

# Comparative Analysis of Text Classification Approaches in Electronic Health Records

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## Abstract

Text classification tasks which aim at harvesting and/or organizing information from electronic health records are pivotal to support clinical and translational research. However these present specific challenges compared to other classification tasks, notably due to the particular nature of the medical lexicon and language used in clinical records.

Recent advances in embedding methods have shown promising results for several clinical tasks, yet there is no exhaustive comparison of such approaches with other commonly used word representations and classification models.

In this work, we analyse the impact of various word representations, text pre-processing and classification algorithms on the performance of four different text classification tasks. The results show that traditional approaches, when tailored to the specific language and structure of the text inherent to the classification task, can achieve or exceed the performance of more recent ones based on contextual embeddings such as BERT.

## 1 Introduction

Clinical text classification is an important task in natural language processing (NLP) (Yao et al., 2019), where it is critical to harvest data from electronic health records (EHRs) and facilitate its use for decision support and translational research. Thus, it is increasingly used to retrieve and organize information from the unstructured portions of EHRs (Mujtaba et al., 2019).

Examples include tasks such as: (1) detection of smoking status (Uzuner et al., 2008); (2) classification of medical concept mentions into family

versus patient related (Dai, 2019); (3) obesity classification from free text (Uzuner, 2009); (4) identification of patients for clinical trials (Meystre et al., 2019).

Most of these tasks involve mapping mentions in narrative texts (e.g. “pneumonia”) to their corresponding medical concepts (and concept ID) generally using the Unified Medical Language System (UMLS) (Bodenreider, 2004), and then training a classifier to identify these correctly (e.g. “pneumonia positive” versus “pneumonia negative”) (Yao et al., 2019).

Text classification performed on medical records presents specific challenges compared to the general domain (such as newspaper texts), including dataset imbalance, misspellings, abbreviations or semantic ambiguity (Mujtaba et al., 2019).

Despite recent advances in NLP, including neural-network based word representations such as BERT (Devlin et al., 2019), few approaches have been extensively tested in the medical domain and rule-based algorithms remain prevalent (Koleck et al., 2019). Furthermore, there is no consensus on which word representation is best suited to specific downstream classification tasks (Si et al., 2019; Wang et al., 2018).

The purpose of this study is to analyse the impact of numerous word representation methods (bag-of-word versus traditional and contextual word embeddings) as well as classification approaches (deep learning versus traditional machine learning methods) on the performance of four different text classification tasks. To our knowledge this is the first paper to test a comprehensive range of word representation, text pre-processing and classification methods combinations on several medical text tasks.

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## 2 Materials & Methods

### 2.1 Datasets and text classification tasks

In order to conduct our analysis we derived text classification tasks from MIMIC-III (Multiparameter Intelligent Monitoring in Intensive Care) (Johnson et al., 2016), and the Shared Annotated Resources (ShARe)/CLEF dataset (Mowery et al., 2014). These datasets are commonly used for challenges in medical text mining and act as benchmarks for evaluating machine learning models (Purushotham et al., 2018).

**MIMIC-III dataset** MIMIC-III (Johnson et al., 2016) is an openly available dataset developed by the MIT Lab for Computational Physiology. It comprises clinical notes, demographics, vital signs, laboratory tests and other data associated with 40,000 critical care patients.

We used MedCAT (Kraljevic et al., 2019) to prepare the dataset and annotate a sample of clinical notes from MIMIC-III with UMLS concepts (Bodenreider, 2004). We selected the concepts with the UMLS semantic type Disease or Syndrome (corresponding to T047), out of which we picked the 100 most frequent Concept Unique Identifier (CUIs, allowing to group mentions with the same meaning). For each concept we then randomly sampled 4 documents containing a mention of each concept, resulting in 400 documents with 2367 annotations in totals. The 100 most frequent concepts in these documents were manually annotated (and manually corrected in case of disagreement) for two text classification tasks:

- Status (affirmed/other, indicating if the disease is affirmed or negated/hypothetical);
- Temporality (current/other, indicating if the disease is current or past).

Such contextual properties are often critical in the medical domain in order to extract valuable information, as evidenced by the popularity of algorithms like ConText or NegEx (Harkema et al., 2009; Chapman et al., 2001).

Annotations were performed by two annotators, achieving an overall inter-annotator agreement above 90%. These annotations will be made publicly available.

