# Learning mixtures of Gaussians

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III. Contributions to the Mathematical Theory of Evolution. By KARL PEARSON, University College, London. Communicated by Professor HENRICI, F.R.

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#### [Plates 1-5.]

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(9.) The whole method may be illustrated by the following numerical example:— Breadth of "Forehead" of Crabs.—Professor W. F. R. WELDON has very kindly given me the following statistics from among his measurements on crabs. They are for 1000 individuals from Naples. The abscisse of the curve are the ratio of "forehead" to body-length, and one unit of abscissa = 004 of body-length. No. 1 of the abscissae corresponds to  $\cdot580 - \cdot583$  of body-length. The ordinates represent the number of individual crabs corresponding to each set of ratios of forehead to bodylength. Thus there was one crab fell into the range  $\cdot580 - \cdot583$ , three fell into the range  $\cdot584 - \cdot587$ , five into the range  $\cdot588 - \cdot591$ , and so on. The average length of animals measured 35 millims, and measurements were recorded to  $\cdot1$  millim.

Abscissæ.	Ordinates.	Abscissæ.	Ordinates.
1	1	16	74
2 3	3	17	84
4	2	19	96
5	10	20 21	85 75
7	13	22	47
8	19	23	$43 \\ 24$
10	20	24	19
11	40	26	9
12 13	60	27	0 0
14	62	29	1
15	54	1	

Observation: data is asymmetric.

Hypothesis: may be a mixture of two Gaussians. Method: numerically matching the moments.



# Learning mixtures of Gaussians in modern times

These days trying to fit data with mixtures of Gaussians is popular in data science.

Modern applications: high-dimensional data



[Richardson and Weiss, Neurips 2018]

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Why mixtures of Gaussians?

- $\checkmark$  fit some natural data well
- $\checkmark$  universal approximators
- $\checkmark$  clustering

## High-dimensional Gaussians

Multivariate normal distribution:

$$\mathcal{N}_{\mu,\Sigma}(x) = rac{\exp\left(-rac{1}{2}(x-\mu)^{\mathsf{T}}\Sigma^{-1}(x-\mu)
ight)}{(2\pi)^{d/2}\sqrt{\det(\Sigma)}} \quad ext{ for } x \in \mathbb{R}^d$$
 $X \sim \mathcal{N}_{\mu,\Sigma}: \ \mathbb{E}\left[X
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Mixture of k Gaussians in  $\mathbb{R}^d$ :  $\sum_{i=1}^k \boldsymbol{w}_i \mathcal{N}_{\mu_i, \Sigma_i}$ mixture weights satisfy  $\boldsymbol{w}_i \geq 0$ ,  $\sum \boldsymbol{w}_i = 1$ Parameters of the model:  $(\boldsymbol{w}_i, \mu_i, \Sigma_i)_{i=1}^k$ :  $\Theta(kd^2)$  parameters







What does it mean to learn/estimate a mixture of Gaussians given data?

Given samples  $x_1, \ldots, x_n$ , find parameters that maximize the likelihood:

$$\prod_{i=1}^n \left(\sum_{j=1}^k w_j \mathcal{N}_{\mu_j, \Sigma_j}(x_i)\right)$$

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ight)$$

- $\checkmark$  Non-convex optimization problem, NP-hard
- ✓ Widely used in practice: expectation-maximization (EM) an iterative algorithm
- $\checkmark\,$  Convergence not well understood, very sensitive to initialization

## Second answer: parameter estimation

Given samples from some unknown mixture of Gaussians  $\sum_{i=1}^{k} w_i \mathcal{N}(\mu_i, \Sigma_i)$ , find each of the parameters within  $\varepsilon$ .

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- ✓ Active area of research in theoretical computer science [Dasgupta 1999]
- ✓ Computational complexity: polynomial in d and 1/ε [Kalai, Moitra, Valiant 2010] [Belkin, Sinha 2010]
- ✓ Any algorithm has sample complexity exponential in k
   [Moitra, Valiant 2010]

Given samples from an unknown mixture of Gaussians f, output a density  $\hat{f}$  that is close to f with high probability, 99%.

