F14-10601, Machine Learning Midterm Review Sheet

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1 introduction

- supervised, unsupervised, active learning, reasoning under uncertainty.
 - supervised: given a set of features and labels, learn a model, predict;
 - unsupervised: discover patterns in data without labels;
 - reasoning: determine a model either from samples or as going along;
 - active: select not only model but also which examples to use.
- probability: random variable, domain, kolmogorov's axioms (foundation):
 - $-0 \le P(A) \le 1;$
 - P(true) = 1, P(false) = 0;
 - $-P(A \vee B) = P(A) + P(B) P(A \wedge B).$

priors, conditional probability, joint distributions, chain rule. bayes rule:

$$P(A|B) = \frac{P(B|A)P(A)}{P(B)} = \frac{P(B|A)P(A)}{\sum_{A} P(B|A)P(A)}$$
(1)

density estimator, and hence the maximum likelihood principle:

$$\hat{P}(\text{Dataset}|M) = \hat{P}(x_1 \wedge x_2 \wedge \dots \wedge x_n|M) = \prod_{k=1}^n \hat{P}(x_k|M)$$
 (2)

the logarithmic maximum likelihood principle:

$$\log \hat{P}(\text{Dataset}|M) = \log \prod_{k=1}^{n} \hat{P}(x_k|M) = \sum_{k=1}^{n} \log \hat{P}(x_k|M)$$
(3)

take ∂ to find the best estimate for binary and guassian distros.

$$\arg\max_{q} q^{n_1} (1 - q)^{n_2} \to q = \frac{n_1}{n_1 + n_2} \tag{4}$$

for gaussian, that's the conventional deviation and variance.

2 classification

• bayes decision rule: determine class label using the bayes rule:

$$q_i(x) \equiv P(y=i|x) = \frac{P(x|y=i)P(y=i)}{P(x)}$$
 (5)

bayes error (risk) with two classes,

$$R(x) = \min\{P_1(x)P(y=1), P_0(x)P(y=0)\}\tag{6}$$

- classifiers: instance based (k nearest neighbors), generative (bayesian networks), discriminative (decision tree), latter two are major types.
- k nearest neighbors (knn): choice of k influences the *smoothness*, but this is determined by the actual distribution, not a pre-defined parameter. :(knn as approximated bayes decision rule.
- a probabilistic interpretation: let V to be volume of the m dimensional ball around z containing the k nearest neighbors for z, then we have

$$p(x)V = P = \frac{K}{N}, \quad p(x) = \frac{K}{NV}, \quad p(x|y=1) = \frac{K_1}{N_1V}, \quad p(y=1) = \frac{N_1}{N_1V},$$
 (7)

and using the bayes rule we get $p(y=1|x) = \frac{K_1}{K}$.

3 decision trees

- discriminative! nodes: attributes, leaves: labels, edges: assignments. split/create subtrees by testing on some attributes.
- determine the best attribute: *entropy*, the definition:

$$H(X) = \sum_{c} -p(X=c)\log_2 p(X=c)$$
 (8)

interpretation: with distribution known, minimum bits required to transmit a value. e.g. $P(X=1)=1 \rightarrow H=0, \ P(X=1)=0.5 \rightarrow H=1.$ expected bits per symbol. – its motivation and origin...

• conditional entropy: definition:

$$H(X|Y) = \sum_{i} P(Y=i)H(X|Y=i)$$
(9)

choose the attribute that produces the most entropy reduction, or the most *information gain*, as defined by

$$IG(X|Y)^* = H(X) - H(X|Y)$$
 (10)

which is always greater than 0, by Jensen's inequality.

- to avoid overfitting: tree pruning:
 - split data into train and test set, then build a tree using training set;
 - for all nodes, remove subtree, assign outcome to be the most common, check testing error and decide whether to keep change or restore.
- for continuous values: threshold, recursively.

