Introduction to Machine Learning Foundations and Applications

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Hypothesis Class Complexity

Motivations

Hypothesis classes are typically infinite $|\mathcal{H}| = \infty$.

Can we still efficiently learn concepts c?

Yes, recall interval problem and axis-aligned rectangle problem was infinite but PAC-Learnable.

We need a notion of complexity for hypothesis class \mathcal{H} beyond cardinality $|\mathcal{H}|$.

Ultimately, we aim to obtain bounds on the generalization error in terms of the empirical risk.



Building Identification



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Notation and definitions:

 \mathfrak{X} input space, \mathfrak{Y} output space \mathfrak{C} concept class, concept c(x): $\mathfrak{X} \rightarrow \mathfrak{Y}$ \mathfrak{K} hypothesis class, hypothesis h(x): $\mathfrak{X} \rightarrow \mathfrak{Y}$.





Hans Rademacher 1892-1969

Issue: Hypothesis classes are typically infinite $|\mathcal{H}| = \infty$. Can we still efficiently learn concepts c?

Recall: Axis-aligned rectangle problem has infinite $|\mathcal{H}| = \infty$ but proved is PAC-Learnable.

Need a notion of complexity for hypothesis class \mathcal{H} beyond cardinality $|\mathcal{H}|$. $h \in \mathcal{H}$, g(x, y) = L(h(x), y)

Let loss function be denoted L: $\mathcal{Y} \times \mathcal{Y} \rightarrow \mathbb{R}$ and let G be family of loss functions associated with \mathcal{H} .

Definition: The **empirical Rademacher complexity** of a family of functions G with g(z): $Z \rightarrow [a,b] \subset \mathbb{R}$ and m fixed samples $S = (z_1, z_2, ..., z_m)$ is given by

$$\hat{\mathcal{R}}_{S}(G) = E_{\sigma} \left[\sup_{g \in G} \frac{1}{m} \sum_{i=1}^{m} \sigma_{i} g(z_{i}) \right] \text{, where } \boldsymbol{\sigma} = (\sigma_{1}, \sigma_{2}, \dots, \sigma_{m}) \text{ are uniform random variables in } \{-1, +1\}.$$

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Definition: The **Rademacher complexity** of a family of functions G on m samples is $\mathcal{R}_m(G) = E_{S \sim D^m} \left[\hat{\mathcal{R}}_S(G) \right]$

- Averaged sum term can be viewed as an inner-product: $\sum \sigma_i \cdot g(z_i) = \boldsymbol{\sigma} \cdot \boldsymbol{g}_S$.
- Rademacher complexity gives a measure of the "richness" of family G in approximating random functions. $\mathcal{R}_m(G) = E_{S \sim D^m, \sigma} \left[\sup_{g \in G} \frac{1}{m} \sigma \cdot \mathbf{g}_S \right].$ Gives a measure of the "correlation" between \mathbf{g}_S and $\boldsymbol{\sigma}$.

Definition: The **empirical Rademacher complexity** of a family of functions G with $g(z): Z \rightarrow [a,b] \subset \mathbb{R}$ and m fixed samples $S = (z_1, z_2, ..., z_m)$ is given by

$$\hat{\mathcal{R}}_{S}(G) = E_{\sigma} \left[\sup_{g \in G} \frac{1}{m} \sum_{i=1}^{m} \sigma_{i} g(z_{i}) \right] \text{, where } \boldsymbol{\sigma} = (\sigma_{1}, \sigma_{2}, \dots, \sigma_{m}) \text{ are random in set } \{-1, +1\}.$$

Example: Rademacher Complexity for family of functions $G = \{g(z) = g_0 \in [-c, c]\}$ (constants).

$$\begin{split} \tilde{\mathcal{R}}_{\mathcal{S}}(\mathcal{G}) &= E_{\sigma} \left[\sup_{g \in \mathcal{G}} \frac{1}{m} \sum_{i=1}^{m} \sigma_{i} g(z_{i}) \right] = E_{\sigma} \left[\max \left\{ \frac{1}{m} \sum_{i=1}^{m} c \sigma_{i}, -\frac{1}{m} \sum_{i=1}^{m} c \sigma_{i} \right\} \right] \\ &= E_{\sigma} \left[\frac{1}{m} c \left| \# \{ \sigma_{i} = +1 \} - \# \{ \sigma_{i} = -1 \} \right| \right] = \frac{c}{m} E_{\sigma} \left[\left| \sum_{i=1}^{m} \sigma_{i} \right| \right] \leq \frac{c \sqrt{m}}{m} = \frac{c}{\sqrt{m}} \end{split}$$