Close in  $L^1$  distance:  $\left\|f - \widehat{f}\right\|_1 = \int_{\mathbb{R}^d} \left|f(x) - \widehat{f}(x)\right| dx = 2 \sup_{A \subseteq \mathbb{R}^d} \left|\int_A f - \int_A \widehat{f}\right|$ 



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Bounds for parameter estimation do not translate to bounds for density estimation: zero-mean 2-dimensional Gaussians with

$$\Sigma_1=egin{pmatrix}1&-0.99\-0.99&1\end{pmatrix}$$
 and  $\Sigma_2=egin{pmatrix}1&-1\-1&1\end{pmatrix}$ 

Close parameters, large  $L^1$  distance.

## Density estimation

Given samples from an unknown density f from some known family C of densities, output a density  $\hat{f}$  that is close to f.

#### REMARKS ON SOME NONPARAMETRIC ESTIMATES OF A DENSITY FUNCTION<sup>1</sup>

By Murray Rosenblatt<sup>2</sup>

University of Chicago



#### On the Learnability of Discrete Distributions

(EXTENDED ABSTRACT)

Michael Kearns AT&T Bell Laboratories

> Ronitt Rubinfeld Cornell University

Yishay Mansour Tel-Aviv University

Robert E. Schapire AT&T Bell Laboratories

Dana Ron Hebrew University

Linda Sellie University of Chicago

# Precise question we study today

#### Question

Let f be an unknown mixture of k Gaussians in  $\mathbb{R}^d$ . How many i.i.d. samples from f is needed to produce, with high probability, a density  $\widehat{f}$  satisfying  $\|f - \widehat{f}\|_1 \leq \varepsilon$ ?

Remarks:

- 1. Algorithm knows k
- 2. Focus is on sample complexity
- 3. Equivalent formulation: given n samples from  $f \in C$ , how small can you make  $\mathbb{E}\left[\|f \widehat{f}\|_1\right]$ ? Minimax risk
- 4. Unbounded for  $L^{p>1}$  or KL

# Popular method in practice for density estimation Kernel density estimation



Unfortunately, sample complexity is exponential in d.

#### Question

Let f be an unknown mixture of k Gaussians in  $\mathbb{R}^d$ . How many i.i.d. samples from f is needed to produce, with high probability, a density  $\hat{f}$  satisfying  $\|f - \hat{f}\|_1 \leq \varepsilon$ ?

- k = 1: sample complexity  $\leq O(d^2/\varepsilon^2)$ compute empirical mean and covariance, and use Gaussian concentration
- d=1: sample complexity  $\leq O(k/\epsilon^2)$ approximate by piecewise polynomials [Chan, Diakonikolas, Servedio, Sun 2014]

#### Question

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$$k = 1$$
: sample complexity  $\leq O(d^2/\epsilon^2)$   
compute empirical mean and covariance, and use Gaussian  
concentration

$$d=1$$
: sample complexity  $\leq O(k/\epsilon^2)$   
approximate by piecewise polynomials  
[Chan, Diakonikolas, Servedio, Sun 2014]

Question: sample complexity  $\leq$  number of parameters divided by  $\epsilon^2$ ? Indeed we will show sample complexity  $\leq kd^2/\epsilon^2 \times \log^2(d)\log(k) = \widetilde{O}(kd^2/\epsilon^2)$ 

#### Definition

Given an i.i.d. sample from an unknown density  $f \in C$ , output  $\hat{f}$  satisfying  $||f - \hat{f}||_1 \leq \varepsilon$  with high probability.  $m_{\mathcal{C}}(\varepsilon) =$  the smallest number of required samples.  $k-\min(\mathcal{C}) =$  class of distributions formed by taking k-mixtures of elements of C

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#### Theorem (Ashtiani, Ben-david, M2017)

For any class C, sample complexity for learning k-mix $(C) \leq O(m_{\mathcal{C}}(\varepsilon) \times k \log k / \varepsilon^2)$ 

