With over 50,000 publications and research papers on the corona-virus family till date, it has become difficult to search across and get useful insights for medical practitioners.

As a *DataScientist* at Google, you are tasked with solving this problem with the help of Machine Learning.

<u>Efficient Estimation of Word Representations in Vector Space</u> (https://arxiv.org/pdf/1301.3781.pdf)

GloVe: Global Vectors for Word Representation (https://nlp.stanford.edu/pubs/glove.pdf)

How can we solve for this problem?

- Can we match keywords from user queries that are present in abstract?
- If we do keyword matching, will we be able to understand the user's intent? For e.g: 'origin' and 'discovery'
- Should we consider the context of the words, then?

Let us build a search engine using Word Embeddings

Datset

- COVID-19 Open Research Dataset, consisting of all publications/research papers related to Covid-19.
- Dataset contains Title, Abstract, Dol among other identifiers.
- To download https://www.kaggle.com/allen-institute-for-ai/CORD-19-research-challenge)

Let us explore the dataset ¶

Importing Necessary Libraries

```
In [ ]: !pip install -q langdetect
!gdown 1VkNpuudQnlj7g5uUCNPJ4MKxFdDdh7bZ
```

```
| 981 kB 15.6 MB/s eta 0:00:01 Building wheel for langdetect (setup.py) ... done Downloading...
```

From: https://drive.google.com/uc?id=1VkNpuudQnlj7g5uUCNPJ4MKxFdDdh7bZ (ht
tps://drive.google.com/uc?id=1VkNpuudQnlj7g5uUCNPJ4MKxFdDdh7bZ)

To: /content/metadata.csv

100% 1.65G/1.65G [00:15<00:00, 107MB/s]

```
In [ ]:
        import spacy
        import string
        import warnings
        import numpy as np
        import pandas as pd
        from pprint import pprint
        from IPython.utils import io
        from tqdm.notebook import tqdm
        from gensim.models import Word2Vec
        from langdetect import DetectorFactory, detect
        from IPython.core.display import HTML, display
        from IPython.display import Image
        from spacy.lang.en.stop_words import STOP_WORDS
        warnings.filterwarnings('ignore')
        tqdm.pandas()
```

Loading the dataset

```
In [ ]: DATA = pd.read_csv("metadata.csv").sample(100000) # taking only 100000 resed
         DATA.reset_index(inplace=True, drop=True)
         print(DATA.columns)
         DATA.head(2)
         Index(['cord_uid', 'sha', 'source_x', 'title', 'doi', 'pmcid', 'pubmed_i
         d',
                 'license', 'abstract', 'publish_time', 'authors', 'journal', 'mag_i
         d',
                 'who_covidence_id', 'arxiv_id', 'pdf_json_files', 'pmc_json_files',
                 'url', 's2_id'],
                dtype='object')
Out[3]:
             cord_uid
                                                          sha source_x
                                                                             title
                                                                        'I Get High
                                                                           With a
                                                                        Little Help 10.3389/fpsyg.
              gnh51fzr
                       fe177de29f14dd45747a3d17c2468300245b0fb4
                                                                  PMC
                                                                         From My
                                                                         Friends...
                                                                        Arenavirus
                                                                           Stable
                                                                           Signal
          1 wbh06gzb ab312eb3286e189488707ea7ac8551401a178940
                                                                  PMC
                                                                                     10.1128/m
                                                                        Peptide Is
                                                                             the
                                                                         Keysto...
```

Among all the features available, we can drill down to using 'Abstract' to create the search engine. Why?

 Abstract gives a complete, yet concise, understanding of the publication's research and findings.

```
In [ ]: DATA.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 100000 entries, 0 to 99999
Data columns (total 19 columns):

# Column Non-Null Count Dtype
--- ---- 0 cord_uid 100000 non-null object
1 sha 35242 non-null object
2 source_x 100000 non-null object
3 title 99946 non-null object
4 doi 62094 non-null object
5 pmcid 36827 non-null object
6 pubmed_id 47157 non-null object
7 license 100000 non-null object
8 abstract 77551 non-null object
9 publish_time 99844 non-null object
10 authors 97680 non-null object
11 journal 91699 non-null object
12 mag_id 0 non-null object
13 who_covidence_id 45679 non-null object
14 arxiv_id 1420 non-null object
15 pdf_json_files 35242 non-null object
16 pmc_json_files 29756 non-null object
17 url 64995 non-null object
18 s2_id 92350 non-null float64
dtypes: float64(2), object(17)
memory usage: 14.5+ MB
```

Approach

For each incoming query, **calculate semantic similarity** with all documents in the dataset and **pick top N publications** from the dataset.