Jensen Inequality (ϕ convex): $\phi(E[X]) \leq E[\phi(X)]$ $(E[|X|])^2 \leq E[|X|^2]$ $E\left[\left|\sum_{i=1}^{m} \sigma_{i}\right|\right] \leq E\left[\left|\sum_{i=1}^{m} \sigma_{i}\right|^{2}\right]^{1/2}$ $= E\left[\sum_{i,j=1}^{m} \sigma_{i}\sigma_{j}\right]^{1/2} = \sqrt{m}$



Notation and definitions:

 \mathfrak{X} input space, \mathfrak{Y} output space \mathfrak{C} concept class, concept c(x): $\mathfrak{X} \rightarrow \mathfrak{Y}$ \mathfrak{H} hypothesis class, hypothesis h(x): $\mathfrak{X} \rightarrow \mathfrak{Y}$.





Hans Rademacher 1892-1969

Theorem: (expectation bounds g: Z \rightarrow [0,1]) For family of loss functions G into [0,1] and any $\delta > 0$ we have with probability $1 - \delta$ that the following bounds hold uniformly for any $g \in G$,



Significance: The expected value E[g] can be bounded by the observed empirical average. This differs at most by the Rademacher Complexity plus a term vanishing as $m \rightarrow \infty$.

We shall use for bound on the generalization error by the empirical risk.

Notation and definitions:

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Hans Rademacher 1892-1969

Definition: The Empirical Rademacher Complexity of a hypothesis class \mathcal{H} is

$$\hat{\mathcal{R}}_{S^{\mathcal{X}}}(H) = E_{\sigma} \left[\sup_{h \in H} \frac{1}{m} \sum_{i=1}^{m} \sigma_i h(x_i) \right] , \text{ (note: we take } h \in \{-1, 1\})$$

Definition: The **Rademacher Complexity** of *a hypothesis class* \mathcal{H} is $\mathcal{R}_m(H) = E_{S \sim D^m} \left[\hat{\mathcal{R}}_{S^{\mathcal{X}}}(H) \right]$, (note: we take $h \in \{-1, 1\}$)

Lemma: For the family of 0-1 loss functions $G = \{(x, y) \to 1_{h(x) \neq y} | h \in H\}$ we have $\hat{\mathcal{R}}_S(G) = \frac{1}{2} \hat{\mathcal{R}}_{S^{\mathcal{X}}}(H)$

• Allows for working more directly with the hypothesis space in constructing bounds.

Notation and definitions:

 \mathfrak{X} input space, \mathfrak{Y} output space \mathfrak{C} concept class, concept c(x): $\mathfrak{X} \rightarrow \mathfrak{Y}$ \mathfrak{H} hypothesis class, hypothesis h(x): $\mathfrak{X} \rightarrow \mathfrak{Y}$.





Hans Rademacher 1892-1969

Theorem: (bound on generalization error for 0-1 loss) For 0 - 1 loss $G = \{(x, y) \rightarrow 1_{h(x) \neq y} | h \in H\}$ and any $\delta > 0$ we have with probability $1 - \delta$ that the following bounds hold uniformly for any $g \in G$,

$$\begin{split} R(h) &\leq \hat{R}_{S}(h) + \mathcal{R}_{m}(H) + \sqrt{\frac{\log(\frac{1}{\delta})}{2m}} \quad \text{, (Rademacher bound)} \\ R(h) &\leq \hat{R}_{S}(h) + \hat{\mathcal{R}}_{S^{\mathcal{X}}}(H) + \underbrace{3\sqrt{\frac{\log(\frac{2}{\delta})}{2m}}}_{\text{empirical estimate complexity}}, \quad \text{(Empirical Rademacher bound)} \end{split}$$

Significance: The generalization error can be bounded by the observed empirical risk. This differs most by the Rademacher Complexity plus a term vanishing as $m \rightarrow \infty$.

