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A Nearest Neighbor Algorithm for Imbalanced Classification

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Due to the inability of the accuracy-driven methods to address the challenging problem of learning from imbalanced data, several alternative measures have been proposed in the literature, like the Area Under the ROC Curve (AUC), the Average Precision (AP), the F-measure, the G-Mean, etc. However, these latter measures are neither smooth, convex nor separable, making their direct optimization hard in practice. In this paper, we tackle the challenging problem of imbalanced learning from a nearest-neighbor (NN) classification perspective, where the minority examples typically belong to the class of interest. Based on simple geometrical ideas, we introduce an algorithm that rescales the distance between a query sample and any positive training example. This leads to a modification of the Voronoi regions and thus of the decision boundaries of the NN classifier. We provide a theoretical justification about this scaling scheme which inherently aims at reducing the False Negative rate while controlling the number of False Positives. We further formally establish a link between the proposed method and cost-sensitive learning. An extensive experimental study is conducted on many public imbalanced datasets showing that our method is very effective with respect to popular Nearest-Neighbor algorithms, comparable to state-of-the-art sampling methods and even yields the best performance when combined with them.

Keywords: Machine Learning; Nearest Neighbor Algorithm; Imbalanced Classification.

1. Introduction

While the machine learning community can benefit nowadays from larger and larger datasets for optimizing provably accurate classifiers, many real world applications still suffer from a lack of data, especially in imbalanced learning, where the positive examples are very scarce compared with the number of negative samples [1–3]. This is typically the case in intrusion detection, health care insurance or bank fraud identification, and more generally anomaly detection, *e.g.*, in medicine or in industrial

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processes. In such a setting, the training set is composed of a few positive examples (*e.g.*, the frauds) and a huge amount of negative samples (*e.g.*, the genuine transactions). Standard learning algorithms struggle to deal with this imbalance scenario because they are typically based on the minimization of (a surrogate of) the 0-1 loss. Therefore, a trivial solution consists in assigning the majority label to any test query, leading to a high performance from an accuracy perspective but completely missing the (positive) examples of interest. To overcome this issue, several strategies have been developed over the years. The first one consists in the optimization of loss functions based on measures that are more appropriate for this context such as the *Area Under the ROC Curve* (AUC), the *Average Precision* (AP), the *G-mean* (GM), the *Balanced-Accuracy* (BA) or the *F-measure* to cite a few [4–6]. The main pitfalls related to such a strategy concern the difficulty to directly optimize non smooth, non separable and non convex measures (see [7] for the specific case of the *F-measure*). A simple and usual solution to fix this problem consists in using off-the-shelf learning algorithms (maximizing the accuracy) and a posteriori pick the model with the highest AP, GM, BA or *F-measure*. Unfortunately, this might be often suboptimal. A more elaborate solution aims at designing differentiable versions of the previous non-smooth measures and optimizing them, *e.g.*, as done by gradient boosting in [8] with a smooth surrogate of the Mean-AP. The second family of methods is based on the modification of the distribution of the training data using sampling strategies [9]. This is typically achieved by removing examples from the majority class, as done, *e.g.*, in *ENN* or *Tomek's Link* [10], and/or by adding examples from the minority class, as in *SMOTE* [11] and its variants, or by resorting to generative adversarial models [12]. One peculiarity of imbalanced learning can be interpreted from a geometric perspective. As illustrated in Fig. 1 (left) which shows the Voronoi cells on an artificial imbalanced dataset (where two adjacent cells have been merged if they concern examples of the same class), the regions of influence of the positive examples are much smaller than that of the negatives. This explains why at test time, in imbalanced learning, the risk to get a false negative (*e.g.*, a fraud that is wrongly classified as a genuine transaction) is high. A large number of false negatives (*FN*) leads to a dramatic decrease of the aforementioned measures that all rely on a fine balance between *FN* and the number of false positives *FP*, building blocks of the so-called *Precision* = $\frac{TP}{TP+FP}$ and *Recall* = $\frac{TP}{TP+FN}$ where *TP* is the number of true positives. Note that increasing the regions of influence of the positives would mechanically reduce *FN*. However, not controlling the expansion of these regions, as illustrated in Fig. 1 (right), may have a dramatic impact on *FP*, and thus on the previous performance measures.

The main contribution of this paper is about the problem of finding the appropriate trade-off (Fig. 1 (middle)) between the two above-mentioned extreme situations (large *FP* or *FN*, both leading to a poor performance at test time). A natural way to increase the influence of positives may consist in using generative models (like GANs [12]) to sample new artificial examples, mimicking the negative training samples. However, beyond the issues related to the parameter tuning, the computation

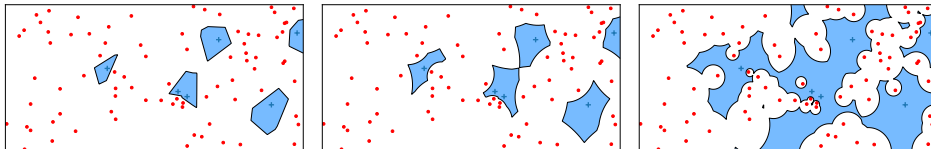


Fig. 1. Toy imbalanced dataset: On the left, the Voronoi regions around the positives are small. The risk to generate false negatives (FN) at test time is large. On the right: by increasing too much the regions of influence of the positives, the probability to get false positives (FP) grows. In the middle: an appropriate trade-off between the two previous situations.

burden and the complexity of such a method, using GANs to optimize the precision and recall is still an open problem (see [13] for a recent study on this topic). We show in this paper that a much simpler strategy can be used by modifying the distance exploited in a k -nearest neighbor (k -NN) algorithm [14] which enjoys many interesting advantages, including its simplicity, its capacity to approximate asymptotically any locally regular density, and its theoretical rootedness [15–17]. k -NN also benefited from many algorithmic advances during the past decade in the field of metric learning, aiming at optimizing under constraints the parameters of a metric, typically the Mahalanobis distance, as done in LMNN [18] or ITML [19] (see [20, 21] for a survey). Unfortunately, existing metric learning methods are dedicated to enhance the k -NN accuracy and do not focus on the optimization of criteria, like the F-measure or G-mean, in scenarios where the positive training examples are scarce. A geometric solution to increase, at a very low cost, the region of influence of the minority class consists in modifying the distance when comparing a query example to a positive training sample. More specifically, we formally show in this paper that the optimization of any (FN, FP) -based performance measure, which are well suited to deal with imbalanced scenarios, is facilitated by scaling the distance to any positive by a coefficient $\gamma \in [0, 1]$ leading to the expansion of the Voronoi cells around the minority examples. An illustration is given in Fig. 1 (middle) which might be seen as a good compromise that results in the reduction of FN while controlling the risk to increase FP . Note that our strategy boils down to modifying the local density of the positive examples. For this reason, we claim that it can be efficiently combined with SMOTE-based sampling methods whose goal is complementary and consists in generating examples on the path linking two (potentially far) positive neighbors. Our experiments will confirm this intuition.

This paper improves substantially on our previous work [22], both with increased details and new algorithmic and experimental contributions: (i) we show an explicit link between the proposed method, called γk -NN, and cost-sensitive learning, (ii) we present a local version of our method that uses clustering to adapt the parameters to the different regions of the input space, and (iii) we rework and extend the experimental study to incorporate new performance measures and to give a qualitative analysis on the well-known image dataset MNIST.

The rest of the paper is organized as follows. Section 2 is dedicated to the

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introduction of our notations and an overview of the main performance measures that will be used to evaluate the compared methods. The related work is presented in Section 3. Section 4 is devoted to the presentation of our method γk -NN. This section includes a theoretical analysis of our method as well as a presentation of a local extension aiming at capturing local specificities of the feature space. We finally establish a link between γk -NN and cost-sensitive learning. The last part of this paper is dedicated to an extensive experimental study performed on 28 imbalanced datasets (see Section 5). In this comparative analysis, we give evidence of the complementarity of our method with sampling strategies. We finally conclude in Section 6.

2. Notations and Evaluation Measures

We consider a training sample $S = \{(\mathbf{x}_i, y_i), i = 1, \dots, m\}$ of size m , drawn from an unknown joint distribution $\mathcal{Z} = \mathcal{X} \times \mathcal{Y}$, where $\mathcal{X} = \mathbb{R}^p$ is the feature space and $\mathcal{Y} = \{-1, 1\}$ is the set of labels. Let us assume that $S = S_+ \cup S_-$ with m_+ positives $\in S_+$ and m_- negatives $\in S_-$ where $m = m_+ + m_-$.

Learning from imbalanced datasets requires to optimize appropriate measures that take into account the scarcity of positive examples. Several of them rely on the following two quantities: the Recall (also called *True Positive Rate (TPR)* or *sensitivity*) which measures the capacity of the model to recall/detect positive examples, and the Precision (also called *Positive Prediction Value (PPV)*) which is the confidence in the prediction of a positive label. They are defined as follows:

$$\text{Recall} = \frac{TP}{TP + FN} \quad \text{and} \quad \text{Precision} = \frac{TP}{TP + FP},$$

where FP (resp. FN) is the number of false positives (resp. negatives) and TP is the number of true positives. Since one can arbitrarily improve the Precision if there is no constraint on the Recall (and vice-versa), they are usually combined into a single measure.

For instance, the *F-measure* [23] (or F_β score), which is widely used in fraud and anomaly detection [24], is defined as the harmonic mean of the Recall and Precision:

$$F_\beta = (1 + \beta^2) \frac{\text{Precision} \times \text{Recall}}{\beta^2 \times \text{Precision} + \text{Recall}},$$

where β is set such that the Recall is considered β times as important as the Precision. Note that F_1 (*i.e.* $\beta = 1$) considers the Precision and Recall equally.

The *G-measure* (G_1) can also be used for imbalanced data classification [25]. Unlike F_1 , it rather considers the geometric mean of Precision and Recall:

$$G_1 = \sqrt{\text{Precision} \times \text{Recall}}.$$

While F_β and G_1 consider both Recall and Precision, the *G-mean (GM)* [26] rather makes use of the Recall (or TPR) and the False Negative Rate (TNR) as follows:

$$GM = \sqrt{TPR \times TNR} = \sqrt{\frac{TP}{TP + FN} \times \frac{TN}{TN + FP}}.$$

In other words, it computes the geometric mean of TPR and TNR . Thus, it gives a higher importance to the negative class, compared to the previous measures. Without being exhaustive, a last performance measure that can be used in an imbalanced setting is the *Balanced Accuracy (BA)* [4] which also relies on TPR and TNR and is defined as the mean accuracy of the two classes:

$$BA = (TPR + TNR)/2$$

In the experimental section of this paper, we will resort to these widely used measures to assess the efficiency of our proposed method to overcome the problem of scarcity of positive samples.

