Machine Learning Cheat Sheet

General

Dataset and Definitions

 \mathcal{D} is a set of training examples, the n-th Training Example (n = 1, 2, ..., N), of this set is:

$$\mathbf{x_n} = \begin{bmatrix} x_{n1} & x_{n2} & \dots & x_{nD} \end{bmatrix}$$

We write all N training examples in a matrix:

$$\mathbf{X} = \begin{bmatrix} \mathbf{x_1}; & \mathbf{x_2}; & ...; & \mathbf{x_N} \end{bmatrix}$$

In supervised learning, you are also given a set of observations corresponding to the training examples: $\mathbf{y} = \begin{bmatrix} y_1 & y_2 & \dots & y_N \end{bmatrix}^{\mathbf{T}}$

We assume a true model behind the data, the

observations are noisy versions of this ground truth: $y = y_{true} + \epsilon$.

The final goal is to predict a \hat{y} given any \mathbf{x} .

Distributions

Multivariate gaussian distribution: $\mathcal{N}(\mathbf{X}|\mu, \Sigma)$

$$\implies p(\mathbf{X} = \mathbf{x}) =$$

$$(2\pi)^{-\mathbf{d}/2} |\Sigma|^{-\mathbf{1}/2} \exp\left[-\frac{1}{2}(\mathbf{x}-\mu)^{\mathrm{T}} \Sigma^{-1}(\mathbf{x}-\mu)\right]$$

Gaussian distribution:
$$\mathcal{N}(X|\mu,\sigma^2)$$
 $\implies p(X=x) = \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left(-\frac{1}{2}(\frac{x-\mu}{\sigma})^2\right)$ Poisson distribution: $\mathcal{P}(X|\lambda)$

$$\implies p(X = k) = \frac{\lambda^k}{k!} \exp(-\lambda)$$

Formulas

Jensen's inequality applied to log: $\log(\frac{\sum_{i=1}^{n} x_i}{\pi}) \ge \frac{\sum_{i=1}^{n} \log(x_i)}{\pi}$

Optimization methods

Grid Search

Simply try values for all parameters at regular intervals. Complexity: $\mathcal{O}(M^DND)$, where M is the number of values tried in each dimension.

Gradient Descent

Update rule: $\boldsymbol{\beta}^{(k+1)} = \boldsymbol{\beta}^{(k)} - \alpha \frac{\partial \mathcal{L}(\boldsymbol{\beta}^{(k)})}{\partial \boldsymbol{\beta}}$

Complexity: $\mathcal{O}(IND)$ where I is the number of iterations we take.

 α is a parameter that needs to be chosen carefully. The gradient for MSE comes out as:

 $\frac{\partial \mathcal{L}}{\partial \boldsymbol{\beta}} = -\frac{1}{N} \tilde{X}^T (\boldsymbol{y} - \tilde{X} \boldsymbol{\beta})$

Newton's method

It uses second order derivatives information to converge faster (we approximate the objective function by a quadratic function rather than a line).

General rule: $\boldsymbol{\beta}^{(k+1)} = \boldsymbol{\beta}^{(k)} - \alpha \mathbf{H}_{\mathbf{k}}^{-1} \frac{\partial \mathcal{L}(\boldsymbol{\beta}^{(\mathbf{k})})}{\partial \boldsymbol{\beta}}$ where $\mathbf{H}_{\mathbf{k}}$ is the $D \times D$ Hessian at step k: $\mathbf{H}_{\mathbf{k}} = \frac{\partial^2 \mathcal{L}(\boldsymbol{\beta}^{(\mathbf{k})})}{\partial \boldsymbol{\beta}^2}$

$$\mathbf{H_k} = \frac{\partial^2 \mathcal{L}(\boldsymbol{\beta}^{(\mathbf{k})})}{\partial \boldsymbol{\beta}^2}$$

Newton's method is equivalent to solving many least-squares problems.

