

Probabilistic Models and Machine Learning



No.4

Generative Models (I): Bayesian Decision Theory

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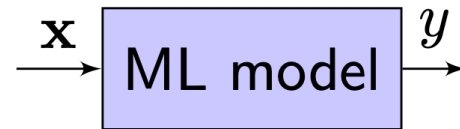
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Outline

- **Discriminative vs. Generative models**
 - **Generative modeling: a statistical perspective to ML**
- **Bayesian decision theory**
 - **Generative models for classification**
 - **Generative models for regression**
- **The Plug-in MAP rule**
- **Some probabilistic models for generative modeling**

Discriminative Models in ML

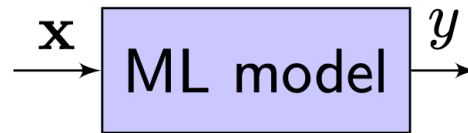


- ▶ Input \mathbf{x} is a random vector, $\mathbf{x} \sim p(\mathbf{x})$
- ▶ Output y is generated by a *deterministic target function* $y = \bar{f}(\mathbf{x})$ for each \mathbf{x}
- ▶ Our goal: estimate $\bar{f}(\cdot)$ in a model space \mathbb{H}
- ▶ Training samples: $\mathcal{D}_N = \{(\mathbf{x}_1, y_1), (\mathbf{x}_2, y_2), \dots, (\mathbf{x}_N, y_N)\}$, where $\mathbf{x}_i \sim p(\mathbf{x})$ and $y_i = \bar{f}(\mathbf{x}_i)$
- ▶ Determine a loss function $l(l, l')$
- ▶ Empirical risk minimization (ERM):

$$f^* = \arg \min_{f \in \mathbb{H}} R_{\text{emp}}(f | \mathcal{D}_N) = \arg \min_{f \in \mathbb{H}} \sum_{i=1}^N l(y_i, f(\mathbf{x}_i))$$

- ▶ The performance depends on the generalization bound

Generative Models in ML

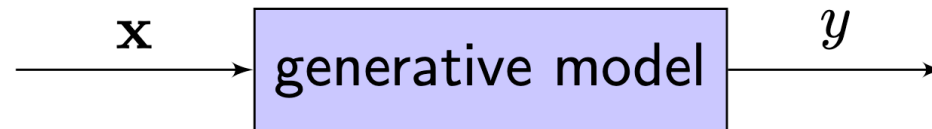


- ▶ Input \mathbf{x} and output y are both random variables, $(\mathbf{x}, y) \sim p(\mathbf{x}, y)$
- ▶ The relation $\mathbf{x} \rightarrow y$ solely relies on $p(y|\mathbf{x})$
- ▶ Our goal: estimate $p(\mathbf{x}, y)$ using a probabilistic model $\hat{p}_\theta(\mathbf{x}, y)$
- ▶ Training samples: $\mathcal{D}_N = \{(\mathbf{x}_1, y_1), (\mathbf{x}_2, y_2), \dots, (\mathbf{x}_N, y_N)\}$, where $(\mathbf{x}_i, y_i) \sim p(\mathbf{x}, y)$
- ▶ The relation $\mathbf{x} \rightarrow y$ may be approximated by:

$$\hat{p}_\theta(y|\mathbf{x})$$

- ▶ The performance depends on the gap between $p(\mathbf{x}, y)$ and $\hat{p}_\theta(\mathbf{x}, y)$: $\text{KL}(p(\cdot) \parallel \hat{p}_\theta(\cdot))$

Generative Models for Classification



- ▶ Input \mathbf{x} : feature vectors (continuous or discrete)
- ▶ Output is discrete $y = \{\omega_1, \omega_2, \dots, \omega_K\}$: class label
- ▶ The joint distribution $p(\mathbf{x}, y) = p(y)p(\mathbf{x}|y)$ breaks down to:
 - ▶ Prior probabilities: $p(y = \omega_k) \triangleq \Pr(\omega_k) \ (\forall k = 1, 2, \dots, K)$
 - ▶ Class-conditional distribution: $p(\mathbf{x}|y = \omega_k) \triangleq p(\mathbf{x}|\omega_k) \ (\forall k = 1, 2, \dots, K)$
- ▶ Probabilistic distribution constraints:
 - ▶ Priors satisfy $\sum_{k=1}^K \Pr(\omega_k) = 1$
 - ▶ If \mathbf{x} is continuous,

$$\int_{\mathbf{x}} p(\mathbf{x}|\omega_k) d\mathbf{x} = 1 \quad (\forall k = 1, 2, \dots, K)$$

- ▶ If \mathbf{x} is discrete,

$$\sum_{\mathbf{x}} p(\mathbf{x}|\omega_k) = 1 \quad (\forall k = 1, 2, \dots, K)$$

Example of class-conditional p.d.f.

