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## An Online Variational Inference and Ensemble Based Multi-label Classifier for Data Streams

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Abstract—Recently, multi-label classification algorithms have been increasingly required by a diversity of applications, such as text categorization, web, and social media mining. In particular, these applications often have streams of data coming continuously, and require learning and predicting done on-thefly. In this paper, we introduce a scalable online variational inference based ensemble method for classifying multi-label data, where random projections are used to create the ensemble system. As a second-order generative method, the proposed classifier can effectively exploit the underlying structure of the data during learning. Experiments on several real-world datasets demonstrate the superior performance of our new method over several well-known methods in the literature.

#### Keywords—variational inference, random projection, ensemble method, online learning, multi-label data stream

#### I. INTRODUCTION

Recently, multi-label classification (MLC) algorithms have been increasingly required by a diversity of applications, such as text categorization, web, and social media mining [1]. It naturally emerged from the multiple meanings of many realworld objects and can be treated as the generalization of multiclass classification (also known as single-label classification), where an instance may have a set of relevant labels instead of only one label. In this paper, we adapt one of our recently published online multiclass classifier (named Online Variational Inference for multivariate Gaussians (VIGO)) [2] to multi-label classification due to its demonstrated superior performance over several well-known methods in the literature. Although many batch MLC algorithms have been proposed in the literature, there is relatively little work being done on online (incremental) MLC [3]. In particular, the demand for effective incremental methods is growing quickly in our big data era, where it is becoming increasingly impractical to store the entire training set in the main memory for batch learning. Moreover, offline methods cannot update its model on-the-fly and must be rebuilt whenever new data arrive, leading to costly operation in real-time applications with streaming data. On the other hand, online methods offer the essential ability of predictive models which can be trained on-the-fly after the arrival of every new data point and be ready to give predictions at any time if requested, by making use of a single/set of observations and then discarding them permanently before the next observations are used.

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Our proposed VIGO [2] employs variational inference (VI) to approximate the distribution of data in each class by a multivariate Gaussian. It is a second-order generative method, which not only gives a point estimate of parameters, but also the distribution of possible solutions. To adapt VIGO to multilabel learning, several difficulties need to be overcome. First, most multi-label datasets are high-dimensional, this can reduce the performance of the VI framework. Second, it is not straightforward to estimate the performance of an online multi-label classifier as an instance can be associated with several labels. Existing performance measures can be inconsistent [4], which leads to the difficulty of building an MLC method that performs well on all performance measures. We propose to use random projections to reduce the dimension of the data before applying VIGO. Since the random projection is data-independent, it is suitable to work with stream data. Moreover, data schemes generated by different random projections are significantly different from each other as well as the original data [5]. This motivated us to construct an ensemble of incremental Bayesian classifiers learned on data schemes generated by randomly projecting the original stream data into a subspace of low dimension. The ensemble scheme further improves the effectiveness of the prediction since an ensemble of classifiers can obtain a better solution than a single classifier. The main contributions of this paper are:

- A flexible generative framework based on variational inference to learn multi-label streaming data proposed
- An ensemble system using random projection proposed to make the online variational inference framework more scalable and efficient
- An empirical demonstration that our method achieves superior performance over many well-known benchmark algorithms.

#### II. RELATED WORK

Generalizing from single-label methods, offline multilabel classifiers can be grouped into 2 main types of approaches: problem transformation and algorithm adaptation [1]. While the former adapts multi-label data to off-the-shell single-label classifiers, the latter adapts a single-label algorithm to produce multi-label outputs. Many multi-label classifiers involve a mix of both approaches.

Regarding the problem transformation group, Binary Relevance (BR) method is used the most. It transforms the multi-label problem to multiple separate binary classification problems, one for each label [7]. As BR does not take into account label correlations explicitly, several methods have been introduced to ease its shortcoming by addressing the correlation between labels in their learning models. The bestknown one is Classifier Chains (CC) [8], which is also trained as many binary models as in BR but chooses them in a random order. After the first classifier is trained with only the original input features, the first output label is then appended to the original instances as a new input feature, and the second classifier is trained on the new input space, and so on. By this way, the chained classifiers attempt to take account of the possible label dependencies. Label Powerset (LP) [7] is another popular approach in this group, where each combination of class labels can be treated as a new label and a single multiclass classifier is needed to obtain the predictions. However, in the LP's framework, the number of class label combinations increase exponentially and the prediction for underrepresented label combinations is very uncertain. To overcome these disadvantages, the sets of labels can be pruned as in Pruned Sets method (PS) [9] so that the learner focuses only on the set of labels with the most important correlations.

