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Extended Abstract

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ABSTRACT

The Skip-Gram Model with negative sampling (SGNS) is a simple and effective algorithm to create word embeddings. SGNS uses Stochastic Gradient Descent (SGD) as its optimizer. While a lot of effort has gone into increasing the throughput of words of the SGNS, not much work has gone into optimizing the convergence time. Therefore, our work focuses on the latter. We propose a new batched version of the SGNS. To improve the quality of our batched version, we only allowed each word to appear once in a batch. Furthermore we deleted every word that appear more frequently than the 99 percentile in our datasets. Finally we also used advanced optimizers and input shuffling to decrease the convergence time. Here are the components used to evaluate our work: to train our model we used the text8 dataset - to measure the quality of our word embeddings we used the wordsim353 dataset, which assess that similar words have vector representations that are close to each other. The advanced optimizer Adam combined with our batched approach outperformed the current state of the art implementation Gensim. Adam converged to a similarity value of 0.66 (state of the art) in 1 epoch, while Gensim took 4 epochs. Hence, this work shows that our batched version with advanced optimizers, and the deletion of outliers do decrease the convergence time of the SGNS. These results must be confirmed with other datasets, but optimizing our model in terms of throughput has the potential of further reducing the overall run time of the SGNS.

CCS CONCEPTS

• **Computer systems organization** → **Embedded systems**; *Redundancy*; Robotics; • **Networks** → Network reliability;

KEYWORDS

ACM proceedings, L^AT_EX, text tagging

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1 INTRODUCTION

Representing words as vectors, i.e word embeddings (WE) is a fundamental aspect of Natural Language Processing (NLP). There are two ways to create such WE, either arbitrarily or with the purpose of capturing the semantic of the words, i.e. vector representations of words that are syntactically or semantically similar will be close to each other in the vector space. By capturing semantic or syntactic meaning WE have shown to facilitate a lot of subsequent NLP tasks, such as entity recognition, machine translation or sentence classification. The first attempt to create WE with neural networks was made by Bengio et al. [?] but more recently Mikolov et al. [?] introduced a software package called w2vec, which uses a simpler network that produces state of the art results. One of the proposed algorithms in this software package is the Skip-Gram Model (SGM). The SGM is an algorithm, that trains a neural network, on the task of predicting the neighboring words in a sentence. The weights of this network are then used as WE. Mikolov et al. [?] then introduced an alteration to the SGM called negative sampling. The goal of the Skip-Gram Model with negative sampling (SGNS) is to distinguish for a given word w , context words c_i (words that appear close to w in a sentence) from words drawn randomly (i.e negative samples) k_i . This is achieved by maximizing the dot product of w and c_i and minimizing the dot product of w and k_i .

The SGM, espacially the SGNS, gained a lot of attention, as it achieved very good results for a very simplistic model. As a consequence, a lot of effort went into optimizing it. Most of this effort was trying to improve the throughput of the model, i.e the number of words that are processed per second. As remarkable progress was made [?], [?]. Both of these approaches and most of the literature are focused on improving the throughput of the model, but not the convergence time. Therefore one could ask if the convergence time of the SGM can be optimized while at the same time maintaining its accuracy.

Our work focused on improving the convergence time of the SGNS. To facilitate our calculations we propose a slightly altered version of the original SNGS, in which we compute the loss for a very large amount of training samples, i.e 2000. The loss of such a batch was computed by taking the sum over the loss of all individual training samples in the specific batch. We combined this batched approach with advanced optimizers and input shuffling to decrease the convergence time of the SGNS. Furthermore we analyzed the distribution of word frequencies of different datasets to create batches of better quality. As a matter of fact a word that appear multiple times in a batch will hinder the quality of the batch. Therefore we focused our work on creating batches that only contained words once. In combination advanced optimizers, input shuffling and our batched approach allowed us to decrease the convergence time of

the SGNS. If combined with an optimized throughput these results could lead to an overall decrease in runtime.

The rest of the paper is structured as follows: section 2 describes the SGM and the SGNS followed by its optimizations in Section 3. The reader is introduced to our Implementation in Section 4. Furthermore we will describe the used datasets, the measure applied to compare the quality of word embeddings and input shuffling in Section 5. Results are presented in Section 6. The last part will focus on the discussion of our results, and possible future work in Section 7 followed by a conclusion in Section 8.

2 BACKGROUND

This section gives in its first subsection a broad overview over the functioning of the Skip-Gram Model, and its second section describes the Technique called Negative Sampling introduced by Mikolov et al. [?] used to make the Skip-Gram model more efficient

2.1 The Skip-Gram Model

The SGM is a very simple model used to learn WE. The idea is to train a neural network, on a fake task and then use some of the weights of the network as embeddings. To understand this fake task center and context words must be defined. The center word is any given word in a sentence, from which we want to learn the WE. The context words of this specific center word are the words left and right in a given window m in the sentence. See Figure 1 as an example, where the context words are highlighted. An important aspect to notice, is that instead of having a fixed window size m , each word will randomly have a window size between 1 and m . The idea behind having different window sizes is that words that are further away in the sentence, have less semantic correlation to the center word.

