Guided Stochastic Gradient Descent Algorithm for Inconsistent Datasets

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Abstract

Stochastic Gradient Descent (SGD) Algorithm, despite its simplicity, is considered an effective and default standard optimization algorithm for machine learning classification models such as neural networks and logistic regression. However, SGD's gradient descent is biased towards the random selection of a data instance. In this paper, it has been termed as data inconsistency. The proposed variation of SGD, Guided Stochastic Gradient Descent (GSGD) Algorithm, tries to overcome this inconsistency in a given dataset through greedy selection of consistent data instances for gradient descent. The empirical test results show the efficacy of the method. Moreover, GSGD has also been incorporated and tested with other popular variations of SGD, such as Adam, Adagrad and Momentum. The guided search with GSGD achieves better convergence and classification accuracy in a limited time budget than its original counterpart of canonical and other variation of SGD. Additionally, it maintains the same efficiency when experimented on medical benchmark datasets with logistic regression for classification.

Keywords: Stochastic Gradient Descent Algorithm, machine learning, classification, logistic regression, neural networks, greedy selection, Guided Stochastic Gradient Descent Algorithm.

1. Introduction

Classification is one of the major techniques in data mining for many applications. Neural networks, logistic regression and support vector machine are some commonly used classification models. Many of these models rely on optimization algorithms for convergence, therefore, accuracy and efficiency of these optimization algorithms have major impact on the quality of classification. Some of the techniques like mathematical Lagrangian method, meta-heuristic algorithms and Gradient Descent Algorithms (GDA) have been employed for solving optimization problems in classification [15, 19, 21, 22]. Optimization problems are

generally NP-hard problems where search techniques can have limitation in searching either for a local or the global optimal solutions. GDA generally falls into local optimum solutions for non-convex functions. GDA is an efficient domain data-driven optimization technique, unlike meta-heuristic algorithms that makes it one of the most popular algorithms for machine learning algorithms such as neural networks (NN) and logistic regression [17, 20, 27].

GDA's two major variants are Stochastic Gradient Descent (SGD) and Batch Gradient Descent (BGD) which will be discussed in detail in Section 2. SGD was derived from BGD to provide an efficient alternative for a gradient descent method. However, the price for efficiency is the untimely random selection of instances which causes high fluctuations in descent or delayed convergence. Additionally, canonical SGD inherits the nature of 'blind' search that does not recognize the layout of a given landscape for the optimum descent. This results in slow rate of convergence in some extreme conditions. Many improvements had thus been proposed in order to combat these drawbacks.

1.1 Related Work

Qian in [18] has shown the valley-shaped search space has the higher slope for hill gradient compared to the valley that makes the slope direction almost perpendicular to the valley. GDA oscillates near the valley that slows down the convergence. Momentum term [18] is added to reduce the oscillation and improve the convergence. Nesterov Accelerated Gradient (NAG) [29] uses an enhanced momentum terms that also approximates the future positions of the gradient to have even better convergence rate. For sparse data set, Adagrad [7] performs larger updates for infrequent, and smaller updates for frequent parameters. However, in general, its auto updating parameters iteratively becomes ineffective and the learning stops. Adadelta/RMSprop [30] is the improvement of Adagrad by restricting parameters to keep it effective for learning. Hence, the learning rate does not aggressively reduce to minimum in a first few iterations. Adam [12] is another method like Adadelta and RMSprop that computes adaptive learning rates for each parameter. RMSprop, Adadelta and Adam are very similar algorithms that do well in similar circumstances. Adams' bias-correction helps it to slightly outperform RMSprop towards the end of optimization as gradients become sparser [20].

All these variations had been successfully used with small scale problems in a single machine as traditional formulation of SGD is not suitable for deep learning, particularly in the environment of distributed online learning. Asynchronous SGD has been widely adopted to solve this problem. Moreover, many variations such as RMSprop, Adadelta and Adam show promising results for large scale distributed machine learning algorithms such as deep NN [2, 5, 31, 32]. This paper only focuses on small scale problems where the proposed algorithm, guided SGD (GSGD) works as an add-on to all the variations of SGD. It was shown that GSGD is compatible with all the popular variations of SGD. The algorithm names of the guided versions are prefixed with G such as GRMSprop and GAdam for RMSprop and Adam respectively. However, GSGD is used in this paper to refer to the guided model in general unless otherwise specified. Guided search is not a new approach to improve the search for optimal solutions in optimization algorithms [9, 23], however, the presented approach is a novel idea where guidance is done through consistent data instances.

1.2 Problem Statement

The variations of SGD exploit the landscape of the search space to have a faster convergence rate. However, they do not address the poor convergence or low classification accuracy due to the inconsistency present in a dataset. The proposed method is a variation of SGD that overcomes the problem of high inconsistency in a dataset. SGD and its current variations do not have any strategy to filter out the inconsistent data, while the GSGD guides them by selective extraction and exploitation of consistent data instances for gradient descent. Thereby, it improves the convergence quality and classification accuracy. In case of consistent dataset, GSGD would not show any improvement.

The scope of this paper is to analyze the behavior of SGD on inconsistent dataset and provide a better solution using GSGD without compromising the efficiency of SGD. Inconsistency in a dataset is caused by those training examples that result in a net error (or cost) to increase during gradient computation. Mathematical explanation of data inconsistencies is given later on, in the paper. Canonical SGD's convergence can be improved if the variance in random selection of the instances can be reduced [11, 13]. This can be achieved by temporarily filtering out inconsistent training examples in gradient computation. High variance causes slow convergence with high fluctuations. Therefore, anomaly/outlier detection and removal

techniques should not be applied without appropriate domain analysis [4]. The proposed algorithm does not perform any data cleaning as preprocessing. It is simply a part of training process that tries to distinguish consistent data with inconsistent one to produce better results. A noise is never removed permanently. Moreover, it does not replace the existing outlier detection techniques. The proposed technique provides a competitive alternative for GDA.

The remainder of the paper is organized as follows: Section 2 describes and formalizes various types of GDAs. Section 3 highlights the impact of inconsistent training examples on gradient descent. Section 4 introduces the proposed variation of SGD algorithm known as guided SGD. Section 5 shows comprehensive experiments to compare original variations with guided variations of SGD and Section 6 discusses the outcome of the experiment. Lastly, Section 7 concludes the paper with some suggestions to the future work.

2. Formalization of Gradient Descent Algorithms

GDA is a domain data driven optimization algorithm that has rightly found its niche in machine learning algorithms such as Neural Networks (NN) and logistic regression to optimize the given parameters. GDA iteratively follows the negative gradient of a given activation function in order to move in the direction of descent, to locate the desired minimum. Some commonly used activation functions are *sigmoid*, *tanh* and *softmax* [27]. Sigmoid and *softmax* have been used for two-class and multi-class classification problems respectively. The optimization problem for linear model ($W^T x = 0$) is formulated as:

$$\widehat{W} = \underset{W}{\operatorname{argmin}} \frac{1}{N} \sum_{n=1}^{N} E(x_n, y_n, W)$$

where $E(x_n, y_n, W)$ is an error or cost function derived from activation function that takes d dimensional N training examples from $(x_1, y_1) \dots (x_N, y_N)$ with $x_n \in R^d$ and $y_n \in R$, and the weight vector $W \in R^d$ for two-class classification using logistic regression. For multi-class classification with T classes, $W \in R^{d \times T}$ and $y_n \in R^T$. There are two major variants of gradient descent based on the manner in which the gradient is calculated, have been described below:

2.1 Batch Gradient Descent

Batch Gradient Descent (BGD) is also known as bulk or true gradient descent. In the batch version of gradient descent the weight vector $W(t)$ is randomly initialized for iteration $t = 0$.

Afterwards, it is iteratively updated such that at each step a short distance is moved in the direction of error's greatest rate of decrease (steepest descent) for each data instance [3, 10]. However, this does not mean that collectively the steepest descent of error is achieved in every iteration for the entire training examples in every step. The linear models for classification have a perceptron or a set of perceptrons as connectors whose weight vector (or matrix) is to be optimized to get the minimum error. The pseudocode of BGD in logistic regression for classification is given in Fig. 1.

BGD algorithm has only two major steps: average gradient computation and weight update. The algorithm begins with the initialization of learning rate η, initial weight vector W_0 and total iterations T. W_0 is a $\mathbf{0}$ vector for logistic regression and η is generally fixed between 0 – 1 but adaptive learning rates are also being used [16, 28]. The size of training dataset is N with d dimensions. $\nabla E(W_t, x_i, y_i)$ is the gradient of error function $E(W_t, x_i, y_i)$ for a given data instance x_i, its output y_i and weight vector W_t. The algorithm stops when the termination criteria is met which is generally the given maximum iterations, or just before the overfitting starts [15].

```
Initialize η and W_0
For t = 1 … T
    Gradient computation
    ∇E̅ = 0;
    For k ∈ {1 … N}
        ∇E̅ = ∇E̅ + ∇E(W_{t−1}, x_k, y_k)
    End for
    ∇E̅ = ∇E̅ / N
    Update weight vector
    W_t = W_{t−1} − η∇E̅
    Track Error Progression
    Ē_t = 0
    For j ∈ {1 … N}
        E_j = E(W_t, x_j, y_j)
        Ē_t = Ē_t + E_j
    End for
    Ē_t = Ē_t / N
    t = t + 1
    Check Termination criteria
End for
```

Fig. 1. Pseudocode for Batch Gradient Descent

The batch gradient descent works well if the training data is not too large because computing the gradient term becomes expensive as the data size increases. The time complexity for a single gradient computation is $O(Nd)$ for linear models where d is the dimension and N is the

size of a given dataset respectively. In case of logistic regression, the overall worst time complexity with T iterations would be $O(T(dN + N) = O(TdN)$, disregarding the early stop due to overlearning.

