Word embeddings trained on published case reports are lightweight, effective for clinical tasks, and completely free of protected health information

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Introduction

Word embeddings, a type of concept embedding [1] trained only on text data, are a low-dimensional, vector representation of semantic meaning that permit efficient analysis of free text. While such embeddings are useful across many domains, their use in analyzing the text of clinical encounter notes presents several challenges. First, commonly available pre-trained embeddings are derived from a broad range of non-medical text sources and thus may not capture the appropriate word sense, linguistic relations, or vocabulary needed in a clinical context [2]. Second, while it may be appealing to train new embeddings on locally available clincally notes within a health system [3], clincal corpora may be smaller than very large public corpora [4], and such embeddings cannot be shared because they would contain protected health information, thus limiting the reproducibility of experiments that would rely on such embeddings.

Prior work on domain-specific word embeddings has focused on knowledge discovery in large corpora of biomedical text [5], using medically-relevant articles from Wikipedia [6] and MEDLINE abstracts [7], and using the MIMIC-III dataset which is de-identified, but requires a data use agreement for researchers and is limited to a single center [8]. None of these approaches has yes to provide a robust evaluation of a set of embeddings specific to the text of clinical encounter notes that is lightweight, free of any protected health information, and readily shareable with other researchers.

To overcome these barriers, we examined the clinical case reports in the PubMed Open Access subset. We hypothesized that 1) embeddings trained on the full text of clinical case reports (which are already fully de-identified as a result of publication) would share similar lexical and syntactic properties with clinical notes and outperform embeddings from a non-medical domain; and 2) that subword n-grams [9] would outperform word-level n-grams [10] due to their ability to produce word vectors for out-of-vocabulary and misspelled terms, which occur frequently in electronic health record data.

Methods

We trained every combination of word embeddings using three different models, four different corpora, and four different dimension sizes. We then tested each set of embeddings with five clinical tasks. All models trained on the open access subset are available for free download: https://github.com/gweissman/clinical_embeddings

Word Embedding Models

We trained embeddings using word2vec [10] and fastText [9] with a skip-gram architecture as implemented in the Python gensim package [11]. We trained GloVe embeddings using GloVe software (version 1.2) [12]. Embeddings were built with 100, 300, 600, and 1200 dimensions.

All models trained with gensim used the following hyperparameters: skip-gram (sg=1), window=7, min_count=5, sorted_vocab=1, seed=2018. The fastText models used the additional hyperparameters: word_ngrams=1, min_n=3, max_n=8. Hyperparameters for GloVe models included: VOCAB MIN COUNT=5, WINDOW SIZE=7, MAX ITER=15.

Text pre-processing

For the OA-CR and MIMIC corpora, the entire corpus was first tokenized by sentence and then by word using the Spacy tokenizer [13]. For the longer OA-ALL and Wiki corpora, the gensim LineSentence method was used to tokenize the corpus for input into the models. Corpora were saved as single text files for input into the GloVe model. For all corpora, all years were normalized to a single token "year"; all real numbers were normalized to "real_number"; and all integers were normalized to "integer". We did not remove stopwords or perform stemming.

Corpus selection and corpus-specific pre-processing

PMC Open Access Subset

The PubMed Open Access Case Reports (OA-CR) corpus was built using case report manuscripts downloaded from PubMed Central with the query "Case Reports[ptyp] AND"2007/01/01"[PDat]:"2017/12/31"[PDat] AND English [lang] AND"humans"[MeSH Terms]" that were available under the OpenAccess Subset as identified using the "oa_file_list.csv" (ftp://ftp.ncbi.nlm.nih.gov/pub/pmc/oa_file_list.csv). Of the 515,592 reports returned by the original query, 27,575 were openly available. Text was extracted from the downloaded XML files from the abstract and body sections. All non-English text was removed using the detect method of the langdetect python package [14]. All text was converted to lowercase and stripped of non-body text, XML tags, breaks and tabs, figure tables and captions, figure references, citations, and URLs. Reports with less than 100 tokens of processed text were removed.

The PubMed OpenAccess all reports (OA-All) corpus was built using all manuscripts downloaded from PubMed Central with the query "2007/01/01" [PDat]: "2017/12/31" [PDat] AND English [lang] AND "humans" [MeSH Terms] that were available under the OpenAccess Subset as described above. Of the 5,834,856 reports returned by the original query, 630,885 were openly available. XML files were processed exactly as described for the OA-CR corpus.