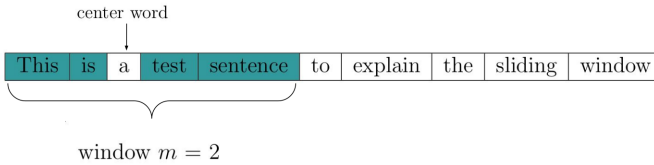


Figure 1: Example of a center word and its context words

With these definitions the fake task can be defined: given a pair consisting of a center and a context words, which will be our training samples, the goal of the network is to predict the probability of each word to appear in the context of the center word. To achieve this goal, Mikolov et al. introduced the following model. The network consists of three layers. See Figure 2 for an illustration. First the network has an input layer, a vector of size V (V being the size of the vocabulary), represent words as an one hot encoding. Secondly, the projection layer weight Matrix M_{in} stores V D -dimensional vectors for the corresponding word in the vocabulary in its rows. The projection layer serves as a look up table for our WE (if a one hot vector is multiplied with M_{in} the corresponding WE will result). Finally, the output layer weight matrix M_{out} also stores V D -dimensional vector representation of words. The idea behind these 2 Matrices is that the projection layer

will represent our words as center word and the output layer as context words.

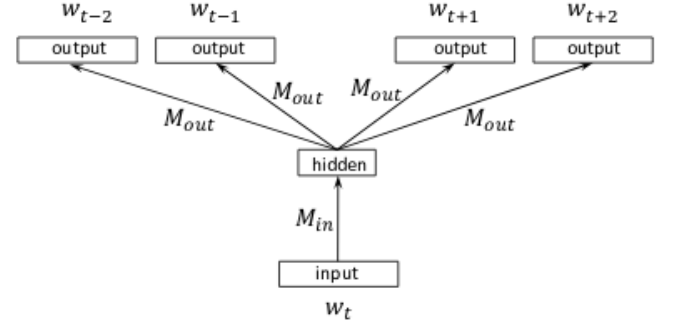


Figure 2: Structure of the network of the Skip Gram Model

Since the task of the network being to predict the probability of each word appearing in the context of a given center word, the following probability should be maximized:

$$\prod_{i=1}^V \prod_{-m < j < m} p(w_{i+j}|w_i) \quad (1)$$

Where V is the number of words in the corpus data, w_i the i^{th} word in the corpus data and m is the context window. This means that the m nearest words to w are considered as context words. Equation 1 can be transformed into sums by using log probabilities:

$$\sum_{i=1}^V \sum_{-m < j < m} \log(p(w_{i+j}|w_i)) \quad (2)$$

where the parameters are the same as in Equation 1.

The basic Skip-Gram Model uses a classical Softmax to calculate the conditional probability $p(w_{i+j}|w_i)$:

$$p(w_{i+j}|w_i) = \frac{\exp(\tilde{v}_{w_{i+j}}^T v_{w_i})}{\sum_{w=1}^V \exp(\tilde{v}_w^T v_{w_i})} \quad (3)$$

Here v_{w_i} and \tilde{v}_{w_i} are the vector representations, stored respectively in the projection Layer (M_{in}) and in the output layer (M_{out}). There is a problem with the classical softmax. For the computation of $\sum_{w=1}^V \exp(\tilde{v}_w^T v_{w_i})$, the denominator in Equation 3, one has to go over the whole corpus data. As very big data sets are needed to train the model, this approach is unsuitable. Mikolov et al. [?] proposed different solutions. The first solution is to use a Hierarchical softmax introduced by Morin and Bengio [?]. In this model, the probability distribution of the output nodes is saved in a binary tree which gives one a logarithmic computation time for each of these probabilities and makes it suitable to compute the softmax. Another possibility is the use of negative sampling which is discussed in the next subsection.

2.2 Negative Sampling

A second alternative solution to the use of a classical softmax is negative sampling. Negative Sampling is a simplification of Noise Contrastive Estimation (NCE) which was introduced by Gutmann

and Hyvärinen [?], and first applied to NLP by Mnih and Teh [?]. This subsection will shortly describe NCE and then how Mikolov et al. [?] simplified it to create the technique called negative sampling. The idea behind NCE is to distinguish data from noise. It does so by reducing the problem to a logistic regression task and does it by maximizing the log probability. The SGM is only interested in good word representation, hence the probability of the word is not meaningful as long as the quality of the word representations remains high. Mikolov et al. [?] simplified NCE and called it Negative Sampling, which will be explained next.

The main goal of negative sampling is to only update the output nodes of certain words, 5-20 in practice. This will obviously save an enormous amount of computation time. The idea is that given a pair $(c, w) \in D$, where c is a word in the context window of w , we select K random words k_i from the corpus data D . We assume those words do not appear in the context of w . By doing so we will only have to update $k + 1$ nodes. Mikolov et al. [?] introduced the following objective function J , that should be maximized:

$$J(c, w) = \log(\sigma(v_c \tilde{v}_w)) + \sum_{k \in K} \sigma(\log(-v_c \tilde{v}_k)) \quad (4)$$

Where v_c and \tilde{v}_w , can be interpreted as before in Equation 3 and $\sigma(x) = \frac{1}{1+e^{-x}}$. To understand why this loss function is working so well, one needs to assume that two vectors are similar if their dot product is high. Therefore, we maximize the dot product of similar words (w, c) (c appears in the context of w) and minimize it for dissimilar words (w, k_i) (those were selected randomly, we therefore assume that they are not similar). We see that to compute our objective function we will only have to compute the sum over the number of negative samples K , which is very small in practice (2-20). To put things in perspective let's imagine our data set consists of 100000 words, we set $K = 2$. Assume each output neuron has a weight vector of dimension $d = 300$. In consequence, when updating our weights we would only update $0.2 * 10^{-2}$ of the 300 million weights in the output layer.

