Open-Set Recognition with Gaussian Mixture Variational Autoencoders

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Abstract

In inference, open-set classification is to either classify a sample into a known class from training or reject it as an unknown class. Existing deep open-set classifiers train explicit closed-set classifiers, in some cases disjointly utilizing reconstruction, which we find dilutes the latent representation's ability to distinguish unknown classes. In contrast, we train our model to cooperatively learn reconstruction and perform class-based clustering in the latent space. With this, our Gaussian mixture variational autoencoder (GMVAE) achieves more accurate and robust open-set classification results, with an average F1 increase of 0.26, through extensive experiments aided by analytical results.

1 Introduction

Until recently, nearly all classification algorithms have been designed for closed-set evaluation. This means that all testing classes are seen in training. However, real-world applications necessitate open-set evaluation where unknown classes, not seen in training, appear during testing. For instance, computer vision systems in self-driving cars must classify and navigate around many different objects. Given the countless number of such possible objects, it is infeasible for all classes to be seen in training (Sünderhauf et al. 2018). Open-set recognition addresses this generalization of the classification task.

While there are several facets of open-set learning, in this paper we focus on training from C known classes for $(C + 1)$-class classification. This $(C + 1)$-th class catches all unknown test samples not belonging to any of the known classes. The training and validation data have no unseen classes from class $C + 1$. To this end, we present a novel supervised, Gaussian mixture variational autoencoder (GMVAE). The bottleneck latent layer simultaneously learns reconstruction and performs class-based clustering (preserving closed-set classification ability). This allows the latent representation to capture complementary structure and classifier information. Furthermore, the latent layer has the explicit capability to form multiple subclusters per class. This challenges the implicit assumption made by many classification methods that a class's embedding is a convex set and

thus is best represented by a single centroid (Bendale and Boult 2016; Hassen and Chan 2020; Lee et al. 2018; Yoshihashi et al. 2019). This provides further flexibility in capturing complementary structure and classifier information.

Our contributions are as follows. In §3, we derive GMVAE to learn the embedding and amend its objective function to make open-set recognition more amenable. We also present a new and simple open-set classification algorithm that utilizes an "uncertainty" threshold on the learned embedding. Following in §4, we present analytical results regarding the number of subclusters and the resulting heuristic procedure for identifying the appropriate number of subclusters in each class. Finally in §5, we conduct open-set classification experiments on three standard datasets. Our findings from experiments are two-fold. First, GMVAE outperforms a state-of-the-art classification-reconstruction-based, deep open-set classifier both in terms of accuracy and robustness to an increasing number of unknown classes. Second, the use of extreme value theory (EVT) to infer class-belongingness (Bendale and Boult 2016; Yoshihashi et al. 2019) may be ill-suited in this classification-reconstruction open-set framework as we find that ours and another simple algorithm consistently beat it.

2 Related Work

While closed-set classification has been well-studied, open-set recognition has been gaining more attention in recent years. Outlier or novelty detection is a precursor but, unlike the problem studied herein, is not generally concerned with distinguishing between the known classes (Geng, Huang, and Chen 2020; Zhou and Paffenroth 2017). Such methods may also rely on the use of synthetic, outlier training datasets (Hendrycks, Mazeika, and Dietterich 2019) whereas we focus on training with only known classes. Earlier works that study $(C + 1)$-class classification utilize, for example, SVM scores (Scheirer et al. 2013; Jain, Scheirer, and Boult 2014) or sparse representation (Zhang and Patel 2017) to fit EVT-based densities to predict classes. The use of deep networks in open-set recognition appears even more recently in studies such as Bendale and Boult (2016) and Yoshihashi et al. (2019). Both use similar procedures of fitting EVT-based densities to the distances between a class's embedding and its centroid to approximate probability of class inclusion. Finally, Oza and Patel (2019) also use a class conditioned

autoencoder for open-set identification but instead apply an EVT-based threshold derived from the training data's reconstruction error.

Herein, our experimental results are benchmarked against the Classification-Reconstruction learning for Open-Set Recognition (CROSR) method (Yoshihashi et al. 2019). We chose this particular benchmark as it achieves state-of-the-art open-set classification accuracies and it relies on the same framework of dual reconstruction-classification learning with a latent space distance-based threshold. In this specific open-set realm, GMVAE reveals the pitfall of using a closed-set, softmax classifier to cluster known classes and showcases the reduction in open-space risk (Scheirer et al. 2013) from utilizing multiple subclusters per class.