• This shows we can use Rademacher complexity in place of $|\mathcal{H}|$ to obtain bounds on **generalization error** to obtain **scaling in m**.

Theorem: (bound on generalization error for 0-1 loss) For 0 - 1 loss $G = \{(x, y) \rightarrow 1_{h(x) \neq y} | h \in H\}$ and any $\delta > 0$ we have with probability $1 - \delta$ that the following bounds hold uniformly for any $g \in G$,

 $R(h) \leq \hat{R}_S(h) + \hat{\mathcal{R}}_{S^{\mathcal{X}}}(H) + 3\sqrt{\frac{\log(\frac{2}{\delta})}{2m}} \text{, (Empirical Rademacher bound)}$

Example: Rademacher Complexity for family of functions $H = {h(x) = h_0 \in [-c, c], c = 1}$ (constants).

$$\tilde{\mathcal{R}}_{\mathcal{S}}(\mathcal{G}) = E_{\sigma} \left[\sup_{g \in \mathcal{G}} \frac{1}{m} \sum_{i=1}^{m} \sigma_i g(z_i) \right] \leq \frac{c\sqrt{m}}{m} = \frac{c}{\sqrt{m}}$$
 (from previous derivation)



Theorem: (bound on generalization error for 0-1 loss) For 0 - 1 loss $G = \{(x, y) \rightarrow 1_{h(x) \neq y} | h \in H\}$ and any $\delta > 0$ we have with probability $1 - \delta$ that the following bounds hold uniformly for any $g \in G$,

$$R(h) \leq \hat{R}_S(h) + \hat{\mathcal{R}}_{S^{\mathcal{X}}}(H) + 3\sqrt{\frac{\log(\frac{2}{\delta})}{2m}},$$
 (Empirical Rademacher bound)

Theorem (Massert's lemma): Let $A \subseteq \mathbb{R}^n$ be a finite set of vectors with $r = \max_{a \in A} ||a||_2$ then $\hat{\mathcal{R}}_S(A) = E_\sigma \left[\sup_{\mathbf{a} \in A} \frac{1}{m} \sum_{i=1}^m \sigma_i a_i \right] \le \frac{r\sqrt{2\log|A|}}{m}$

Hypothesis class \mathcal{H} and m samples consider the set $A = \{(h(x_1), h(x_2), ..., h(x_m)) : h \in \mathcal{H}\}$.

Finite hypothesis class we have $|A| \leq |\mathcal{H}|$.

Note: Result similar to prior complexity bound for finite consistent case

$$R(h_S) \le \frac{1}{m} \Big(\log |H| + \log \frac{1}{\delta} \Big)$$

Massert's Lemma significantly generalizes this result since |A| is now allowed to grow with m for $|\mathcal{H}| = \infty$.

Alternatively, combinatorial measures like complexity |A| may be easier to estimate than Rademacher complexity.

Growth Function

Definition: The growth function $\Pi_H: \mathbb{N} \to \mathbb{N}$ for a hypothesis class \mathcal{H} is defined as $\Pi_H(m) = \max_{\{x_1, x_2, \dots, xm\} \subseteq X} |\{(h(x_1), h(x_2), \dots, h(x_m)): h \in \mathcal{H}\}|$

Counts maximum number of distinct m-vectors $(h(x_1), h(x_2), ..., h(x_m))$ that can be generated by the hypothesis class \mathcal{H} .

Upper bound on the number of distinct ways m points can be classified by \mathcal{H} .

Example: $\mathcal{X} = \{-2, -1, 1\}, \ \mathcal{Y} = \{-1, 1\}, \ \mathcal{H} = \{h_1(x) = \text{sign}(x), \ h_2(x) = \text{sign}(x - 1.5)\}, \ h_1: -1, -1, 1; \ h_2: -1, -1, -1.$ For m=2, most variation for x₁=-1, x₂=1, with $\Pi_H(2) = |\{(-1, +1), (-1, -1)\}| = 2$. In general, we have $\Pi_H(m) = 2$.

