500+ Times Faster Than Deep Learning

(A Case Study Exploring Faster Methods for Text Mining StackOverflow)

Suvodeep Majumder, Nikhila Balaji, Katie Brey, Wei Fu, Tim Menzies
Computer Science, NC State, USA
smajumd3,nbalaji,kebrey,wfu@ncsu.edu;tim@menzies.us

ABSTRACT

Deep learning methods are useful for high-dimensional data and are becoming widely used in many areas of software engineering. Deep learners utilize extensive computational power and can take a long time to train—making it difficult to widely validate and repeat and improve their results. Further, they are not the best solution in all domains. For example, recent results show that for finding related Stack Overflow posts, a tuned SVM performs similarly to a deep learner, but is significantly faster to train.

This paper extends that recent result by clustering the dataset, then tuning very learners within each cluster. This approach is over 500 times faster than deep learning (and over 900 times faster if we use all the cores on a standard laptop computer). Significantly, this faster approach generates classifiers nearly as good (within 2% F1 Score) as the much slower deep learning method. Hence we recommend this faster methods since it is much easier to reproduce and utilizes far fewer CPU resources.

More generally, we recommend that before researchers release research results, that they compare their supposedly sophisticated methods against simpler alternatives (e.g. applying simpler learners to build local models).

KEYWORDS

Deep learning, parameter tuning, DE, KNN, local versus global, KMeans, SVM, CNN

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1 INTRODUCTION

Recently, deep learning methods like convolutional neural networks (CNN) have become a popular choice for text mining in SE. Such deep learning works well with high dimensional data [35] but are very expensive in terms of time and required CPU.

In response to the computational problems of deep learners, researchers have tried alternate methods. For example, Fu et al. [12] recently revisited a study by Xu et al. [52] on finding the semantic relatedness of Stack Overflow posts. In that study, a single Stack Overflow question along with its complete answer was called a “knowledge unit” (KU). If any two KUs are semantically related, they are considered as linkable knowledge. Otherwise, they are considered isolated. When Xu et al. applied CNN [21] (a specific kind of deep learner), that took 14 hours to train. Fu et al. showed that this computational cost was avoidable by replacing a more complex learner (CNN) with a simpler technique augmented by some hyperparameter optimization. Specifically, Fu et al. shows that support vector machine (SVM) tuned via differential evolution (DE) could perform as well as CNN, while training 84 times faster.

This paper further extends the Fu et al. Using very simple widely used data mining method (KMeans), we can train even faster that Fu et al. and 500 times faster than deep learning (and over 900 times faster if we use all the cores on a standard laptop computer). The core to our approach is (1) building multiple local models then (2) tuning per local model. This paper evaluates this divide and conquer approach by:

(1) Exploring the Xu et al. task using SVM and K-nearest-neighbor (KNN) classifiers;
(2) Repeating step 1 using hyperparameter tuning—specifically, differential Evolution (DE)—to select control parameters for those learners;
(3) Repeats steps 1 and 2 using local modeling; i.e. clustering the data then apply tuning and learning to each cluster;
(4) Evaluating these local models in terms of both their training time and performance

Table 1 lists all the learners explored here.

| Type | Abbreviation | Learner Description |
|------|--------------|---------------------|
| G    | SVM          | Support Vector Machine |
| G    | KNN          | K-nearest Neighbors |
| G    | DE_KNN       | K-nearest Neighbors tuned using differential evolution (DE) |
| G    | DE_SVM       | Support Vector Machines Tuned using differential evolution |
| G    | CNN          | Convolution Neural Network |
| L    | KMeans_KNN   | Cluster + multiple KNNs |
| L    | KMeans_DE_KNN| Cluster + multiple KNNs tuned via DE |
| L    | KMeans_SVM   | Cluster + multiple SVMs |
| L    | KMeans_DE_SVM| Cluster multiple SVMs tuned via DE |

Table 1: Combinations of learning algorithms. L= “local models” and G= “global models” which denotes learning from clusters or all of the data (respectively).
This approach lets us ask and answer the following questions:

- **RQ1**: Can we reproduce Fu et al.’s results for tuning SVM with differential evolution (DE)?