2.2 Stochastic Gradient Descent

Stochastic gradient descent (SGD) addresses the issue of high computational cost by having much faster convergence, in the order of $O(T(d + N)$ only. SGD only differs in how much data is used to compute the gradient of the objective function. Depending on the amount of data, a trade-off is made between the accuracy in weight update and the time it takes to perform an update [20, 26].

The pseudocode of SGD is given in Fig. 2. SGD also has only two major steps: individual gradient computation and weight update, where it calculates a gradient $\nabla E(W_t, x_i, y_i)$ of an instance i randomly selected at iteration t, instead of a true gradient $\overline{\nabla E}$. The definition of all parameters are same as of BGD algorithm. BGD converges smoothly towards the global minimum for strictly convex error function and possibly to a local minimum for non-convex function. On the other hand, SGD continuously fluctuates to converge, where it helps W_t to jump to a new and potentially better local minima for non-convex error function [1]. The fluctuations in SGD happens because of high variance in random selection of a single gradient instead of the true gradient of the entire training dataset [11]. The algorithm stops when the termination criteria is met which is again the maximum number of iterations or before the algorithm begins to over learn [15].

```
Initialize η and W_0
For t = 1 … T
    For i ∈ {1 … N} selected in a random sequence one at a time.
        W_t = W_{t−1} − η∇E(W_{t−1}, x_i, y_i)
        Track Error Progression
        Ē_t = 0
        For j = 1 … N
            E_j = E(W_t, x_j, y_j)
            Ē_t = Ē_t + E_j
        End for
        Ē_t = Ē_t / N
        t = t + 1
        Check Termination criteria
    End for
End for
```

Fig. 2. Pseudocode for Stochastic Gradient Descent

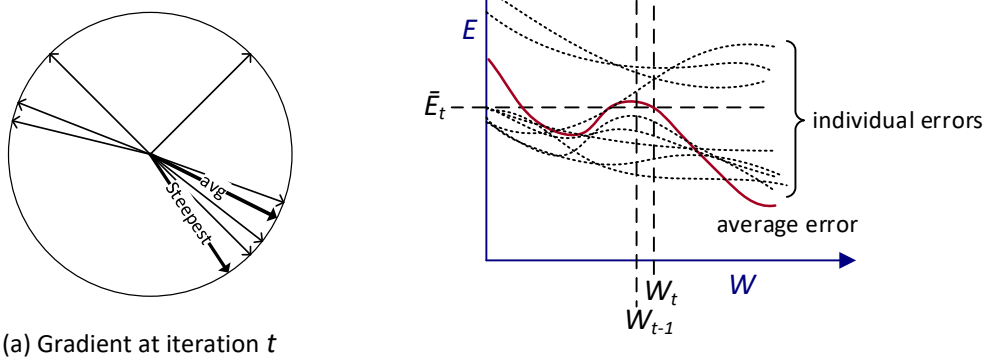
3. Impact of Data Inconsistency on Gradient Descent

The gradient descent algorithms do not differentiate between consistent and inconsistent training examples. BGD relies on true gradient to nullify the effects of inconsistent data in every iteration. However, determining true gradient in every iteration makes the algorithm very expensive and unsuitable to the applications that require online or deep learning. On the other hand, SGD randomly picks a training example without considering its consistency. That helps in faster convergence and avoidance of many local optimum solutions, however, it may also result in irregular convergence. The analysis of stepwise convergence and effects of inconsistent training example in every iteration in gradient descent is discussed below.

The objective of BGD is to reduce the average error $\bar{E}_t$ on every iteration t. At a given weight vector W_t, on the landscapes of Weights vs Error function, different instances produce different gradients as illustrated in Fig. 3 (a). The average gradient is not necessarily the steepest gradient at iteration t. Depending on the sample selection, complexity and noise level of the data, the expectation on every iteration is to have an overall training error $\Delta\bar{E}_t = \bar{E}_t - \bar{E}_{t-1} \leq 0$ which can be given as:

$$\Delta\bar{E}_t = \frac{1}{N}\big(\Delta E(W_t, W_{t-1}, x_1, y_1) + \cdots + \Delta E(W_t, W_{t-1}, x_i, y_i) + \cdots + \Delta E(W_t, W_{t-1}, x_N, y_N)\big) \leq 0 \tag{1}$$

where $\Delta E(W_t, W_{t-1} x_i, y_i) = E(W_t, x_i, y_i) - E(W_{t-1}, x_i, y_i)$. A rough sketch for this behavior of individual instances are shown in Fig. 3 (b), where most of the instances have error decreasing at iteration t while couple of instances have error increasing.



(a) Gradient at iteration t

(b) Behavior of individual instances

Fig. 3. Gradients for BGD

The parameters of error functions $E\left(W_t, x_j, y_j\right)$, $\Delta E\left(W_t, W_{t-1}, x_j, y_j\right)$ and $\nabla E\left(W_t, x_j, y_j\right)$ are replaced with subscripted parameters $E_j(W_t)$, ΔE_j and ∇E_j respectively for rest of the paper for simplicity. From Eq. (1) instances $1,2,\dots,i$ are considered consistent as their respective errors are decreasing i.e. $\{\forall\, \Delta E_j \leq 0 \mid j \leq i\}$ and vice-versa for instances $i+1,\dots,N$ where $\{\forall\, \Delta E_j > 0 \mid j > i\}$. If $\bar{E}_t \propto \frac{1}{i}\sum_{j=1}^{i} \nabla E_j$ and $\left|\sum_{j=1}^{i} \Delta E_j\right| > \left|\sum_{j=i+1}^{N} \Delta E_j\right|$ then the error is said to be decreasing with $W_t = W_{t-1} - \eta\overline{\nabla E}$ for a non-chaotic error function. If the error for training examples $\{1,\dots,i\}$ has the highest Lipschitz constant L then the Lipschitz condition for continuity can be written as:

$$\left|\Delta E_j\right| = \left|E_j(W_t) - E_j(W_{t-1})\right| \leq L|W_t - W_{t-1}| \text{ for } \forall j \leq i \tag{2}$$

where $E_j(W_t) - E_j(W_{t-1}) \leq 0$

The highest Lipschitz constant for $\forall j > i$ has been determined by the summation of Lipschitz condition from Eq. (2).

$$\sum_{j=1}^{i}\left|E_j(W_t) - E_j(W_{t-1})\right| \leq iL|W_t - W_{t-1}| \text{ for } \forall j \leq i \tag{3}$$

Similarly,

$$\left|\Delta E_j\right| = \left|E_j(W_t) - E_j(W_{t-1})\right| \leq L'|W_t - W_{t-1}| \text{ for } \forall j > i \tag{4}$$

$$\sum_{j=i+1}^{N}\left|E_j(W_t) - E_j(W_{t-1})\right| \leq (N-i)L'|W_t - W_{t-1}| \tag{5}$$

where L' is the highest Lipschitz constant for $\forall j > i$ and $E_j(W_t) - E_j(W_{t-1}) > 0$. To have overall descending (or same) error in an iteration t, the following condition must hold:

$$(N-i)L'|W_t - W_{t-1}| \leq iL|W_t - W_{t-1}|$$

$$\therefore L' \leq \frac{i}{N-i}L \tag{6}$$

In an ideal case, the landscape of training data behaves consistently with every instance, i.e. $i = N$. So, any small positive value of L ($L > 0$) would ensure error decreases in every next iteration until convergence is almost complete. Training examples' high inconsistency can be tolerated with the true gradient as long as Eq. (6) is true.