#### Corollary

 $\begin{array}{l} \textit{Sample complexity for learning mixtures of Gaussians} \\ \leq O((d^2/\varepsilon^2) \times k \log k/\varepsilon^2) = O(kd^2 \log(k)/\varepsilon^4) \end{array}$ 

$$Y(\mathcal{C}) = \left\{ \{x \in \mathbb{R}^d : f_1(x) > f_2(x)\} : f_1, f_2 \in \mathcal{C} 
ight\}$$

1.  $m_{\mathcal{C}}(\varepsilon) \leq O(\text{VC-dim}(Y(\mathcal{C}))/\varepsilon^2)$  [Devroye and Lugosi 2001]

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- 1.  $m_{\mathcal{C}}(\varepsilon) \leq O(\text{VC-dim}(Y(\mathcal{C}))/\varepsilon^2)$  [Devroye and Lugosi 2001]
- 2. When C = mixtures of Gaussians, VC-dim $(Y(C)) \leq O(k^4 d^4)$  [Khovanskii 1991], [Karpinski and Macintyre 1997], [Anthony and Bartlett 1999]
- 3. Gives an upper bound of  $O(k^4 d^4/\epsilon^2)$  for the sample complexity of mixtures of Gaussians.

We will improve this to  $kd^2/\varepsilon^2$ .

Lower bounds

## Lower bounds?

Best known lower bound was  $\Omega(kd/\epsilon^2)$ . [Suresh, Orlitsky, Acharya, and Jafarpour 2014]

Theorem (Ashtiani, Ben-David, Harvey, Liaw, M, Plan'18)

Any algorithm that learns mixtures of Gaussians has sample complexity  $\Omega(kd^2/\epsilon^2)$ .

Suffices to show lower bound of  $\Omega(d^2/\epsilon^2)$  for a single Gaussian.

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General idea: find lots of distributions that are hard to distinguish but far in  $L^1$  distance. [LeCam 1973], [Hasminskii 1976], [Assoud 1983]

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General idea: find lots of distributions that are hard to distinguish but far in  $L^1$  distance. [LeCam 1973], [Hasminskii 1976], [Assoud 1983]

Hasminskii+Fano's inequality: find  $2^{\Omega(d^2)}$  Gaussians with pairwise KL-divergence  $\leq \varepsilon^2$  and pairwise  $L^1$  distance  $> \varepsilon$ .

$$ext{KL}(f_1 \parallel f_2) \coloneqq \int f_1(x) \log rac{f_1(x)}{f_2(x)} ext{d}x \qquad [Kullback-Leibler]$$

Need to build  $2^{\Omega(d^2)}$  Gaussians with pairwise KL-divergence  $\leq \epsilon^2$  and pairwise  $L^1$  distance  $> \epsilon$ .

We will use zero-mean Gaussians, so just need to specify the covariance matrices.

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First construction [Ashtiani, Ben-David, Harvey, Liaw, M, Plan'18]. Repeat  $2^{d^2}$  times: start with an identity covariance matrix, then choose a random subspace of dimension d/9 and slightly increase the eigenvalues corresponding to this eigenspace:  $\Sigma = I + \frac{\varepsilon}{\sqrt{d}} UU^{\mathsf{T}}$ , with  $U \in \mathbb{R}^{d \times d/9}$  orthonormal. Then prove that with large probability, any two of these have  $L^1$  distance  $> \varepsilon$ .

Need to build  $2^{\Omega(d^2)}$  Gaussians with pairwise KL-divergence  $\leq \epsilon^2$  and pairwise  $L^1$  distance  $> \epsilon$ .

We will use zero-mean Gaussians, so just need to specify the covariance matrices.