For algorithm adaptation group, some well-known approaches are k-nearest neighbor based multi-label classification [10], kernel-based Rank-SVM adapting methods [11], and tree-based hierarchical multi-label classification [12]. Recently, Rank and Threshold method (RT) [13] replicates multi-label instances into single-label instances, then train a multi-class classifier and uses a threshold to give multi-label outputs. It can be considered as both problem transformation and method adaptation.

To deal with big or stream data, several online MLC have been presented. In [3], Read et al. showed how to use problem transformation methods as the building block and combine them with off-the-shelf online single-label classifiers to create streaming multi-label learners. The same authors also adapt incremental multiclass Hoeffding tree [14] to the multi-label scenario [3]. Furthermore, the multi-label Hoeffding tree [3] is combined with the Pruned Set classifier [9] to prune the label combination at each leaf node. These well-known incremental multi-label learners will be used as the benchmark algorithms in our experiments.

#### III. BACKGROUND

#### A. VI for multivariate Gaussian

Let a vector  $\mathbf{x} = (x_1, x_2, ..., x_D) \in \mathbb{R}^D$  be a sample to be classified, and  $Y = \{1, 2, ..., L\}$  be the set of all labels. In a Bayesian framework, a multiclass classifier gives a prediction  $\hat{y}$  to  $\mathbf{x}$  as:

$$\hat{y} \sim \underset{l \in \{1,2,\cdots,L\}}{\operatorname{argmax}} p(y_l) \cdot p(\mathbf{x}|y_l)$$
(1)

where  $p(y_l)$  is the prior probability of class *l*, and  $p(\mathbf{x}|y_l)$  is the class conditional probability.

 $p(y_l)$  is often approximated by:

$$p(y_l) \approx \frac{c_l}{c} \sim c_l$$
 (2)

which is the proportion of samples from class *l* in the set of samples received so far,  $c_l =$ #samples having label *l* to date and  $c = \sum c_l$ .

 $p(\mathbf{x}|\mathbf{y}_l)$  is approximated by a multivariate Gaussian  $\mathcal{N}(\mathbf{x}|\mathbf{\mu}, \mathbf{\Lambda}^{-1})$ . Here we temporarily ignore the sub-index *l* of  $\mathcal{N}(\mathbf{x}|\mathbf{\mu}_l, \mathbf{\Lambda}_l^{-1})$  for simplicity. In our variational inference framework, the parameters  $(\mathbf{\mu}, \mathbf{\Lambda})$  are treated as random variables. We look for the posterior distribution  $p(\mathbf{\mu}, \mathbf{\Lambda}|\mathbb{X}) \approx q(\mathbf{\mu}, \mathbf{\Lambda})$  after giving the model a suitable conjugate prior. Here we assume that the training set  $\mathbb{X} = \{\mathbf{x}_j | j = 1, ..., n\}$  is drawn independently from  $\mathcal{N}(\mathbf{x}|\mathbf{\mu}, \mathbf{\Lambda}^{-1})$ . Below are the resulted distributions of  $\mathbf{\mu}$  and  $\mathbf{\Lambda}$  (see [15] for details).

