Boosting Deep Learning Risk Prediction with Generative Adversarial Networks for Electronic Health Records

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Abstract—The rapid growth of Electronic Health Records (EHRs), as well as the accompanied opportunities in Data-Driven Healthcare (DDH), has been attracting widespread interests and attentions. Recent progress in the design and applications of deep learning methods has shown promising results and is forcing massive changes in healthcare academia and industry, but most of these methods rely on massive labeled data. In this work, we propose a general deep learning framework which is able to boost risk prediction performance with limited EHR data. Our model takes a modified generative adversarial network namely ehrGAN, which can provide plausible labeled EHR data by mimicking real patient records, to augment the training dataset in a semi-supervised learning manner. We use this generative model together with a convolutional neural network (CNN) based prediction model to improve the onset prediction performance. Experiments on two real healthcare datasets demonstrate that our proposed framework produces realistic data samples and achieves significant improvements on classification tasks with the generated data over several stat-of-the-art baselines.

Index Terms—electronic health record; generative adversarial network; health care; deep learning;

I. INTRODUCTION

The worldwide exponential surge in volume, detail, and availability of *Electronic Health Records (EHRs)* promises to usher in the era of personalized medicine, enhancing each stage of the healthcare chain from providers to patients. This field of research and applications, commonly mentioned as *Data-Driven Healthcare (DDH)* [1], has been under rapid development and attracted many researchers and institutions to utilize state-of-the-art machine learning and statistical models on a broad set of clinical tasks which are difficult or even impossible to solve with traditional methods [2], [3]. Among these frontier models, deep learning tends to be the most exciting and promising solution to those difficult and important tasks.

Recent success and development in deep learning is revolutionizing many domains such as computer vision [4], natural language processing [5], and healthcare, with notable innovations and applicable solutions. A series of excellent work have been conducted in seek of novel deep learning solutions

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in different healthcare applications including but not limited to computational phenotyping [6], [7], risk prediction [8], [9], medical imaging analysis [10], and clinical natural language processing [11]. For EHR data, researchers attempted different types of neural networks to exploit the natural temporarily, such as recurrent neural networks (RNNs) [12], [13], deep state space model [14], and convolutional neural networks (CNNs) [15], [16]. These works have made us closest ever towards the ultimate goal of improving health quality, reducing cost, and most importantly saving lives.

While existing achievements on deep learning models for healthcare are encouraging, the peculiar properties of EHRs, such as heterogeneity, longitudinal irregularity, inherent noises, and incomplete nature, make it extremely difficult to apply most existing mature models to healthcare compared with other welldeveloped domains with clean data. Properly-designed deep neural networks have the prospect of handling these issues if equipped with massive data, but the amount of clinical data, especially with accurate labels and for rare diseases and conditions, is somewhat limited and far from most models' requirements [17]. This comes from the following reasons: The diagnosis and patient labeling process highly relies on experienced human experts and is usually very time-consuming; Getting detailed results of lab tests and other medical features, though has become more feasible with modern facilities than ever, are still quite costly; Not to mention the privacy issues and regulations which makes it even harder to collect and keep enough medical data with desired details. These unique challenges lying in healthcare domain prevent existing deep learning models from exerting their strength with enough available and high-quality labeled data.

One way to overcome the challenges arising from limited data in machine learning field is *semi-supervised learning* (SSL) [18]. Semi-supervised learning is a class of techniques that makes use of unlabeled or augmented data together with a relatively small set of labeled data to get better performance. Though some previous work utilized SSL methods on EHR data [19], most of them focus on clinical text data [20], [21], and only limited work attempt to perform semi-supervised



learning method on structured quantitative EHR data [22].