3. Related Work

In this section, we present the main strategies that have been proposed in the literature to address the problem of learning from imbalanced datasets. We first present methods specifically dedicated to enhance a k -NN classifier. Then, we give an overview of the main sampling strategies used to balance the two classes. All these methods will be used in the experimental comparison in Section 5.

3.1. Distance-based Methods

Several strategies have been devised to improve k -NN. The oldest method is certainly the one presented in [27] which consists in associating to each neighbor a voting weight that is inversely proportional to its distance to a query point \mathbf{x} . The assigned label \hat{y} of \mathbf{x} is defined as:

$$\hat{y} = \sum_{\mathbf{x}_i \in \text{kNN}(\mathbf{x})} y_i \times \frac{1}{d(\mathbf{x}, \mathbf{x}_i)},$$

where $\text{kNN}(\mathbf{x})$ stands for the set of the k nearest neighbors of \mathbf{x} . A more refined version consists in taking into account both the distances to the nearest neighbors and the distribution of the features according to the class $p(\mathbf{x}_i | y_i)$ [28]. Despite these modifications in the decision rule, the sparsity of the positives remains problematic and it is possible that no positives fall in the neighborhood of a new query \mathbf{x} . To tackle this issue, a variant of k -NN, called $kPNN$ [29], is to consider the region of the space around a new query \mathbf{x} which contains exactly k positive examples. By doing so, the authors are able to use the density of the positives to estimate the probability of belonging in the minority class.

A more recent version has been shown to perform better than the two previous approaches: $kRNN$ [30]. If the idea remains similar (*i.e.* estimating the local sparsity of minority examples around a new query), the posterior probability of belonging in the minority class is adjusted so that it considers both the local and global disequilibrium for the estimation. In [31], the authors use both the label and the distance to the neighbors (\mathbf{x}_i, y_i) to define a scaled metric d' from the Euclidean

distance d , as follows:

$$d'(\mathbf{x}, \mathbf{x}_i) = \left(\frac{m_i}{m}\right)^{1/p} d(\mathbf{x}, \mathbf{x}_i),$$

where m_i is the number of examples in the class y_i . As we will see later, this method falls in the same family of strategies as our contribution, aiming at scaling the distance to the examples according to their label. However, three main differences explain the superiority of our method, observed in the experiments: (i) $kRNN$ fixes d' in advance while we will automatically adapt the scaling factor to optimize the considered performance measure; (ii) because of (i), d' needs to take into account the dimension p of the feature space (and so will tend to d as p grows) while our method captures the intrinsic dimension of the space by selecting the best weight; (iii) d' is useless when combined with sampling strategies (indeed, $\frac{m_i}{m}$ would tend to be uniform) while our method will allow us to scale differently the distance to the original positive examples and the ones artificially generated.

Another way to assign weights to each class, which is close to the sampling methods presented in the next section, is to duplicate the positive examples according to the Imbalance Ratio $IR = m_-/m_+$. Thus, it can be seen as a *uniform* over-sampling technique, where all positives are replicated the same number of times. However, note that this method requires to work with $k > 1$.

A last family of methods that try to improve k -NN is related to *metric learning* [20, 21]. LMNN [18] or ITML [19] are two famous examples which optimize under constraints a Mahalanobis distance $d_{\mathbf{M}}(\mathbf{x}, \mathbf{x}_i) = \sqrt{(\mathbf{x} - \mathbf{x}_i)^\top \mathbf{M}(\mathbf{x} - \mathbf{x}_i)}$ parameterized by a positive semi-definite (PSD) matrix \mathbf{M} . Such methods seek a linear projection of the data in a latent space where the Euclidean distance is applied. As we will see in the following, our scaling method is a special case of metric learning which looks for a diagonal matrix (but applied only when comparing a query to a positive example) and which behaves well whatever the considered performance measure.

3.2. Sampling Strategies

One way to overcome the issues induced by the lack of positive examples is to compensate artificially the imbalance between the two classes. Sampling strategies [9] have been proven to be very efficient to address this problem. In the following, we overview the most used methods in the literature.

The Synthetic Minority Over-sampling Technique [11] (SMOTE) over-samples a dataset by creating new synthetic positive data. For each minority example \mathbf{x} , it randomly selects one of its k nearest positive neighbors and then creates a new random positive point on the line between this neighbor and \mathbf{x} . This is done until some desired ratio is reached.

Borderline-SMOTE [32] is an improvement of the SMOTE algorithm. While the latter generates synthetic points from all positive points, BorderLine-SMOTE only focuses on those having more negatives than positives in their neighborhood.

More precisely, new samples are generated if the number n of negatives in the k -neighborhood is such that $k/2 \leq n \leq k$.

The Adaptive Synthetic [33] (ADASYN) sampling approach is also inspired from SMOTE. By using a weighted distribution, it gives more importance to classes that are more difficult to classify, *i.e.* where positives are surrounded by many negatives, and thus generates more synthetic data for these classes.

Two other strategies combine an over-sampling step with an under-sampling procedure. The first one uses the Edited Nearest Neighbors [34] (ENN) algorithm on the top of SMOTE. After SMOTE has generated data, the ENN algorithm removes samples that are misclassified by their k nearest neighbors. The second one combines SMOTE with Tomek's link [10]. The latter is a pair of points $(\mathbf{x}_i, \mathbf{x}_j)$ from different classes for which there is no other point \mathbf{x}_k verifying $d(\mathbf{x}_i, \mathbf{x}_k) \leq d(\mathbf{x}_i, \mathbf{x}_j)$ or $d(\mathbf{x}_k, \mathbf{x}_j) \leq d(\mathbf{x}_i, \mathbf{x}_j)$. In other words, \mathbf{x}_i is the nearest neighbor of \mathbf{x}_j and vice-versa. If so, one removes the example of $(\mathbf{x}_i, \mathbf{x}_j)$ that belongs to the majority class. Note that both strategies tend to eliminate the overlapping between classes.

Interestingly, we can note that all the previous sampling methods try to overcome the problem of learning from imbalanced data by resorting to the notion of k -neighborhood. This is justified by the fact that k -NN has been shown to be a good estimate of the density at a given point in the feature space.

In our contribution, we also leverage k -NN but with a different approach. Instead of generating (many) new examples (which would have a negative impact from a complexity perspective), we locally modify the density around the positive points. We achieve this by rescaling the distance between a test sample and the positive training examples. We show that such a strategy can be efficiently combined with sampling methods, whose goal is complementary, by potentially generating new examples in regions of the space where the minority class is not present.

4. Proposed Approach

In this section, we present our γk -NN method which works by scaling the distance between a query point and positive training examples by a factor.

4.1. An Adjusted k -NN algorithm

Statistically, when learning from imbalanced data, a new query \mathbf{x} has more chance to be close to a negative example due to the rarity of positives in the training set, even around the mode of the positive distribution. We have seen two families of approaches that can be used to counteract this effect: (i) creating new synthetic positive examples, and (ii) changing the distance according to the class. The approach we propose falls into the second category.

We suggest to modify how the distance to the positive examples is computed, in order to compensate for the imbalance in the dataset. We artificially bring a new query \mathbf{x} closer to any positive data point $\mathbf{x}_i \in S_+$ in order to increase the effective area of influence of positive examples. The new measure d_γ that we propose is

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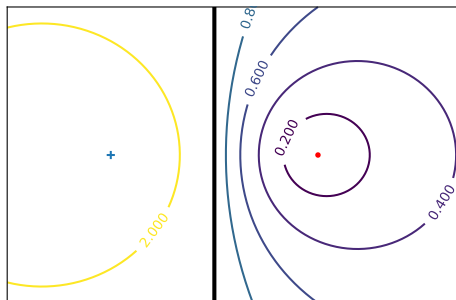


Fig. 2. Evolution of the decision boundary based on d_γ , for a 1-NN classifier, on a 2D dataset with one positive (resp. negative) instance represented by a blue cross (resp. red point). The value of γ is given on each boundary ($\gamma = 1$ on the thick line).

defined using an underlying distance d (e.g., the Euclidean distance) as follows:

$$d_\gamma(\mathbf{x}, \mathbf{x}_i) = \begin{cases} d(\mathbf{x}, \mathbf{x}_i) & \text{if } \mathbf{x}_i \in S_-, \\ \gamma \cdot d(\mathbf{x}, \mathbf{x}_i) & \text{if } \mathbf{x}_i \in S_+. \end{cases} \quad (1)$$

As we will tune the γ parameter, this new way to compute the similarity to a positive example is close to a Mahalanobis-distance learning algorithm, looking for a PSD matrix, as previously described. However, the matrix \mathbf{M} is restricted here to be $\gamma^2 \cdot \mathbf{I}$, where \mathbf{I} refers to the identity matrix. Moreover, while metric learning typically works by optimizing a convex loss function under constraints, our γ is simply tuned such as maximizing the non convex performance measure. Lastly, and most importantly, it is applied only when comparing the query to positive examples. As such, d_γ is not a proper distance. However, this is what allows it to compensate for the class imbalance. In the binary setting, there is no need to have a γ parameter for the negative class, since only the relative distances are used. In the multi-class setting with K classes, we would have to tune up to $K - 1$ values of γ .

Before formalizing the γk -NN algorithm that will leverage the distance d_γ , we illustrate in Fig. 2, on 2D data, the decision boundary induced by a nearest neighbor binary classifier that uses d_γ . We consider an elementary dataset with only two points, one positive and one negative. The case of $\gamma = 1$, which is a traditional 1-NN is shown in a thick black line. Lowering the value of γ below 1 brings the decision boundary closer to the negative point, and eventually tends to surround it very closely. Fig 3 shows, with more complex (toy) datasets, that γ controls how much we want to push the boundary towards negative examples. Fig 3 (right) should be imagined as a zoomed-in boundary between the classes, where one class is 20 times less represented. It shows that, due to sampling, the 1-NN boundary wrongly causes regions of false negatives, while γk -NN is able to correct the bias.