Remark: For finite hypothesis class always have $\Pi_H(m) \leq |\mathcal{H}|$.

Example: $\mathcal{X} = \mathbb{R}$, $\mathcal{Y} = \{-1,1\}$, $\mathcal{H} = \{h(x) = \text{sign}(p(x)) \text{ with } p(x) \text{ polynomial degree } n\}$. Now $|\mathcal{H}| = \infty$ and we have $\Pi_H(m) \leq r(m)2^{n+1}$, r = poly. Follows from Lagrange interpolation.⁻¹

Example: $\mathcal{X} = \mathbb{R}$, $\mathcal{Y} = \{-1,1\}$, $\mathcal{H} = \{h(x) = \text{sign}(x - a) \text{ with } a \in \mathbb{R}\}$ half-space classifiers. Now $|\mathcal{H}| = \infty$ and we have $\Pi_H(m) = m + 1$. $\#\{h(x_i) = -1\}, i \in \{1, \dots, m\}$ linear classifier

polynomial classifier

h(x)

-1





Growth Function

Definition: The growth function $\Pi_H: \mathbb{N} \to \mathbb{N}$ for a hypothesis class \mathcal{H} is defined as $\Pi_H(m) = \max_{\{x_1, x_2, \dots, xm\} \subseteq X} |\{(h(x_1), h(x_2), \dots, h(x_m)): h \in \mathcal{H}\}|$

- **Counts maximum number** of distinct m-vectors $(h(x_1), h(x_2), ..., h(x_m))$ that can be generated by the hypothesis class \mathcal{H} .
- Upper bound on the number of distinct ways m points can be classified by *H*.

Theorem (Massert's Lemma): The Rademacher complexity is bounded by the growth function as

$$\mathcal{R}_m(\mathcal{H}) \le \sqrt{\frac{2\log\left(\Pi_{\mathcal{H}}(m)\right)}{m}}$$

Theorem (bound on generalization error for 0-1 loss): For any $\delta > 0$ we have with probability $1 - \delta$ that the following bounds hold uniformly for any $h \in \mathcal{H}$,

$$R(h) \le \hat{R}_S(h) + \sqrt{\frac{2\log\left(\Pi_{\mathcal{H}}(m)\right)}{m}} + \sqrt{\frac{\log(\frac{1}{\delta})}{2m}}$$

Note: Bound is now distribution D independent depending only on combinatorial features of \mathcal{H} .



VC-Dimension

Definition: For a sample set $S = (x_1, x_2, ..., x_m)$ of size m, a **dichotomy** is one of the possible ways to label the set (y_1, y_2, \dots, y_m) .



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Alexev Chervonenkis

Definition: A set S of size m is said to be **shattered** by the hypothesis class \mathcal{H} if for each dichotomy **y** there is an $h \in \mathcal{H}$ so that $(h(x_1) = y_1, h(x_2) = y_2, ..., h(x_m) = y_m)$.

Example: $\mathcal{X} = \{-2, -1, 1\}, \ \mathcal{U} = \{-1, 1\}, \ H = \{h_1(x) = \text{sign}(x), \ h_2(x) = \text{sign}(x - 1.5)\}.$ h_1 : -1,-1,1; h_2 : -1,-1,-1. Now for $x_1 = -2$ with dichotomy $y_1 = 1$ can not be obtained from either h_1 or h_2 this hypothesis class fails to shatter even \mathcal{X}^{m} for m = 1.



Example: $\mathcal{X} = \mathbb{R}$, $\mathcal{Y} = \{-1,1\}$, $\mathcal{H} = \{h : h(x) = sign(x-a) \cdot sign(b-x) \text{ for some } a, b \in \mathbb{R}\}$ the set of intervals [a,b]. Now for m = 2 for any two points $x_1, x_2 \in \mathbb{R}$ we have \mathcal{H} shatters \mathcal{X}^2 by taking [a,b] to contain points with $y_i = 1$ and exclude any point with $y_i = -1$.