But, before we start building this out, we will have to process the abstract and represent them as contextual word embeddings.

What is the need for representing words as texts?

 Machine Learning algorithms and Deep Learning Architectures are capable of processing only numbers.

But, we have already learnt to represent words as discrete representations, why learn another technique?

- Discrete representations treat words as completely **independent entities**.
- Cannot capture relationship between words, thus providing **no context**.

So how do we capture contexts?

- Using Continous word representations.
- Capable of capturing context of a word, based on syntactic and semantic similarity.
- · Capture word to word relationships.
- Techniques for representations:
 - Singular Value Decomposition (SVD) based methods
 - Iteration based methods

What are SVD based methods and how are embeddings generated?

- · Key Concepts:
 - SVD theorem says its always possible to decompose a matrix A_{m*n} as a product of U_{m*m} , $\Sigma m*n$, V_{n*n} , where U^TU = I and V^TV = I.
 - We loop over the corpus and accumulate word co-occurence counts in the form of a matrix called 'X'.
 - Perform **SVD** on **X** to get $U\Sigma V^T$ decomposition.
 - Use rows of U as the word embeddings for all words in our dictionary.

Let us discuss few choices of generating "X":

{A} Word-Document Matrix:

- Assumption Words that are related will often appear in the same documents and vice versa.
- Loop over the corpus and for each time word i appears in document j, we add one to entry X_{ii} .
- The X generated is of the size $\{R^{|V|\times M}\}$ and it scales with the number of documents.

{B} Co-Occurence Matrix:

- Similar to Word-Document Matrix, however, the **matrix X contains co-occurrences of words** thereby becoming an **affinity matrix**.
- X is generating by counting the **number of times each word appears inside a window** of a particular size **around the word of interest**.
- · Calculate this count across all the words in the corpus.

SVD with an example:

Corpus contains 3 sentences:

- 1. "I enjoy flying."
- 2. "I like NLP."
- 3. "I like deep learning."

Based on window size of 1, the count matrix will be:

SVD Count Matrix

$$X = \begin{bmatrix} I & like & enjoy & deep & learning & NLP & flying \\ like & 2 & 1 & 0 & 0 & 0 & 0 & 0 \\ like & 2 & 0 & 0 & 1 & 0 & 1 & 0 & 0 \\ enjoy & 1 & 0 & 0 & 0 & 0 & 0 & 1 & 0 \\ learning & 0 & 1 & 0 & 0 & 0 & 0 & 1 \\ learning & 0 & 0 & 0 & 1 & 0 & 0 & 0 & 1 \\ NLP & 0 & 1 & 0 & 0 & 0 & 0 & 0 & 1 \\ flying & 0 & 0 & 1 & 0 & 0 & 0 & 0 & 1 \\ . & 0 & 0 & 0 & 0 & 1 & 1 & 1 & 0 \end{bmatrix}$$

Performing SVD on the Co-Occurence Matrix

Pick an index k from U matrix that will capture variance based on desired percentage using

$$\frac{\sum_{i=1}^{k} \sigma_i}{\sum_{i=1}^{|V|} \sigma_i}$$

Finally, the submatrix $U_{1:|V|,1:k}$ will form our word embedding matrix.

$$|V| \begin{bmatrix} \hat{X} \end{bmatrix} = |V| \begin{bmatrix} 1 & 1 & 1 & 1 \\ u_1 & u_2 & \dots & 1 \end{bmatrix} k \begin{bmatrix} \sigma_1 & 0 & \dots & \sigma_2 & \dots \\ 0 & \sigma_2 & \dots & \vdots & \vdots & \ddots \end{bmatrix} k \begin{bmatrix} - & v_1 & - \\ - & v_2 & - & \vdots & \vdots & \ddots \end{bmatrix}$$

Word embedding matrix

Can we use SVD for building for our search engine?