**Result 1**

*Our DE with SVM perform no worse than Fu et al.*

- **RQ2**: How do the local models compare with global models in both tuned and untuned versions in terms model training time?

**Result 2**

*Local models perform comparably to their global model counterparts, but are 570 times faster in model training time.*

(To be precise, that 570 figure comes from running on a single core. If we distribute the execution cross the eight cores of a standard laptop computer, our training times become 965 times faster.)

- **RQ3**: How does the performance of local models compare with global models and state-of-the-art deep learner when used with SVM and KNN?

**Result 3**

*Local models performance very nearly as well (within 2% F1 Score) as their global counterpart and the state-of-the-art deep Learner.*

Based on these experiments and discoveries, our contribution and outcome from the paper are:

- A dramatically faster solution to the Stack Overflow text mining task first presented by Xu et al. This new method runs three orders of magnitude faster than prior work.
- Support for “not everything needs deep learning”; i.e. sometimes, applying deep learning to a problem may not be the best approach.
- Support for a simplicity-first approach; i.e. simple method like KMeans_DE_SVM can perform as good some of the state of the art models but with a (much) faster training time.
- Support for local modeling. Such local models can significantly reduce training time by clustering data then restricting learning to on each cluster.
- A reproduction package - which can be used to reproduce, improve or refute our results¹.

The rest of the paper is organized into the following sections Section 2 provides background information that directly relates to our research questions, in addition to laying out the motivation behind our work. In Section 3 a detailed description of our experimental setup and data, along with our performance criteria for evaluation is presented. It is followed by Section 4 the results of the experiments and answers to our research questions are detailed. Section 5 discusses threats to validity. Finally Section 6 concludes the paper with implications and scope for future work.

¹URL blinded for review.

## 2 Background

### 2.1 Motivation

Why obsesses on making software analytics faster? Why not just buy more cloud CPU time? Such a “just throw money at it” approach might not impress researchers like Fisher et al. [11] who define “software analytics” as a work flow that distills large quantities of low-value data down to smaller sets of higher value data. Due to the complexities and computational cost of some kinds of SE analytics, “the luxuries of interactivity, direct manipulation, and fast system response are gone” [11]. They characterize modern cloud-based analytics as a throwback to the 1960s-batch processing mainframes where jobs are submitted and then analysts wait, wait, wait for results with “little insight into what’s really going on behind the scenes, how long it will take, or how much it’s going to cost” [11]. Fisher et al. document the issues seen by 16 industrial data scientists, one of whom remarks “Fast iteration is key, but incompatible with the say jobs are submitted and processed in the cloud. It’s frustrating to wait for hours, only to realize you need a slight tweak to your feature set.”

Fisher’s experience matches with our own. We find that the slower the data mining method, the worse the user experience and the fewer the people willing to explore that method. These result is particularly acute in research where data miners have to be run many times to (a) explore the range of possible behaviors resulting from these methods or (b) generate the statistically significant results that can satisfy peer review. To understand the CPU problems with validation, consider the standard validation loop:

```
1. FOR L = 20 projects DO
2. FOR R = 20 times DO # repeats to satisfy central limit theorem
3. Randomly divide project.data to B = 10 bins;
4. FOR i = 1..B DO
5. test = bin[i]
6. train = project.data - test
7. FOR F = 5 different data filters DO
8. train = filter(train) # e.g. over-sample rare classes
9. model = leaner(train)
10. print report(apply(model,test))
```

Note the problem with this loop- it must call a data miner (at line 9) L * R * B * F = 20,000 times. While some data miners are very fast (e.g. Naive Bayes), some are not (e.g. deep learning). Worse, several “local learning” results [28–30, 51] report that software analytics results are specific to the data set being processed- which means that analysts may need to rerun the above loop anytime new data comes to hand.