SGD does not use true gradient but rather picks up an example randomly. The increase or decrease in error value depends on the presence of inconsistent examples in a given iteration. The error value is determined by the same principles as discussed above from Eq. (1) to Eq. (6); where only the true gradient $\bar{E}_t$ is replaced by an individual gradient ∇E_k, where $k \in \{1,2,\dots,N\}$. ∇E_k can be directly proportional to either $\frac{1}{N}\sum_{j=1}^{i}\nabla E_j$ or $\frac{1}{N}\sum_{j=i+1}^{N}\nabla E_j$ which makes the error descend or ascend respectively. Generally, for consistent data $i \gg (N-i)$; descent happens more often over the iterations, and of course with a better convergence rate. More specifically, for $\left|\sum_{j=1}^{i}\Delta E_j\,(W_t)\right| > \left|\sum_{j=i+1}^{N}\Delta E_j(W_t)\right|$ the error $\bar{E}_t$ decreases if $\frac{\nabla E_k}{\|\nabla E_k\|} \propto \frac{1}{i}\sum_{j=1}^{i}\frac{\nabla E_j}{\|\nabla E_j\|}$ where $k \leq i$; otherwise the error increases in a normal circumstances when some data instances are not chaotic in nature. Using Lipschitz condition for continuity if descend happens only in t' iterations out of total T iterations then following Eq. (2) and Eq. (4), the equation for the net descend after T iterations would be:

$\sum_{t=1}^{t'}\sum_{j=1}^{i(t)}\left|E_j(W_t) - E_j(W_{t-1})\right| \leq LNt'|W_t - W_{t-1}|$ for net gradient descent

$\sum_{t=t'+1}^{T}\sum_{j=i(t)+1}^{N}\left|E_j(W_t) - E_j(W_{t-1})\right| \leq L'N(T-t')|W_t - W_{t-1}|$ for net gradient ascent

$$\therefore L' \leq \frac{t'L}{(T-t')} \text{ for } t' < T \tag{7}$$

Thus, net descent in SGD happens after T iterations when Eq. (7) holds true. Otherwise, the algorithm will diverge for the given iterations. In comparison with BGD, gradient descent does not happen in every iteration, however, some fluctuations are seen while the algorithm runs. Convergence for consistent data is smoother compared to inconsistent data. This shows the importance of consistent examples in gradient descent which is the primary focus of this paper.

One issue with SGD is its high variance caused by the random selection of gradient instead of average gradient [11]. Therefore, one logic variation of canonical SGD is the combination of BGD with SGD known as min-batch gradient descent that reduces the variance to some extent and still manages better convergence rate. However, most recent SGD variations discussed in introduction section has transpired that neither decreasing step-sizes nor mini-batching are necessary to resolve the non-vanishing variance issue inherent in the canonical SGD methods [13, 25].

4. Guided Stochastic Gradient Descent Algorithm

After describing BGD and SGD it is evident that high data inconsistency plays a major role in the degree of error descent per iteration and convergence. The proposed algorithm is named as guided stochastic gradient descent (GSGD) algorithm that realizes the importance of inconsistency in a dataset and tries to improve the convergence by separating consistent and inconsistent data instances to some extent. GSGD hides inconsistent data for "a while" in an anticipation that they will become consistent after few iterations. In addition to individual gradient computation and the weight update, GSGD selects consistent instances gradually and does the refinement of the weight update.

Inconsistent data instances are simply the data instances within the neighborhood of instance j; which individually performs better, while the average error value $\bar{E}_t$ performs worse than the average error of the previous iteration $\bar{E}_{t-1}$, and vice-versa. The pseudocode of the algorithm is given in Fig. 4 and Fig. 5. The definition of variables used in the GSGD algorithm are same as in SGD and BGD unless otherwise specified. The algorithm begins by initializing weight vector (W_0), learning rate (η), total iterations (T) and neighborhood size (ρ). Neighborhood allocation allows GSGD to run through ρ training examples recursively before further refinement of W is done with the consistent data. Consistent data is selected from ρ training examples at a time. Each data instance x_i with label y_i is randomly selected to update the weight vector (W_t) at iteration t like SGD. The neighborhood of an instance k is kept in sets ψ^+ and ψ^- where they track inconsistent instances. After collecting the data about inconsistency for ρ iterations, only consistent instances are extracted and kept in ψ^+ and ψ^-. ψ_k^+ in the pseudocode refers to k^{th} element of either ψ^+ or ψ^-, i.e., $\left\{\psi_k^+ \in \psi^+ | k \in \{1 \dots \rho\}\right\}$ and $\left\{\psi_k^- \in \psi^- | k \in \{1 \dots \rho\}\right\}$. Lastly, W_t is further updated with consistent instances ω_t.

Its flowchart is given in Fig. 6 where the algorithm starts with a random selection of a data instance whose gradient is computed to update the weight vector. After ρ iterations, weight vector is further refined with consistent neighboring data instances. ρ is also the neighborhood size which is generally assigned to a constant value 10.

```
//Initialize η, ρ, T and W_0
for t = 1 … T
    ∀ψ_k^+ = 0 | k ∈ {1 … ρ}
    ∀ψ_k^- = 0 | k ∈ {1 … ρ}
    for i ∈ {1 … N} //pick i in a random sequence
        W_t = W_{t-1} − η∇E(W_{t-1}, x_i, y_i) //canonical weight update
        Ē_t = 0
        for j = 1 … N
            E_j = E(W_t, x_j, y_j)
            Ē_t = Ē_t + E_j
            [ψ^+, ψ^-] = collectInconsistentInstances(i, j, ρ, E_j, Ē_{t-1}, ψ^+, ψ^-)
        End for
        Ē_t = Ē_t / N

        [ψ^+, ψ^-, ω_t] = extractConsistentInstances(i, ρ, ψ^+, ψ^-, Ē_t, Ē_{t-1})

        //further refinement of W_t with consistent data
        For each k ∈ ω_t
            W_t = W_t − η∇E(W_t, x_k, y_k) //canonical weight update
        End for
        t = t + 1
        //Check Termination criteria
    End for
End for
```

Fig. 4. Pseudocode for Guided Stochastic Gradient Algorithm

```
[ψ^+, ψ^-] = collectInconsistentInstances(i, j, ρ, E_j, Ē_{t-1}, ψ^+, ψ^-)
    //Determine inconsistent data instances
    if ⌊(j + ρ − 1)/ρ⌋ == ⌊(i + ρ − 1)/ρ⌋
        k = j − ρ⌊(j + ρ − 1)/ρ − 1⌋
        if E_j − Ē_{t-1} > 0
            ψ_k^+ = ψ_k^+ + (E_j − Ē_{t-1})
        else
            ψ_k^- = ψ_k^- − (E_j − Ē_{t-1})
        End if
    End if
End

[ψ^+, ψ^-, ω_t] = extractConsistentInstances(i, ρ, ψ^+, ψ^-, Ē_t, Ē_{t-1})
    ω_t = ∅
    Extract consistent data points with respect to x_i
    if mod(i, ρ) == 0
        if Ē_t < Ē_{t-1};
            ψ^+ = sort(ψ^+, "Descend");
            ω_t = {k | ψ_k^+ ∈ ψ^+ ∧ ψ_k^+ = 0} //get indices with no noise
        else
            ψ^- = sort(ψ^-, "Descend");
            ω_t = {k | ψ_k^- ∈ ψ^- ∧ ψ_k^- = 0} //get indices with no noise
        End if
        ∀ψ_k^+ = 0 | k ∈ {1 … ρ}
        ∀ψ_k^- = 0 | k ∈ {1 … ρ}
    End if
End
```

Fig. 5. Subroutines for Guided Stochastic Gradient Algorithm

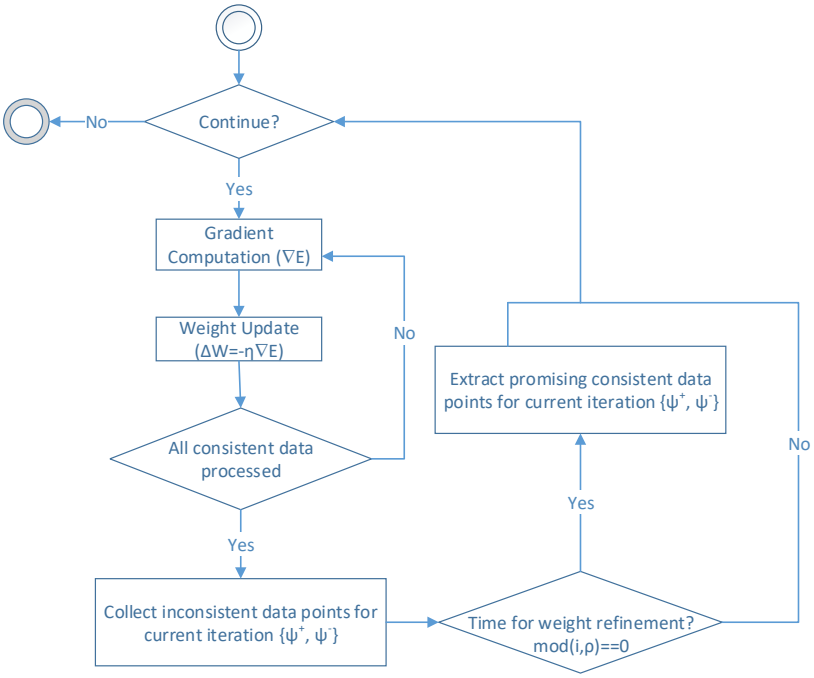


Fig. 6. Flowchart of Guided Stochastic Gradient Algorithm

The worst time complexity of GSGD is same as SGD even with the addition of sorting and further refinement of the weight vector. Sorting (with quick sort) and further refinement have worst time complexity of ρ^2 and $d\rho$ respectively. Hence, overall the worst time complexity of GSGD is $O\big(T(d+N+\rho^2+d\rho)\big) = O(T(d+N)$ which is same as SGD because $\rho \ll N$ is a constant term. It is also interesting to note that GSGD is same as SGD for the first ρ iterations. Therefore, it is very important to choose an appropriate value of ρ as its very large value will simply make the algorithm original SGD. As such, more analysis has been done on it in Section 5.