Complexity: $\mathcal{O}(IND^2 + ID^3)$, with the D^3 cost coming from the inversion of H_k .

Regression

Simple linear regression: $y_n \approx \beta_0 + \beta_1 x_{n1}$ Multiple linear regression:

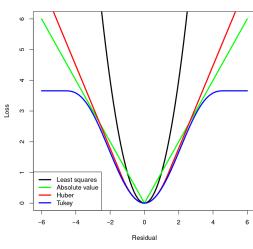
 $y_n \approx f(\mathbf{x_n}) := \beta_0 + \beta_1 \mathbf{x_{n1}} + \beta_2 \mathbf{x_{n2}} + \dots + \beta_D \mathbf{x_{nD}}$

Linear basis function model

 $y_n = \beta_0 + \sum_{i=1}^M \beta_i \phi_i(\mathbf{x_n}) = \tilde{\phi}^{\mathbf{T}}(\mathbf{x_n^T}) \boldsymbol{\beta}$. The optimal β is given by $\beta = (\tilde{\Phi}^T \tilde{\Phi})^{-1} \tilde{\Phi}^T y$ where $\tilde{\Phi}$ is a matrix with N rows and the n-th row is $[1, \phi_1(x_n)^T, ..., \phi_M(x_n)^T].$

Ridge regression: $\beta_{ridge} = (\tilde{\pmb{\Phi}}^T \tilde{\pmb{\Phi}} + \lambda \pmb{I})^{-1} \tilde{\pmb{\Phi}}^T \pmb{y}$

Cost functions



Cost function / Loss: $\mathcal{L}(\beta) = \mathcal{L}(\mathcal{D}, \beta)$ Mean square error (MSE): $\frac{1}{2N} \sum_{n=1}^{N} [y_n - f(\mathbf{x_i})]^2$ MSE matrix formulation: $\frac{1}{2N}(\mathbf{y} - \mathbf{X}\beta)^{\mathbf{T}}(\mathbf{y} - \mathbf{X}\beta)$ Mean absolute error (MAE): $\frac{1}{2N} \sum_{n=1}^{N} |y_n - f(\mathbf{x_i})|$

 $\text{Huber loss: } \mathcal{L}_{\delta}(a) = \begin{cases} \frac{1}{2}a^2 & \text{for } |a| \leq \delta, \\ \delta(|a| - \frac{1}{2}\delta), & \text{otherwise.} \end{cases}$

Root mean square error (RMSE): $\sqrt{2 * MSE}$ Epsilon insensitive ("hinge loss", used for SVM):

$$\mathcal{L}_{\epsilon}(y, \hat{y}) = \begin{cases} 0 & \text{if } |y - \hat{y}| \leq \epsilon, \\ |y - \hat{y}| - \epsilon, & \text{otherwise.} \end{cases}$$

Least squares

Complexity: $\mathcal{O}(ND^2 + D^3)$ The gradient of the MSE set to zero is called the normal equation:

$$-\mathbf{y}^{\mathbf{T}}\mathbf{X} + \mathbf{X}^{\mathbf{T}}\mathbf{X}\boldsymbol{\beta} = \mathbf{0}$$

We can solve this for β directly (by matrix manipulation) $\beta = (\tilde{X}^T \tilde{X})^{-1} \tilde{X}^T u$

Classification

Logistic Function $\sigma = \frac{exp(x)}{1 + exp(x)}$

Classification with linear regression: Use y = 0 as class \mathcal{C}_{∞} and y=1 as class \mathcal{C}_{\in} and then decide a newly estimated y belongs to \mathcal{C}_{∞} if y < 0.5.

Logistic Regression

$$\tilde{\mathbf{X}}^T [\sigma(\tilde{\mathbf{X}}\beta) - y] = 0$$

Generalized linear model

The GLM consists of three elements:

- A probability distribution from the exponential family.
- A linear predictor $\hat{y} = \mathbf{X}\beta$.
- A link function g such that $E(y) = = q^{-1}()$.