The parameter  $\mu$  has a Gaussian distribution  $\mu \sim \mathcal{N}(\mu | m, H^{-1})$  with:

$$\mathbf{m} = \frac{\beta_0 \mathbf{m}_0 + N\bar{\mathbf{x}}}{\beta_0 + N} \tag{3}$$

$$\mathbf{H} = (\boldsymbol{\beta}_0 + \boldsymbol{N}) \mathbb{E}[\boldsymbol{\Lambda}] \tag{4}$$

The parameter  $\Lambda$  has a Wishart distribution  $\Lambda \sim \mathcal{W}(\Lambda | \mathbf{W}, v)$  with:

$$v = v_0 + N + 1 \tag{5}$$

$$\mathbf{W}^{-1} = \mathbf{W}_0^{-1} + (\beta_0 + N)\mathbf{H}^{-1} + \mathbf{S} + \frac{\beta_0 N}{\beta_0 + N}\mathbf{J}$$
(6)

Here

$$\bar{\mathbf{x}} = \frac{1}{N} \sum_{n=1}^{N} \mathbf{x}_n \tag{7}$$

$$\mathbf{S} = \sum_{n=1}^{N} (\mathbf{x}_n - \bar{\mathbf{x}}) (\mathbf{x}_n - \bar{\mathbf{x}})^T$$
(8)

$$\bar{\mathbf{x}} = \frac{1}{N} \sum_{n=1}^{N} \mathbf{x}_n \tag{9}$$

are the sufficient statistics.  $\beta_0$ ,  $\mathbf{m}_0$ ,  $\nu_0$ ,  $\mathbf{W}_0$  are the prior information.

Thus, we have Algorithm 1 for multivariate Gaussian distribution estimation based on variational inference (VIG) [15]. The stop condition of the iterative re-estimation procedure is the minor change ( $<\varepsilon$ ) of the lower bound value  $\mathcal{L}(q) = \ln p(\mathbb{X}) - KL(q||p)$  [15].

Algorithm 1. VIG ( $\mathbb{X}$ , $\mathbf{m}_0$ , $\beta_0$ , $v_0$ , $\mathbf{W}_0$ )
<b>Input</b> : Dataset X, $\varepsilon$ , $\mathbf{m}_0$ , $\beta_0$ , $v_0$ , $\mathbf{W}_0$ , $\mathbb{E}[\mathbf{\Lambda}] = v_0 \mathbf{W}_0$
Output: m, H, W, v
for $i = 1$ to 2 do
Calculate $\mathbf{m}$ , $\mathbf{H}$ as in (3), (4)
Calculate $v$ , <b>W</b> as in (5), (6)
while $\mathcal{L}_i(q) - \mathcal{L}_{i-1}(q) < \varepsilon$ do
Calculate $\mathbf{m}$ , $\mathbf{H}$ as in (3), (4)
Calculate $v$ , <b>W</b> as in (5), (6)
$i \coloneqq i + 1$

The default setting for Algorithm 1 is as follows:  $\varepsilon = 1e - 10$ ; *D*-dimension vector  $\mathbf{m}_0^{(j)} = (0, ..., 0)^T$ ;  $\beta_0^{(j)} = 0$ ;  $v_0^{(j)} = D$ ;  $D \times D$  dimension matrix  $\mathbf{W}_0^{(j)} = \mathbf{I}$  (an identity matrix), j = 1, ..., L. When the point estimate of a parameter is needed, we often pick the mean of their distribution. For example,  $\mathbb{E}[\mathbf{\mu}] = \mathbf{m}$  and  $\mathbb{E}[\mathbf{\Lambda}] = v\mathbf{W}$  can be used as the point estimate of  $\mathbf{\mu}$ , and  $\mathbf{\Lambda}$ , respectively.

Based on VIG, we built the online classifier VIGO [2]. When a small batch of size *n* of new observations  $\in$  class *l* comes, they can be used as dataset X. From the present state of the model, we extract the prior  $\mathbf{m}_0$ ,  $\beta_0$ ,  $v_0$ ,  $\mathbf{W}_0$ . After that, VIGO update the predictive model  $\mathcal{N}(\mathbf{x}|\mathbf{\mu}, \mathbf{\Lambda}^{-1})$  for class *l* and assign new values for  $\mathbf{m}_0$ ,  $\beta_0$ ,  $v_0$ ,  $\mathbf{W}_0$ . While  $\mathbf{m}_0$ ,  $v_0$ ,  $\mathbf{W}_0$  can take the value of updated  $\mathbf{m}$ , v,  $\mathbf{W}$ , the new value for  $\beta_0$  is set as