The MIMIC clinical notes (MIMIC) corpus was built using the NOTEEVENTS.csv file in the MIMIC-III v1.4 dataset [15,16]. 278,269 notes were extracted and processed. All text was converted to lowercase and stripped of end fields, generic fields, de-identified notation, underscores used as separation lines, and breaks and tabs. Documents with less than 50 tokens of processed text were removed.

The Wikipedia (Wiki) corpus was built using a Wikipedia dump (downloaded 2018-11-02, 15.6GB). The download contained 18,906,413 articles, which were processed using the WikiCorpus method from the gensim python package [11]. Articles with less than 50 words were removed.

Multi-word expressions

Multiword expressions were identified in both an intrinsic and extrinsic manner. Intrinsically, pointwise mutual information (PMI) was calculated for every bigram and trigram in the OA-CR corpus using the nltk python package [17]. Bigrams and trigrams were filtered for presence in at least 10 manuscripts. Bigrams and trigrams scoring in the 50th to 95th percentile of PMI were kept based on manual review of clinical relevance. Extrinsically, the National Library of Medicine specialist lexicon [18] was used to identify MWEs by the presence of lexicon terms in the OA-CR corpus. In total, 398,217 n-grams were classified as MWEs. The KeywordProcessor method in the flashtext python package [19] was used to join all MWEs with a '_' between words in all corpora for training embeddings and for all tasks.

Tasks

Lexicographic Coverage

Models were evaluated on their ability to provide a vector representations for all the words in a given text. Coverage of all 362,430 unique words in 53,425 MIMIC ICU discharge summary notes [15,16]. Coverage is reported as proportion of all tokens for which a model could provide a word vector.

Semantic Similarity

Semantic similarity was measured by computing the correlation between the cosine distance between word pairs and the manually curated similarity scores of those pairs in the UMNSRS similarity dataset [20]. Correlations are reported using Spearman's ρ . Only word pairs for which both terms had a vector representation were considered in the comparison.

Clustering Purity

Clustering purity was measured by creating a document-level vector representation of the discharge summaries from six difference ICUs in the MIMIC dataset. Each document was represented by calculating the centroid across all individual word vectors. Words that did not produce an embedding were ignored. Duplicate reports and addenda qwere removed, leaving 49,698 ICU discharge notes. A K-means procedure with six clusters was used to cluster the document-level vector. Clustering purity was calculated by summing the number of correctly assigned notes, where ICU type is assigned to a class based on the ICU type that received the maximum number of notes for that class, and dividing by the total number of notes. Additionally, we report the 95% bootstrapped confidence interval of 1,000 replicates for the purity measure.

Linguistic Regularity

A known feature of continuous space language models is the preservation of an offset vector that captures some semantic regularity [21]. A useful set of clinically-relevant word embeddings should capture relationships related to medical care and using appropriate medical terminology. We curated a list of 100 pairs of medical terms with a relationship $is_a_treatment_for$ across inpatient, outpatient, medical, and surgical contexts likely to be discussed in a clinical encounter note. For example, metformin is a treatment for diabetes just as lisinopril is a treatment for hypertension. Therefore, an embedding that captures clinically relevant semantic information should produce $v_{metformin} - v_{diabetes} = v_{lisinopril} - v_{hypertension}$.

We used two approaches to measure how well this treatment relationship was preserved in the embedding space for each model. First, we computed the vector difference for each pair, then computed the cosine similarity between that and the centroid of all vectors. The standard deviation of this similarity is reported as a measure of the regularity of this relationship. Embeddings with low measures of variance have a more regular representation of the treatment relationship in these pairs.

Second, we used a previously reported analogy completion task [22]. For every combination of pairs, representing an analogy a:b::c:d, we calculated the cosine similarity between d and the single closest vocabulary word to a-b+c.

Mortality Prediction

We also tested each set of embeddings to represent the text of clinical notes in a mortality prediction model. The first physician encounter note charted within 24 hours of hospital admission was used to predict in-hospital mortality in a convolutional neural network. The network includes an input layer, a convolutional layer, a max pooling layer, a dense layer with 128 nodes, and an output layer with a single node using a sigmoid activation. Of the 4,170 notes, 522 (12.5%) were associated with a patient death during the hospitalization. These data were randomly split into 80% samples for training and 20% for testing. Sampling was stratified to maintain balance in the outcome between the two sets. We reported performance of each model using the mean Brier score over 1,000 runs with a 95% bootstrapped confidence interval score on the testing sample.

