target sentence from a CP.

MMR Re-ranking: The last experiment used Maximal Marginal Relevance (MMR) (Carbonell and Goldstein, 1998) to rank the sentences. All sentences were represented as TF.IDF vectors of extended information as described in previous paragraph. The final score of a sentence was the combination of the similarity with the citance, and similarity with the other sentences of the summary, according to the formula shown in Figure 1. A value of $\lambda = 0.97$ was chosen.

For all experiments, the systems were designed to return 3 sentences, as specified in the shared task. All short sentences (under 50 characters) were ignored, to avoid including headings or mistakes made by the sentence segmentation algorithm.

6 The Taln.UPF System

In the TALN.UPF system, the following text analysis tools were first used to pre-process the sanitized CL-Summ corpus:

1. **Tokenizer and POS tagger:** The GATE⁵ tool and its ANNIE (A Nearly-New Information Extraction system) NLP tools for English were used to tokenize and tag the corpus.

2. **Sentence TF.IDF vector calculator:** A TF.IDF vector was generated for each sentence. The IDF values of the terms of each document were computed by considering the CPs and RP as a complete corpus.

6.1 Task 1A: Identifying RP text spans for each citance

The TALN.UPF system implemented an algorithm to map every citation in the CP to one or more (up to three) *reference text spans* from the RP. Sentences from the CP that overlapped partially or totally with the citation text were selected and referred to as the *citation context* (CtxSent1,..., CtxSentN). For every sentence in the RP, the system associated a *score* equal to the sum of the TF.IDF vector cosine similarities computed between that sentence and each sentence belonging to the citation context (CtxSent1,..., CtxSentN). Finally, the top n sentences from the RP with the highest score were chosen as the reference span. Conflicts in choice were resolved by preferring sentences that occurred in the same document section in the RP. Furthermore, scores for referencing sentences were weighted by the prominence of document sections selected reference spans; for instance, if there 6.5% of all reference spans were sourced from the Abstract section, then the score for a sentence from the Abstract of the RP is multiplied by 0.065. The system evaluated the performance of the algorithm with varying values of n , the number of sentences included in the reference span.

⁵<https://gate.ac.uk/ie/annie.html>

6.2 Task 1B: Identifying the discourse facet of the cited text spans

Task 1B was cast as a sentence classification problem. From the corpus, the system selected sentences from the CPs that overlapped partially or totally with a manually annotated reference text span. These CP sentences were then classified with the discourse facet of the overlapping manually-annotated reference text span. This resulted in a set of 266 CP sentences as distributed in Table 1.

Docset	Citing papers
<i>Aim</i>	46
<i>Hypothesis</i>	1
<i>Implication</i>	25
<i>Results</i>	29
<i>Method</i>	165
TOTAL:	266

Table 1: Discourse facet of the sentences of cited papers belonging to a manually annotated reference text span.

Every sentence was modelled as a word vector of unigrams, bigrams, trigrams and lemmatized versions of all three. Sentence classification performance was compared across 3 classifiers – *Naive Bayes (NB)*, *SVM* using a linear kernel and *Logistic Regression (LR)*. Results from a 10-fold cross validation over the set of CP sentences listed in Table 1 are shown in Table 4. LR performed best with an averaged F_1 of 0.719.

7 Evaluation and Results

Results for Task 1: All three teams submitted their self-assessed results, using ROUGE (Lin, 2004) for Task 1A. ROUGE-L, which compares system output against a set of target summaries using the longest common subsequence of words, was used. Since ROUGE uses actual content words, and not offsets, we expect non-zero results when systems choose a sentence that is somewhat similar to (but not identical) to one chosen by annotators.

For Task 1A, the MQ and TALN.UPF systems were unsupervised, while clair_umich system was supervised. The former two systems were evaluated over all 10 topics in a single run, while clair_umich reported cross validated performance over the 10 topics. Table 2 shows the overall self-reported results, averaged over all topics, while Table 3 shows

	P	R	F_1
MQ	0.212	0.335	0.223
clair_umich	0.444	0.574	0.487
TALN.UPF	0.194	0.344	0.225

Table 2: Task 1A performance for the participating systems expressed as ROUGE-L score averaged over all topics.