We next summarize CROSR. The latent representation is a concatenation $[y, z]$ where y is the activation vector of a closed-set, softmax classifier and z is the reconstructive latent representation. To learn an effective y and z concurrently, Yoshihashi et al. (2019) introduced Deep Hierarchical Reconstruction Nets (DHRNets). Conceptually, the DHRNet architecture is a deep classifier f with autoencoder networks $h_l, \widetilde{h}_l$ appended at the internal layers x_l. Thus, bottleneck representations can be extracted from multi-stage features of the classifier. The autoencoders' reconstructions then form a reverse network to reconstruct the original input. Mathematically, the main-body network $f(x) = (y, z)$ is comprised of

$$x_{l+1} = f_l(x_l) \quad l\text{-th layer of the DHRNet classifer}$$
$$z_l = h_l(x_l) \quad \text{encoder network for } l\text{-th layer}$$
$$\widetilde{x}_l = g_l(\widetilde{x}_{l+1} + \widetilde{h}_l(z_l)) \quad \text{decoder network } \widetilde{h}_l$$
$$\text{and reconstruction network } g_l \text{ for } l\text{-th layer}$$

where networks are a series of convolutions and up or downsampling layers. For training, Yoshihashi et al. (2019) minimizes the sum of the cross-entropy classification error and the L_2 reconstruction errors.

With latent representation $[y, z]$ in hand, CROSR applies EVT by fitting a Weibull distribution to the hypersphere defined by $d(x, C_i) = ||[y, z] - \mu_i||_2$ where μ_i is the respective mean within class C_i. A proxy for probability of class inclusion is then given by $\mathbb{P}(x \in C_i) = 1 - \text{WeibullCDF}(d(x, C_i); \rho_i) = \exp\left\{-\left(\frac{d(x, C_i)}{\eta_i}\right)^{m_i}\right\}$ and thresholding is then used to classify a sample as "unknown." Here m_i and η_i are parameters of the distribution fitted from class C_i's training data.

In contrast to DHRNets, Gaussian mixture variational autoencoders (Dilokthanakul et al. 2016) are deep generative models which estimate the density of training data under assumptions on its latent prior. This could lead to more complex latent structures than in classification-based models, especially with the inclusion of multiple subclusters per class. However, inference in this unsupervised setting is challenging, especially with open-set recognition. We address this by extending this deep generative model to supervised learning including capturing subclusters within classes.

3   Gaussian Mixture Variational Autoencoders

In this section we present our complete, novel procedure for open-set recognition. It follows the same two phases as previous works: first, learn a latent representation to (sub)cluster known classes, and second, apply an open-set classification algorithm on that embedding. Our GMVAE model is an extension of the Gaussian mixture variational autoencoder presented in Dilokthanakul et al. (2016) and explained next.

Variational autoencoders (VAEs) assume data is generated from a uni-modal Gaussian prior. In Dilokthanakul et al. (2016), the authors instead choose a mixture of Gaussians as an intuitive extension. In order to maintain standard backpropagation via the reparametrisation trick, the standard VAE architecture was altered. The generative model, factorizing as $p_{\beta,\theta}(x, z, w, v) = p(w)p(v)p_\beta(z|w, v)p_\theta(x|z)$, generates a sample x from the latent variables z, w, and v with the following process

$$w \sim \mathcal{N}(0, I), \quad v \sim \text{Mult}(\pi)$$
$$(z|w, v) \sim \prod_{k=1}^{K} \mathcal{N}\left(\mu_k(w; \beta), \text{diag}\left(\sigma_k^2(w; \beta)\right)\right)^{v_k}$$
$$(x|z) \sim \mathcal{N}\left(\mu(z; \theta), \text{diag}\left(\sigma^2(z; \theta)\right)\right) \quad \text{or} \quad \mathcal{B}\left(\mu(z; \theta)\right)$$

where K is the user-defined number of mixture components and $\mu_k(\cdot; \beta)$, $\sigma_k^2(\cdot; \beta)$, $\mu(\cdot; \theta)$, and $\sigma^2(\cdot; \theta)$ are neural networks parametrized by β and θ, respectively. The recognition model is then factorized as $q(z, w, v|x) = q_{\phi_z}(z|x)q_{\phi_w}(w|x)p_\beta(v|z, w)$ where ϕ_z and ϕ_w parametrize neural networks that output means and diagonal covariances of the Gaussian posterior variational distributions. Using Bayes' rule, the v-posterior term $p_\beta(v|z, w)$ can be written in terms of factors of the generative model. To train, the log-evidence lower bound (ELBO) $\mathbb{E}_{q(z,w,v|x)}[p_{\beta,\theta}(x, z, w, v)/q(z, w, v|x)]$ is maximized. In §3.1 and 3.2, we present the derivation and differences of our GMVAE. Finally we introduce our new open-set classification algorithm that utilizes an "uncertainty" threshold in §3.3.

3.1   Gaussian Mixture Variational Autoencoders with Multiple Subclusters Per Class

Our GMVAE model nontrivially extends the unsupervised learning framework of Dilokthanakul et al. (2016) to essentially a Gaussian mixture prior for each class. For notation, there are C known classes with each class composed of K_c subclusters where $c = 1, 2, ..., C$. The samples $x \in \mathbb{R}^d$ and labels $y \in \mathbb{R}^C$ as one-hot vectors comprise the labeled, known data set $(x, y) \in \mathcal{X}$. The GMVAE's generative process $p_{\beta,\theta}(x, v, w, z|y) = p_\theta(x|z)p_\beta(z|w, y, v)p(w)p(v|y)$ is conditioned on class and given by

$$w \sim \mathcal{N}(0, I), \quad (v|y) \in \mathbb{R}^{K_c} \sim \text{Mult}(\pi(y))$$