Since GSGD uses the same gradient computation formula, it has been easily incorporated into commonly used amendments for SGD; namely Adam, Adagdelta, Adagrad, RMSprop, Momentum and Nesterov. These amendments include some additional factors like momentum, regularization or variance reduction techniques [18]. To apply these variations to GSGD, only canonical weight update section in Fig. 4 needs to be replaced with the chosen variation of weight update. For example weight update for RMSprop is $W_t = W_{t-1} - \eta \frac{\overline{\nabla E}}{\sqrt{(r_t+\varepsilon)}}$, where $\varepsilon = 1.0 \times 10^{-8}$, and $r_t = \beta r_{t-1} + (1-\beta)\overline{\nabla E_t}^{\,2}$ where $r_1 = \overline{\nabla E_1}$ and $\beta = 0.9$. The guided version of RMSprop is given in Fig. 7.

```
//Initialize η, ρ, T and W_0
for t = 1 ... T
    ∀ψ_k^+ = 0 | k ∈ {1 ... ρ}
    ∀ψ_k^- = 0 | k ∈ {1 ... ρ}
    β = 0.9 //RSMprop specific constant
    for i ∈ {1 ... N} //pick i in a random sequence
        r_t = β r_{t-1} + (1 − β) ∇E(W_{t-1}, x_i, y_i)^2
        W_t = W_{t-1} − η∇E(W_{t-1}, x_i, y_i) / √(r_t + ε)  //RMSprop weight update
        Ē_t = 0
        for j = 1 ... N
            E_j = E(W_t, x_j, y_j)
            Ē_t = Ē_t + E_j
            [ψ^+, ψ^-] = collectInconsistentInstances(i, j, ρ, E_j, Ē_{t-1}, ψ^+, ψ^-)
        End for
        Ē_t = Ē_t / N

        [ψ^+, ψ^-, ω_t] = extractConsistentInstances(i, ρ, ψ^+, ψ^-, Ē_t, Ē_{t-1})

        //further refinement of W_t with consistent data
        For each k ∈ ω_t
            W_t = W_t − η∇E(W_t, x_k, y_k) / √(r_t + ε)  //RMSprop weight update
        End for
        t = t + 1
        //Check Termination criteria
    End for
End for
```

Fig. 7. Pseudocode for Guided RMSprop

The highlighted lines show the only changes required to be done in GSDG to make it guided RMSprop. Rest of the code are same. Similarly, guided version for Adam, Adagdelta, Adagrad, Momentum and Nesterov have been created.

Finally, the convergence of GDAs has been discussed with constant learning rate and constraints associated with them. Using Taylor series for BGD the error function can be written as:

$$E(W_t) \leq E(W_{t-1}) + (W_t - W_{t-1})\nabla E = E(W_{t-1}) + (W_{t-1} - \eta\nabla E - W_{t-1})\nabla E$$

$$\therefore E(W_t) \leq E(W_{t-1}) - \eta\|\nabla E\|^2 \tag{8}$$

This shows that the error converges towards local optimum solution (global for convex function) gradually in every iteration. Large value of η would result in divergence.

Similarly, error function for SGD can be written as:

$$E(W_t) \leq E(W_{t-1}) + (W_t - W_{t-1})\nabla E$$

$$= E(W_{t-1}) + \left(W_{t-1} - \eta\nabla E\left(W_{t-1}, x_k, y_k\right) - W_{t-1}\right)\nabla E$$

Here instead of the true gradient ∇E, a randomly selected gradient for k^{th} instance $\nabla E(W_{t-1}, x_k, y_k)$ has been used to update the weight W_t.

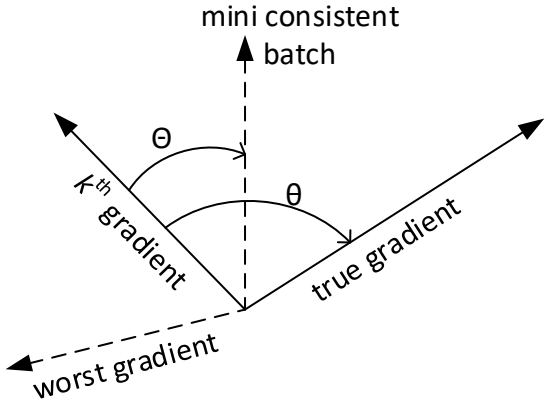


Fig. 8. Deviation of individual gradients from the true gradient

$$E(W_t) \leq E(W_{t-1}) - \eta \nabla E\left(W_{t-1}, x_k, y_k\right) \nabla E = E(W_{t-1}) - \eta \|\nabla E\|^2 cos\theta$$

where θ is the angle between k^{th} gradient and the true gradient as shown in Fig. 8.

$$\therefore E(W_t) \leq \begin{cases} E(W_{t-1}), \text{when consistent } (\theta \leq \frac{\pi}{2}) \\ E(W_{t-1}) + \eta \|\nabla E\|^2, \text{when inconsistent } (\theta > \frac{\pi}{2}) \end{cases} \quad (9)$$

SGD would converge when the dataset is consistent, though convergence cannot be guaranteed due to high inconsistency present in the data set. This is the reason many fluctuations are seen during gradient descent. GSGD tries to reduce θ by selecting promising instances that can bring k^{th} gradient closer to the true gradient as shown by Θ in Fig. 8. Since GSGD is an offshoot of SGD, its convergence is similar to SGD given in Eq. (9). The error value based on the algorithm defined in Fig. 4 and Fig. 5 can be given as:

$$E(W_t) \leq E(W_{t-1}) + \left(W_{t-1} - \eta \sum_{i=1}^{|\omega_t|} \nabla E\left({W_{t-1}}^{(i)}, x_i, y_i\right) - W_{t-1}\right) \nabla E$$

where ${W_{t-1}}^{(i)}$ refers to the intermediate weight value while processing the consistent set ω_t.

$$\therefore E(W_t) \leq E(W_{t-1}) - \eta \|\nabla E\|^2 cos\Theta$$

$$\text{and, } E(W_t) \leq \begin{cases} E(W_{t-1}), \text{when consistent } (\Theta \leq \frac{\pi}{2}) \\ E(W_{t-1}) + \eta \|\nabla E\|^2, \text{when inconsistent } (\Theta > \frac{\pi}{2}) \end{cases} \quad (10)$$

By selecting promising instances the k^{th} gradient moves closer to the true gradient. GSGD will have better convergence with $\Theta \leq \theta$.

5. Experiment

Since GSGD is simply a "guided" version of SGD, hence a comprehensive analysis was performed with guided version of canonical SGD, Adam, Adagrad, RMSprop, Adadelta, Momentum and Nesterov with their corresponding original versions. The coding was completely written in Matlab 2016. The experimental setup and execution have been discussed below:

5.1 Test Data

The test data have been collected from various medical benchmark data sets available in UCI library [6]. The presence of inconsistency in datasets differs from each other; for instance, Breast Cancer Diagnostic has high consistency, Cancer and New Thyroid has medium level of consistency while rest of data sets are very chaotic.

To have a comparative analysis on the performance of algorithms, the quality metric is designed that consists of classification accuracy (CA) on a limited time budget, and number of function calls (NFC) for 30 successive runs on each data set. The algorithms are given sufficient and equal amount of time to converge where the algorithms are also monitored to not to overlearn.

The selected datasets belong to either two-class or multi-class classification problems. Since the GSGD differs from SGD mainly in prioritizing selection of consistent data for gradient computation, this may be argued as merely a kind of data preprocess with noise filtering. Hence, the algorithms have also been tested for both orientation of datasets i.e., without noise filtering and with noise filtering through outlier detection and removal. This demonstrates that even after noise filtering, GSGD performs better than SGD. Statistical inter-quartile range (IQR) outlier detection method available in [8] have been used to remove stochastic noise from datasets where CAs are low. This technique has been commonly used in medical domain with increased descriptive classification accuracy but with low predictive accuracy [14, 24]. The datasets are not intended to be free from the noise but to reflect the effect of some outlier removals in the tested algorithms.

5.2 Parameter Settings

The experimental setup ensures that initial conditions like random order of training examples are same for both SGD and GSGD. Both algorithms are placed as subroutines inside the common loop of iterations. In this manner, same data instances are fed into the algorithms to make the comparison more reliable.