Paper ID	MQ	clair_umich	TALN.UPF
C90-2039	0.235	0.635	0.180
C94-2154	0.288	0.536	0.200
E03-1020	0.239	0.478	0.198
H05-1115	0.350	0.375	0.233
H89-2014	0.332	0.546	0.275
J00-3003	0.196	0.559	0.263
J98-2005	0.101	0.344	0.196
N01-1011	0.221	0.498	0.254
P98-1081	0.200	0.367	0.211
X96-1048	0.248	0.535	0.240

Table 3: Task 1A ROUGE-L F_1 scores for individual topics.

micro-level results, giving the ROUGE-L F_1 scores of each individual reference document from the CL-Summ dataset.

For Task 1B, the TALN.UPF system also followed a supervised approach with 10-fold cross validation. Self-reported results are shown in Table 4. The ROUGE-L scores have been calculated using the system output of a set of selected sentences as the system summary, and comparing their overlap against the target summaries are the sentences given by the annotators.

Results for Task 2: The MQ team performed an additional test to see whether information from the citances were useful for building an extractive summary, as this was also the case with the Biomed-Summ task (Mollá et al., 2014). They implemented extractive summarization systems with and without citance information. The summarizers without information from the citances scored each sentence as the sum of the TF.IDF values of the sentence elements. They tried the TF.IDF approach described in Section 5.

The summarizers with information from the citances scored each candidate sentence i on the basis of $rank(i, c)$ obtained in Task 1A, which yields val-

Discourse facet	NB	SVM	LR
<i>Aim</i>	0.725	0.734	0.732
<i>Method</i>	0.706	0.826	0.828
<i>Implication</i>	0.049	0.000	0.200
<i>Results</i>	0.509	0.533	0.533
<i>Hypothesis</i>	0.024	0.000	0.000
WEIGHED AVG. F_1	0.623	0.698	0.719

Table 4: Task 1B self-evaluation for TALN.UPF: F_1 classification performance comparison.

ues between 0 (first sentence) and n (last sentence), and represents the rank of sentence i in citance c :

$$score(i) = \sum_{c \in citances} 1 - \frac{rank(i, c)}{n}$$

The summaries were again evaluated using ROUGE-L, where the model summaries are the abstracts of the corresponding papers. Since paper X96-1048 of the SciSumm data did not have an abstract, it was omitted from this experiment.

An example excerpt from a target summary (Abstract) for the reference paper J03-3003 is:

We describe a statistical approach for modeling dialogue acts in conversational speech, i.e., speech-act-like units such as STATEMENT, QUESTION, BACKCHANNEL, AGREEMENT, DISAGREEMENT, and APOLOGY. Our model detects and predicts dialogue acts based on lexical, collocational, and prosodic cues, as well as on the discourse coherence of the dialogue act sequence. The dialogue model is based on treating the discourse structure of a conversation as a hidden Markov model and the individual dialogue acts as observations emanating from the model states. Constraints on the likely sequence of dialogue acts are modeled via a dialogue act n-gram... We achieved good dialogue act labeling accuracy (65% based on errorful, automatically recognized words and prosody, and 71% based on word transcripts, compared to a chance baseline accuracy of 35% and human accuracy of 84%) and a small reduction in word recognition error.

The MQ System’s output baseline summary for the same reference paper is 20 sentences long; below is an excerpt:

Dialogue Act Modeling for Automatic Tagging and Recognition of Conversational Speech. In all these cases, DA labels would enrich the available input for higher-level processing of the spoken words. The relation between utterances and speaker turns is not one-to-one: a single turn can contain multiple utterances, and utterances can span more than one turn (e.g., in the case of backchanneling by the other speaker in midutterance). The most common of these are the AGREEMENT/ACCEPTS. One frequent example in our corpus was the distinction between BACKCHANNELS and AGREEMENTS (see Table 2), which share terms such as “right” and “yeah”. Networks compare to decision trees for the type of data studied here. Neural networks are worth investigating since they offer potential advantages over decision trees.

Table 5 shows the breakdown of ROUGE-L F_1 scores per document.

8 Discussion

We look at each of the three systems in detail in the following discussion.