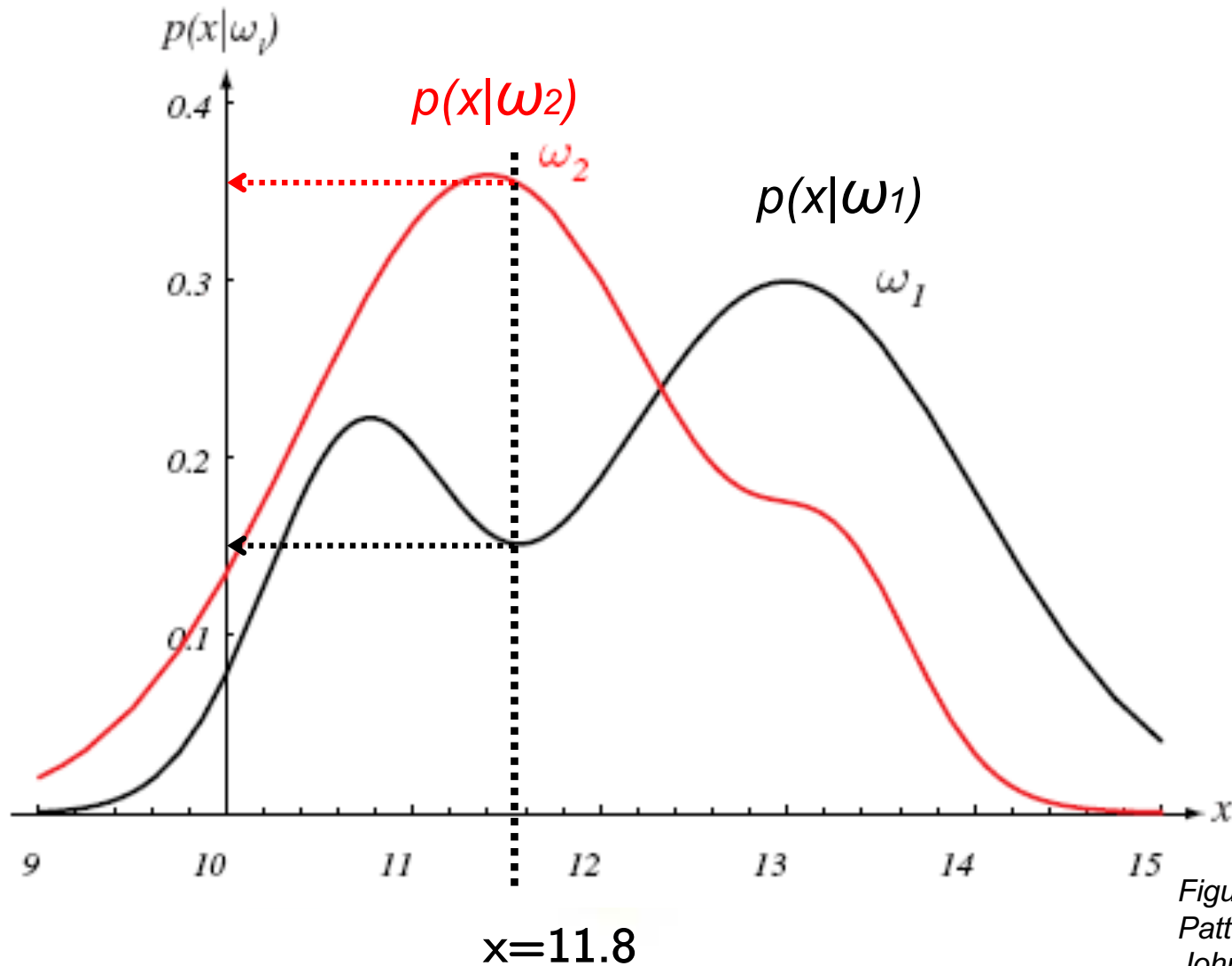


Figure from Duda et. al.,
Pattern classification
John Wiley & Sons®, Inc.