One question remains: how are the negative samples selected? Mikolov et al. [?] used the following unigram distribution, to define the probability $P(w)$ of a word w being drawn as a negative sample:

$$P(w) = \frac{f(w)^{\frac{3}{4}}}{\sum_{t=0}^T f(w_t)^{\frac{3}{4}}} \quad (5)$$

where $f(w)$ is the frequency of w_t . The value of $\frac{3}{4}$ is set empirically. Raising the unigram distribution to the power of $\frac{3}{4}$ makes it less likely for a word to be drawn if it often appears in the dataset in comparison to the basic unigram distribution. With all of the above knowledge it's understandable that the SGNS will outperform the classical softmax in computation time. Now the question arises if the accuracy is maintained but according to Mikolov et al. [?] the negative sampling method "is an extremely simple training method that learns accurate representations". As a matter of fact, Mikolov et al. [?] reported a 6% accuracy improvement in comparison to a Hierarchical Softmax model. Therefore, it is a good solution to the problem raised by the classical softmax. We now have enough

background knowledge about the SGM to look at how it can be optimized. The next section introduces earlier approaches to optimize the SGM.

3 RELATED WORK

In the original model, the optimization is done with Stochastic Gradient Descent (SGD), which is a sequential algorithm. This process does not favor parallelization. To deal with this specific problem Mikolov et al. [?] used a Hogwild! approach proposed by Recht et al. [?]. The idea is to allow multiple threads to access a shared memory, in this case, the single model. In the original SGM, the threads are constructed as follows: at the beginning of training, the dataset is split into N evenly sized chunks and each of these chunks will be processed by a one thread. The threads run parallel and have access to the shared memory. Therefore, overwriting errors are bound to happen. According to Recht et al. [?] the overwriting errors won't lead to a significant accuracy loss if the data is sparse enough. But in the case of NLP, the problem seems to be a bit more significant, and especially for word embedding, as many words share the same context words. There were several attempts at solving this issue, and optimize the throughput of the SGM.

3.1 Parallelization by the use of caching

This idea was proposed by Vuurens et al. [?]. The architecture used here is the SGM with a hierarchical softmax. The general idea is to cache the most frequently used weight vectors, and update the model after a certain number of seen training samples, while taking in consideration the changes that happened during these number of seen training samples (the paper used the number 10). The unfrequent weight vectors are handled as in the original paper, i.e. updated in shared memory after each training sample. This technique allows it to avoid collision for the most frequent words, and the author therefore hoped to be able to increase the number of concurrent threads. The paper produced interesting results as they managed to decrease execution time by increasing the number of cores used for the calculation. This is very powerful because in the original implementation the execution time regressed after using more than 8 cores. It seems to indicate that too much overwriting was happening, as the number of concurrent threads surpasses a certain threshold. The model did not suffer any accuracy loss in comparison to the original SGM model.

This work proposes a very good way to parallelize the SGM, as in particular, it allows using more cores during the computation. As this approach focused on the Hierarchical softmax, in contrast to our work which uses negative sampling, the next subsection covers optimizations of the Skip-Gram Model with negative sampling (SGNS).

3.2 Parallelization in shared and Distributed Memory

In contrast to the above approach, this one, introduced by Ji et al. [?] focused on the SGNS. The focus of this work was to speed up the computation of the loss function of the SGNS described in Equation 2.2. Where the dot product of the center word with negative samples

as well as the context word is computed. The dot product is a level 1-BLAS operation and is not as efficiently implemented into hardware, as matrix multiplication which are a level 3-BLAS operation. To achieve such a transformation, Ji et al. decided to use the same negative samples for given amount of training samples. The training samples that will have the same negative samples are chosen as follows: given a word w_c one can construct all the training samples where w_c is a context word. The following training batch X will result: $X = \{(w_{in}^i, w_c) | w_c \text{ appears in the context of } w_{in}^i\}$. The matrix multiplication can be represented in the following way:

$$\begin{bmatrix} w_c \\ w_{n_1} \\ \vdots \\ w_{n_k} \end{bmatrix} * \begin{bmatrix} w_{in}^1 \\ \vdots \\ w_{in}^{2m} \end{bmatrix}$$

where, $w_{n_1} \dots w_{n_k}$ are the shared negative samples. The above Matrix multiplication will compute the loss for each training sample and the weights of the used vectors will be updated after each batch. Ji et al.'s method achieves a 3.6 fold increase in throughput, by only losing 1