We can now present γk -NN (see Algorithm 1) that is parameterized by the γ parameter. It has the same overall complexity as k -NN. The first step to classify

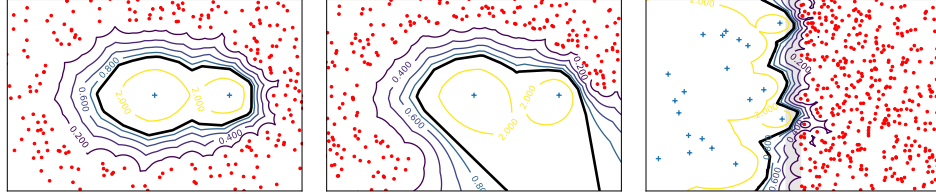


Fig. 3. Behavior of the decision boundary according to the γ value for the 1-NN classifier on toy datasets. Positive points are shown as blue crosses and negatives ones as red dots. The black line represents the standard decision boundary for the 1-NN classifier, *i.e.* when $\gamma = 1$.

a query \mathbf{x} is to find its k nearest negative neighbors and its k nearest positive neighbors. Then, the distances to the positive neighbors are multiplied by γ , to obtain d_γ . These $2k$ neighbors are then ranked and the k closest ones are used for classification (with a majority vote, as in k -NN). It should be noted that, although d_γ does not define a proper distance, we can still use any existing fast nearest neighbor search algorithm, because the actual search is done only using the original distance d (but twice, once for S_+ , once for S_-).

Algorithm 1: Classification of a new example with γk -NN.

Input : a query \mathbf{x} to be classified, a set of labeled samples $S = S_+ \cup S_-$,
 a number of neighbors k , a positive real value γ , a distance function d

Output: the predicted label of \mathbf{x}

$\mathcal{NN}^-, \mathcal{D}^- \leftarrow nn(k, \mathbf{x}, S_-)$ // nearest negative neighbors with their distances

$\mathcal{NN}^+, \mathcal{D}^+ \leftarrow nn(k, \mathbf{x}, S_+)$ // nearest positive neighbors with their distances

$\mathcal{D}^+ \leftarrow \gamma \cdot \mathcal{D}^+$

$\mathcal{NN}_\gamma \leftarrow firstK(k, sortedMerge((\mathcal{NN}^-, \mathcal{D}^-), (\mathcal{NN}^+, \mathcal{D}^+)))$

$y \leftarrow \oplus$ if $|\mathcal{NN}_\gamma \cap \mathcal{NN}^+| \geq \frac{k}{2}$ else \ominus // majority vote based on \mathcal{NN}_γ

return y

4.2. Theoretical analysis

In this section, we formally analyze what could be a good range of values for γ in our γk -NN algorithm. To this aim, we study what impact γ has on the probability to get a false positive (and false negative) at test time and explain why it is important to choose $\gamma < 1$ when the imbalance in the data is significant. The following analysis is made for $k = 1$ but note that the conclusion still holds for $k > 1$.

Proposition 1. (*False Negative probability*) Let $d_\gamma(\mathbf{x}, \mathbf{x}_+) = \gamma d(\mathbf{x}, \mathbf{x}_+)$, $\forall \gamma > 0$, be our modified distance used between a query \mathbf{x} and any positive training example \mathbf{x}_+ , where $d(\mathbf{x}, \mathbf{x}_+)$ is some distance function. Let $FN_\gamma(\mathbf{z})$ be the probability for a positive example \mathbf{z} to be a false negative using Algorithm (1). The following result

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holds: if $\gamma \leq 1$,

$$FN_\gamma(\mathbf{z}) \leq FN(\mathbf{z})$$

Proof. (sketch of proof) Let ϵ be the distance from \mathbf{z} to its nearest-neighbor $N_{\mathbf{z}}$. \mathbf{z} is a false negative if $N_{\mathbf{z}} \in S_-$ that is all positives $\mathbf{x}' \in S_+$ are outside the sphere $\mathcal{S}_{\frac{\epsilon}{\gamma}}(\mathbf{z})$ centered at \mathbf{z} of radius $\frac{\epsilon}{\gamma}$. Therefore,

$$\begin{aligned} FN_\gamma(\mathbf{z}) &= \prod_{\mathbf{x}' \in S_+} \left(1 - P(\mathbf{x}' \in \mathcal{S}_{\frac{\epsilon}{\gamma}}(\mathbf{z}))\right), \\ &= \left(1 - P(\mathbf{x}' \in \mathcal{S}_{\frac{\epsilon}{\gamma}}(\mathbf{z}))\right)^{m_+} \end{aligned} \quad (2)$$

while

$$FN(\mathbf{z}) = \left(1 - P(\mathbf{x}' \in \mathcal{S}_\epsilon(\mathbf{z}))\right)^{m_+}. \quad (3)$$

Solving (2) \leq (3) implies $\gamma \leq 1$. \square

This result means that satisfying $\gamma < 1$ allows us to increase the decision boundary around positive examples (as illustrated in Fig. 3), yielding a smaller risk to get false negatives at test time. An interesting comment can be made from Eq.(2) and (3) about their convergence. As m_+ is supposed to be very small in imbalanced datasets, the convergence of $FN(\mathbf{z})$ towards 0 is pretty slow, while one can speed-up this convergence with $FN_\gamma(\mathbf{z})$ by increasing the radius of the sphere $\mathcal{S}_{\frac{\epsilon}{\gamma}}(\mathbf{z})$, that is taking a small value for γ .

Proposition 2. (*False Positive probability*) Let $FP_\gamma(\mathbf{z})$ be the probability for a negative example \mathbf{z} to be a false positive using Algorithm (1). The following result holds: if $\gamma \geq 1$,

$$FP_\gamma(\mathbf{z}) \leq FP(\mathbf{z})$$

Proof. (sketch of proof) Using the same idea as before, we get:

$$\begin{aligned} FP_\gamma(\mathbf{z}) &= \prod_{\mathbf{x}' \in S_-} \left(1 - P(\mathbf{x}' \in \mathcal{S}_{\gamma\epsilon}(\mathbf{z}))\right), \\ &= \left(1 - P(\mathbf{x}' \in \mathcal{S}_{\gamma\epsilon}(\mathbf{z}))\right)^{m_-} \end{aligned} \quad (4)$$

while

$$FP(\mathbf{z}) = \left(1 - P(\mathbf{x}' \in \mathcal{S}_\epsilon(\mathbf{z}))\right)^{m_-}. \quad (5)$$

Solving (4) \leq (5) implies $\gamma \geq 1$. \square

As expected, this result suggests to take $\gamma > 1$ to increase the distance $d_\gamma(\mathbf{z}, \mathbf{x}_+)$ from a negative test sample \mathbf{z} to any positive training example \mathbf{x}_+ and thus reduce the risk to get a false positive. It is worth noticing that while the two conclusions from Propositions 1 and 2 are contradictory, the convergence of $FP_\gamma(\mathbf{z})$ towards 0

Table 1. Cost matrix for a binary classification task.

	Predicted positive	Predicted negative
Actual positive	c_{TP}	c_{FN}
Actual negative	c_{FP}	c_{TN}

is much faster than that of $FN_\gamma(\mathbf{z})$ because $m_- \gg m_+$ in an imbalance scenario. Therefore, fulfilling the requirement $\gamma > 1$ is much less important than satisfying $\gamma < 1$. For this reason, we will impose our Algorithm (1) to take $\gamma \in [0, 1]$.

4.3. Link with cost-sensitive learning

In this section, we show that it is possible to establish a link between the cost-sensitive learning framework [35] and our algorithm γk -NN. The goal of cost-sensitive learning is to assign different costs to each entry of the confusion matrix as depicted in Table 1 for a binary setting where we will denote the 4 costs as c_{TP} , c_{FN} , c_{FP} and c_{TN} . Cost sensitive methods are widely used, including in imbalanced scenarios, to give more importance (*i.e.* higher weights/costs) to the examples of the positive/minority class. By doing so, a learned classifier will focus more on decreasing the loss associated to the positive samples. We show here that, despite not being learned by optimizing a loss function, γk -NN can still be seen in the lens of cost-sensitive learning.

Let us assume that the correct predictions are not penalized, *i.e.* $c_{TN} = c_{TP} = 0$ and that c_{FP} and c_{FN} are such that $c_{FP} + c_{FN} = 1$ (without loss of generality, as only their relative values matter here). Let \mathbf{x}^- (resp. \mathbf{x}^+) be the nearest negative (resp. positive) neighbor of an example \mathbf{x} . Suppose that we have a model $\eta(\mathbf{x}) = \mathbb{P}(y = 1 \mid \mathbf{x})$ that gives the probability for \mathbf{x} to be positive. Then the positive label will be assigned to \mathbf{x} if $\eta(\mathbf{x}) > 1/2$, without considering the costs of misclassification. Taking these latter into account changes the classification rule. Indeed, to minimize the cost-sensitive risk, an example \mathbf{x} must be predicted positive if:

$$\begin{aligned} c_{FP} \mathbb{P}(y = 0 \mid \mathbf{x}) &\leq c_{FN} \mathbb{P}(y = 1 \mid \mathbf{x}), \\ \Leftrightarrow c_{FP} (1 - \eta(\mathbf{x})) &\leq c_{FN} \eta(\mathbf{x}), \\ \Leftrightarrow \eta(\mathbf{x}) &\geq c_{FP}. \end{aligned}$$

On the other hand, our algorithm γk -NN classifies an example \mathbf{x} as positive if $d(\mathbf{x}, \mathbf{x}^-) > \gamma d(\mathbf{x}, \mathbf{x}^+)$. Given this classification rule, we can show that γk -NN resorts to an approximation $\hat{\eta}(\mathbf{x})$ of $\eta(\mathbf{x})$ for a given weighted problem, as follows:

$$\begin{aligned} d(\mathbf{x}, \mathbf{x}^-) &> \gamma d(\mathbf{x}, \mathbf{x}^+), \\ \Leftrightarrow d(\mathbf{x}, \mathbf{x}^-)(1 + \gamma) &> \gamma(d(\mathbf{x}, \mathbf{x}^+) + d(\mathbf{x}, \mathbf{x}^-)), \\ \Leftrightarrow \frac{d(\mathbf{x}, \mathbf{x}^-)}{d(\mathbf{x}, \mathbf{x}^+) + d(\mathbf{x}, \mathbf{x}^-)} &> \frac{\gamma}{1 + \gamma}. \end{aligned}$$

Setting $c_{FP} = \frac{\gamma}{\gamma + 1}$ (and therefore $c_{FN} = 1 - \frac{\gamma}{\gamma + 1} = \frac{1}{\gamma + 1}$) and $\hat{\eta}(\mathbf{x}) =$

$\frac{d(\mathbf{x}, \mathbf{x}^-)}{d(\mathbf{x}, \mathbf{x}^+) + d(\mathbf{x}, \mathbf{x}^-)}$ finishes to establish the link between γk -NN and cost-sensitive learning. Note that if $\gamma = 1$ then $c_{FP} = c_{FN} = \frac{1}{2}$ implying that we retrieve a standard k -NN classifier which treats positive and negative samples equally without cost sensitivity.