No. While SVD based methods give us word vectors that are more than sufficient to encode semantic and syntactic information they are associated with many **dis-advantages**:

- SVD based methods do not scale well for large matrices. Given the size of our data, it will be **computationally expensive** to perform.
- The matrix can become extremely sparse since most words do not co-occur.

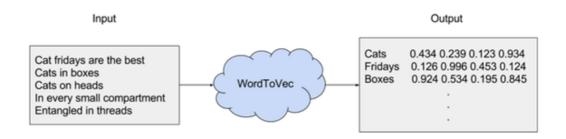
How do we circumvent the drawbacks of SVD based methods?

- Instead of computing and storing global information about some huge data, we can build
 a model that can encode contextual information and produce word vectors.
- One such architecture called the Word2Vec, an iterative method, to generate word embeddings.

What is Word2Vec?

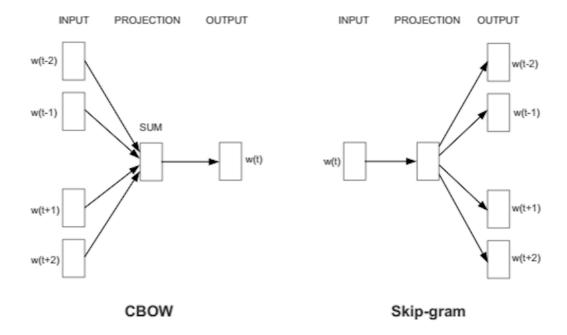
Word2Vec works with the idea that two words are similar if they share or occur in similar context.

- Word2vec is a shallow neural network that is trained to reconstruct linguistic contexts of words.
- Word2vec takes in large corpus of text as input and produces a vector of several hundred dimensions.
- Each unique word in the corpus is assigned a corresponding vector in the space.
- Word vectors are positioned in the vector space such that words that share common contexts in the corpus are located close to one another in the space.



How does Word2Vec learn word embeddings?

- Word2vec utilizes either of the two model architectures to produce a distributed representation of words:
- Continuous Bag-of-Words (CBoW)

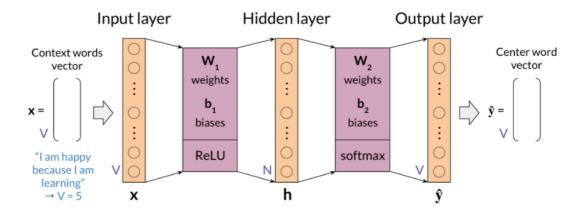


- In the Continuous Bag-of-Words (CBoW) architecture, the model predicts the current word from a window of surrounding context words.
- In the **Skip-gram** architecture, the model **uses current word to predict** the surrounding window of **context words**.

Let us deep-dive into CBoW!!

- CBoW uses a shallow feed forward neural network with single hidden layer to produce word embeddings (vectors).
- Given a word, CBoW uses before and after words, called as the context words, to create a vector representation.

Architecture of CBoW:



- Input Layer
 - Takes in context words as inputs.
 - Is of the size **V**, size of the vocabulary (unique words in corpus).
- Hidden Layer

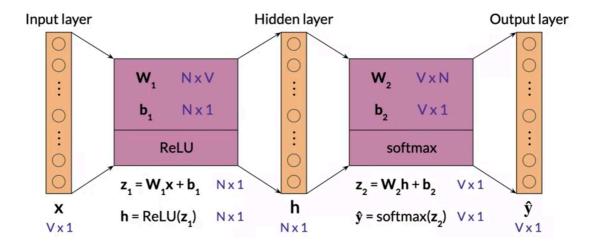
- Is of the size N, a hyper-parameter of the network that defines the shape of word representations.
- ReLU used as the activation function.
- Output Layer
 - Outputs **probability** for all the words in the vocabulary.
 - Is of the size V, size of the vocabulary (unique words in corpus).
 - Uses **softmax** function to output probabilities.

: Center Word
: Context Word
c=0 The cute cat jumps over the lazy dog.
c=1 The cute cat jumps over the lazy dog.
c=2 The cute cat jumps over the lazy dog.

- Training data is generated by iterating through the corpus using a **sliding window** action to **define the target word and its context words**.
- The window size (size of the sliding window) determines the number of before and words to consider when predicting the target word.

Working of CBoW

CBoW models the problem as a multi-class classification, wherein given the context words, the model outputs the target word and in doing so learns the representation for the words.