8.1 MQ System Performance: BioMedSumm Vs. CL-Summ

Since the tasks of CL-Summ parallel those of Biomedsumm it is interesting to compare results across the two domains. MQ, which performed both Tasks 1 and 2, is used for this comparison. Table 6 compares MQ’s results over the BiomedSumm corpus against those from the CL-Summ corpus, over the runs which featured iterative improvements.

The results show an improvement in both domains, with the exception that MMR does not improve over the run that uses TF.IDF over context in CL-Summ, whereas there is an improvement in BiomedSumm. The absolute values are better in the BiomedSumm data, and looking at the confidence intervals it can be presumed that the difference between the best and the worst run is statistically significant in the BiomedSumm data. The results over the CL-Summ data are worse in general but there are no statistically significant differences. Overall, the improvement of results in CL-Summ mirrors that of the BiomedSumm data, so it can be suggested that on adding more information to the models that compute TF.IDF, the results improve. It is expected that alternative approaches, which gather related information to be added for computing the vector models will produce even better results. The results with MMR is mixed across the two domains, but the dif-

Paper ID	TF.IDF	Task 1A TF.IDF	Task 1A MMR	Paper ID	TF.IDF	Task 1A TF.IDF	Task 1A MMR
C90-2039_TRAIN	0.347	0.315	0.293	J00-3003_TRAIN	0.221	0.382	0.367
C94-2154_TRAIN	0.095	0.123	0.120	J98-2005_TRAIN	0.221	0.216	0.233
E03-1020_TRAIN	0.189	0.189	0.196	N01-1011_TRAIN	0.187	0.268	0.284
H05-1115_TRAIN	0.134	0.306	0.321	P98-1081_TRAIN	0.241	0.210	0.206
H89-2014_TRAIN	0.294	0.319	0.320	Average	0.214	0.259	0.260

Table 5: ROUGE-L F_1 results for summaries generated by the MQ system.

Run	CL-Summ				BiomedSumm			
	P	R	F_1	CI	P	R	F_1	CI
TF.IDF	0.198	0.316	0.211	0.185–0.240	0.326	0.273	0.279	0.265–0.293
topics	0.201	0.324	0.217	0.191–0.245	0.357	0.288	0.300	0.285–0.316
context	0.214	0.339	0.225	0.197–0.255	0.372	0.291	0.308	0.293–0.323
MMR	0.212	0.335	0.223	0.195–0.251	0.375	0.290	0.308	0.293–0.323

Table 6: ROUGE-L results of the MQ system runs for Task 1A.

ference is small and may not be statistically significant.

8.2 Tweaking Parameters — clair_umich Baseline

In the clair_umich baseline, for any citing sentence, the TF.IDF cosine similarity was computed with all the sentences in the source paper, and any sentences that had a cosine similarity higher than a given threshold were added to the matched sentences. Table 7 shows the precision / recall values for different cosine thresholds.

Similarity Threshold	Precision	Recall	F_1
0.01	0.027	0.641	0.051
0.05	0.048	0.426	0.087
0.1	0.060	0.235	0.095
0.2	0.079	0.081	0.080
0.3	0.062	0.032	0.042
0.4	0.022	0.085	0.012
0.5	0.007	0.002	0.003

Table 7: Precision / Recall for different values of the cosine threshold for the baseline clair_umich system.

The F_1 scores reach a maximum at a threshold of 0.1. The recall at the threshold of 0.1 is 0.23, while the precision is only 0.06. This suggests that more efforts may tackle this problem by first removing

these spurious matches that have high lexical similarity.

8.3 Tweaking Parameters — Taln.UPF

Taln.UPF’s algorithm for Task 1A chooses the top n sentences of the cited paper with the highest *score* as reference text spans. They also experimented with various settings for n to evaluate and tune the performance of their approach. Table 8 shows that on average the best result (TOP 4) is obtained when the top 4 sentences that are most similar to the citation context are selected to make up the reference span.

Paper ID	Top 2	Top 3	Top 4	Top 5
C90-2039	0.087	0.097	0.153	0.134
C94-2154	0.000	0.096	0.110	0.101
E03-1020	0.087	0.099	0.106	0.093
H05-1115	0.017	0.112	0.106	0.093
H89-2014	0.214	0.196	0.178	0.152
J00-3003	0.121	0.103	0.084	0.072
J98-2005	0.145	0.105	0.083	0.068
N01-1011	0.125	0.107	0.128	0.167
P98-1081	0.104	0.105	0.086	0.072
X96-1048	0.205	0.175	0.153	0.156
Average:	0.111	0.120	0.121	0.116

Table 8: Variation of the F_1 score when the reference text span is identified by considering the 2/3/4/5 sentences of the cited paper with highest similarity to the citation context.