**ShARe/CLEF (MIMIC-II) dataset** The ShARe/CLEF annotated dataset proposed by Mowery et al. (2014) is based on 433 clinical records from the MIMIC-II database (Saeed et al., 2002). It was generated for community distribution as part of the Shared Annotated Resources (ShARe) project (Elhadad et al., 2013), and contains annotations including disorder mention spans, with several contextual attributes. For our analysis we derived two tasks from this dataset, focusing on two attributes, comprising 8075 annotations for each:

- Negation (yes/no, indicating if the disorder is negated or affirmed);
- Uncertainty (yes/no, indicating if the disorder is hypothetical or affirmed).

**Text classification tasks** For both annotated datasets, we extracted from each document the portions of text containing a mention of the concepts of interest, keeping 15 words on each side of the mention (including line breaks). Each task is then made up of sequences comprising around 31 words, centered on the mention of interest, with its corresponding meta-annotation (status, temporality, negation, uncertainty), making up four text classification tasks, denoted:

- MIMIC | Status;
- MIMIC | Temporality;
- ShARe | Negation;
- ShARe | Uncertainty.

Table 1 summarizes the class distribution for each task.

Task	Class 1	Class 2	Total
<b>MIMIC   Status</b> (1: affirmed, 2: other)	1586 (67%)	781 (33%)	2367
<b>MIMIC   Temporality</b> (1: current, 2: other)	2026 (86%)	341 (14%)	2367
<b>ShARe   Negation</b> (1: yes, 2: no)	1470 (18%)	6605 (82%)	8075
<b>ShARe   Uncertainty</b> (1: yes, 2: no)	729 (9%)	7346 (91%)	8075

Table 1: Class distribution

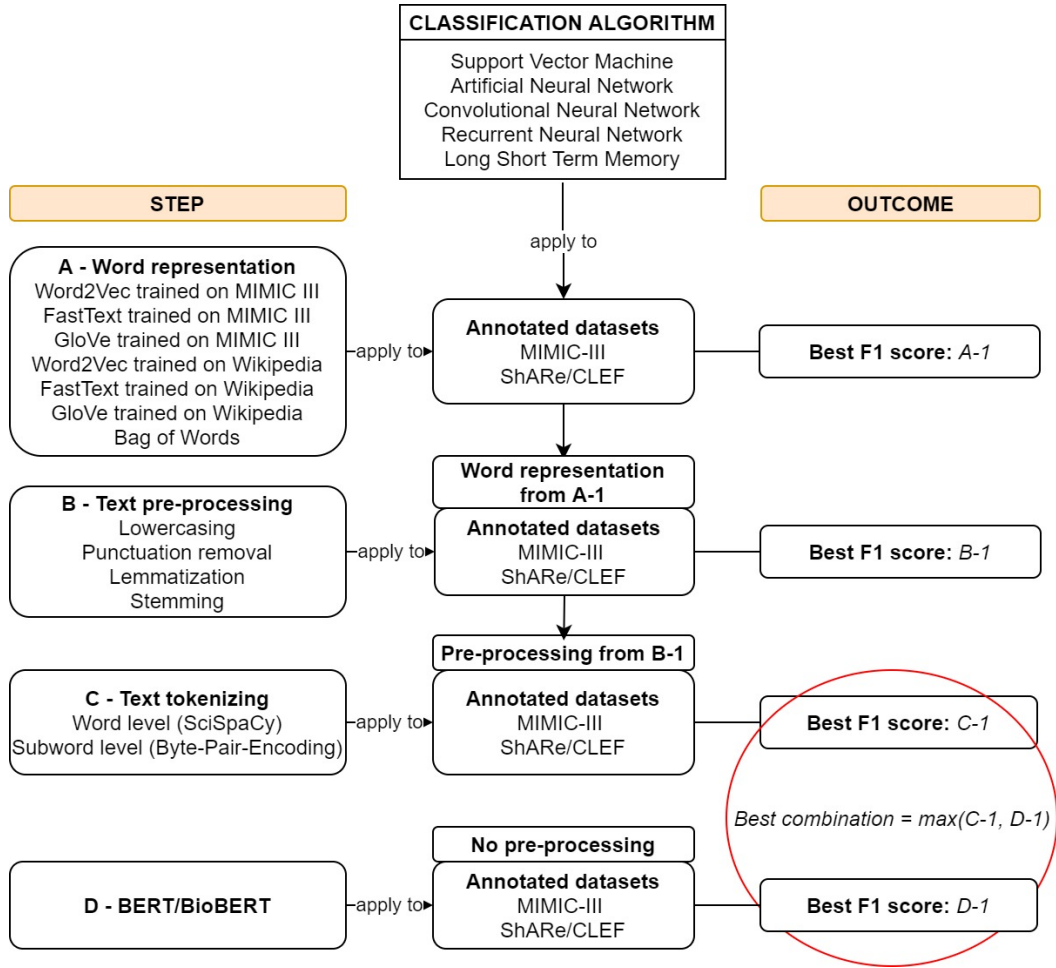


Figure 1: Main workflow

## 2.2 Evaluation steps and main workflow

We used the four different text classification tasks described in Section 2.1 in order to explore various combinations of word representation models (see Section 2.3), text pre-processing and tokenizing variations (Section 2.4) and classification algorithms (Section 2.5). In order to evaluate the different approaches we followed the steps detailed in Table 2 and Figure 1 for all four classification tasks.

