

# TextEssence: A Tool for Interactive Analysis of Semantic Shifts Between Corpora

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## Abstract

Embeddings of words and concepts capture syntactic and semantic regularities of language; however, they have seen limited use as tools to study characteristics of different corpora and how they relate to one another. We introduce TextEssence, an interactive system designed to enable comparative analysis of corpora using embeddings. TextEssence includes visual, neighbor-based, and similarity-based modes of embedding analysis in a lightweight, web-based interface. We further propose a new measure of embedding confidence based on nearest neighborhood overlap, to assist in identifying high-quality embeddings for corpus analysis. A case study on COVID-19 scientific literature illustrates the utility of the system. TextEssence is available from <https://github.com/drgriffis/text-essence>.

## 1 Introduction

Distributional representations of language, such as word and concept embeddings, provide powerful input features for NLP models in part because of their correlation with syntactic and semantic regularities in language use (Boleda, 2020). However, the use of embeddings as a lens to investigate those regularities, and what they reveal about different text corpora, has been fairly limited. Prior work using embeddings to study language shifts, such as the use of diachronic embeddings to measure semantic change in specific words over time (Hamilton et al., 2016; Schlechtweg et al., 2020), has focused primarily on quantitative measurement of change, rather than an interactive exploration of its qualitative aspects. On the other hand, prior work on interactive analysis of text collections has focused on analyzing individual corpora, rather than facilitating inter-corpus analysis (Liu et al., 2012; Weiss, 2014; Liu et al., 2019).

We introduce TextEssence, a novel tool that combines the strengths of these prior lines of research

by enabling interactive comparative analysis of different text corpora. TextEssence provides a multi-view web interface for users to explore the properties of and differences between multiple text corpora, all leveraging the statistical correlations captured by distributional embeddings. TextEssence can be used both for categorical analysis (i.e., comparing text of different genres or provenance) and diachronic analysis (i.e., investigating the change in a particular type of text over time).

Our paper makes the following contributions:

- We present TextEssence, a lightweight tool implemented in Python and the Svelte JavaScript framework, for interactive qualitative analysis of word and concept embeddings.
- We introduce a novel measure of *embedding confidence* to mitigate embedding instability and quantify the reliability of individual embedding results.
- We report on a case study using TextEssence to investigate diachronic shifts in the scientific literature related to COVID-19, and demonstrate that TextEssence captures meaningful month-to-month shifts in scientific discourse.

The remainder of the paper is organized as follows. §2 lays out the conceptual background behind TextEssence and its utility as a corpus analysis tool. In §3 and §4, we describe the nearest-neighbor analysis and user interface built into TextEssence. §5 describes our case study on scientific literature related to COVID-19, and §6 highlights key directions for future research.

## 2 Background

Computational analysis of text corpora can act as a lens into the social and cultural context in which those corpora were produced (Nguyen et al., 2020). Diachronic word embeddings have been shown to reflect important context behind the corpora they

are trained on, such as cultural shifts (Kulkarni et al., 2015; Hamilton et al., 2016; Garg et al., 2018), world events (Kutuzov et al., 2018), and changes in scientific and professional practice (Vylomova et al., 2019). However, these analyses have proceeded independently of work on interactive tools for exploring embeddings, which are typically limited to visual projections (Zhordaniya et al.; Warmerdam et al., 2020). TextEssence combines these directions into a single general-purpose tool for interactively studying differences between any set of corpora, whether categorical or diachronic.

## 2.1 From words to domain concepts

When corpora of interest are drawn from specialized domains, such as medicine, it is often necessary to shift analysis from individual words to *domain concepts*, which serve to reify the shared knowledge that underpins discourse within these communities. Reified domain concepts may be referred to by multi-word surface forms (e.g., “Lou Gehrig’s disease”) and multiple distinct surface forms (e.g., “Lou Gehrig’s disease” and “amyotrophic lateral sclerosis”), making them more semantically powerful but also posing distinct challenges from traditional word-level representations.

