

CSCI567 Machine Learning (Spring 2021)

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Logistics

Outline

- 1 Logistics
- 2 Review of last lecture
- 3 Density estimation

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Logistics

Logistics

- Today is the tracking day for the project.
- When you submit course deliverables, check them to ensure you have uploaded the correct files.
- In **April 23, 2021**'s lecture we will have **Alice Xiang**, *Senior Research Scientist and AI Ethics Lead at Sony AI* to talk about **Fairness in AI**. If possible, please plan to join the class live!

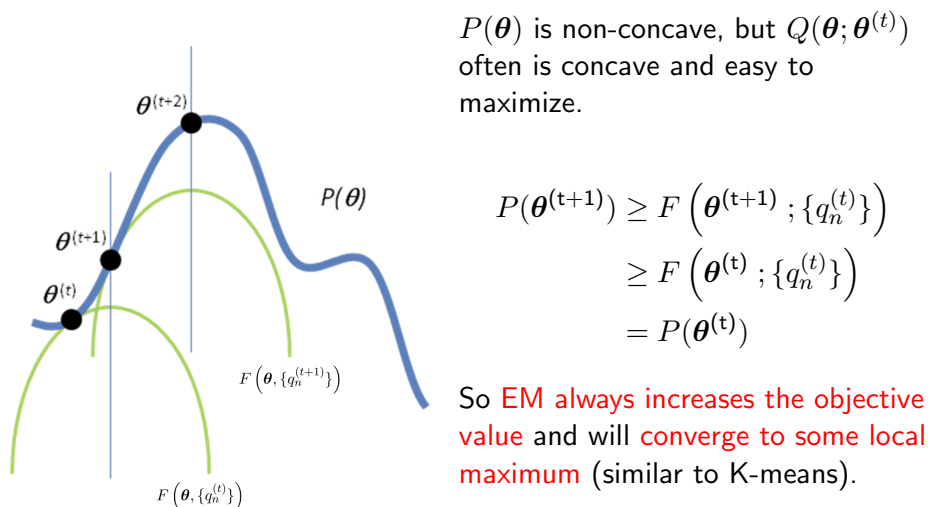
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Pictorial explanation



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General EM algorithm

Step 0 Initialize $\theta^{(1)}$, $t = 1$

Step 1 (E-Step) update the posterior of latent variables

$$q_n^{(t)}(\cdot) = p(\cdot | \mathbf{x}_n; \theta^{(t)})$$

and obtain **Expectation** of complete likelihood

$$Q(\theta; \theta^{(t)}) = \sum_{n=1}^N \mathbb{E}_{z_n \sim q_n^{(t)}} [\ln p(\mathbf{x}_n, z_n; \theta)]$$

Step 2 (M-Step) update the model parameter via **Maximization**

$$\theta^{(t+1)} \leftarrow \operatorname{argmax}_{\theta} Q(\theta; \theta^{(t)})$$

Step 3 $t \leftarrow t + 1$ and return to Step 1 if not converged

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Applying EM to learn GMMs

EM for clustering:

Step 0 Initialize $\omega_k, \mu_k, \Sigma_k$ for each $k \in [K]$

Step 1 (E-Step) update the “soft assignment” (fixing parameters)

$$\gamma_{nk} = p(z_n = k | \mathbf{x}_n) \propto \omega_k N(\mathbf{x}_n | \mu_k, \Sigma_k)$$

Step 2 (M-Step) update the model parameter (fixing assignments)

$$\omega_k = \frac{\sum_n \gamma_{nk}}{N} \quad \mu_k = \frac{\sum_n \gamma_{nk} \mathbf{x}_n}{\sum_n \gamma_{nk}}$$

$$\Sigma_k = \frac{1}{\sum_n \gamma_{nk}} \sum_n \gamma_{nk} (\mathbf{x}_n - \mu_k)(\mathbf{x}_n - \mu_k)^T$$

Step 3 return to Step 1 if not converged

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- 1 Logistics
- 2 Review of last lecture
- 3 Density estimation
 - Parametric methods
 - Nonparametric methods

Density estimation

Observe what we have done indirectly for clustering with GMMs:

Given a training set x_1, \dots, x_N , **estimate a density function p that could have generated this dataset** (via $x_n \stackrel{i.i.d.}{\sim} p$).