Step	Description	Outcome (best F1)
A	Run all bag-of-word and traditional embeddings + classification algorithms and select the best combination (using baseline methods for text pre-processing and tokenization)	A-1
B	Using A-1 as the new baseline model, test different pre-processing methods (lowercasing, punctuation removal, lemmatization, stemming)	B-1
C	Using B-1 as the new baseline model, compare various tokenizers (word and subword level)	C-1
D	Test contextual embedding approaches: BERT (base, uncased) and BioBERT	D-1

Table 2: Evaluation steps

For each step we measured the impact by evaluating the best possible combination, based on the average F1 score (weighted average score derived from 10-fold cross validation results).

## 2.3 Word representation models

Word embeddings as opposed to bag-of-words (BoW) present the advantage of capturing semantic and syntactic meaning by representing words as real valued vectors in a dimensional space (vectors that are close in that space will represent similar words). Contextual embeddings go one step further by capturing the context surrounding the word, whilst traditional embeddings assign a single representation to a given word.

For our analysis we considered four off-the-shelf embedding models, pre-trained on public domain data, and compared them to the same embedding models trained on biomedical corpora, as well as a BoW representation.

For the traditional embeddings we chose three

commonly used algorithms, namely Word2Vec (Mikolov et al., 2013), GloVe (Pennington et al., 2014) and FastText (Bojanowski et al., 2017).

We used publicly available models pre-trained on Wikipedia and Google News for all three (Yamada et al., 2018).

To obtain medical specific models we trained all three on MIMIC-III clinical notes (covering 53,423 intensive care unit stays, including those used in the classification tasks) (Johnson et al., 2016). The following hyperparameters, aligned to off-the-shelf pre-trained models, were used: dimension of 300, window size of 10, minimum word count of 5, uncased, punctuation removed.

For the contextual embeddings we used BERT base (Devlin et al., 2019), and BioBERT (Lee et al., 2019) which are pre-trained respectively on general domain corpora and biomedical literature (PubMed abstracts and PMC articles).

Finally we used a BoW representation as a baseline approach.

## 2.4 Text pre-processing and tokenizers

In addition to pre-training several embedding models, we tested two different text tokenization methods, using the following types of tokenizers: (1) SciSpaCy (Neumann et al., 2019), a traditional tokenizer based on word detection; and (2) byte-pair-encoding (BPE) adapted to word segmentation that works on subword level (Gage, 1994; Sennrich et al., 2016).

For the word level tokenizer we chose SciSpaCy as it is specifically aimed at biomedical and scientific text processing. We further tested additional text pre-processing: lowercasing, punctuation removal, stopwords removal, stemming and lemmatization.

For the subword BPE tokenizer we used byte level byte-pair-encoding (BBPE) (Wang et al., 2019; Wolf et al., 2020). In this case the only pre-processing performed is lowercasing, whilst everything else including line breaks and spaces is left as is. This approach allows to limit the vocabulary size and is especially useful in the medical domain where a large number of words are very rare. We limited the number of words to 30522, a standard vocabulary size also used in BERT (Devlin et al., 2019).

## 2.5 Text classification algorithms

On all four classification tasks, we tested various machine learning algorithms which are widely used

for clinical data mining tasks and achieve state-of-the-art performance (Yao et al., 2019), namely artificial neural network (ANN), convolutional neural network (CNN), recurrent neural network (RNN), bi-directional long short term memory (Bi-LSTM), and BERT (Devlin et al., 2019; Wolf et al., 2020). We compared these with a statistics-based approach as a baseline, using a Support Vector Machine (SVM) classifier, a popular method used for classification tasks (Cortes and Vapnik, 1995).

For Bi-LSTM and RNN, we tested both a standard approach and one that is configured to simulate attention on the medical entity of interest. This custom approach consisted in taking the representation of the network at the position of the entity of interest, which in most cases corresponds to the center for each sequence. We refer to this latter approach as custom Bi-LSTM and custom RNN.

