Margin-aware Adversarial Domain Adaptation with Optimal Transport
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Abstract

In this paper, we propose a new theoretical analysis of unsupervised domain adaptation (DA) that relates notions of large margin separation, adversarial learning and optimal transport. This analysis generalizes previous work on the subject by providing a bound on the target margin violation rate, thus reflecting a better control of the quality of separation between classes in the target domain than bounding the misclassification rate. The bound also highlights the benefit of a large margin separation on the source domain for adaptation and introduces an optimal transport (OT) based distance between domains that has the virtue of being task-dependent, contrary to other approaches. From the obtained theoretical results, we derive a novel algorithmic solution for domain adaptation that introduces a novel shallow OT-based adversarial approach and outperforms other OT-based DA baselines on several simulated and real-world classification tasks.

1. Introduction

Learning to classify elements of a data set is one of the most widespread tasks in machine learning. Classically, it is done after assuming that data used for testing and those used for training a classification model stem from the same probability distribution (Sen et al., 2020). However, such an assumption is usually violated in real-world applications: product reviews classification where the difference of product types change the words distributions (Blitzer et al., 2007) or image classification where the variation of acquisition methods and conditions between training and test data introduce non-negligible distribution shifts (Hutchison et al., 2010), to name a few. While manual labeling may be considered as a tractable solution, such an approach is time consuming, can necessitate an often costly intervention of experts for labeling (e.g. in medical imaging applications) and totally discards the information available on a different, yet related, labeled training set. Such a setting has motivated the emergence of domain adaptation (Pan & Yang, 2010; Weiss et al., 2016), a branch of statistical learning that takes the distribution shift into account, and in which the test data is assumed to be partially or totally unlabeled. In the literature, the training set and test set distributions are respectively termed source and target domains. In this paper, we focus on the challenging setting of unsupervised domain adaptation (Margolis, 2011), were no labels are available for the target data.

Since the inception of the domain adaptation field, several theoretical contributions were proposed to analyze this problem in the statistical learning framework (Ben-David et al., 2007b; Mansour et al., 2009; Cortes & Mohri, 2014; Germain et al., 2016; Zhang et al., 2019). The general idea behind any such analysis usually consists in bounding the target domain error rate by a source error rate plus an estimable term reflecting a certain distance between domains, called the alignment or divergence term, plus a non estimable term that is assumed to be small for adaptation to be possible. To this end, the seminal work of (Ben-David et al., 2007b) considered the bounds for 0-1 loss in binary classification setting by introducing a divergence term that takes into account the complexity of the hypothesis space. Their results were further generalized for any loss function verifying the triangle inequality in (Mansour et al., 2009) and to a case when the hypothesis space is a RKHS in (Cortes & Mohri, 2014; Cortes et al., 2019). A somewhat different result was recently proposed (Zhang et al., 2019) where the authors provided generalization bounds for DA in the case of multi-class classification with source domain error defined by the margin violation rate. Finally, several DA bounds where proposed in (Redko et al., 2016; Courty et al., 2017; Shen et al., 2017) for the specific case when the considered alignment term is given by the Wasserstein distance (Santambrogio, 2016).

At the algorithmic level, there have been a plethora of algorithms that deal with the unsupervised domain adaptation problem, and they can be roughly divided to shallow (Kouw
introduces required preliminary knowledge and notations. Section 3 is dedicated to our theoretical contributions presenting a novel bound on the margin violation error on the target domain, its thorough analysis and relation to other existing bounds. Then, in Section 4, we use it to derive an algorithm that is further specialized to linear classifiers, resulting in a convex programming formulation. Finally, in the last section, we evaluate our algorithm on a toy data set and on a benchmark real-world problem.

2. Preliminary Knowledge

In this section, we present the problem setup of our study with the notations used throughout the paper. We also provide background knowledge on learning bounds in DA to allow a further comparison to them in the rest of the manuscript.

2.1. Problem setup and notations

We consider a binary classification setting, in which source and target data are respectively drawn from $S$ and $T$, the joint distributions over the product space of instances and labels $X \times Y$, where $X \in \mathbb{R}^d$ and $Y = \{-1, 1\}$. We denote their corresponding marginal distributions as $D_S$ and $D_T$ and use bold upper-case letters for matrices (e.g., $D$) and bold lower-case letters for vectors (e.g., $x$). Although both domains are assumed to be labeled, only the labels of the source instances are observable during the learning stage. This setting is often referred to as unsupervised domain adaptation.

To proceed, let $\mathcal{H}$ and $\mathcal{H}'$ denote two compact classes of hypotheses acting on $X$ and taking values in $[-1, 1]$. For further developments, we define several quantities to assess classifiers’ performances on different domains. Let $l^{\rho, \beta}$ be the loss function defined by

$$
l^{\rho, \beta}(t) := \begin{cases} 
1 - \frac{(t-\rho)}{\beta} , & \text{if } \rho \leq t \leq \beta + \rho \\
[t < \rho] , & \text{otherwise} 
\end{cases}
$$

where $1 > \rho, \beta > 0$, and $[\cdot]$ denotes the Iverson bracket for indicator functions. From its definition, we note that $l^{\rho, 0}(t) = [t < \rho]$ and that it verifies the following inequality for all $\rho, \beta > 0$ and $t \in \mathbb{R}$:

$$
l^{\rho, 0}(t) = [t < \rho] < l^{\rho, \beta}(t) < l^{\rho + \beta, 0}(t) = [t < \rho + \beta] \quad (1)
$$

illustrated in Figure 1.