However, for m \ge 3 we *can not match* all **dichotomies**. Take for example $x_1 < x_2 < x_3$ with the labels $y_1 = +1$, $y_2 = -1$, $y_3 = +1$ then there is no interval containing both x_1 and x_3 but excluding x_2 . Therefore, there exists dichotomies when m = 3 that no $h \in \mathcal{H}$ can classify correctly.

VC-Dimension

Definition: The Vapnik-Chervonenkis dimension is defined as $VCdim(\mathcal{H}) = \max\{m : \Pi_H(m) = 2^m\}$

 The VC-dimension measures the size of the largest set that can be shattered by the hypothesis class \mathcal{H} .



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- When $VCdim(\mathcal{H}) = d$ this means there exists a set of size d that can be fully **shattered** by \mathcal{H} .
- For finite $|\mathcal{H}| < \infty$ hypothesis space we have $VCdim(\mathcal{H}) \leq \log(|\mathcal{H}|)$.

Example: $\mathcal{X} = \mathbb{R}$, $\mathcal{Y} = \{-1,1\}$, $\mathcal{H} = \{h : h(x) = sign(x - a) \cdot sign(b - x) \text{ for some } a, b \in \mathbb{R}\}$ the set of intervals [a,b]. For m = 2 for any two points $x_1, x_2 \in \mathbb{R}$ we have \mathcal{H} shatters \mathcal{X}^2 by taking [a,b] to contain points with v_i = 1 and exclude any point with $y_i = -1$.

However, for m \ge 3 we can not match all dichotomies. Take for example $x_1 < x_2 < x_3$ with the labels $y_1 = +1$, $y_2 = -1$, $y_3 = +1$ then there is no interval containing both x_1 and x_3 but excluding x_2 . **Therefore**, $VCdim(\mathcal{H}) = 2$.

Example: $\mathcal{X} = \mathbb{R}$, $\mathcal{Y} = \{-1, 1\}$, $\mathcal{H} = \{h : h(x) = sign(p(x)) \text{ polynomial } p(x) \text{ of degree } n\}$. We have \mathcal{H} shatters \mathcal{X}^m for m = n + 1. This follows from Lagrange interpolation. However, can not shatter for m > n + 1, so d = $VCdim(\mathcal{H}) = n + 1$.

VC-Dimension

Definition: The **Vapnik-Chervonenkis dimension** is defined as $VCdim(\mathcal{H}) = \max\{m : \Pi_H(m) = 2^m\}$

- The VC-dimension measures the size of the largest set that can be **shattered** by the hypothesis class \mathcal{H} .
- When $VCdim(\mathcal{H}) = d$ this means there exists a set of size d that can be fully **shattered** by \mathcal{H} .

Theorem (bound on generalization error for 0-1 loss): When $VCdim(\mathcal{H}) = d$, for any $\delta > 0$ we have with probability $1 - \delta$ that the following bounds hold uniformly for any $h \in \mathcal{H}$,

$$R(h) \le \hat{R}_S(h) + \sqrt{\frac{2d\log\left(\frac{em}{d}\right)}{m}} + \sqrt{\frac{\log(\frac{1}{\delta})}{2m}}$$

• Note the ratio of m/d governs the bound. This corresponds to the overall form

$$R(h) \le \hat{R}_S(h) + O\left(\sqrt{\frac{\log(m/d)}{(m/d)}}\right)$$

• This shows we need sample size m >> d to obtain small bound. Provides useful complexity when $|\mathcal{H}| = \infty$.



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VC Dimension

Example: $VCdim(\mathcal{H})$ axis-aligned rectangles.

Claim: $VCdim(\mathcal{H}) = 4$.

Two steps:

(i) lower bound $VCdim(\mathcal{H}) \ge 4$ (ii) upper bound $VCdim(\mathcal{H}) < 5$

Lower bound: Place 4 points into a diamond configuration. All cases can clearly be handled.

Upper bound: Place 5 points with 4 on rectangle and the 5th point in the interior. No axis-aligned rectangle that can correctly classify these points for all labels. Hence, $VCdim(\mathcal{H}) < 5$.

Characterizes the complexity of the infinite dimensional hypothesis space \mathcal{H} .

VC-dimension bounds provide a sampling complexity for learning the axis-aligned rectangle.