A variety of embedding algorithms have been developed for learning representations of domain concepts and real-world entities from text, including weakly-supervised methods requiring only a terminology (Newman-Griffis et al., 2018); methods using pre-trained NER models for noisy annotation (De Vine et al., 2014; Chen et al., 2020); and methods leveraging explicit annotations of concept mentions (as in Wikipedia) (Yamada et al., 2020).<sup>1</sup> These algorithms capture valuable patterns about concept types and relationships that can inform corpus analysis (Runge and Hovy, 2020).

TextEssence only requires pre-trained embeddings as input, so it can accommodate any embedding algorithm suiting the needs and characteristics of specific corpora (e.g. availability of annotations or knowledge graph resources). Furthermore, while the remainder of this paper primarily refers to concepts, TextEssence can easily be used for word-level embeddings in addition to concepts.

## 2.2 Why static embeddings?

Contextualized, language model-based embeddings can provide more discriminative features for

<sup>1</sup>The significant literature on learning embeddings from knowledge graph structure is omitted here for brevity.

NLP than static (i.e., non-contextualized) embeddings. However, static embeddings have several advantages for this comparative use case. First, they are less resource-intensive than contextualized models, and can be efficiently trained several times without pre-training to focus entirely on the characteristics of a given corpus. Second, the scope of what static embedding methods are able to capture from a given corpus has been well-established in the literature, but is an area of current investigation for contextualized models (Jawahar et al., 2019; Zhao and Bethard, 2020). Finally, the nature of contextualized representations makes them best suited for context-sensitive tasks, while static embeddings capture aggregate patterns that lend themselves to corpus-level analysis. Nevertheless, as work on qualitative and visual analysis of contextualized models grows (Hoover et al., 2020), new opportunities for comparative analysis of local contexts will provide fascinating future research.

## 3 Identifying Stable Embeddings for Analysis

While embeddings are a well-established means of capturing syntax and semantics from natural language text (Boleda, 2020), the problem of comparing multiple sets of embeddings remains an active area of research. The typical approach is to consider the nearest neighbors of specific points, consistent with the “similar items have similar representations” intuition of embeddings. This method also avoids the conceptual difficulties and low replicability of comparing embedding spaces numerically (e.g. by cosine distances) (Gonen et al., 2020). However, even nearest neighborhoods are often unstable, and vary dramatically across runs of the same embedding algorithm on the same corpus (Wendlandt et al., 2018; Antoniak and Mimno, 2018). In a setting such as our case study, the relatively small sub-corpora we use (typically less than 100 million tokens each) exacerbate this instability. Therefore, to quantify variation across embedding replicates and identify informative concepts, we introduce a measure of *embedding confidence*.<sup>2</sup>

We define embedding confidence as the mean overlap between the top  $k$  nearest neighbors of a

<sup>2</sup>An *embedding replicate* here refers to the embedding matrix output by running a specific embedding training algorithm on a specific corpus. Ten runs of word2vec on a given Wikipedia dump produce ten replicates; using different Wikipedia dumps would produce one replicate each of ten different sets of embeddings.

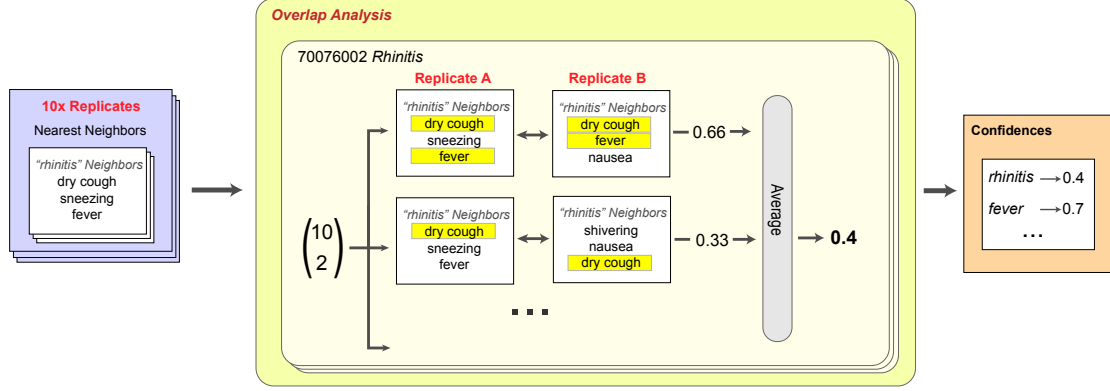


Figure 1: Illustration of embedding confidence calculation, using 10 embedding replicates.