8.4 Error Analysis for the Participating Systems

Several drawbacks were observed in the approach and evaluation for the MQ system. We illustrate this in the example in Figure 8.4 for Task 1A (for H89-2014).

(1) *“The statistical methods can be described in terms of Markov models.”*
 (2) *“An alternative approach taken by Jelinek, (Jelinek, 1985) is to view the training problem in terms of a “hidden” Markov model: that is, only the words of the training text are available, their corresponding categories are not known.”*
 (3) *“In this regard, word equivalence classes were used (Kupiec, 1989).”*
 (TS) *The target sentence was: “The work described here also makes use of a hidden Markov model.”*

Figure 2: Overzealous vocabulary matching problems with ROUGE as observed by MQ.

The first sentence of the sample output was very similar to the target sentence. It was not the best match, but it was a close match, and an evaluation metric such as ROUGE would reward it. On the other hand, the second sentence, even though it talked about HMMs, it was not strictly about the approach used by the paper and therefore it should not be rewarded with a good score. However, ROUGE is too lenient for this example, highlighting issues identified by the MQ system, as they followed a purely lexical approach.

In the *clair.umich* system, a number of errors made by the baseline system are due to the selection source sentences that match the words but differ slightly in their information content. An example is shown in Figure 3. Here, even though the false positive sentences contain the same lexical items (“noun”, “co-occur”, “graph”), they differ slightly in the facts presented. The detection of such subtle differences in meaning is challenging for an automated system.

Another set of difficulties arise when the citing sentence says something that is implied by the sentence in the RP, as evident in Figure 4. Here, the citing text mentions a proof from the RP, but to match the sentence in the RP, the system needs to understand that the act of showing something in a scientific paper constitutes a proof.

TALN.UPF’s top n algorithm for finding the ref-

Citing text: “use the BNC to build a co-occurrence graph for nouns, based on a co-occurrence frequency threshold”

True positives:

- “Following the method in (Widdows and Dorow, 2002), we build a graph in which each node represents a noun and two nodes have an edge between them if they co-occur in lists more than a given number of times.”

False positives:

- “Based on the intuition that nouns which co-occur in a list are often semantically related, we extract contexts of the form Noun, Noun,... and/or Noun, e.g. “genomic DNA from rat, mouse and dog”.”
- “To detect the different areas of meaning in our local graphs, we use a cluster algorithm for graphs (Markov clustering, MCL) developed by van Dongen (2000).”
- “The algorithm is based on a graph model representing words and relationships between them.”

Figure 3: Lexically similar false positive sentences.

Citing text: “The line of our argument below follows a proof provided in ... for the maximum likelihood estimator based on nite tree distributions”

False negatives:

- “We will show that in both cases the estimated probability is tight.”

Figure 4: Implied example.

erence spans makes errors in when selecting 3rd and 4th sentences. In particular, in 4 document sets (H05-1115, H89-2014, J00-3003 and X96-1048) they notice that the best F₁ score is obtained by selecting only the top 2 sentences since the 3rd, 4th and 5th most similar sentences do not overlap with the gold standard. The classification algorithm used in Task 1B also encounters sparse occurrences of the correct *implication* class, causing it to be particularly difficult.

9 Shortcomings and Limitations

The participating teams helped us to address the inherent errors in the CL corpus, which were identified in the process of annotating and parsing the corpus for use in the task-based evaluations:

- **Text encoding:** Often, the text was not in UTF-8 format as expected. The TALN.UPF team solved this by running the universal charset tool provided by Google Code over all the text and annotations in order to determine the right file encoding to use. It was found that some of the files were also in *WINDOWS-1252* and *GB18030* formats, thus making difficult the implementation of an automated homogeneous textual processing pipeline.
- **Content:** Some of the older PDF files, when parsed to text or XML, presented several text formatting issues: hyphenation problems, words not separated by blank spaces, page headers and footnotes included in the textual flow, misspelled words, spaces within words, sentences in the wrong place and so on. Unfortunately these errors were OCR parsing errors, and not within our control. We recommended that participants configure their string matching to be lenient enough to alleviate such problems.
- **Errors in citation / reference offsets:** In the original annotations, citation / reference offset numbers were character-based, and relative to an XML encoding which was not shared in the task, and did not match with the offset numbers on the text-only, cleaned version of the document. Although the text versions of the source documents were shared with the intention to help the participants, this often made their tasks more difficult if their system was geared towards numerical and not system matching. A solution was found for reference offsets by revising them to sentence ID numbers based on available XML files from the clair_umich team's donated pre-processing stage; however, the citation offsets remain character-based. As a consequence, in order to retrieve the annotated texts, other systems, such as TALN-UPF, manually searched through citing documents to

identify the correct offset. The clair_umich system created an automatic program to generate sentence offsets.

- **Discontiguous texts:** The use of “...” follows the BioMedSumm standard practice of indicating discontiguous texts, meaning that there was a gap between two text spans (citation spans or reference spans). The gap might be because text moves onto a new page. Sometimes there was a formula, page number or figure between two text spans which is not a part of the annotation. However, this notation caused mismatches for sentences which used text from different parts of the same sentence.
- **Small corpus:** The corpus comprised only a set of 10 topics, each with up to 10 citing documents. In this small corpus, participants were asked to conduct a 10-fold cross validation. The small size of the data set meant that there were no statistically significant results. Overall trends should be regarded as indicative only.
- **Errors in file construction:** An automatic, open-source software was used to map the citation annotations from the adopted annotation software, Protege, to a text file. However, participants identified several errors in the output – especially in cases where there was one-to-many mapping between citations and references. Besides this, several annotation texts had no annotation ID (Cintance Number field).

10 Conclusion

Three systems participated in the CL Pilot Task, consisting of Tasks 1A, 1B and 2. All three teams used versions of TF.IDF as baselines. For the citation span identification task, MQ and TALN.UPF implemented unsupervised algorithms, while the clair_umich system decided on a supervised approach. Overall, in this first task, clair_umich's supervised algorithm performed best, using lexical, syntactic and knowledge-based features to calculate the overlap between sentences in the citation span and the reference paper. The clair_umich system incorporated WordNet synsets for expanding and comparing cited text with reference papers, and used syntactic features to further enrich overlap calculations. In contrast, the TALN.UPF and MQ system were purely lexical-based. The MQ system was

a simple port of the system originally built for the BioMedSumm task – but with some domain-specific features discarded for this task. We believe that the lack of domain knowledge, coupled with OCR-related and PDF parsing errors, affected its performance for the CL task.

TALN.UPF attempted the second part of the task (Task 1B) for identifying the discourse facet being cited. They compared the performance of three sentence classifiers and found that the best performance was obtained using logistic regression on lemmatized word features.

Task 2 was attempted by the MQ team. They compared the baseline summaries of reference papers against gold standard summaries, based on TF.IDF calculations. In comparison with MQ’s results on the BioMedSumm task, the results were inconclusive to state whether or not the system’s features were actually aiding in generating better gold standard summaries. This may be an artifact of the small size of the corpus, but it does suggest that different domains of scientific research have different styles and features in their scientific summaries. Methods that worked in the biomedical domain do not seem to have fared well in computational linguistics.

We deem our pilot task a success, as it has spurred the development of tools and approaches for scientific summarization for our own domain of computational linguistics. However, with the limited size of the corpus and lack of a proper test corpus, we only have indicative results and do not conjecture about the optimal methods for summarizing CL research papers. Importantly, the resources from this task we feel are important artifacts for the community going forward. In particular, the annotated computational linguistics corpus – and pre-processed versions generously shared by the `clair_umich` and `TALN.UPF` teams – are freely available for researchers to use. The results of the pilot are encouraging: there seems to be ample interest from the community and it seems possible to answer more detailed methodological questions with more detailed analyses over larger datasets. In a future task in 2016, we plan a systematic annotation of a training, development and test sets, and have planned for more than one gold standard annotation. We hope also to be able to provide open-sourced tools and resources to support

the efforts of participating teams.

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