The parameter settings for SGD and GSGD are shown below in Table 1. Each algorithm was allowed to run for the maximum of 1000 iterations for 30 times. Due to small sized dataset 3-fold cross validation is done with 60:40 training-testing ratio. Neighborhood size (ρ) should not be too large as it will make the algorithm less efficient. Additionally, it is important to choose the appropriate value of ρ to avoid consistent instances becoming 'stale' and turning into inconsistent. GSGD's convergence is similar to SGD's where gradient descent happens more than once (for $\rho > 1$) in every iteration.

Table 1. Parameter setting for GSGD and SGD

Parameter	**Value**
Max iterations	1000
Attempts	30 consecutive runs
Validation	3-fold cross validation
Training:Testing ratio	60:40
Learning rate (η)	0.2
Neighborhood size (ρ)	10

5.3 Parameter tuning

The degree of descent depends on learning rate, slope of the gradient, and also inconsistency present in a data set as shown in Eq. (7). Gradient descent with consistent instances happen after every ρ iterations. To evaluate the influence of different values of ρ in the degree of convergence, contour plot have been drawn for the difference of in-sample error between GSGD and SGD in Fig. 9 (a). The plot is based on the average of 30 runs with the same parameter settings defined in Table 1. The x-axis shows the number of iterations and y-axis shows the range of values for ρ from 2 – 50. The lower the value of the contour plot, the better the performance of GSGD compared to SGD, and vice-versa. The selected data set was breast cancer diagnostic as it has produced smooth convergence for both SGD and GSGD, shown in Fig. 11. It is observed that $\rho \leq 10$ performs better compared to higher values of ρ.

Very low value of ρ may seem to perform better with low in-sample error but its CA is also not very high. Fig. 9 (b) shows that $6 < \rho < 11$ have produced good CAs. Hence, parameter $\rho = 10$ or little less is recommended to have a good convergence.

Non-parametric test was also performed with Wilcoxon test for ranking (two-tailed) to show null hypothesis with $p > 0.05$ (there is no significance difference between tested algorithms) is wrong. The test is conducted on 250th and 750th iteration where insignificant difference are

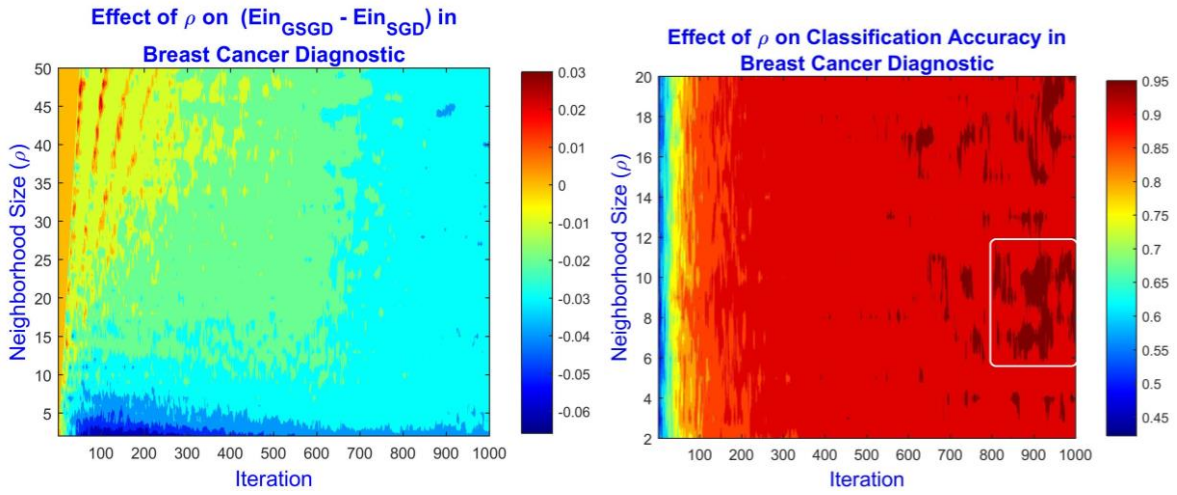


Fig. 9. Effect of ρ in degree of convergence

denoted by † and ‡ respectively in the result tables in the Appendix. Only in few instances it was shown that the guided and original algorithms have no significant difference. It may happen when the influence of inconsistent data is negligible or not many inconsistent data are present.

5.4 Experiment Execution

The comparative experimental results have been carried out between guided and original versions of the SGD algorithms. SGD has been tested against GSDG and similarly, Adadelta, Adagrad, Adam, Momentum, Nesterov, RMSprop have been tested against GAdadelta, GAdagrad, GAdam, GMomentum, GNesterov, and GRMSprop respectively. One to one comparison shows the influence of guided approach on a given original algorithm. The test results for the above mentioned algorithms are shown in the Appendix in Table A.1 – Table A.7 respectively. The datasets with removal of outliers are indicated as filtered datasets. The analysis is based on statistical testing of average, median, best and worst CAs together with Wilcoxon's no significant difference indicator († and/or ‡). The first column shows the tested

datasets, the second column indicates whether the algorithm is guided or original, and the third column indicates win%, which represents number of times guided (or original) algorithm got better result than the later (or former) out of 30 runs. Rest of the columns shows average, best, median and worst solutions with NFC after '@' symbol. The better results between the two approaches are shown in bold letters. Guided approach has outperformed original approaches in almost all the algorithms. The empirical results definitely establishes that avoiding inconsistent training examples regularly produces better results. Furthermore, among the tested variations of SGD, Adam and RMSprop has mostly produced the best results.

6. Discussion

GSGD has not only outperformed SGD on its canonical version but also in all other popular variations of SGD. Due to the extent of experimental results, the outcomes have been summarized in Fig. 10 where each bar shows the average classification accuracy of the best solutions obtained by tested variations of SGD on a given problem. The highlighted values on the top of bars show notable improvement with the proposed guided approach such as Pima Indian Diabetes (filtered and non-filtered), where the improvement in classification accuracy

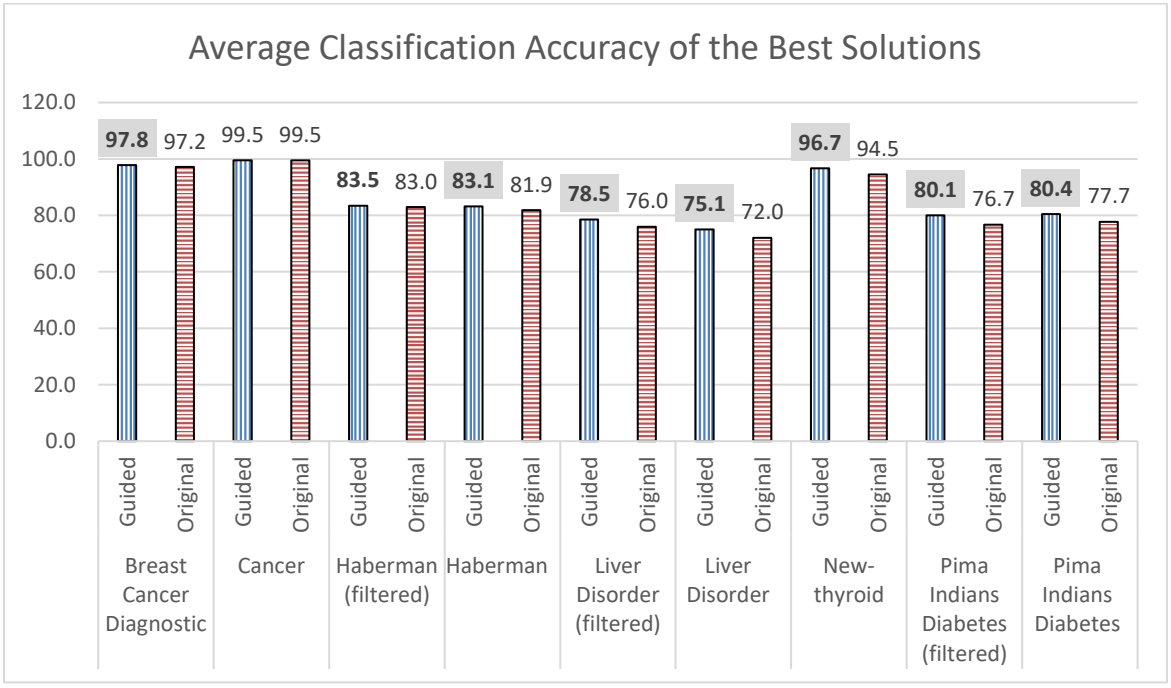


Fig. 10. Average classification accuracy of the best solutions for all the variations of SGD (and GSGD)

is by 3.4% and 2.7% respectively. Other notable improvement is shown in Liver Disorder (non-filtered and filtered) and New-thyroid, where classification accuracy is improved by 3.1%, 2.5% and 2.2% respectively. The selected graphs of convergence for canonical SGD and GSGD are shown in Fig. 11. The graphs show in-sample error (or training error) E_{in} with validation error E_v. Tracking of E_v has been done to avoid any over-fitting. Guided validation error is prefixed with a letter 'G'. The convergence of Haberman, Liver Disorder and Pima