given item between multiple embedding replicates. Formally, let  $E^1 \dots E^m$  be  $m$  embedding replicates trained on a given corpus, and let  $kNN^i(c)$  be the set of  $k$  nearest neighbors by cosine similarity of concept  $c$  in replicate  $E^i$ . Then, the embedding confidence  $EC@k$  is defined as:

$$EC@k(c, E^1 \dots E^m) = \frac{1}{m(m-1)} \sum_i \sum_{j \neq i} |kNN^i(c) \cap kNN^j(c)|$$

This calculation is illustrated in Figure 1.

We can then define the set of *high-confidence* concepts for the given corpus as the set of all concepts with an embedding confidence above a given threshold. A higher threshold will restrict to highly-stable concepts only, but exclude the majority of embeddings. We recommend an initial threshold of 0.5, which can be configured based on observed quality of the filtered embeddings.

After filtering for high-confidence concepts, we summarize nearest neighbors across replicates by computing *aggregate nearest neighbors*. The aggregate neighbor set of a concept  $c$  is the set of high-confidence concepts with highest average cosine similarity to  $c$  over the embedding replicates. This helps to provide a more reliable picture of the concept’s nearest neighbors, while excluding concepts whose neighbor sets are uncertain.

#### 4 The TextEssence Interface

The workflow for using TextEssence to compare different corpora is illustrated in Figure 2. Given the set of corpora to compare, the user (1) trains embedding replicates on each corpus; (2) identifies the high-confidence set of embeddings for each corpus; and (3) provides these as input to TextEssence.

TextEssence then offers three modalities for interactively exploring their learned representations: (1) Browse, an interactive visualization of the embedding space; (2) Inspect, a detailed comparison of a given concept’s neighbor sets across corpora; and (3) Compare, a tool for analyzing the pairwise relationships between two or more concepts.

##### 4.1 Browse: visualizing embedding changes

The first interface presented to the user is an overview visualization of one of the embedding spaces, projected into 2-D using t-distributed Stochastic Neighbor Embedding (t-SNE) (van der Maaten and Hinton, 2008). High-confidence concepts are depicted as points in a scatter plot and color-coded by their high-level semantic grouping (e.g. “Chemicals & Drugs,” “Disorders”), allowing the user to easily navigate to an area of interest. The user can select a point to highlight its aggregated nearest neighbors in the high-dimensional space, an interaction similar to TensorFlow’s Embedding Projector (Smilkov et al., 2016) that helps distinguish true neighbors from artifacts of the dimensionality reduction process.

The Browse interface also incorporates novel interactions to address the problem of visually comparing results from several corpora (e.g., embeddings from individual months). The global structures of the corpora can differ greatly in both the high-dimensional and the low-dimensional representations, making visual comparison difficult. While previous work on comparing projected data has focused on aligning projections (Liu et al., 2020; Chen et al., 2018) and adding new comparison-focused visualizations (Cutura et al., 2020), we chose to align the projections using a simple Procrustes transformation and enable the

