Beyond Black Box Densities: Parameter Learning for the Deviated Components

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Abstract

As we collect additional samples from a data population for which a known density function estimate may have been previously obtained by a black box method, the increased complexity of the data set may result in the true density being deviated from the known estimate by a mixture distribution. To model this phenomenon, we consider the *deviating mixture model* $(1-\lambda^*)h_0 + \lambda^*(\sum_{i=1}^k p_i^* f(x|\theta_i^*))$, where h_0 is a known density function, while the deviated proportion λ^* and latent mixing measure $G_* = \sum_{i=1}^k p_i^* \delta_{\theta_i^*}$ associated with the mixture distribution are unknown. Via a novel notion of distinguishability between the known density h_0 and the deviated mixture distribution, we establish rates of convergence for the maximum likelihood estimates of λ^* and G^* under Wasserstein metrics. Simulation studies are carried out to illustrate the theory.

1 Introduction

Most data-driven learning processes typically consist of iterative steps that involve model training and fine-tuning, with more data in-take leading to further model re-training and refinement. As more samples come in and exhibit more complex patterns, the initial model may be obsolete, risks being discarded or absorbed into a richer class of models to adapt better to increased complexity. It takes much resources to train complex models on a rich data population. Moreover, many successful models in modern real-world applications have become so complex that make them hard to properly evaluate and interpret; aside from the predictive performance they may as well be considered as black boxes. Nonetheless, as data populations evolve and so must the learning models, several desiderata remain worthy: the ability to adapt to new complexity while retaining aspects of old "wise" model, and the ability to interpret the changes.

In this paper we will investigate a class of complex models for density estimation that are receptive to *adaptation*, *reuse* and *interpretablity*: we posit that there is an existing distribution h_0 which may have been obtained a priori by some means for the data population of interest, e.g., via kernel density estimation (KDE) [22] or mixture models [20] or some modern black box methods, such as generative adversarial networks (GANs) [13, 1] or normalizing flows [9]. Nonetheless, as more samples come in and/or as the data population changes, it is possible that the true density may deviate from h_0. While h_0 is potentially difficult to interpret, it is the deviation from the known h_0 that we wish to learn and interpret. We will use a mixture distribution to represent this deviation, leading to what we call a *deviating mixture model* for the underlying data population:

$$p_{\lambda^* G_*}(x) := (1-\lambda^*)h_0(x) + \lambda^* F(x, G_*), \tag{1}$$

for $x \in \mathbb{R}^d$, where $F(x, G_*) := \sum_{i=1}^{k_*} p_i^* f(x|\theta_i^*))$ represents a mixture distribution for the density components deviating from h_0. Such deviating components are from a known family of density function f. The unknown parameters for this model are the mixing proportion $\lambda^* \in [0,1]$, and the

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mixing measure $G_* = \sum_{i=1}^{k_*} p_i^* \delta_{\theta_i^*}$, where $k_* \geq 1$ number of *deviated* components. The choice of mixture distribution $F(x, G_*)$ allows us to express complex deviation from h_0, yet the overall model remains amenable to the interpretation of its parameters: λ_* represents the amount of deviation from the existing candidate h_0, while the mixing measure G_* represents heterogeneous patterns of the deviation. The primary contribution of this paper is a rigorous investigation into the rather challenging questions of identifiability and parameter learning rates that arise from a standard maximum likelihood estimation procedure.

Relations to existing works. This modeling framework owes its roots to several significant bodies of work in both statistics and machine learning literature. In classical statistics, a dominant approach to address increased complexity of data populations is via hypothesis testing: one can test alternative (possibly composite) hypothesis represented by a class of distributions against the null hypothesis represented by h_0. Due to the constraint for obtaining simple and theoretically valid test statistics in order to accept or reject the null hypothesis, the testing approaches were mostly restricted to simple choices of distribution for the null and alternative hypotheses [6, 10, 7, 4, 8]. More similar to (1) is the class of *contaminated mixture models* for density estimation: in this framework, the data are assumed to be sampled from a mixture of P_0 and Q where either P_0 or Q can be an unknown distribution that needs to be estimated. While this approach offers more flexibility in terms of modeling, it does not always guarantee the identifiability of the mixing weight or mixture components P_0, Q [25, 23, 19]. Without identifiability, it is virtually impossible to interpret the model parameters for the data domains. To avoid the identifiability issue, several researchers added the semi-parametric or parametric structures on P_0 and Q, such as P_0 and Q are mixture distributions [2, 12]. However, to the best of our knowledge, the convergence theory of these models remains poorly understood, except for some simple settings (see also [3, 5]). The main distinction between our modeling framework of deviating mixture models and the existing research on contaminated mixtures lies in our assumption that one of the mixture components, namely h_0 is known, allowing us to focus on the inference of the deviation from h_0, for which a considerable learning theory for the parameters of interest can be established and will be presented in this paper.

Finally, estimating parameters of mixture distributions is an essential problem in mixture models. The convergence properties have been studied using identifiability notions and Wasserstein distances [21, 18, 15]. Our technical approach requires a generalization of the identifiability notion to take account of structural property of the existing component h_0, which helps to shed light on a considerably more complex convergence behavior of the deviated components.

Contributions. The primary contribution of this paper is a rich theory of *identifiability* and rates of convergence for *parameters* and density estimation that arise in the *deviating mixture model* (1), under various settings of the existing component h_0, and that of the deviating components (via f and G_*). The convergence of density estimation in Hellinger distance under the MLE procedure is well studied in [26]. Thus, the bulk of our technical innovation lies in establishing a collection of *inverse bounds* which relate the Hellinger distance of densities in model (1) in terms of that of their parameters. To this end, we introduce a novel notion of *distinguishability* between h_0 and family of density f. The inverse bounds will be characterized under such distinguishability conditions (or the lack thereof). Moreover, we will be able to provide many examples to demonstrate the broad applicability of our theory, including cases where the existing component h_0 is obtained by a black box method (e.g., deep learning model) and a more traditional method (e.g., via KDE's or mixture models).

Organization. The remainder part of this paper is structured as follows. In Section 2, we review the MLE method and the identifiability conditions, where the notion of distinguishability is presented. In Section 3, the main results of inverse bounds and convergence rates for parameters estimation of model (1) are shown. In Section 4, multiple simulation experiments are carried out to support the theory. Finally, Section 5 is used to discuss and conclude. Proofs of all the results in the main text are deferred to the Supplementary Material.