$$(z|w, y, v) \sim \prod_{c=1}^{C} \prod_{k=1}^{K_c} \mathcal{N}\left(\mu_{ck}(w; \beta), \text{diag}\left(\sigma_{ck}^2(w; \beta)\right)\right)^{y_c \cdot v_k}$$

$$(x|z) \sim \mathcal{B}(\mu(z; \theta)).$$

It is common to take $\pi(y)$ to simply be uniform for each class. The recognition model is factorized as $q_\phi(v, w, z|x, y) = p_\beta(v|z, w, y)q_{\phi_w}(w|x, y)q_{\phi_z}(z|x)$ where $\phi = (\phi_x, \phi_w)$. We parametrize variational factors with networks ϕ that output mean and diagonal covariance of variational distributions and specify their form to be Gaussian posteriors:

$$(z|x) \sim \mathcal{N}\left(\mu(x; \phi_z), \text{diag}\left(\sigma^2(x; \phi_z)\right)\right)$$
$$(w|x, y) \sim \mathcal{N}\left(\mu(x, y; \phi_w), \text{diag}\left(\sigma^2(x, y; \phi_w)\right)\right).$$

There is a p_β factor in the q_ϕ factorization because the p_β factor can be written in terms of generative factors, lowering the number of trainable parameters. Using Bayes', we can rewrite $p_\beta(v|z, w, y)$ as

$$p_\beta(v|z, w, y) = \frac{p_\beta(z|w, y, v)p(v|y)}{\sum_{v'} p_\beta(z|w, y, v')p(v'|y)}. \quad (1)$$

The details are provided in the technical appendix. Another benefit is that $p_\beta(v|z, w, y)$ can be computed for all v with simply one forward pass. The GMVAE's ELBO is then given by

$$\mathcal{L}(K) = \mathbb{E}_{q_\phi(v, w, z|x, y)}\left[\log \frac{p_{\beta, \theta}(x, v, w, z|y)}{q_\phi(v, w, z|x, y)}\right]$$

$$= \mathbb{E}_{q_{\phi_z}(z|x)}\left[\log p_\theta(x|z)\right] \quad \text{(reconstruction)}$$

$$- \mathbb{E}_{q_{\phi_w}(w|x, y)q_{\phi_z}(z|x)}\left[\log q_{\phi_z}(z|x) - \sum_{j=1}^{K_c} p_\beta(v = j|z, w, y) \log p_\beta(z|w, y, v = j)\right]$$

$$\text{(latent covering)}$$

$$- KL(q_{\phi_w}(w|x, y)||p(w)) \quad \text{(w-prior)}$$
$$- \mathbb{E}_{q_{\phi_w}(w|x, y)q_{\phi_z}(z|x)}\left[KL(p_\beta(v|z, w, y)||p(v|y))\right]$$
$$\text{(subcluster v-prior).}$$

Since $K = (K_1, K_2, ..., K_C)$ is user-defined, the ELBO dependence on K is made explicit and used later in the analyses. The reconstruction term promotes a latent representation meaningful to reconstruct the samples. The latent covering term attempts to subcluster the latent representation based on classes. The w-prior and subcluster v-prior terms drive those posteriors closer to their respective priors.

3.2 Modification of the ELBO: Removing v-Prior

In this subsection, we propose removing the v-prior term from the original ELBO to make GM-VAE more amenable to open-set recognition for two reasons. First, minimizing the v-prior term $\mathbb{E}_{q_{\phi_w}(w|x, y)q_{\phi_z}(z|x)}\left[KL(p_\beta(v|z, w, y)||p(v|y))\right]$ is in direct conflict with the goal of distinct subclustering within a class. Our goal is to create disjoint subclusters in a class's latent representation so as to further provide reconstruction more flexibility and alleviate the assumption that a class's embedding is a convex set. However, notice that the v-prior term is minimized when $p_\beta(v|z, w, y) = p(v|y)$ for every z, w, and y. Combined with (1) and a uniform $p(v|y)$, this in turn implies that $p_\beta(z|w, y, v = i) = p_\beta(z|w, y, v = j)$ for every w, y, i, and j. Equivalent generative model distributions leads to mode collapse in the latent subclusters due to the maximization of the latent covering term. Put differently, the v-prior term discourages one-hot subcluster v posteriors. However, this is exactly what is needed to robustly identify subclusters.

Second, as proven in Proposition 2 in §4, without the v-prior term the optimal GMVAE loss for $C = 1$ is non-increasing with respect to K. This is an analytical result which provides a heuristic procedure for identifying the appropriate number of subclusters K_c to use for each class. Given these two reasons, for all the experiments in §5, we used the following modified ELBO:

$$\mathcal{L}_{\text{no } v\text{-prior}}(K) = \mathbb{E}_{q_{\phi_z}(z|x)}\left[\log p_\theta(x|z)\right]$$
$$- KL(q_{\phi_w}(w|x, y)||p(w))$$
$$- \mathbb{E}_{q_{\phi_w}(w|x, y)q_{\phi_z}(z|x)}\left[\log q_{\phi_z}(z|x) - \sum_{j=1}^{K_c} p_\beta(v = j|z, w, y) \log p_\beta(z|w, y, v = j)\right].$$

In a sense, it is as if we do not impose a prior on the subcluster distributions. While we could have also negated the v-prior term, simply removing it actually yields the best experimental results.