Results

Table 1: Summary of corpora.

Corpus	Documents	Tokens	Words - Word2Vec	Words - fastText	Words - GloVe
Open Access Case Reports	27,449	49,590,835	333,360	333,360	435,835
MIMIC-III	$220,\!453$	148,089,760	160,411	160,411	267,629
Wikipedia	$4,\!555,\!827$	2,542,552,916	3,338,426	3,338,426	3,338,427
Open Access - All	628,404	$1,\!848,\!856,\!520$	3,748,342	3,748,342	3,755,370

Lexicographic Coverage

The fastText models had perfect coverage, as embeddings for out-of-vocabulary words are built from character-level n-grams. Although the total vocabulary for the GloVe models was larger, all word2vec models had as much if not greater coverage.

Table 2: Lexical coverage for each model measured as the proportion of 362,430 unique tokens in 53,425 MIMIC ICU discharge summary notes for which each model could produce a word vector.

Training corpus	FastText	GLoVE	Word2Vec
MIMIC-III	1	0.379	0.428
Open Access - All	1	0.571	0.571
Open Access Case Reports	1	0.435	0.473
Wikipedia	1	0.401	0.401

Semantic Similarity

Models trained on the smaller clinical corpora consistently outperformed those trained on the larger Wikipedia corpus (Figure 1). Word2vec models outperformed FastText models in all corpora except in the higher dimension Wiki models. GloVe models consistently performed worst, except in the Wiki models where the higher dimension models scored the best for the corpus.

Clustering Purity

The embeddings trained with fastText produced the highest clustering purity across nearly all categories (Figure 2). With the exception of the GloVe models, which performed poorly for this task, those models trained on clinical corpora outperformed those trained with the Wikipedia corpus. Orthographically similar words in fastText embeddings appeared to be more tightly clustered than those trained with word2vec or GloVe (Figure 3).

Linguistic Regularity

Models trained with the case report corpus had the best performance, followed by those trained with all open access manuscripts (Figure 4).

Mortality Prediction

Clinically trained models performed better across higher dimensions (Figure 6). The word2vec model with 300 dimensions trained on all OA manuscripts had the best performance (Brier Score 0.1330). Models trained

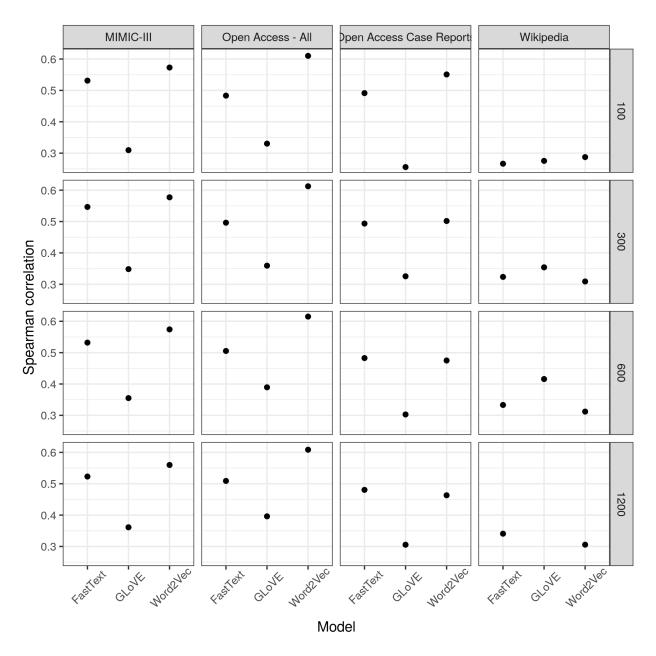


Figure 1: Spearman correlation between the cosine similarity of the words in each pair and the manually annotated similarity.