The reader interested in cost-sensitive k -NN classifiers can refer to [36, 37].

4.4. *Towards a local approach of γk -NN*

In what have been presented so far, we consider a single γ for the whole input space. While this has the advantage of having a single parameter to tune, it removes the ability to capture non-stationary class imbalance. Indeed, it is possible that a γ value is optimal in one part of the space but not in another.

We thus propose a non-stationary version of our algorithm, called local- γk -NN. Conceptually, we could have a $\gamma_{\mathbf{x}}$ for every position \mathbf{x} in the space. However, such an over parameterized model would loose the simplicity of the proposed approach and increase the risk of overfitting. To deal with these two issues, we rather partition the input space into $q \in \mathbb{N}^*$ sub-spaces, $\{C_j\}_{j=1}^q$, using a clustering algorithm (*e.g.*, k-means). Then a value γ_j , for all $j = 1, \dots, q$ is tuned according to the performance measure of interest and using only the available data in the subspace C_j . To classify a test query that falls in cluster j we use γk -NN (with γ_j) in this cluster. We will show in the experimental Section 5.4 two possible variants of this local approach.

5. Experiments

This part is devoted to an extensive experimental evaluation of γk -NN on public datasets with comparisons to classic distance-based methods and state-of-the-art sampling strategies able to deal with imbalanced data. All the results are reported for nearest neighbor classification with $k = 1$ and 3 by considering the four different evaluation measures introduced in Section 2 (F_1 , G_1 , GM and BA). We also conduct, in Section 5.3, a qualitative analysis on the behavior of our approach on the famous MNIST image dataset [38]. Finally, we conclude our experimental study by an evaluation of the performance of the local version of γk -NN (in Section 5.4).

5.1. *Experimental setup*

For these experiments, we use 28 public datasets from the well-known UCI^a and KEEL^b repositories. The main properties of these datasets are summarized in Table 2, including the imbalance ratio IR defined as: $IR = m_- / m_+$.

^a<https://archive.ics.uci.edu/ml/datasets.html>

^b<https://sci2s.ugr.es/keel/datasets.php>

Table 2. Information about the studied public datasets sorted by imbalance ratio IR. The *target* column refers to the label chosen as the minority class (*i.e.* positive examples) in the dataset. The short name of each dataset is given first and will be used, for the sake of readability, in some graphs of this study. (*) The target for YEAST is *ME2 vs MIT, ME3, EXC, VAC, ERL*.

DATASETS	SIZE	DIM	TARGET	IR	DATASETS	SIZE	DIM	TARGET	IR
BAL - BALANCE	625	4	<i>L</i>	1.2	PAG - PAGEBLOCKS	5472	10	<i>2,3,4,5</i>	8.8
AUT - AUTOMPG	392	7	<i>2,3</i>	1.7	SAT - SATIMAGE	6435	36	<i>4</i>	9.3
ION - IONOSPHERE	351	34	<i>b</i>	1.8	YEA - YEAST-0-5-6-7-9vs4	528	8	(*)	9.35
PIM - PIMA	768	8	<i>positive</i>	1.87	LIB - LIBRAS	360	90	<i>1</i>	14
GLA - GLASS	214	9	<i>1</i>	2.1	Y17 - YEAST-1vs7	459	7	<i>VAC vs NUC</i>	14.3
GER - GERMAN	1000	23	<i>2</i>	2.3	ARR - ARRHYTHMIA	452	278	<i>6</i>	17
YE1 - YEAST1	1484	8	<i>NUC</i>	2.46	SOL - SOLAR-FLARE-M0	1389	32	<i>M0</i>	19
HAB - HABERMAN	306	3	<i>positive</i>	2.78	OIL - OIL	937	49	<i>minority</i>	22
VE3 - VEHICLE3	846	18	<i>Class 3 Opel</i>	2.99	YE4 - YEAST4	1484	8	<i>ME2</i>	28.1
HAY - HAYES	132	4	<i>3</i>	3.4	WI4 - REDWINEQUALITY4	1599	11	<i>4</i>	29.2
SEG - SEGMENTATION	2310	19	<i>WINDOW</i>	6	YE5 - YEAST5	1484	8	<i>ME1</i>	32.73
AB8 - ABALONES	4177	10	<i>8</i>	6.4	YE6 - YEAST6	1484	8	<i>EXC</i>	41.4
YE3 - YEAST3	1484	8	<i>ME3</i>	8.1	A17 - ABALONE17	4177	10	<i>17</i>	71
EC3 - ECOLI3	336	7	<i>imU</i>	8.6	A20 - ABALONE20	4177	10	<i>20</i>	159.7

All the datasets are normalized using a min-max normalization such that each feature lies in the range $[-1, 1]$. We randomly draw 80%-20% splits of the data to generate the training and test sets respectively. Hyperparameters are tuned with a 10-fold cross-validation over the training set. We repeat the process over 5 runs and average the results in terms of the four performance measures. In a first series of experiments, we compare our method γk -NN to 6 other distance-based baselines:

- the classic k -Nearest Neighbor algorithm (k -NN),
- the weighted version of k -NN using the inverse distance as a weight to predict the label (wk -NN) [27],
- the class weighted version of k -NN (cwk -NN) [31],
- the k -NN version where each positive is duplicated according to the IR of the dataset ($dupk$ -NN),
- $kRNN$ where the sparsity of minority examples is taken into account [30] by modifying the way the posterior probability of belonging to the positive class is computed.
- the metric learning method LMNN [18].

The hyperparameter μ of $LMNN$, weighting the impact of impostor constraints (see [18] for more details), is tuned in the range $[0, 1]$ using a step of 0.1. Our γ parameter is tuned in the range $[0, 1]^c$ using a step of 0.1. For $kRNN$, we use the parameters values as described in [30].

In a second series of experiments, we compare our method to five oversampling strategies described in Section 3.2: SMOTE, Borderline-SMOTE, ADASYN, SMOTE with ENN, SMOTE with Tomek's link. The number of generated positive

^cWe experimentally noticed that using a larger range for γ leads in fact to a potential decrease of performances due to overfitting phenomena. This behavior is actually in line with the analysis provided in Section 4.2.

examples is tuned over the set of ratios $\frac{m_+}{m_-} \in \{0.1, 0.2, \dots, 0.9, 1.0\}$ and such that the new ratio is greater than the original one before sampling. The other parameters of these methods are the default ones used by the package *ImbalancedLearn* of *Scikit-learn*. We report the performance of the best oversampler that we denote as OS*. In order to evaluate how both strategies are complementary, we also combine γk -NN with oversamplers, and use the notation (OS+ γk -NN)* to indicate the best combination obtained by a 10-cross validation. In this latter scenario, we propose to learn a different γ value to be used with the synthetic positives. Indeed, some of the synthetic examples may be generated in some true negative areas and, in this situation, it might be more appropriate to decrease their influence. The γ parameter for these examples is tuned in the range $[0, 2]$ using a step of 0.1. Note the upper bound of the range is now set to 2. This allows γk -NN to adapt to the different sampling strategies of the oversamplers and enables the possibility to move synthetic positive examples away from dense regions of negatives by selecting $\gamma > 1$.

5.2. Analysis of the results

The results on the public datasets using the six baselines are provided in Tables 4,5,6 and 7 for the four different performance measures F_1 , BA , GM and G_1 respectively. These tables report the complete results when $k = 1$ (in k -NN) and provide only the mean results over the 28 datasets for $k = 3$, for the sake of concision and because the behavior for this latter value is similar. Overall, our γk -NN approach performs much better than its competitors by achieving an improvement of 0.7 to 5 points on average, compared to the other state-of-the-art algorithms when $k = 1$. It is worth noticing that the results are even better when $k = 3$. But the certainly most striking result comes from the capacity of γk -NN associated with the Balanced Accuracy (BA) in Table 5 and G-mean (GM) in Table 6 to address large imbalanced learning tasks. While the other methods struggle to get good results, γk -NN with BA and GM gets the best performance 19 and 20 times respectively over the 22 largest imbalanced datasets (from YEAST1 to ABALONE20). Even the metric learning algorithm LMNN fails to be competitive while it optimizes a representation of the data specifically dedicated to deal with nearest neighbor classification. Indeed, LMNN suffers from the lack of positive data to learn an efficient projection when dealing with highly imbalanced tasks. On the other hand, γk -NN does not seem particularly sensitive to the imbalance ratio.

The second series of experiments focuses on the use of sampling strategies and the potential interest of combining γk -NN with a synthetic generation of additional positive examples. Fig. 4 compactly summarizes, for the four measures of interest and for both $k = 1$ (on the left) and $k = 3$ (on the right), the impact of sampling strategies. Two main comments can be made from these results. First, γk -NN is complementary to the oversamplers. Indeed, for both $k = 1$ and $k = 3$ and for 3 out of 4 measures (G_1 excluded), using γk -NN in addition with a sampler leads

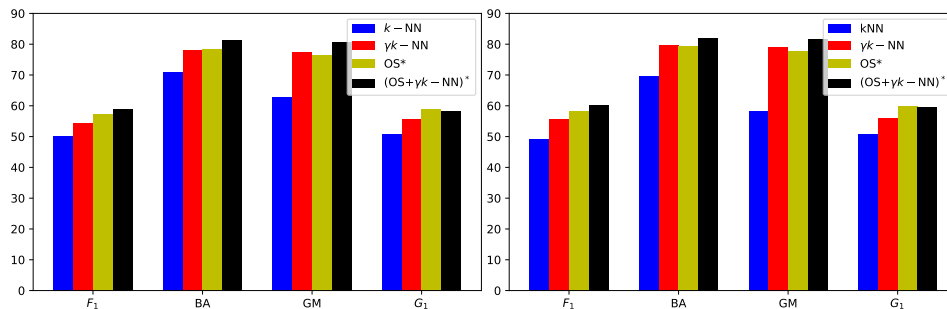


Fig. 4. Comparison of k -NN, γk -NN, the best oversampler among SMOTE, BorderSmote, SMOTE+ENN, SMOTE+Tomek’s links and ADASYN, and the best coupling oversampler + γk -NN, in terms of mean of F-measure (F_1), Balanced Accuracy (BA), Geometric Mean (GM) and G-measure (G_1) over all the datasets, for $k = 1$ (left) and $k = 3$ (right).

to better results and gives evidence of the fact that γk -NN and oversamplers do not work the same way, focusing on different subparts of the feature space. While γk -NN aims at expanding the decision boundaries in favor of the positives in the neighborhood of the test query, oversamplers rather tend to fill in the empty parts of the space by generating synthetic positive examples. Second, γk -NN (red bars) alone is shown to be very competitive while benefiting from its simplicity. Indeed, we remind the reader that the performance of OS^* (resp. $(OS+\gamma k-NN)^*$) are obtained from the costly selection of the best oversampler (resp. γk -NN + oversampler) for each dataset. Therefore, the green and black bars in Fig. 4 give an optimistic usage of an oversampling strategy because it is generated from the average obtained over a large set of oversamplers (SMOTE, Borderline-SMOTE, ADASYN, SMOTE+ ENN and SMOTE with Tomek’s link) that can be seen as an additional hyperparameter. On the other hand, in γk -NN, only one parameter (γ) is required to be tuned.