Second construction (combinatorial) [Devroye, M, Reddad 2018]. For d = 3, consider the following inverse covariance matrices:

$$\begin{pmatrix} 0 & -\delta & -\delta \\ -\delta & 0 & -\delta \\ -\delta & -\delta & 0 \end{pmatrix}, \begin{pmatrix} 0 & \delta & \delta \\ \delta & 0 & -\delta \\ \delta & -\delta & 0 \end{pmatrix}, \begin{pmatrix} 0 & \delta & -\delta \\ \delta & 0 & \delta \\ -\delta & \delta & 0 \end{pmatrix}, \begin{pmatrix} 0 & -\delta & \delta \\ -\delta & 0 & \delta \\ \delta & \delta & 0 \end{pmatrix}$$

For general d, build  $2^{d^2/10}$  inverse covariance matrices so that any two of them are different in at least  $d^2/3$  coordinates (Gilbert-Varshamov bound in coding theory).

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Next: upper bound

# Our upper bound

#### $m_{\mathcal{C}}(\varepsilon) = \text{sample complexity for learning a density from class } \mathcal{C}.$

Theorem (Ashtiani, Ben-David, Harvey, Liaw, M, Plan'18)

If C = mixtures of k Gaussians in  $\mathbb{R}^d$ , then  $m_{\mathcal{C}}(\varepsilon) = \widetilde{O}(kd^2/\varepsilon^2).$ 

# Covering number argument

#### Lemma (Yatracos 1985)

Suppose there exist  $f_1, \ldots, f_M \in C$  such that for any  $f \in C$ , there exists some *i* with  $||f - f_i||_1 \leq \varepsilon$ . Then  $m_{\mathcal{C}}(\varepsilon) = O(\log(M)/\varepsilon^2)$ .

Proved by a clever combination of Hoeffding's inequality and the union bound.

## Covering number argument

## Lemma (Yatracos 1985)

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A bound on the covering number of a distribution class bounds its sample complexity.

covering number =  $\varepsilon$ -net number = packing number =

 $\varepsilon$ -Kolmogorov entropy= metric entropy

## Gaussians are not bounded

Unfortunately, Gaussian distributions have infinite covering number, even if the mean is bounded.



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Our novel idea to solve this: Use some of the data to reduce the search space significantly. To formalize this idea, we introduce the notion of compression.

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$$\left\| N(\widehat{\mu},\widehat{\sigma}) - N(\mu,\sigma) \right\|_{1} \leq \varepsilon$$



$$\left\| N(\widehat{\mu},\widehat{\sigma}) - N(\mu,\sigma) \right\|_{1} \leq \varepsilon$$

One-dimensional Gaussians admit  $(100/\varepsilon, 2)$ -compression.

# Compression implies learnability

#### Definition (compression)

Class C admits  $(n(\varepsilon), \tau(\varepsilon))$ -compression if, for any  $f \in C$ , after  $n(\varepsilon)$ i.i.d. samples from f are generated, with high probability there exist  $\tau(\varepsilon)$  of the samples from which  $\hat{f}$  can be constructed satisfying  $\|f - \hat{f}\|_1 \leq \varepsilon$ .

## Lemma (compression implies learnability)

If C admits  $(n, \tau)$ -compression,  $m_{\mathcal{C}}(\varepsilon) = O\left(n + \frac{\tau \log n}{\varepsilon^2}\right)$ .

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Algorithm: Exhaustive search + Yatracos' algorithm

Running time:  $n^{\tau}$ 

1. Compressing *d*-dimensional Gaussians

d-dimensional Gaussians admit  $\widetilde{O}(d, d^2)$ -compression.

#### 2. Compressing mixtures

If C admits  $(n, \tau)$ -compression, then k-mix(C) admits  $\widetilde{O}(kn, k\tau)$ -compression.

3. Compression implies learnability

If C admits  $(n, \tau)$ -compression,  $m_{\mathcal{C}}(\varepsilon) = \widetilde{O}(n + \tau/\varepsilon^2)$ .