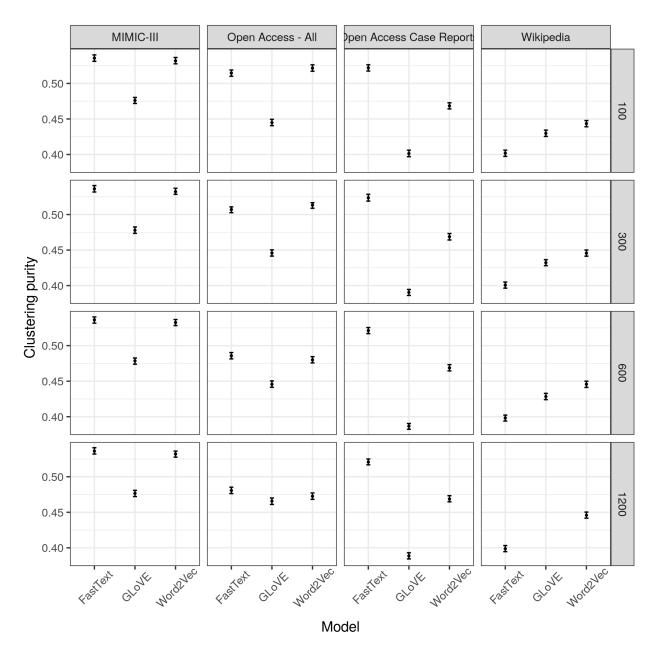


Figure 2: Clustering purity based on a k-means procedure using document-level vectors for discharge summaries from six intensive care units in the MIMIC-III dataset.

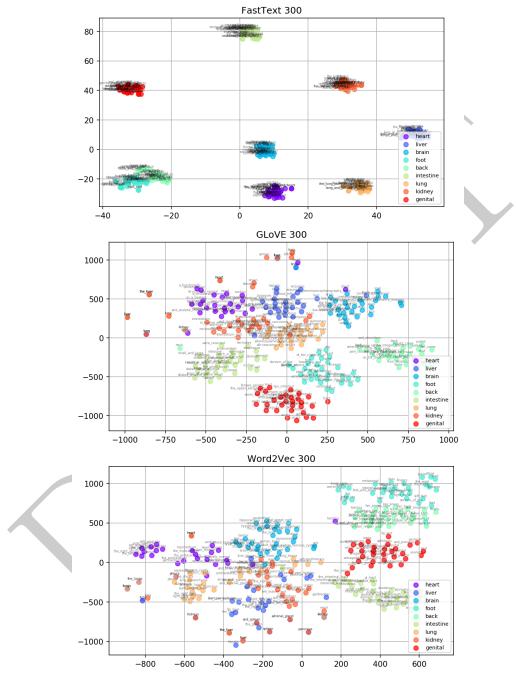


Figure 3: tSNE plot of 300-dimensional embeddings trained with clinical case reports from the PubMed Open Access Subset using fastText (top), GloVe (middle), and word2vec (bottom).

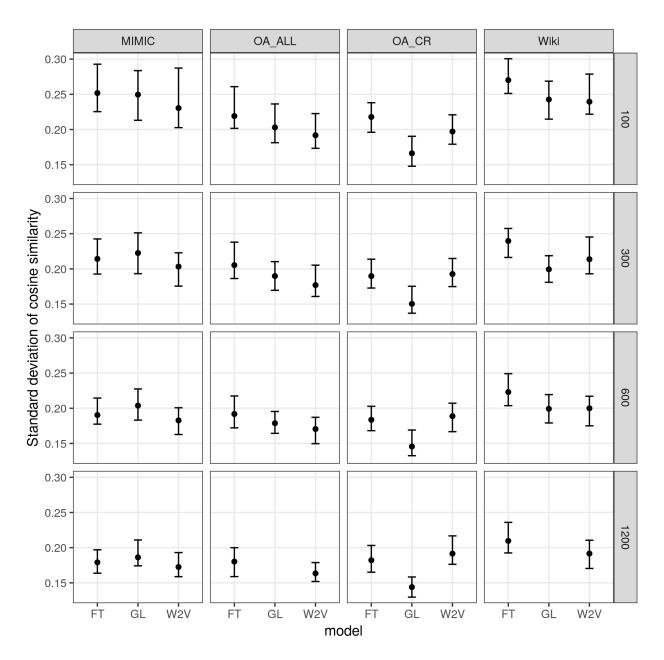


Figure 4: The variance in cosine similarity across vector differences of 100 word pairs related by is_a_treatment_for. Embeddings with lower standard deviation capture a more regular treatment relationship using the same vector difference.