For any domain $\mathcal{P}$ with marginal feature distribution $D_\mathcal{P}$ and any hypotheses $h, f$, we define their disagreement associated to the loss $l^{\rho, \beta}$ as

$$
\epsilon_{\mathcal{P}}^{\rho, \beta}(h, f) := \mathbb{E}_{x \sim D_\mathcal{P}} [l^{\rho, \beta}(f(x)h(x))].
$$
All previous analyses of the DA problem aimed at bounding the target domain error by its source counterpart, an estimable divergence term and a non estimable term representing the a priori adaptability of the problem. Consequently, the major differences between the different available results lie in the considered definition of the error, the divergence measure they introduce and the form of the non estimable term.

To this end, the first rigorous theoretical analysis of domain adaptation, presented in (Ben-David et al., 2007b) (and later in (Ben-David et al., 2010)), introduces the alignment term defined as the divergence and the ideal joint error for two hypotheses in $\mathcal{H}$, respectively defined for a binary hypotheses class $\mathcal{H}$ as follows:

$$d_{\Delta H} := 2 \sup_{h,h' \in \mathcal{H}} \left| \epsilon^0_S(h,h') - \epsilon^0_T(h,h') \right|,$$

$$\lambda := \inf_{f \in \mathcal{H}} \left( \epsilon^0_S(f) + \epsilon^0_T(f) \right).$$

Note that the $\Delta H$-divergence between the two domains does not require strict equality between distributions over all measurable sets, but only over supports of hypotheses of the $\Delta H$ space defined as the space of symmetric differences between hypotheses in $\mathcal{H}$: if $h, h' \in \mathcal{H}$ take their values in $\{0, 1\}$, it is simply equal to hypothesis $\{|h - h'|\}$, i.e., the hypothesis that takes value 1 if and only if $h$ and $h'$ disagree.

The authors show that this quantity is estimable from finite samples and that the sample complexity of such estimation depends on the VC-dimension of the considered hypothesis space $\mathcal{H}$. As for $\lambda$, it cannot be estimated as it involves the target domain’s labels, to which one has no access in the setting of unsupervised domain adaptation.

**Remark 1.** If $h, h' \in \mathcal{H}$ take their values in $\{-1, 1\}$, their disagreement at a point $x$ is equal to $-h(x).h'(x)$, as this latter equals 1 if and only if $h(x)$ and $h'(x)$ have opposite labels.

**Remark 2.** A similar bound for 0-1 loss with the Wasserstein distance and the same $\lambda$ term was proved in (Shen et al., 2017).

This result was further generalized in (Mansour et al., 2009) where the authors considered an arbitrary symmetric loss function $l$ verifying the triangle inequality. This assumption makes their result more general than the one in (Ben-David et al., 2010) proved only for 0-1 loss. After considering the expected $l$--disagreement between two hypotheses on a marginal distribution $D_T$ defined as

$$\mathcal{L}_{D_T}(h, h') := E_{x \sim D_T} [l(h(x), h'(x))],$$

the authors define their discrepancy distance as

$$\text{disc}_l(D_S, D_T) := \sup_{h, h' \in \mathcal{H}} \left| \mathcal{L}_{D_S}(h, h') - \mathcal{L}_{D_T}(h, h') \right|.$$

Such a generalization to arbitrary loss function was later also provided for the bounds with the Wasserstein distance between joint (Courty et al., 2017) and marginal domains distributions (Redko et al., 2016). While being more general than the bound of (Ben-David et al., 2010)

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1As mentioned in (Mansour et al., 2009), the bounds based on the discrepancy distance are in general incomparable to those of (Ben-David et al., 2007b).
new insights on domain adaptation problem by introducing the margin violation rate and scoring functions that give the confidence level of belonging to a class of interest rather than functions with binary output. However, as they bound the 0-1 loss on the target domain, i.e., \( \epsilon_T^{0,0}(h, f) \), their bound does not indicate the behaviour of the margin violation rate on this latter. As for \( \lambda^{(\beta)} \), it remains conceptually similar to the \( \lambda \) term of the other bounds with the only difference consisting in the definition of the error terms.

We now proceed to the presentation of our main theoretical contributions.

3. Bounding the Target Margin Violation Risk

This section is dedicated to our theoretical contributions. We begin by bounding the margin violation rate on the target domain for a classifier \( h \) picked from a given hypothesis class \( \mathcal{H} \). This bound introduces a new divergence term for which we provide a convex proxy afterwards. All of the proofs can be found in the supplementary material.

3.1. Bound with non convex divergence between distributions

The theorem below aims at providing a first theoretical result for DA that includes only interconnected terms depending on the margin of the considered hypothesis. Such interdependence allows to better highlight the possible trade-offs between the different terms in the bound and to gain new insights into the conditions leading to a successful adaptation.