3.3 Open-Set Classification Algorithms

With recent literature in open-set recognition, it has nearly become universal to model class-belongingness by fitting a Weibull distribution to the tail-end, inlier distances between a class's latent representations and its centroid (Bendale and Boult 2016; Hassen and Chan 2020; Yoshihashi et al. 2019). Indeed, the benchmark method CROSR (Yoshihashi et al. 2019) achieves state-of-the-art accuracies through this EVT framework. However, our experiments demonstrate that two much simpler algorithms can significantly outperform CROSR's EVT-based classification algorithm. While fitting an EVT distribution to the inlier distances may be an effective way to model a decision boundary, we believe it is inherently at odds with distances related to softmax classifiers. EVT makes use of tail-end data and thus is robust

to underestimating probability of class inclusion for positive samples far away from its class's centroid. However, this procedure may render inaccurate predictions with embeddings that do not optimize for low intra-spread within each known class. For instance, CROSR's embedding is composed of the closed-set, softmax classifier's activation vector; this encourages elements of that vector to tend towards positive and negative infinity. This gives rise to known embeddings being systematically far away from its class's centroid. Accordingly, we have empirically observed the expected effect where the CROSR'S EVT procedure over-recognizes unknown samples as known.

Next we present the two simple open-set classification algorithms we implemented. While GMVAE outputs a Gaussian distribution in latent space, we simply choose the mean $\mu(x; \phi_z)$ as the effective latent representation. Algorithm 1 is derived from the so-called outlier score from Hassen and Chan (2020) but is most aptly described as nearest centroid thresholding on distance to the nearest centroid. This algorithm is modified to incorporate multiple subclusters per class.

Algorithm 1: Nearest centroid thresholding on distance to the nearest centroid

Input: Training samples $\mathcal{X}_c$ for each known class $c = 1, 2, ..., C$ and test sample $\widehat{x}$

1. For each class c, compute K_c centroids of $\mu(\mathcal{X}_c; \phi_z)$ using k-means clustering. Denote centroid $\overline{z}_{ck}$ as k-th centroid of class c.
2. Let $(c^*, k^*) = \arg\min_{c,k} ||\mu(\widehat{x}; \phi_z) - \overline{z}_{ck}||_2$ and $d = \min_{c,k} ||\mu(\widehat{x}; \phi_z) - \overline{z}_{ck}||_2$
3. If $d < \tau$, predict class as c^*; else, predict class as unknown $C + 1$

Experimental results show that thresholding on distance to the nearest centroid more robustly fits a hypersphere decision boundary around the respective centroid. However, a similar shortcoming shared with CROSR's EVT method is that distance is a rotationally symmetric measure. It does not include any sense of orientation. We stand to reason that in any nearest centroid-based algorithm, the open space between centroids poses the most risk from an open-set classification standpoint. This leads into the second algorithm which utilizes a novel threshold on an "uncertainty" quantity U. We define U as the ratio between the distance to the nearest centroid to the average distance to all other centroids. At its base, this ratio captures how similar a sample is with respect to the known classes. So if $U = 1$, the test sample's latent representation is equidistant from all centroids which can be interpreted as unclassifiable. If $U = 0$, the test sample's latent representation is exactly a centroid meaning there is no ambiguity in classification. In this way, Algorithm 2 includes a notion of orientation between centroids as U penalizes the open space directly between centroids more heavily. This is reminiscent of the nearest neighbors distance ratio of Mendes Júnior et al. (2017).

Algorithm 2: Nearest centroid thresholding on uncertainty U

Input: Training samples $\mathcal{X}_c$ for each known class $c = 1, 2, ..., C$ and test sample $\widehat{x}$

1. For each class c, compute K_c centroids of $\mu(\mathcal{X}_c; \phi_z)$ using k-means clustering. Denote centroid $\overline{z}_{ck}$ as k-th centroid of class c.
2. Let $(c^*, k^*) = \arg\min_{c,k} ||\mu(\widehat{x}; \phi_z) - \overline{z}_{ck}||_2$, $N = \sum_{c=1}^C K_c$, and

$$U = \frac{\min_{c,k} ||\mu(\widehat{x}; \phi_z) - \overline{z}_{ck}||_2}{\frac{1}{N-1} \sum_{(c,k) \neq (c^*, k^*)} ||\mu(\widehat{x}; \phi_z) - \overline{z}_{ck}||_2}$$

3. If $U < \tau$, predict class as c^*; else, predict class as unknown $C + 1$

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4 Identifying the Number of Subclusters in Each Class

Since the number of subclusters in each class is user-defined, identifying the appropriate number is critical for model usage. A natural procedure that immediately arises is to iteratively apply GMVAE to each class's data alone for an increasing number of subclusters K_c. Given the reconstruction and clustering objectives, the empirical model loss terms should naturally inform us of the optimal number of subclusters. This is akin to increasing k in k-means clustering and studying the resulting inertia plot. To this end, in this section we first present analytical results regarding the effect of $K = K_1$ on the optimal $C = 1$ (single class), original and modified GMVAE losses. In particular, we show monotonicity of the optimal GMVAE losses with respect to $K = K_1$. This then provides a foundation for our heuristic procedure for identifying the ideal number of subclusters in each class.