Cases for 4 points





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VC-Dimension: Hyperplanes

Example: Learning separating hyperplane in \mathbb{R}^N (related to SVM). For data $\{(x_i, y_i)\}$ with $x_i \in \mathbb{R}^N$ and $y_i \in \{-1, 1\}$. Ideally, find **w**, b so that sign $(w^T x_i + b) = y_i$.

Hypothesis class:

 $\mathcal{H} = \{h: h(\mathbf{x}) = \operatorname{sign}(\mathbf{w}^{\mathrm{T}}\mathbf{x} + b) \text{ with } \mathbf{w} \in \mathbb{R}^{\mathsf{N}}, b \in \mathbb{R}\}.$

What is the $VCdim(\mathcal{H})$?

Claim: $VCdim(\mathcal{H}) = N + 1$

Two steps:

(i) lower bound $VCdim(\mathcal{H}) \ge N + 1$.

(ii) upper bound $VCdim(\mathcal{H}) < N + 2$.



Lower bound: For N + 1 points, let $\mathbf{x}_0 = (0, 0, ..., 0)$ origin, $\mathbf{x}_i = (0, ..., 1, ..., 0, 0) = \mathbf{e}_i$, with ith component one. For any labels $y_i \in \{-1, 1\}$, let $\mathbf{w} = (y_1, y_2, ..., y_N)$ and $b = \frac{y_0}{2}$ which defines the classifier $h(\mathbf{x}_i) = \operatorname{sign}(\mathbf{w}^T \mathbf{x}_i + \mathbf{b}) = \operatorname{sign}(y_i + \frac{y_0}{2}) = y_i$. This verifies any N + 1 labels can be classified correctly.

Hyperplanes \mathcal{H} shatters this N + 1 point-set so $VCdim(H) \ge N+1$.

VC-Dimension: Hyperplanes

Example: Learning separating hyperplane in \mathbb{R}^N (related to SVM). For data $\{(x_i, y_i)\}$ with $x_i \in \mathbb{R}^N$ and $y_i \in \{-1, 1\}$. Ideally, find **w**, b so that sign $(\mathbf{w}^T \mathbf{x}_i + b) = y_i$.

Upper bound: $VCdim(\mathcal{H}) < N+2$. Must show for any N + 2 points for some labels there is no hyperplane classifier.

Theorem (Radon): In \mathbb{R}^N a set of N + 2 points always can be partitioned into two disjoint subsets \mathcal{X}_1 and \mathcal{X}_2 that have intersecting convex hulls $C(\mathcal{X}_1) \bigcap C(\mathcal{X}_2) \neq \emptyset$.

Implication: Let labels of \mathcal{X}_1 be say +1 and \mathcal{X}_2 be -1 then there is no separating hyperplane (it would separate the convex hulls).





Proof (Radon): Consider the set of N + 1 linear equations in N + 2 unknowns: $\sum_{i=1}^{N+2} \alpha_i \mathbf{x}_i = \mathbf{0}$ and $\sum_{i=1}^{N+2} \alpha_i = \mathbf{0}$ Non-trivial null-space so equations have non-zero solution $\beta_1, \dots, \beta_{d+2}$ with $\sum_{i=1}^{N+2} \beta_i = \mathbf{0}$. Let $I_1 = \{i : \beta_i > 0\}$ and $I_2 = \{i : \beta_i \le 0\}$, then both non-empty. Let $\mathbf{x}^* = \sum_{i_1 \in I_1} \frac{\beta_{i_1}}{\beta} \mathbf{x}_{i_1} = \sum_{i_2 \in I_2} \frac{-\beta_{i_2}}{\beta} \mathbf{x}_{i_2}$ with $\beta = \sum_{i_1 \in I_1} \beta_{i_1}$. We have $\sum_{i_1 \in I_1} \frac{\beta_{i_1}}{\beta} = \sum_{i_2 \in I_2} \frac{-\beta_{i_2}}{\beta} = 1$ and $\frac{\beta_{i_1}}{\beta} \ge 0$, $\frac{-\beta_{i_2}}{\beta} \ge 0$, so $x^* \in C(\mathcal{X}_1) \bigcap C(\mathcal{X}_2) \neq \emptyset$ so convex hulls intersect.