**Theorem 1.** Assume that for any \( h' \in \mathcal{H} \), we have \( \mathbb{P}_{x \sim D_T} [h'(x) = 0] = \mathbb{P}_{x \sim D_T} [h'(x) = 1] = 0 \). Let \( \rho, \beta, \alpha > 0 \) be such that \( \rho + \beta < \alpha < 1 \). Then, for any \( h \in \mathcal{H} \), the following bound holds:

\[
\epsilon_T^{0,0}(h) \leq \epsilon_S^{\rho,\alpha}(h) + d_{h,\mathcal{H}_T}(D_S, D_T) + \lambda_\alpha
\]

where

\[
d_{h,\mathcal{H}_T}(D_S, D_T) := \sup_{h' \in \mathcal{H}_T} \left| \epsilon_S^{\rho,\beta}(h, h') - \epsilon_T^{\rho,\beta}(h, h') \right|
\]

and

\[
\lambda_\alpha := \inf_{f \in \mathcal{H}_T} \epsilon_T^{0,0}(f) + \epsilon_S^{0,0}(f) + \mathbb{P}_{x \sim D_S} [\lvert f(x) \rvert < \alpha].
\]

**Proof idea.** The proof uses the fact that for any \( s, t, u, v \geq 0, st \leq uv \Rightarrow s \leq u \) or \( t \leq v \) in order to bound probabilities of margin violation. The rest follows from the techniques used to establish the bound in (Ben-David et al., 2010) and on inequality (1).

While being similar in shape to the previous DA bounds (Ben-David et al., 2007b; Mansour et al., 2009; Zhang et al., 2019), the obtained result has several fundamental distinctions. First, our bound concerns the margin violation rate on the target domain, rather than the misclassification rate used in Equation (3) (Zhang et al., 2019) and thus, it offers a better estimation of the quality of separation between the classes in the target domain. Second, its estimable part given by the alignment term \( d_{h,\mathcal{H}_T}(D_S, D_T) \) does not introduce the decision function associated to a hypothesis (its sign in case of binary classification with labels encoded as \(-1\) and \(1\)) as in Equation (3) (Zhang et al., 2019), and thus avoids discontinuities in this term making it more suitable for optimization algorithms. One can further show that it is an integral probability metric (Müller, 1997) as for a fixed \( h \in \mathcal{H} \), the set \( \{ x \mapsto t^{\rho,\beta}(h(x)h((x))); h' \in \mathcal{H}' \} \) is a class of bounded measurable functions implying that \( d_{h,\mathcal{H}_T}(D_S, D_T) \) is a pseudometric on the space of probability distributions over \( \mathcal{X} \). Finally, the non estimable term \( \lambda_\alpha \) is non symmetric with respect to the the source and target domains’ roles as, in addition to having low errors in both domains, it requires only a large absolute margin on the source domain, reflected by \( \mathbb{P}_{x \sim D_S} [\lvert f(x) \rvert < \alpha] \) where \( f_\alpha \) is the function achieving the minimum of \( \lambda_\alpha \). This latter thus reflects an intuitively understandable behaviour: if one has a large margin of separation on \( S \), i.e., there exists \( \alpha \) large enough with \( \mathbb{P}_{x \sim D_S} [\lvert f_\alpha(x) \rvert < \alpha] \) small enough, the space of classifiers that are good for the source becomes bigger. Consequently, it becomes more likely to find among them a classifier that is not only good on the source, but on the target domain too. This claim is also supported by the fact that for fixed \( \rho, \beta > 0 \), the source error term is the violation rate of margin \( \frac{\rho + \beta}{\alpha} \): as \( \alpha \) increases, that rate decreases, implying that less concentration on the performance in the target domain is needed. Of course, no gain can be obtained from augmenting \( \alpha \) if it considerably increases \( \lambda_\alpha \).

Additionally, compared to non estimable terms from previous works, \( \lambda_\alpha \) is at least equal to its counterpart \( \lambda \) from (Ben-David et al., 2010), and is not directly comparable to \( \lambda^{(\rho)} \) from (Zhang et al., 2019). However, it verifies the following inequality:

\[
\lambda_\alpha \leq \min_{f \in \mathcal{H}_T} \epsilon_S^{0,0}(f) + \epsilon_T^{0,0}(f)
\]

This is not very different from \( \lambda^{(\rho)} \), as this latter minimizes the joint expectation of the \( t^{\rho,\beta} \) loss, while the right hand side in the previous bound minimizes the joint rate of violating margin \( \alpha \). Finally, \( \beta \), for the moment, appears as the cost of the Lipschitz character of the loss function used in defining the discrepancy term. Its role will become clear when we bound our alignment term by a convex one, using the Lipschitz property.

In the following corollary, we formerly link our bound to that of (Ben-David et al., 2007b).
Corollary 1. If $\mathcal{H} = \mathcal{H}'$ is a class of binary hypotheses taking values in $\{-1, 1\}$, the bound from Theorem 1 is equivalent to the one in (Ben-David et al., 2010).

This corollary shows that our bound generalizes that of (Ben-David et al., 2010) to more informative scoring functions, and to the margin violation rate criterion, without requiring the loss function to verify the triangle inequality as in (Mansour et al., 2009).