With two unrestrictive neural network assumptions, we are able to prove two propositions regarding the effect of K on the optimal original and modified GMVAE losses. The assumptions and proofs can be found in the technical appendix. The first proposition demonstrates that when there truly is only one subcluster within a class, and we know its distribution, then the optimal original loss is constant with respect to K. Since $C = 1$, we write x instead of (x, y).

Proposition 1. *Let us assume that $x \in \mathcal{X}$ is distributed as $x \sim p_{data} = \mathcal{B}(\mu_x)$, $C = 1$, and Assumption 1 holds. Then the optimal original GMVAE loss is constant with respect to K. In fact, we have that $\min -\mathbb{E}_{\mathcal{X}}[\mathcal{L}(K)] = -\mathbb{E}_{\mathcal{X}}[\log p_{data}]$ for every $K \geq 1$ and a globally optimal solution reads*

$$\mu(x; \phi_z^*) = \mu_{c=1,k}(w; \beta^*) = \mu_z$$
$$\sigma^2(x; \phi_z^*) = \sigma_{c=1,k}^2(w; \beta^*) = \sigma_z^2$$
$$\mu(x, y; \phi_w^*) = \vec{0}, \quad \sigma^2(x, y; \phi_w^*) = \vec{1}, \quad \mu(z; \theta^*) = \mu_x$$

for any constant vectors μ_z, σ_z.

The second proposition makes no data assumptions and shows that the optimal modified loss with the v-prior removed is non-increasing with respect to K.

Proposition 2. *Let us assume $C = 1$ and Assumptions 1 and 2 hold. We have* $\min\left\{-\mathbb{E}_{\mathcal{X}}[\mathcal{L}_{no\ v\text{-}prior}(K; \phi_z, \phi_w, \beta, \theta)]\right\} \geq \min\left\{-\mathbb{E}_{\mathcal{X}}[\mathcal{L}_{no\ v\text{-}prior}(K+1; \phi_z, \phi_w, \beta, \theta)]\right\}$ *for all $K \geq 1$.*

These proofs do not inform us on the transient dynamics of training nor even reaching the global optimum. As such, in the following experimental results section, we apply these propositions in practice by comparing the latent covering loss given reconstruction loss for each $K \geq 1$. This answers: How well does K subclusters "cover" the embedding for a given reconstruction level? When the latent covering loss's decreases begin to diminish (the propositions validate this expected monotonicity), then it is an indication that additional subclusters are only marginally beneficial and perhaps should not be included. It is worth noting from these propositions that there is no theoretical harm in over-specifying the number of subclusters K in each class. However, the user should be aware of the balance between computational difficulty and meaningful subclusters (in terms of reconstruction structure).

5 Experimental Results

The experimental results demonstrate several findings. First, EVT may not be appropriate in conjunction with closed-set, softmax classifiers as simple nearest centroid procedures consistently beat it. Second, even without the added benefit of subclustering, GMVAE for $K = \vec{1}$ often leads to a latent representation more amenable for open-set recognition compared to CROSR. Finally, subclustering within classes represents a means of bolstering dual supervised-reconstruction embeddings.

Each dataset has the following composition. The training data has only labeled samples from the C known classes. The validation set also only has samples from the same C classes. The validation set is used to determine the threshold τ. Finally, the test set has samples from the C known classes and samples from additional Q unknown classes, which are all treated as class $C + 1$.

For each of the experiments below, we perform an ablation study. Four combinations of model and classification algorithms were applied: (i) CROSR with CROSR's EVT (CROSR+EVT), (ii) CROSR with Algorithm 1 (CROSR+NC-D), (iii) GMVAE with Algorithm 1 (GMVAE+NC-D), and (iv) GMVAE with Algorithm 2 (GMVAE+NC-U). CROSR+NC-D and GMVAE+NC-D are meant to directly compare the two latent representations' amenability to open-set recognition. We did not study CROSR with Algorithm 2 because our "uncertainty" measure is really a proxy for confidence and it has been shown that it is erroneous to equate softmax classifiers with confidence (Nguyen, Yosinski, and Clune 2015). Correctly adapting "uncertainty" to CROSR is outside of this paper's scope. For each combination, we calculate the macro-averaged F1 scores (the threshold τ is algorithmically picked based on the validation set) for an increasing number Q of unknown classes (and samples). The first two experiments are for $K = \vec{1}$ and in the last two, we manufacture classes with multiple subclusters to apply $K = (2, 2)$.

