Marky: A Lightweight Web Tracking Tool for Document Annotation

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Abstract. Document annotation is an elementary task in the development of Text Mining applications, notably in defining the entities and relationships that are relevant to a given domain. Many annotation software tools have been implemented. Some are particular to a Text Mining framework while others are typical stand-alone tools. Regardless, most development efforts were driven to basic functionality, i.e. performing the annotation, and to interface, making sure operation was intuitive and visually appellative. The deployment of large-scale annotation jamborees and projects showed the need for additional features regarding inter- and intra-annotation management. Therefore, this paper presents Marky, a new Web-based document annotation tool that integrates a highly customisable annotation environment with a robust project management system. Novelty lays on the annotation tracking system, which supports *per* user and *per* round annotation change tracking and thus, enables automatic annotation correction and agreement analysis.

Keywords: Text mining, document annotation, annotation guidelines, interannotator agreement, Web application.

1 Introduction

Text Mining (TM) has a wide range of applications that require differentiated processing of documents of various natures [1]. Overall, the goal is to be able to recognise and contextualise information of relevance, notably named entities and relationships among them. Language knowledge plays a key role characterising meaningful elements in sentence composition, such as nouns and verbs. Domain implementation implies to be generally familiar with the written language and specifically aware of the terminology and "writing structure" employed in the context under analysis. For example, TM practitioners of written English are required to learn about the structure of scientific papers, and the specificities of the terminology used, in order to apply TM methods and algorithms to biomedical research documents.

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Ontologies and controlled vocabularies are crucial in capturing the semantics of a domain, and machine learning models have proven successful in employing these resources to automatically recognise and extract information of interest. Currently, there are many commercial and free TM frameworks and software tools available. Apache Unstructured Information Management Architecture (UIMA) [2] and General Architecture for Text Engineering (GATE) [3] are two meaningful examples of open source initiatives. Apart from the natural language processors and machine learning recognisers, the most sophisticated components of TM tools are the document annotator and the document viewer. Typically, user-system interaction relies on these components and therefore, attractiveness, intuitiveness, ergonomics and flexibility are major development directives. UIMA's U-Compare [4] and GATE's Teamware [5], as others alike, are offered as an integrated framework option. Solutions not bound to TM frameworks also exist. For example, MyMiner [6], EGAS [7] and PubTator [8] offer free Web-based solutions, benefiting from feedback on user experience collected at jamborees and annotation evaluations. Arguably, the data staging area is the component of the annotation life-cycle less developed so far. Namely, existing tools come short in features such as: monitoring intra-annotator and inter-annotator annotation patterns, assessing the suitability of annotation guidelines, and identifying unanticipated semantics, or other annotation issues, while still conducting annotation rounds. These features are equally important to large-scale annotation projects and smaller, more application-specific projects. Notably, they are quite important when the annotators involved in the project present different levels of domain expertise and/or are not so familiar with the concept and implications of document annotation.

This paper presents Marky, a freely accessible Web-based annotation tool that aims to provide for customised document annotation while supporting project management. Notably, the novelty lays on the annotation tracking system, which ensures that all actions occurring within the annotation project are recorded and may be reverted at any point. This ability is crucial to assess inter-annotator agreement and observe intraannotator patterns and thus, this tracking system is expected to improve the overall quality of project's results.

The next sections detail Marky design and its main functionalities. Attention is called to the following key activities: the creation of annotation projects, which involves the definition of the entities of interest and the main guidelines of annotation; the deployment of annotation rounds, which includes intra-annotator and interannotator statistics analysis; and the use of the annotation tracking system.

2 Marky Web Application

Marky is a Web-based multi-purpose document annotation application. The application was developed using the CakePHP framework (http://cakephp.org/) [9], which follows the Model–View–Controller (MVC) software pattern. Crafting application tasks into separate models, views, and controllers has made Marky lightweight, maintainable and modular. Notably, the modular design separates back-end development (e.g. the inclusion of natural language tools) from front-end development (e.g. documents and annotations visual representation), and allows developers to make changes in one part of the application without affecting the others.