VC-Dimension: Hyperplanes

Example: Learning separating hyperplane in \mathbb{R}^N (related to SVM). For data $\{(x_i, y_i)\}$ with $x_i \in \mathbb{R}^N$ and $y_i \in \{-1, 1\}$. Ideally, find **w**, b so that sign $(w^T x_i + b) = y_i$.

Hypothesis class:

 $\mathcal{H} = \{h: h(\mathbf{x}) = sign(\mathbf{w}^{T}\mathbf{x} + b) \text{ with } \mathbf{w} \in \mathbb{R}^{N}, b \in \mathbb{R}\}.$

What is the $VCdim(\mathcal{H})$?

Claim: $VCdim(\mathcal{H}) = N + 1$

Shows in separable case that we have bound on generalization error

$$R(h) \le \widehat{R}(h) + \sqrt{\frac{2(N+1)\log\frac{em}{N+1}}{m}} + \sqrt{\frac{\log\frac{1}{\delta}}{2m}}$$

Turns out we can do even better in bounding sampling complexity for SVM. Want independent of feature dimension N, for bounded features (future lectures).

Will discuss further these results later when we cover Support Vector Machines.





Building Identification

VC Dimension: Lower Bounds

Lower Bounds: Given assumptions of PAC-Learning and $VCdim(\mathcal{H})$. What is lower bound on generalization error given m samples?

Theorem: Under assumptions of PAC for $d = VCdim(\mathcal{H}) > 1$ given any learning algorithm \mathcal{A} there always exists a distribution D and concept $f \in \mathcal{C}$ so that for m samples

$$\Pr_{S \sim D^m} \left[R_D(h_S, f) > \frac{d-1}{32m} \right] \ge 1/100$$

Shows that at least 1% of the time you will always have generalization error bigger than $\frac{d-1}{32m}$.

Characterizes the worse-case generalization errors given complexity of \mathcal{H} .

Consequence: If $VCdim(\mathcal{H}) = \infty$ then task is **not PAC-Learnable**.



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VC Dimension: Lower Bounds

Example: Consider hypothesis class of all polynomials $\mathcal{H} = \{h: h(x) = sign(p(x)) \text{ any polynomial of finite degree} \}.$

Complexity: $VCdim(\mathcal{H}) = \infty$ (recall for n degree polynomial VCdim = n+1).

Consequence: Concepts from \mathcal{H} are **not** PAC-Learnable.

Why? At least 1% of the time you will always have generalization error bigger than $\frac{d-1}{32m}$ so make d = [31.7m + 1] (since $VCdim(\mathcal{H}) = \infty$ can take any d > 1) then we have $\Pr_{S \sim D^m} \left[R_D(h_S, f) > \frac{d-1}{32m} \right] \ge 1/100 \longrightarrow \Pr_{S \sim D^m} \left\{ R_D(h_S, f) > 0.99 \right\} \ge 1/100$

Shows no matter how many samples m used, 1% of the time the generalization error is greater than 99%.

Not enough information from finite data alone to distinguish unknown function in \mathcal{H} without further assumptions (i.e. could miss local variations). Need other approaches (i.e. regularization, level of smoothness).

Consequence, if $VCdim(\mathcal{H}) = \infty$ then task is **not PAC-Learnable.**

Complexity: Rademacher, Growth Functions, VC-Dimension Complexity Bounds Theory and Practice

Significance: Complexity measures give some guarantees to assess generalization errors based on observed empirical risk.

Neural Networks and Deep Learning

In practice, often challenging since models have large complexity and we want to avoid^{ship} overfitting data by only minimizing empirical risk. Training methods often also have further regularizations or stochasticity (Kernel-SVM, N-Nets, SGD, Dropout).

Many extensions to the introduced ideas here but PAC + complexity bounds provide a good starting point for theory and intuition.

How can we use complexity measures in practice to inform our design of learning algorithms, training methods, and assess expected performance?



Support Vector Machines





Generative Methods





Manifold Learning





Image Classification





MIT and Boston Dynamics



washingtonpost.com

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Machine Learning: Foundations and Applications

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