

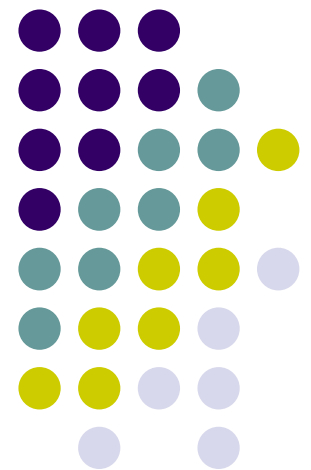
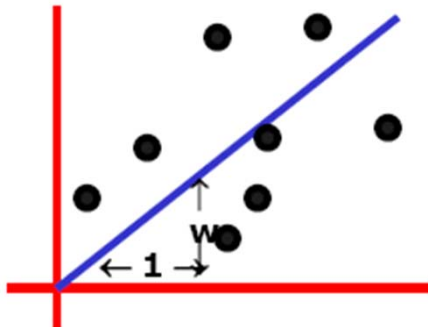
Advanced Introduction to Machine Learning

10715, Fall 2014

Linear Regression and Sparsity

Eric Xing

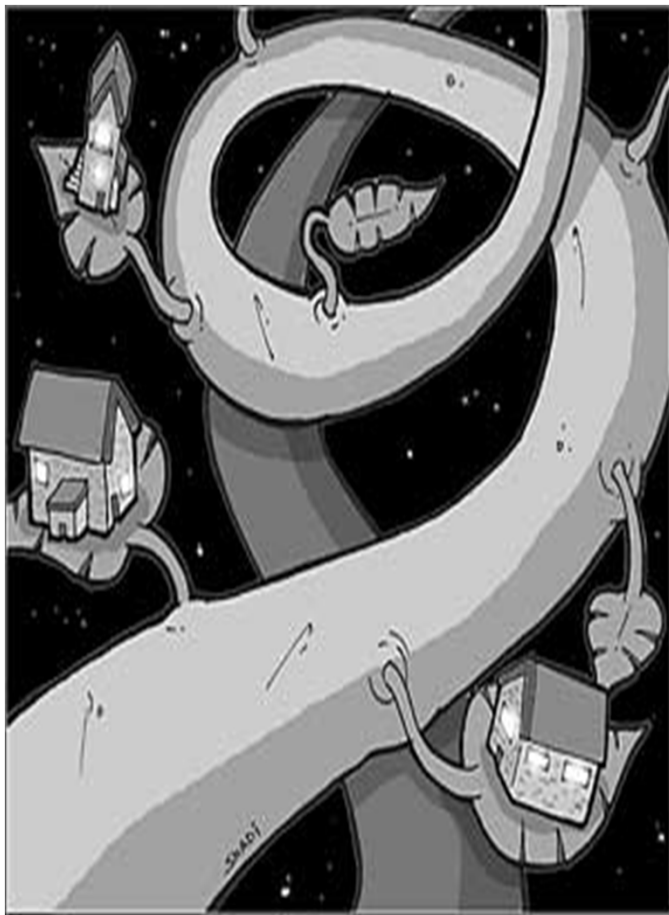
Lecture 2, September 10, 2014



Reading:

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Machine learning for apartment hunting



- Now you've moved to Pittsburgh!!

And you want to find the **most reasonably priced** apartment satisfying your **needs**:

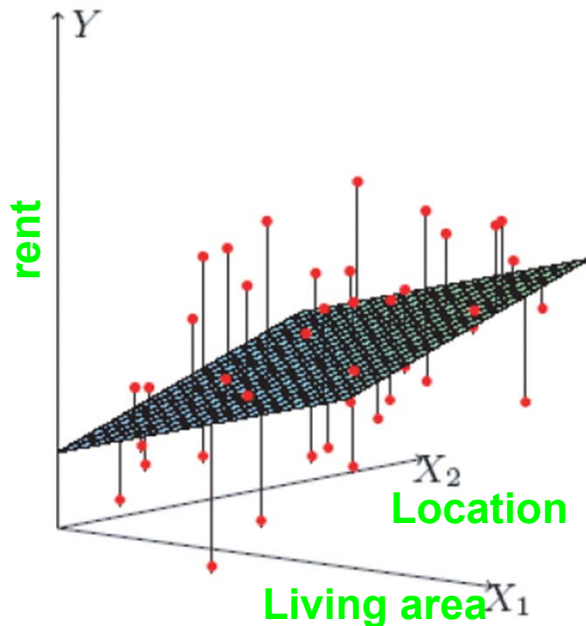
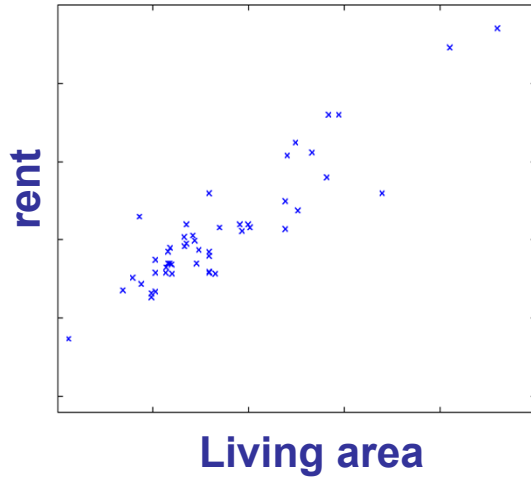