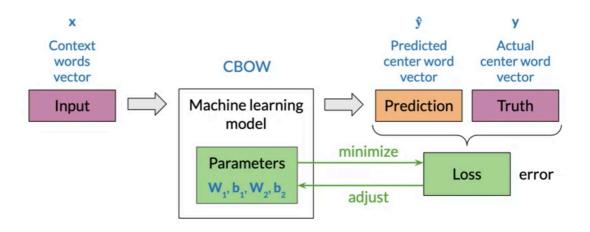


- One-hot encoded representation (V rows X 1 column) of the context word/words are sent in as input.
- Weight matrix **W**₁ is initialized.
- Weighted sum of the inputs and the weights (W₁) with bias b₁ is passed through ReLU activation function to produce output at hidden layer 'h'.
 - Weight matrix (W₁) is of shape N x V, where N is the dimension along which we want to represent our word vectors and V is size of vocabulary.
 - $z_1 = W_1 x + b_1$
 - h = ReLU(z₁) --> Output at hidden layer

- Similarly, weighted sum of output of hidden layer and weight matrix W₂ with bias b₂ is
 passed through softmax activation function to produce probabilities of all words in the
 vocabulary.
 - The weight matrix W₂ takes a shape of V rows and N columns.
 - $z_2 = W_2x + b_2$
 - $\hat{y} = \text{softmax}(z_2)$ --> Final output/prediction
- The word with the **highest probability** is considered as the model's prediction of the **target word**, i.e. **argmax(P(ŷ))**.
- The weight matrix \mathbf{W}_1 is **considered as the vector representation** of the words.

How does the model learn the right word representations?

- CBOW uses **cross-entropy loss** as the objective function during training
- During back-propogation, the network tries to reduce this cost function.



Generalized cost function for batches of training examples:

$$J_{batch} = -\frac{1}{m} \sum_{i=1}^{m} \sum_{j=1}^{V} y_j^{(i)} \log \hat{y}_j^{(i)}$$

Partial derivates of the cost function w.r.t weights and biases:

$$\frac{\partial J_{batch}}{\partial \mathbf{W_1}}, \frac{\partial J_{batch}}{\partial \mathbf{W_2}}, \frac{\partial J_{batch}}{\partial \mathbf{b_1}}, \frac{\partial J_{batch}}{\partial \mathbf{b_2}}$$

• Finally, the network parameters \mathbf{W}_1 and \mathbf{W}_2 are adjusted using gradient descent.

Is CBoW used in practice?

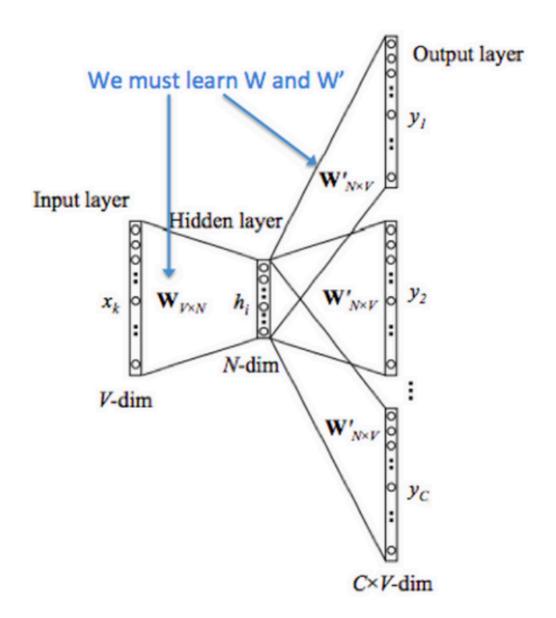
- When the context window > 1, CBoW **overfits on frequent words** as the input is an average of one-hot encoded vectors of the context word.
- CBoW does not produce good representations for rare words in the corpus.

We can over-come these dis-advantages of CBoW by using an alternative architecture of Word2Vec called the Skip-Gram.

What is Skip-Gram?