4 naive-bayes classifier

• generative! naive-bayes assumes that the attributes are conditionally independent given the class label, or mathematically,

$$P(X|y) = \prod_{j} p_j(x^j|y) \tag{11}$$

and the full classification rule becomes.

$$\hat{y} = \arg\max_{v} p(y = v|X) = \arg\max_{v} \frac{p(X|y = v)p(y = v)}{p(X)}$$

$$= \arg\max_{v} \prod_{j} p_{j}(x^{j}|y = v)p(y = v)$$
(12)

and finally the conditional likelihood,

$$L(X_i|y_i = v, \Theta_v) = \prod_j p(x_i^j|y_i = v, \theta_v^j)$$
(13)

• for continuous values: covariance matrix \rightarrow diagonal matrix:

$$P(X|y=v) = \prod_{j} \frac{1}{\sqrt{2\pi}\sigma_v^j} \exp\left[-\frac{(x^j - \mu_v^j)^2}{2(\sigma_v^j)^2}\right]$$
(14)

separate means and variances for each class, the estimate would be:

$$\mu_v^j = \frac{1}{k_v} \sum_{X_i: y_i = v} x_i^j, \quad (\sigma_v^j)^2 = \sum_{X_i: y_i = v} \left(x_i^j - \mu_v^j \right)^2$$
 (15)

where $(\cdot)_i$ is the instance index, $(\cdot)^j$ feature index. $(\cdot)_v$ class label.

• problems: naive! assumption often violated. also: go through the article classification example in the homework or face annihilation.

5 regression

• linear regression with least squares:

$$w = \arg\min_{w} \sum_{i} (y_i - wx_i)^2$$
 where $y = wx + \epsilon$ (16)

 ϵ as noise, y as label, x_i as features, w as weights. taking the derivative

$$\frac{\partial}{\partial w} \sum_{i} (y_i - wx_i)^2 = 0 \quad \to \quad w = \frac{\sum_{i} x_i y_i}{\sum_{i} x_i^2}$$
 (17)

gives the best estimate for w.

• introducing a bias term: $y = w_0 + w_1 x + \epsilon$, then

$$w_0 = \frac{\sum_i (y_i - w_1 x_i)}{n}, \quad w_1 = \frac{\sum_i x_i (y_i - w_0)}{\sum_i x_i^2}$$
 (18)

• multivariate regression + non-linear basis function, with w, ϕ vectors,

$$y = w_0 \phi_0(x) + w_1 \phi_1(x) + \dots + w_k \phi_k(x) = \sum_{j=0}^n w_j \phi_j(x) = w^T \phi(x) \quad (19)$$

and let's minimize the loss-function, with w, ϕ vectors,

$$J(w) = \sum_{i} (y^i - w^T \phi(x^i))^2$$
(20)

$$\frac{\partial}{\partial w}J(w) = 2\sum_{i}(y^{i} - w^{T}\phi(x^{i}))\phi(x^{i})^{T} \Rightarrow 0$$
 (21)

we get, by pesudo-inverse, with j as feature index, and i instance index:

$$w = (\Phi^T \Phi)^{-1} \Phi^T y \quad \text{where} \quad \Phi = \phi_i(x^i) \tag{22}$$

- probabilistic interpretation: mle instead of least squares.
- locally weighted models, use weight function Ω_x and solve

$$\min_{w} \sum_{i} \Omega_x(x^i) (y^i - w^T \phi(x^i))^2 \tag{23}$$

where the weights can be determined, e.g., by gaussian function.

6 logistic-regression

- discriminative vs. generative.
- regression for classification: sigmoid function as activation function.
- determine parameters using maximum likelihood estimation.

$$L(y|X;w) = \prod_{i} \left(\frac{1}{1 + \exp w^{T}X}\right)^{1 - y_{i}} \left(\frac{\exp w^{T}X}{1 + \exp w^{T}X}\right)^{y_{i}}$$
(24)

$$LL(y|X;w) = \sum_{i} \left[y_i w^T X_i - \ln\left(1 + \exp w^T X_i\right) \right]$$
 (25)

$$\frac{\partial}{\partial w^{j}} LL(y|X;w) = \sum_{i} X_{i}^{j} \left[y_{i} - p \left(y^{i} = 1 | X_{i}; w \right) \right]$$
 (26)

bad news: $no\ close\ form\ solution.$ good news: $concave\ function.$

• gradient ascent: z = x(y - g(w; x)) and its update rule (small ε):

$$w^{j} \leftarrow w^{j} + \varepsilon \sum_{i} X_{i}^{j} \left[y_{i} - \left(1 - g(X_{i}; w) \right) \right]$$
 (27)

until likelihood has no further improvement.