square-ft., # of bedroom, distance to campus ...

Living area (ft ²)	# bedroom	Rent (\$)
230	1	600
506	2	1000
433	2	1100
109	1	500
...		
150	1	?
270	1.5	?



The learning problem



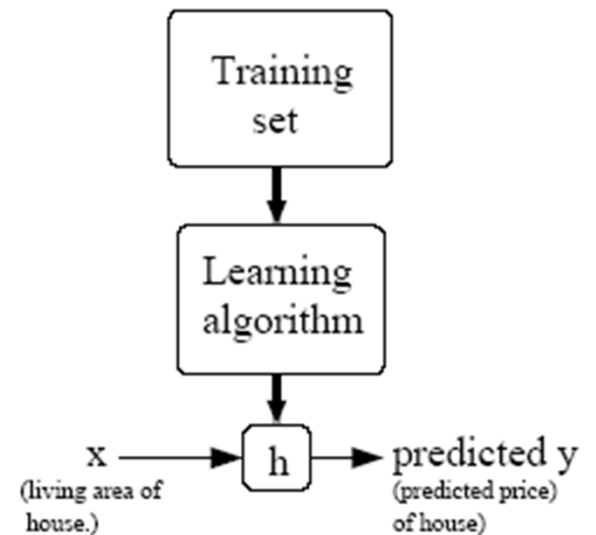
- Features:
 - Living area, distance to campus, # bedroom ...
 - Denote as $\mathbf{x}=[x^1, x^2, \dots x^k]$
- Target:
 - Rent
 - Denoted as y
- Training set:

$$\mathbf{X} = \begin{bmatrix} \text{---} & \mathbf{x}_1 & \text{---} \\ \text{---} & \mathbf{x}_2 & \text{---} \\ \vdots & \vdots & \vdots \\ \text{---} & \mathbf{x}_n & \text{---} \end{bmatrix} = \begin{bmatrix} x_1^1 & x_1^2 & \dots & x_1^k \\ x_2^1 & x_2^2 & \dots & x_2^k \\ \vdots & \vdots & \vdots & \vdots \\ x_n^1 & x_n^2 & \dots & x_n^k \end{bmatrix}$$

$$\mathbf{Y} = \begin{bmatrix} y_1 \\ y_2 \\ \vdots \\ y_n \end{bmatrix} \quad \text{or} \quad \begin{bmatrix} \text{---} & \mathbf{y}_1 & \text{---} \\ \text{---} & \mathbf{y}_2 & \text{---} \\ \vdots & \vdots & \vdots \\ \text{---} & \mathbf{y}_n & \text{---} \end{bmatrix}$$

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Our goal:





Linear Regression

- Assume that Y (target) is a linear function of X (features):

- e.g.:

$$\hat{y} = \theta_0 + \theta_1 x^1 + \theta_2 x^2$$

- let's assume a vacuous "feature" $X^0=1$ (this is the **intercept** term, why?), and define the feature vector to be:

- then we have the following general representation of the linear function:

- Our goal is to pick the optimal θ . How!