$$\beta_0 \coloneqq \beta_0 + |\mathbb{X}| = \beta_0 + n \tag{10}$$

where  $|\cdot|$  denotes the cardinality of a set. This is to weigh the number of past instances in (3). Furthermore, the sufficient statistics  $\bar{\mathbf{x}}$ ,  $\mathbf{S}$ ,  $\mathbf{J}$  can be calculated in a sequential manner [2], hence there is no need of storing data in the VIGO framework. When a new instance  $\mathbf{x}_t$  arrives, it can be used for updating  $\bar{\mathbf{x}}$ ,  $\mathbf{S}$ ,  $\mathbf{J}$  of its related class, after that it is discarded permanently. VIGO will use the sufficient statistics to update the predictive model  $\mathcal{N}(\mathbf{x}|\mathbf{\mu}, \mathbf{\Lambda}^{-1})$  for the related class when needed (for example, when a wrong prediction is made). After the sufficient statistics are used, they will be reset to store fresh information from the incoming data.

#### B. Random projection (RP)

When applying VIGO to multilabel classification, the high-dimensional data which is very common in multi-label problems can make the approximation process in variational inference less effective, resulting in the degradation of the system performance. Therefore, it is helpful to reduce the data dimension beforehand. We propose to use random projection for the dimensionality reduction as its projection matrix is generated randomly without requiring a validation set, which is often not available in the stream context. It is also inexpensive to generate and is data-independent. Another motivation for using random projection is to make discrete (e.g. binary) attributes become continuous, which is beneficial for VIGO.

In this work, we generate a random projection  $\mathbf{z} \in Q$ -dimensional space  $\mathbb{R}^{Q}$  (called down-space) of a vector  $\mathbf{x} \in D$ -dimensional space (called up-space) as

$$T: \mathbb{R}^D \to \mathbb{R}^Q: \mathbf{z} = T(\mathbf{x}) = \frac{1}{\sqrt{Q}} \mathbf{x} \mathbf{R}$$
 (11)

where  $\mathbf{R} = \{r_{ij}\}$  is a  $D \times Q$  random matrix, the expectation of each entry  $\mathbb{E}(r_{ij}) = 0$ , and its variance  $\operatorname{var}(r_{ij}) = 1$ , and  $Q \ll D$  [16-20]. As in [20], we choose Gaussian RP:  $r_{ij} \sim \mathcal{N}(0; 1)$ , and  $Q = 2\log_2(D)$ . This kind of randomized mapping has been shown to perturb the original data introducing only bounded distortions, approximately preserving their metric structure [16-18].

As opposed to other sampling methods like bootstrapping, the datasets generated by different random matrices can be quite different (diverse) [21]. We, therefore, employ an ensemble system based on a set of K random projections to benefit from this diversity.

#### IV. PROPOSED METHOD

We build an online multi-label classifier based on variational inference and random projections. First, *K* VIGO classifiers  $V_k$ , k = 1, ..., K are initialized. Each  $V_k$  contains *L* sub-models  $V^{(l,k)}$ , l = 1, ..., L, where  $V^{(l,k)}$  is a VIG to approximate the multivariate Gaussian distribution for class *l*. Then, we generate *K* random matrices  $\{\mathbf{R}^{(k)}\}_{k=1,...,K}$ .

When the sample  $\mathbf{x}_t$  arrives, we project it to the  $k^{\text{th}}$  down-space to obtain its projection  $\mathbf{z}_t^{(k)}$ :

$$\mathbf{z}_{t}^{(k)} = \frac{1}{\sqrt{Q}} \mathbf{x}_{t} \mathbf{R}^{(k)}, k = 1, \dots, K$$
(12)

The related classifier  $V_k$  processes  $\mathbf{z}_t^{(k)}$  and give the confidence  $\{p_k(y_l|\mathbf{x}_t)\}$  that  $\mathbf{x}_t$  belongs to class label l (l = 1, ..., L):

$$p_k(y_l | \mathbf{x}_t) \in [0, 1] \text{ and } \sum_{l=1}^{L} p_k(y_l | \mathbf{x}_t) = 1$$
 (13)