3.3 Accelleration of word2vec by using GPU's

Seulki and Youngmin [?] focused on getting a better throughput on the SGM with the use of GPU's. As the SGM is a sequential algorithm, it is not easy to parallelize it, especially if one wants to parallelize the training of individual training samples. As the algorithm goes sequentially over a sentence, the samples next to each other, in order of execution, will almost every time have the same input word. Consequently, it's very hard to parallelize at this level. Since the dimensions of the word embeddings are completely independent of each other Seulki and Youngmin [?] tried to parallelize their update. They achieved this by mapping each dimension to a CUDA thread while mapping each sentence to a CUDA block. As each CUDA block runs independently, the training of the sentences is parallelized, and the fact that sentences have different length is of no problem. If the execution time of the GPU kernel is greater than the time used to read the sentences, it could be a smart choice to use multiple GPU's. According to Seulki and Youngmin [?], if multiple GPU's are used, there is a need for synchronizing the model, which will hinder run time performance. They achieved their best results with 2 concurrent GPU'S. The acomplished results were very good as they managed a 20x speedup compared to a single threaded CPU execution, which is a 3x increase in comparison to the original C code from Mikolov et al. [?]. The speedup was achieved, while maintaining the accuracy of the model. The problem with this and all the above optimization is that the code is not easily available or of use. Therefore, we need an optimized implementation of the SGM that is easily available. This is provided by Gensim [?], which will be outlined in the next subsection.

3.4 Gensim

Gensim [?] is a pure Python library that holds a state of the art implementation of the SGM. Gensim is written in Cython, which first allowed Gensim to have the same runtime as the original C code. Furthermore, it made use of BLAS's and precomputed sigmoid tables, while also trying to parallelize the training of different

sentences. This finally yielded in a 4x speedup in run time compared to the original C code from Mikolov et al. Gensim as a python library, and because it was already used in related work [?] made the comparison easy and of value.

This concludes our overview of related work. The next Section will cover our implementation of the Skip-Gram Model with negative sampling.

4 BATCHED SKIP-GRAM MODEL

This section will give an overview of our implementation. The proposed implementation is a slightly altered version of the original Skip-Gram Model with negative sampling (SGNS). The idea behind this altering, is to compute the loss for multiple words and context pairs at the same time, this is inspired from mini-batch training, a model often used in other machine learning problem. Instead of computing the loss function of each pair, we computed the sum over a certain number of pairs (i.e 2000) as a loss. The exact process will be described in the following paragraph.

4.1 Forwarding

As the computation of the new loss function represents the challenging part of our implementation the forwarding method is explained step by step. Each time step is illustrated to make the explanation clearer.

Let $X = (v_1, c_1), (v_2, c_2), (v_3, c_3)$, be the training batch, where (v_i, c_i) is a training sample constituted by a word and one of its context words.

4.1.1 Input. The forwarding method will accept two vectors v and c , and a matrix A as an input. The first vector represents all the center words in a batch, the second one the context words. The Matrix represents the negative samples. The two vectors are of the same length, defined as n . The matrix must be of dimension $n \times k$, with k being the number of negative samples per pair. This means the i^{th} row will store the negative samples for the i^{th} word context pair.

The input can be illustrated as follows:

$$v = \begin{bmatrix} v_1 & v_2 & v_3 \end{bmatrix}, c = \begin{bmatrix} c_1 \\ c_2 \\ c_3 \end{bmatrix} \text{ and } A = \begin{bmatrix} k_{1,1} & k_{2,1} & k_{3,1} \\ k_{1,2} & k_{2,2} & k_{3,2} \\ k_{1,3} & k_{2,3} & k_{3,3} \end{bmatrix}$$

4.1.2 Concatenation of samples. The concatenation of the vector c and the Matrix A will result in a Matrix \tilde{A} , with:

$$\tilde{A} = \begin{bmatrix} c_1 & k_{1,1} & k_{2,1} & k_{3,1} \\ c_2 & k_{1,2} & k_{2,2} & k_{3,2} \\ c_3 & k_{1,3} & k_{2,3} & k_{3,3} \end{bmatrix}$$

4.1.3 Embeddings. Now it's necessary to access the word embeddings. For this purpose a Matrix E_v of dimensionality $n \times d$ where d is the dimension of our word embedding is created. E_v stores the word embedding of the i^{th} word from our input vector v in it's i^{th} row. The same is done for the other Matrix \tilde{A} . This will result in a $n \times k + 1 \times d$ Array E_c .

$E_v = \begin{bmatrix} \tilde{v}_{11} & \dots & \tilde{v}_{1d} \\ \tilde{v}_{21} & \dots & \tilde{v}_{2d} \\ \tilde{v}_{31} & \dots & \tilde{v}_{3d} \end{bmatrix}$, where $\tilde{v}_i = [\tilde{v}_{i1} \dots \tilde{v}_{id}]$ is the embedding of v_i .