For ANN and statistics-based models, which are limited by the size of the dataset and embeddings (300 dimensions x 31 words x 2300 or 8000 sequences), we chose to represent sequences by averaging the embeddings of the words composing each sequence. This representation method is commonly used and has proven efficient for various NLP applications (Kenter et al., 2016).

Furthermore, each of these models was tested using different sets of parameters (e.g. varying the support function, dropout, optimizer, as reported in Table 3), the ones producing the best performance were selected for further testing and are summarized in Table 3.

	SVM	ANN	CNN	RNN Bi-LSTM
<b>Kernel or activation function</b>	<b>Radial basis</b> Linear Poly Sigmoid	ReLU + sigmoid	ReLU (with max pooling)	N/A
<b>Layers</b>	N/A	2	3	2
<b>Filters</b>	N/A	N/A	128	N/A
<b>Hidden units dimensions</b>	N/A	100	N/A	300
<b>Dropout</b>	N/A	<b>0.5</b> 0	<b>0.5</b> 0	<b>0.5</b> 0
<b>Optimizer</b>	N/A	<b>Adam</b> Stochastic Gradient Descent	<b>Adam</b> Stochastic Gradient Descent	<b>Adam</b> Stochastic Gradient Descent
<b>Learning rate</b>	N/A	0.001	0.001	0.001
<b>Epochs</b>	tolerance: 0.001	5000	200	50

Table 3: Classifiers and corresponding parameters evaluated. Parameters highlighted in **bold** were the ones selected based on performance.



### 3 Results

#### 3.1 Performance comparison for all embedding and algorithm combinations (steps A & D)

In this section we compare the performance of the different embeddings and classification approaches. We report the weighted average F1/precision/recall (weighted average value obtained from the 10-fold cross-validation results) for selected combinations on the four classification tasks in Tables 4 and 5 (full results in Appendix A.1).

For all word embedding methods tested (Word2Vec, GloVe, FastText), the ones trained on biomedical data show the best performance (see Table 4).

For classification algorithms, the best performance is obtained when using the custom Bi-LSTM model configured to target the biomedical concept of interest (see Table 5). Both contextual embeddings (BERT and BioBERT), whether trained on biomedical or general corpora, outperform any other combination of embedding/classification algorithm tested, and give results very close to the customized Bi-LSTM, as shown in Table 5.

This indicates that for tasks incorporating information about the position of the entity of interest in the text (e.g. ShARE which reports disorder mentions span offsets), the custom Bi-LSTM approach performs better than BioBERT, without necessitating any text pre-processing.

On the other hand, when looking at pure text classification, BioBERT shows better performance than a Bi-LSTM approach, and consequently may be preferred for tasks where the sequence of interest is not easily centered on a specific entity.

Finally, whilst the performance of BERT and BioBERT is relatively similar, BioBERT converges faster across all tasks tested.

#### 3.2 Impact of text pre-processing (step B)

In addition to exploring various embeddings, we tested the impact of text pre-processing on classification task performance. In order to do so, we selected the best performing word embedding obtained in the previous step (Word2Vec trained on MIMIC-III, using SciSpacy tokenizer), and compared performances between all text cleaning variations (lowercasing, punctuation removal, stemming, lemmatization).

For each variant investigated, the same pre-processing settings were applied to prepare the annotated corpus as well as to the entire MIMIC-III dataset, which was then used to re-train Word2Vec. This ensured the same vocabulary was used across the embedding and sequences to classify for each experiment.

The results, summarized in Table 6, suggest that text pre-processing has a minor impact for all classification algorithms tested. Notably, stemming and lemmatization have a slightly negative impact on performance.

#### 3.3 Impact of tokenizers (step C)

We tested the impact of tokenization on the performance of text classification tasks, focusing on SciSpacy and BBPE tokenizers, as they allow us to compare whole word versus subword unit methods. The results for the MIMIC | Status task (and using Word2Vec trained on MIMIC-III) are shown in Table 7, and indicate that the performances are roughly similar when using the BBPE tokenizer compared to SciSpacy.

Furthermore we compared both approaches in terms of speed and vocabulary size. Tokenizing text took on average 2.5 times longer with SciSpacy (250 seconds to tokenize 100,000 medical notes for SciSpacy versus 99 seconds for BBPE, excluding model loading time). For the models trained on MIMIC-III corpus, SciSpacy comprised 474,145 words, and BBPE 29,452 subword units.

#### 3.4 Embeddings analysis: word similarities comparison

Finally, in order to analyse the differences between embeddings trained on general and medical corpora, we compared the semantic information captured by Word2Vec (using SciSpacy tokenizer and without any preliminary text pre-processing).