### 3.2. A convex domain divergence based on optimal transport

Although the previous bound offers several novel insights on the behaviour of the target error with respect to different components of the bound, its estimable part is non convex as a function of hypothesis $h \in \mathcal{H}$. In order to convexify its two components, i.e., the error on the source domain and the divergence term, it is sufficient to use a convex loss proxy for the first, while for the second we propose to leverage optimal transport (OT) theory (Santambrogio, 2016). To this end, let us introduce two projection operators $\pi_1 : (x_1, x_2) \mapsto x_1$ and $\pi_2 : (x_1, x_2) \mapsto x_2$ defined over $\mathcal{X} \times \mathcal{X}$. The set $\Pi$ of transport plans between $\mathcal{D}_S$ and $\mathcal{D}_T$ is the set of probability distributions $D$ over $\mathcal{X} \times \mathcal{X}$ that verify the following two properties:

$$\pi_1 \# D = \mathcal{D}_S, \quad \pi_2 \# D = \mathcal{D}_T,$$

where $\#$ denotes the pushforward measure. With the previous notations, the convex bound for $d_{h,\mathcal{H}}^{\alpha, \beta}(\mathcal{D}_S, \mathcal{D}_T)$ is given in the following proposition.

**Proposition 1** (Convex bound for alignment term). For any $\rho, \beta > 0$, we have

$$d_{h,\mathcal{H}}^{\rho, \beta}(\mathcal{D}_S, \mathcal{D}_T) \leq \frac{1}{\beta} \inf_{D \in \Pi} \Delta_{\mathcal{H}'}(h, D),$$

where

$$\Delta_{\mathcal{H}'}(h, D) := \sup_{h' \in \mathcal{H}', s, t \sim \mathcal{D}} \mathbb{E} \left[ |h'(x_s) - h'(x_t)| \right].$$

**Proof idea.** From the $\frac{1}{\beta}$-lipschitzness of $b_{\rho, \beta}$, we use the dual form of the Wasserstein distance between distributions $h h' \# \mathcal{D}_S$ and $h h' \# \mathcal{D}_T$, then its primal form (with an infimum), and finally the inf-sup inequality. 

To see the convexity of the introduced alignment term, we note that $d_{h,\mathcal{H}}^{\rho, \beta}$ can be bounded by $\Delta_{\mathcal{H}'}(h, D)$ for any transport plan $D$. Furthermore,

$$\{ h \mapsto \mathbb{E}_{x_s, x_t \sim \mathcal{D}} \left[ |h'(x_s) - h'(x_t)| \right], h' \in \mathcal{H}' \}$$

and

$$\{ D \mapsto \mathbb{E}_{x_s, x_t \sim \mathcal{D}} \left[ |h'(x_s) - h'(x_t)| \right], h' \in \mathcal{H}' \}$$

defined for a fixed $D \in \mathcal{D}$ and a fixed $h \in \mathcal{H}$, respectively are two families of convex functions in $h$ and in $D$. Thus, taking the supremum over $h' \in \mathcal{H}'$ is as well convex in $h$ (resp. in $D$). We note that the function $h \mapsto \inf_{D \in \Pi} \Delta_{\mathcal{H}'}(h, D)$ is not necessarily convex, but when we derive our algorithm, we show that only convexity of $\Delta_{\mathcal{H}'}(\cdot, \cdot)$ is needed.

The bound in Proposition 1 also has the form of a robust version of the Wasserstein distance between 1D distributions $h h' \# \mathcal{D}_S$ and $h h' \# \mathcal{D}_T$ and admits the following robust interpretation: for a fixed joint distribution $D \in \mathcal{D}$, taking the supremum over $h' \in \mathcal{H}'$ is trying to separate the two domains, while taking $D$ that achieves the infimum resists to this separation.

Combined with Theorem 1, Proposition 1 allows to immediately deduce a domain adaptation bound involving the introduced OT-based divergence.

**Proposition 2** (Optimal transport bound on the target risk). With the assumptions and notations of Theorem 1 and Proposition 1, we have for any $h \in \mathcal{H}$:

$$\epsilon_T^0(h) \leq \epsilon_T^{\alpha, \beta, 0}(h) + \frac{1}{\beta} \inf_{D \in \Pi} \Delta_{\mathcal{H}'}(h, D) + \lambda_\alpha. \quad (4)$$

Compared to other DA bounds involving the Wasserstein distance (Redko et al., 2016; Courty et al., 2017; Shen et al., 2017), our divergence term takes into account the considered hypothesis classes, making it a pseudo-metric that is less strict than the Wasserstein distance between marginal distributions of the domains. To support this claim, we bound our optimal transport based alignment term by the Wasserstein distance between the two domains.

**Proposition 3** (Bounding by the p-Wasserstein distance). Let $p \geq 1$, and assume that all of the hypotheses from $\mathcal{H}$ and $\mathcal{H}'$ verify the $L-$Lipschitz continuity with respect to $\|\cdot\|_p$ norm for some $L > 0$. Then, the following holds

$$\sup_{h' \in \mathcal{H}' \atop D \in \Pi} \inf_{h \in \mathcal{H}} \Delta_{\mathcal{H}'}(h, D) \leq 2L W_p(\mathcal{D}_S, \mathcal{D}_T)$$

where

$$W_p(\mathcal{D}_S, \mathcal{D}_T) := \inf_{D \in \Pi} \mathbb{E}_{x_s, x_t \sim \mathcal{D}} \left[ \|x_s - x_t\|_p \right]$$

is the $p$-Wasserstein distance.