Examples of pattern classification(I)

- **Speech recognition:**
 - **Pattern:** voice spoken by a human being
 - **Classes:** language words/sentences used by the speaker
 - **Input features:** speech signal characteristics measured by a microphone → a sequence of feature vectors
 - **Each vector:** continuous, high-dimensional, real-valued numbers
- **Natural language understanding:**
 - **Pattern:** written or spoken languages of human
 - **Classes:** all possible semantic meanings or intentions
 - **Input features:** the used words or word-sequences (sentences)
 - **Discrete, scalars or vector**

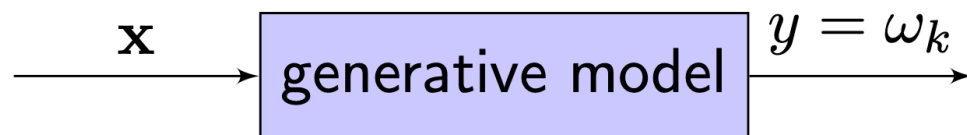


Examples of pattern classification(II)

- **Image understanding:**
 - **Pattern:** given images
 - **Classes:** all known object categories
 - **Input features:** color or gray scales in all pixels
 - **Continuous, multiple vectors/matrix**
 - **Examples:** face recognition, OCR (optical character recognition).
- **Gene finding in bioinformatics:**
 - **Pattern:** a newly sequenced DNA sequence
 - **Classes:** all known genes
 - **Input features:** all nucleotides in the sequence
 - **Discrete; 4 types (adenine, guanine, cytosine, thymine)**
- **Protein classification in bioinformatics:**
 - **Pattern:** protein primary 1-D sequence
 - **Classes:** all known protein families or domains
 - **Input features:** all amino acids in the sequence: discrete; 20 types



Bayesian Decision Theory (I): Classification



- ▶ Given any \mathbf{x} , determine the best $g(\mathbf{x}) \in \{\omega_1, \dots, \omega_K\}$
- ▶ The decision rule: $\mathbf{x} \Rightarrow \omega_k \quad (\forall k = 1, 2, \dots, K)$
- ▶ Bayesian Decision Theory: the best decision is

$$\begin{aligned} g^*(\mathbf{x}) &= \arg \max_k p(\omega_k | \mathbf{x}) = \arg \max_k \frac{\Pr(\omega_k) p(\mathbf{x} | \omega_k)}{p(\mathbf{x})} \\ &= \arg \max_k \Pr(\omega_k) \cdot p(\mathbf{x} | \omega_k) \end{aligned}$$

which is called **maximum a posterior (MAP) rule** or Bayes decision rule.

- ▶ Proof: why this is optimal?

Optimality of the MAP rule (I)

Theorem 1

Assume $p(\mathbf{x}, \omega)$ is known, when \mathbf{x} is used to predict ω , the MAP rule leads to the lowest expected risk (using 0-1 loss).

Proof:

- ▶ The 0-1 loss function: $l(\omega, \omega') = \begin{cases} 0 & \text{when } \omega = \omega' \\ 1 & \text{otherwise} \end{cases}$
- ▶ The expected risk of any rule $\mathbf{x} \rightarrow g(\mathbf{x}) \in \{\omega_1, \dots, \omega_K\}$:

$$\begin{aligned} R(g) &= \mathbb{E}_{p(\mathbf{x}, \omega)} [l(\omega, g(\mathbf{x}))] = \int_{\mathbf{x}} \sum_{k=1}^N l(\omega_k, g(\mathbf{x})) p(\mathbf{x}, \omega_k) d\mathbf{x} \\ &= \int_{\mathbf{x}} \underbrace{\left[\sum_{k=1}^N l(\omega_k, g(\mathbf{x})) p(\omega_k | \mathbf{x}) \right]}_{\sum_{\omega_k \neq g(\mathbf{x})} p(\omega_k | \mathbf{x})} p(\mathbf{x}) d\mathbf{x} \end{aligned}$$

Optimality of the MAP rule (II)

- ▶ Due to $\sum_{k=1}^N p(\omega_k|\mathbf{x}) = 1$, we have

$$\sum_{\omega_k \neq g(\mathbf{x})} p(\omega_k|\mathbf{x}) = 1 - p(g(\mathbf{x})|\mathbf{x})$$

- ▶ We have

$$R(g) \downarrow \implies \forall \mathbf{x}, \left[1 - p(g(\mathbf{x})|\mathbf{x})\right] \downarrow \implies \forall \mathbf{x}, p(g(\mathbf{x})|\mathbf{x}) \uparrow$$