Notation. We denote by $\mathcal{E}_k(\Theta) = \{\sum_{i=1}^k p_i f(x|\theta_i) : \sum_{i=1}^k p_i = 1, p_i > 0, \theta_i \in \Theta \, \forall 1 \leq i \leq k\}$ the family of mixtures with exactly k components and $\mathcal{O}_K(\Theta) = \{\sum_{i=1}^K p_i f(x|\theta_i) : \sum_{i=1}^K p_i = 1, p_i \geq 0, \theta_i \in \Theta \, \forall 1 \leq i \leq K\}$ the family of mixtures with no more than K components. $\mathcal{E}_{k,c_0}(\Theta) = \{\sum_{i=1}^k p_i f(x|\theta_i) : \sum_{i=1}^k p_i = 1, p_i \geq c_0, \theta_i \in \Theta \, \forall 1 \leq i \leq k, k \leq K\}$ is the family of mixtures with exactly K components and mixing proportions being bounded below by c_0, and $\mathcal{O}_{K,c_0}(\Theta) = \{\sum_{i=1}^{k'} p_i f(x|\theta_i) : \sum_{i=1}^{k'} p_i = 1, p_i \geq c_0, \theta_i \in \Theta \, \forall 1 \leq i \leq k', k' \leq K\}$. $\|\cdot\|_2$ is the usual l^2 norm

for vectors in $\mathbb{R}^d$ and matrices in $\mathbb{R}^{d\times d}$. We write $g(x) \gtrsim h(x)$ if $g(x) > ch(x)$ for all x, where c is a constant does not depend on x (similar for $g(x) \lesssim h(x)$). For any $\lambda \in \mathbb{R}$ and $B \subset \mathbb{R}$, denote by $1_{\{\lambda\in B\}}$ the function that takes value 1 if $\lambda \in B$, and 0 otherwise. For any two densities p and q, we denote $h(p, q)$ by the Hellinger distance and $V(p, q)$ by the Total Variation distance between them.

2 Identifiability and distinguishability theory

The principal goal of the paper is to establish the efficiency of parameter learning for the deviating mixture model (1) via the standard maximum likelihood estimation (MLE) method. To achieve this goal, the parameters have to be identifiable to begin with. Thus, our theory builds on and extends a standard notion of identifiability of families of density $\{f(x|\theta) : \theta \in \Theta\}$ that has been considered in previous work [21, 15].

Definition 2.1. The family $\{f(x|\theta), \theta \in \Theta\}$ (or in short, f) is <u>identifiable in the order r</u>, for some $r \geq 1$, if $f(x|\theta)$ is differentiable up to the order r in θ and the following holds:

A1. For any $k \geq 1$, given k different elements $\theta_1, \ldots, \theta_k \in \Theta$. If we have $\alpha_\eta^{(i)}$ such that for almost all x

$$\sum_{l=0}^{r} \sum_{|\eta|=l} \sum_{i=1}^{k} \alpha_\eta^{(i)} \frac{\partial^{|\eta|} f}{\partial \theta^\eta}(x|\theta_i) = 0$$

then $\alpha_\eta^{(i)} = 0$ for all $1 \leq i \leq k$ and $|\eta| \leq r$.

Many commonly used families f for mixture modeling satisfy the first order identifiability condition, including location-scale Gaussian distributions, e.g., $f(x|\theta) = N(x|\mu, \sigma^2)$ where μ and σ^2 represent the mean and variance parameters, and location-scale Student's t-distributions. In model (1), however, due to the presence of the existing component h_0, the deviated mixture components need to be *distinguishable* from h_0. This motivates a more general notion of identifiability, namely, *distinguishability* that we now define. This condition specifies a property jointly for both the existing component h_0 and the family of density functions f that make up the deviated components.

Definition 2.2. For any natural numbers $k, r \geq 1$, we say that the family of density functions $\{f(\cdot|\theta), \theta \in \Theta\}$ with complexity level k (or in short, (f, k)) is <u>distinguishable</u> <u>up to the order r</u> from h_0 if the following holds:

A2. For any k distinct components $\theta_1, \ldots, \theta_k$, if we have real coefficients $\alpha_\eta^{(i)}$ for $0 \leq i \leq k$ such that

$$\alpha^{(0)} h_0(x) + \sum_{l=0}^{r} \sum_{|\eta|=l} \sum_{i=1}^{k} \alpha_\eta^{(i)} \frac{\partial^{|\eta|} f}{\partial \theta^\eta}(x|\theta_i) = 0,$$

for almost surely $x \in \mathcal{X}$, then $\alpha^{(0)} = \alpha_\eta^{(i)} = 0$ for $1 \leq i \leq k'$ and $|\eta| \leq r$.

A simple but non-trivial example of the distinguishability condition can be derived directly from the definitions.

Example 2.3. (a) When $h_0(x) = \sum_{i=1}^{k_0} p_i^0 f(x|\theta_i^0)$ for some given weights $(p_1^0, \ldots, p_{k_0}^0)$ and parameters $(\theta_1^0, \ldots, \theta_{k_0}^0)$ where $k_0 \geq 1$, then (f, k) is distinguishable in the order r from h_0 as long as $k < k_0$ and the family of density f is identifiable in the order r.
(b) Given the choice of h_0 in (a), (f, k) is not distinguishable in the order r from h_0 when $k \geq k_0$.

More significantly, we can establish a broad class of h_0 and families f for which distinguishability holds. This is exemplified by the following theorem, where f represents a family of location or location-scale Gaussian kernels, and h_0 is subject to a relatively weak condition.

Theorem 2.4. *(a) Suppose that $-\log h_0(x) \gtrsim \|x\|_2^{\beta_1}$ or $-\log h_0(x) \lesssim \|x\|_2^{\beta_2}$ for all $\|x\|_2 > x_0$, for some $x_0 > 0$, $\beta_1 > 2$, and $\beta_2 < 2$. Then, for f being family of location-scale Gaussian and any $k > 0$, (f, k) is distinguishable from h_0 up to the first order, where the derivatives in Assumption A2 are taken with respect to both location and scale parameters, and (f, k) is also distinguishable from h_0 up to any order, where the derivatives in Assumption A2 are taken only with location parameters.*

(b) Suppose that h_0 is the pdf of a pushforward measure of $N(0, I_d)$ by a piecewise linear function with a finite and positive number of breakpoints. Then, the same conclusions as in part (a) hold.