Generative model is also considered as a promising solution. As one type of SSL algorithms, it aims at learning the joint probability distribution over observations and labels from the training data, and can be further used for downstream algorithms and applications such as data modeling [23], classifier and predictor training [24], and data augmentations [25]. Though generative model approaches have been well explored for years, deep generative models haven't caught enough attentions due to its complexity and computation issues until the recent development of generative adversarial network (GAN) [26]. GAN simultaneously trains a deep generative model and a deep discriminative model, which captures the data distribution and distinguishes generated data from original data respectively, as a mini-max game. GANs have been mainly used on image, video and text data to learn useful features with better understandings [27], [28] or sample data for specific demand [29], [30]. The generators can produce images of unprecedented visual quality [31], while the discriminators learn features with rich semantics that can benefit other learning paradigms such as semi-supervised learning and transfer learning [32]. Generally speaking, existing methods include feature augmentation [33], virtual adversarial training [34], and joint training [35]. However, it is noting that all these related models are only applied to and designed for vision or natural language processing (NLP) domains. However, few GANs have been applied for generating sequential or time series EHR data, where large amount of reliable data, either from real dataset or augmentations, are in great demand for powerful predictive models. To extend existing GAN framework to EHR data is not straightforward. Moreover, to facilitate GANs with semisupervised learning for onset prediction is also difficult. These two unsolved challenges are well addressed in our proposed SSL framework.

In this paper, we investigate and propose general deep learning solutions to the challenges on high dimensional temporal EHR data with limited labels. We propose a generative model, ehrGAN, for EHR data via adversarial training, which is able to generate convincing and useful samples similar to realistic patient records in existing data. We further propose an SSL framework which achieves boosted risk prediction performance by utilizing the augmented data and representations from the proposed generative models. We conduct experiments on two real clinical tasks and demonstrate the efficacy of our generative model and prediction framework.

II. THE PROPOSED METHOD

In this section, we first introduce the deep learning predictive model in the proposed framework, then describe <code>ehrGAN</code>, a modified generative adversarial network specifically designed for EHR data, and present the data augmented semi-supervised learning schema which performs boosted onset predictions.

A. Basic Deep Prediction Model

The basic model used in the paper is a convolutional neural network (CNN) model with 1D convolutional layer over the

temporal dimension and max over-time pooling layer. The input to the model is the EHR records of patient p, which is represented as a temporal embedding matrix $x^p \in \mathbb{R}^{T_p \times M}$, where T_p , which can be different among patients, is the number of medical events in patient p's record, and M is the dimension of the learned embedding. The rows of x^p are the embedding vectors for the medical events, arranged in the order of time of occurrence. The embedding for medical events is trained by Word2vec model [36] on the same corpus of EHR data. We apply convolutional operation only over the temporal dimension but not over embedding dimension. We us a combination of filters with different lengths to capture temporal dependencies in multiple levels, and our preliminary experiments validated the performance improvement from such strategy. After the convolutional step, we apply a max-pooling operation along the temporal dimension to keep the most important features across the time. This temporal pooling converts the inputs with different temporal lengths into a fixed length output vector. Finally a fully connected soft-max layer is used to produce prediction probabilities. This CNN-based deep prediction model described above is shown to be the most competitive baseline among all other tested baselines in our experiments and serves as the basic prediction component in our proposed work.

B. ehrGAN: Modified GAN Model for EHR Data

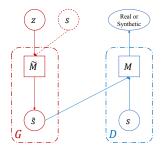
The original GAN [26] is trained by solving

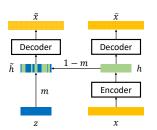
$$\min_{G} \max_{D} \underset{\boldsymbol{x} \sim p_{data}(\boldsymbol{x})}{\mathbb{E}} \left[\log D(\boldsymbol{x}) \right] + \underset{\boldsymbol{z} \sim p_{\boldsymbol{z}}(\boldsymbol{z})}{\mathbb{E}} \left[\log \left(1 - D(G(\boldsymbol{z})) \right) \right]$$

where $p_{data}(\boldsymbol{x})$ is the true data distribution; $D(\boldsymbol{x})$ is the discriminator that takes a sample as the input and outputs the probability of the sample drawing from real dataset; $G(\boldsymbol{z})$ is the generator that maps a noise variable $\boldsymbol{z} \in \mathbb{R}^d$ drawn from a given distribution $p_{\boldsymbol{z}}(\boldsymbol{z})$ back to the input space. The training procedure consists of two loops optimizing G and D iteratively. After the mini-max game reaches its Nash equilibrium [37], G defines an implicit distribution $p_g(\boldsymbol{x}) = p_{data}(\boldsymbol{x})$ that recovers the true data distribution.