- ▶ Since $g(\mathbf{x}) \in \{\omega_1, \dots, \omega_K\}$, we choose:

$$g^*(\mathbf{x}) = \arg \max_k p(\omega_k|\mathbf{x})$$



The MAP decision rule example

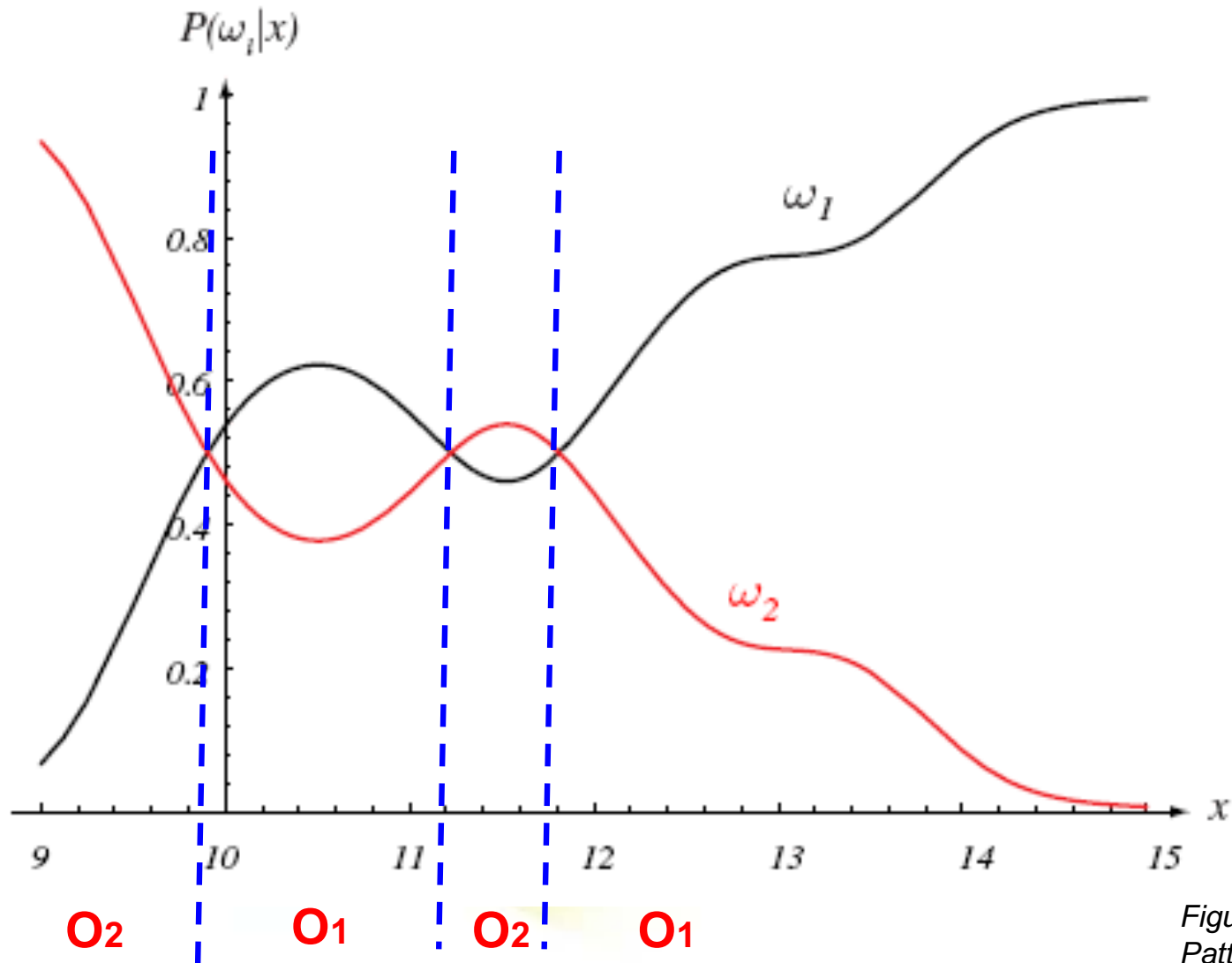
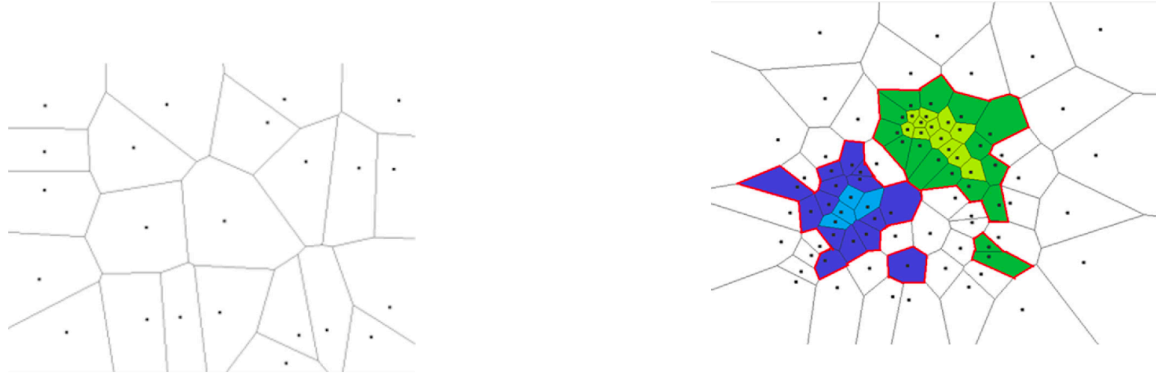