**Proof idea.** The proof relies on the triangle inequality verified by the absolute value, the $L-$lipschitzness and the boundedness of the classifiers, as well as the monotonicity of the inf and the sup operators.

This inequality comes essentially from the fact that the space of $L-$Lipschitz functions is bigger than the considered hypothesis spaces $\mathcal{H}$ and $\mathcal{H}'$, and the supremum over $h \in \mathcal{H}$...
is due to the independence of the Wasserstein distance between distributions of classifier \( h \). It formally shows that the attachment of our alignment term to the task at hand, via \( h \in \mathcal{H} \) and the supremum over \( h' \in \mathcal{H}' \), makes it far less strict than the Wasserstein distance between data marginal distributions between the domains.

4. Domain Adaptation Algorithm

With our considered setting and notations, the goal of our domain adaptation task is to find \( h \in \mathcal{H} \) such that \( \epsilon_h^\rho,\delta(h) \), the margin violation rate on the target domain, is as small as possible. As we assume that no access to the labels of \( T \) is given, we look for a hypothesis that minimizes the estimable part of our bound of Proposition 2.

4.1. Minimizing the estimable part of the bound

To derive the objective function for the estimable part of the bound, we have to consider two terms. First is the margin violation error on the source domain \( \epsilon_h^\rho,\delta(h) \) (where \( \rho' = \frac{\epsilon + \delta}{\alpha} \)) whose minimization is known to be a NP-hard problem (Arora et al., 1997). Hence, we replace it by its commonly used proxy, the hinge risk \( \mathbb{E}_{x,y \sim S}[\max(\rho' - y, h(x))] \). The second term is the domain alignment term, which is the infimum over \( D \in \mathcal{D} \) of convex functions of \( h \), as we mentioned earlier. The resulting optimization problem is then given as follows:

\[
\min_{h \in \mathcal{H}, D, D'} \mathbb{E}_{x,y \sim S}[\max(\rho' - y, h(x))] + \frac{1}{\beta} \Delta_{\mathcal{H}'}(h, D) \tag{5}
\]

We would like to stress out that the cost function in this case contains a supremum over the potentially infinite hypothesis space \( \mathcal{H}' \), hence, it might be difficult to solve problem (5), even though convexity is verified. However, we show in the next section that a particular choice of \( \mathcal{H} \) and \( \mathcal{H}' \) allows one to deal efficiently with this term.

Below, we specify the proposed method to a particular case of linear classifiers thus introducing a shallow adversarial DA approach.

4.2. Application to linear classification

We consider our algorithm’s formulation in the linear classification case, where \( \mathcal{H} \) is the space of \( \ell^2 \) bounded linear classifiers, and \( \mathcal{H}' \) is the space of \( \ell^1 \) bounded classifiers.

**Proposition 4.** Let \( \mathcal{H} \) be the space of linear classifier with bounded \( \ell^2 \) norm, and \( \mathcal{H}' \) be the space of linear classifiers with bounded \( \ell^1 \) norm. Let \( l \) denote the hinge loss, \( D_{st} := x_i x_j^T - x_i x_j^T \) and \( \|D_{st}w\|_1 := \|D_{st}w\|_1 \) for \( 1 \leq i \leq d \), where \( w \in \mathbb{R}^d \). Then, Problem (5) can be equivalently expressed as the following convex program:

\[
\min_{w \in \mathbb{R}^d, x_i, x_j \sim D} \mathbb{E}_{x,y \sim S}[l(y, w^T x)] + \delta \mathbb{E}_{x_i, x_j \sim D}[\|D_{st}w\|_1] + \zeta \|w\|_2^2 \tag{6}
\]

where \( \delta, \zeta > 0 \) are two hyper-parameters related to the bounds on \( \mathcal{H} \) and \( \mathcal{H}' \).

**Proof idea.** The supremum over \( \mathcal{H}' \) is a supremum of a convex function on the \( \ell_1 \) unit ball, hence over its vertices which appears via the \( \infty \)-norm. The regularization is to take into account the boundedness of classifiers from \( \mathcal{H} \).

The previous proposition introduces two hyper-parameters linked to regularization of the classifier and to the alignment term. The further the domains are from each other, the more concentration we need on the alignment term, which is achieved by increasing \( \delta \). Also, we note that this is a strongly convex optimization problem, due to the strong convexity of the regularization \( \|w\|_2^2 \) which is an important feature for numerical optimization with gradient descent, thus justifying our choice of the space \( \mathcal{H} \).