• regularization: additional constraints over w^j , e.g.: gaussian. in which case we have a prior: $p(y=1,\theta|X) \propto p(y=1|X;\theta)p(\theta)$.

$$LL'(y|X;w) = \sum_{i} \left[y_i w^T X_i - \ln\left(1 + \exp w^T X_i\right) \right] - \sum_{i} \frac{(w^i)^2}{2\sigma^2}$$
 (28)

and the update rule becomes

$$w^{j} \leftarrow w^{j} + \varepsilon \sum_{i} X_{i}^{j} \left[y_{i} - \left(1 - g(X_{i}; w) \right) \right] - \varepsilon \frac{w^{j}}{\sigma^{2}}$$
 (29)

• logistic regression for more than two classes: for i < k

$$p(y = i|X; \theta) = \frac{\exp z_i}{1 + \sum_{i=1}^{k-1} \exp z_i}$$
(30)

and for i = k,

$$p(y=i|X;\theta) = 1 - \sum_{i=1}^{k-1} p(y=i|X;\theta) = \frac{1}{1 + \sum_{j=1}^{k-1} \exp z_j}$$
(31)

and the derivative becomes

$$: \frac{\partial}{\partial w_m^j} LL(y|X; w) = \sum_i X_i^j \left[\delta_m(y_i) - p\left(y^i = m|X_i; w\right) \right]$$
(32)

$$w^{j} \leftarrow w^{j} + \varepsilon \sum_{i} X_{i}^{j} \left[\delta_{m}(y_{i}) - p\left(y^{i} = m | X_{i}; w\right) \right]$$
 (33)

• data transformation: linear \rightarrow generalized.

7 the perceptron

- logistic regression as *soft* linear classifier. signum smoothed to exp.
- stochastic gradient descent (SGD): one random sample i at a time: ¹

$$w^{j} \leftarrow w^{j} + \varepsilon X_{i}^{j} \left[y_{i} - p(y^{i} = 1 | X_{i}; w) \right]$$
(34)

• non-stochastic gradiate descent: taking the average over all samples.

$$w^{j} \leftarrow w^{j} + \varepsilon \frac{1}{n} \sum_{i} X_{i}^{j} \left[y_{i} - p(y^{i} = 1 | X_{i}; w) \right]$$
 (35)

trying to make expected value of $y|x^j = \cdot$ equal: data \leftarrow predicted.

- linear classifiers (boundary linear): naive bayes, logistic regression. questions: learning *easy*? performance?
- perceptron experiment: 1) A \rightarrow B x; 2) B predicts y; 3) A reveals y. = compute $\hat{y}_i = sign(\vec{v}_k \cdot \vec{x}_i)$, if wrong, $\vec{v}_{k+1} = \vec{v}_k + y_i \vec{x}_i$, where $y = \pm 1$.
 - rule 1: examples near the origin: $\forall \vec{x_i}, |\vec{x_i}|^2 \leq R^2$;
 - rule 2: examples separatable: $\exists \vec{u}, |u| = 1, \text{ s.t.}, \forall \vec{x}_i, (\vec{u} \cdot \vec{x}_i) y_i > \gamma.$

and two lemmas: $\forall k \colon \vec{v}_k \cdot \vec{u} \geq k \gamma$ (by rule 2); $\forall k, \, |v_k|^2 \leq k R^2$ (by rule 1). $\rightarrow k < R^2/\gamma^2$, meaning that the less than R^2/γ^2 mistakes are made.

- the Δ trick. add *label* features to the original features.
- the voted perceptron: online to batch learning. when picking at random, P(error) = k/m. where m = number of samples. but keeping \vec{v}_k could be expensive, so: sum of \vec{v}_k weighted by m_k/m .
- sparse vectors.

8 neural networks

- basic unit of a neural net: input: $x^i, w^i \to y = s(f(w^T X_i))$.
- matrix inversion vs. gradient descent: no iterations, no need to specify parameters, closed form solution; however, not universally applicable.
- assume $f(w^T X_i) = w^T X_i$: back to regression and gradient descent:

$$\frac{\partial}{\partial w_j} \sum_{i}^{n} \left(y_i - w^T X_i \right)^2 = -2 \sum_{i}^{n} x_i^j \left(y_i - w^T X_i \right) \tag{36}$$

¹See also: http://leon.bottou.org/publications/psgz/nimes-1991.ps.gz

• otherwise, assume sigmoid function:

$$\frac{\partial}{\partial w^j} \sum_{i} (y_i - g(w^T X_i))^2 = \sum_{i} 2 (y_i - g(w^T X_i)) g'(w^T X_i) X_i^j \qquad (37)$$

where g'(x) = -g(x)(1 - g(x)), and $g(x) = 1/(1 + \exp x)$.