We optimize over the training set using Adam until the

loss, evaluated on the known validation set, plateaus. For the MNIST and Fashion MNIST datasets (grayscale images), the reconstruction distribution used was the unnormalized, continuous Bernoulli distribution. For the CIFAR-10 dataset (RGB images), a truncated $[0, 1]$ Gaussian models the reconstruction. The latent space dimension of z equals 10, 50, 5, and 20 for the four experiments. A table of GMVAE network architectures for each experiment can be found in the technical appendix. We will publish our code upon acceptance of this paper.

5.1 Fashion MNIST Withholding 4 Classes

The six known classes are t-shirts/tops, trousers, pullovers, dresses, coats, and shirts, while the four unknown classes are sandals, sneakers, ankle boots, and bags. Fashion MNIST's standard training set is randomly split into the validation set (6,000 samples of known classes) and training set (30,000 samples). Fashion MNIST's standard testing set (10,000 samples) is kept the same. We use the same CROSR network architecture as Yoshihashi et al. (2019) for their MNIST experiment.

Known validation F1 scores versus τ are plotted in Figure 1 for CROSR+NC-D, GMVAE($K = \vec{1}$)+NC-D, and GMVAE($K = \vec{1}$)+NC-U. For the purposes of comparing the distance-based F1 scores, the smallest τ such that all validation samples are classified as "unknown" $C + 1$ is standardized to 1. The procedure of Yoshihashi et al. (2019) is followed and a threshold of 0.5 is used for all CROSR+EVT experiments. For the other three model and classification algorithm combinations, we have empirically observed that a consistently good threshold τ to pick is where the known validation F1 curve saturates or plateaus (plotted with dashed lines). This can be thought of as increasing the d or U hypersphere surrounding each class's centroid until diminishing classification accuracy returns. Any larger τ can be thought of as overfitting the known validation set and runs the risk of underclassifying "unknown" samples. Let $\widetilde{\tau} = \min\{\tau : F1'(\tau) \geq \epsilon_1\}$, then we define this saturation as $\min\{\tau : \tau > \widetilde{\tau} \quad \text{and} \quad F1'(\tau) \leq \epsilon_2\}$. All of the following experiments' test F1 scores use this procedure with $\epsilon_1 = 1.5$ and $\epsilon_2 = 0.4$ for picking the threshold τ. The derivative is approximated using the forward difference.

Test F1 scores versus the number of unknown classes Q are plotted in Figure 2. While GMVAE is not as accurate in the closed-set regime, it outperforms CROSR as Q increases. CROSR's open-set accuracies, in turn, diminish as Q increases, CROSR+EVT in particular. GMVAE's F1 scores are more robust to increasing Q. For all $Q \geq 0$, GMVAE+NC-U's F1 scores are on average 0.06 greater than those of CROSR+EVT.

5.2 CIFAR-10 Withholding 4 Classes

The six known classes are airplanes, automobiles, birds, cats, deer, and dogs. The four unknown classes are frogs, horses, ships, and trucks. CIFAR-10's standard training set is randomly split into the validation set (6,000 samples of known classes) and training set (24,000 samples). CIFAR-10's standard testing set (10,000 samples) is kept the same.
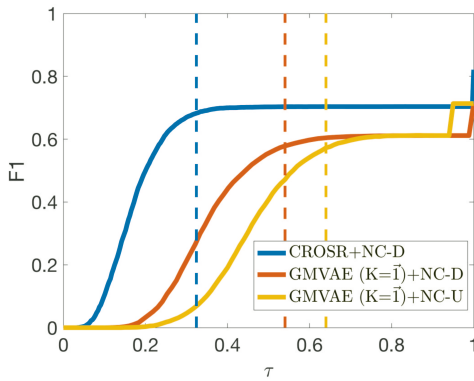
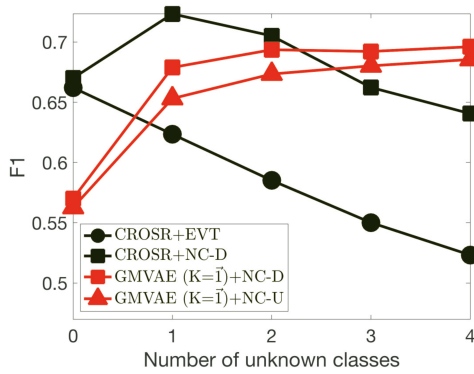Figure 1: Fashion MNIST known validation F1 scores versus τ and the corresponding picked thresholds.

Figure 2: Fashion MNIST open-set test F1 scores.

Figure 3: $K = \vec{1}$ CIFAR-10 open-set test F1 scores.

Figure 4: t-SNE plot of (left) both components $[y, z]$, (center) only y, and (right) only z of CROSR's training latent representations for the first CIFAR-10 experiment. Stars are the respective component's class centroids.

For both CIFAR-10 experiments, we use the same CROSR architecture as Yoshihashi et al. (2019) for their CIFAR-10 experiment.

Test F1 scores are plotted in Figure 3. GMVAE consistently beats CROSR and again CROSR+EVT performs worst. Algorithm 2 augments GMVAE and we deduce this is because unknown CIFAR-10 samples are more difficult to distinguish and thus more likely to be embedded to the interior of known latent clusters where "uncertainty" has more influence. For all $Q \geq 0$, GMVAE+NC-U F1 scores are on average 0.25 greater than those of CROSR+EVT.