Generally, both D and G are parameterized as deep neural networks. In the context of EHR data, similar to the basic prediction models, our choice of G and D falls into the family of 1D convolutional neural networks (CNNs) and 1D deconvolutional neural networks (DCNNs). The overview of the model is shown in Figure 1(a). In the following parts, we will discuss model details in terms of the design of discriminator and generator, and some specific training techniques.

- 1) Discriminator: We adopt the structure of the basic prediction model to the discriminator, due to its simplicity and excellent classification performance. We replace the top prediction layer by a single sigmoid unit to output the probability of the input data being drawn from the real dataset.
- 2) Generator: The goal of the generator in GAN is to translate a latent vector z into the synthetic sample \tilde{x} . The generator is encoded by a de-convolutional neural network with two consecutive fully connected layers, the latter of which is reshaped and followed by two de-convolutional layers to perform upsampling convolution. Empirically, this generator





(a) The structure of ehrGAN model. M and \tilde{M} are the models in discriminator (D) and generator (G). s and \tilde{s} represent real and synthetic samples.

(b) The structure of the generator in ehrGAN. \boldsymbol{z} and \boldsymbol{m} are drawn randomly. $\tilde{\boldsymbol{x}}$ is the generated synthetic sample based on the real sample \boldsymbol{x} .

Figure 1. Illustration of the proposed ehrGAN model.

is able to generate good samples. However, this version of generator can not be directly used in semi-supervised learning setting as the model is trained only to differentiate real or synthetic data instead of the classes. To solve this problem, we introduce a variational version of the generator, which also provides some new understandings of GANs.

a) Generator with variational contrastive divergence: The design of the variational generator is based on the recently proposed variational contrastive divergence (VCD) [38]. Instead of directly learning a generator distribution defined by G(z), we learn a transition distribution of the form $p(\tilde{x}|x)$ for the generated sample \tilde{x} , with $x \sim p_{data}(x)$. The marginal distribution of of the generator is then given by $p_g(\tilde{x}) = \mathbb{E}_{x \sim p_{data}(x)} p(\tilde{x}|x)$. Intuitively, the transition distribution $p(\tilde{x}|x)$ encodes a generating process. In this process, based on an example drawn from the training data distribution, a neighboring sample \tilde{x} is generated. To be more specific, the generator is equipped with encoder-decoder CNN networks. For each real sample x, we can get the representation h from encoder, and the reconstruction \bar{x} by feeding h into decoder. h can be mixed with a random noise vector z of the same dimensionality by a random binary mask vector mto obtain $\tilde{h} = m * z + (1 - m) * h$, where * represents element-wise multiplication. The synthetic sample \tilde{x} can be obtained by feeding h to the same decoder. An illustration of this VCD-based generator is shown in Figure 1(b). The generator attempts to minimize the objective as

$$\underset{\boldsymbol{x} \sim p_{data}(\boldsymbol{x})}{\mathbb{E}} \left[\rho \cdot \underset{\tilde{\boldsymbol{x}} \sim p_{g}(\tilde{\boldsymbol{x}}|\boldsymbol{x})}{\mathbb{E}} \left[-\log D(\tilde{\boldsymbol{x}}) \right] + (1 - \rho) \cdot \|\bar{\boldsymbol{x}} - \boldsymbol{x}\|_{2}^{2} \right] \quad (1)$$