$E_c = \begin{bmatrix} \tilde{c}_1 & \tilde{k}_{1,1} & \tilde{k}_{2,1} \\ \tilde{c}_2 & \tilde{k}_{1,2} & \tilde{k}_{2,2} \\ \tilde{c}_3 & \tilde{k}_{1,3} & \tilde{k}_{2,3} \end{bmatrix}$, where each entry of the matrix is a vector of dimension d

4.1.4 Batch multiplication and negation of samples. Now we need to compute the dot product of each word vector with its pair and the negative samples, exactly as done as in the original loss function of Mikolov et al. shown in Equation 2.2. For this task some definitions are necessary: let A_j be the j^{th} row of the matrix A , let $E_c(i, j)$ be the d dimensional embedding of the word stored in $\tilde{A}(i, j)$. The dot product is computed with a so-called batch multiplication¹ which will result in a matrix S where $S(i, j) = E_c(i, j) \cdot E_{v_j}$. This will result in a $n \times k + 1$ Matrix S . Now we only have to negate the last k rows. The sum of each row represents the loss computed in Equation 2.2, for each word context pair. Since computation time is too long with this approach the loss function is altered

$$S = \begin{bmatrix} \tilde{v}_1 \cdot \tilde{c}_1 & -\tilde{v}_1 \cdot \tilde{k}_{1,1} & -\tilde{v}_1 \cdot \tilde{k}_{2,1} & -\tilde{v}_1 \cdot \tilde{k}_{3,1} \\ \tilde{v}_2 \cdot \tilde{c}_2 & -\tilde{v}_2 \cdot \tilde{k}_{1,2} & -\tilde{v}_2 \cdot \tilde{k}_{2,2} & -\tilde{v}_2 \cdot \tilde{k}_{3,2} \\ \tilde{v}_3 \cdot \tilde{c}_3 & -\tilde{v}_3 \cdot \tilde{k}_{1,3} & -\tilde{v}_3 \cdot \tilde{k}_{2,3} & -\tilde{v}_3 \cdot \tilde{k}_{3,3} \end{bmatrix}$$

4.1.5 Loss function. Summing the matrix S and multiplying it with -1 (to make the problem a minimizing problem) results in the loss for our entire batch. As some words may appear more than once in the batch this will more be an average of it than as with Mikolov et al. [?] the exact update per pair.

$$L = - \sum_{(i,j) \in k \times n} S(i, j)$$

This finishes the description of our batched approach.

4.2 Optimization of the quality of batches

There are two main ways to create a batch. Either arbitrary or with specific rules. Both of these approaches have benefits, the first one do not need any preprocessing and the second should in theory provide better learning. In our case the quality of a batch is dependent on the number of multiple appearances of words in the batch. The more frequently a word appears in a batch, the more the quality of the word embedding diminishes. Therefore the next sections will go into details on how we constructed our batches.

4.2.1 Analyzing word frequencies. For this matter we chose to analyze the distribution of word frequencies. A box plot of the distribution, on the text8 dataset, can be found in Figure 3. This shows that very few words appear very frequently, and are therefore responsible for the building of the most context pairs. For example the word *the* creates 50% of the pairs in the text8 dataset. In consequence if we create arbitrary batches, most words will appear multiple times in a batch and the learning will be hindered. A solution would be to only allow words to appear once in a batch, in the

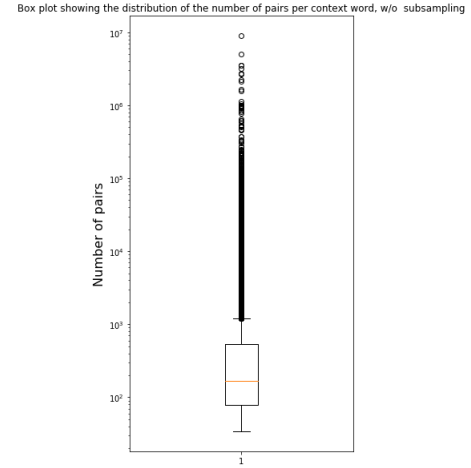


Figure 3: Distribution of word frequencies in the text8 dataset

next subsection we present such a solution with a variable batch size.

4.2.2 Criteria for batches. To solve the above problem we only chose to block multiple appearance in a batch on the center words. The context words can be the same for two different center words. Each batch will contain the highest possible number of pairs where each center word does not appear twice. As with all batches, each pair will only appear once in a batch. Therefore the batch size will be very high at the beginning of the training and will then decrease very rapidly. In consequence the average batch size is small and training time is increased, in comparison to arbitrary created batches. To remedy this problem we tried to reduce the number of very small batches by altering the dataset, while maintaining the quality of learning. We call it, deletion of outliers and explain it in the next subsection.

4.2.3 Deletion of outliers. To solve the above matter, we looked at the distribution of word frequencies (Figure 3). We experimented with different settings (mean, median, average, outlier) and found that deleting all words that had a frequency greater than the 99 percentile was best. This technique drastically reduces the data set size, it also deletes words that appear very frequently, it therefore reduces the number of very small batches.

The next section will describe our experimental setting and the possibilities that exists to evaluate the quality of word embeddings.