In a generalized linear model (GLM), each outcome of the dependent variables y is assumed to be generated from a particular distribution in the exponential family, a large range of probability distributions that includes the normal, binomial. Poisson and gamma distributions, among others.

Cost functions

Root Mean square error (RMSE):

$$\sqrt{\frac{1}{N}\sum_{n=1}^{N}\left[y_n-\hat{p_n}\right]^2}$$

0-1 Loss: $\frac{1}{N} \sum_{n=1}^{N} \delta(y_n, \hat{y_n})$

 $-\frac{1}{N}\sum_{n=1}^{N}y_n\log(\hat{p_n})+(1-y_n)\log(1-\hat{p_n})$

Maximum Likelihood

The Likelihood Function maps the model parameters to the probability distribution of v:

 \mathcal{L}_{lik} : parameter space $\rightarrow [0;1] \quad \beta \mapsto \mathbf{p}(\mathbf{y} \mid \beta)$ An underlying p is assumed before. If the observed y are IID, $p(\mathbf{y} \mid \beta) = \prod_{\mathbf{n}} \mathbf{p}(\mathbf{y_n} \mid \beta)$.

 \mathcal{L}_{lik} can be viewed as just another cost function. Maximum likelihood then simply chooses the parameters β such that observed data is most likely. $\beta = \arg\max_{\mathbf{a} \parallel \beta} L(\beta)$

Assuming different p is basically what makes this so flexible. We can chose e.g.:

Gaussian p $\mathcal{L}_{lik} = -\mathcal{L}_{MSE}$ Poisson p $\mathcal{L}_{lik} = -\mathcal{L}_{MAE}$

Bayesian methods

Bayes rule: p(A, B) = p(A|B)p(B) = p(B|A)p(A)The **prior** $p(\mathbf{f}|\mathbf{X})$ encodes our prior belief about the "true" model ${\bf f}$. The likelihood $p({\bf y}|{\bf f})$ measures the probability of our (possibly noisy) observations given the prior.

Least-squares tries to find model parameters β which maximize the likelihood. Ridge regression maximizes the **posterior** $p(\beta|\mathbf{y})$

Graphical Models

TODO: Bayes Net: Directed acyclic graph

TODO: Belief propagation

Graph between the observations and the variables is a bi-partite graph.

Kernel methods

Basically, Kernels are a mean to measure distance, or "similarity" of two vectors. We define:

$$(\mathbf{K})_{\mathbf{i},\mathbf{j}} = \kappa(\mathbf{x}_{\mathbf{i}}, \mathbf{x}_{\mathbf{j}}) = \tilde{\phi}(\mathbf{x}_{\mathbf{i}})^{\mathbf{T}} \tilde{\phi}(\mathbf{x}_{\mathbf{j}}).$$

The ϕ are not that important in the end, because we only use the Kernel as is. Sometimes it's even impossible to write them down explicitly.

Linear	$\kappa(\mathbf{x_i}, \mathbf{x_j}) = \mathbf{x_i}^T \mathbf{x_j}$
Polynomial	$\kappa(\mathbf{x_i}, \mathbf{x_j}) = (\mathbf{x_i}^T \mathbf{x_j} + \mathbf{c})^d$
RBF	$\kappa(\mathbf{x_i}, \mathbf{x_j}) = \exp\left(-\frac{ \mathbf{x_i} - \mathbf{x_j} ^2}{2\sigma^2}\right)$

Properties of a Kernel:

- K should be symmetric.
- K should be positive semi-definite $\forall x_i, x_i$.