Theorem (Ashtiani, Ben-David, Harvey, Liaw, M, Plan'18)

If C is mixtures of k Gaussians in  $\mathbb{R}^d$  then  $m_{\mathcal{C}}(\varepsilon) = \widetilde{O}(kd^2/\varepsilon^2).$ 

1. Compressing *d*-dimensional Gaussians

d-dimensional Gaussians admit  $\widetilde{O}(d, d^2)$ -compression.

$$\mathcal{N}(\mu, \Sigma) = \mathcal{N}(\mu, v_1 v_1^\mathsf{T} + v_2 v_2^\mathsf{T}).$$



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1. Compressing *d*-dimensional Gaussians

d-dimensional Gaussians admit  $\widetilde{O}(d, d^2)$ -compression.

In general, use  $d \log^2(d/\varepsilon)$  data points to encode the mean, and  $d \log^2(d/\varepsilon)$  data points for each eigenvector.

Lemma (Litvak, Pajor, Rudelson, Tomczak-Jaegermann 2005)

If we take  $O(d \log d)$  samples from  $\mathcal{N}(0, I_d)$ , their convex hull with high probability contains  $\frac{1}{20}B_2^d$ 

# Main result

Theorem (Ashtiani, Ben-David, Harvey, Liaw, M, Plan, NeurIPS 2018)

Given  $\widetilde{O}(kd^2/\epsilon^2)$  samples from an unknown mixture of k Gaussians in d dimensions, we can output a density that is  $\epsilon$ -close in  $L^1$  to the underlying density with high probability. Moreover, any algorithm achieving this task requires at least  $\Omega(kd^2/\epsilon^2)$  many samples.

improve previous upper bounds of  $\widetilde{O}(kd^2/\varepsilon^4)$  and  $O(k^4d^4/\varepsilon^2)$ , and the lower bound of  $\Omega(kd/\varepsilon^2)$ .

Upper bound. a novel technique for distribution learning based on compressions, high-dimensional geometry + Yatracos' algorithm.

Lower bound. a packing argument, Fano's inequality.

Open questions



- 1. Polynomial time algorithm?
- 2. What if k is not known?
- 3. Sample complexity for general classes?

## Thanks to my co-authors



Hassan Ashtiani (McMaster)



Shai Ben-David (Waterloo)



Nick Harvey (UBC)



Chris Liaw (UBC)



Yaniv Plan (UBC)



## **Research direction 1**

What is the sample complexity for learning a class C?

- $\checkmark\,$  Relate this to some notion of dimension of the class?
- $\checkmark\,$  Apply the compression idea to other classes?
- ✓ Probabilistic graphical models, e.g. the Ising model [Devroye, M, Reddad'18]
- $\checkmark$  Distributions generated by neural networks



Picture taken from the work of Karras, Aila, Laine, and Lehtinen 2017

Research direction 2: computational complexity

Which classes are learnable in polynomial time?

Polynomial time algorithm for mixtures of Gaussians?

Exists for mixtures of spherical Gaussians. [Suresh, Orlitsky, Acharya, and Jafarpour 2014] Research direction 2: computational complexity

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Research direction 3: robustness

Design learners that are robust against noisy data.

Our algorithm works in agnostic learning.

What if a small fraction of the data is corrupted in an adversarial way?

#### Research direction 4: online learning

What if data is not revealed at once, but is received in an online manner? Can we compete against a batch algorithm that sees all the data at once?

Research direction 5: model selection

Can we learn the class C itself from data?

What if the number of Gaussian components, k, is not known?

# Popular method in practice for density estimation Kernel density estimation-continued

Table 4.2 Sample size required (accurate to about 3 significant figures) to ensure that the relative mean square error at zero is less than 0.1, when estimating a standard multivariate normal density using a normal kernel and the window width that minimizes the mean square error at zero

Dimensionalit y	Required sample size
1	4
2	19
3	67
4	223
5	768
6	2 790
7	10 700
8	43 700
9	187 000
10	842 000

Silverman 1998, Density estimation for Statistics and Data Analysis

## **VC-dimension**

For a family  $\mathcal{Y}$  of subsets of X, the VC-dimension of  $\mathcal{Y}$  is the size of the largest set  $A \subseteq X$ , such that for any  $B \subseteq A$  there exists some  $Y \in \mathcal{Y}$  with  $Y \cap A = B$ .