The proof of Theorem 2.4 is in Appendix C.1. Note that in part (a), h_0 can be a pdf of any distribution possessing a lighter or heavier tail than Gaussian distributions, and in part (b), h_0 represents the pushforward of a Gaussian distribution by any piecewise linear function (recall that family of piecewise linear functions is dense in the Banach space of continuous functions with compact support). In the sequel we shall demonstrate several examples of interest that are applicable to Theorem 2.4 where h_0 may have been estimated by some popular "black box" methods.

Kernel based representation. Suppose that h_0 was obtained from a m- sample $Y_1, \ldots, Y_m \in \mathbb{R}^d$ by a classical kernel density estimation (KDE) method [22] or a RKHS-based method [24], so that

$$h_0(x) = \frac{1}{m} \sum_{j=1}^{m} k_\sigma(x, Y_j) \quad \forall x \in \mathbb{R}^d, \tag{2}$$

where k_σ is a kernel function with bandwidth σ. Popular choices of kernels include the Gaussian kernel $k_\sigma(x, x') = \left(\frac{1}{\sqrt{2\pi}\sigma}\right)^d \exp\left(-\frac{\|x - x'\|_2^2}{2\sigma^2}\right)$ and the multivariate Student's kernel $k_\sigma(x, x') = \left(\frac{1}{\sqrt{\pi}\sigma}\right)^d \frac{\Gamma((\nu+d)/2)}{\Gamma(\nu/2)} \left(1 + \frac{\|x-x'\|_2^2}{\nu\sigma^2}\right)^{-\frac{\nu+d}{2}}$. The corresponding distinguishability guarantee is as follows.

Corollary 2.5. *Suppose h_0 is defined by Eq. (2), where k_σ is Gaussian kernel and $m > K$, or k_σ is the multivariate Student's kernel. Then, for f being family of location-scale Gaussian, (f, K) is distinguishable from h_0 up to the first order, where the derivatives in Assumption A2 are taken with respect to both location and scale parameters, and (f, K) is also distinguishable from h_0 up to any order, where the derivatives in Assumption A2 are taken only with location parameters.*

In application, it is common that the condition $m > K$ is satisfied. It is also matches with the scenario that we consider in the paper, where h_0 is already trained using a big data set, and there is a small number of deviated components.

Neural networks. Deep neural networks represent a powerful, albeit black box, approximation device for constructing rich classes of distribution for generative models [13, 1]. Accordingly, h_0 is the pdf function of a Gaussian distribution being push-forwarded by a map T, which is represented by a neural network (NN). Suppose that the NN representing T has a positive and finite number of layers L, and so

$$T(x) = a(W_L a(W_{L-1}(\ldots a(W_1 x + b_1)) + b_{L-1}) + b_L), \tag{3}$$

where $W_1, \ldots, W_L \in \mathbb{R}^{d \times d}$ are the weights and $b_1, \ldots, b_L \in \mathbb{R}^d$ are the biases. The activation function a is chosen to be rectified linear unit (ReLU) function defined by $a(x) = \max\{x, 0\}$, and is applied elementwise to any vector in $\mathbb{R}^d$. The corresponding guarantee on the distinguishability condition is as follows.

Corollary 2.6. *Suppose that h_0 is the pdf a pushforward measure of $N(0, I_d)$ by a map T defined by Eq. (3). Then, for f being family of location-scale Gaussian and any $k > 0$, (f, k) is distinguishable from h_0 up to the first order, where the derivatives in Assumption A2 are taken with respect to both location and scale parameters, and (f, k) is also distinguishable from h_0 up to any order, where the derivatives in Assumption A2 are taken only with location parameters.*

3 Convergence rates of density estimation

In this section, we first establish the rate of density estimation for the deviating mixture models in Section 3.1. We then describe a general procedure to obtain the convergence rate of parameter estimation based on that of density estimation via inverse bounds in Section 3.2. Finally, we provide comprehensive inverse bounds under several settings of the deviating mixture models in Section 3.3.

3.1 MLE for deviating mixture model

Given n i.i.d. sample $X_1, X_2, \ldots, X_n$ from $p_{\lambda^* G_*}$ as in model (1), where G_* has k_* components, we want to estimate λ^* and G_* from the data. We refer to the problem as in *exact-fitted setting* if k_* is known, and we refer to it as in *over-fitted setting* if k_* is unknown but is known to be bounded by

some number K. We denote the MLE for exact-fitted setting by

$$\widehat{\lambda}_n, \widehat{G}_n \in \underset{\lambda \in [0,1], G \in \mathcal{E}_{k_*}(\Theta)}{\arg\max} \sum_{i=1}^{n} \log(p_{\lambda G}(X_i)),$$

and for the over-fitted setting, we replace $\mathcal{E}_{k_*}(\Theta)$ in the equation above by $\mathcal{O}_K(\Theta)$, where $K \geq k_*$.

In order to state a rate of convergence for the density estimators $p_{\widehat{G}_n}$ under the Hellinger distance h [26], we need a condition on the complexity of the function class

$$\overline{\mathcal{P}}_k^{1/2}(\Theta, \epsilon) = \left\{ \bar{p}_{\lambda G}^{1/2} : G \in \mathcal{O}_k(\Theta),\ h(\bar{p}_{\lambda G}, p_{\lambda^* G_*}) \leq \epsilon \right\}, \tag{4}$$

where for any $G \in \mathcal{O}_K(\Theta)$, we write $\bar{p}_{\lambda G} = (p_{\lambda G} + p_{\lambda^* G_*})/2$. The definition of $\overline{\mathcal{P}}_k(\Theta, \epsilon)$ originates from [26]. We measure the complexity of this class through the bracketing entropy integral

$$\mathcal{J}_B(\epsilon, \overline{\mathcal{P}}_k^{1/2}(\Theta, \epsilon), \nu) = \int_{\epsilon^2/2^{13}}^{\epsilon} \sqrt{\log N_B(u, \overline{\mathcal{P}}_k^{1/2}(\Theta, u), \nu)} du \vee \epsilon, \tag{5}$$

where $N_B(\epsilon, X, \eta)$ denotes the ϵ-bracketing number of a metric space (X, η) and ν is the Lebesgue measure. We require the following assumption.