4.3. Learning in similarity induced spaces

In the previous section, we chose \( \mathcal{H}' \) as the space of linear classifiers with bounded \( \ell^1 \) norm in order to tackle the difficulty of computing the supremum over \( h' \in \mathcal{H}' \), transforming it into a maximum over the vertices of the unit \( \ell^1 \) ball. However, the main idea behind this choice is not only to simplify the computation of the supremum, but also to have a theoretical justification for our algorithm when the data lies in a space induced by an \( (\epsilon, \gamma, \tau) \)-good similarity function, introduced in the seminal paper (Balcan et al., 2008). Indeed, this latter theoretically guarantees the existence of a low-error linear classifier with bounded \( \ell^1 \) norm in the similarity space defined by transformation

\[
\Psi(x) = (K(x, x_1), ..., K(x, x_L)) \tag{7}
\]

where \( \{x_1, ..., x_L\} \) is a finite set of landmarks usually defined as a subset of the original data set.

Hence, if we consider all of the points from both the source and target domains to be landmarks, then the existence of a good similarity function able of separating the classes implies the existence of an \( \ell_1 \)-bounded classifier that performs well on both domains, i.e., an ideal joint hypothesis with a low error in the similarity induced space. Thus, in the next experimental section, we solve Problem (6) after applying the mapping \( \Psi \), but with a regularization term \( \|w\|_2^2 \).

Note that this latter that does not depend on the similarity matrix \( \{K_{ij}\} := K(x_i, x_j) \) for \( x_i, x_j \in \mathcal{X} \), unlike in the case of kernelized approaches where it is equal to \( w^T Kw \), with \( K \) being the kernel matrix. We note further that in
We use smooth proxies of the positive part \( (\cdot)_+ \), the absolute value \( |\cdot| \) and the infinite norm \( \|\cdot\|_\infty \) (for more details, we refer the interested reader to the supplementary material).

4.4. Optimization procedure of the discrete problem

In the empirical case, one has access to finite data sets \( S_m = \{ (x_1, y_1), \ldots, (x_m, y_m) \} \sim S^m \), and \( T_n = \{ (x'_1, y'_1), \ldots, (x'_n, y'_n) \} \sim T^n \), where the labels of \( T_n \) are not used for learning classifier \( w \). We define \( \hat{\Pi} \), the empirical counterpart of \( \Pi \), as the set:

\[
\hat{\Pi} = \{ \Gamma \in \mathbb{R}^{m \times n}; \Gamma 1_m = \frac{1}{m} 1_m; \Gamma^T 1_m = \frac{1}{n} 1_n \}
\]

where \( 1_p = (1, 1, \ldots, 1) \in \mathbb{R}^p \). Denoting \( D_{ij} = x_i x_j^T - x'_i x'_j^T \in \mathbb{R}^d \times d \), the empirical cost function of Problem (6) becomes:

\[
\frac{1}{m} \sum_{1 \leq i \leq m} l(y_{s,i}, w^T x_{s,i}) + \delta \sum_{1 \leq i \leq m} \sum_{1 \leq j \leq n} \gamma_{ij} |D_{ij} w| + \zeta \|w\|_2^2
\]

which is a function of \( w \in \mathbb{R}^d \) and \( \Gamma \in \hat{\Pi} \) having elements \( \gamma_{ij} \).

Similar to (Courty et al., 2017), the objective function of our minimization problem is convex in two sets of variables: the classifier \( w \) and a transport matrix \( \Gamma \). Following their procedure, we use block coordinate descent (Grippo & Sciandrone, 2000) which alternates between the two steps:

1. For a fixed transport matrix \( \Gamma \), minimize over \( w \). To this end, we use the L-BFGS quasi-Newton method.

2. For a fixed linear classifier \( w \), the minimization over \( \Gamma \) only involves the term multiplied by \( \delta \) in (8), and due to the positivity of all coordinates of vector \( \sum_{ij} \gamma_{ij} |D_{ij} w| \), this minimization is equivalent to:

\[
\min_{q \in \Delta_d} \max_{\Gamma \in \hat{\Pi}} \left( -\sum_{ij} \gamma_{ij} q^T |D_{ij} w| \right)
\]

where \( \Delta_d \) is the probability simplex in dimension \( d \). In this case, we use the min-max algorithm from (Blankenship & Falk, 1976, Algorithm 2.2) to find the optimal transport matrix \( \Gamma \).

We use smooth proxies of the positive part \((\cdot)_+\), the absolute value \( |\cdot| \) and the infinite norm \( \|\cdot\|_\infty \) (for more details, we refer the interested reader to the supplementary material).

5. Empirical Evaluation

In this section, we evaluate our method on two domain adaptation problems: a toy set with controllable adaptation difficulty and a real-world sentiment analysis problem. For all experiments, we use the version of our algorithm specialized to linear classifiers as described in problem (6). We further use a similarity function \( K \) to be specified for each data set considered as in Equation (7) to calculate the features from the raw data. Finally, we denote our method by MADAOT following the abbreviation of our paper’s title. The code for the different experiments is available on this link\(^2\).

5.1. Hyper-parameter tuning

Hyper-parameter tuning is a longstanding problem in unsupervised domain adaptation that was mainly addressed by the introduction of the reversed validation procedure proposed in (Zhong et al., 2010; Bruzzone & Marconcini, 2010). Although this latter may seem as the most suitable cross-validation procedure for the unlabeled scenario, it was shown to fail at selecting the best hyper-parameters for several methods (Wilson & Cook, 2019, Section 8.2), (Bousmalis et al., 2016). One possible reason for this failure is its dependence on accurate estimation of the ratio between the marginal distributions that was proved to require a very large number of samples to be approximated correctly (Ben-David & Urner, 2012).