We combine the outputs of K base learners

$$\begin{bmatrix} p_1(y_1|\mathbf{x}_t) & \cdots & p_1(y_L|\mathbf{x}_t) \\ \vdots & \ddots & \vdots \\ p_K(y_1|\mathbf{x}_t) & \cdots & p_K(y_L|\mathbf{x}_t) \end{bmatrix}$$
(14)

using Sum rule to obtain the final hypothesis.

$$\hat{Y}_t = \left\{ l \in L \middle| \theta_l(\mathbf{x}_t) = \frac{1}{K} \sum_{k=1}^K p_k(y_l | \mathbf{x}_t) \ge \theta \right\} \quad (15)$$

where  $\hat{Y}_t$  is the predicted label set of  $\mathbf{x}_t$ ,  $\theta_l(\mathbf{x}_t) \in [0; 1]$  is the confidence score for the label  $l, \theta \in [0; 1]$  is a predefined threshold. It is easy to see that  $\sum_{l=1}^{L} \theta_l(\mathbf{x}_t) = 1$ . As in [3], we update the threshold  $\theta$  in the predictive equation (15) by windows of the same size  $\omega$ . Initially,  $\theta$  is set = 0.5 for the first window  $W_0$ . Subsequently, the threshold on window  $W_i$  is the one best approximating the true label cardinality of window  $W_{i-1}$ . In detail, we calculate  $|Y_W|$  – the total number of true labels of all instances in  $W_{i-1}$  and store the confidence scores given for each label of the label set of all instances in  $W_{i-1}$  (this is an array of length  $L \times \omega$ ). We then sort the confidence score as the threshold for window  $W_i$ .

Finally, we use the projected datum  $\mathbf{z}_t^{(k)}$  for training  $V_k$ . Since the dimension of  $\mathbf{z}_t^{(k)}$  is much smaller than that of  $\mathbf{x}_t$ , the learning process will be more time-efficient.

Another concern is that in MLC problems, we have a lot of different measures, and it is not trivial to decide whether the predictive models need to be updated when a multi-label classifier makes wrong or right prediction, unlike in VIGO [2]. To overcome this, a "batch"  $B^{(l,k)}$  will be used to update the sub-model  $V^{(l,k)}$  if it collects information from enough |B|instances and is cleared totally after that to make space for new incoming instances. It is worth mentioning that for each "batch"  $B^{(l,k)}$ , we only maintain its sufficient statistics, every coming instance is handled on-the-fly and discarded right after that. The default value for |B| is 200, which is equal the default grace period of the state-of-the-art Hoeffding tree [14] (i.e. the number of instances a leaf should observe between split attempts). Additionally, to build a stable model and make the base classifiers more diverse, before updating in batches of size |B|, each sub-model  $V^{(l,k)}$  will be updated in batches of size  $|B_1| = 30$  until it has collected at least  $|B_{init}| + k$ instances (building step).

Clearly, our proposed MVI is an RT type method [13] which transforms a multi-class classifier into a multi-label classifier using rank and threshold. Although this type of transformation is simple, we believe that the superior performance of the VI framework, which is boosted further under an ensemble based framework, will make MVI highly efficient and competitive. Similar to VIGO, MVI is expected to work well on all range of data: from rare data to big data. Although outputting high-order information, variational

inference for multivariate Gaussian has been demonstrated to quickly converge to the approximate solution, taking a low time cost [2]. The pseudo-code of MVI is given in Algorithm 2.