A3. Given a universal constant $J > 0$, there exists $N > 0$, possibly depending on d and k, such that for all $n \geq N$ and all $\epsilon > (\log n/n)^{1/2}$,

$$\mathcal{J}_B(\epsilon, \overline{\mathcal{P}}_k^{1/2}(\Theta, \epsilon), \nu) \leq J\sqrt{n}\epsilon^2.$$

Theorem 3.1. *Assume that Assumption A3 holds, and let $k \geq 1$. There exists a constant $C > 0$ depending only on d, k such that for all $n \geq 1$,*

$$\sup_{G_* \in \mathcal{O}_k(\Theta), \lambda^* \in [0,1]} \mathbb{E}_{\lambda^*, G_*} h(p_{\widehat{\lambda}_n \widehat{G}_n}, p_{\lambda^* G_*}) \leq C\sqrt{\log n/n}.$$

Therefore, in order to get convergence rate for density functions based on MLE procedure, we only need to check assumption A3. This assumption holds true for a wide range class of parametric model [26]. For our model, we give an example that it holds when h_0 has an exponential tail (satisfied for KDE's and Neural networks above) and f is location-scale Gaussian distribution.

Proposition 3.2. *Suppose f is location-scale Gaussian family and $\Theta = [-a, a]^d \times \Omega$, where Ω is a subset of S_d^{++} whose eigenvalues are bounded in $[\underline{\lambda}, \overline{\lambda}]$, $a, \underline{\lambda}, \overline{\lambda} > 0$, and h_0 is bounded with tail $-\log h_0(x) \gtrsim \|x\|_2^{\beta}$ for some $\beta > 0$. Then, the family of densities $\{p_{\lambda G} : \lambda \in [0,1], G \in \mathcal{O}_k(\Theta)\}$ satisfies assumption A3.*

3.2 Parameter learning rates of deviated components

The core of this paper lies in establishing a collection of *inverse bounds*, provided that some distinguishability condition developed in Section 2 holds. The inverse bounds basically say that a small distance between $p_{\lambda G}$ and $p_{\lambda^* G_*}$ under the total variation distance entails that (λ, G) and (λ^*, G_*) are similar under appropriate distances, where (λ^*, G_*) is fixed. To this end, we employ Wasserstein metrics [27] and their extensions.

Wasserstein distances. Wasserstein distances are natural and useful for assessing the convergence of latent mixing measures in mixture models [21, 16, 14]. Given two measures $G = \sum_{i=1}^{k} p_i \delta_{\theta_i}$ and $G' = \sum_{j=1}^{k'} p'_j \delta_{\theta'_j}$ on a space Θ endowed with a metric ρ, the Wasserstein metric of order $r \geq 1$ is:

$$W_r(G, G') = [\inf_q \sum_{i,j} q_{ij} \rho^r(\theta_i, \theta'_j)]^{1/r},$$

where the infimum is taken over all joint distribution on $[1, \ldots, k] \times [1, \ldots, k']$ such that $\sum_i q_{ij} = p'_j, \sum_j q_{ij} = p_i$. Note that if G_n is a sequence of discrete measures that converges to G in a Wasserstein distance, then for every atom of G, there is a subset of atoms of G_n converges to it. Therefore, the convergence in Wasserstein metrics implies convergence of parameters in mixture models. In this paper, space Θ is often chosen to be a compact subset of $\mathbb{R}^d$ and ρ is the usual l^2 distance. In the case of location-scale Gaussian mixtures, space Θ is a compact subset of $\mathbb{R}^d \times$

S_d^{++}, where S_d^{++} is the set of positive definite and symmetric matrices in $\mathbb{R}^{d\times d}$, and for every $(\mu,\Sigma),(\mu',\Sigma')\in\Theta$, the distance ρ is defined by $\rho((\mu,\Sigma),(\mu',\Sigma'))=\|\mu-\mu'\|_2+\|\Sigma-\Sigma'\|_2$.

From inverse bounds to parameter learning rates. Suppose that some distinguishablity condition is satisfied, then we will establish an inverse bound providing a guarantee that a small distance between $p_{\lambda^* G_*}$ and $p_{\lambda G}$ entails a small distance between λ and λ^* and between G and G_*. More concretely, define a divergence between two measures λG and $\lambda^* G_*$ via

$$\overline{W}_r(\lambda G,\lambda^* G_*):=|\lambda-\lambda^*|+(\lambda+\lambda^*)W_r^r(G,G_*).$$

for all $r\geq 1$, and the inverse bounds will have the form that $V(p_{\lambda G},p_{\lambda^*,G_*})\gtrsim\overline{W}_r(\lambda G,\lambda^* G_*)$, for some r that depends on the level of distinguishable level of the model. Since total variational distance is upper bounded by Hellinger distance, if Assumption A3. holds, then combining the aforementioned inverse bound with Theorem 3.1 we immediately obtain

$$\mathbb{E}_{\lambda^*,G_*}\overline{W}_r(\widehat{\lambda}_n\widehat{G}_n,\lambda^* G_*)\leq C\sqrt{\frac{\log n}{n}}.$$

This further implies that the convergence rate of $\hat{\lambda}_n$ to λ^* is of order $(\log(n)/n)^{1/2}$ and the convergence rate of $W_r(\hat{G}_n,G_*)$ to 0 is of order $(\log(n)/n)^{1/2r}$.

3.3 Inverse bounds in distinguishable setting

We shall establish inverse bounds provided a distinguishability condition for model (1) holds under either exact-fitted and over-fitted settings regarding the true number of components k_*.

Theorem 3.3. *Assume that k_* is known and (f,k_*) is distinguishable in the first order from h_0. Then, for any $G\in\mathcal{E}_{k_*}(\Theta)$, there exist positive constant C_1 and C_2 depending only on λ^*,G_*,h_0,Θ such that the following holds:*

(a) When $\lambda^*=0$, then $V(p_{\lambda^* G_*},p_{\lambda G})\geq C_1\lambda$.

(b) When $\lambda^*\in(0,1]$, then $V(p_{\lambda^* G_*},p_{\lambda G})\geq C_2\overline{W}_1(\lambda G,\lambda^* G_*)$.

In this theorem, we can see that if the number of components k_* of G_* is known, the first order distinguishability is sufficient to establish the lower bound of the total variation distance in terms of latent mixing measures' distance. In application, the true number of components k_* might not be known and we often fit the model (1) with $G\in\mathcal{O}_K(\Theta)$ for some $K\geq k_*$. The next result shows that similar bounds can also be established in this case, where we require distinguishability of f and h_0 in a higher order.