Hence, we choose to present our algorithm’s performance for two cases. In the first one, we do not use target labels during training phase, but we use them as a validation set to select the best hyper-parameters (defined in Proposition 4) via a 5-fold cross-validation procedure for 10 values of \( \delta \) ranging from \( 10^{-2} \) to \( 10^2 \), and 10 values for \( \zeta \) from \( \{10^{-6}, 10^{-5}\} \), both on a logarithm scale. This is a rather standard procedure in unsupervised domain adaptation used in several other papers on the subject (Courty et al., 2015; Bousmalis et al., 2016). We use this procedure for the first dataset. As for the real-world data set, we run all experiments by setting \( \delta = 1 \) and \( \zeta = 10^{-5} \).

5.2. Inter-twinning moons data set

We carry on our experiments on the moons data set used in (Courty et al., 2015). For this data set, the source domain’s data sample is represented by two inter-twinning moons centered at the origin \((0, 0)\), and composed of 300 instances. The source domain’s data are then rotated around their center by a certain angle to get the target domain data. Obviously, the greater is the angle, the further from each other the two domains are and the harder is the adaptation. Similar to (Courty et al., 2015), we cope with the non-linearity of this

\(^2\)https://github.com/sofiendhouib/MADAOT.
Table 1. Average accuracy (percentage) over 10 realizations for the moons toy set.

<table>
<thead>
<tr>
<th>Angle (°)</th>
<th>10</th>
<th>20</th>
<th>30</th>
<th>40</th>
<th>50</th>
<th>70</th>
<th>90</th>
</tr>
</thead>
<tbody>
<tr>
<td>SVM (Courty et al., 2015)</td>
<td>100</td>
<td>89.6</td>
<td>76</td>
<td>68.8</td>
<td>60</td>
<td>26.6</td>
<td>17.2</td>
</tr>
<tr>
<td>OT-GL (Courty et al., 2015)</td>
<td>100</td>
<td>100</td>
<td>98.7</td>
<td>80.4</td>
<td>62.2</td>
<td>49.2</td>
<td></td>
</tr>
<tr>
<td>JDOT (Courty et al., 2017)</td>
<td>98.9</td>
<td>95.5</td>
<td>90.6</td>
<td>86.5</td>
<td>81.5</td>
<td>70.5</td>
<td>60</td>
</tr>
<tr>
<td>MADAOT</td>
<td>99.5</td>
<td>99.3</td>
<td>99.6</td>
<td>98.9</td>
<td>98.9</td>
<td>77</td>
<td>64.1</td>
</tr>
</tbody>
</table>

Table 2. Accuracy on the Amazon Reviews data set.

<table>
<thead>
<tr>
<th>Task</th>
<th>B → D</th>
<th>B → E</th>
<th>B → K</th>
<th>D → B</th>
<th>D → E</th>
<th>D → K</th>
<th>E → B</th>
<th>E → D</th>
<th>E → K</th>
<th>K → B</th>
<th>K → D</th>
<th>K → E</th>
</tr>
</thead>
<tbody>
<tr>
<td>SVM (CV)</td>
<td>79.5</td>
<td>69.2</td>
<td>71.4</td>
<td>78.5</td>
<td>71.7</td>
<td>76.6</td>
<td>71</td>
<td>73.8</td>
<td>85.3</td>
<td>72.3</td>
<td>73.4</td>
<td>84.4</td>
</tr>
<tr>
<td>DANN</td>
<td>80.6</td>
<td>74.7</td>
<td>76.7</td>
<td>74.7</td>
<td>73.8</td>
<td>76.5</td>
<td>71.8</td>
<td>72.6</td>
<td>85</td>
<td>71.8</td>
<td>73</td>
<td>84.7</td>
</tr>
<tr>
<td>OT-GL</td>
<td>75.7</td>
<td>75.4</td>
<td>77.9</td>
<td>75.9</td>
<td>70</td>
<td>78</td>
<td>71.2</td>
<td>69.6</td>
<td>81.4</td>
<td>73.1</td>
<td>74.1</td>
<td>81.7</td>
</tr>
<tr>
<td>JDOT$_{SVM}$</td>
<td>75</td>
<td>77.1</td>
<td>77.9</td>
<td>70.9</td>
<td>78.3</td>
<td>78</td>
<td>65.9</td>
<td>71.7</td>
<td>79.4</td>
<td>66.8</td>
<td>66.1</td>
<td>77.5</td>
</tr>
<tr>
<td>JDOT$_{NN}$</td>
<td>79.5</td>
<td>78.1</td>
<td>79.4</td>
<td>76.3</td>
<td><strong>78.8</strong></td>
<td><strong>82.1</strong></td>
<td>74.9</td>
<td>73.7</td>
<td>87.2</td>
<td>72.8</td>
<td><strong>76.5</strong></td>
<td>84.5</td>
</tr>
<tr>
<td>MADAOT</td>
<td><strong>82.4</strong></td>
<td>75</td>
<td><strong>80.4</strong></td>
<td><strong>80.9</strong></td>
<td>73.5</td>
<td>81.5</td>
<td><strong>77.2</strong></td>
<td><strong>78.1</strong></td>
<td><strong>88.1</strong></td>
<td><strong>75.6</strong></td>
<td>75.9</td>
<td>87.1</td>
</tr>
</tbody>
</table>