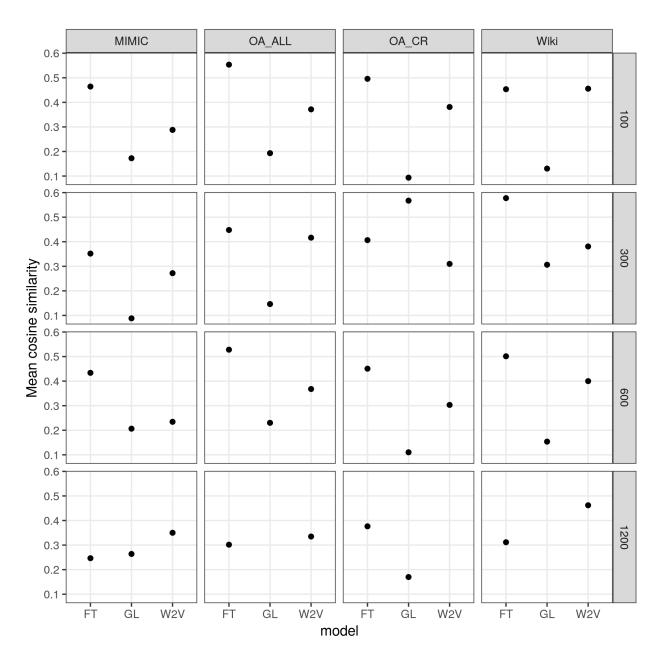


Figure 5: Mean cosine similarity between a-b+c and d across every combination of 100 pairs of treatment:disease (4,950 analogies).

on Wikipedia performed best only at 100 dimensions. For 100 dimensional size, Wiki models outperformed the other corpora, but it performed worse for higher dimensions. Both OA-CR and OA-All Word2Vec models outperformed MIMIC Word2Vec models, but not fastText models. GloVe models performed worst across all corpora.

Discussion

Word embeddings trained on the OA-CR corpus performed on par with, and often better than, embeddings trained on larger text corpora in a number of semantic meaning and NLP tasks. Optimal performance in a specific task, however, varied by corpus and model. The ability to learn from clinical language text represents a major advance in the clinical informatics community, and while it has been shown that embeddings trained on clinical-domain-specific text perform better than general-language corpora [3], we show that the lightweight embeddings trained on publically available case reports can be effective in clinically relevant natural language processing tasks.

Our general language corpus, Wiki, was trained on the largest text corpus, both in terms of documents and words. However, it performed worst in the lexicographic coverage task, scoring lower than even the OA-CR models, which were built on a corpus that was two orders of magnitude smaller. Though the Wiki corpus was the largest, the OA-All embedding models contained more words. This is explained by our requirement that a word must appear at least five times in a corpus to be included in the model. An obscure word that appears in a manuscript is more likely to appear at least five times than an obscure word in a Wikipedia article. The OA-All models performed the best on the lexicographic coverage task. Finally, we note that the GLoVE OA-CR and MIMIC models have more words in them but perfom worse on the lexicographic coverage task. This can be explained by the use of the same tokenizer in preparing the Word2Vec and FastText models and preparing the coverage test dataset, which was not used in preparing the GLoVE models.

We used the MIMIC-III clinical notes as a proxy for a local, health-system based clinical text corpus. The MIMIC corpus size is smaller than other clinical corpora used [2]. Additionally, it is important to note that the lexicographic coverage, ICU clustering, and mortality prediction tasks all utilized MIMIC-III notes for testing. The parallel performance of the models built using OpenAccess text highlight the versatility of those models in capturing clinical encounter text.

Our findings also highlight several opportunities for future work. In the UMNSRS semantic relatedness task, our highest performing model performed as well as the highest performing model reported by [2], which was also built from manuscripts in the PMC database but was not limited to those in the OpenAccess subset. Additionally, the performance of the models in the MIMIC-III based mortality prediction task is better than the performance reported in mortality prediction of severe sepsis patients [23], but worse than the performance reported in acute kidney injury patients [24]. However, our models made use solely of the text of the admission note, and could be improved by the inclusion of other data points related to the admission.