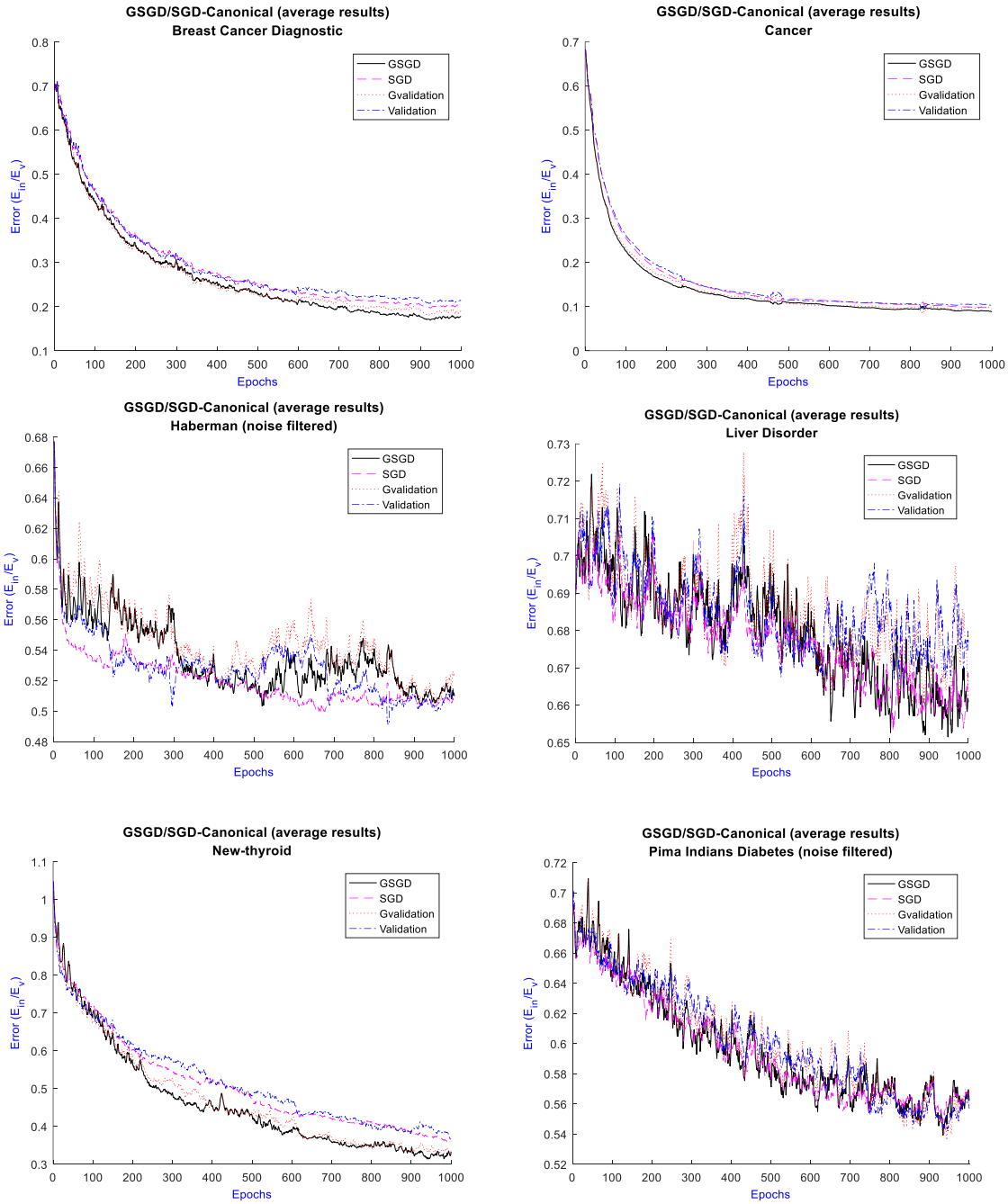


Fig. 11. Convergence graph of canonical SGD and GSGD for selected test problems

Indians diabetes data sets are quite chaotic including filtered ones. Their best CAs are also quite low compared to the other problems, because of high bias (high training and testing error) and high variance with the chosen model. Additionally, they have high data inconsistency and insufficient data. It was also observed that even after removing the outliers and extreme values, GSGD has performed better than SGD. For rest of the problems, it is evident that the guided approach's E_{in} and E_v are lower than the original approach throughout the training process. The purpose of GSGD is not to be used as an alternative or addition to the noise removal techniques, but to guide SGD when data inconsistency is high. Even if SGD would be used with a more complex model, most probably the testing error will not be improved unless sufficient and consistent data are provided.

7. Conclusion

This paper has introduced a guided search approach to improve canonical SGD, Adadelta, Adagrad, Adam, Momentum, Nesterov and RMSprop. The test results demonstrate that the guided approach has completely outperformed original approaches when used in logistic regression. In some cases, average classification accuracy has been improved by more than 3%. The proposed approach has realized that the inconsistent training examples leads to poorer classification accuracy. Additionally, its worst time complexity is also same as SGD's $O(T(d+N))$ and better than BGD's $O(TdN)$. Hence, it can be very useful for training large data sets efficiently. The future work would be to apply guided approach in deep learning and in asynchronous gradient descent where its efficiency should not be compromised. Guided approaches' weakness could be the usage of additional memory and time to filter out inconsistent training examples. However, its effect is still needed to be seen on online learning/ensemble learning. Finally, a standard metrics and measurements are required to be accommodated for the presence of inconsistency in a training data set.

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Bibliography

[1] Y. S. Abu-Mostafa, M. Magdon-Ismail, and H.-T. Lin, *Learning From Data*. S.l.: AMLBook, 2012.

[2] Y. Bengio, R. Ducharme, P. Vincent, and C. Jauvin, "A Neural Probabilistic Language Model," *Journal of Machine Learning Research*, vol. 3, no. Feb, pp. 1137–1155, 2003.

[3] C. M. Bishop, *Neural Networks for Pattern Recognition*. New York, NY, USA: Oxford University Press, Inc., 1995.

[4] V. Chandola, A. Banerjee, and V. Kumar, "Anomaly Detection: A Survey," *ACM Comput. Surv.*, vol. 41, no. 3, pp. 15:1–15:58, Jul. 2009.

[5] J. Dean et al., "Large Scale Distributed Deep Networks," in *Advances in Neural Information Processing Systems 25*, F. Pereira, C. J. C. Burges, L. Bottou, and K. Q. Weinberger, Eds. Curran Associates, Inc., 2012, pp. 1223–1231.

[6] D. Dheeru and E. Karra Taniskidou, *UCI Machine Learning Repository*. University of California, Irvine, School of Information and Computer Sciences, 2017.

[7] J. Duchi, E. Hazan, and Y. Singer, "Adaptive Subgradient Methods for Online Learning and Stochastic Optimization," *Journal of Machine Learning Research*, vol. 12, no. Jul, pp. 2121–2159, 2011.

[8] Eibe Frank, Mark A. Hall, and Ian H. Witten, "The WEKA Workbench," in *Online Appendix for "Data Mining: Practical Machine Learning Tools and Techniques,"* 4th ed., Morgan Kaufmann, 2016.

[9] M. T. Hoque, M. Chetty, and L. S. Dooley, "A new guided genetic algorithm for 2D hydrophobic-hydrophilic model to predict protein folding," in *2005 IEEE Congress on Evolutionary Computation*, 2005, vol. 1, pp. 259-266 Vol.1.

[10] Howard Anton, Irl Bivens, and Stephen Davis, "Partial Derivatives," in *Calculus, Early Transcendentals*, 10th ed., USA: John Wiley & Sons, Inc., 2012, pp. 906–999.

[11] R. Johnson and T. Zhang, "Accelerating Stochastic Gradient Descent Using Predictive Variance Reduction," in *Proceedings of the 26th International Conference on Neural Information Processing Systems*, USA, 2013, pp. 315–323.

[12] D. P. Kingma and J. Ba, "Adam: A Method for Stochastic Optimization," *arXiv:1412.6980 [cs]*, Dec. 2014.

[13] J. Konečný, J. Liu, P. Richtárik, and M. Takáč, "Mini-Batch Semi-Stochastic Gradient Descent in the Proximal Setting," *IEEE Journal of Selected Topics in Signal Processing*, vol. 10, no. 2, pp. 242–255, Mar. 2016.

[14] J. Laurikkala, M. Juhola, and E. Kentala, "Informal Identification of Outliers in Medical Data," in *Fifth International Workshop on Intelligent Data Analysis in Medicine and Pharmacology*, Berlin, Germany, 2000, pp. 20–24.

[15] Z. Michalewicz and D. B. Fogel, *How to Solve It: Modern Heuristics*, 2nd ed. Springer, 2004.

[16] M. Moreira and E. Fiesler, *Neural Networks with Adaptive Learning Rate and Momentum Terms*. INSTITUT DALLE MOLLE D'INTELLIGENCE ARTIFICIELLE PERCEPTIVE, 1995.