Algorithm 2. MVI Input: ensemble size K, down-space dimension Q, threshold  $\varepsilon$ , |B|,  $|B_1|$ ,  $|B_{init}|$  $\text{Initialize: } \mathbf{m}_{0}^{(l,k)}, \beta_{0}^{(l,k)}, \mathbf{v}_{0}^{(l,k)}, \mathbf{W}_{0}^{(l,k)}, c_{l} = 1, l = 1, \dots, L,$ k = 1, ..., K;// Random matrix generation Generate  $\mathbf{R}^{(k)} = \left\{ r_{ij}^{(k)} \right\}, k = 1, \dots, K;$ for t = 1, 2,Instance  $\mathbf{x}_t$  arrives;  $\mathbf{z}_t^{(k)} = \frac{1}{\sqrt{Q}} \mathbf{x}_t \mathbf{R}_k$ , k = 1, ..., K; // Label pediction  $p(y_l) \sim c_l; l = 1, ..., L$ for k = 1, ..., KApply  $V^{(l,k)}$  on  $\mathbf{z}_t^{(k)}$  to output  $p_k(\mathbf{z}_t^{(k)}|y_l)$ ,  $l = 1, \ldots, L;$  $p_k(y_l|\mathbf{x}_t) \sim p(y_l)p_k(\mathbf{z}_t^{(k)}|y_l);$ end for Normalize  $\{p_k(y_l|\mathbf{x}_t)\}$  as in (13); Predict label set of  $\mathbf{x}_t$  using combining rule (15); // Update base classifiers Reveal the true label set of  $\mathbf{x}_t$  from the environment:  $Y_t$ ; for  $l \in Y_t$  $c_l = c_l + 1;$ for k = 1, ..., KUpdate sufficient statistics  $\bar{\mathbf{z}}^{(l,k)}$ ,  $\mathbf{S}^{(l,k)}$ ,  $\mathbf{I}^{(l,k)}$ of  $V^{(l,k)}$ ;  $|B^{(l,k)}| = |B^{(l,k)}| + 1$ ; if  $\beta_0^{(l,k)} < |B_{\text{init}}| + k$ Update  $(V^{(l,k)}, |B_1|);$ else Update  $(V^{(l,k)}, |B|);$ end if end for end for end for . . . . . . . . . . **procedure** Update  $(V^{(l,k)}, size);$ if  $|B^{(l,k)}| = size$ Run VIG  $(B^{(l,k)}, \mathbf{m}_{0}^{(l,k)}, \beta_{0}^{(l,k)}, v_{0}^{(l,k)}, \mathbf{W}_{0}^{(l,k)})$  to update  $\mathbf{m}^{(l,k)}, \mathbf{H}^{(l,k)}, v^{(l,k)}, \mathbf{W}^{(l,k)}$  for  $V^{(l,k)}$ ;  $\mathbf{m}_{0}^{(l,k)} = \mathbf{m}^{(l,k)}$ ;  $v_{0}^{(l,k)} = v^{(l,k)}$ ;  $\mathbf{W}_{0}^{(l,k)} = \mathbf{W}^{(l,k)}$ ;  $\beta_{0}^{(l,k)} = \beta_{0}^{(l,k)} + size$ ; Reset  $\bar{\mathbf{z}}^{(l,k)}, \mathbf{S}^{(l,k)}, \mathbf{J}^{(l,k)}; |B^{(l,k)}| = 0$ ; end if

end procedure

#### V. EXPERIMENTS

#### A. Experimental design

We use the prequential (test then train) methodology [22] to evaluate the performance of our proposed methods in stationary and dynamic environments. In our experiment, each algorithm runs on each dataset one time. Samples from a dataset come in a sequential manner to simulate a data stream. An arriving data point is tested to give a set of predicted relevant labels, then it is used to update the learner's model. Based on the prediction results on the whole stream, different performance measures are calculated. Our proposed method and benchmark algorithms are compared in these performance measures and time cost.

In our experiments, we use the subset of the common realworld multi-label datasets listed in [3] (see TABLE I). MVI is compared with several well-known online multi-label methods which are available in MEKA: a recent prevalent library of multi-label classifiers [13]. They are binary relevance BR (Binary Relevance) [7], CC (Classifier Chains) [8], label combination methods PS (Prune Sets) [9], ML (Majority Labelset) [13], rank and RT (Rank + Threshold) [13]. All these methods use the state-of-the-art incremental Hoeffding trees [14] as the base classifiers except ML, which always predicts the most common labelset. Parameters are set as default in MEKA. Default parameters of MVI are K =50,  $|B_{init}| = 1000$ , |B| = 200, and  $|B_1| = 30$ .

Compared with the traditional single-label learning, performance evaluation in multi-label learning is much more complicated as each sample can have several relevant labels at the same time. Therefore, a number of performance measures focusing on different aspects have been proposed in the literature. To ensure a fair and honest evaluation, we use a set of well-known measures, which are used in recent papers about multi-label methods [3, 4]. They are Example-based F1/Accuracy, label-based Micro F1/Macro F1, and ranking-based Average Precision/Ranking Loss. The  $\uparrow$  (or  $\downarrow$ ) symbol next to the measure means that the greater/smaller its value the better.