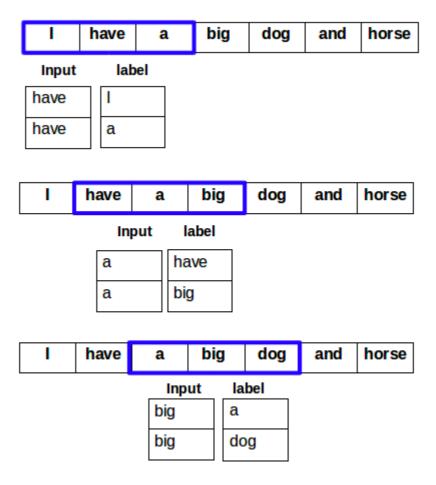
- Skip-Gram is simply an **inversion of the CBoW** architecture.
- Takes in input as center/context word and predict its surrounding words.

Architecture of Skip-Gram



How do we generate training examples for Skip-Gram?

- Similar to CBoW, we generate training examples by **iterating over the sentence**.
- Based on window size, single word is chosen as input and its surrounding words would be the target.



Working of Skip-Gram

- Input (x) is one-hot encoded representation of the center word of size |V| (vocabulary's size).
- Weighted sum of the input and weight matrix W_{VxN} is passed on through the hidden layer that has linear activation.
- Output layer applies a dot product between the output from hidden layer h₁ and W^{*}
 _{NxV} resulting in the output vector U.
- The **output vector U** is passed through **softmax activation function** to produce probabilities of observing each context word (based on window size) y'^{c-m} , ..., y'^{c-1} , y'^{c+1} , ..., y'^{c+m} .

How does Skip-Gram learn the right representations?

- Output probabilities are matched against true probabilities.
 - y^{c-m} , ..., y^{c-1} , y^{c+1} , ..., y^{c+m} , the one hot vectors of the surrounding/context words.
- Skip-Gram is trained with objective function of \log likelihood, i.e. for given center word ' w_i , predict context words within a fixed window of size m.

$$\frac{1}{T} \sum_{t=1}^{T} \sum_{-c \le j \le c, j \ne 0} \log p(w_{t+j}|w_t)$$

- Objective function tries to maximize the probability of any context word given the current word w_t .
- Model also makes a bayesian assumption that given the center word, all the output words are completely independent.
- Given the objective function, the model parameters **W** and **W'** are fine-tuned using gradient descent via backpropagation.
- Similar to CBoW, the **weight matrix** W_1 is considered as the **word vector**.

Negative Sampling for Skipgram training:

Refer: https://jalammar.github.io/illustrated-word2vec/ (https://jalammar.github.io/illustrated-word2vec/ (https://jalammar.github.io/ (https://jalammar.github.io/

Playground: https://projector.tensorflow.org/)

![image.png]

(data:image/png;base64,iVBORw0KGgoAAAANSUhEUgAAAmcAAADvCAIAAAA4rOTaAAAM

Now that we've understood how Word2Vec can help generate meaningful and contextual word embeddings, let us get back to solving our business problem.

Let us focus on only features that we'd need to build the semantic search engine

Out[21]:

	paper_id	title	abstract	
0	fe177de29f14dd45747a3d17c2468300245b0fb4	'I Get High With a Little Help From My Friends	Psychoactive drugs have been central to many h	10.3389/fp
1	ab312eb3286e189488707ea7ac8551401a178940	Arenavirus Stable Signal Peptide Is the Keysto	The rodent arenavirus glycoprotein complex enc	10.112
2	NaN	Two-Port Fetoscopic Repair of Myelomeningocele	OBJECTIVE The aim of this study was to assess	10
3	NaN	Learning Curves in COVID-19: Student Strategie	In New Zealand, similar to the rest of the wor	
4	NaN	Glass Fragment Injury to the Craniocervical Ju	BACKGROUND: Nonmissile penetrating injuries to	
4				•

We will look into the data and check if we have any null values.

```
In [ ]: df_covid.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 100000 entries, 0 to 99999
Data columns (total 4 columns):
   Column Non-Null Count Dtype
#
---
             -----
    paper id 35242 non-null object
0
    title 99946 non-null object
1
    abstract 77551 non-null object
2
             62094 non-null object
 3
    doi
dtypes: object(4)
memory usage: 3.1+ MB
```