Table 8 explores word similarities by showing the top ten similar words for medical (“cancer”) and non-medical (“concentration” and “attention”) terms.

Notably, it highlights the numerous misspellings, abbreviations and domain-specific meanings contained in the medical lexicon, suggesting that general corpora such as Wikipedia may not be appropriate when working on data from medical records (and by implication, for other specific domains).

Model	Tokenizer	Embedding	F1-score (average from 10-fold cross validation)			
			MIMIC Status	MIMIC Temporality	ShARe Negation	ShARe Uncertainty
Bi-LSTM (custom)	SciSpacy	Wiki   Word2Vec	92.8%	97.3%	98.4%	96.7%
Bi-LSTM (custom)	SciSpacy	Wiki   GloVe	93.4%	97.2%	98.4%	97.2%
Bi-LSTM (custom)	SciSpacy	Wiki   FastText	93.6%	96.9%	98.6%	96.4%
Bi-LSTM (custom)	SciSpacy	MIMIC   Word2Vec	<b>94.5%</b>	<b>97.9%</b>	<b>98.7%</b>	<b>97.3%</b>
Bi-LSTM (custom)	SciSpacy	MIMIC   GloVe	93.9%	<b>97.9%</b>	<b>98.7%</b>	96.9%
Bi-LSTM (custom)	SciSpacy	MIMIC   FastText	93.7%	97.6%	98.5%	97.2%
BERT	WordPiece	BERTbase	91.5%	97.3%	98.2%	93.6%
BioBERT	WordPiece	BioBERT	93.4%	97.3%	98.5%	94.2%
SVM	SciSpacy	Wiki   Word2Vec	76.9%	94.8%	88.5%	85.9%
SVM	SciSpacy	Wiki   GloVe	78.6%	94.9%	88.8%	87.1%
SVM	SciSpacy	Wiki   FastText	78.1%	94.4%	88.7%	86.3%
SVM	SciSpacy	BoW	82.7%	96.0%	90.2%	91.7%
SVM	SciSpacy	MIMIC   Word2Vec	80.6%	95.1%	89.8%	90.2%
SVM	SciSpacy	MIMIC   GloVe	79.1%	94.1%	89.4%	87.6%
SVM	SciSpacy	MIMIC   FastText	79.6%	93.7%	88.9%	88.0%

Table 4: Comparison of embeddings (steps A & D)

Model	Tokenizer	Embedding	F1-score (average from 10-fold cross validation)			
			MIMIC Status	MIMIC Temporality	ShARe Negation	ShARe Uncertainty
Bi-LSTM	SciSpacy	MIMIC   Word2Vec	88.4%	97.1%	96.2%	94.1%
Bi-LSTM (custom)	SciSpacy	MIMIC   Word2Vec	<b>94.5%</b>	<b>97.9%</b>	<b>98.7%</b>	<b>97.3%</b>
BERT	WordPiece	BERTbase	91.5%	97.3%	98.2%	93.6%
BioBERT	WordPiece	BioBERT	93.4%	97.3%	98.5%	94.2%
ANN	SciSpacy	MIMIC   Word2Vec	80.9%	96.5%	88.6%	86.7%
CNN	SciSpacy	MIMIC   Word2Vec	84.6%	97.3%	92.0%	87.5%
RNN	SciSpacy	MIMIC   Word2Vec	77%	96.8%	94.0%	87.1%
RNN (custom)	SciSpacy	MIMIC   Word2Vec	89.5%	96.7%	97.9%	96.5%
SVM	SciSpacy	MIMIC   Word2Vec	80.6%	95.1%	89.8%	90.2%
ANN	SciSpacy	BoW	79.8%	94.8%	89.3%	89.3%
SVM	SciSpacy	BoW	82.7%	96%	90.2%	91.7%

Table 5: Comparison of classification algorithms (steps A & D)

Task	Embedding		Text pre-processing	F1-score (average from 10-fold cross validation)					
				SVM	ANN	RNN	RNN (custom)	CNN	Bi-LSTM (custom)
MIMIC   Status	MIMIC	Word2Vec	Lowercase (L)	<b>80.6%</b>	<b>80.9%</b>	77.0%	<b>89.5%</b>	84.6%	<b>94.5%</b>
MIMIC   Status	MIMIC	Word2Vec	L + punctuation removal (LP)	80.1%	80.0%	<b>80.2%</b>	86.1%	<b>84.7%</b>	94.4%
MIMIC   Status	MIMIC	Word2Vec	LP + lemmatizing	<b>80.6%</b>	79.6%	78.0%	86.3%	83.8%	94.1%
MIMIC   Status	MIMIC	Word2Vec	LP + stemming	80.4%	79.7%	79.4%	86.1%	84.1%	94.1%