We believe the underlying reason why CROSR's F1 scores in Figures 3 and 8 are so poor is because the activation vector y monopolizes the embedding since the reconstruction latent component z fails to cluster the classes. This is confirmed with latent t-SNE plots. We first show a t-SNE plot of the CROSR latent representation components in Figure 4 to bring into question the explicit use of classifier activation vectors in an open-set recognition embedding. We see that the reconstruction latent variable z does little to cluster the known classes and so open-set classification is dominated by the known classifier's activation vector y.

In contrast to CROSR, GMVAE's latent representation $\mu(x; \phi_z)$ in Figure 5 separates classes better (in comparison to the right figure in Figure 4). GMVAE's embedding is able to effectively capture both class and reconstruction infor-
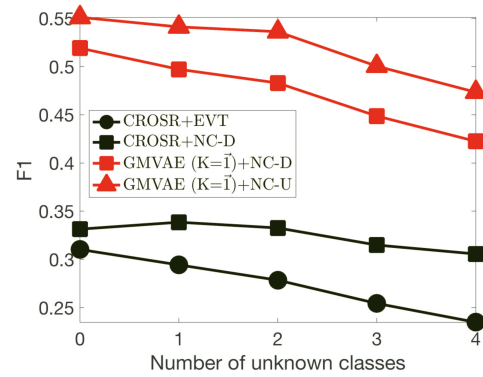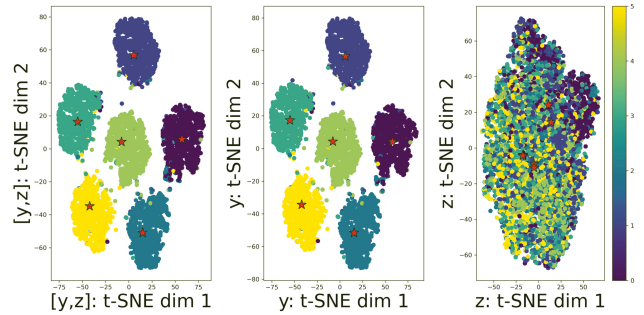
mation simultaneously, leading to more amenable open-set recognition. As CIFAR-10 images are highly hetergeneous within classes, we expect class overlap from reconstruction.
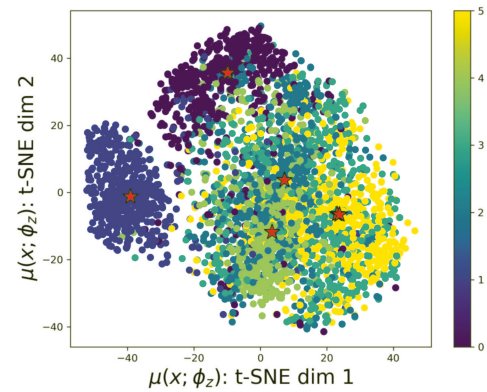

Figure 5: t-SNE plot of $\mu(x; \phi_z)$ of GMVAE's training latent representations for the first CIFAR-10 experiment. Stars are the class centroids.

5.3 MNIST with "Even" and "Odd" Classes

The two known classes are "even," comprised of digits 0 and 2, and "odd," comprised of digits 1 and 3. The six unknown classes are digits 4 and greater. MNIST's standard training set is randomly split into the validation set (4,000 samples of known classes) and training set (about 18,000 samples). MNIST's standard testing set (10,000 samples) is kept the same. We use the same CROSR architecture as Yoshihashi et al. (2019) for their MNIST experiment.

This is a clearcut example where each class has two subclusters. To determine that $K = (2, 2)$ is indeed the optimal GMVAE selection, we implement the procedure in §4 in Figure 6. On the left, the mean difference between the $K = 1$ and $K = 2$ latent covering loss is 0.86 while the mean difference between $K = 2$ and $K = 3$ is 0.22. This is indicative of two true subclusters within "even." Similarly on the right, the mean difference between $K = 1$ and $K = 2$ latent covering loss is 1.23 while the mean difference between $K = 2$ and $K = 3$ is -0.09. This is again indicative of two true subclusters within "odd." For these plots, the early epochs are truncated.
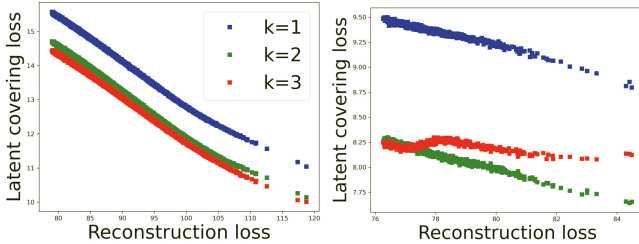

Figure 6: The latent covering loss plotted against reconstruction loss for increasing K for the (left) "even" and (right) "odd" classes of MNIST.