• learn parameters of two-layer neural networks: *backpropagation*. take partial derivative for each of the weighted parameters:

$$err_i = (y_i - g(w^{(:)}^T g(w^{(:,j)}^T x_i))^2$$
 (38)

where $w^{(:)}$ is the weight vector for the last layer; $w^{(:,j)}$ for the second-last layer, and the inner $g(\cdot)$ produces a vector. for the hidden layer:

$$\frac{\partial}{\partial w^{k,j}} err_i = 2\underbrace{(y_i - g(w^T g(w^{j^T} x_i))(-g'(w^T g(w^{j^T} x_i))(w^j g'(w^{j^T} x_i))x_i^k}_{\Delta_i^j: \text{ the mystically } set \text{ term on the slides}}$$
(39)

where w^j is the j-th element of $w^{(:)}$. update rule revision is trivial. note that for consistency we might sum err over all samples.

- neural network encoding: an example.
- historical background:
 - 1st-gen (1960): perceptrons, hand-coded features;
 - 2nd-gen (1985): error signal + backpropagation.

back-propagation drawbacks: 1) labeled training data; 2) slow; 3) stuck in poor local optima. solution: 1) iteratively learn different layers; 2) weights adjusted to reproduce input. 3) learn features for unlabeled input.

9 support vector machine

- regression classifiers recalled: possibly many classifiers.

 max margin classifiers: boundary that leads to the largest margin from points on both sides, aka, support vector machine.
- boundary: $w^Tx + b = 0$: $w^Tx + b \ge +1 \to +1$, $w^Tx + b \le -1 \to -1$. margin m defined in terms of w and b how? observations:
 - \vec{m} is orthogonal to ± 1 planes;
 - for x^+ on +1 and x^- closest point to x^+ on -1: $x^+ = \lambda w + x^-$

together with $w^T x^{\pm} + b = \pm 1$, $|x^+ - x^-| = m$, $\to m = |\lambda w| = 2/\sqrt{w^T w}$, now search w that correctly classifies all points and maximize m.

- quadratic programming: $\min_{u} \left(\frac{1}{2} u^T R u + d^T u \right)$, constraints: $au \leq b$ and eu = f. where u, b, f, d vector, R diagonal, a, e matrix: usable on sym.
- non-linearly separable case: errors involved. if min $w^T w$ and min err, hard to solve; if min $(w^T w + err)$, hard to code in qp. \rightarrow solution: minimize distance ε_i between misclassified points and their correct planes:

$$\min_{w} \left(\frac{1}{2} w^T w + \sum_{i}^{n} C \varepsilon_i \right) \tag{40}$$

where $\forall x_i \in \text{Class}_{1,2}$: $w^T x + b \ge \pm 1 \mp \varepsilon_i$, C constant, with $\varepsilon_i \ge 0$.

• dual formulation: lagrange multiplier for sym optimization:

$$\min \frac{1}{2}(w^T w) \quad \text{subject to} \quad (w^T x_i + b) y_i \ge 1$$
 (41)

$$\min_{w,b} \max_{\alpha} \frac{w^T w}{2} - \sum_{i} \alpha_i \left[\left(w^T x_i + b \right) y_i - 1 \right] \quad \text{subject to} \quad \alpha_i \ge 0 \quad (42)$$

taking the ∂ -wrt w, α and b: $w = \sum_i \alpha_i x_i y_i$, $b = y_i - w^T x_i$, $\sum_i \alpha_i y_i = 0$, then substituting w back in to get the dual formulation:

$$\max_{\alpha} \left(\sum_{i} \alpha_{i} - \frac{1}{2} \sum_{i,j} \alpha_{i} \alpha_{j} y_{i} y_{j} x_{i} x_{j}^{T} \right), \sum_{i} \alpha_{i} y_{i} = 0 \quad \text{subject to} \quad \alpha_{i} \geq 0$$