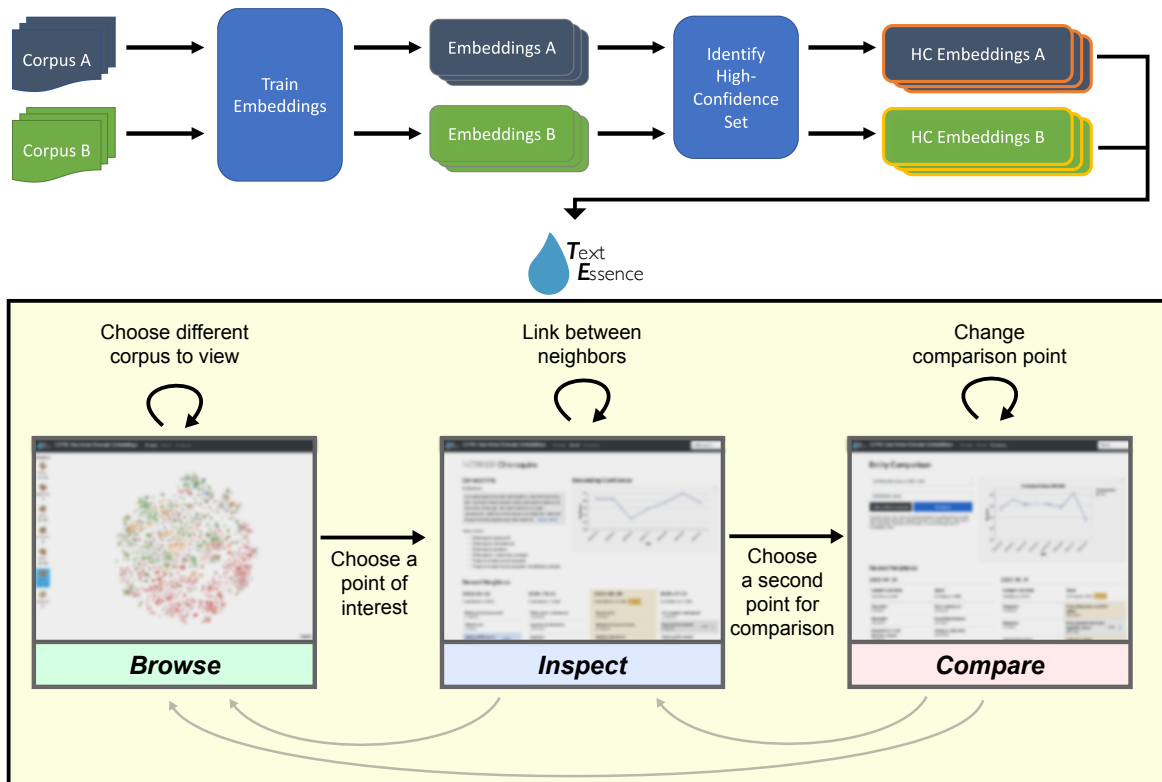


Figure 2: Workflow for comparing corpus embeddings with TextEssence. The system enables three different kinds of interactions: (1) Browse the embedding space for each corpus; (2) Inspect a single concept in each corpus; and (3) Compare two or more concepts across corpora. Each view transitions to the others using the current concept.

user to compare them using animation. When the user hovers on a corpus thumbnail, lines are shown between the positions of each concept in the current and destination corpora, drawing attention to the concepts that shift the most. Upon clicking the thumbnail, the points smoothly follow their trajectory lines to form the destination plot. In addition, when a concept is selected, the user can opt to *center* the visualization on that point and then transition between corpora, revealing how neighboring concepts move relative to the selected one.

#### 4.2 Inspect: tracking individual concept change

Once a particular concept of interest has been identified, the Inspect view presents an interactive table depicting how that concept’s aggregated nearest neighbors have changed over time. This view also displays other contextualizing information about the concept, including its definitions (derived from the UMLS (Bodenreider, 2004) for our case study<sup>3</sup>), the terms used to refer to the con-

<sup>3</sup>We included definitions from all English-language sources in the UMLS, as SNOMED CT includes definitions

cept (limited to SNOMED CT for our case study), and a visualization of the concept’s embedding confidence over the sub-corpora analyzed. For information completeness, we display nearest neighbors from every corpus analyzed, even in corpora where the concept was not designated high-confidence (note that a concept must be high-confidence in at least one corpus to be selectable in the interface). In these cases, a warning is shown that the concept itself is not high-confidence in that corpus; the neighbors themselves are still exclusively drawn from the high-confidence set.

#### 4.3 Compare: tracking pair similarity

The Compare view facilitates analysis of the changing relationship between two or more concepts across corpora (e.g. from month to month). This view displays paired nearest neighbor tables, one per corpus, showing the aggregate nearest neighbors of each of the concepts being compared. An adjacent line graph depicts the similarity between the concepts in each corpus, with one concept specified as the reference item and the others serving as only for a small subset of concepts.