[17] K. Murota, *Discrete Convex Analysis: Monographs on Discrete Mathematics and Applications 10*. Philadelphia, PA, USA: Society for Industrial and Applied Mathematics, 2003.

[18] N. Qian, "On the momentum term in gradient descent learning algorithms," *Neural Networks*, vol. 12, no. 1, pp. 145–151, Jan. 1999.

[19] F. Rothlauf, *Design of Modern Heuristics: Principles and Application*, 1st ed. Springer Publishing Company, Incorporated, 2011.

[20] S. Ruder, “An overview of gradient descent optimization algorithms,” *arXiv:1609.04747 [cs]*, Sep. 2016.
[21] S. J. Russell and P. Norvig, *Artificial Intelligence - a modern approach*, 2nd ed. New Jersey: Pearson Education Inc, 2003.
[22] A. Sharma, “Information Discovery from Constraints for Evolutionary Computational Model (Phd Dissertation),” University of Canberra, 2014.
[23] A. Sharma and D. Sharma, “ICHEA – A Constraint Guided Search for Improving Evolutionary Algorithms,” in *In: Huang, T., Zeng, Z., Li, C., Leung, C.S. (eds.) ICONIP 2012, Part I. LNCS*, Doha, Qatar, 2012, vol. 7663, pp. 269–279.
[24] H. E. Solberg and A. Lahti, “Detection of Outliers in Reference Distributions: Performance of Horn’s Algorithm,” *Clinical Chemistry*, vol. 51, no. 12, pp. 2326–2332, Dec. 2005.
[25] C. Wang, X. Chen, A. J. Smola, and E. P. Xing, “Variance Reduction for Stochastic Gradient Optimization,” in *Advances in Neural Information Processing Systems 26*, C. J. C. Burges, L. Bottou, M. Welling, Z. Ghahramani, and K. Q. Weinberger, Eds. Curran Associates, Inc., 2013, pp. 181–189.
[26] D. R. Wilson and T. R. Martinez, “The general inefficiency of batch training for gradient descent learning,” *Neural Networks*, vol. 16, no. 10, pp. 1429–1451, Dec. 2003.
[27] Yaser S Abu-Mostafa, “Learning From Data - The Lectures,” *Learning From Data, California Institute of Technology*, 2016. [Online]. Available: https://work.caltech.edu/lectures.html. [Accessed: 30-Aug-2016].
[28] C.-C. Yu and B.-D. Liu, “A backpropagation algorithm with adaptive learning rate and momentum coefficient,” in *Proceedings of the 2002 International Joint Conference on Neural Networks, 2002. IJCNN ’02*, 2002, vol. 2, pp. 1218–1223.
[29] Yurii Nesterov, “A method for unconstrained convex minimization problem with rate of convergence O(1/k2),” in *Doklady AN SSSR (In Russian). (English translation. Soviet Math. Dokl*, 1983.
[30] M. D. Zeiler, “ADADELTA: An Adaptive Learning Rate Method,” *arXiv:1212.5701 [cs]*, Dec. 2012.
[31] S. Zheng et al., “Asynchronous Stochastic Gradient Descent with Delay Compensation,” in *International Conference on Machine Learning*, 2017, pp. 4120–4129.
[32] S. Zheng and J. T. Kwok, “Follow the Moving Leader in Deep Learning,” in *International Conference on Machine Learning*, 2017, pp. 4110–4119.

Appendix

Experimental results of guided and original SGD to compare classification accuracies

Table A.1. Classification accuracies of the canonical SGD against GSGD algorithm

Canonical SGD versus Guided SGD (GSGD)						
Datasets	**Algorithm**	**Win%**	**Average**	**Best**	**Median**	**Worst**
Breast Cancer Diagnostic	**Guided**	**96.70%**	**96.8@684**	**98.7@542**	96.5@663	95.2@809
	Original	3.30%	95.9@471	98.2@610	95.6@515	93.8@575
Cancer	**Guided**	**13.30%**	**98.0@81**	99.5@12	98.4@31	96.2@21
	Original	10.00%	97.9@58	99.5@24	98.4@29	95.1@32
Haberman (filtered) ‡	**Guided**	**70.00%**	**77.8@356**	**83.3@484**	78.1@279	71.9@169
	Original	0.00%	76.8@343	82.5@199	77.2@375	68.4@637
Haberman‡	**Guided**	**70.00%**	**77.3@446**	82.0@109	77.9@481	72.1@6
	Original	0.00%	75.0@339	82.0@91	75.4@400	68.9@1
Liver Disorder (filtered)	**Guided**	**60.00%**	**71.4@557**	76.7@679	71.3@616	65.9@805
	Original	16.70%	70.2@368	76.7@416	70.2@385	63.6@544
Liver Disorder	**Guided**	**66.70%**	**66.5@445**	**73.9@196**	65.9@435	62.3@406
	Original	20.00%	65.4@365	71.0@516	65.6@380	60.9@665
New-thyroid	**Guided**	**86.70%**	**96.4@376**	98.8@980	96.5@421	89.5@424
	Original	3.30%	91.6@438	98.8@567	91.9@475	77.9@687
Pima Indians Diabetes (filtered)	**Guided**	**100.00%**	**77.2@695**	**82.2@907**	77.4@733	74.2@576
	Original	0.00%	75.1@511	79.4@614	75.1@546	72.5@606
Pima Indians Diabetes	**Guided**	**93.30%**	**77.8@629**	**81.1@255**	78.0@631	74.6@210
	Original	0.00%	76.1@458	80.1@628	76.4@496	72.3@20

Table A.2. Classification accuracies of the Adadelta against GAdadelta algorithm

Original Adadelta versus Guided Adadelta (GAdadelta)						
Datasets	**Algorithm**	**Win%**	**Average**	**Best**	**Median**	**Worst**
Breast Cancer Diagnostic	**Guided**	**100.00%**	**92.9@471**	**96.0@679**	93.2@531	89.4@38
	Original	0.00%	90.6@162	93.4@90	90.7@108	84.6@12
Cancer	**Guided**	23.30%	**97.9@135**	99.5@847	97.8@46	96.2@48
	Original	23.30%	97.8@196	99.5@122	97.8@159	95.6@211
Haberman (filtered)	**Guided**	**16.70%**	**76.3@7**	80.7@1	76.3@2	71.1@1
	Original	0.00%	76.0@2	80.7@1	76.3@1	71.1@1
Haberman	**Guided**	**26.70%**	**74.2@14**	**79.5@32**	74.6@3	68.0@1
	Original	0.00%	74.0@4	78.7@30	73.8@1	68.0@1
Liver Disorder (filtered)	**Guided**	**96.70%**	**70.0@289**	**77.5@227**	70.5@227	58.9@286
	Original	3.30%	58.8@64	73.6@276	60.5@30	0.0@0
Liver Disorder	**Guided**	**96.70%**	**66.9@307**	**76.1@341**	66.7@273	59.4@14
	Original	3.30%	59.5@37	68.8@141	58.7@9	50.7@2
New-thyroid	**Guided**	**40.00%**	**77.8@13**	**93.0@28**	78.5@5	64.0@1
	Original	0.00%	75.4@4	90.7@3	75.6@3	64.0@1
Pima Indians Diabetes (filtered)	**Guided**	**96.70%**	**70.9@427**	**78.7@162**	70.7@474	63.4@24
	Original	0.00%	65.9@4	70.0@1	65.9@1	61.7@1
Pima Indians Diabetes	**Guided**	**100.00%**	**73.1@490**	**78.5@574**	72.6@505	69.1@147
	Original	0.00%	66.4@15	73.3@33	66.5@6	61.6@1

Table A.3. Classification accuracies of the Adagrad against GAdagrad algorithm

Original Adagrad versus Guided Adagrad (GAdagrad)						
Datasets	**Algorithm**	**Win%**	**Average**	**Best**	**Median**	**Worst**
Breast Cancer Diagnostic	**Guided**	**93.30%**	**94.2@569**	**96.0@480**	94.3@555	92.1@418
	Original	0.00%	93.3@413	95.6@313	93.4@411	90.3@387
Cancer	Guided	**23.30%**	**97.7@90**	99.5@322	97.8@32	96.2@18
	Original	6.70%	97.6@70	99.5@245	97.8@31	96.2@24
Haberman (filtered)	Guided	**76.70%**	**78.4@252**	84.2@202	78.9@172	71.9@1
	Original	0.00%	76.5@33	84.2@483	75.4@1	70.2@1
Haberman	**Guided**	**86.70%**	**76.3@240**	**83.6@125**	76.2@178	66.4@257
	Original	0.00%	74.0@119	80.3@82	73.8@8	65.6@1
Liver Disorder (filtered) †	**Guided**	**83.30%**	**71.9@545**	**78.3@223**	71.7@596	62.8@928
	Original	6.70%	69.4@338	74.4@302	70.2@296	58.1@197
Liver Disorder	**Guided**	**83.30%**	**68.6@540**	**76.8@175**	68.5@462	60.1@371
	Original	6.70%	65.5@228	72.5@66	65.2@171	57.2@23
New-thyroid	**Guided**	**100.00%**	**90.1@519**	**97.7@86**	90.1@603	83.7@897
	Original	0.00%	82.6@493	87.2@619	82.6@503	76.7@488
Pima Indians Diabetes (filtered)	**Guided**	**96.70%**	**72.4@531**	**78.0@459**	72.5@579	68.6@584
	Original	3.30%	69.7@296	74.6@99	69.3@282	64.8@365
Pima Indians Diabetes†	**Guided**	**100.00%**	**72.7@669**	**76.9@138**	72.5@746	69.4@807
	Original	0.00%	68.9@299	73.6@378	68.4@319	63.2@469