Theorem 3.4. *Assume that k_* is unknown and strictly upper bounded by a given K. Assume additionally that (f,K) is distinguishable in second order from h_0. Then, for any $G\in\mathcal{O}_K(\Theta)$, there exist positive constant C_1 and C_2 depending only on λ^*,G_*,h_0,Θ such that the following holds:*

(a) When $\lambda^*=0$, then $V(p_{\lambda^* G_*},p_{\lambda G})\geq C_1\lambda$.

(b) When $\lambda^*\in(0,1]$, then $V(p_{\lambda^* G_*},p_{\lambda G})\geq C_2\overline{W}_2(\lambda G,\lambda^* G_*)$.

Thanks to the distinguishability up to second order, no matter how large the number of over-fitted components K is, we always get the $\overline{W}_2$ lower bound for the total variation distances. Next, we relax the assumption of Theorem 3.4 by working on the setting where f is not second order identifiable. This is an instance of the so-called *weakly identifiable* setting — One popular example of weakly identifiable f is location-scale Gaussian distribution, which admits the partial differential equation (PDE) structure $\frac{\partial^2 f}{\partial\mu^2}(x|\mu,\Sigma)=2\frac{\partial f}{\partial\Sigma}(x|\mu,\Sigma)$, for all $x\in\mathbb{R}^d$ where $f(x|\mu,\Sigma)$ stands for location-scale Gaussian density function with location μ and covariance Σ. In order to illustrate the result of our bound for that weak identifiability setting of f, we specifically consider f to be location-scale Gaussian distribution. In this case, the parameter space Θ is a compact subset of $\mathbb{R}^d\times S_d^{++}$, where S_d^{++} is the set of positive definite and symmetric matrices in $\mathbb{R}^{d\times d}$ equipped with the usual Frobenius norm. To put our result in context, we shall adopt a notion used in analyzing the convergence rate of parameter estimation in location-scale Gaussian mixtures in [16]. For any $k\geq 1$, let $\overline{r}(k)$ be the minimum value of r such that the following system of polynomial equations:

$$\sum_{j=1}^{k+1}\sum_{n_1,n_2}\frac{c_j^2 a_j^{n_1}b_j^{n_2}}{n_1!n_2!}=0 \text{ for each } \alpha=1,\ldots,r, \tag{6}$$

does not have any nontrivial solution for the unknown variables $(a_j, b_j, c_j)_{j=1}^{k+1}$, where the ranges of n_1 and n_2 in the second sum consist of all natural pairs satisfying the equation $n_1 + 2n_2 = \alpha$. A solution to the above system is considered *nontrivial* if all of variables c_j are non-zeroes, while at least one of the a_j is non-zero. Some examples of known values of $\overline{r}$ are $\overline{r}(1) = 4$ and $\overline{r}(2) = 6$, and $\overline{r}(k) \geq 7$ for all $k \geq 3$. Using this notion, we can characterize the convergence of parameters of model (1) for the location-scale Gaussian family via the following theorem for inverse bounds.

Theorem 3.5. *Assume that $G^* \in \mathcal{E}_{k^*,c_0}(\Theta)$, and k_* is unknown and strictly upper bounded by a given K. In addition, f is location-scale Gaussian distribution and (f, K) with varied location, fixed variance parameters is distinguishable in any order from h_0. Then, for any $G \in \mathcal{O}_{K,c_0}(\Theta)$, there exist positive constant C_1 and C_2 depending only on $\lambda^*, G_*, h_0, \Theta$ such that the following holds:*

(a) When $\lambda^* = 0$, then $V(p_{\lambda^* G_*}, p_{\lambda G}) \geq C_1 \lambda$.

(b) When $\lambda^* \in (0, 1]$, then $V(p_{\lambda^* G_*}, p_{\lambda G}) \geq C_2 \overline{W}_{\overline{r}(K-k_*)}(\lambda G, \lambda^* G_*)$.

3.4 Inverse Bounds in Partially Distinguishable Setting

What happens if the distinguishability condition required by Def. 2.2 no longer holds generally? Recall in Example 2.3 (b) that this situation is not uncommon, specifically when

$$h_0(x) = f(x; G_0) = \sum_{i=1}^{k_0} p_i^0 f(x|\theta_i^0), \tag{7}$$

where $G_0 := \sum_{i=1}^{k_0} p_i^0 \delta_{\theta_i^0}$. In some specific cases of this setting, in fact, we fail to attain distinguishability, and the model may not even be identifiable in the classical sense, i.e. $p_{\lambda G} = p_{\lambda^* G_*}$ does not guarantee to have $\lambda G = \lambda^* G_*$. Since h_0 is the pdf of a mixture distribution — a popular choice for modeling complex forms of probability densities given its amenability to interpretation compared to black box type models — it is of interest to study the implication of parameter estimation for the deviated components in this setting, provided that the distinguisability condition may be at least partially achieved in some suitable sense. As we shall see, our theory demands a more refined analysis. To facilitate the presentation, denote $\mathcal{A} := \{1 \leq i \leq k_* : \theta_i^* \in \{\theta_1^0, \ldots, \theta_{k_0}^0\}\}$. Also, set $\bar{k} := |\mathcal{A}|$, which stands for the cardinal of the set $\mathcal{A}$. Our results will be divided into three separate regimes of $\bar{k}$ and λ^*: (i) $\lambda^* = 0$, (ii) $\bar{k} < k_0$ and $\lambda^* \in (0, 1]$, and (iii) $\bar{k} = k_0$ and $\lambda^* \in (0, 1]$. We only choose to present results of the second regime (ii) in the main text because of limited spaces and because of its representativeness as it shows all the intriguing behaviours of the model in this partially distinguishable setting. The first and third regime are deferred to Appendix A.

3.4.1 Regime B: $\bar{k} < k_0$ and $\lambda^* \in (0, 1]$

First, we consider the exactly-specified setting of model (1), namely, k_* is known. When $\bar{k} < k_0$, we can check that we still have dishtinguishability of h_0 and linear combinations of $\{f(x|\theta_i^*)\}_{i=1}^{k_*}$ and its derivatives. Therefore, as long as f is first order identifiable, one can invoke the proof of Theorem 3.3 to establish the same lower bound $V(p_{\lambda G}, p_{\lambda^* G_*})$ in terms of $\overline{W}_r(\lambda G, \lambda^* G_*)$ for some $r \geq 1$. Thus, our focus in this subsection is the settings when k_* is unknown.