Interesting classes - 1 Probabilistic graphical models



Example (The Ising model). Each  $X_i \in \{-1, +1\}$  and

$$\mathbb{P}\left[X_1 = x_1, \ldots, X_d = x_d
ight] \propto \exp\left(\sum_{ij \in E(G)} w_{i,j} x_i x_j
ight)$$

## Theorem (Devroye, M, Reddad'18)

Let  $\mathcal{I}_G = Ising models on G$ . Then,  $m_{\mathcal{I}_G}(\varepsilon) = \Theta(|E(G)|/\varepsilon^2)$ .

# Interesting classes - 2





[Karras, Aila, Laine, and Lehtinen 2017]

# Proof of mixture lemma

#### Compressing mixtures

If C admits  $(n, \tau)$ -compression, then k-mix(C) admits  $(nk \log(k), k\tau + k \log k)$ -compression.

Let  $\mathbf{P} = \frac{1}{3}P_1 + \frac{1}{3}P_2 + \frac{1}{3}P_3$ , where each  $P_i$  is 2-compressible.

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Let  $\mathbf{P} = \frac{1}{3}P_1 + \frac{1}{3}P_2 + \frac{1}{3}P_3$ , where each  $P_i$  is 2-compressible.



## Lemma (Yatracos 1985)

Suppose there exist  $f_1, \ldots, f_M \in C$  such that for any  $f \in C$ , there exists some *i* with  $||f - f_i||_1 \leq \varepsilon/5$ . Then  $m_{\mathcal{C}}(\varepsilon) = O(\log(M)/\varepsilon^2)$ .

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Let  $Y \coloneqq \left\{ \{x : f_i(x) > f_j(x)\} \text{ for } i, j = 1, \dots, m \right\}$  and let S be an i.i.d. sample of size  $50 \log(M) / \varepsilon^2$  from f. For density f, let  $f(A) \coloneqq \int_A f$ .  $|S \cap A| \sim \text{binomial}(|S|, f(A))$ . By Hoeffding [1963] and a union bound over  $A \in Y$ ,

$$\operatorname{err}(f) \coloneqq \sup_{A \in Y} \left| f(A) - \frac{|S \cap A|}{|S|} \right| \le \varepsilon/5$$

with probability  $1 - 2M^2 \exp(-|S|\epsilon^2/25) \ge 99\%$ . Thus there exists some *i* with  $\operatorname{err}(f_i) \le 2\epsilon/5$ . So  $\min_j \operatorname{err}(f_j) \le 2\epsilon/5$ , and it can be shown that the argmin here is within L1 distance  $\epsilon$  of f.

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$$\min_{j=1,...,M} \sup_{A \in Y} \left| f_j(A) - \frac{|S \cap A|}{|S|} \right|$$

# An application of density estimation detecting breast cancer

- ✓ Training data consists of normal (non-cancerous) X-ray images.
- $\checkmark$  A probability density function  $f: \mathbb{R}^d \to \mathbb{R}$  is learned
  from the data.
- $\checkmark$  When a new input x is presented, a high value for f(x) indicates a normal image, while a low value indicates a novel input, which might be characteristic of an abnormality.

[Tarassenko, Hayton, Cerneaz, Brady 1995: Novelty detection for the identification of masses in mammograms]

An example of density estimation Generating random faces for computer games

- $\checkmark\,$  Training data consists of actual faces.
- $\checkmark\,$  A probability density function  $f:\mathbb{R}^d\to\mathbb{R}$  is learned from the data.
- $\checkmark\,$  New random faces are generated using the learned distribution.

An example of density estimation Generating random faces for computer games

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A popular approach: generative adversarial networks (GANs), based on deep neural networks.

## Density estimation in action



## Top: generated images using generative adversarial networks Bottom: a small part of the training data

Picture from Karras, Aila, Laine, and Lehtinen (NVIDIA and Aalto University), October 2017