Fig. 5 illustrates, for the F_1 and GM measures and $k = 1$, a dataset-wise view of the advantage of combining γk -NN with an oversampler compared to a standard k -NN. A point (representing one of the 28 datasets) below the line $y = x$ means that k -NN is outperformed. Moreover, a move of a point (illustrated by a right arrow) from left to right illustrates that the joint approach leads to better results. We can see that even for the least favorable measure (*i.e.* F_1 on the left), most of the datasets are below the line and benefit from γk -NN associated to an oversampler.

In Fig 6 (left), we illustrate how having two γ parameters (γ on reals and γ on synthetics) gives the flexibility to independently control the influence of the actual and artificial positives respectively. The other figures (center and right) represent two examples of heatmaps of the F-measure (note that the trend is the same for the other 3 measures). We can note that while the γ parameter tuned for the real positives tends to be smaller than 1 (according to the analysis of Section 4.2), the γ parameter required to deal with the synthetic positives is sometimes smaller (right), sometimes greater than 1 (center), depending on the underlying density and the pe-

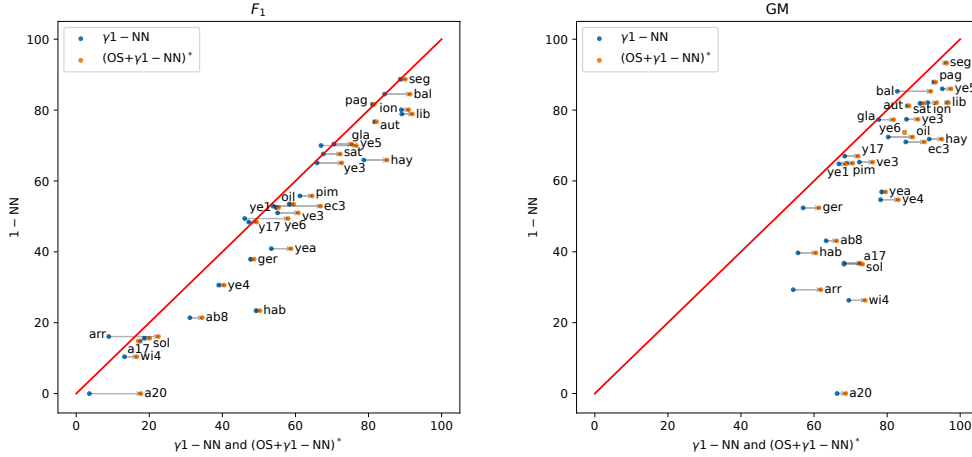
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Fig. 5. Comparison of k -NN with (i) γk -NN (points in blue) and (ii) γk -NN coupled with the best sampling strategy (points in orange) for each dataset, in terms of F-measure (left) and Geometric Mean (right) and for $k = 1$. Points below the line $y = x$ means that k -NN is outperformed. A move from left to right illustrates that the joint approach is better.

Table 3. Comparison of γk -NN and k -NN on MNIST for $k = 3$.

	F_1	BA	GM	G_1
k -NN	97.18	98.31	98.30	97.19
γk -NN	97.21	98.97	98.97	97.21

culiarity of the feature space.

Recall that Propositions 1 and 2 in Section 4.2 tell us that selecting a γ parameter smaller than 1 for the real positives should tend to reduce the false negative (FN) rate while still optimizing the performance measure. To illustrate our theoretical study, we plot in Fig. 7 the percentage of FN generated by the 7 compared methods. As expected, we can note that whatever the performance measure and the value of k ($k = 1$ on the left and $k = 3$ on the right), the number of FN is much smaller than that of the competitors explaining why γk -NN gets the best results.

5.3. A qualitative analysis on the MNIST dataset

In order to visualize the qualitative impact of γk -NN, we conduct in this section some additional experiments on the MNIST dataset. To generate a minority class, we build 10 datasets $MNIST_i$ (one for each digit, $i = 0, \dots, 9$) from the original one by considering the label i as the minority class and all the other classes representing the remaining digits as the majority class.

As previously done, a 10-fold CV is performed to find the optimal value γ . The mean results of the comparison of γk -NN with k -NN are reported in Table 3

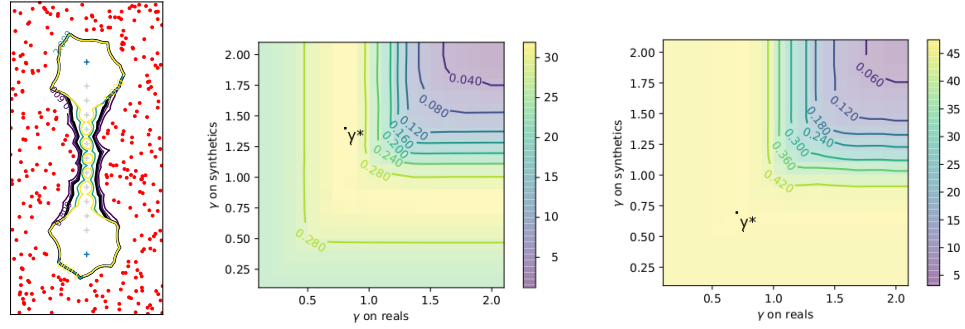


Fig. 6. Left: Illustration on a toy dataset of the effect of varying γ for the generated positive points (in grey) while keeping a fixed $\gamma = 0.4$ for the real positives. Center and Right: Two examples of heatmap for the F-Measure that show the pair of γ (on real and synthetic positives) corresponding to the best joint approach $(OS + \gamma k\text{-NN})^*$ on ABALONES (center) and GERMAN (right).

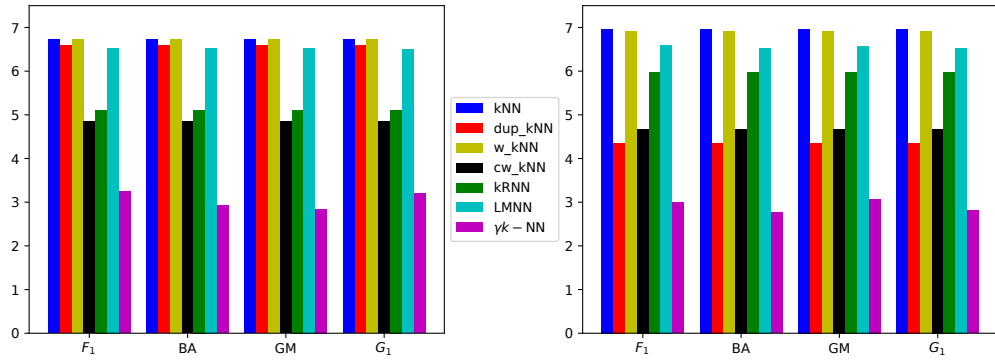


Fig. 7. Percentage of false negatives (FN) generated by the 7 compared methods w.r.t. the four performance measures with 1-NN (left) and 3-NN (right).

where $k = 3$. We can see that whatever the performance measure, $\gamma k\text{-NN}$ allows us to outperform $k\text{-NN}$. As expected, the gain on this well-known MNIST dataset is not significant due to the already very high accuracy reached by the standard $k\text{-NN}$. However, the main objective here is elsewhere. We aim at showing the quality of both the space and the neighborhoods induced by $\gamma k\text{-NN}$. To illustrate this purpose and visualize how using d_γ (as defined in Eq.(1)) bends the feature space, we leverage t-SNE to embed the $MNIST_i$ points in 2D. Note that even if d_γ is not an actual distance (the symmetry is not satisfied), it can still be used with t-SNE that only embeds points while preserving relative pair-wise dissimilarities.

Following the definition of d_γ , we scale the Euclidean distance when the second point in the pair is a positive one. Fig. 8 compares, on the $MNIST_2$ dataset, the output of t-SNE when using d (left) and d_γ (right). The analysis of this embedding shows that d_γ is able to gather minority examples together in a denser cluster while the Euclidean distance leads to a space where the positive samples are more

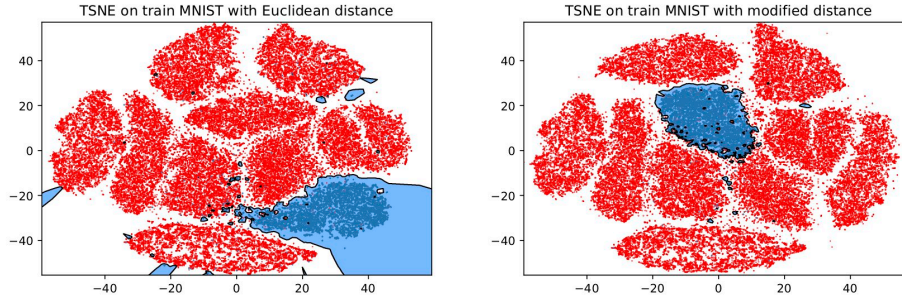
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Fig. 8. Visualization on $MNIST_2$ of the influence of the Euclidean distance d (left) and d_γ (right) with t-SNE. The red (resp. blue) points correspond to negatives (resp. positives). The blue areas represent the subparts of the space leading to a positive classification by a 3-NN.

k -NN			True	γk -NN		

Fig. 9. First three columns: 3 nearest neighbors using k -NN; fourth column: the test query; last three figures: 3 nearest neighbors using γk -NN.

scattered, some being in the middle of negative regions. This impact of γk -NN on the decision boundaries, that we see with this t-SNE experiment, is also illustrated in Fig. 9 which shows some examples for which, the γk -NN predictions are different from that of k -NN according to their 3-nearest neighbors (on the original dataset).