Note that the above problem is not solvable by (1) waiting for faster CPUs or (2) parallelization. We can no longer rely on Moore’s Law [34] to double our computational power very 18 months. Power consumption and heat dissipation issues effectively block further exponential increases to CPU clock frequencies [34]. As to parallelization, that would require the kinds of environments that Fisher et al. discuss; i.e. environments where it is frustrating to wait for hours, only to realize you need a slight tweak to your feature setting.

Accordingly, in our research, whenever we have a slow and competent result, we explore methods to make that result faster. The rest of this paper offers a case study where local learner significantly speed up than deep learning.
Deep learning is a type of machine learning algorithm based on multiple layers of neural networks, where each layer is created with multiple neurons. These layers are interconnected via weights, which are tuned as the model trains. These connections and weights are very specific to models and its performance. According to Le-Cun et al. [26], deep learning methods are representation-learning methods with multiple levels of representation, obtained by composing simple but non-linear modules that each transforms the representation at one level (starting with the raw input) into a representation at a higher, slightly more abstract level.

Deep learning has been applied to many areas including image processing, natural language processing, genomics etc:

- Wan and Wang [46], deep learning tries to resolve the "Semantic Gap" issue in content based image retrieval process [41].
- Deep learning has also been used to predict DNA-RNA binding proteins [1] which helps to solve problems faced by less sophisticated methods (particularly the problem of the automatic extraction of meaningful features from raw data).
- Deep learning has also recently become established its presence in software engineering effort estimation [8] and text mining [7, 35, 49, 50, 53, 54].

Deep learning is a computationally expensive method. It often takes hours to weeks to train the models. This makes cross validation and stability of model check very expensive if not impractical. For example, the methods of Xu et al. take 14 hours to terminate or GU et al. [17] reported in his paper that his deep learning model took almost 240 hours to train [17]. Note that if those computations were repeated (say) 20 times for statistical purposes, then those systems would take 11 and 200 days to complete. Worse still, hyperparameter optimization\(^2\) might require 100s to 1000s of repeated runs – which would take another three to 50 years to terminate\(^3\).

For the above reasons, reproducing deep learning results is a significant problem:

- Most deep learning paper’s baseline methods in SE are either not publicly available or too complex to implement [25, 49].
- It is not yet common practice for deep learning researchers to share their implementations and data [7, 17, 25, 47, 49, 50], where a tiny difference may lead to a huge difference in the results.
- Due to the nature of complexity in implementation and unavailability of original implementation, data or environment it is not possible to implement a deep learner for baseline purpose, and this is one of the reasons for SE community and this paper to directly utilizes the results published for comparison [12, 25].

Hence, much research (including this paper) is forced to compare their new methods with published numbers in deep learning papers (rather than re-running the rig of the other researchers.)

As discussed below, local learning is one method for reducing that runtime.

\(^2\) Hyperparameter optimizers are algorithms that learn the control parameters of a learner. For more details on such algorithms, see §2.8.

\(^3\) Note that such long runtimes have been observed in the SE literature. In 2013, a team from UCL needed 15 years of CPU time to complete the hyperparameter optimization study of four software clone detection tools [46].

2.3 Local Learning

When running a data mining algorithm, all the training data can be used to build one training model. Alternatively, the training data can be somehow divided into small pieces and one model learned per piece.

Local models have shown promising results in different studies. Many researchers have found building specialized local models for specific regions of the data provides a better overall performance, thus according to the studies instead of trying to find a generalized model we should try to find best models specific to different region of data. For example, Menzies et al. [28] shows that for defect prediction and effort estimation, lessons learned from models build on small part of data set from PROMISE repository were superior to the generalized model built on all the data. That said, recent studies have suggested for at least for defect prediction, the benefit of local model may be learner specific [20].

This paper explored local learning for a somewhat different perspective. The claim made in this paper is not that local models always performs significantly better. Rather we recommend it since it can lead to significantly faster inference with very little compromise in the performance of the learned model.