where D is the discriminator function and the hyperparameter ρ controls how close the synthetic sample should be to the corresponding real sample. The usage of VCD-based <code>ehrGAN</code> brings two benefits. First, while original GANs are known to have $mode\ collapsing\$ issues, i.e., G is encouraged to generate only a few modes, <code>ehrGAN</code> eliminates $mode\ collapsing\$ issue by its design, as the diversity of the generated samples inherently approximates that of the training data. Second and more importantly, the learned transition distribution $p(\tilde{x}|x)$ contains rich structures of the data manifold around true examples x, which can help our semi-supervised learning framework obtain effective classification models.

3) Training techniques: We train the proposed ehrGAN by optimizing the generator and discriminator iteratively with stochastic gradient descent (SGD). The training procedure (shown in Algorithm 1) is similar to that of standard GANs. We take several techniques to stabilize the training of GANs similar to those in [28], [35], and relieve the training instability and sensitivity to hyper-parameters. Firstly, we switch the order of discriminator and generator training, and perform k=5 optimization steps for the generator for every one steps for the discriminators. Secondly, we add an l_2 -norm regularizer in the cost function of discriminator. Finally, batch normalization and label smoothing techniques are used.

Algorithm 1 The optimization procedure of ehrGAN

```
1: for enough iterations until convergence do
2: for k inner steps do
3: sample N noise variables \{\boldsymbol{z}^1,\dots,\boldsymbol{z}^N\}, and N binary mask vectors \{\boldsymbol{m}^1,\dots,\boldsymbol{m}^N\};
4: update generator G by one step gradient ascent of \frac{1}{N}\sum_{i=1}^N\log D(G(\boldsymbol{z}^i,\boldsymbol{m}^i))
5: end for
6: sample N training data \{\boldsymbol{x}^1,\dots,\boldsymbol{x}^N\}, N noise variables \{\boldsymbol{z}^1,\dots,\boldsymbol{z}^N\}, and N binary mask vectors \{\boldsymbol{m}^1,\dots,\boldsymbol{m}^N\};
7: update discriminator D with one step gradient descent of -\frac{1}{N}\sum_{i=1}^N\log D(\boldsymbol{x}^i) - \frac{1}{N}\sum_{i=1}^N\log \left(1 - D(G(\boldsymbol{z}^i,\boldsymbol{m}^i))\right)
8: end for
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C. Semi-supervised Learning with GANs

We next introduce our method of conducting semi-supervised learning (SSL) with a learned ehrGAN, in a way which is similar to our previous model for images [38]. The basic idea is to use the learned transition distribution to perform data augmentation. To be concrete, within the SSL setting we minimize the follow loss function:

$$\frac{1}{N} \sum_{i=1}^{N} \mathcal{L}(\boldsymbol{x}^{i}, \boldsymbol{y}^{i}) + \mu \cdot \frac{1}{N} \sum_{i=1}^{N} \mathbb{E}_{\tilde{\boldsymbol{x}}^{i} \sim p(\tilde{\boldsymbol{x}}|\boldsymbol{x}^{i})} \mathcal{L}(\tilde{\boldsymbol{x}}^{i}, \boldsymbol{y}^{i})$$
(2)

where \mathcal{L} refers to the binary crossentropy loss on each data sample, and μ leverages the ratio of the numbers of training data and augmented data from GANs. In other words, this model assumes that a well trained generator with distribution $p(\tilde{x}|x)$ should be able to generate samples that are likely to align within the same class of x, which can in turn provide valuable information to the classifier as additional training data. This method is called SSL-GAN (Semi-supervised learning with a learned ehrGAN) in this paper.

III. EXPERIMENTAL RESULTS

In this section we apply our models to two real clinical datasets extracted from heart failure and diabetes cohorts. It is a particularly interesting to investigate how well GANs can generate EHRs samples as the real ones. Also, understanding how the proposed method can boost the performance of onset prediction is crucial for many healthcare applications. We start this section by introducing the datasets and experimental settings, and provide the evaluation analysis, followed by the discussions on the selections of parameters.