5.4. On local- γk -NN using clustering

We now evaluate our local algorithm local- γk -NN (as presented in Section 4.4), which partitions the input space into q clusters (C_1, C_2, \dots, C_q) and uses a parameter γ_j for each cluster j . The partitioning is performed using k-means, run on the training set. Note that we consider two ways of obtaining the γ_j values. The first version (V1) consists in applying the 10 fold cross-validation (CV) procedure in each cluster C_j to tune γ_j . At test time, a new point \mathbf{x}' is first assigned to the

nearest cluster C_k based on the closest centroid using the Euclidean distance, and the corresponding γ_k value is used to scale the distances to the positives.

We propose a second version (V2) to compensate for the fact that V1 is at risk of generating very different values of γ for two neighboring clusters. While the test decision is similar to V1, the γ_j values are obtained differently, by computing several clusterings. In V2, the 10 fold CV also includes the clustering, so 10 additional partitionings are performed. Each training point \mathbf{x} will thus fall in 9 clusters (in the 9 different clusterings for which the point is not in the validation fold). Each point thus has 9 “best” γ values that we average to get a single value $\gamma_{\mathbf{x}}$ for every single point. In the end, γ_j is computed as $\frac{1}{|C_j|} \sum_{\mathbf{x} \in C_j} \gamma_{\mathbf{x}}$, *i.e.* the average γ value of the training points falling into cluster j .

The results are provided for the 4 performance measures in Fig. 10, 11, 12 and 13. Despite an inherent increase of the time complexity, it is worth noting that V2 is better than the original γk -NN (on average over the 28 datasets), while V1 does not lead to improvements probably due to overfitting phenomena. Note also that in a huge majority of the datasets (around 90%), the V2 version of local- γk -NN equals or outperforms γk -NN.

6. Conclusion

In this paper, we have proposed a new approach, γk -NN, that addresses the problem of learning from imbalanced datasets. It is based on the k -NN algorithm but it modifies the distance to the positive examples by expanding the decision boundaries around these minority samples. It has been shown to outperform its competitors in terms of several performance measures. Furthermore, we gave evidence of the complementarity of γk -NN with oversampling strategies. Our algorithm, despite its simplicity, is highly effective and its local version local- γk -NN has shown to be even more efficient by taking the spatial specificity of the distributions into account.

Two main lines of research deserve future investigations. First, we plan to extend the idea of the local variant of γk -NN by proposing a multi-view learning approach, where the different results of γk -NN obtained with different subsets of features (the different views) would be combined in some way. Second, we can note that tuning γ is equivalent to building a diagonal matrix (with γ^2 in the diagonal) and applying a Mahalanobis distance only between a query and a positive example. This comment opens the door to a new family of metric learning algorithms dedicated to optimizing a PSD matrix under (FP, FN) -based constraints that could leverage recent metric learning approaches for imbalanced data [39].

Acknowledgments

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 Table 4. Results for $k = 1$ with F_1 as performance measure over 5 runs. The standard deviation is indicated after the \pm sign and the best results on each dataset is indicated in bold. Only the mean value when $k = 3$ is shown in the last line.

DATASETS	k -NN	DUP k -NN	w k -NN	cw k -NN	κ RNN	LMNN	γk -NN
BALANCE	84.5 \pm 2.2	84.5 \pm 2.2	84.5 \pm 2.2	84.4 \pm 1.7	88.2\pm1.2	84.1 \pm 4.6	84.4 \pm 1.7
AUTOMPG	76.7 \pm 7.4	76.7 \pm 7.4	76.7 \pm 7.4	76.2 \pm 6.2	82.5\pm2.8	77.5 \pm 3.1	81.8 \pm 3.0
IONOSPHERE	80.1 \pm 4.2	80.1 \pm 4.2	80.1 \pm 4.2	83.3 \pm 3.0	0.83 \pm 3.0	81.3 \pm 3.3	89.0\pm4.5
PIMA	55.8 \pm 4.7	55.8 \pm 4.7	55.8 \pm 4.7	61.1 \pm 3.7	62.3\pm3.9	55.9 \pm 2.8	61.2 \pm 5.4
GLASS	70.4 \pm 8.7	70.4 \pm 8.7	70.4 \pm 8.7	73.2 \pm 6.4	76.2\pm8.6	68.9 \pm 7.7	70.5 \pm 7.5
GERMAN	37.9 \pm 5.0	37.9 \pm 5.0	37.9 \pm 5.0	41.1 \pm 3.6	43.7 \pm 4.0	41.0 \pm 3.8	47.7\pm1.9
YEAST1	52.5 \pm 2.6	52.5 \pm 2.6	52.5 \pm 2.6	53.3 \pm 3.6	52.5 \pm 2.1	51.3 \pm 3.7	54.8\pm3.8
HABERMAN	23.4 \pm 6.7	23.4 \pm 6.7	23.4 \pm 6.7	35.5 \pm 10	33.2 \pm 7.6	24.6 \pm 6.9	49.2\pm4.4
VEHICLE3	51.0 \pm 3.4	51.0 \pm 3.4	51.0 \pm 3.4	51.2 \pm 3.7	56.1\pm3.3	54.7 \pm 3.6	55.1 \pm 3.4
HAYES	65.9 \pm 10	65.9 \pm 10	65.9 \pm 10	87.5\pm4.9	76.6 \pm 8.3	76.8 \pm 14	78.7 \pm 10
SEGMENTATION	88.7 \pm 2.9	88.7 \pm 2.9	88.7 \pm 2.9	88.9 \pm 2.8	87.2 \pm 1.6	91.2\pm2.3	88.7 \pm 2.9
ABALONE8	21.4 \pm 1.3	21.4 \pm 1.3	21.4 \pm 1.3	31.3\pm0.9	23.8 \pm 1.1	21.9 \pm 1.7	31.1 \pm 2.0
YEAST3	65.1 \pm 2.5	65.1 \pm 2.5	65.1 \pm 2.5	63.0 \pm 2.6	69.4\pm1.2	63.6 \pm 1.0	65.9 \pm 2.3
ECOLI3	52.9 \pm 9.7	52.9 \pm 9.7	52.9 \pm 9.7	54.1 \pm 7.8	61.2\pm5.8	57.2 \pm 11	53.9 \pm 7.0
PAGEBLOCKS	81.6\pm2.6	81.6\pm2.6	81.6 \pm 2.6	81.1 \pm 2.4	81.3 \pm 4.4	81.5 \pm 3.2	81.2 \pm 2.2
SATIMAGE	67.6 \pm 3.6	67.6 \pm 3.6	67.6 \pm 3.6	68.0 \pm 3.4	68.8 \pm 2.7	69.0\pm4.5	67.6 \pm 3.6
YEAST-0.5-6-7-9vs4	40.9 \pm 11	40.9 \pm 11	40.9 \pm 11	49.7 \pm 4.1	51.9 \pm 7.3	45.5 \pm 15	53.4\pm8.3
LIBRAS	78.9 \pm 8.7	78.9 \pm 8.7	78.9 \pm 8.7	78.9 \pm 8.7	73.7 \pm 6.0	78.8 \pm 5.4	89.1\pm8.1
YEAST-1VS7	48.4 \pm 6.0	48.4 \pm 6.0	48.4 \pm 6.0	23.8 \pm 5.3	49.1\pm8.8	40.4 \pm 16.6	47.2 \pm 5.0
ARRYTHMIA	16.1 \pm 20	16.1 \pm 20	16.1 \pm 20	15.6 \pm 20	15.6 \pm 20	20.2\pm21	8.9 \pm 12
SOLAR-FLARE-M0	15.6 \pm 5.6	15.6 \pm 5.6	15.6 \pm 5.6	18.4 \pm 1.9	21.4\pm6.3	15.3 \pm 10	18.6 \pm 2.1
OIL	53.5 \pm 11.2	53.5 \pm 11	53.5 \pm 11	57.2 \pm 9.7	55.5 \pm 5.2	61.3\pm12	58.3 \pm 12
YEAST4	30.6 \pm 11.3	30.6 \pm 11	30.6 \pm 11	29.2 \pm 1.9	41.4\pm4.0	32.4 \pm 12	39.0 \pm 8.4
REDWINEQUALITY4	10.4 \pm 5.7	10.4 \pm 5.7	10.4 \pm 5.7	12.4 \pm 2.6	12.9 \pm 7.0	12.0 \pm 6.9	13.2\pm5.6
YEAST5	70.0 \pm 11	70.0 \pm 11	70.0 \pm 11	56.4 \pm 8.2	62.6 \pm 8.3	70.1\pm12	67.0 \pm 9.8
YEAST6	49.4 \pm 13	49.4 \pm 13	49.4 \pm 13	26.2 \pm 2.3	49.9\pm8.6	47.1 \pm 19	46.1 \pm 10
ABALONE17	14.8 \pm 9.7	14.8 \pm 9.7	14.8 \pm 9.7	10.5 \pm 4.0	16.3 \pm 6.9	14.3 \pm 6.7	17.3\pm9.4
ABALONE20	00.0 \pm 0.0	00.0 \pm 0.0	00.0 \pm 0.0	05.2 \pm 3.5	06.6\pm6.7	00.0 \pm 0.0	03.6 \pm 4.7
MEAN ($\kappa=1$)	50.1	50.1	50.1	50.6	53.7	51.4	54.4
MEAN ($\kappa=3$)	49.3	54.0	50.0	52.2	53.2	51.9	55.8

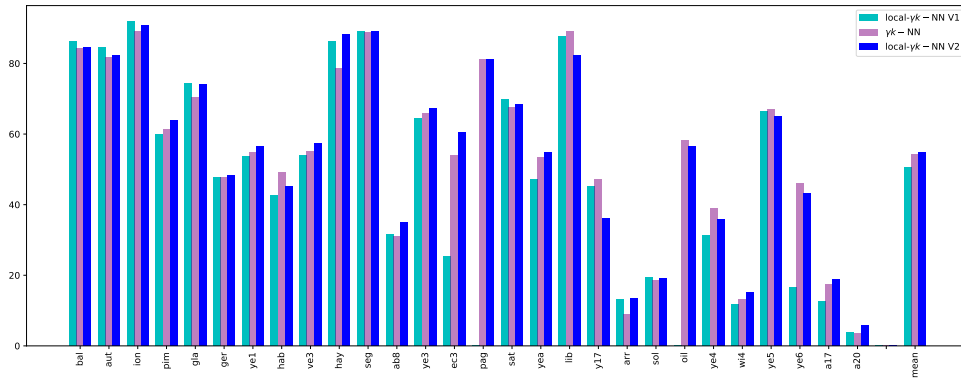

 Fig. 10. Comparison of $\gamma 1$ -NN with the two versions of local- $\gamma 1$ -NN, in terms of F-measure.