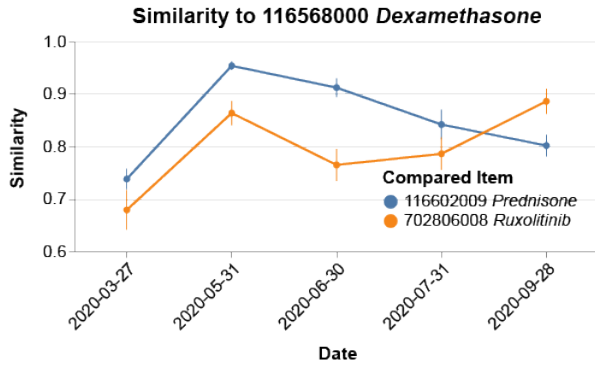


Figure 3: Similarity over time of two drugs to 116568000 *Dexamethasone* in our case study. April, August, and October are omitted as *Dexamethasone* was not high confidence for these months. Similarity values are mean over embedding replicates within each month; error bars indicate standard deviations.

Month	Docs	Words	Entities	Hi-Conf.
March	41,750	158M	38,451	15,100
April	10,738	41M	25,142	1,851
May	73,444	125M	40,297	5,051
June	24,813	34M	19,749	2,729
July	24,786	35M	19,334	2,800
August	28,642	31M	19,134	2,407
September	33,732	38M	20,947	4,381
October	38,866	44M	21,470	1,990

Table 1: 2020 monthly snapshots of *CORD-19* dataset (documents added each month only; not cumulative). Entities denotes the number of SNOMED CT codes for which embeddings were learned; Hi-Conf. is the subset of these that had confidence above the 0.5 threshold.

comparison items (similar to Figure 3). Similarity between two concepts for a specific corpus is calculated by averaging the cosine similarity between the corresponding embeddings in each replicate.

## 5 Case Study: Diachronic Change in *CORD-19*

The scale of global COVID-19-related research has led to an unprecedented rate of new scientific findings, including developing understanding of the complex relationships between drugs, symptoms, comorbidities, and health outcomes for COVID-19 patients. We used TextEssence to study how the contexts of medical concepts in COVID-19-related scientific literature have changed over time. Table 1 shows the number of new articles indexed in the COVID-19 Open Research Dataset (*CORD-19*; Wang et al. (2020a)) from its beginning in March 2020 to the end of October 2020; while additions of new sources over time led to occasional jumps

Concept	Month(s)	Representative neighbors
44169009 Anosmia	Mar-Sep	2553606007 Gustatory, 51388003 Pharyngeal pain, 60707004 Taste
	Oct	15387003 Vomiting, 73879007 Nausea, 49727002 Cough
116568000 Dexamethasone	Mar	19130008 Injury, 417746004 Traumatic injury
	May-Jul	116602009 Prednisone, 108675009 Infliximab
	Sep	702806008 Ruxolitinib
83490000 Hydroxy-chloroquine	All	80229008 Antimalarial agent, 96034006 Azithromycin
	Aug	198051006 Nosocomial infection, 233765002 Respiratory failure without hypercapnia

Table 2: Representative nearest neighbors (manually selected from top 10) for three concepts in *CORD-19*, grouped by period of observation. Complete nearest neighbor tables are omitted for brevity, but may be viewed using our released code and data.

in corpus volumes, all are sufficiently large for embedding training. We created disjoint sub-corpora containing the new articles indexed in *CORD-19* each month for our case study.

*CORD-19* monthly corpora were tokenized using ScispaCy (Neumann et al., 2019), and concept embeddings were trained using JET (Newman-Griffis et al., 2018), a weakly-supervised concept embedding method that does not require explicit corpus annotations. We used SNOMED Clinical Terms (SNOMED CT), a widely-used reference representing concepts used in clinical reporting, as our terminology for concept embedding training, using the March 2020 interim release of SNOMED CT International Edition, which included COVID-19 concepts. We trained JET embeddings using a vector dimensionality  $d = 100$  and 10 iterations, to reflect the relatively small size of each corpus. We used 10 replicates per monthly corpus, and a high-confidence threshold of 0.5 for EC@5.

### 5.1 Findings

TextEssence captures a number of shifts in *CORD-19* that reflect how COVID-19 science has devel-

oped over the course of the pandemic. Table 2 highlights key findings from our preliminary investigation into concepts known *a priori* to be relevant. Please note that while full nearest neighbor tables are omitted due to space limitations, they can be accessed by downloading our code and following the included guide to inspect CORD-19 results.