$$(43)$$

- non-separable training set may be separable with additional dimensions. but: 1) high computation burden; 2) more parameters. svm solves them by 1) kernel tricks 2) dual formulation only assigns parameters to samples.
- quadratic kernels. the features with added dimensions are:

$$\Phi(x) = 1, \underbrace{\sqrt{2}x^{1}, \dots, \sqrt{2}x^{m}}_{m}, \underbrace{(x^{1})^{2}, \dots, (x^{m})^{2}}_{m}, \underbrace{\sqrt{2}x^{1}x^{2}, \dots, \sqrt{2}x^{m-1}x^{m}}_{\frac{1}{2}m(m-1)}$$
(44)

but $\forall i, j, \Phi(x_i) \cdot \Phi(x_j)^T$ costs $m + m + \frac{1}{2}m(m-1) = O(m^2)$ operations. however $(x_i \cdot x_j + 1)^2$ produces the same thing with only m operations. it also works for higher order scenarios. just find the correct kernels. comments: kernel is a trick! we're just doing projection to higher-dimension space, and happen to find that kernel makes it easier to compute dot-product! or, think of it as refining dot-product!

• for non-linearly separable case: bound on α .

10 model and feature selection

- model selection: features (linear, logistic, svm); parameters (prior, regularization, decision tree, clustering).
- greed heuristic model selection algorithm:
 - 1. start with a empty feature set F_0 ,
 - 2. $\forall X$, run alogrithm on $F_0 \cup X$; lowest training error $X_j \to F_{i+1}$.

optimal at each stage, but not necessarily global optimal. good for most classifiers so far. but stop when? use an independent data-set. data waste! better: leave-one-out cross validation. faster: k-fold cross validation.

- regularization: include all features but punish large parameters (logistic), zero probability outcomes (naive bayes). chi-square test (decision tree). e.g.: bayesian learning: λ selected how? **cross validation!**
- minimum description length (mdl): the information theoretic interpretation of regularization: minimize length of data and length of hypothesis.
- the confidence interval bounds:

$$Accuracy_s(h) \pm Z_n \sqrt{\frac{Accuracy_s(h)(1 - Accuracy_s(h))}{n}}$$
 (45)

11 computational learning theory

- problem setting: instances X, hypotheses H, target functions C, sequence of training instances, noise-free labels c(x): outputs $h = \arg\min_{h \in H} error_{train}(h)$. $error_{train}(h)$ and $error_{true}(h)$. overfitting, confidence interval bounding.
- some definitions:
 - **consistent**: $\forall (x, c(x)) \in D, h(x) = c(x);$
 - version space: all $h \in H$ consistent with D;
 - ε-exhausted: $\forall h \in VS_{H,D}, error_{true} < ε$.

and a theorem:

$$-Pr(\exists h \in H, \text{s.t.}, error_{train} = 0 \land error_{true} > \epsilon) \le |H|e^{-\epsilon m}$$

how many training examples? $m \ge \epsilon^{-1} (\ln |H| - \ln \delta)$; if $error_{train} = 0$, with at least $1 - \delta$: $error_{true} \le m^{-1} (\ln |H| - \ln \delta)$.

- pac-learnable: learning in polynomial time.
- more definitions:
 - **dichotomy**: partition into two disjoint subsets.

- **shattered**: S shattered by H if \exists consistent $h \in H \to \text{dichotomy}$.
- vc-dimension: size of largest finite subset of X shattered by H.

to guarantee $\forall h \in H$ perfectly fits training data, need instances

$$m \ge \frac{1}{\epsilon} \left(4\log_2 \frac{2}{\epsilon} + 8\text{VC}(H)\log_2 \frac{13}{\epsilon} \right)$$
 (46)

- to show that VC(H) = d, we need to
 - There exists a *d*-sized subset shatterable with arbitrary labels;
 - There is no subset of size d+1 the can be shattered.
- agnostic learning: when the true classifier $c \notin H$, we can still bound the true error by training error, and see how close we are to $h^* \in H$ that has a lowest true error, with a probability at least 1δ .

$$error_{true}(h) \leq error_{train}(h) + \sqrt{\frac{\ln(|H|) + \ln(1/\delta)}{2m}}$$

$$error_{true}(h) \leq error_{train}(h) + \sqrt{\frac{VC(|H|)(\ln\frac{2m}{VC(H)} + 1) + \ln(4/\delta)}{m}}$$

$$(47)$$

• mistake bounds, optimal mistake bounds, (c.f.: the find-s example),

$$VC(C) \le \min_{A} M_A(C) \le \log_2(|C|) \tag{49}$$

where $M_A(C)$ is the max number of mistakes when learning C using A.