Table A.4. Classification accuracies of the Adam against GAdam algorithm

Original Adam versus Guided GAdam (GAdam)						
Datasets	**Algorithm**	**Win%**	**Average**	**Best**	**Median**	**Worst**
Breast Cancer Diagnostic	**Guided**	**56.70%**	**97.9@657**	99.1@871	98.2@684	95.2@867
	Original	13.30%	97.5@507	99.1@397	97.8@534	95.2@620
Cancer†‡	**Guided**	**23.30%**	**98.0@279**	100.0@9	98.1@197	96.7@220
	Original	0.00%	97.9@258	100.0@9	97.8@216	96.2@11
Haberman (filtered) †‡	**Guided**	**46.70%**	**80.1@353**	**84.2@388**	80.3@262	74.6@585
	Original	10.00%	79.5@280	83.3@343	79.8@329	72.8@351
Haberman†	**Guided**	**33.30%**	**79.5@451**	86.9@518	79.5@344	73.0@761
	Original	10.00%	79.2@303	86.9@395	79.5@311	71.3@47
Liver Disorder (filtered) †	**Guided**	**40.00%**	**74.3@555**	79.8@201	74.4@588	69.8@653
	Original	30.00%	73.9@372	79.8@556	74.0@337	70.5@517
Liver Disorder	**Guided**	**60.00%**	**71.8@598**	76.1@821	71.7@664	63.8@321
	Original	13.30%	70.8@372	76.1@269	71.0@363	63.0@130
New-thyroid	**Guided**	**23.30%**	**98.6@378**	100.0@699	98.8@232	95.3@877
	Original	13.30%	98.4@353	100.0@436	98.8@375	94.2@398
Pima Indians Diabetes (filtered) †	**Guided**	80.00%	77.3@662	**80.8@802**	77.2@648	74.2@433
	Original	16.70%	76.4@494	79.8@491	76.7@490	73.2@652
Pima Indians Diabetes	**Guided**	70.00%	77.8@692	**81.4@734**	77.7@725	74.6@832
	Original	16.70%	76.8@545	81.1@678	76.7@564	73.6@619

Table A.5. Classification accuracies of the Momentum against GMomentum algorithm

Original Momentum versus Guided Momentum (GMomentum)						
Datasets	**Algorithm**	**Win%**	**Average**	**Best**	**Median**	**Worst**
Breast Cancer Diagnostic	**Guided**	**100.00%**	**96.0@718**	**98.7@705**	95.8@739	93.0@938
	Original	0.00%	95.2@473	97.8@388	95.2@477	91.6@457
Cancer	Guided	6.70%	97.9@80	99.5@18	97.8@29	96.7@45
	Original	6.70%	97.9@76	99.5@16	97.8@29	96.7@31
Haberman (filtered) ‡	**Guided**	**76.70%**	**79.3@364**	**86.0@190**	79.8@333	71.1@8
	Original	0.00%	78.2@405	84.2@463	78.1@462	71.1@8
Haberman‡	**Guided**	**83.30%**	**77.8@454**	**85.2@159**	77.9@327	70.5@1
	Original	3.30%	75.1@282	83.6@215	74.6@360	68.9@5
Liver Disorder (filtered)	**Guided**	**63.30%**	**68.9@631**	**80.6@506**	69.0@630	62.8@204
	Original	20.00%	68.2@407	78.3@455	69.8@441	62.0@637
Liver Disorder	**Guided**	**63.30%**	**66.2@638**	**72.5@907**	65.9@700	60.1@969
	Original	20.00%	65.2@382	71.0@345	65.6@371	57.2@164
New-thyroid	**Guided**	**90.00%**	**96.8@566**	**100.0@387**	97.7@538	93.0@179
	Original	0.00%	91.5@402	98.8@258	91.9@407	82.6@30
Pima Indians Diabetes (filtered)	**Guided**	**90.00%**	**75.6@643**	**81.5@965**	75.4@695	71.8@806
	Original	3.30%	74.2@444	78.4@432	74.0@465	70.4@544
Pima Indians Diabetes	**Guided**	**80.00%**	**75.7@567**	**80.5@755**	75.6@580	72.3@551
	Original	6.70%	74.7@418	79.2@163	74.4@424	71.3@429

Table A.6. Classification accuracies of the Nesterov against GNesterov algorithm

Original Nesterov versus Guided Nesterov (GNesterov)						
Datasets	**Algorithm**	**Win%**	**Average**	**Best**	**Median**	**Worst**
Breast Cancer Diagnostic	**Guided**	**100.00%**	**95.3@407**	**98.2@558**	95.2@379	92.1@591
	Original	0.00%	94.2@284	97.4@519	94.3@244	90.7@167
Cancer	Guided	13.30%	97.7@60	98.9@18	97.8@28	96.7@122
	Original	6.70%	97.7@46	98.9@33	97.5@26	96.7@20
Haberman (filtered)	Guided	6.70%	74.9@3	79.8@1	75.4@1	70.2@1
	Original	3.30%	74.9@2	79.8@1	75.4@1	70.2@1
Haberman	**Guided**	**30.00%**	**74.5@28**	**82.8@297**	73.8@2	68.9@3
	Original	3.30%	74.0@3	79.5@23	73.8@1	68.9@3
Liver Disorder (filtered)	**Guided**	**100.00%**	**70.7@612**	**77.5@712**	71.3@716	64.3@723
	Original	0.00%	60.0@90	69.8@31	59.3@48	50.4@4
Liver Disorder	**Guided**	**100.00%**	**66.4@556**	**71.7@174**	66.3@442	60.1@248
	Original	0.00%	59.9@47	68.8@118	59.4@22	54.3@1
New-thyroid	**Guided**	**43.30%**	**78.9@132**	**87.2@27**	79.7@9	68.6@31
	Original	3.30%	76.8@5	86.0@9	76.2@2	65.1@1
Pima Indians Diabetes (filtered)	**Guided**	**96.70%**	**74.1@485**	**77.7@692**	74.4@546	70.0@163
	Original	0.00%	67.6@32	74.2@22	67.9@5	61.0@1
Pima Indians Diabetes	**Guided**	**100.00%**	**75.2@493**	**81.8@824**	74.8@478	71.0@891
	Original	0.00%	67.5@25	74.9@82	66.5@17	61.6@1

Table A.7. Classification accuracies of the RMSprop against GRMSprop algorithm

Original RMSprop versus Guided RMSprop (GRMSprop)						
Datasets	**Algorithm**	**Win%**	**Average**	**Best**	**Median**	**Worst**
Breast Cancer Diagnostic	Guided	**70.00%**	**96.3@706**	98.2@371	96.5@767	94.3@714
	Original	13.30%	95.8@560	**98.7@668**	95.8@622	93.8@665
Cancer	**Guided**	**13.30%**	**97.9@212**	99.5@317	97.8@72	96.2@26
	Original	3.30%	97.8@169	99.5@636	97.8@59	96.2@79
Haberman (filtered) †‡	**Guided**	**60.00%**	**81.2@408**	86.0@145	81.6@315	74.6@5
	Original	3.30%	80.6@253	86.0@479	80.7@278	74.6@5
Haberman †‡	**Guided**	**50.00%**	**78.1@451**	82.0@299	78.7@378	73.0@926
	Original	3.30%	77.5@249	82.0@242	78.3@239	72.1@57
Liver Disorder (filtered)	**Guided**	**73.30%**	**74.3@515**	79.1@790	75.6@524	67.4@211
	Original	13.30%	73.3@411	79.1@162	73.6@464	65.9@530
Liver Disorder	**Guided**	**70.00%**	**73.4@564**	**78.3@428**	73.6@583	68.1@517
	Original	10.00%	71.7@452	76.1@306	72.1@502	63.0@164
New-thyroid	Guided	46.70%	98.4@488	100.0@841	98.8@468	95.3@191
	Original	3.30%	97.9@277	100.0@284	97.7@224	95.3@207
Pima Indians Diabetes (filtered)	**Guided**	**93.30%**	**77.3@640**	**81.5@268**	77.7@567	73.2@442
	Original	6.70%	75.8@529	80.5@371	76.0@551	70.0@392
Pima Indians Diabetes	**Guided**	**93.30%**	**77.1@694**	**82.7@744**	77.0@741	73.6@424
	Original	3.30%	75.5@574	81.8@646	75.6@646	71.7@351