Over-fitted setting with strongly identifiable f. Moving to the over-fitted settings of model setup (1), i.e., k_* is unknown and strictly upper bounded by a given K, as long as $K \geq k_0$, (f, K) is not distinguishable from h_0. Therefore, the results of Theorem 3.3 are not always applicable to the setting when $K \geq k_0$. Besides, in the over-fitted setting, the identifiability of model (1) no longer holds. Indeed, for any $\lambda > \lambda^*$, if we take

$$\overline{G}_*(\lambda) = (1 - \lambda^*/\lambda)\, G_0 + (\lambda^*/\lambda) G_*, \tag{8}$$

then $p_{\lambda^* G_*} = p_{\lambda \overline{G}_*(\lambda)}$. We present this pathological behavior in the following result.

Theorem 3.6. *Assume that h_0 takes the form (7) and $\bar{k} < k_0$. Besides that, $K \geq k_0$ and f is second order identifiable. Then, for any $G \in \mathcal{O}_K(\Theta)$, there exist positive constants C_1 and C_2 depending only on $\lambda^*, G_*, h_0, \Theta$ such that the following hold:*

(a) If $K \leq k_* + k_0 - \bar{k} - 1$, then $V(p_{\lambda^*, G_*}, p_{\lambda, G}) \geq C_1 \overline{W}_2(\lambda G, \lambda^* G_*)$,

(b) If $K \geq k_* + k_0 - \bar{k}$, then

$$V(p_{\lambda^*, G_*}, p_{\lambda, G}) \geq C_2 \left(1_{\{\lambda \leq \lambda^*\}} \overline{W}_2(\lambda G, \lambda^* G_*) + 1_{\{\lambda > \lambda^*\}} W_2^2(G, \overline{G}_*(\lambda)) \right).$$

(c) As a special case, if $K = k_* + k_0 - \bar{k}$, we have

$$V(p_{\lambda^*,G_*}, p_{\lambda,G}) \geq C_3 1_{\{\lambda > \lambda^* + \delta\}} W_1(G, \overline{G}_*(\lambda)),$$

for all $\delta > 0$, where C_3 depends on $\lambda^*, G_*, h_0, \Theta, \delta$.

A few comments are in order. Firstly, as we can see, the magnitude of λ compared to λ^* will decide the solution of (λ, G) to the identifiable equation $p_{\lambda G} = p_{\lambda^* G^*}$, therefore lead to different lower bounds such in part (b) of the theorem. In particular, if $\lambda \leq \lambda^*$, the solution is $(\lambda, G) = (\lambda^*, G_*)$, and for any $\lambda > \lambda^*$, the solution is $G = \overline{G}_*(\lambda)$ given in Eq. (8). Specifically, when λ is strictly larger than λ^* by some amount $\delta > 0$, then the latter case is well separated from the former, and we have an exact-fitted result when $K = k_0 + k_* - \bar{k}$. Secondly, we can view $p_{\lambda G}$ as a mixture distributions with latent mixing measures $\widehat{G} = (1-\lambda)\sum_{i=1}^{k_0} p_i^0 \delta_{\theta_i^0} + \sum_{i=1}^{K} p_i \delta_{\theta_i}$ having at most $K + k_0$ elements, while $p_{\lambda^* G_*}$ as a mixture with latent measure $\widehat{G}_* = \sum_{i=1}^{\bar{k}} \left[(1-\lambda^*)p_i^0 + \lambda^* p_i^*\right] \delta_{\theta_i^0} + \sum_{i=\bar{k}+1}^{k_0} (1 - \lambda^*) p_i^0 \delta_{\theta_i^0} + \sum_{i=\bar{k}+1}^{k_*} \lambda^* p_i^* \delta_{\theta_i^*}$ having exactly $k_0 + k_* - \bar{k}$ elements. Because $k_0 + k_* - \bar{k} < K + k_0$, a direct application of Theorem 3.2 in [17] gives us $V(p_{\lambda^*,G_*}, p_{\lambda,G}) \gtrsim W_2^2(\widehat{G}_*, \widehat{G})$. But this bound is not as tight as what in Theorem 3.6(c), since $W_1 \gtrsim W_2^2$. The bounds established in the theorem are possible as we carefully explore the structure of $\widehat{G}_*$ and $\widehat{G}$.

Over-fitted setting with weakly identifiable f. Similar to Theorem 3.5, when f is the location-scale Gaussian, the weak identifiability can worsen the power of the bound in the over-fitted case.

Theorem 3.7. *Assume that h_0 takes the form (7). Besides that, $K \geq k_0$ and f is location-scale Gaussian distribution. Then, for any $\lambda \in [0,1]$ and $G \in \mathcal{O}_{K,c_0}(\Theta)$ for some $c_0 > 0$, there exist positive constants C_1, C_2, C_3, C_4 depending only on $\lambda^*, G_*, G_0, \Theta$ (C_3 and C_4 also depend on δ) such that the following holds:*

(a) When $K \leq k_* + k_0 - \bar{k} - 1$, then $V(p_{\lambda^*,G_*}, p_{\lambda,G}) \geq C_1 \overline{W}_{\overline{r}(K-k_*)}(\lambda G, \lambda^* G_*)$.