5 EVALUATION

This section gives an overview of the used datasets, the used metric to evaluate our models, the configuration of our model, and finally an explanation about input shuffling

5.1 Dataset

In this implementation we only used the text8² dataset. We chose this dataset for two reasons. First of all, it's a small dataset, that

¹Documentation: <https://pytorch.org/docs/stable/torch.html#torch.bmm>

²<http://mattmahoney.net/dc/enwik8.zip>

allowed us to do a lot of computations. Secondly, this dataset was used in related work [?] hence giving us a very good benchmark. The text8 dataset consists of 1702 lines of 1000 words, with a vocabulary of roughly 63000 words. Conveniently, there is no punctuation in the dataset. Therefore, we had to choose between building arbitrary sentences and keeping the dataset as it is. We chose the first option because it gives us a faster computation time, and did not show any significant loss in quality empirically. We chose a length of sentences of 20. Furthermore, we applied a technique called subsampling to reduce the data set size, which is explained in the following subsection.

5.1.1 Subsampling. Subsampling is a technique introduced by Mikolov et al. [?] to reduce the size of the dataset while at the same increasing the quality of the word embeddings (WE). The idea behind subsampling is the removal of very frequent words such as: "the, as, it, of". These words do not give an intrinsic value to the words that appear in its context. Therefore, the goal of subsampling is to delete such words from the dataset. Subsampling will decrease the computation time, as it will reduce the number of training samples, and increases the accuracy of the model. Details about the probability of the deletion of a word can be found in the original paper from Mikolov [?].

5.1.2 Min count. Another idea introduced by Mikolov et al. [?] is to delete every word that appear less than 5 times and is a good technique because of three reasons: First certain words of our data sets do not appear in a common lexicon (twigleg, melqu), or are names and acronyms. Secondly, each document often has spelling mistakes. Lastly, a word that only appears one time in our dataset will be very dependent on its original initialization. For all the above reasons, we applied this technique. Similar to subsampling, it will decrease computation time and should in theory improve the quality of the word embeddings.

5.2 Evaluating word embeddings

Evaluating word embedding is not an easy task, such as evaluating the accuracy of a common classifier for example. We cannot split our data set into train and test set. As the task that the network is learning is of no interest to us. Therefore, to ensure the quality of word embeddings other techniques have to be used. To define quality we first need to define a measure of similarity between two vectors. For this task we will use the cosine distance. Two vectors will be judged similar if they point in the same direction. To measure the quality of our word embedding, in question of word similarity we must use a dataset to compare our self too. The next section describes the chosen dataset.

5.2.1 wordsim353. We chose wordsim353³ for this task, as it's the most used in the related literature. The data set consists of 353 pairs of words rated by humans on their similarity. The similarity score is in the range of 1 and 10, with 10 being the best score. An example for two of such pairs can be found in Table ?? . We will rank our embeddings on the Pearson correlation coefficient between the cosine distance and the scores attributed by humans, as this is the common procedure. This concludes our description

of the evaluation methods of word embeddings. The next section will present the configuration of our network used to optimize the SGNS.

5.3 Configuration of the network

The SGNS has a lot of possible parameters, that can be tuned. We experimented with different models and finally decided for one that we tried to optimize. The network was configured the following way:

- Negative Samples: - 10
- Context Window: - 5
- Dimension of the embedding: - 100
- Batch size: - 2000
- Alpha: - (1e-5,1)

This concludes our overview of the possible parameters of our model of the SGNS.

To summarize the evaluation, we deleted words that appear very often (subsampling) and very infrequently (min count) from our dataset, i.e text8 dataset. To evaluate the resulting word embeddings we defined the cosine distance, which is 1 minus the cosine of the angle of two vectors. We will measure the quality of the embeddings by analyzing the correlation of the cosine distance of pairs of vectors to humanly judged similarity score of the corresponding words, and used the wordsim353 to collect these scores. The reader was also given an explanation about the configuration of our network and introduced to the concept of input shuffling. With all this knowledge we can now present the achieved experimental results in the next section.

6 RESULTS

As stated in the previous section we ran multiple experiments for each optimizer with the above configuration and datasets. . This Section will give an overview of the experimental results. Each subsection will give an explanation over the achieved result with a specific optimizer.

6.1 Simple Batch Version

6.1.1 SGD. The first challenge for each optimizer was to find a correct learning rate. As SGD is the optimizer used in Gensim [?] we first tried the same learning rate as the default value in Gensim [?], i.e 0.01, and then performed a random search to find a better one. As expected a bell curve shape resulted, a learning rate that is too high leads to diversion and a learning rate that is too low leads to a training time that is too slow. The best value that we found for the learning rate is 0.0075. With this setting SGD converged in 11 epochs. The second experiment was to add input shuffling. As seen in Figure 4, for almost every learning rate the convergence time decreased. Our model, with the best setting, now converges in only 7 epochs. Another interesting fact to point out from Figure 4 is that with input shuffling we achieved better results with higher learning rates. As for learning rates of 0.01 and 0.025 we did converge in 11 epochs with input shuffling but did not converge in 20 epochs without it.