A. Datasets and Settings

The datasets came from a real-world longitudinal Electronic Health Record database of 218,680 patients and 14,969,489 observations of 14,690 unique medical events, between the year 2011 to 2015 from a health insurance company. In these datasets, a set of diseases related ICD-9 codes were recorded to indicate medical conditions as well as drug prescriptions. We identify two following cohorts and predict whether a patient is from case or control group as a binary classification task. The labels of both case and control groups are identified by domain experts according to ICD-9 codes.

- Congestive heart failure (*Heart Failure*), with 3,357 patients in case group and 6,714 patients in control group;
- Diabetes (*Diabetes*), with 2,248 patients in case group and 4,496 patients in control group.

We import ICD-9 diagnosis and medications as the input features, eliminate those which show less than 5 times in this dataset, and get 8,627 unique medical features. We segmented the time dimension into disjoint 90-day windows and combined all the observations within each window. We split datasets into training, validation and test with ratio 7:1:2, and limit the length of each record sequence between 50 and 250 and form it to the embedding matrix. All sequences are 0-padded to match the longest sequence. The embedding is trained by Word2vec [36] on the entire dataset with dimension of 200. The ehrGAN is trained on only the training subset. For the CNN discriminator, we employ filters of sizes $\{3,4,5\}$ with 100 feature maps. For the generator, the dimension of the latent variable z is 100. It is first projected and reshaped by the generator and up-sampled by two one-dimensional CNN layers with filers size 100 and 3. The output of the generator is an embedding matrix with size 200×150 . These hyperparameters are selected based on preliminary experimental results. To generate samples with different length, we paddle a special embedding mark at the end of each training record. The masks m in the VCD-based generator is uniformly sampled with probability 0.5. The Adam algorithm [39] with learning rate 0.001 for both discriminator and generator is utilized for optimization. Gradients are clipped if the norm of the parameter vector exceeds 5 [5]. After we get the generated data, we can map it into EHR record by finding the nearest-neighbor with cosine distance for each feature. The selection of optimal values for hyper-parameters μ and ρ will be discussed later in Section III-E.

Table I PREDICTION PERFORMANCE COMPARISON.

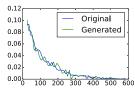
	Heart l	Failure	Diabetes		
	Accuracy	AUROC	Accuracy	AUROC	
CNN	0.8630	0.9329	0.9644	0.9789	
GRU	0.8578	0.9129	0.9304	0.9659	
LSTM	0.8511	0.9103	0.9448	0.9553	
LR	0.8494	0.9052	0.9066	0.9681	
SVM	0.8443	0.9017	0.8944	0.9462	
RF	0.8571	0.9225	0.9476	0.9658	

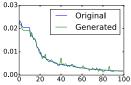
B. Risk Prediction Comparison on Basic Models

First, we show the performance of our basic predict model (*CNN*), which explores the CNN model with pre-trained medical feature embedding and is a strong baseline even before boosted. We compare it with logistic regression (*LR*), linear support vector machine (*SVM*), random forest (*RF*) and two other deep models, recurrent neural network models using gated recurrent units (*GRU* [40]) and long short-term memory (*LSTM* [13]). For *LR*, *SVM* and *RF*, we use the same setting as mentioned in previous work [16]. We follow the similar settings from existing work [13] for *GRU* and *LSTM*.

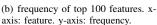
Table I shows the classification accuracy and AUROC (area under receiver operating characteristic curve) of all basic baseline models on the two prediction tasks. *CNN* is among the best methods in *Heart Failure* task and significantly outperforms baselines in *Diabetes* task. The performance improvement mainly comes from the learned embeddings in heart failure task, and from CNN model structures in diabetes task. The other two deep models *GRU* and *LSTM* work well but can not beat *CNN*.