- We seek θ that minimize the following **cost function**:

$$J(\theta) = \frac{1}{2} \sum_{i=1}^n (\hat{y}_i(\bar{x}_i) - y_i)^2$$

The Least-Mean-Square (LMS) method



- The Cost Function:

$$J(\theta) = \frac{1}{2} \sum_{i=1}^n (\mathbf{x}_i^T \theta - y_i)^2$$

- Consider a **gradient descent** algorithm:

$$\theta_j^{t+1} = \theta_j^t - \alpha \left. \frac{\partial}{\partial \theta_j} J(\theta) \right|_t$$

The Least-Mean-Square (LMS) method



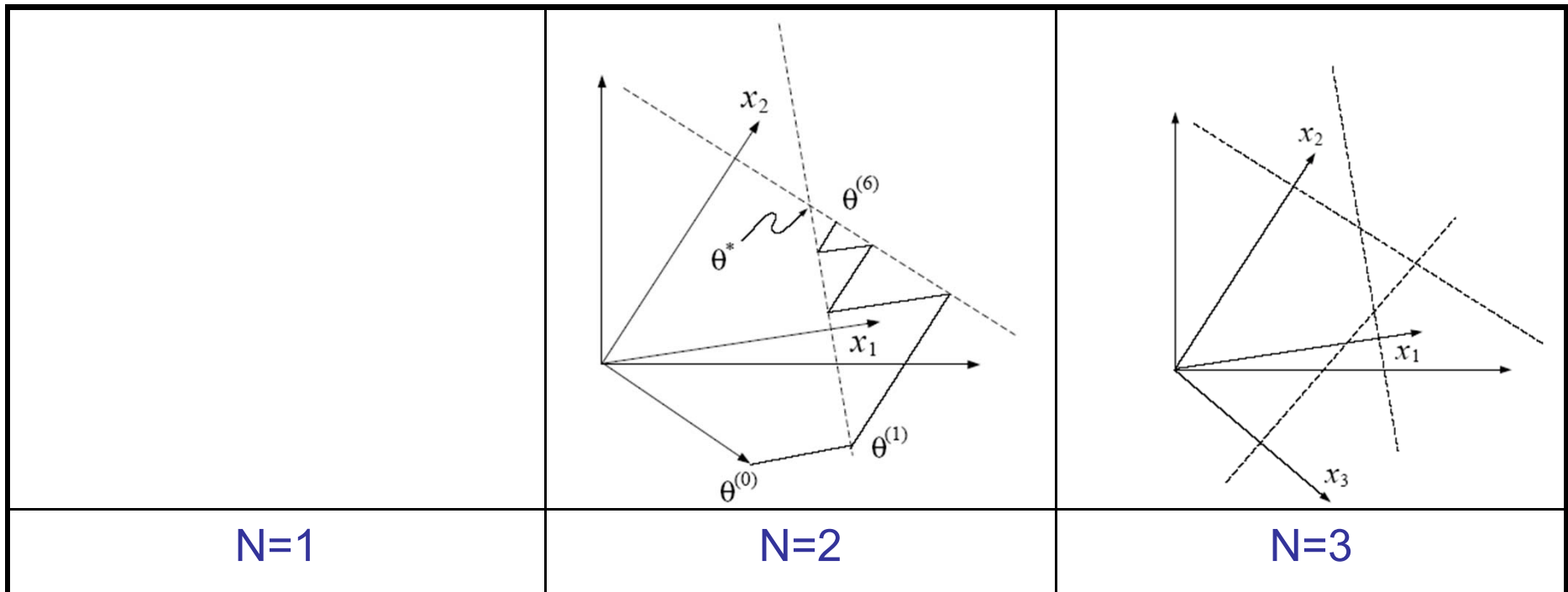
- Now we have the following descent rule:

$$\theta_j^{t+1} = \theta_j^t + \alpha \sum_{i=1}^n (y_i - \bar{\mathbf{x}}_i^T \theta^t) x_i^j$$

- For a single training point, we have:
 - This is known as the LMS update rule, or the Widrow-Hoff learning rule
 - This is actually a "**stochastic**", "**coordinate**" descent algorithm
 - This can be used as a **on-line** algorithm



Geometry and Convergence of LMS



$$\theta^{t+1} = \theta^t + \alpha(y_i - \bar{\mathbf{x}}_i^T \theta^t) \bar{\mathbf{x}}_i$$

Claim: when the step size α satisfies certain condition, and when certain other technical conditions are satisfied, LMS will converge to an “optimal region”.