Table 6: Comparison of text pre-processing methods (step B)

Task	Embedding		Tokenizer	F1-score (average from 10-fold cross validation)					
				SVM	ANN	RNN	RNN (custom)	CNN	Bi-LSTM (custom)
MIMIC   Status	MIMIC	Word2Vec	Scispace	<b>80.6%</b>	<b>80.9%</b>	<b>77.0%</b>	<b>89.5%</b>	<b>84.6%</b>	94.5%
MIMIC   Status	MIMIC	Word2Vec	BBPE	78.8%	80.5%	76.5%	86.0%	84.3%	<b>94.7%</b>

Table 7: Comparison of tokenizing methods (step C)

Term: “cancer”		Term: “concentration”		Term: “attention”	
Word2Vec Medical	Word2Vec General	Word2Vec Medical	Word2Vec General	Word2Vec Medical	Word2Vec General
ca (0.78)	prostate (0.85)	hmf (0.51)	concentrations (0.71)	paid (0.43)	attentions (0.65)
carcinoma (0.78)	colorectal (0.82)	concentrations (0.49)	arbeitsdorf (0.67)	approximation (0.34)	notoriety (0.63)
cancer- (0.75)	melanoma (0.8)	formula (0.47)	vulkanwerft (0.65)	followup (0.33)	attracted (0.63)
caner (0.71)	pancreatic (0.8)	mct (0.47)	sophienwalde (0.64)	proximity (0.32)	criticism (0.63)
adenocarcinoma (0.71)	leukemia (0.79)	polycose (0.47)	lagerbordell (0.64)	short-term (0.31)	publicity (0.57)
ca- (0.64)	entity/breast.cancer (0.79)	virtue (0.45)	sterntal (0.64)	mangagement (0.31)	praise (0.57)
melanoma (0.64)	leukaemia (0.78)	corn (0.45)	dürrgoy (0.62)	atetntion (0.31)	aroused (0.56)
cancer;dehydration (0.63)	tumour (0.77)	dosage (0.44)	straflager (0.61)	attntion (0.3)	acclaim (0.55)
cancer/sda (0.61)	cancers (0.76)	planimetry (0.44)	maidanek (0.61)	attention (0.3)	interest (0.55)
rcc (0.61)	ovarian (0.75)	equation (0.44)	szebnie (0.61)	non-rotated (0.3)	admiration (0.55)

Table 8: Comparison of word similarities between general and domain-specific embeddings

## 4 Discussion

This study compared the impact of various embedding and classification methods on four different text classification tasks. Notably we investigated the impact of pre-training embedding models on clinical corpora versus off-the-shelf models trained on general corpora.

The results suggest that using embeddings pre-trained for the specific task (clinical corpora in our case) leads to better performance with any classification algorithm tested. However, pre-training such embeddings is not necessarily feasible due to either data or technical constraints. In this case our results highlight that using off-the-shelf embeddings trained on large general corpora such as Wikipedia still produce acceptable performance. In particular BERTbase outperformed most algorithms tested, even when these were combined with clinical embeddings.

Additionally, BioBERT was not pre-trained on medical notes but on texts from a related domain (biomedical articles and abstracts as opposed to clinical records), and therefore excludes specificities inherent to the medical domain such as misspellings or technical jargon. Despite this, BioBERT's performance is only marginally below that of the best model (custom Bi-LSTM) combined with clinical embeddings.

The various experiments conducted on text pre-processing only lead to small variations in terms of performance, and even negatively impact the performance of several algorithms, for the text classification task and embedding model tested. Given the additional constraints required to perform this step (need to train embeddings on pre-processed texts and to clean input data) and the mixed results in performance, pre-processing does not appear to be essential.

Novel tokenization methods based on subword dictionaries, whilst not improving the performance, eliminate several shortcomings presented by SciS-pacy and similar methods, notably its speed and vocabulary size.

In light of these limitations and the very small difference in performance for the task tested, BBPE appears to be a suitable alternative to traditional tokenizers and allows to reduce significantly computational costs.

Finally, custom Bi-LSTM outperforms BioBERT when it simulates attention on the entity of interest. However, this configuration requires information on the entity mention span, and then to center each document on this span. For some datasets, such information may either be readily available, or can be obtained by performing an additional named-entity extraction step. Unfortunately, many text classification tasks do not usually have this information, or may not rely on the specific entities/keywords required (e.g. sentiment analysis tasks). When Bi-LSTM is not customized, then both BERT models (trained on general and specific domains) produce the best performance, and consequently should be preferred for texts not easily allowing such customization.