#### B. Results

The summarized result in Fig. 1 shows that MVI has the best average rank for all measures except Macro F1. For this performance metric, the proposed method ranks second after BR. The more detailed result can be found in TABLE II, where the value of performance measures is supplied together with its per-dataset ranking in parenthesis.

For all 6 measures, MVI, BR and RT are the top 3 methods regarding their average ranks. MVI learns well on 3 datasets: small ENRON, SLASHDOT, and big IMDB. Especially, on SLASHDOT, the performance of MVI is far better than that of all other benchmark methods. BR do well on TMC and OHSUMED.

Overall, our proposed method performs well on all 6 mentioned measure. BR and RT achieve quite competitive results, meanwhile CC, PS have a big variation in performance. Clearly, the simplest method ML performs weakly.

The time cost of all the mentioned algorithms is depicted in TABLE III. MVI has a moderate average time cost (201.86 seconds), which is much smaller than that of BR (534.88 seconds), CC (542.58 seconds). In particular, the proposed method is quicker than the two competitive methods BR and RT. Although MVI are slower than ML and PS, compared to these methods, MVI's classification performance is much better.

#### VI. CONCLUSIONS

We have presented an online multi-label classification algorithm based on variational inference and random projection. Besides achieving superior performance over several well-known benchmark algorithms, MVI has a moderate time cost which can be further reduced by parallelization. As the proposed method is a flexible secondorder generative framework which gives not just a point estimate but a distribution of solutions, our future work will be to investigate its application in other advanced tasks of multi-label learning such as cost-sensitive learning, imbalanced learning and adaptive learning.

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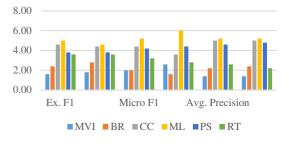
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TABLE I. INFORMATION OF DATASETS USED IN EVALUATION

Dataset	Ν	L	А	$\phi_{ m LC}$
ENRON	1702	53	1001 <i>b</i>	2.0
IMDB OHSUMED	120919	28	1001 <i>b</i>	2.0
	13929	23	1002 <i>n</i>	1.7
SLASHDOT	3728	22	1079 <i>b</i>	1.2
TMC	28596	22	500 <i>b</i>	2.2

n indicates numeric attributes, and b - binary attributes



Average ranks of MVI and benchmark methods

TABLE II.	PERFORMANCE MEASURES OF MVI AND BENCHMARK ALGORITHMS	S
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Fig. 1.