Marky reaches for state-of-the-art and free Web technologies to offer the best possible user experience and provide for efficient project management. The HTML5 (http://www.w3.org/TR/html5/) and CSS3 (http://www.css3.info/) technologies support the design of intuitive interfaces whereas Ajax and JQuery (http://jquery.com/) technologies account for user-system interaction, notably document traversal and manipulation, event handling, animation, and efficient use of the network layer. Additionally, the Rangy library (http://code.google.com/p/rangy/) is used in common DOM range and selection tasks to abstract from the different browser implementations of these functionalities (namely, Internet Explorer versus DOM-compliant browsers). MySQL database engine supports data management.

This section describes the annotation life-cycle and the core management and analysis functionalities currently provided by the application.

2.1 Project Life-Cycle

A project accounts for the following main components: documents or corpus, species or concepts of interest, annotations and users (administrator and annotators). The project administrator and the team of annotators have one shared goal: to carry out the work adequately to meet the project's objectives.

At initiation, the annotation goal of the project is established and the team is defined. The documents to be annotated are automatically retrieved from an online source (e.g. PubMed Central) or uploaded by the administrator. The concepts of interest, in particular the different types of concepts and their association, are identified manually (Fig. 1) and their semantics is formalised in a set of annotation guidelines.

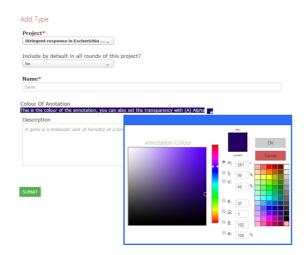


Fig. 1. Defining the types of concepts to be annotated in the project

This formalisation typically occurs after the preliminary meetings with annotators, and is meant to guide the annotation process, help annotators decide on (or disambiguate) textual mentions, and leverage annotators' domain expertise.

Often, the annotation project involves several annotators and is conducted in multiple rounds to guarantee the quality of the final annotations (Fig 2). Marky keeps track of the work done by each annotator at every round. Annotation rounds may prompt unanticipated issues, which may lead to changes in annotation guidelines, and even in the set of annotation types. Therefore, each round of annotation has associated its own set of guidelines and concept types.

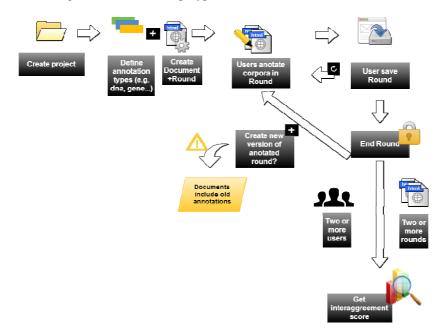


Fig. 2. The life-cycle of an annotation project in Marky

Round and user results are compared for the concept types in common. The improvement in the rates of agreement is quantified using the F-score, a common metric in IAA evaluations [10] presented below:

$$F - score = 2 \times \frac{precision \times recall}{precision + recall}$$

$$Precision = \frac{number of identical entities in set A and set B}{number of entities in set A}$$

$$Recall = \frac{number of identical entities in set A and set B}{number of entities in set B}$$

such that Set A refers to the set of annotations produced by annotator A and set B refers to the set of annotations produced by annotator B, and recall(set A, set B)=precision(set B, set A) [11].

2.2 Annotation Function

Marky offers an interactive interface allowing annotators to identify various kinds of concepts or entities within documents, according to task definition. The annotation component handles both plain text and HTML documents, and relies on state-of-theart Web technologies, such as HTML5, CSS3, Ajax and JQuery, to offer an intuitive What-You-See-Is-What-You-Get editor.