## 5 Conclusion

In this article we have explored the performance of various word representation approaches (comparing bag-of-words to traditional and contextual embeddings trained on both specific and general corpora, combined with various text pre-processing and tokenizing methods) as well as classification algorithms on four different text classification tasks, all based on publicly available datasets.

A detailed performance comparison on these four tasks highlighted the efficacy of contextual embeddings when compared to traditional methods when no customization is possible, whether these embeddings are trained on specific or general corpora.

When combined with appropriate entity extraction tasks and specific domain embeddings, Bi-LSTM outperforms contextual embeddings. Across all classification algorithms, text pre-processing and tokenization approaches showed limited impact for the task and embedding tested, suggesting a rule of thumb to opt for the least time and resource intensive method.

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## References

- Olivier Bodenreider. 2004. [The Unified Medical Language System \(UMLS\): integrating biomedical terminology](#). *Nucleic Acids Research*, 32(suppl\_1):D267–D270.
- Piotr Bojanowski, Edouard Grave, Armand Joulin, and Tomas Mikolov. 2017. [Enriching Word Vectors with Subword Information](#). *Transactions of the Association for Computational Linguistics*, 5:135–146.
- Wendy W. Chapman, Will Bridewell, Paul Hanbury, Gregory F. Cooper, and Bruce G. Buchanan. 2001. [A Simple Algorithm for Identifying Negated Findings and Diseases in Discharge Summaries](#). *Journal of Biomedical Informatics*, 34(5):301–310.
- Corinna Cortes and Vladimir Vapnik. 1995. [Support-vector networks](#). *Machine Learning*, 20(3):273–297.
- Hong-Jie Dai. 2019. [Family member information extraction via neural sequence labeling models with different tag schemes](#). *BMC Medical Informatics and Decision Making*, 19(10):257.
- Jacob Devlin, Ming-Wei Chang, Kenton Lee, and Kristina Toutanova. 2019. [BERT: Pre-training of Deep Bidirectional Transformers for Language Understanding](#). *arXiv:1810.04805 [cs]*. ArXiv: 1810.04805.
- N Elhadad, W. W Chapman, T O’Gorman, M Palmer, and G Savova. 2013. The ShARe schema for the syntactic and semantic annotation of clinical texts.
- Philip Gage. 1994. A new algorithm for data compression.
- Henk Harkema, John N. Dowling, Tyler Thornblade, and Wendy W. Chapman. 2009. [ConText: an algorithm for determining negation, experience, and temporal status from clinical reports](#). *Journal of Biomedical Informatics*, 42(5):839–851.
- Alistair E. W. Johnson, Tom J. Pollard, Lu Shen, Liwei H. Lehman, Mengling Feng, Mohammad Ghassemi, Benjamin Moody, Peter Szolovits, Leo Anthony Celi, and Roger G. Mark. 2016. [MIMIC-III, a freely accessible critical care database](#). *Scientific Data*, 3(1):1–9.
- Tom Kenter, Alexey Borisov, and Maarten de Rijke. 2016. [Siamese CBOW: Optimizing Word Embeddings for Sentence Representations](#). In *Proceedings of the 54th Annual Meeting of the Association for Computational Linguistics (Volume 1: Long Papers)*, pages 941–951, Berlin, Germany. Association for Computational Linguistics.
- T.A. Koleck, C. Dreisbach, P.E. Bourne, and S. Bakken. 2019. [Natural language processing of symptoms documented in free-text narratives of electronic health records: A systematic review](#). *Journal of the American Medical Informatics Association*, 26(4):364–379.
- Zeljko Kraljevic, Daniel Bean, Aurelie Mascio, Lukasz Roguski, Amos Folarin, Angus Roberts, Rebecca Bendayan, and Richard Dobson. 2019. [MedCAT – Medical Concept Annotation Tool](#). *arXiv:1912.10166 [cs, stat]*. ArXiv: 1912.10166.
- Jinhyuk Lee, Wonjin Yoon, Sungdong Kim, Donghyeon Kim, Sunkyu Kim, Chan Ho So, and Jaewoo Kang. 2019. [BioBERT: a pre-trained biomedical language representation model for biomedical text mining](#). *Bioinformatics*, page btz682. ArXiv: 1901.08746.