data set by using a Gaussian kernel as similarity function $K$, where the width parameter is chosen as the mean Euclidean distance between the source instances, as suggested in (Kar & Jain, 2011). As for the baselines, our algorithm is compared to SVM classifier with Gaussian kernel trained on the source domain (without adaptation) and two optimal transport based domain adaptation algorithms OT-GL (Courty et al., 2015) and JDOT (Courty et al., 2017). Note that we report only the variation of the method proposed in (Courty et al., 2015) with the group-Lasso regularization as this latter was showed to be the most efficient for this data set. Finally, as the results for JDOT on moons were not presented in the original paper, we run it with the hyper-parameter ranges suggested by the authors. The final results averaged over 10 tests on independent data sets of 1000 instances are presented in Table 1. From it, we can make several conclusions. First, all considered DA baselines manage to achieve an almost perfect score on the angles from $10°$ to $40°$, while SVM without adaptation has a $30\%$ drop in accuracy for these angles. This shows that moons data set presents a challenging adaptation task that goes beyond the generalization capacities of a standard supervised learning algorithm. As for the DA baselines, their performance is rather not surprising as for these angles the adaptation problem remains fairly easy. Starting from $50°$ (Table 1(left)) and up to $90°$ (Table 1(left)), out method provides a better performance that those obtained with both JDOT and OT-GL with the most significant improvement obtained for the angle of $50°$. One should note that OT-GL method relies on the information about the source labels encoded in the group-lasso term but even this does not help to maintain its performance for larger angles. We conclude by saying that the theoretical features of the introduced OT-based distance used by our algorithm are highlighted by its efficiency in this experiment compared to strong OT baselines.

5.3. Sentiment analysis data set

Below, we consider the famous Amazon product reviews dataset (Blitzer et al., 2007) related to the sentiment analysis task. For this data set, we choose 4 of its subsets corresponding to different product categories, namely: books, dvd, electronics and kitchen (denoted by B, D, E, K, respectively). This leads to 12 domain adaptation tasks of varying difficulty as the proximity and the number of semantic relationship between the different domains vary a lot. As the original data is represented by over 100 000 features given by uni- and bigrams, we follow the pre-processing of (Chen et al., 2011) (resulting in between 20000 and 40000 features) and consider a linear kernel as a similarity function $K$. For each task, we use predefined sets of 2000 instances of source and target data samples for training, and keep 4000 instances of the target domain for testing. We compare our method to SVM with cross-validated hyper-parameters as a baseline, to a state-of-the-art adversarial DA approach DANN (Ganin et al., 2016) and to JDOT (NN) with a neural network used as a classifier, as done in (Courty et al., 2017). As our method uses only linear classifiers, we also run two baseline shallow algorithms for comparison, JDOT with a linear SVM and OT-GL (Courty et al., 2015) with 1-Nearest Neighbor classifier. The results of our experiments are reported in Table 2. From this table, we see that MADAOT outperforms other methods on 8 out of 12 tasks, and has the second best performance on 2 others. This is rather surprising considering that both DANN and JDOT (NN) rely on neural networks to learn the final classifier and this latter...
is expected to have a higher discriminative power than the class of linear classifiers. Consequently, we attribute this performance gain to the efficiency of our task-dependent OT-based alignment term that manages to better align the two distributions compared to the minimization of the original 2-Wasserstein distance considered in OT-GL and JDOT. Furthermore, the minimax formulation of our alignment term addresses the curse of dimensionality problem related to OT as the sample complexity of this latter is known to scale exponentially in dimension. Several approaches were proposed to address this problem recently in order for the calculation of the Wasserstein distance to make sense for high-dimensional data and our findings show that applying it in the DA context can lead to an improved performance.

6. Conclusion and Future Perspectives

In this paper, we presented a novel theoretical analysis of unsupervised domain adaptation problem for binary classification that considers the margin violation loss on the target domain as the error measure. We proved a new bound on this latter that involves source margin violation error, a novel convex alignment term given by a task-dependent variant of the Wasserstein distance between the source and target domains and a non-estimable term that offers new insights on domain adaptation problem and on the importance of the notion of margin violation for its a priori success. Our analysis generalizes several prior works on this subject and includes them as special cases. Our algorithm, derived from the established learning bounds, has proved to be efficient on both simulated and real-world problems compared to several state-of-the-art methods.

The future research directions of this paper are many. First, we would like to study the direct maximization of our non convex alignment term introduced in Theorem 1 using a deep adversarial approach. Even though our method offers a remarkable performance compared to several deep learning baselines, its efficiency can be further improved by this latter extension. Second, we plan to investigate in more detail the theoretical properties of our data dependent optimal transport term by establishing a concentration inequality for this latter. This would allow to theoretically highlight the success of our algorithm for high-dimensional data.

References