More specifically, in this study we observe a three order of magnitude improvement over a prior text mining results by Xu et al [52]. Hence we recommend local learning since:

- It rarely performs worse than learning from all the data.
- Sometimes it can lead to better performance [5, 28, 30, 38].
- It can lead to significantly faster training times.

The general framework for local learning is shown as algorithm in Figure 1. This algorithm uses a cluster based model with the training dataset to find diversity in the dataset. This helps to create clusters with similar data. Now different classification models are accessed and improved via hyperparameter tuning on each of the clusters with local data as training dataset. While accessing the model for performance, the test data is sent through the clustering model to predict its probable cluster by calculating similarity measure, next the test data is classified using the local model built on that cluster.

The rest of this paper explores the local learning framework of Figure 1 using the following learning methods:

- k-means for the clustering;
- Differential evolution for the fitting;
- SVM or K-th-nearest neighbor for the classification.

For details on these methods, see later in this paper.

2.4 Word Embedding

In the case study of this paper, we explore local learning for text mining methods that use word embedding. This is the process of converting words to vectors in order to compare their similarity by comparing cosine similarity between vectors. One method for doing this is a continuous skip-gram model (word2vec), which is an unsupervised two-layer neural network that converts words into semantic vector representations [32] and is also used by Fu et al. [12] and Xu et al. [52] in their paper.
Another component of the case study explored here is support vector machines. SVMs are a type of supervised machine learning algorithm which analyzes data using classification or regression analysis. In SVM models the examples are points in space mapped corresponding word vector representation. There are many ways to create word embedding models. Mikolov et al. [32] proposed hierarchical softmax and negative sampling techniques etc which improve the computation efficiency. The word2vec model learns vector representation of a word (center word) by predicting surrounding words in a context window (c) by maximizing the mean of log probability of the surrounding words \( w_{i+j} \), given the center word \( w_i \) -

\[
\frac{1}{n} \sum_{c=1}^{n} \sum_{-c \leq j \leq c, j \neq 0} \log(w_{i+j}|w_i)
\]

(1)

The probability \( p(w_{i+j}|w_i) \) is a conditional probability defined by a softmax function -

\[
p(w_{i+j}|w_i) = \frac{\exp(v_w^T v_{w_{i+j}})}{\sum_{j=1}^{c} \exp(v_w^T v_{w_{i+j}})}
\]

(2)

Here the \( v_w \) and \( v_w^T \) are respectively the input and output vectors of a word \( w \) in the neural network, and \( W \) is the vocabulary of all words in the word corpus. \( p(w_{i+j}|w_i) \) is normalized probability of word \( w_{i+j} \) appearing in a specific context for a center word \( w_i \). To improve the computation efficiency, Mikolove et al. [32] proposed hierarchical softmax and negative sampling techniques etc which can also be used for creating word embedding models.

This paper uses the word2vec models trained by Fu et al. [12] that converted the Stack Overflow text data into the corresponding vectors. For our experiments, we use 100,000 randomly selected knowledge units tagged with “java” from Stack Overflow posts table (include titles, questions and answers). This data was pruned by removing superfluous HTML tags (while keeping short code snippets in code tag), then fitted into the gensim word2vec module [39]. This is a python wrapper over original word2vec package where, for word \( w_i \) in the post is sent to the trained word2vec model to get the corresponding word vector representation \( v_i \). After converting, all the KUs the output vectors are then used for training and testing the models.

2.5 SVM in Text Mining

Another component of the case study explored here is support vector machines. SVMs are a type of supervised machine learning algorithm which analyzes data using classification or regression analysis. In SVM models the examples are points in space mapped in such a way that separate categories/classes are divided by a clear gap (hyperplane in instance space) [44]. SVMs execute by transforming the original data space to a higher dimensional space where hyperplane between data from different classes is easier to detect.

SVMs are particularly useful for in text mining can have a very large number of features [13]. In most cases the document vectors are sparse and linearly separable in some hyper-dimensional space [23].