**44169009** *Anosmia* While associations of anosmia (loss of sense of smell) were observed early in the pandemic (e.g., Hornuss et al. (2020), posted in May 2020), it took time to begin to be utilized as a diagnostic variable (Talavera et al., 2020; Wells et al., 2020). *Anosmia*'s nearest neighbors reflect this, staying stably in the region of other otolaryngological concepts until October (when Talavera et al. (2020); Wells et al. (2020), *inter alia* were included in CORD-19), where we observe a marked shift in utilization to play a similar role to other common symptoms of COVID-19.

**116568000** *Dexamethasone* The corticosteroid dexamethasone was recognized early as valuable for treating severe COVID-19 symptoms (Lester et al. (2020), indexed July 2020), and its role has remained stable since (Ahmed and Hassan (2020), indexed October 2020). This is reflected in the shift of its nearest neighbors from prior contexts of traumatic brain injury (Moll et al., 2020) to a stable neighborhood of other drugs used for COVID-19 symptoms. However, in September 2020, 702806008 *Ruxolitinib* emerges as *Dexamethasone*'s nearest neighbor. This reflects a spike in literature investigating the use of ruxolitinib for severe COVID-19 symptom management (Gozzetti et al., 2020; Spadea et al., 2020; Li and Liu, 2020). As the similarity graph in Figure 3 shows, the contextual similarity between dexamethasone and ruxolitinib steadily increases over time, reflecting the growing recognition of ruxolitinib's new utility (Caocci and La Nasa (2020), indexed May 2020).

**83490000** *Hydroxychloroquine* Hydroxychloroquine, an anti-malarial drug, was misleadingly promoted as a potential treatment for COVID-19 by President Trump in March, May, and July 2020, leading to widespread misuse of the drug (Englund et al., 2020). As a result, a number of studies re-investigated the efficacy of hydroxychloroquine as a treatment for COVID-19 in hospitalized patients (Ip et al. (2020); Albani et al. (2020); Rahmani et al. (2020), all indexed August 2020). This shift is reflected in the neighbors of *Hydroxychloroquine*, adding investigative outcomes such as nosocomial

(hospital-acquired) infections and respiratory failure to the expected anti-malarial neighbors.

## 6 Conclusion and Future Work

TextEssence is an interactive tool for comparative analysis of word and concept embeddings. Our case study on scientific literature related to COVID-19 demonstrates that TextEssence can be used to study diachronic shifts in usage of domain concepts, and a previous study on medical records (Newman-Griffis and Fosler-Lussier, 2019) showed that the technologies behind TextEssence can be used for categorical comparison as well.

The utility of TextEssence is not limited to analysis of text corpora. In settings where multiple embedding strategies are available, such as learning representations of domain concepts from text sources (Beam et al., 2020; Chen et al., 2020), knowledge graphs (Grover and Leskovec, 2016), or both (Yamada et al., 2020; Wang et al., 2020b), TextEssence can be used to study the different regularities captured by competing algorithms, to gain insight into the utility of different approaches. TextEssence also provides a tool for studying the properties of different terminologies for domain concepts, something not previously explored in the computational literature.

While our primary focus in developing TextEssence was on its use as a qualitative tool for targeted inquiry, diachronic embeddings have significant potential for knowledge discovery through quantitative measurement of semantic differences. However, vector-based comparison of embedding spaces faces significant conceptual challenges, such as a lack of appropriate alignment objectives and empirical instability (Gonen et al., 2020). While nearest neighbor-based change measurement has been proposed (Newman-Griffis and Fosler-Lussier, 2019; Gonen et al., 2020), its efficacy for small corpora with limited vocabularies remains to be determined. Our novel mbedding confidence measure offers a step in this direction, but further research is needed.

Our implementation and experimental code is available at <https://github.com/drgriffis/text-essence>, and the database derived from our CORD-19 analysis is available at <https://doi.org/10.5281/zenodo.4432958>. A screenshot of TextEssence in action is available at <https://youtu.be/1xEEfsMwL0k>.

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