• upper bound of VC(H) as a function of |H|.

12 clustering

- finding internal structure, underlying rules, recurring patterns and topics... unsupervised learning: no provided labels. **distance**.
- type i: hierarchical: (bottom-up) repetitively find closest pairs and merge. question: how to compute the distance between clusters?
 - single link: distance of two closest members,
 - complete link: distance of two farthest members,
 - average link: average distance of pairs. (robust against noise)

summary: 1) no need to specify the number of clusters, 2) structure interpreted subjectively, 3) poor scalibility, 4) local optima 5) intuitive. using the dendrogram to detect outliers.

- type ii: partitional: specify number of clusters, and place the instances. (top-down) graph-based clustering: the steps:
 - construct the neighborhood graph,
 - assign weights to edges (similarity),
 - partition the nodes using the graph structure.

how to find the number of clusters? knee/elbow finding!

- cluster validation:
 - internal validation: coherence. average intra/inter-cluster similarity.
 requiring the definition of similarity and distance metric.
 - internal validation: stability. subject to minor perturbation. requiring measurement of clusters distance: label matrices. (subsampling)
 - external validation: relation to existent categories. the p-value: randomly chosen cluster element exists in some category G.

13 more clustering

- distance: minkowski metric, hamming distance, correlation coefficient,
- k-means algorithm: repeat:
 - 1. decide k and initialize their centers randomly,
 - 2. decide memberships by assigning objects to nearest cluster centroid,
 - 3. move the cluster centers closer to the objects assigned to them.

parallelized easily (map-reduce); sensitive to starting points.

• mixture models (multi-modal density model): gaussian mixture models where all variables are observable:

$$p(x_n|\mu, \Sigma) = \sum_{i=1}^{k} \pi_i N(x|\mu_i, \Sigma_i)$$
 (50)

and the likelihood for a bunch of data D would be

$$l(\theta; D) = \log \prod_{n} p(z_{n}, x_{n}) = \log \prod_{n} p(z_{n} | \pi) p(x_{n} | z_{n}; \mu, \sigma)$$

$$= \sum_{n} \log \prod_{k} \pi_{k}^{z_{n}^{k}} + \sum_{n} \log \prod_{k} N(x_{n}; \mu_{k}, \sigma_{k})^{z_{n}^{k}}$$

$$= \sum_{n} \sum_{k} z_{n}^{k} \log \pi_{k} - \sum_{n} \sum_{k} z_{n}^{k} \frac{1}{2\sigma_{k}^{2}} (x_{n} - \mu_{k})^{2}$$
(51)

which gives the parameter estimates as follows:

$$\hat{\pi} = \sum_{i} z_{i}^{k} / |D|; \quad \hat{\mu} = \frac{\sum_{n} z_{n}^{k} x_{n}}{\sum_{n} z_{n}^{k}}; \quad \hat{\sigma} = ?$$
 (52)

• exceptation-maximization (em) algorithm: we're not going through this... but know that K-means is a hard version of em.

14 bias-variance

• as a way to understand overfitting and underfitting: decomposition. setting: true function f, noise ε , $h_D(\cdot)$ hypothesis learned from D, error:

error =
$$E_{D,\varepsilon} \left\{ (f(x) + \varepsilon - h_D(x))^2 \right\}$$
 (53)

more notations: $f \equiv f(x) + \varepsilon$, $\hat{y} = \hat{y}_D \equiv h_D(x)$, and fact: $h \equiv E_D \{h_D(x)\}$.

error =
$$E_{D,\varepsilon} \left\{ (f - \hat{y})^2 \right\} = E_{D,\varepsilon} \left[(f - h)^2 \right] + E_{D,\varepsilon} \left[(h - \hat{y})^2 \right]$$
 (54)

where the first term is $BIAS^2$ (you tried so hard but still couldn't get to the original f), second VARIANCE (your each individual attempt may deviate from your best performance.)