Overall, the GLoVE models performed worse than the Word2Vec and FastText models, and the Wiki corpus models performed worst among the corpora. Embedding dimension size did not affect the results of the semantic relatedness and clustering purity tasks. It did have an effect on mortality prediction. The effect of dimension size on the linguistic regularity task was ambiguous, however, the observed effect on the standard deviation measure could be the result of the increased embedding space.

In visualizing the embeddings there is a stark contrast between the tight clusters of the FastText model and the more dispersed clusters of the Word2Vec and GLoVE models. Looking at the words that cluster around a term it was evident that the FastText clusters were comprised of words in which the term was a subword of the word, for example 'heart' produced 'hearth' and 'whole_heart', while the other models did not produce these words. As FastText learns embeddings for n-grams of words, the subword will learn every context of all the words of which it is a subword. We hypothesize that this sharing of contexts place the words in close proximity in the embedding space. This also represents a limitation of our multiword expression procedure, as a single word will always be an n-gram of the joined multiword expression.

It remains an active area of research as to how best to include text data in machine learning tasks. We

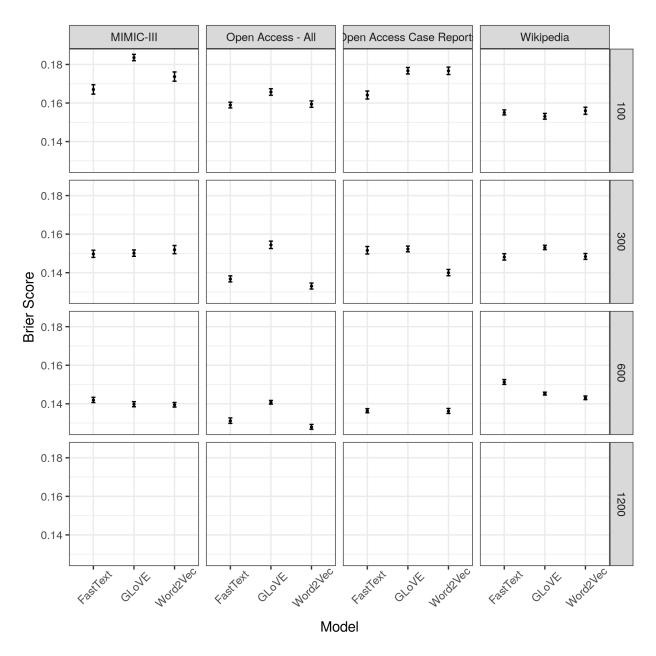


Figure 6: Performance of a mortality prediction model using only the first physician note in the first 24 hours of hospitalization. Performance is reported on the testing set using the Brier Score.

employ two different methods for converting clinical note text to manageable input for learning models. In the clustering purity task, we reduce the text of a note to a single vector and cluster the notes based on the single vector. In the mortality prediction task, we reduce every sentence of a note to a single vector, and use all sentence vectors as input to a neural network. Both methods performed well for our study.

In this study we show that embeddings trained on manuscripts from the PMC OpenAccess database capture clinical semantic meaning, and that a substantially smaller subset made up of only case reports can perform very well for clinical NLP tasks. We evaluated the models for their ability to capture clinical semantic meaning and be used for downstream predictive tasks. By outlining how we built our models and measuring their performance in a variety of ways, we hope to make word embeddings more accessible to the broader clinical community. Furthermore, by evaluating FastText, GLoVE, and Word2Vec, as well as for multiple dimension sizes, we hope that our results serve as a benchmark for embeddings built from other corpora. As the focus on data in medicine continues to grow, the ability to make use of information collected in text will be imperative.