Dropping duplicates

We will focus on only publications in English and drop anything that is non-English.

```
In [ ]: # set seed
        DetectorFactory.seed = 0
        # hold label - language
        languages = []
        # loop through each text
        for ii in tqdm(range(0,len(df_covid))):
            text = df_covid.iloc[ii]['abstract'].split(" ")
            lang = "en"
            try:
                if len(text) > 50:
                    lang = detect(" ".join(text[:50]))
                elif len(text) > 0:
                    lang = detect(" ".join(text[:len(text)]))
            except Exception as e:
                all_words = set(text)
                try:
                    lang = detect(" ".join(all_words))
                except Exception as e:
                    lang = "unknown"
                    pass
            # Appending to Language Label
            languages.append(lang)
```

0% | 0/29458 [00:00<?, ?it/s]

Let's look at the numbers of articles for each language.

```
In [ ]: languages_dict = {}
    for lang in set(languages):
        languages_dict[lang] = languages.count(lang)

    print("Total: {}\n".format(len(languages)))
    print(languages_dict)

Total: 29458

{'fr': 51, 'pt': 4, 'en': 29261, 'ca': 1, 'nl': 11, 'unknown': 2, 'de': 6
    2, 'ro': 3, 'hu': 1, 'it': 3, 'es': 59}
```

Since most of the publications are in English, we can safely drop the rest

We also do not need the language feature column.

```
In [ ]: df_covid = df_covid.drop(['language'], axis = 1)
    df_covid.head()
```

Out[27]:

	abstract	title	paper_id	
10.3389/fpsyg.2	Psychoactive drugs have been central to many h	'I Get High With a Little Help From My Friends	fe177de29f14dd45747a3d17c2468300245b0fb4	0
10.1128/mt	The rodent arenavirus glycoprotein complex enc	Arenavirus Stable Signal Peptide Is the Keysto	ab312eb3286e189488707ea7ac8551401a178940	1
10.1016/j.jviromet.	Coxsackievirus B3 (CVB3) infection has been fo	A rapid and quantitative assay for measuring n	f9e71114eecf03cedc124d3f670b22fdf3a476b4	8
10.1186/s13099-(BACKGROUND: Diarrhea remains the leading cause	Burden and etiology of moderate and severe dia	dc1559d0835d272fcc0c12043b27813c89251295	15
10.1038/s41581-	Chronic kidney disease is a major public healt	Educating primary healthcare providers about k	8de77b96a986a8f94fd68b022640ba0a53b4ac50	17
—				4 6

Use text pre-processing techniques to get rid of punctuations and stop words

```
In [ ]: punctuations = string.punctuation
    stopwords = list(STOP_WORDS)
```

Creating custom list of stop-words based on our corpus

Cleaning abstracts

```
def pre_processor(sentence):
                mytokens = sentence.split(' ')
                mytokens = [word.lower() for word in mytokens if word not in stopwords a
                mytokens = " ".join([i for i in mytokens])
                return mytokens
           df_covid["processed_abstract"] = df_covid["abstract"].progress_apply(pre_process_apply)
           df_covid.head()
              0%|
                              | 0/29261 [00:00<?, ?it/s]
Out[30]:
                                                                   title
                                                   paper_id
                                                                                abstract
                                                              'I Get High
                                                                            Psychoactive
                                                                 With a
                                                                              drugs have
                  fe177de29f14dd45747a3d17c2468300245b0fb4
                                                              Little Help
                                                                                            10.3389/fpsyg.2
                                                                          been central to
                                                               From My
                                                                               many h...
                                                               Friends...
                                                              Arenavirus
                                                                 Stable
                                                                              The rodent
                                                                  Signal
                                                                              arenavirus
             1 ab312eb3286e189488707ea7ac8551401a178940
                                                                                               10.1128/mk
                                                              Peptide Is
                                                                             glycoprotein
                                                                    the
                                                                           complex enc...
                                                               Keysto...
                                                             A rapid and
                                                                          Coxsackievirus
                                                             quantitative
                                                                              B3 (CVB3)
             8
                   f9e71114eecf03cedc124d3f670b22fdf3a476b4
                                                               assay for
                                                                                         10.1016/j.jviromet.
                                                                            infection has
                                                              measuring
                                                                               been fo...
                                                                    n...
                                                                 Burden
                                                                         BACKGROUND:
                                                                    and
                                                              etiology of
                                                                                Diarrhea
                 dc1559d0835d272fcc0c12043b27813c89251295
                                                                                          10.1186/s13099-0
                                                               moderate
                                                                             remains the
                                                             and severe
                                                                         leading cause...
                                                                  dia...
                                                              Educating
                                                                          Chronic kidney
                                                                primary
                                                                             disease is a
```

To train the Word2Vec model, we'll have to convert the sentences into list of words

healthcare

providers

about k...