(b) When $K \geq k_* + k_0 - \bar{k}$, then

$$V(p_{\lambda^*,G_*}, p_{\lambda,G}) \geq C_2 \left(1_{\{\lambda \leq \lambda^*\}} \overline{W}_{\overline{r}(K-k_*)}(\lambda G, \lambda^* G_*) + 1_{\{\lambda > \lambda^*\}} W^{\overline{r}(K-k_*)}_{\overline{r}(K-k_*)}(G, \overline{G}_*(\lambda)) \right).$$

(c) For $\delta > 0$, when $K = k_* + k_0 - \bar{k}$, we have

$$V(p_{\lambda^*,G_*}, p_{\lambda,G}) \geq C_3 1_{\{\lambda > \lambda^* + \delta\}} W_1(G, \overline{G}_*(\lambda)),$$

and when $K > k_* + k_0 - \bar{k}$, we have

$$V(p_{\lambda^*,G_*}, p_{\lambda,G}) \geq C_4 1_{\{\lambda > \lambda^* + \delta\}} W^{\overline{r}(K-k_0-k_*+\bar{k})}_{\overline{r}(K-k_0-k_*+\bar{k})}(G, \overline{G}_*(\lambda)).$$

In this theorem, we once again observe the pathological behavior of the lower bound by Wasserstein distances caused by the unidentifiability of the model (1). In part (c), when there is a well separation between two region of solutions of equation $p_{\lambda G} = p_{\lambda^* G_*}$, we can improve the order of Wasserstein distances for both exact-fitted case and over-fitted case. In application, if $\hat{\lambda}_n$ and $\hat{G}_n$ are the MLE of model (1) estimated by n i.i.d. data, then the convergence of $(\hat{\lambda}_n, \hat{G}_n)$ depends on the limit of $\hat{\lambda}_n$ (or its subsequence) comparing to λ^*. If $K = k_0 + k_* - \bar{k}$, any subsequence of $(\hat{\lambda}_n)$ having limit greater than λ^* can achieve W_1 convergence rate of the distance between $\hat{G}_n$ and $\overline{G}_*(\hat{\lambda}_n)$. If $K > k_0 + k_* - \bar{k}$, any subsequence of $(\hat{\lambda}_n)$ having limit greater than λ^* can achieve $W^{\overline{r}(K-k_0-k_*+\bar{k})}_{\overline{r}(K-k_0-k_*+\bar{k})}$ convergence rate of the distance between $\hat{G}_n$ and $\overline{G}_*(\hat{\lambda}_n)$, where $\overline{r}(K - k_0 - k_* + \bar{k})$ is smaller than $\overline{r}(K - k_*)$ in part (b).

4 Experiments

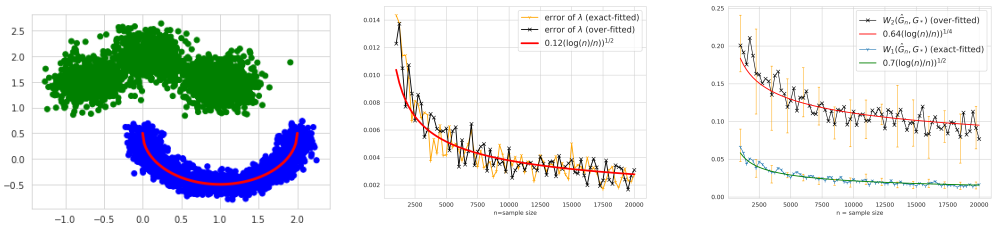
We now would like to demonstrate the convergence rates in Section 3 via two synthetic experiments: one for distinguishable setting and one for partially distinguishable setting. For the partially distinguishable one, the experiments are in Appendix B.

Distinguishable setting. We conduct an experiment where the original data distribution comes from an uniform distribution on a curve (half circle) in $\mathbb{R}^2$ convoluted with Gaussian noises (red curve and

blue points in Fig. 1(a)), and train a Normalizing Flow neural network [11] (Masked Autoregressive architecture) with 5 layers to get a good density estimation h_0 for this dataset. Then we assume that there are new data coming in, and the original distribution h_0 is deviated by a mixture of distributions in the location Gaussian family $f(x|\theta)$. So the true generating density now is

$$p_{\lambda^* G_*}(x) = (1-\lambda^*)h_0(x) + \lambda^* \sum_{i=1}^{3} p_i^* f(x|\theta_i^*), \tag{9}$$

where $\lambda^* = 0.5, G_* = \sum_{i=1}^{3} p_i^* \delta_{\theta_i^*}$, where $p_1^* = 0.3, p_2^* = 0.3, p_3^* = 0.4, \theta_1^* = (-0.7, 1.5), \theta_2^* = (0.1, 2.0), \theta_3^* = (1.0, 1.5)$. Samples from the deviated component are green points in Fig. 1(a). It can be seen from Proposition 2.4(a) that h_0 is distinguishable with family f. For each n, we simulate n data points from true model (9), estimate $\hat{\lambda}_n, \hat{G}_n$ by the EM algorithm (it is possible because Normalizing Flows provides exact density computation), and measure its convergence to the true λ^*, G_*. We conduct 16 replications for each sample size. The average error estimations with a 75% error bar can be seen in Fig. 1. The W_1 error in the exact-fitted case is of order $(\log(n)/n)^{1/2}$ and W_2 error in the over-fitted case is of order $(\log(n)/n)^{1/4}$. Meanwhile, thanks to the distinguishability, the estimation errors in both cases of λ are all of the order $(\log(n)/n)^{1/2}$. These simulation results are matched with the theoretical results found in Theorem 3.3 and Theorem 3.4. From the result, we see that the deviating mixture model successfully learns the deviated components and reuses the pre-trained black box model h_0, which helps to reduce computational costs.



(a) Synthetic data set (b) Convergence rates of $\hat{\lambda}_n$ (c) Convergence rates of $W(\hat{G}_n, G_*)$

Figure 1: Convergence rates for parameter estimation in the distinguishable case.

5 Discussion

In this work, we have presented the deviating mixture model and studied its parameter learning rates under MLE procedure. With a novel notion of distinguishability between distributions, we are able to prove inverse bounds for our model under several distinguishability settings, which allow us to deduce the parameter learning rates from the convergence rate of density functions. The distinguishability condition is shown to be satisfied for multiple families of distributions including those that come from black box models.

We now discuss practical implication of the theory. The deviating mixture model is designed to capture the deviated mixture components, and learning its parameters can reveal meaningful information about subpopulations in the data. When there is distinguishability in the model, our theory implies that we can learn the deviated proportion with the parametric rate and deviated components with a rate depending on the identifiable of f. However, our theory does not support using deviating mixture model when the existing distribution h_0 itself is a mixture distributions in family f and possesses parameters similar to deviated part, as the learning rate can be slow, and the deviated proportion estimator may not converge to the true value.

We believe that this work is the first attempt in the effort of understanding a broader class of mixture models combining with black box models, and its interpretability. There is room for future work building on top of this work. From a theoretical viewpoint, one can consider showing minimax lower bounds for the learning rates of the deviating mixture model, or show uniform inverse bounds for the model when λ^* and G_* are considered as signals that will change with samples. From a modeling viewpoint, it is potential to explore mixtures of black box models and develop a new suitable notion of identifiability and inverse bounds so that the learning process is efficient.