This is exactly the problem of *density estimation*, another important unsupervised learning problem.

Useful for many downstream applications

- we have seen clustering already, will see more today
- these applications also *provide a way to measure quality of the density estimator*

Parametric methods: generative models

Parametric estimation assumes a generative model parametrized by θ :

$$p(x) = p(x; \theta)$$

Examples:

- **GMM**: $p(x | \theta) = \sum_{k=1}^K \omega_k N(x | \mu_k, \Sigma_k)$ where $\theta = \{\omega_k, \mu_k, \Sigma_k\}$
- **Multinomial**: a discrete variable with values in $\{1, 2, \dots, K\}$ s.t.

$$p(x = k; \theta) = \theta_k$$

where θ is a distribution over K elements.

Size of θ is independent of the training set size, so it's *parametric*.

Parametric methods: estimation

Again, we apply **MLE** to learn the parameters θ :

$$\operatorname{argmax}_{\theta} \sum_{n=1}^N \ln p(x_n; \theta)$$

For some cases this is intractable and we can use **EM** to approximately solve MLE (e.g. GMMs).

For some other cases this admits a *simple closed-form solution* (e.g. multinomial).

MLE for multinomial

The log-likelihood is

$$\begin{aligned}\sum_{n=1}^N \ln p(x = x_n; \theta) &= \sum_{n=1}^N \ln \theta_{x_n} \\ &= \sum_{k=1}^K \sum_{n: x_n=k} \ln \theta_k = \sum_{k=1}^K z_k \ln \theta_k\end{aligned}$$

where $z_k = |\{n : x_n = k\}|$ is **the number of examples with value k** .

The solution is simply

$$\theta_k = \frac{z_k}{N} \propto z_k,$$

i.e. **the fraction of examples with value k** .

Nonparametric methods

Can we estimate *without assuming a fixed generative model*?

Yes, **kernel density estimation (KDE)** is a common approach

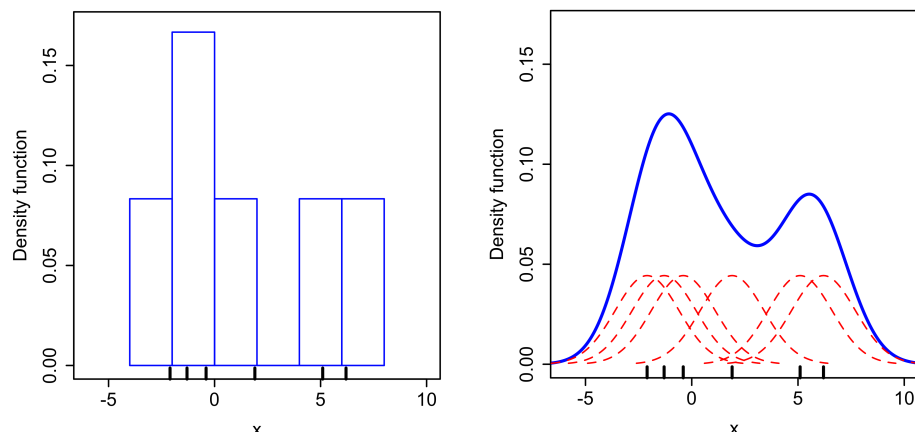
- here “**kernel**” means something different from what we have seen for “**kernel function**” (in fact it refers to several different things in ML)
- the approach is **nonparametric**: it keeps the entire training set
- we focus on the 1D (continuous) case

High level idea

picture from Wikipedia

Construct something similar to a **histogram**:

- for each data point, create a “**bump**” (via a Kernel)
- sum up or average all the bumps