2.6 KNN in Text Mining

If SVM is our most sophisticated classifier, our simplest is K-Nearest Neighbor [55]. KNN is a non-parametric [15] method used for classification and regression problems. Here \( k \) is the input and refers to the number of closest examples that the model will look for among the training data in the feature space. The output the model gives represents the class to which the test data belongs to and it depends on the majority vote of its k-nearest neighbor [31].

2.7 KMeans Clustering

One way to reduce the computational cost of KNN is, before running a classifier, group data into similar sets of clusters. KMeans is an unsupervised machine learning algorithm that is used for clustering data [22]. In KMeans, \( k \) is an input that refers to the number of clusters the data set should be divided into. The algorithm initializes \( k \) number of centroids from the data and labels each as a cluster. For each data point it checks which centroid it is closest to, and assigns it to that cluster. After one pass to the data, centroids are recalculated and the process repeats until cluster stability is achieved. This experiments uses the scikit-learn module sklearn.cluster.KMeans [37].

Picking an appropriate \( k \) value for KMeans is a challenge. If \( k \) is too small, models will over-fit and fails to capture any larger patterns in the data. On the other hand, if using a large \( k \) will increase the variability and hence the level of uncertainty within each cluster. accordingly, this study uses the GAP statistics to determine the optimal number of \( k \) or centroids for KMeans [33, 45]. The GAP statistic looks at the difference between the dispersion of the clustered data, and the dispersion of a null reference distribution, for increasing \( k \) values. It finds the largest \( k \) where the gap is bigger than the next gap minus a value that accounts for simulation error. Figure 2 describes the GAP statistic computation [45].

\[
\text{def GAP( tData, nrefs=3, cMax = 15): # default settings}
\]

where hyperplane between data from different classes is easier to detect.

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Table 2: 'Tuning Range' of Parameters for SVM and KNN

| Parameters | Default Parameter | Tuning Range | Description |
|------------|-------------------|--------------|-------------|
| SVM C      | 1.0               | [1,50]       | Penalty parameter of error term |
| Kernel     | 'rbf'             | ['linear', 'poly', 'rbf', 'sigmoid'] | Specify the kernel type to be used in the algorithms |
| gamma      | 1/n_features      | [0,1]        | Kernel coefficient for 'rbf', 'poly' and 'sigmoid' |
| coef0      | 0                 | [0,1]        | Independent term in kernel function. It is only used in 'poly' and 'sigmoid' |

| Parameters | Default Parameter | Tuning Range | Description |
|------------|-------------------|--------------|-------------|
| KNN n_neighbors | 5 | [2,10] | Number of neighbors |
| weights | 'uniform' | ['uniform', 'distance'] | weight function used for predictions |

Figure 3: Tuner Procedure - as mentioned in Fu et al.'s paper. It is based on Storn’s DE optimizer.

### 2.8 Parameter Tuning with DE

All learners come with "magic parameters" that control their performance. For example, with SVM, there are several parameter that control the SVM kernel. One way to select those parameters is to use hyperparameter optimization via algorithms like Differential Evolution. DE is a stochastic population-based optimization algorithm [43]. DE starts with a frontier of randomly generated candidate solutions. For example, when exploring tuning, each member of the frontier would be a different possible set of control settings for (say) an SVM or KNN.

After initializing this frontier, a new candidate solution is generated by extrapolating by some factor $f$ between other items on the frontier. Such extrapolations are performed for all attributes at probability $cf$. If the candidate is better than one item of the frontier, then the candidate replaces the frontier item. The search then repeats for the remaining frontier items. For the definition of "better", this study uses the same performance measures as Fu et al.; i.e. "better" means maximizing the objective score of the model based F1 Score.