Test F1 scores are plotted in Figure 7. Here, CROSR+NC-D outperforms GMVAE+NC-D but not GMVAE+NC-U. However, CROSR+EVT again performs worst. There is a significant increase in GMVAE open-set accuracy and robustness to increasing Q from utilizing the "uncertainty" threshold. This algorithm complements the use of class subclusters as unknown classes' latent representations are strategically more likely embedded in the open space between centroids where U is larger. For all $Q \geq 0$, GMVAE+NC-U F1 scores are on average 0.29 greater than those of CROSR+EVT.

5.4 CIFAR-10 with "Animals" and "Vehicles" Classes

The two known classes are "animals," comprised of cats and dogs, and "vehicles," comprised of cars and trucks. The unknown classes are the other 6 classes. CIFAR-10's standard training set is randomly split into the validation set (4,000 samples of known classes) and training set (16,000 samples). CIFAR-10's standard testing set (10,000 samples) is kept the same. Determining that $K = (2, 2)$ is again the optimal GMVAE selection is qualitatively the same as the
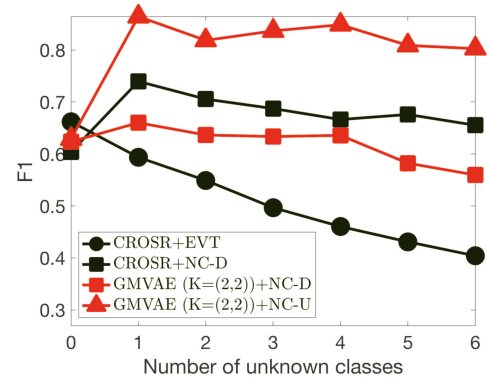

Figure 7: "Even" and "odd" MNIST open-set test F1 scores.

previous experiment. The parallel figures are placed in the technical appendix.

Test F1 scores are plotted in Figure 8. Discussed in §3.3, as a result of CROSR's softmax classifier, the centroids are not representative and thus its open-set classification suffers. Again, because of the class subclusters, the "uncertainty" threshold provides a significant increase in open-set recognition capability. For all $Q \geq 0$, GMVAE+NC-U F1 scores are on average 0.44 greater than those of CROSR+EVT.
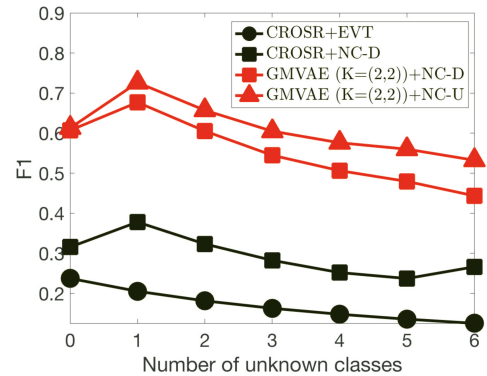

Figure 8: $K = (2, 2)$ CIFAR-10 open-set test F1 scores.

6 Conclusion

We developed GMVAE, an extension of Gaussian mixture variational autoencoders, as a better means of dual reconstruction-classification learning for open-set recognition. To augment this model we also introduced a novel "uncertainty" threshold that consistently beats other algorithms. Multiple image recognition experiments demonstrate that GMVAE outperforms CROSR, a previously state-of-the-art deep open-set classifier utilizing this same dual reconstruction-classification framework. The use of multiple subclusters per class and not relying on closed-set, softmax classifiers in the embedding, we believe, are instrumental in these results. Non-convex clustering of known classes remains an interesting open avenue of research within open-set recognition.

Acknowledgments

Ethics Statement

The immediate motivation for open-set recognition falls under automation. The ability of classifiers to predict unknown classes would focus and streamline human interaction with the system. This is perhaps most evident with computer vision tasks such as those found in automated driving. A procedure for identifying unknowns is critical when it is impossible to include all feasible classes in training. However, this in turn leads to the larger, ethics-centered question of how conservatively to proceed given an "unknown" classification. For instance, with autonomous driving, this requires a dilemmic balance between stopping to avoid hitting a potential life and perhaps consistently disrupting traffic flow.

While the focus of open-set recognition has primarily been image recognition, we also apply GMVAE to cancer treatment predictions. Cancer treatment regimens often consist of a combination, or "cocktail," of drugs. The landscape of cancer drug cocktails evolves with discoveries of novel cocktails with improved treatment and lessening side effects. Predicting cancer treatments can, therefore, be naturally formulated in terms of an open-set learning problem. Again, both physicians and patients may benefit from the automated efficiencies of this application but there might certainly be unintended negative effects. Any deep network can suffer from erroneously learning from demographic data and thus run the risk of being inappropriately biased. Our system is no different. While this may not present issues in innocuous datasets such as CIFAR-10, leveraging any biases in medical data could put large populations at risk for applications in medical treatment.

Finally, we have empirically observed that better open-set recognition often accompanies poorer closed-set classification. It seems natural to expect a trade-off between classifying known classes and robustly identifying unknown classes. And so, the consequences of failure of either open or closed-set classification can be unbounded in application. The further development of more robust and accurate deep open-set classifiers is therefore of significant importance as automation increases in the near future.

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