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Abstract The bacterial stringent response, triggered by nutritional deprivation, causes an accumulation of the signalit the replication arrest that occurs during the stringent response in <i>Escherichia coli</i> . Wild type cells undergo hydroximate to contain an integer number of chromosomes and a replication ongin-to-terminus ratio of 1. number of hiromosomes upon arrest. Nucleoids of these cells are decondensed; in the absence of the abilit condensed, similar to that seen after treatment with the translational inhibitor <u>thiromaphenotical</u> . After induct orresponding to the origins of replication segregate, the termini remain colocalized in wild-type cells. In cor- not show colocalized termini, suggesting that the <u>stimpent responsal arrests</u> . Chromosome segregation and accumulation is relieved in laced and limit unating, although other aspects of the <u>stringent esponsal</u> and Sept binding to non-origin to be necessary to enforce a full <u>stringent arrest</u> , affecting both initiation or the inst indication thaten is there into the stringent arrest.	a RelA-depende The growth rat by to synthesize ion of the strin trast, cells arre specific point. I nhibition of colo in to be intact. V f replication and	ent arrest after treatment with semme te prort to starwaton determines the propose, nucleoids become highly gent response, while regions steed by riampicin and cephalexin do Release from starwation causes rapid my formation by propo We propose that DNA methylation i chromosome segregation. This is
Author Summary		
Management of cell growth and division in response to environmental conditions is important for all cells. In accumulation of the nucleotice process . Armon and starvation causes a programmed change in transcription, causes an arrest of cell cycle in bacters, whose mechanism has not been throughly investigated. Here, we response is in effect, complete chromosomal replication but do not initiate new rounds and arrest with an chromosomes corresponds to the growth rate prior to arrest, in polypolid arrested cells, the chromosomal whereas the termini regions remain colocalized. The <i>E</i> , coll Enromosome remains decondensed and unsegre and segregation, concomitant with fintemesome condensation, upon release. The protein Ent , a DNA binc necessary for enforcing this arrest.	known as the show that E. co iteger number of regions at whic gated during ar	"stringent response"; ppGpp also oli cells, when the stringent of chromosomes. The number of th replication initiates are segregated, rrest and rapidly resumes replication
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Fig. 3. Document annotation in action

At each round, the annotator has a list of documents to annotate. Documents are rendered in a Web form that supports term annotation as well as annotation visual presentation (Fig 3). By right-clicking on one or more words, the annotator marks a term or concept of interest (coloured in accordance to its type). While the annotation round is open, any annotation can be edited or removed. After the administrator closes the round, annotation statistics are calculated and round assessment is conducted, to evaluate the quality of the current version of annotations and decide upon launching a new round or not.

2.3 Annotation Tracking Function

By monitoring annotation changes, the project administrator may supervise the compliance of annotators with annotation guidelines and thus, adjust these guidelines and alert annotators about incorrect or dubious curation patterns. Notably, it is highly unlikely that two annotators annotate the very same text fragments, or completely agree on text-concept associations. Mostly, variability arises from differences in domain expertise, annotation skills, and interpretation of annotation criteria.

Per round, annotator statistics describe the volume of annotations achieved by the annotator regarding the different types of concepts in analysis (Fig 4). These data are useful to assess what concepts are most annotated and are rarely annotated by a given annotator, or within the round.

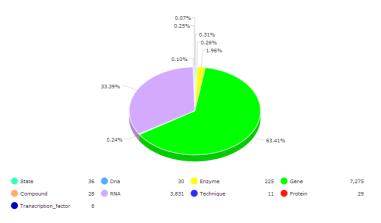


Fig. 4. Reporting annotator statistics in round

By assessing the agreement between annotators in the round, the project administrator evaluates the annotation discrepancies regarding the present set of guidelines (Fig 5). Typically, it is important to see whether certain concepts are being systematically missed or only some annotators are able to identify them. Moreover, annotators may not agree on term classification, i.e. terms are classified differently (and accounted in the statistics of different concept types), or term boundaries, i.e. annotations may only match partially.

Annotation guidelines may thus be revised so to contemplate any new semantics contexts and help to solve/minimise the annotation discrepancies observed. After deploying a new round, and besides analysing intra-round behaviour, the project administrator may compare the results between rounds to assess whether the revised set of guidelines was successful or new corrections/additions are still in need. Typically, the number of rounds performed depends considerably on time and cost constraints, leading the team to commit to a satisfactory score of agreement.

3 Conclusions

Marky is a free Web-based generic annotation tool that aims to provide highly customised annotation while supporting project management life-cycle. Indeed, Marky detaches from existing annotation tools in that it incorporates an annotation tracking system to monitor compliance with annotation guidelines and inter-annotator agreement. The ability to redo or undo annotations automatically is of help while consolidating annotation guidelines and minimises the manual work required from the annotators. Notably, the active monitoring of annotation patterns helps the administrator to "leverage" (at some extent) the expertise of the annotators by pin pointing interpretation/semantics issues that require further discussion and contextualisation.



Fig. 5. Reporting inter-annotator agreement

Regarding annotation functionalities, Marky design has favoured the use of stateof-the-art Web technologies as means to ensure wide user-system interaction and tool interoperability. Currently, Marky offers the same extent of manual operation of other tools. The ability to plug in named entity recognisers, or deploy the automatic recognition of dictionary entries will be sought after in the near future.

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