This process is repeated for $lives$ number of repeated traversals of the frontier. For full details of DE, see Figure 3. As per Storn’s advice [43] we use

$$f = 0.75, cf = 0.3, lives = 60$$

For SVM, this study uses the SVM module from Scikit-learn [37], a Python package for machine learning, where the following parameters shown in Table 2 are selected for tuning:

- **Parameter C** is to set the amount of regularization, which controls the trade-off between the errors on training data and the model complexity. A small value for $C$ will generate a simple model with more training errors, while a large value will lead to a complicated model with fewer errors.
- **gamma** defines how far the influence of a single training example reaches, with low values meaning 'far' and high values meaning 'close'.
- **coef0** is an independent parameter used in sigmoid and polynomial kernel function to scale the data.

Similarly KNN uses different parameters to control how it learns and how it predicts. This study uses KNN module from Scikit-learn, and the parameters in Table 2. For KNN:

- **Parameter n_neighbors** is number of neighbors to be used for query to check and classification using a majority vote [18].
- **weight** is another parameter which is tuned as part of hyperparameter tuning. If set to ‘uniform’, then all the points in each neighborhood are weighted equally. If set to ‘distance’, then the weights are in reverse of their distance from the new example (so farther away the point, the less weight it has in deciding the class).

### 3 EXPERIMENTAL DESIGN

#### 3.1 Data

This experiment uses the same training and testing dataset as Xu et al. [52] and Fu et al. [12]. For the reasons discussed in Section 2.2, this study compares our results to those reported by Xu et al.

Our dataset includes 6,400 training examples and 1,600 testing examples. Each class is equally represented in both the training and test datasets, with 1600 of each class in the training dataset and 400 of each class in the test dataset, so no handling of class imbalance is necessary.

Both test and training data are in Pandas dataframe [27] format, which includes a post Id, related post id, and a link type which is determined by a score between the 2 posts. The link between 2 posts can be of 4 types depending on the score between the sentences as per Table 3.

For finding word embedding [32] of Stack Overflow [3] post data, this paper uses the word2vec model trained by Fu et al. and described in Section 2.4. The final data that passed to our classifiers is similar to Table 4.
Table 3: Classification of data

| Scores | Class ID | Type          |
|--------|----------|---------------|
| 1.0    | 1        | duplicate     |
| 0.8    | 2        | direct link   |
| 0.8    | 3        | indirect link |
| 0.00   | 4        | isolated      |

Table 4: Training/Test data in Pandas.dataframe format.

| ID | Post Id | Related Post Id | Link Type Id | Post Id Vec | Related Post Id Vec | Output |
|----|---------|-----------------|--------------|-------------|---------------------|--------|
| 0  | 283     | 297             | 1            | [...]       | [...]               | [...]  |
| 1  | 56      | 68              | 2            | [...]       | [...]               | [...]  |
| 2  | 5       | 16              | 3            | [...]       | [...]               | [...]  |
| 3  | 9083    | 6841            | 4            | [...]       | [...]               | [...]  |

3.2 Method

Training a models and hyper-parameter tuning technique can take much time, depending on the complexity of the training dataset. This makes it harder to perform cross validation or repeatability of prior results. This study check if dividing up the data into small clusters and then train and tune models within each cluster, reduces the overall learner’s training time. In order to do this the data is clustered first, then a model is built for each cluster. This process is shown Figure 4.

For the clustering algorithm KMeans algorithm from Scikit-learn has been used. For the parameters, this study used the k-means++ algorithm [2] for the initialization of cluster centroids. In order to choose $k$, the number of clusters, the GAP statistic method, discussed in Figure 2 has been utilized.

For each cluster, a learner is built that is tuned specifically for that cluster. This study looks at two different learners, SVM and KNN. As discussed above, to tune the learners, DE is being used, specifically Fu et al.’s DE implementation. The study uses F1 Score [42] to evaluate the intermediate models in DE, as F1 Score is calculated as the trade-off between precision and recall. This will help us to get models which have both good precision and recall.

To use the model, the test data is first sent to KMeans to find the cluster which it should belong to by calculating vector distance from all cluster’s centroid and returning the cluster with minimum distance. Then the model predict the class using that cluster’s learner.