Figure from Duda et. al.,
Pattern classification
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Classification Error Probability

- ▶ Any rule $\mathbf{x} \rightarrow g(\mathbf{x}) \in \{\omega_1, \dots, \omega_K\}$ partitions input space into K regions: O_1, O_2, \dots, O_K : if $\mathbf{x} \in O_k$, implies $g(\mathbf{x}) = \omega_k$.



- ▶ The expected risk is the probability of classification error:

$$\begin{aligned} R(g) &= \Pr(\text{error}) = 1 - \Pr(\text{correct}) = 1 - \sum_{k=1}^K \Pr(\mathbf{x} \in O_k, \omega_k) \\ &= 1 - \sum_{k=1}^K \Pr(\omega_k) \int_{\mathbf{x} \in O_k} p(\mathbf{x}|\omega_k) d\mathbf{x} \end{aligned}$$

- ▶ Bayes error: $R(g^*)$ of the MAP rule (the lowest possible error)

Example of Error Probability in 2-class case

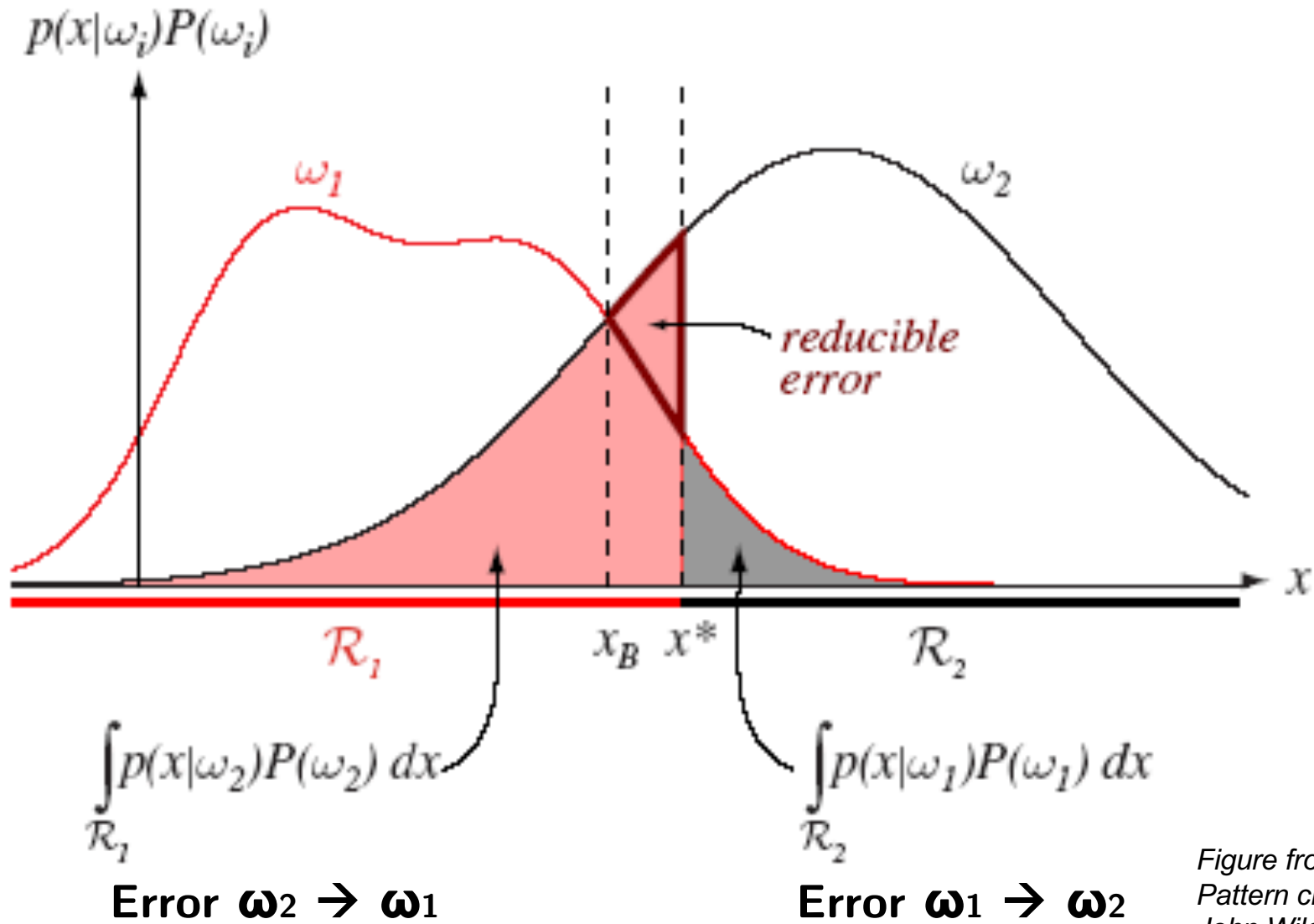


Figure from Duda et. al.,
Pattern classification
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Bayes Error

- Bayes error: error probability of the Bayes (MAP) decision rule.
- Since Bayes decision rule guarantees the minimum error, the Bayes error is the lower bound of all possible error probabilities.
- It is difficult to calculate the Bayes error, even for the very simple cases because of discontinuous nature of the decision regions in the integral, especially in high dimensions.
- Some approximation methods to estimate an upper bound.
 - Chernoff bound
 - Bhattacharyya bound
- Evaluate on an independent test set.



Example: the MAP rule for independent binary features