Kernel

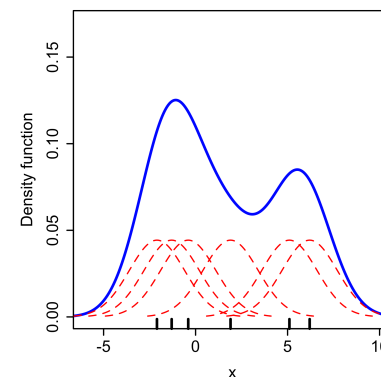
KDE with a **kernel K** : $\mathbb{R} \rightarrow \mathbb{R}$:

$$p(x) = \frac{1}{N} \sum_{n=1}^N K(x - x_n)$$

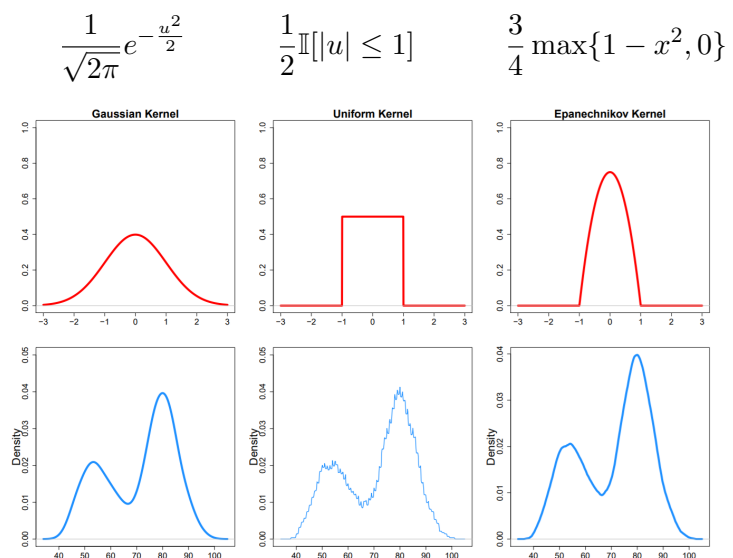
e.g. $K(u) = \frac{1}{\sqrt{2\pi}} e^{-\frac{u^2}{2}}$, the **standard Gaussian density**

Kernel needs to satisfy:

- **symmetry**: $K(u) = K(-u)$
- $\int_{-\infty}^{\infty} K(u) du = 1$, makes sure **p is a density function**.



Different kernels $K(u)$



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Bandwidth

If $K(u)$ is a kernel, then for any $h > 0$

$$K_h(u) \triangleq \frac{1}{h} K\left(\frac{u}{h}\right) \quad (\text{stretching the kernel})$$

can be used as a kernel too (verify the two properties yourself)

So general KDE is determined by both the kernel K and the bandwidth h

$$p(x) = \frac{1}{N} \sum_{n=1}^N K_h(x - x_n) = \frac{1}{Nh} \sum_{n=1}^N K\left(\frac{x - x_n}{h}\right)$$

- x_n controls the center of each bump
- h controls the width/variance of the bumps

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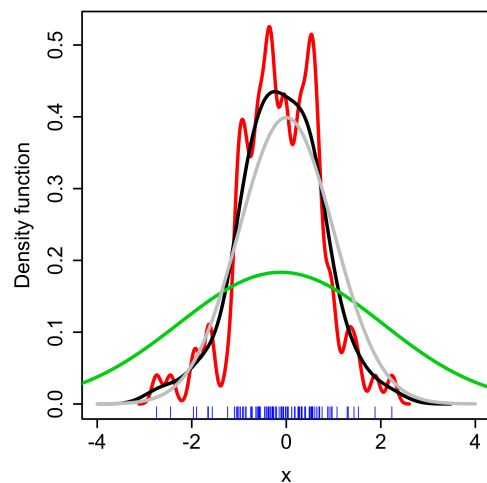
Effect of bandwidth

picture from Wikipedia

Larger h means larger variance and also smoother density

Gray curve is ground-truth

- Red: $h = 0.05$
- Black: $h = 0.337$
- Green: $h = 2$



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Bandwidth selection

For selecting h

- there are theoretically-motivated approaches
- one can also do cross-validation based on downstream applications

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