In this study a 10-fold cross validation [24] has been used, which was repeated 10 times for the training data. Thus, the results are the mean of 100 models. Each learner (SVM or KNN) on each cluster has been trained on 90% of the data from that cluster and tuned on rest of the 10% and then tested on the untouched test data set.

3.3 Performance Criteria

For evaluating the described model, this study collects and present the same metrics for performance evaluation as Xu et al. and Fu et al., in order to compare results. These metrics are precision, recall and F1 Score. This multiclass classification problem have 4 classes denoted as Duplicate(C1), Direct Link(C2), Indirect Link(C3) and Isolated(C4). The result presents class-wise metrics as well as the mean for the whole model.

From the confusion matrix describe in Table 5, it can be observed that the correct predictions are the one with label $C_{ij}$ for a class $C_i$. And from this we will be able to calculate our evaluation matrix.
Using this nomenclature, the evaluation metric \( F1 \) Score can be defined as follows:

\[
\text{precision} = \frac{C_{ii}}{\sum_{j} C_{ij}} \quad (3)
\]

\[
\text{recall} = \frac{C_{ii}}{\sum_{i} C_{ji}} \quad (4)
\]

\[
F1 = \frac{2 \ast \text{recall} \ast \text{precision}}{(\text{precision} + \text{recall})} \quad (5)
\]

### 3.4 Statistical Analysis

In order to compare results of the local models with other models, there are two useful tests: significance tests [4] and effect size tests [40] [6].

- Significance tests assess if results are distinct enough to be considered different;
- Effect size tells us whether that difference is large enough to be interesting.

After Wu et al., this study uses the Scott-Knott test, which ranks treatments using a recursive bi-clustering algorithm. At each level, treatments are split where expected value of the treatments has most changed from before. Results in each rank are considered the same according to both significance and effect size tests. As per the recommendations of Wu et al., the Scott-Knott test uses the non-parametric bootstrap [10] method and Cliff’s Delta. Note that these two tests are also used and endorsed by other SE researchers [14].

### 4 RESULTS

**RQ1: Can we reproduce Fu et al.’s results for tuning SVM with differential evolution (DE)?**

This study uses the same differential evolution with SVM for both global and local models. Thus to compare with Fu’s DE with SVM as global model, the first task as part of this experiment was to recreate Fu et al.’s work so that this study have a baseline to measure against. Hence, this research question is a “sanity check” that must be passed before moving on to the other, more interesting research questions.

For RQ2, this experiment built one model for each clusters using either normal or tuned versions of SVM or KNN (where tuning was performed with DE): For the default SVM and KNN the experiment uses the default parameters, described in Table 1.

As discussed above, this study have used the GAP statistic [33] [45] for finding the best number of clusters, using minimum and maximum number of clusters as 3 and 15, respectively. As part of the experiment we learned that 13 clusters achieves best results (measured as per the GAP statistic).

This study measures the time taken for this model to train which includes time taken by GAP statistic, K Means training time, and SVM/KNN with DE training time.

Figure 5 compare the model training time in log scale of all models with the results from XU et al.’s CNN approach. Its apparent from the figure 5 that for this domain KNN and SVM has the fastest runtimes. That said, as describe below, we cannot recommend these methods since, as shown below, they achieve poor F1 Scores.
The figure also shows, when we cluster data and then create local models on each cluster it achieves fast training times. For example, in the case of KMeans\_DE\_SVM, we achieve an eight-fold speed improvement from using DE\_SVM. Further, we see a speed improvement of over 570 faster training time from Xu’s CNN. 

**RQ3:** How does the performance of local models compare with global models and state-of-the-art Deep Learner when used with SVM and KNN?

The final part of our research question was to check if the local models performance is comparable to Fu et al.’s DE\_SVM and the Xu’s state of the art CNN. To evaluate the performance of the models this study compares F1 performance measures described in Section 3.3. As mentioned in the section, a 10 fold * 10 repeat cross validation was performed, so all the results are mean of 100 models created.