- ▶ 2-class (ω_1 and ω_2) classification: $\Pr(\omega_1)$ and $\Pr(\omega_2)$
- ▶ Using n independent binary features $\mathbf{x} = [x_1, x_2, \dots, x_n]^T$, $x_i \in \{0, 1\}$ $i = 1, 2, \dots, n$
- ▶ Denote $p_i \triangleq \Pr(x_i = 1|\omega_1)$ and $q_i \triangleq \Pr(x_i = 1|\omega_2)$, we have:

$$p(\mathbf{x}|\omega_1) = \prod_{i=1}^n p_i^{x_i} (1 - p_i)^{1-x_i} \quad p(\mathbf{x}|\omega_2) = \prod_{i=1}^n q_i^{x_i} (1 - q_i)^{1-x_i}$$

- ▶ The MAP rule: given \mathbf{x} , classify as ω_1 if $\Pr(\omega_1) \cdot p(\mathbf{x}|\omega_1) \geq \Pr(\omega_2) \cdot p(\mathbf{x}|\omega_2)$, otherwise ω_2 .
- ▶ Take logarithm to derive a **linear** decision boundary:

$$g(\mathbf{x}) = \sum_{i=1}^n \lambda_i x_i + \lambda_0 = \begin{cases} \geq 0 & \implies \omega_1 \\ < 0 & \implies \omega_2 \end{cases}$$

where $\lambda_i = \ln \frac{p_i(1-q_i)}{q_i(1-p_i)}$ and $\lambda_0 = \sum_{i=1}^n \ln \frac{1-p_i}{1-q_i} + \ln \frac{\Pr(\omega_1)}{\Pr(\omega_2)}$

Generative Models for Regression



- ▶ Input: n -dimensional vector \mathbf{x} , output: $y \in \mathbb{R}$
- ▶ The joint distribution $p(\mathbf{x}, y)$ is known, \mathbf{x} is used to predict y .
- ▶ What is the best decision rule for $\mathbf{x} \rightarrow y = g(\mathbf{x})$?

$$g^*(\mathbf{x}) = \mathbb{E}(y|\mathbf{x}) = \int_{\mathbf{x}} y \cdot p(y|\mathbf{x}) dy$$

Theorem 2

Assume $p(\mathbf{x}, y)$ is known, the conditional mean $\mathbb{E}(y|\mathbf{x})$ leads to the lowest expected risk (using mean square loss).