Table 5. Results for $k = 1$ with BA as performance measure over 5 runs. The standard deviation is indicated after the \pm sign and the best results on each dataset is indicated in bold. Only the mean value when $k = 3$ is shown in the last line.

DATASETS	k -NN	DUP k -NN	w k -NN	cw k -NN	κ RNN	LMNN	γk -NN
BALANCE	85.4±2.1	85.4±2.1	85.4±2.1	84.2±1.9	88.9±1.2	85.5±4.1	84.2±1.9
AUTOMPG	81.6±6.1	81.6±6.1	81.6±6.1	81.1±5.2	86.4±2.6	81.9±2.6	86.0±2.9
IONOSPHERE	83.6±3.0	83.6±3.0	83.6±3.0	85.9±2.1	85.6±2.1	84.6±2.4	91.2±3.6
PIMA	66.6±3.4	66.6±3.4	66.6±3.4	69.4±3.3	70.4±3.4	66.7±1.7	69.7±3.7
GLASS	78.2±6.5	78.2±6.5	78.2±6.5	80.3±4.8	83.3±6.7	77.1±6.4	76.5±4.3
GERMAN	57.1±3.0	57.1±3.0	57.1±3.0	58.0±2.4	59.0±3.1	58.6±2.6	57.7±3.0
YEAST1	66.6±1.9	66.6±1.9	66.6±1.9	66.5±3.2	66.2±1.7	65.8±2.6	67.2±3.8
HABERMAN	49.8±5.4	49.8±5.4	49.8±5.4	52.7±9.0	53.3±6.2	50.4±5.4	61.1±4.7
VEHICLE3	67.3±2.3	67.3±2.3	67.3±2.3	67.4±2.8	71.2±2.7	70.1±3.2	72.6±1.1
HAYES	75.7±6.0	75.7±6.0	75.7±6.0	90.7±4.6	82.4±5.4	82.9±8.7	91.9±4.2
SEGMENTATION	93.5±2.2	93.5±2.2	93.5±2.2	95.1±2.1	95.5±0.9	94.9±1.5	96.2±0.9
ABALONES	54.5±0.7	54.5±0.7	54.5±0.7	61.3±0.8	55.7±0.7	54.9±0.9	62.6±1.8
YEAST3	79.4±2.9	79.4±2.9	79.4±2.9	85.5±2.4	83.4±2.8	80.4±2.5	85.7±2.9
ECOLI3	74.4±6.9	74.4±6.9	74.4±6.9	82.3±5.9	81.2±5.2	73.3±3.5	85.5±8.1
PAGEBLOCKS	88.5±1.5	88.5±1.5	88.5±1.5	91.4±2.1	90.4±2.3	88.7±1.9	92.9±1.3
SATIMAGE	83.0±2.0	83.0±2.0	83.0±2.0	87.5±1.7	86.6±1.6	83.8±1.8	89.1±1.2
YEAST-0.5-6-7-9vs4	65.4±5.2	65.4±5.2	65.4±5.2	78.3±3.7	71.6±3.3	68.1±7.4	79.3±3.6
LIBRAS	83.9±5.0	83.9±5.0	83.9±5.0	83.9±5.0	83.4±4.8	83.9±4.7	96.7±3.4
YEAST-1vs7	71.7±3.2	71.7±3.2	71.7±3.2	67.5±7.7	73.1±5.3	68.1±1.0	72.5±7.0
ARRHYTHMIA	57.5±16	57.5±16	57.5±16	56.8±17	57.2±16	59.7±16	54.7±6.5
SOLAR-FLARE-M0	55.1±2.5	55.1±2.5	55.1±2.5	65.1±1.8	58.2±2.9	55.0±4.3	67.3±4.2
OIL	76.6±8.8	76.6±8.8	76.6±8.8	80.7±6.2	83.4±4.0	79.3±9.7	84.6±4.4
YEAST4	64.9±8.3	64.9±8.3	64.9±8.3	78.7±2.1	77.5±3.5	66.6±8.5	79.3±3.5
REDWINEQUALITY4	53.5±2.7	53.5±2.7	53.5±2.7	58.7±3.9	55.7±4.4	54.3±3.2	69.2±5.8
YEAST5	87.2±7.4	87.2±7.4	87.2±7.4	91.4±5.5	90.9±5.7	86.2±6.7	95.1±2.8
YEAST6	77.7±10	77.7±10	77.7±10	84.9±9.3	85.7±9.3	79.0±14	79.2±7.0
ABALONE17	56.8±4.9	56.8±4.9	56.8±4.9	64.2±7.3	63.2±6.8	57.7±3.7	67.0±4.2
ABALONE20	49.7±0.1	49.7±0.1	49.7±0.1	58.8±6.9	55.5±6.4	49.7±0.1	68.8±11
MEAN ($k=1$)	70.9	70.9	70.9	75.3	74.8	71.7	78.0
MEAN ($k=3$)	69.6	75.4	69.9	75.5	74.2	71.7	79.7

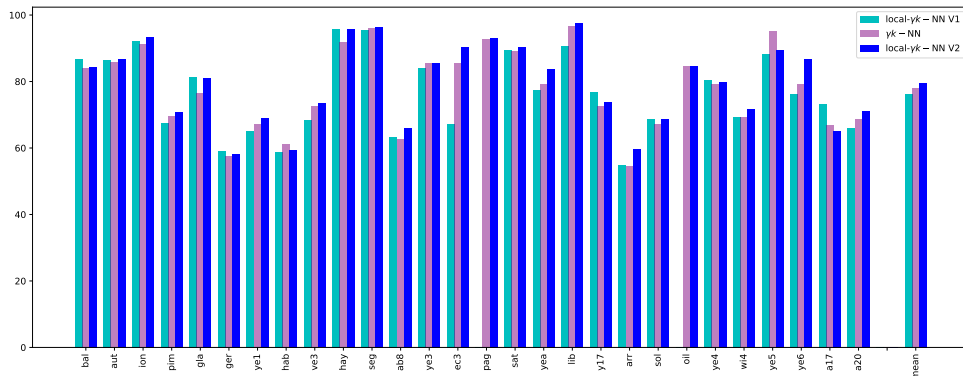


Fig. 11. Comparison of $\gamma 1$ -NN with the two versions of local- $\gamma 1$ -NN, in terms of Balanced Accuracy (BA).

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 Table 6. Results for $k = 1$ with GM as performance measure over 5 runs. The standard deviation is indicated after the \pm sign and the best results on each dataset is indicated in bold. Only the mean value when $k = 3$ is shown in the last line.

DATASETS	k -NN	DUP k -NN	w k -NN	cw k -NN	kRNN	LMNN	γk -NN
BALANCE	85.3±2.1	85.3±2.1	85.3±2.1	82.8±2.2	88.8±1.2	85.4±4.2	82.8±2.2
AUTOPMG	81.2±6.4	81.2±6.4	81.2±6.4	80.7±5.4	86.2±2.5	81.6±2.6	85.7±3.1
IONOSPHERE	82.1±3.5	82.1±3.5	82.1±3.5	84.9±2.3	84.7±2.3	84.1±2.2	91.1±3.7
PIMA	65.0±3.8	65.0±3.8	65.0±3.8	69.2±3.2	70.3±3.4	65.1±2.3	68.9±3.9
GLASS	77.3±7.2	77.3±7.2	77.3±7.2	79.7±5.3	82.6±7.2	76.1±6.8	77.7±6.0
GERMAN	52.4±4.2	52.4±4.2	52.4±4.2	55.4±3.0	57.6±3.4	55.1±3.2	57.0±3.0
YEAST1	64.8±1.8	64.8±1.8	64.8±1.8	66.3±3.1	65.6±1.7	63.8±2.7	66.8±4.2
HABERMAN	39.7±6.4	39.7±6.4	39.7±6.4	51.4±9.8	49.6±6.8	40.9±5.9	55.6±7.9
VEHICLE3	65.3±3.1	65.3±3.1	65.3±3.1	66.6±3.1	70.5±3.2	68.4±4.1	72.4±1.6
HAYES	71.8±8.6	71.8±8.6	71.8±8.6	90.2±5.1	80.6±6.6	80.3±11.9	91.5±4.6
SEGMENTATION	93.3±2.3	93.3±2.3	93.3±2.3	95.0±2.2	95.5±0.9	94.9±1.5	96.2±0.9
ABALONES	43.1±1.6	43.1±1.6	43.1±1.6	59.6±0.9	47.2±1.6	43.6±2.1	63.3±2.2
YEAST3	77.4±3.9	77.4±3.9	77.4±3.9	85.2±2.9	82.3±3.6	78.9±3.4	85.3±3.5
ECOLI3	71.0±9.0	71.0±9.0	71.0±9.0	81.9±6.5	79.9±6.1	73.2±9.7	85.1±8.5
PAGEBLOCKS	87.9±1.6	87.9±1.6	87.9±1.6	91.2±2.2	90.1±2.5	88.1±2.0	92.9±1.3
SATIMAGE	81.9±2.3	81.9±2.3	81.9±2.3	87.3±1.8	86.2±1.7	83.2±2.7	89.0±1.2
YEAST-0.5-6-7-9vs4	56.9±9.2	56.9±9.2	56.9±9.2	77.5±4.4	67.1±4.7	58.9±14.2	78.5±4.2
LIBRAS	82.1±6.0	82.1±6.0	82.1±6.0	82.1±6.0	81.8±5.7	82.0±5.8	96.6±3.6
YEAST-1vs7	67.0±5.0	67.0±5.0	67.0±5.0	65.8±9.4	69.0±7.5	59.8±16.5	68.4±10.5
ARRHYTHMIA	29.3±36.7	29.3±36.7	29.3±36.7	29.2±36.6	29.2±36.6	37.9±34.1	54.3±8.4
SOLAR-FLARE-M0	36.5±6.1	36.5±6.1	36.5±6.1	63.9±1.9	43.5±5.9	36.3±8.6	68.2±4.3
OIL	72.4±12.4	72.4±12.4	72.4±12.4	78.6±7.9	82.3±5.0	75.8±12.7	80.3±2.2
YEAST4	54.7±13.5	54.7±13.5	54.7±13.5	78.1±2.4	75.3±4.4	57.6±14.1	78.2±4.3
REDWINEQUALITY4	26.3±14.0	26.3±14.0	26.3±14.0	50.1±7.1	35.1±17.9	28.8±15.4	69.5±6.6
YEAST5	86.0±8.4	86.0±8.4	86.0±8.4	91.1±6.0	90.4±6.2	85.0±7.3	95.1±2.8
YEAST6	73.7±13.7	73.7±13.7	73.7±13.7	83.7±11.1	84.1±11.2	73.7±21.1	84.8±6.2
ABALONE17	36.8±11.3	36.8±11.3	36.8±11.3	56.9±11.2	52.4±12.1	39.4±9.4	68.2±6.4
ABALONE20	00.0±0.0	00.0±0.0	00.0±0.0	45.4±11.8	27.6±23.4	00.0±0.0	66.3±12.6
MEAN (K=1)	62.9	62.9	62.9	72.5	69.8	64.2	77.5
MEAN (K=3)	58.4	71.6	59.1	70.71	67.6	61.7	78.9