	ENRON	IMDB	OHSUMED	SLASHDOT	TMC	Avg. Rank↓
(a) Ex. F1↑						
MVI	0.4790(1)	0.3132 (1)	0.2835 (3)	0.3552 (1)	0.6153 (2)	1.60
BR	0.4566 (2)	0.2332 (5)	0.3225 (2)	0.1652 (2)	0.6256(1)	2.40
CC	0.3933 (4)	0.0915 (6)	0.2425 (4)	0.0278 (6)	0.6146 (3)	4.60
ML	0.2021 (6)	0.2483 (3)	0.1432 (6)	0.1484 (4)	0.2166 (6)	5.00
PS	0.2778 (5)	0.2420 (4)	0.3316(1)	0.1438 (5)	0.5696 (4)	3.80
RT	0.4422 (3)	0.3095 (2)	0.1986 (5)	0.1628 (3)	0.5378 (5)	3.60
(b) Ex.Accuracy	y↑					
MVI	0.3625 (1)	0.2342 (1)	0.2175 (3)	0.3218 (1)	0.5048 (3)	1.80
BR	0.3417 (2)	0.1710 (5)	0.2579 (2)	0.1451 (4)	0.5199(1)	2.80
CC	0.2979 (4)	0.0710(6)	0.2110 (4)	0.0263 (6)	0.5185 (2)	4.40
ML	0.1680 (6)	0.2081 (3)	0.1270 (6)	0.1458 (2)	0.1786 (6)	4.60
PS	0.2246 (5)	0.2032 (4)	0.2902 (1)	0.1407 (5)	0.4804 (4)	3.80
RT	0.3239 (3)	0.2272 (2)	0.1548 (5)	0.1452 (3)	0.4220 (5)	3.60
(c) Micro F1↑						
MVI	0.4773 (1)	0.3240 (2)	0.3196 (4)	0.3774 (1)	0.6372 (2)	2.00
BR	0.4443 (3)	0.2580 (3)	0.3770 (1)	0.1811 (2)	0.6384 (1)	2.00
CC	0.3990 (4)	0.1263 (6)	0.3206 (3)	0.0509 (6)	0.6346 (3)	4.40
ML	0.1427 (6)	0.2379 (4)	0.1374 (6)	0.1411 (4)	0.1988 (6)	5.20
PS	0.2547 (5)	0.2309 (5)	0.3297 (2)	0.1379 (5)	0.5731 (4)	4.20
RT	0.4723 (2)	0.3267 (1)	0.2251 (5)	0.1611 (3)	0.5340 (5)	3.20
(d) Macro F1↑						
MVI	0.1048(1)	0.0551 (3)	0.0914 (4)	0.1652(1)	0.3494 (4)	2.60
BR	0.0650(2)	0.0859(1)	0.2980 (1)	0.0840 (2)	0.4612 (2)	1.60
CC	0.0440 (5)	0.0452 (4)	0.2589 (2)	0.0468 (4)	0.4456 (3)	3.60
ML	0.0090 (6)	0.0245 (6)	0.0135 (6)	0.0133 (6)	0.0232 (6)	6.00
PS	0.0464 (4)	0.0277 (5)	0.1478 (3)	0.0397 (5)	0.1905 (5)	4.40
RT	0.0474 (3)	0.0561 (2)	0.0345 (5)	0.0562 (3)	0.5173 (1)	2.80
(e) Avg. Precisio	on↑					
MVI	0.5656(1)	0.4695 (2)	0.4643 (2)	0.5281 (1)	0.7669(1)	1.40
BR	0.5265 (3)	0.4153 (3)	0.5170(1)	0.3763 (2)	0.7468 (2)	2.20
CC	0.3709 (4)	0.1841 (6)	0.3684 (5)	0.2547 (6)	0.6423 (4)	5.00
ML	0.2409 (6)	0.3003 (4)	0.2700 (6)	0.2986 (4)	0.2860 (6)	5.20
PS	0.2934 (5)	0.2959 (5)	0.4220 (3)	0.2919 (5)	0.5953 (5)	4.60
RT	0.5438 (2)	0.4754 (1)	0.3809 (4)	0.3442 (3)	0.7122 (3)	2.60
(f) Ranking Los	is↓					
MVI	0.1694 (2)	0.1777 (2)	0.2083 (1)	0.1779 (1)	0.0660(1)	1.40
BR	0.1922 (3)	0.2185 (3)	0.2135 (2)	0.2346 (2)	0.1040 (2)	2.40
CC	0.3803 (4)	0.4934 (6)	0.3652 (5)	0.4048 (6)	0.2254 (4)	5.00
ML	0.5095 (6)	0.4514 (4)	0.4376 (6)	0.3705 (4)	0.5848 (6)	5.20
PS	0.4594 (5)	0.4532 (5)	0.3350 (4)	0.3757 (5)	0.2826 (5)	4.80
RT	0.1196(1)	0.1719(1)	0.2566 (3)	0.2655 (3)	0.1118 (3)	2.20

TABLE III. TIME COST  $\downarrow$  (IN Seconds) OF MVI and Benchmark Algorithms

	ENRON	IMDB	OHSUMED	SLASHDOT	TMC	Average
MVI	24.69	765.91	73.08	16.69	128.94	201.86
BR	58.33	2246.40	184.11	50.54	135.01	534.88
CC	58.58	2257.23	193.73	48.43	154.94	542.58
ML	0.19	1.85	0.36	0.15	0.52	0.62
PS	2.68	57.40	28.64	8.69	15.78	22.64
RT	68.82	1184.84	84.08	26.74	86.32	290.16