Neural Networks

TODO: Intuition and notation, backpropagation, regularization techniques

Support Vector Machines

Search for the hyperplane separating the data such that the gap is biggest. It minimizes the following cost function:

$$\mathcal{L}_{SVM}(\beta) = \sum_{\mathbf{n=1}}^{\mathbf{N}} [\mathbf{1} - \mathbf{y_n} \tilde{\phi}_{\mathbf{n}} \beta]_{+} + \frac{\lambda}{2} \sum_{\mathbf{j=1}}^{\mathbf{M}} \beta_{\mathbf{j}}^2$$

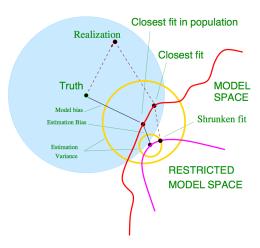
This is convex but not differentiable.

Concepts

Occam's Razor

It states that among competing hypotheses, the one with the fewest assumptions should be selected. Other, more complicated solutions may ultimately prove correct, butin the absence of certaintythe fewer assumptions that are made, the better.

Bias-Variance Decomposition



Bias-variance come directly out of the test error:

$$\begin{split} \overline{teErr} &= E[(\text{observation} - \text{prediction})^2] = E[(y-\hat{y})^2] \\ &= E[(y-y_{true} + y_{true} - \hat{y})^2] \\ &= \underbrace{E[(y-y_{true})^2]}_{\text{var of measurement}} + E[(y_{true} - \hat{y})^2] \\ &= \sigma^2 + E[(y_{true} - E[\hat{y}] + E[\hat{y}] - \hat{y})^2] \\ &= \sigma^2 + \underbrace{E[(y_{true} - E[\hat{y}])^2]}_{\text{pred bias}^2} + \underbrace{E[(E[\hat{y}] - \hat{y})^2]}_{\text{pred variance}} \end{split}$$

	bias	variance
regularization	+	-
reduce model complexity	+	-
more data	-	

Consistency

An estimator is said to be consistent, if it eventually recovers the true parameters that generated the data as the sample size goes to infinity. Of course, this only makes sense if the data actually comes from the specified model, which is not usually the case. Nevertheless, it can be a useful theoretical property.

Convexity

f is called convex f:
$$\forall x_1, x_2 \in X, \forall t \in [0, 1]$$
: $f(tx_1 + (1 - t)x_2) < tf(x_1) + (1 - t)f(x_2)$.

Efficiency

An estimator is called efficient if it achives the Cramer-Rao lower bound: $\mathrm{Var}(\beta) \geq 1/I(\beta)$, where I is the fisher information.

Curse of dimensionality

The main reason behind the curse is that, with the dimensionality increase, every data point becomes arbitrarily far from every other data point and therefore the choice of nearest neighbor becomes random. In high dimension, data only covers a tiny fraction of the input space, making generalization extremely difficult. Or in other words, the volume of the space increases so fast that the available data become sparse.

Primal vs. Dual

Instead of working in the column space of our data, we can work in the row space:

$$\hat{\mathbf{y}} = \mathbf{X}\boldsymbol{\beta} = \mathbf{X}\mathbf{X}^{\mathbf{T}}\boldsymbol{\alpha} = \mathbf{K}\boldsymbol{\alpha}$$

where $\beta \in \mathbb{R}^{\mathbf{D}}$ and $\alpha \in \mathbb{R}^{\mathbf{N}}$ and (like magic) \mathbf{K} shows up, the Kernel Matrix.

Representer Theorem: For any β minimizing

$$\min_{\beta} \sum_{n=1}^{N} \mathcal{L}(y_n, \mathbf{x_n^T} \beta) + \sum_{\mathbf{d}=1}^{\mathbf{D}} \lambda \beta_{\mathbf{d}}^{\mathbf{2}}$$

there exists an α such that $\beta = \mathbf{X}^{\mathbf{T}} \alpha$.

When we have an explicit vector formulation of β , we can use the matrix inversion lemma to get it. E.g. for ridge regression:

$$\beta = (X^T X + \lambda I_D)^{-1} X^T y = X^T \underbrace{(X X^T + \lambda I_N)^{-1} y}_{}$$

k-fold cross-validation, definition of Test-Error, Train-Error

TODO: statistical/computational tradeoff

Definitions of: Optimality, Identifiability

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