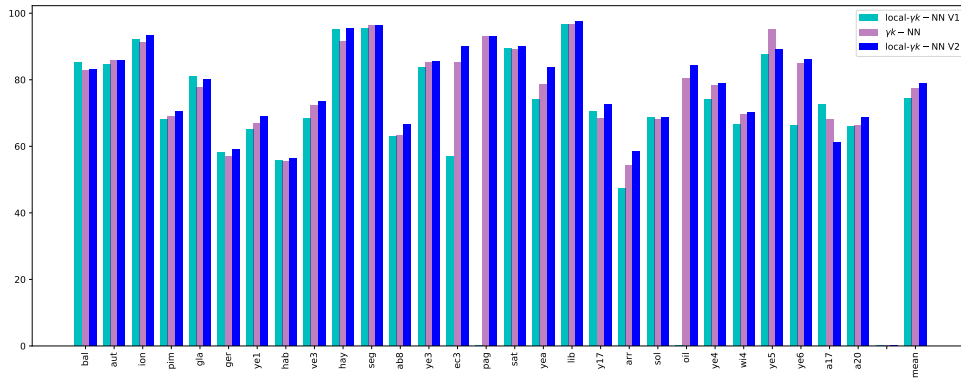
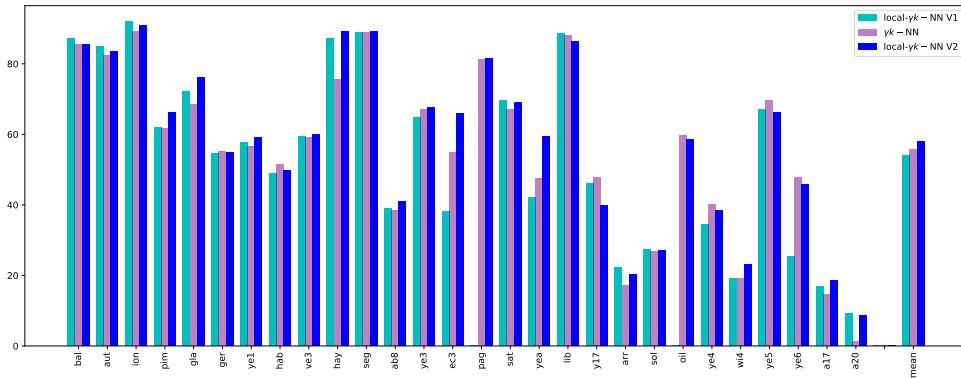

 Fig. 12. Comparison of $\gamma 1$ -NN with the two versions of local- $\gamma 1$ -NN, in terms of Geometric Mean (GM).

Table 7. Results for $k = 1$ with G_1 as performance measure over 5 runs. The standard deviation is indicated after the \pm sign and the best results on each dataset is indicated in bold. Only the mean value when $k = 3$ is shown in the last line.

DATASETS	k -NN	DUP k -NN	w k -NN	cw k -NN	κ RNN	LMNN	γk -NN
BALANCE	84.5 \pm 2.2	84.5 \pm 2.2	84.5 \pm 2.2	85.4 \pm 1.5	88.4\pm1.2	84.2 \pm 4.6	85.4 \pm 1.5
AUTOPMG	76.9 \pm 7.4	76.9 \pm 7.4	76.9 \pm 7.4	76.4 \pm 6.3	82.9\pm2.9	77.7 \pm 3.1	82.3 \pm 3.2
IONOSPHERE	81.4 \pm 3.9	81.4 \pm 3.9	81.4 \pm 3.9	84.2 \pm 3.0	83.8 \pm 2.9	82.0 \pm 3.1	89.2\pm4.3
PIMA	56.1 \pm 4.6	56.1 \pm 4.6	56.1 \pm 4.6	61.3 \pm 3.6	62.6\pm3.8	56.1 \pm 2.6	61.8 \pm 4.9
GLASS	71.0 \pm 8.2	71.0 \pm 8.2	71.0 \pm 8.2	73.8 \pm 5.9	77.4\pm7.9	69.6 \pm 7.6	68.5 \pm 5.0
GERMAN	38.1 \pm 5.0	38.1 \pm 5.0	38.1 \pm 5.0	41.1 \pm 3.6	43.8 \pm 4.0	41.1 \pm 3.8	55.1\pm1.1
YEAST1	52.6 \pm 2.6	52.6 \pm 2.6	52.6 \pm 2.6	53.8 \pm 3.5	52.9 \pm 2.0	51.4 \pm 3.6	56.7\pm2.4
HABERMAN	23.8 \pm 7.0	23.8 \pm 7.0	23.8 \pm 7.0	36.0 \pm 10.3	33.3 \pm 7.6	25.1 \pm 7.4	51.4\pm4.9
VEHICLE3	51.1 \pm 3.4	51.1 \pm 3.4	51.1 \pm 3.4	51.5 \pm 3.7	56.5 \pm 3.6	54.8 \pm 3.7	59.1\pm1.2
HAYES	68.9 \pm 8.6	68.9 \pm 8.6	68.9 \pm 8.6	88.0\pm4.6	77.9 \pm 7.6	79.1 \pm 11.3	75.6 \pm 10.2
SEGMENTATION	88.8 \pm 2.9	88.8 \pm 2.9	88.8 \pm 2.9	89.0 \pm 2.8	87.5 \pm 1.5	90.9\pm2.9	88.8 \pm 2.9
ABALONES	21.4 \pm 1.3	21.4 \pm 1.3	21.4 \pm 1.3	33.1 \pm 0.9	23.9 \pm 1.2	21.9 \pm 1.7	38.5\pm1.2
YEAST3	65.4 \pm 2.3	65.4 \pm 2.3	65.4 \pm 2.3	64.9 \pm 1.8	69.7\pm1.0	63.5 \pm 1.2	67.0 \pm 2.6
ECOLI3	53.0 \pm 9.8	53.0 \pm 9.8	53.0 \pm 9.8	56.7 \pm 8.0	61.7\pm6.2	54.2 \pm 10.0	54.9 \pm 10.7
PAGEBLOCKS	81.7\pm2.6	81.7\pm2.6	81.7\pm2.6	81.3 \pm 2.5	81.3 \pm 4.4	81.5 \pm 3.2	81.2 \pm 2.2
SATIMAGE	67.7 \pm 3.6	67.7 \pm 3.6	67.7 \pm 3.6	69.0 \pm 3.2	69.3\pm2.7	69.0 \pm 4.5	67.2 \pm 4.0
YEAST-0-5-6-7-9vs4	42.2 \pm 10.4	42.2 \pm 10.4	42.2 \pm 10.4	51.9 \pm 4.3	52.7\pm7.5	46.5 \pm 14.8	47.6 \pm 14.4
LIBRAS	80.2 \pm 8.5	80.2 \pm 8.5	80.2 \pm 8.5	80.2 \pm 8.5	74.3 \pm 6.0	75.9 \pm 4.5	88.0\pm8.3
YEAST-1vs7	48.7 \pm 5.8	48.7 \pm 5.8	48.7 \pm 5.8	29.2 \pm 7.3	49.3 \pm 8.8	50.0\pm12.5	47.9 \pm 7.1
ARRYTHMIA	17.1 \pm 21.7	17.1 \pm 21.7	17.1 \pm 21.7	16.7 \pm 21.4	16.7 \pm 21.4	20.5\pm22.0	17.1 \pm 21.7
SOLAR-FLARE-M0	16.5 \pm 6.8	16.5 \pm 6.8	16.5 \pm 6.8	24.3 \pm 1.7	21.5 \pm 6.4	11.7 \pm 7.9	26.7\pm3.2
OIL	54.6 \pm 10.8	54.6 \pm 10.8	54.6 \pm 10.8	58.0 \pm 9.4	57.1 \pm 4.8	66.6\pm10.0	59.7 \pm 10.1
YEAST4	31.2 \pm 11.6	31.2 \pm 11.6	31.2 \pm 11.6	35.7 \pm 2.1	43.6\pm4.1	35.0 \pm 13.2	40.2 \pm 8.7
REDWINEQUALITY4	10.8 \pm 5.8	10.8 \pm 5.8	10.8 \pm 5.8	15.2 \pm 3.7	13.3 \pm 7.1	12.2 \pm 7.0	19.2\pm5.5
YEAST5	70.5 \pm 11.0	70.5 \pm 11.0	70.5 \pm 11.0	60.2 \pm 8.1	64.9 \pm 8.1	72.0\pm7.4	69.7 \pm 9.6
YEAST6	50.0 \pm 13.9	50.0 \pm 13.9	50.0 \pm 13.9	35.3 \pm 5.3	52.9\pm9.9	46.6 \pm 18.1	47.9 \pm 11.5
ABALONE17	15.0 \pm 9.7	15.0 \pm 9.7	15.0 \pm 9.7	15.0 \pm 5.7	18.3\pm7.8	17.0 \pm 7.2	14.7 \pm 2.4
ABALONE20	00.0 \pm 0.0	00.0 \pm 0.0	00.0 \pm 0.0	8.3\pm5.2	7.6 \pm 7.5	00.0 \pm 0.0	1.2 \pm 2.4
MEAN (K=1)	50.7	50.7	50.7	52.7	54.5	52.0	55.8
MEAN (K=3)	50.8	55.2	51.4	54.0	54.6	52.7	55.9


 Fig. 13. Comparison of γ_1 -NN with the two versions of local- γ_1 -NN, in terms of G-measure (G_1).

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