6.1.2 Adam. Adam is the most advanced of all the optimizers used in our experiments and did yield the best results as seen in Figure

³<http://www.cs.technion.ac.il/~gabir/resources/data/wordsim353/wordsim353.zip>

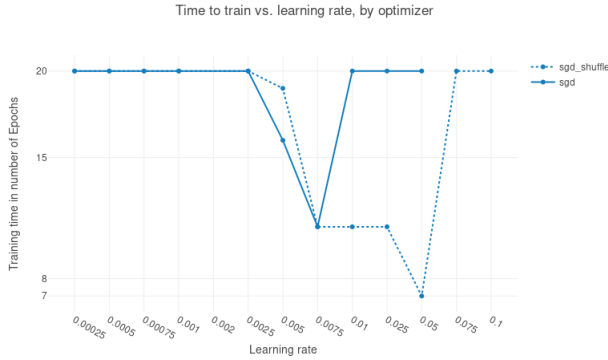


Figure 4: Training time Stochastic Gradient Descent with input Shuffling

5. Adam converged in 3 epochs without shuffling and a learning rate of 0.05 and 2 with shuffling and a learning rate of 0.001, as shown in Figure 5. Those are the best results achieved in any of our experiment. Exactly as Adagrad, Adam did not react to input shuffling the same way as SGD did.

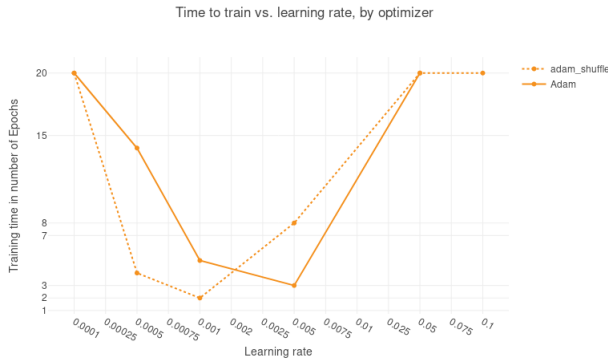


Figure 5: Training time Adam with input Shuffling

6.2 Advanced Batches Version

The advanced batched version is largely dependent on the definition of outlier. As shown in Table if the value for outliers is too small then the model convergence into score that is too small. If the threshold is higher training time is increased. We found that 99% is a good reference value. With this setting our model converged in 1 epoch for the text8 dataset, which is our best result, and 7 for the enwik9 corpus. A recap of all our results with the text8 dataset can be found in Table.

7 DISCUSSION

This section shortly discusses the empirical results and then extensively compares the findings of this work to the existing literature while trying to explain some of the differences. It is concluded by

Table 1: Results on the text8 dataset

| Model | Convergence Time | Word Similarity |
|--------------------------|------------------|-----------------|
| SGD | 11 | 0.65 |
| SGD w/shuffling | 7 | 0.66 |
| Adam | 3 | 0.66 |
| Adam w/ shuffling | 2 | 0.66 |
| Gensim | 4 | 0.66 |
| Adam w/o outliers | 1 | 0.66 |

a subsection describing the limitations and possible extensions of this work.

7.1 Our work

This subsection discusses our findings. First it will analyze the possible reasons behind the influence of input shuffling to the learning rate. Second it will conclude with the discussion of unexpected results.

7.1.1 Shuffling and learning rate with SGD. In this subsubsection we will use the term SGD for SGD, Momentum and NAG, and advanced optimizers for Adagrad and Adam. The models was able to use a higher learning rates when SGD optimizers where combined with input shuffling in comparison to as when not, as shown in Figures 4, ?? and ??. Therefore, arises the questions why these phenomena happen, especially as it did not happen with advanced optimizers.

First, we suspect our batched version of the Skip-Gram model with negative sampling (SGNS), to be at the origin of this phenomena. In consequence to this batched version, when input shuffling is not used a lot of words will appear multiple times in the same batch, as explained in Section ??. Therefore, the gradients will be an average of all the training samples. When the input is shuffled, less words will appear multiple time which will make the gradients more precise.

Second, advanced optimizers have a way to counter-attack this issue, namely adaptive learning rates, explained in Section ??. Adaptive learning rates counter attack the issue in the following way: high frequency words will appear more often multiple times in a batch, they will also have lower learning rates. Therefore the impact of appearing in the same batch will be reduced.

The two above arguments could explain why a higher learning rate combined with SGD achieved better results with input shuffling but not when advanced optimizers are used.

7.2 Comparison to Gensim

This subsection will compare our finding extensively to Gensim [?]. As explained in Section 3.4, Gensim is optimized to have a very high throughput, this made it possible to achieve a lot of computations. Furthermore, Gensim provides access to the loss and the resulting word embeddings, which facilitated the comparison process. This subsection is structured as follows: first it will describe the used configuration of Gensim, second it will compare Gensim to our implementation of the SGNS with the use of SGD and finally compare our model with Adam to Gensim. A graphic showing the comparison of our models to Gensim can be found in Figure 6.

7.2.1 Configuration of Gensim. To compare our-self in a correct manner we used the same dataset, with the same preprocessing parameters, i.e subsampling and min count. The hyper parameters of the network (window size, embedding size, number of negative sample, exponent to which the unigram distribution, which decides how a negative sample is used, is raised) are equivalent to our parameters described in Section 5.3. The only difference with our model is the learning rate. Gensim has a starting learning rate of 0.025 and linearly decrease it at every epoch to 0.0001.