C. Analysis of Generated Data





(a) Data length distributions. x-axis: data length; y-axis: probability.







(c) Co-occurrence frequency of top 20 diagnosis features. Green: generated data; Blue: original data.

Figure 2. Data analysis on Heart Failure dataset.

Before testing our semi-supervised prediction models with augmented data, we need to inspect whether the generated data from ehrGAN are able to simulate original data well enough, especially for the patient records in the case cohorts. Having the generated data similar to original one is an important precondition to improve our model performance instead of hurting it. We compared the length and features of the original data (\mathcal{D}_o) and generated data \mathcal{D}_g for the two case groups. We only show the results on Heart Failure. The results on Diabetes are similar and can be found in the full version of this paper. Figure 2(a) shows that the generated dataset has similar length distributions to original dataset. Figure 2(b) shows that the generated dataset keeps similar frequencies of top 100 features. Comorbidities (cooccurrences) in patient records are quite useful in clinical prediction tasks. We select 20 most frequent diagnosis features from these two case cohorts, and

show the comorbidity heatmaps in Figure 2(c). We can find that both the feature frequencies and the comorbidity clusters are well simulated in our generated dataset. The list of top 10 diagnosis features is listed in Table II. Our generated model is able to capture the occurrence patterns from case cohorts and keep those patterns very similar to those in the corresponding original datasets. These analyses not only verify the quality of our generated data, but also help us get better understandings on patterns in cohorts for different tasks.

Table II
TOP 10 MOST FREQUENT ICD-9 DIAGNOSIS CODES OF HEART FAILURE
COHORT GROUP IN GENERATED DATA.

Rank in \mathcal{D}_g	Rank in \mathcal{D}_o	ICD-9 Code	Diagnosis Descriptions
1	2	250.0	Diabetes mellitus, unspecified
2	1	401.1	Hypertension, benign
3	3	427.31	Atrial fibrillation
4	4	401.9	Hypertension, unspecified
5	5	272.4	Other and unspecified hyperlipidemia
6	6	496	Chronic airway obstruction
7	14	585.6	End stage renal disease
8	7	272.0	Pure hypercholesterolemia
9	9	285.9	Anemia, unspecified
10	8	244.9	Hypothyroidism, unspecified

D. Evaluation of the Boosted Model

To evaluate the performance of the boosted model with semi-supervised learning setting, we conduct extensive experiments on the following six approaches. *CNN-BASIC*: The basic model described in Section II-A, trained only on the training subset; *CNN-FULL*: The basic model trained with the same amount of labeled data as *SSL-GAN*; *CNN-RAND*: The basic model trained with the same amount of data as *SSL-GAN* with random labels for additional data and true labels for training subset; *SSL-SMIR*: Squared-loss mutual information regularization [41]; *LGC*: Semi-supervised learning approach with local and global consistency [42]; *SSL-GAN*: The proposed method with ehrGAN based data augmentation.

It is notable that SSL-SMIR and SSL-LGC are strong and robust SSL baselines. CNN-FULL, SSL-SMIR, and SSL-LGC

Table III
PERFORMANCE COMPASSION OF DIFFERENT PREDICTION MODELS.

	HF	50	Dia50		
	Accuracy	AUROC	Accuracy	AUROC	
CNN-BASIC	0.8096	0.8784	0.8990	0.9156	
CNN-RAND	0.7418	0.7856	0.7734	0.8011	
CNN-FULL	0.8631	0.9212	0.9335	0.9528	
SSL-SMIR	0.8207	0.8842	0.9089	0.9197	
SSL-LGC	0.8119	0.8767	0.8844	0.9102	
SSL-GAN	0.8574	0.9075	0.9135	0.9354	

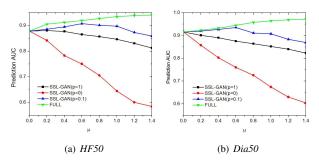


Figure 3. AUROC score comparison with different values of μ .