- Stéphane M. Meystre, Paul M. Heider, Youngjun Kim, Daniel B. Aruch, and Carolyn D. Britten. 2019. [Automatic trial eligibility surveillance based on unstructured clinical data](#). *International Journal of Medical Informatics*, 129:13–19.
- Tomas Mikolov, Kai Chen, Greg Corrado, and Jeffrey Dean. 2013. [Efficient Estimation of Word Representations in Vector Space](#). *arXiv:1301.3781 [cs]*. ArXiv: 1301.3781.
- Danielle L Mowery, Sumithra Velupillai, Brett R South, Lee Christensen, David Martinez, Liadh Kelly, Lorraine Goeuriot, Noemie Elhadad, Guer-gana Savova, and Wendy W Chapman. 2014. Task 2: ShARe/CLEF eHealth Evaluation Lab 2014. page 12.
- Ghulam Mujtaba, Liyana Shuib, Norisma Idris, Wai Lam Hoo, Ram Gopal Raj, Kamran Khowaja, Khairunisa Shaikh, and Henry Friday Nweke. 2019. [Clinical text classification research trends: Systematic literature review and open issues](#). *Expert Systems with Applications*, 116:494–520.
- Mark Neumann, Daniel King, Iz Beltagy, and Waleed Ammar. 2019. [ScispaCy: Fast and Robust Models for Biomedical Natural Language Processing](#). *Proceedings of the 18th BioNLP Workshop and Shared Task*, pages 319–327. ArXiv: 1902.07669.
- Jeffrey Pennington, Richard Socher, and Christopher Manning. 2014. [Glove: Global Vectors for Word Representation](#). In *Proceedings of the 2014 Conference on Empirical Methods in Natural Language Processing (EMNLP)*, pages 1532–1543, Doha, Qatar. Association for Computational Linguistics.
- Sanjay Purushotham, Chuizheng Meng, Zhengping Che, and Yan Liu. 2018. [Benchmarking deep learning models on large healthcare datasets](#). *Journal of Biomedical Informatics*, 83:112–134.
- M. Saeed, C. Lieu, G. Raber, and R. G. Mark. 2002. MIMIC II: a massive temporal ICU patient database to support research in intelligent patient monitoring. *Computers in Cardiology*, 29:641–644.
- Rico Sennrich, Barry Haddow, and Alexandra Birch. 2016. [Neural Machine Translation of Rare Words with Subword Units](#). *arXiv:1508.07909 [cs]*. ArXiv: 1508.07909.
- Yuqi Si, Jingqi Wang, Hua Xu, and Kirk Roberts. 2019. [Enhancing clinical concept extraction with contextual embeddings](#). *Journal of the American Medical Informatics Association*, 26(11):1297–1304.
- Ozlem Uzuner, Ira Goldstein, Yuan Luo, and Isaac Kohane. 2008. [Identifying patient smoking status from medical discharge records](#). *Journal of the American Medical Informatics Association: JAMIA*, 15(1):14–24.
- Özlem Uzuner. 2009. [Recognizing Obesity and Comorbidities in Sparse Data](#). *Journal of the American Medical Informatics Association : JAMIA*, 16(4):561–570.
- Changhan Wang, Kyunghyun Cho, and Jiatao Gu. 2019. [Neural Machine Translation with Byte-Level Subwords](#). *arXiv:1909.03341 [cs]*. ArXiv: 1909.03341.
- Y. Wang, L. Wang, M. Rastegar-Mojarad, S. Moon, F. Shen, N. Afzal, S. Liu, Y. Zeng, S. Mehrabi, S. Sohn, and H. Liu. 2018. [Clinical information extraction applications: A literature review](#). *Journal of Biomedical Informatics*, 77:34–49.
- Thomas Wolf, Lysandre Debut, Victor Sanh, Julien Chaumond, Clement Delangue, Anthony Moi, Pierric Cistac, Tim Rault, Rémi Louf, Morgan Funtowicz, and Jamie Brew. 2020. [HuggingFace’s Transformers: State-of-the-art Natural Language Processing](#). *arXiv:1910.03771 [cs]*. ArXiv: 1910.03771.
- Ikuya Yamada, Akari Asai, Hiroyuki Shindo, Hideaki Takeda, and Yoshiyasu Takefuji. 2018. [Wikipedia2Vec: An Optimized Tool for Learning Embeddings of Words and Entities from Wikipedia](#). *arXiv:1812.06280 [cs]*. ArXiv: 1812.06280.
- Liang Yao, Chengsheng Mao, and Yuan Luo. 2019. [Clinical text classification with rule-based features and knowledge-guided convolutional neural networks](#). *BMC Medical Informatics and Decision Making*, 19(3):71.

## A Appendices

- A.1** Test set comparison of word representation, text pre-processing, tokenization and classification methods across tasks