7.2.2 Gensim vs. SGD. As stated earlier, we are not going to compare this work to Gensim in run time. Gensim is heavily optimized and written in cython⁴, which is 23x faster than plain Numpy. Since we used PyTorch the difference is not that big, but still remains. As shown in Figure 6 the convergence time was not the same between our implementation and Gensim. There are different possible reasons why this could be the case:

First, our batched approach could hinder performance in terms of convergence time since our loss function is not exactly the same. We compute the loss for multiple training samples by taking the sum over each score which is individually used as a loss by Gensim. Secondly, a difference to our implementation is the fact that Gensim checks whether negative samples are not equal to the context word. If that is the case Gensim selects a new random sample. Therefore, the learning of the input and output context is optimized.

Finally, another possibility is the decay of the learning rate used by Gensim. In fact, decaying the learning rate has been proven in a lot of work to decrease the convergence time. Gensim linearly decreases the learning rate, as we did not use this technique, the decay of the learning rate could help explain the noted differences. The first hypothesis, the fact that we used a batched approach, may be confirmed by the fact that when combined with input shuffling SGD does perform closer to Gensim, going from 11 to 7 epochs to converge, as input shuffling reduces the number of co-occurrence of the same word in a batch. Now the question arises if the 3 epochs, that Gensim is better, can be explained by the selection of better negative samples and the learning rate decay.

7.2.3 Gensim vs. Adam. The Adam optimizer did outperform the Gensim application in quality of word embeddings (only slightly: 0.01 correlation coefficient better) and convergence time. Adam converged in 2 epochs while Gensim in 4. To confirm the achieved results we ran each computation 40 times. The results can be seen in Figure 6.

7.3 The quality of batches can be optimized

As shown in Table 1, the creation of batches where each word only appears once in a batch outperforms the simple batch version. This shows empirically that our suspicion that words that appear multiple time in a batch hinder performance. Furthermore the deletion of outliers, speeds up training without hindering performance. A reason could be that words that appear very often have mostly the same context words and therefore will have a very similar representation as they learn from the same words.

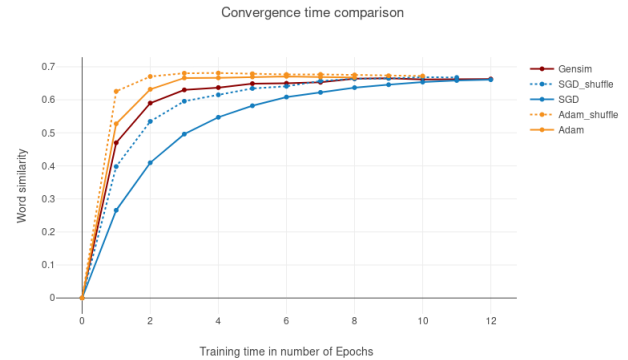


Figure 6: Convergence time of SGD and Adam compared to Gensim

7.4 Future Work

This work shows that the convergence time of the SGNS could be improved by using input shuffling, advanced optimizers and a batched version with specific rules to the creation of the batches. As with every work, there still exists possible extensions. First and foremost an aspect of our implementation that can be prejudicial is that we only extensively tested our model with one small dataset. The fact that we only used one dataset as well that it's a small one is problematic.

Problem with a small dataset:

By using a very small dataset we do not use the model in the condition it is most needed for, as the dataset used in practice usually consists of more than 1 billion words. There is a small argument that can be made for machine translation as the use of small parallel corpus is not unusual in this field.

Problem with using only one dataset:

It has been shown that some optimizers perform better with specific shapes of loss functions. To make a compelling argument it's necessary to show that our model with the use of input shuffling and Adam as its optimizer also outperforms Gensim with other data sets.

Finally, as it wasn't the goal of our implementation to outperform Gensim in run time, one could improve an already existing, optimized version, with input shuffling and advanced optimizers and should achieve a better run time than Gensim.

8 CONCLUSION

This work provides an overview of the Skip Gram Model with negative Sampling (SGNS) and the numerous successful attempts of optimizing the throughput of the model. As this is the case, no effort went into optimizing the convergence time of the SGNS, therefore this work focused on the convergence time. We decided to use advanced optimizers, input shuffling and a batched approach with specific rules for the batch creation to achieve a better convergence time. We propose a batched version of the SGNS, where the idea is to compute the loss over the sum of a high number of training samples, instead of computing it for each individually. The quality of each individual batch is hindered by words appearing multiple

⁴<https://rare-technologies.com/word2vec-in-python-part-two-optimizing/>

time in a batch, we therefore decided to create batches where this would not be the case. Furthermore we decided to cut the dataset size by deleting each word that appears more often than the 99 percentile, this technique maintained the quality of the dataset, while drastically improving the computation time. We used the text8 dataset to train our model and word similarity as a quality measure for the word embeddings (WE). To compare our work, Gensim was used, which holds a state of the art implementation of the SGNS. We did achieve a better convergence time than Gensim with Adam as an optimizer and the use of input shuffling. Gensim converged in 4 epochs to a word similarity of 0.66 and our model only took 2 epochs to achieve the same quality. The latter model was improved by deleting each word that appears more often than the 99 percentile and using batches where words only appear once. Those results still need to be confirmed with more datasets. Finally, if this work would be combined with an optimized throughput it could improve the state of the art run time of the SGNS.