10.1038/s41581-

major public

healt...

8de77b96a986a8f94fd68b022640ba0a53b4ac50

17

```
In [ ]: | abstracts = df_covid['processed_abstract'].values
        ## Let us use Spacy for faster tokenization
        nlp = spacy.load('en_core_web_sm', disable=["tagger", "ner"])
        nlp.add pipe('sentencizer')
        def tokenize_sentences(sentence):
            sentence_corpus = []
            doc = nlp(sentence)
            sentences = [sent.text.strip() for sent in doc.sents]
            for sent in sentences:
                processed_sent_list = sent.split(" ")
                sentence_corpus.append(processed_sent_list)
            return sentence_corpus
        df covid['tokenized abstract'] = df covid['processed abstract'].progress app
        corpus data = df covid['tokenized abstract'].to list()
        word2vec_corpus = [item for items in corpus_data for item in items]
          0%|
                       | 0/29261 [00:00<?, ?it/s]
```

Let us create our own Word2Vec model by training on our corpus

We will be using the **Skip-Gram** architecture along with **Negative Sampling** as the objective function.

Why Negative Sampling instead of Softmax?

- Denominator in softmax is a normalizing factor computed over entire vocabulary.
- For large vocabulary, softmax computation is highly expensive.

We over-come the above drawbacks of softmax using Negative-Sampling.

- Negative-Sampling maximizes similarity of words in the same context and minimize when they occur in different contexts.
- Instead of minimizing over all words, Negative-Sampling randomly samples handful of words (2 < k < 20) and use them to optimize the objective.
- Each **sample** picked **(word, context)** can be **represented** as a **probability** of whether or not this pair is **near each other** in training data.

$$\arg \max_{\theta} \sum_{(w,c)\in D} \log \sigma(v_c.v_w) + \sum_{(w,c)\in D'} \log (\sigma(-v_c.v_w))$$

- We **optimize** over these two **probability distributions** to learn word vectors.
- Large values of k are chosen for small dataset and vice-versa.

Using Gensim's API to train Word2Vec

Parameters:

- input : Corpus data
- min_count : Ignores all words with total frequency lower than this.
- size : Dimensionality of the output word vectors.
- workers: Use these many worker threads to train the model (=faster training with multicore machines).
- sg: Training algorithm: 1 for skip-gram; otherwise CBOW.
- **negative**: If > 0, negative sampling will be used, the int for negative specifies how many "noise words".

```
In [ ]: model = Word2Vec(word2vec_corpus, min_count=3, size= 100, workers=4, window
```

Now that we have word embedding for our corpus, we can measure the cosine distance between the centroid of each abstract and the query to find the similarity

Let us calculate the centroid for each abstract using the vectors of all the words incorporating the abstract.

```
In []: a = [0.0]*100
    df_covid["centroid"] = [a]*df_covid.shape[0]

for index, row in df_covid.iterrows():
    abstract = row['processed_abstract']
    total_sim = 0
    words = abstract.split(" ")
    centroid = np.array([0.0]*100)
    for word in words:
        try:
        b = model[word]
    except:
        continue
    centroid = np.add(centroid, b)

    df_covid.at[index,'centroid'] = centroid.tolist()
```

Let us create our base function, which when given a query will retrieve and rank documents based on similarity.

```
In [ ]: def rank_docs(model, query, df_covid, num) :
            cosine_list = []
            a = []
            query = query.split(" ")
            for q in query:
                    a.append(model[q])
                except:
                    continue
            for index, row in df_covid.iterrows():
                centroid = row['centroid']
                total_sim = 0
                for a_i in a:
                    cos_sim = np.dot(a_i, centroid)/(np.linalg.norm(a_i)*np.linalg.r
                    total_sim += cos_sim
                cosine_list.append((row['title'], row['doi'], total_sim))
            cosine_list.sort(key=lambda x:x[2], reverse=True) ## in Descending order
            papers_list = []
            for item in cosine_list[:num]:
                papers_list.append((item[0], item[1], item[2]))
            return papers_list
```