Figure 6 shows our F1 Score results (mean result across all 4 class of Table 6). The numbers on top of each bar show the results of statistical tests. Bars with the same rank are statistically indistinguishable. Note that these results should be discussed with respect to the runtime results shown above:

- In Figure 5, we saw that CNN was our slowest learning method. We would excuse this slowness and endorse its use if the performance scores for this extremely slow method was outstandingly high. This is not the case: as seen Figure 6, CNN is beaten by many of the treatments shown here.
- In Figure 5, DE\_SVM was our second slowest method. Again, we would endorse this approach if nothing else comes close to it in terms of performance. While KMeans\_DE\_SVM is significantly different (as shown by our Scott-Knott results), the median performance delta is very small indeed (median F1 Score scores of 94 vs 92). This is an important pragmatic consideration since, as shown in Figure 5, the learning time of KMeans\_DE\_SVM was 743/88 = 840\% faster than that of DE\_SVM.

In summary, we recommend one of our local learning method (KMeans\_DE\_SVM) since:

- It is 840\% faster than the prior state-of-the-art results (Fu et al.’s 2017 DE\_SVM method);
- On a single core machine, it is 570 times faster than the prior state-of-the-art before that (Xu et al.’s 2016 CNN method);
- On a standard laptop with 8 cores, KMeans\_DE\_SVM runs 965 times faster than Xu et al.’s CNN method;
- Our local learner performs better than Xu et al. and only a tiny fraction worse than Fu et al. Given these small performance deltas, from a pragmatic engineering perspective, we find it hard to justify the extra computational cost of DE\_SVM over KMeans\_DE\_SVM.

## 5 Threats to Validity

As with any empirical study, biases can affect the final results. Therefore, any conclusions made from this work must be considered with the following issues in mind:

- **Sampling bias:** threatens any classification experiment; i.e., what matters there may not be true here. For example, the data sets used here is a Stack Overflow dataset and were supplied by one individual. Although this study uses multiple word2vec models to validate with a 10-fold * 10 repeat validations. The text data is of similar format of question pairs.
- **Learner bias:** For building the model for finding semantic relatedness of question pairs in this study, we elected to use Support Vector Machine and K-Nearest Neighbor for classification and KMeans for clustering. This study chose these methods because its results were comparable to the more complicated algorithms and has been successful in text classification field. Classification and clustering is a large and active field and any single study can only use a small subset of the known algorithms.
- **Evaluation bias:** This paper uses precision, recall and F1 Score as performance measures. Other methods like Concordant - Discordant ratio [16], Gini Coefficient [36], Kolmogorov Smirnov chart [19] that are used for this purpose can also be used for performance evolution in future studies.
- **Input bias:** For the localization algorithm, this study randomly selects input values for a range to determine the number of clusters, also for hyperparameter tuning using DE, a subset of parameters has been selected for tuning and their range is either the whole range that the parameter accepts or a range that is selected for the study.

## 6 Conclusion

This paper investigates the value of local learning from the perspective of reducing the training time compared to methods for predicting knowledge unit’s relatedness on Stack Overflow. This study clustered the data first; then build local models on each cluster separately; then use the clustering algorithms with the local classification algorithm together to predict. As shown above:
Clustering the data first and then building local models on those subsets of data shows significant reduction in runtime.

Using KMeans on the Word Embedding model first to cluster the data into smaller subsets and then running SVM and KNN with their DE versions showed runtime reduction as large as 570 times for KMeans_DE_SVM version with CNN and almost 8 times improvement than its global DE_SVM version with sequential run. And with parallel run the performance improvement for KMeans_DE_SVM version with CNN is 965x and almost 14 times with its global counter part.

The performance in term of precision, recall and F-score is almost similar to its global counter part and sometime better then the XU’s CNN model.

For future work, we suggest trying several variations of this experiment -
- This study tunes the local models separately, tuning all the local models along with the clustering algorithm together might produce some interesting results.
- Trying to find correlation between number of cluster and training time and performance might give us effect of number of clusters on the prediction model.

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