are trained with additional samples from a held-off subset. The parameters setting of SSL-SMIR and SSL-LGC follows those in the original papers [42], [41] and bag of words feature are used. We choose the values of ρ and μ in SSL-GAN with best performance by Section III-E. We summarize the classification performance in Table III in different settings with different amounts of labeled data. Here, we show the results on HF50 and Dia50 (50% of the training set is used), and leave results on HF67 and Dia67 (2/3 of the training set is used) in our full paper. First of all, our model consistently beats CNN-BASIC and CNN-RAND. Due to the messed up label information, the performance of CNN-RAND is even worse than CNN-BASIC. Second, compared with the CNN-FULL method, our model can also achieve comparable results. The two standard SSL methods SSL-SMIR and SSL-LGC do not perform very well, only achieving similar performances as CNN-BASIC, and our method easily beats them. Overall, these evaluations show the strong boosting power of the proposed SSL-GAN model.

 $\label{eq:table_IV} \textbf{AUROC} \ \textbf{SCORE} \ \textbf{COMPARISON} \ \textbf{WITH} \ \textbf{DIFFERENT} \ \rho.$

Setting	$\mu = 0$	$\rho = 0$	$\rho=0.001$	$\rho = 0.01$	$\rho = 0.1$	$\rho = 0.2$	$\rho = 1$
HF50	0.8784	0.8654	0.8823	0.8911	0.9075	0.8876	0.7503
Dia50	0.9156	0.8754	0.9188	0.9237	0.9354	0.9025	0.7603

E. Selections of Parameters

- 1) The effectiveness of ρ : In this part, we discuss how the selection of ρ in Equation 1 affects the performance. We fix other parameters and vary ρ from 0 to 1, and report the AUROC score with different settings in Table IV. We see that on both datasets, with a properly chosen ρ the generator is able to provide good generations to improve learning. $\rho=0.1$ is an optimal selection for the model (results of $\rho>0.2$ are no better than $\rho=0.2$ and thus omitted here). On the other hand, with $\rho=0$, which corresponds to sample from an autoencoder, hurts performance. $\rho=1$ completely messes up training as the generated samples are not guaranteed to have the same label as the samples conditioned on. This shows that the transition distribution is able to generate samples that are sufficiently different from training samples to boost the performance.
- 2) The effectiveness of μ : How to optimally utilize the augmented data from GANs to support the supervised learning is an important problem. In our task, this is controlled by the parameter μ in Equation 2, which leverages the ratio of

labeled data and augmented data. Generally, including more augmented data will help while too many augmented data may even hurt the performance. With a fixed value of ρ , we vary μ from 0.2 to 1.4 and test the prediction performance on two datasets. We also include the setting with fully labeled data (FULL), and μ represents the number of real labeled data used instead of from GANs in this setting. The prediction AUROC scores of different methods are shown in Figure 3. It is obvious for the method with fully labeled data that the prediction performance continues improving with μ increased. For SSL-GAN, when $\rho = 0.1$ (the optimal setting), we can see it achieved the best performance when $\mu = 0.6$. After that point, the performance decreased a little, which indicates that more augmented data can not help further. For the setting with $\rho = 1$ and $\rho = 0$, the prediction power continues falling as including more augmented data is harmful for both cases. Similar trend are also observed under the measure of accuracy.

IV. CONCLUSION

In this paper, we focus on exploiting deep learning technique and its applications in healthcare. We first present ehrGAN, a generation model via adversarial training, and discuss several techniques for learning such a model for EHR data. We demonstrate that the proposed model can produce realistic data samples by mimicking the input real data, and the learned latent representation space can continuously encode plausible samples. To boost risk prediction performance, we utilize the learned model to perform data augmentation by semi-supervised learning. Experimental results on two datasets show that the proposed model improves the generalization power and the prediction performance compared with strong baselines.

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