Method to use the base function and retrieve top matching documents

```
In [ ]: def query(query, top_matches=10):
    model_to_use = model
    df_covid_to_use = df_covid
    return rank_docs(model_to_use, query, df_covid_to_use, top_matches)
```

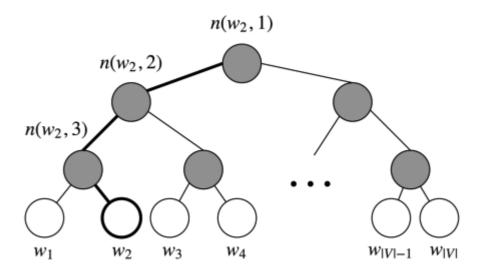
```
In [ ]: | query('origin of corona virus')
Out[37]: [('Role of Nonstructural Proteins in the Pathogenesis of SARS-CoV-2',
           '10.1002/jmv.25858',
           2.381886614774109),
          ('Re-emergence of a genetic outlier strain of equine arteritis virus: Imp
         act on phylogeny',
           '10.1016/j.virusres.2014.12.009',
           2.3744738115509545),
          ('Emerging and Neglected Viruses of Zoonotic Importance in Croatia',
            '10.3390/pathogens10010073',
           2.370784610584278),
          ('The global emergence of severe acute respiratory syndrome coronavirus 2
         in human',
           '10.1007/s13337-020-00613-y',
           2.369730359573394),
          ('An Introduction to SARS Coronavirus 2; Comparative Analysis with MERS a
         nd SARS Coronaviruses: A Brief Review',
           '10.18502/ijph.v49is1.3667',
           2.364636515916258),
          ('Virology features of a family cluster of SARS-CoV-2 infections in Shang
         hai, China',
           '10.1016/j.bsheal.2021.05.003',
           2.3634767705386515),
          ('Canine Distemper Spillover in Domestic Dogs from Urban Wildlife',
           '10.1016/j.cvsm.2011.08.005',
           2.3596838753219522),
          ('Next-Generation Sequencing for Porcine Coronaviruses',
            '10.1007/978-1-4939-3414-0_19',
           2.3580303371054683),
          ('Zwiesel bat banyangvirus, a potentially zoonotic Huaiyangshan banyangvi
         rus (Formerly known as SFTS)-like banyangvirus in Northern bats from Germa
         ny',
           '10.1038/s41598-020-58466-w',
           2.357634998523873),
          ('Update of the current knowledge on genetics, evolution, immunopathogene
         sis, and transmission for coronavirus disease 19 (COVID-19)',
           '10.7150/ijbs.48812',
           2.3548538950775884)]
```

Conclusion

- Continous text representations are used to capture syntactic and semantic similarity that, discrete text representations dont capture.
- SVD based methods suffer from scaling and in practice, iterative method like Word2Vec is used.
- Word2Vec models the data as multi-class classification problem and in doing so, learns the word representations.
- Two architectures of Word2Vec (i) **CBoW** and (ii) **Skip-Gram**.
- Training methods available (i) Negative Sampling and (ii) Hierarchical Softmax
- Gensim library is used for building Word2Vec from scratch.
- Finally, we built a search engine based on semantic similarity between query and abstracts of the publications.

Post Read

1. Hierarchical Softmax:



- Another training method to overcome the computation expense of softmax.
- Uses binary tree to construct vocabulary and leaves are the words.
- Each node of the graph is associated to a vector, which the model will have to learn.
- The probability of a word 'w' is equal to the probability of a random walk starting in the root and ending in leaf node corresponding to 'w'.
- With a goal to maximize likelihood (or minimize negative log-likelihood), in hierarchical softmax, the vectors of the nodes in the tree are updated that are in path from root to leaf node.

2. How are the weights adjusted in Word2Vec?:

Adjusting Weight Parameters (https://thinkinfi.com/continuous-bag-of-words-cbow-single-word-model-how-it-works/)

3. Enhancements over Word2Vec

- <u>Introduction to GloVe (https://towardsdatascience.com/light-on-math-ml-intuitive-guide-to-understanding-glove-embeddings-b13b4f19c010)</u>
- Introduction to FastText (https://amitness.com/2020/06/fasttext-embeddings/)