Optimality of Conditional Mean for Regression

Proof:

- ▶ The expected risk of any rule $\mathbf{x} \rightarrow g(\mathbf{x}) \in \mathbb{R}$:

$$\begin{aligned} R(g) &= \mathbb{E}_{p(\mathbf{x}, y)} [l(\omega, g(\mathbf{x}))] = \int_{\mathbf{x}} \int_y (y - g(\mathbf{x}))^2 p(\mathbf{x}, y) d\mathbf{x} dy \\ &= \int_{\mathbf{x}} \underbrace{\left[\int_y (y - g(\mathbf{x}))^2 p(y|\mathbf{x}) dy \right]}_{Q(g|\mathbf{x})} p(\mathbf{x}) d\mathbf{x} \end{aligned}$$

- ▶ Functional derivative:

$$\frac{\partial Q(g|\mathbf{x})}{\partial g(\cdot)} = 0 \implies \int_y (g(\mathbf{x}) - y) p(y|\mathbf{x}) dy = 0$$

$$\implies g^*(\mathbf{x}) = \int_y y \cdot p(y|\mathbf{x}) dy = \mathbb{E}(y|\mathbf{x}) \quad \blacksquare$$

Plug-in MAP Decision Rule for classification

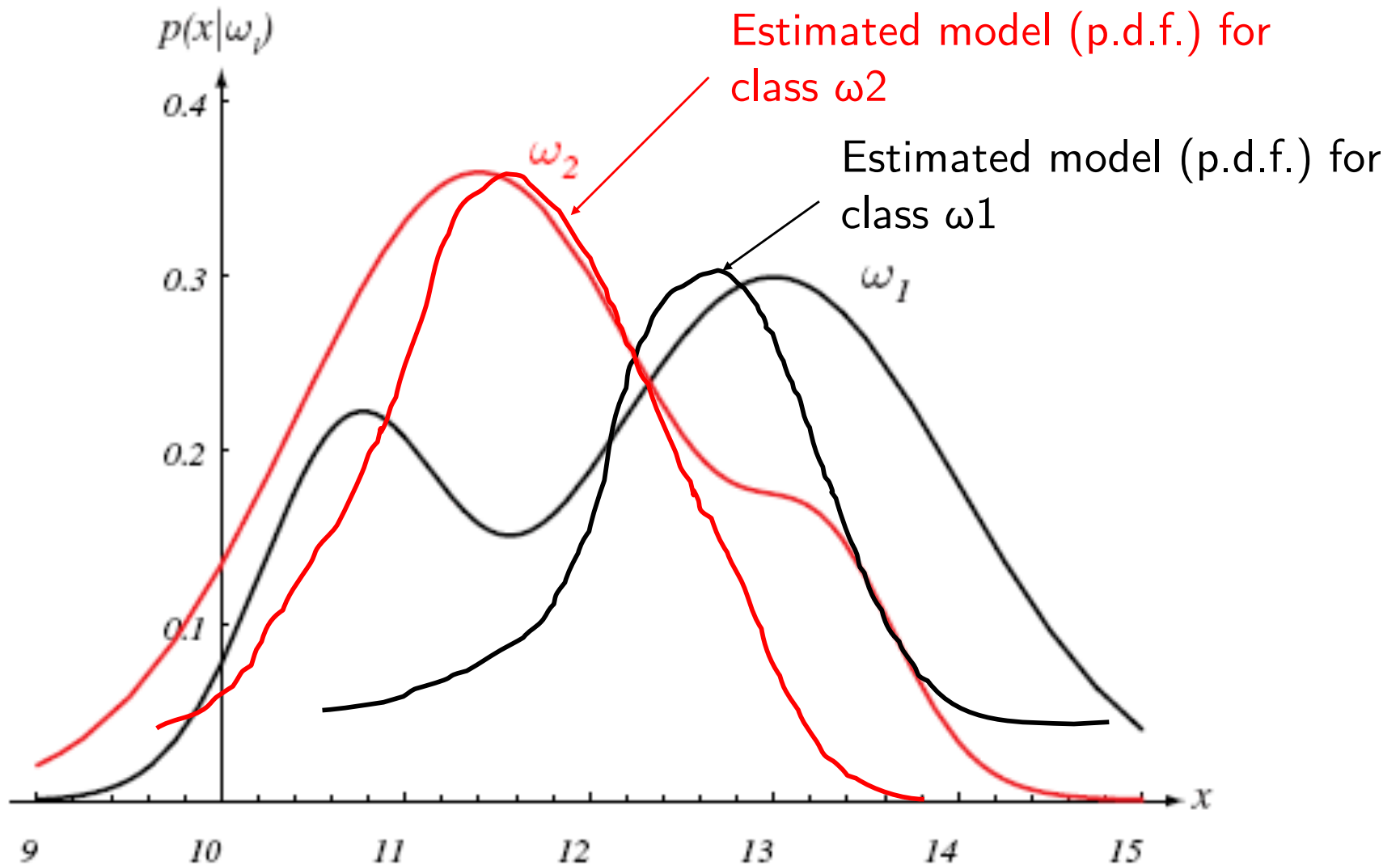
- ▶ The true distributions $\Pr(\omega_k)$ and $p(\mathbf{x}|\omega_k)$ are unknown.
- ▶ Training data: $\mathcal{D}_N = \{(\mathbf{x}_1, y_1), (\mathbf{x}_2, y_2), \dots, (\mathbf{x}_N, y_N)\}$
- ▶ Choose two probabilistic models:
 - ▶ $\hat{p}_\lambda(\omega_k)$ to approximate $\Pr(\omega_k)$
 - ▶ $\hat{p}_{\theta_k}(\mathbf{x})$ to approximate $p(\mathbf{x} | \omega_k)$ ($\forall k = 1, 2, \dots, K$)
- ▶ Parameter estimation: estimate $\{\lambda, \theta_1, \dots, \theta_K\}$ using \mathcal{D}_N
- ▶ The optimal MAP rule:

$$\omega^* = \arg \max_k \Pr(\omega_k) \cdot p(\mathbf{x}|\omega_k)$$

- ▶ The Plug-in MAP decision rule:

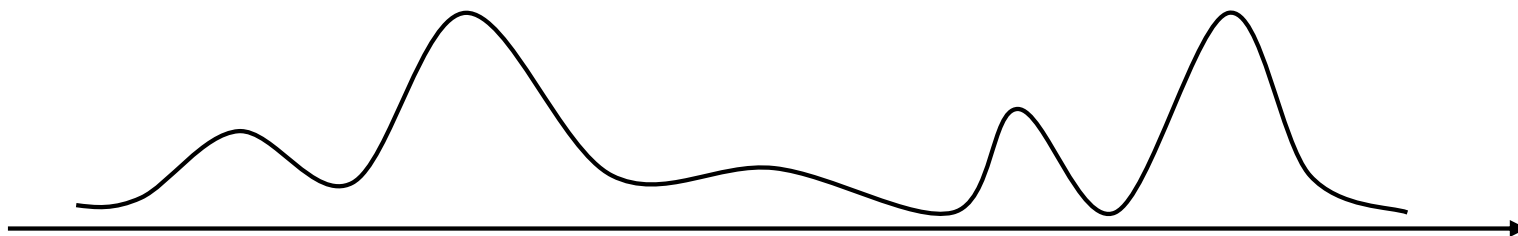
$$\omega^* = \arg \max_k \hat{p}_\lambda(\omega_k) \cdot \hat{p}_{\theta_k}(\mathbf{x})$$

Data modeling



Useful generative models (I)

- A proper generative model must be chosen based on the nature of observation data (the underlying structure of data).
- Some useful generative models for a variety of data types:
 - Normal (Gaussian) distribution
 - ➔ uni-modal continuous feature scalars
 - Multivariate normal (Gaussian) distribution
 - ➔ uni-modal continuous feature vectors
 - Gaussian Mixture models (GMM)
 - ➔ continuous feature scalars/vectors with multi-modal distribution nature
 - ➔ For speaker recognition/verification
distribution of speech features over a large population



Useful generative models (II)

- **Some useful generative models (cont' d)**
 - **Markov chain model: discrete sequential data**
 - **N-gram model in language modeling**
 - **Hidden Markov Models (HMM): ideal for various kinds of sequential observation data; provides better modeling capability than simple Markov chain model.**
 - **Model speech signals for recognition (one of the most successful story of data modeling)**
 - **Model language/text data for part-of-speech tagging, shallow language understanding, etc.**
 - **Model biological data (DNA & protein sequence): profile HMM.**
 - **Lots of other application domains.**



Useful generative models (III)

- **Some useful generative models (cont'd)**
 - **Markov Random Field (a.k.a. undirected graphical model):**
 - **multi-dimensional spatial data**
 - **Conditional random fields (CRF)**
 - **Bayesian networks (a.k.a. directed graphical model)**
 - **High-dimensional data (discrete or continuous)**
 - **Latent Dirichlet allocation (LDA)**
 - **Automatically learn dependency from data**