References

- 1 Beam AL, Kompa B, Fried I et al. Clinical Concept Embeddings Learned from Massive Sources of Multimodal Medical Data. arXiv:180401486 [cs, stat] Published Online First: April 2018.http://arxiv.org/abs/1804.01486
- 2 Pakhomov SVS, Finley G, McEwan R et al. Corpus domain effects on distributional semantic modeling of medical terms. Bioinformatics 2016;32:3635–44. doi:10.1093/bioinformatics/btw529
- 3 Wang Y, Liu S, Afzal N et al. A comparison of word embeddings for the biomedical natural language processing. *Journal of Biomedical Informatics* 2018;87:12–20. doi:10.1016/j.jbi.2018.09.008
- 4 Roberts K. Assessing the Corpus Size vs. Similarity Trade-off for Word Embeddings in Clinical NLP.;10.
- 5 Chiu B, Crichton G, Korhonen A et al. How to Train good Word Embeddings for Biomedical NLP. In: Proceedings of the 15th Workshop on Biomedical Natural Language Processing. Berlin, Germany:: Association for Computational Linguistics 2016. 166–74. doi:10.18653/v1/W16-2922
- 6 Chen Z, He Z, Liu X et al. Evaluating semantic relations in neural word embeddings with biomedical and general domain knowledge bases. BMC Medical Informatics and Decision Making 2018;18. doi:10.1186/s12911-018-0630-x
- 7 Major V, Surkis A, Aphinyanaphongs Y. Utility of General and Specific Word Embeddings for Classifying Translational Stages of Research. *AMIA Annual Symposium Proceedings* 2018;**2018**:1405–14.
- 8 Boag W, Doss D, Naumann T et al. What's in a Note? Unpacking Predictive Value in Clinical Note Representations. AMIA Summits on Translational Science Proceedings 2018;2017:26–34.
- 9 Bojanowski P, Grave E, Joulin A et al. Enriching Word Vectors with Subword Information. 2016.
- 10 Mikolov T, Chen K, Corrado G et al. Efficient Estimation of Word Representations in Vector Space. 2013.
- 11 Řehůřek R, Sojka P. Software Framework for Topic Modelling with Large Corpora. In: *Proceedings of the LREC 2010 Workshop on New Challenges for NLP Frameworks*. Valletta, Malta:: ELRA 2010. 45–50.
- 12 Pennington J, Socher R, Manning C. Glove: Global Vectors for Word Representation. In: *Proceedings* of the 2014 Conference on Empirical Methods in Natural Language Processing (EMNLP). Doha, Qatar:: Association for Computational Linguistics 2014. 1532–43. doi:10.3115/v1/D14-1162
- 13 Honnibal M, Johnson M. An improved non-monotonic transition system for dependency parsing. In: *Proceedings of the 2015 conference on empirical methods in natural language processing.* Lisbon, Portugal:: Association for Computational Linguistics 2015. 1373–8.https://aclweb.org/anthology/D/D15/D15-1162
- 14 Shuyo N. Language detection library for java. 2010.http://code.google.com/p/language-detection/
- 15 Johnson AE, Pollard TJ, Shen L et al. MIMIC-III, a freely accessible critical care database. Scientific Data 2016;3:160035. doi:10.1038/sdata.2016.35

- 16 Goldberger AL, Amaral LAN, Glass L et al. PhysioBank, physiotoolkit, and physionet. Circulation 2000;101:e215–20. doi:10.1161/01.CIR.101.23.e215
- 17 Bird S, Klein E, Loper E. Natural language processing with python. 1st ed. O'Reilly Media, Inc. 2009.
- 18 Medicine (U.S.) NL of. *UMLS knowledge sources: Metathesaurus, semantic network, [and] specialist lexicon.* U.S. Department of Health; Human Services, National Institutes of Health, National Library of Medicine 2003. https://books.google.com/books?id=xTtrAAAAMAAJ
- 19 Singh V. Replace or Retrieve Keywords In Documents at Scale. ArXiv e-prints 2017.
- 20 Pakhomov S, McInnes B, Adam T *et al.* Semantic Similarity and Relatedness between Clinical Terms: An Experimental Study. *AMIA Annual Symposium Proceedings* 2010;**2010**:572–6.
- 21 Mikolov T, Yih W-t, Zweig G. Linguistic regularities in continuous space word representations. In: Proceedings of the 2013 conference of the north american chapter of the association for computational linguistics: Human language technologies. 2013. 746–51.
- 22 Newman-Griffis D, Lai A, Fosler-Lussier E. Insights into Analogy Completion from the Biomedical Domain. In: BioNLP~2017. Vancouver, Canada,: Association for Computational Linguistics 2017. 19–28. doi:10.18653/v1/W17-2303
- 23 Zhang Z, Hong Y. Development of a novel score for the prediction of hospital mortality in patients with severe sepsis: The use of electronic healthcare records with lasso regression. *Oncotarget* 2017;8:49637.
- 24 Lin K, Hu Y, Kong G. Predicting in-hospital mortality of patients with acute kidney injury in the icu using random forest model. *International journal of medical informatics* 2019;**125**:55–61.