- tradeoffs: *large bias, small variance*: few features, highly regularized, highly pruned decision trees, large k-NN. *large variance, small bias*: many features, less regularization, unpruned trees, small k-NN.
- generalization: (which is not confusing at all):
 - optimal prediction: $y^* = \arg\min_{y'} L(t, y')$;
 - main prediction of **learner**: $y_m = y_{m,D} = \arg\min_{y'} E_D \{L(y,y')\};$
 - bias of **learner**: BIAS = $L(y^*, y_m)$;
 - variance of **learner**: VARIANCE = $E_D[L(y_m, y)]$;
 - noise: NOISE = $E_t[L(t, y^*)]$

bias: more expressive (complex)! variance: less expressive (complex)!

- how to approximate E_D $\{h_D(x)\}$? background: bootstrap sampling: input: dataset D, output: variants: D_1, D_2, \ldots, D_T . algorithm: for each $D_{(\cdot)}$, pick |D| instances uniformly from D. then, train on $D_{(\cdot)}$ and test on remaining data to get $h_1(x), h_2(x), \ldots$ variance = ordinary variance of $h_{(\cdot)}(x)$, bias = $\bar{h}_{(\cdot)}(x) y$.
- bagging: bootstrap aggregation. **input**: dataset D, and YFCL (wtf); **output**: a classifier $h_{D,BAG}$. **algorithm**: get $D_{(\cdot)}$ and train YFCL on it to get $h_{(\cdot)}$, then to classify do majority vote with $h_{(\cdot)}$. generally bagged decision trees outperform linear classifiers if the data is large and clean.
- more details on the confusing generalization: for (x^*, y^*) a single instance.
 - noisy channel: $y_i = \text{noise}(f(x_i))$, may change $y \to y'$;

- learned hypothesis: $h = h_D$ from some dataset D;
- main prediction of **learner**: $y_m(x^*) = \arg\min_{y'} E_{D,P} \{L(h(x^*), y')\};$
- bias of **learner**: $B(x^*) = L(y_m(x^*), f(x^*));$
- variance of **learner**: $V(x^*) = E_{D,P} \{ L(h_D(x^*), y_m(x^*)) \}$
- noise: $N(x^*) = L(y^*, f(x^*)).$

and yes, we can do case analysis. but no.

15 boosting

- make weaker classifiers better: lemma ideas:
 - simple: models \rightarrow one with lowest training error, but what if many?
 - stacked learners: train new classifier on predicted and true labels.
- ensemble methods: divide and conquer. a final classifier = a linear combination of votes of different classifiers weighted by their strength. weighted data: i-th training example counts as D(i) examples.
- adaboost: given (x_i, y_i) where $x_i \in X$, $y_i \in Y = \{-1, +1\}$, i = 1, 2, ..., m. initialize $D_1(i) = m^{-1}$, and iterator over $t \in \{1, 2, ..., T\}$:
 - 1. train weak learner using distribution D_t and get $h_t: X \to \mathbb{R}$;
 - 2. choose $\alpha_t \in \mathbb{R}$, and update

$$D_{t+1}(i) = \frac{D_t(i) \exp(-\alpha_t y_i h_t(x_i))}{Z_t = \sum_{i=1}^m D_t(i) \exp(-\alpha_t y_i h_t(x_i))}$$
(55)

and output the final classifier:

$$H(x) = \operatorname{sign} f(x) = \operatorname{sign} \left(\sum_{t=1}^{T} \alpha_t h_t(x) \right)$$
 (56)

• problem: how to choose α_t ? final training error is bounded by:

$$\frac{1}{m} \sum_{i=1}^{m} \delta(H(x_i) \neq y_i) \le \frac{1}{m} \sum_{i=1}^{m} \exp(-y_i f(x_i)) = \prod_{t} Z_t$$
 (57)

so minimizing Z_t minimizes the training error. we define that

$$\epsilon_t = \sum_{i=1}^m D_t(i)\delta(h_t(x_i) \neq y_i) \quad \text{s.t.,} \quad Z_t = (1 - \epsilon_t)e^{-\alpha_t} + \epsilon_t e^{\alpha_t}$$
 (58)

hence take $\partial Z_t/\partial \alpha_t$ tells you that $\alpha_t = \frac{1}{2} \ln \frac{1-\epsilon_t}{\epsilon_t}$, bang!

• if each classifier is better than random, this gives you zero training error exponentially fast. also: robust to overfitting, testing error decreases after.