References

[1] Martin Arjovsky, Soumith Chintala, and Léon Bottou. Wasserstein generative adversarial networks. In *International conference on machine learning*, pages 214–223. PMLR, 2017.

[2] L. Bordes, S. Mottelet, and P. Vandekerkhove. Semiparametric estimation of a two-component mixture model. *Annals of Statistics*, 34, 2006.

[3] Cristina Butucea and Pierre Vandekerkhove. Semiparametric mixtures of symmetric distributions. *Scandinavian Journal of Statistics*, 41(1):227–239, 2014.

[4] T. Cai, X. J. Jeng, and J. Jin. Optimal detection of heterogeneous and heteroscedastic mixtures. *Journal of the Royal Statistical Society: Series B (Statistical Methodology)*, 73, 2011.

[5] T Tony Cai, Jiashun Jin, and Mark G Low. Estimation and confidence sets for sparse normal mixtures. *The Annals of Statistics*, 35(6):2421–2449, 2007.

[6] H. Chen and J. Chen. Tests for homogeneity in normal mixtures in the presence of a structural parameter. *Statistica Sinica*, 13:351–365, 2003.

[7] J. Chen and P. Li. Hypothesis test for normal mixture models: the em approach. *Annals of Statistics*, 37:2523–2542, 2009.

[8] J. Chen, P. Li, and Y. Fu. Inference on the order of a normal mixture. *Journal of the American Statistical Association*, 107:1096–1105, 2012.

[9] Laurent Dinh, Jascha Sohl-Dickstein, and Samy Bengio. Density estimation using real NVP. In *5th International Conference on Learning Representations, ICLR 2017, Toulon, France, April 24-26, 2017, Conference Track Proceedings*, 2017.

[10] D. Donoho and J. Jin. Higher criticism for detecting sparse heterogeneous mixtures. *Annals of Statistics*, 32, 2004.

[11] Conor Durkan, Artur Bekasov, Iain Murray, and George Papamakarios. nflows: normalizing flows in PyTorch, November 2020.

[12] Sébastien Gadat, Jonas Kahn, Clément Marteau, and Cathy Maugis-Rabusseau. Parameter recovery in two-component contamination mixtures: The $\theta\{2\}$ strategy. In *Annales de l'Institut Henri Poincaré, Probabilités et Statistiques*, volume 56, pages 1391–1418. Institut Henri Poincaré, 2020.

[13] Ian Goodfellow, Jean Pouget-Abadie, Mehdi Mirza, Bing Xu, David Warde-Farley, Sherjil Ozair, Aaron Courville, and Yoshua Bengio. Generative adversarial nets. *Advances in neural information processing systems*, 27, 2014.

[14] Aritra Guha, Nhat Ho, and XuanLong Nguyen. On posterior contraction of parameters and interpretability in bayesian mixture modeling. *Bernoulli*, 27(4):2159–2188, 2021.

[15] Philippe Heinrich and Jonas Kahn. Strong identifiability and optimal minimax rates for finite mixture estimation. *Annals of Statistics*, 46(6A):2844–2870, 2018.

[16] N. Ho and X. Nguyen. Convergence rates of parameter estimation for some weakly identifiable finite mixtures. *Annals of Statistics*, 44:2726–2755, 2016.

[17] N. Ho and X. Nguyen. On strong identifiability and convergence rates of parameter estimation in finite mixtures. *Electronic Journal of Statistics*, 10:271–307, 2016.

[18] N. Ho and X. Nguyen. Singularity structures and impacts on parameter estimation in finite mixtures of distributions. *arXiv:1609.02655*, 2016.

[19] J. Katz-Samuels, G. Blanchard, and C. Scott. Decontamination of mutual contamination models. *Journal of Machine Learning Research*, 20, 2019.

[20] Geoffrey J. McLachlan and David Peel. *Finite mixture models*, volume 299 of *Probability and Statistics – Applied Probability and Statistics Section*. Wiley, New York, 2000.

[21] X. Nguyen. Convergence of latent mixing measures in finite and infinite mixture models. *Annals of Statistics*, 4(1):370–400, 2013.

[22] Emanuel Parzen. On estimation of a probability density function and mode. *The annals of mathematical statistics*, 33(3):1065–1076, 1962.

[23] Rohit Kumar Patra and Bodhisattva Sen. Estimation of a two-component mixture model with applications to multiple testing. *Journal of the Royal Statistical Society: Series B (Statistical Methodology)*, 78(4):869–893, 2016.

[24] B. Schölkopf and A. Smola. *Learning with Kernels*. MIT Press, Cambridge, MA, 2002.

[25] C. Scott. A rate of convergence for mixture proportion estimation, with application to learning from noisy labels. In *AISTATS*, 2015.

[26] Sara van de Geer. *Empirical Processes in M-estimation*, volume 6. Cambridge university press, 2000.

[27] C. Villani. *Optimal Transport: Old and New. Grundlehren der Mathematischen Wissenschaften [Fundamental Principles of Mathemtical Sciences]*. Springer, Berlin, 2009.

[28] Sidney J Yakowitz and John D Spragins. On the identifiability of finite mixtures. *The Annals of Mathematical Statistics*, 39(1):209–214, 1968.

Checklist

1. For all authors...
   (a) Do the main claims made in the abstract and introduction accurately reflect the paper's contributions and scope? [Yes] See Section 1
   (b) Did you describe the limitations of your work? [Yes] See Section 5
   (c) Did you discuss any potential negative societal impacts of your work? [No]
   (d) Have you read the ethics review guidelines and ensured that your paper conforms to them? [Yes]
2. If you are including theoretical results...
   (a) Did you state the full set of assumptions of all theoretical results? [Yes]
   (b) Did you include complete proofs of all theoretical results? [Yes]
3. If you ran experiments...
   (a) Did you include the code, data, and instructions needed to reproduce the main experimental results (either in the supplemental material or as a URL)? [Yes]
   (b) Did you specify all the training details (e.g., data splits, hyperparameters, how they were chosen)? [Yes] It can be seen in the source code.
   (c) Did you report error bars (e.g., with respect to the random seed after running experiments multiple times)? [Yes] See Figure 1, 2, 3.
   (d) Did you include the total amount of compute and the type of resources used (e.g., type of GPUs, internal cluster, or cloud provider)? [No] The experiments are run on CPU's only.
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