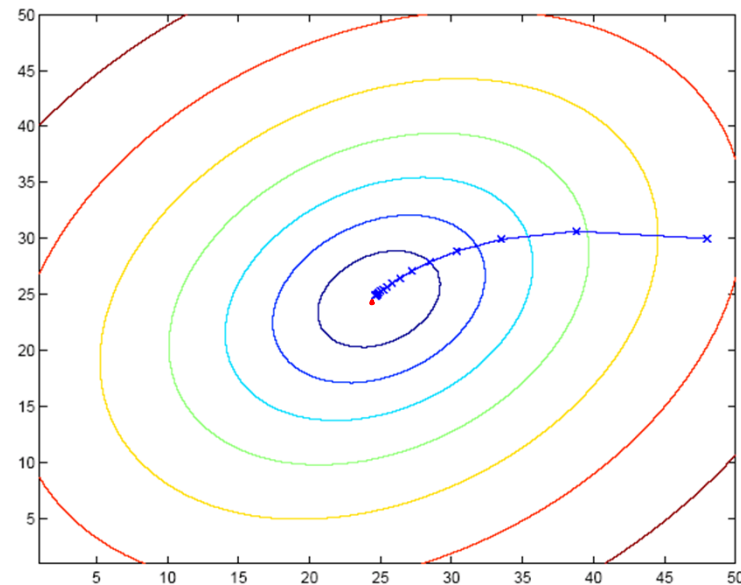


Steepest Descent and LMS

- Steepest descent
 - Note that:

$$\nabla_{\theta} J = \left[\frac{\partial}{\partial \theta_1} J, \dots, \frac{\partial}{\partial \theta_k} J \right]^T = - \sum_{i=1}^n (y_n - \mathbf{x}_n^T \theta) \mathbf{x}_n$$

$$\theta^{t+1} = \theta^t + \alpha \sum_{i=1}^n (y_n - \mathbf{x}_n^T \theta^t) \mathbf{x}_n$$



- This is as a **batch** gradient descent algorithm



The normal equations

- Write the cost function in matrix form:

$$\begin{aligned} J(\theta) &= \frac{1}{2} \sum_{i=1}^n (\mathbf{x}_i^T \theta - y_i)^2 \\ &= \frac{1}{2} (X\theta - \bar{\mathbf{y}})^T (X\theta - \bar{\mathbf{y}}) \\ &= \frac{1}{2} (\theta^T X^T X\theta - \theta^T X^T \bar{\mathbf{y}} - \bar{\mathbf{y}}^T X\theta + \bar{\mathbf{y}}^T \bar{\mathbf{y}}) \end{aligned}$$

$$\mathbf{X} = \begin{bmatrix} \text{---} & \mathbf{x}_1 & \text{---} \\ \text{---} & \mathbf{x}_2 & \text{---} \\ \vdots & \vdots & \vdots \\ \text{---} & \mathbf{x}_n & \text{---} \end{bmatrix}$$
$$\bar{\mathbf{y}} = \begin{bmatrix} y_1 \\ y_2 \\ \vdots \\ y_n \end{bmatrix}$$

- To minimize $J(\theta)$, take derivative and set to zero:

$$\begin{aligned} \nabla_{\theta} J &= \frac{1}{2} \nabla_{\theta} \text{tr}(\theta^T X^T X\theta - \theta^T X^T \bar{\mathbf{y}} - \bar{\mathbf{y}}^T X\theta + \bar{\mathbf{y}}^T \bar{\mathbf{y}}) \\ &= \frac{1}{2} (\nabla_{\theta} \text{tr} \theta^T X^T X\theta - 2\nabla_{\theta} \text{tr} \bar{\mathbf{y}}^T X\theta + \nabla_{\theta} \text{tr} \bar{\mathbf{y}}^T \bar{\mathbf{y}}) \\ &= \frac{1}{2} (X^T X\theta + X^T X\theta - 2X^T \bar{\mathbf{y}}) \\ &= X^T X\theta - X^T \bar{\mathbf{y}} = \mathbf{0} \end{aligned}$$

$$\Rightarrow \boxed{X^T X\theta = X^T \bar{\mathbf{y}}}$$

The normal equations

$$\Downarrow$$
$$\theta^* = (X^T X)^{-1} X^T \bar{\mathbf{y}}$$



Some matrix derivatives

- For $f : \mathbb{R}^{m \times n} \mapsto \mathbb{R}$, define:

$$\nabla_A f(A) = \begin{bmatrix} \frac{\partial}{\partial A_{11}} f & \cdots & \frac{\partial}{\partial A_{1n}} f \\ \vdots & \ddots & \vdots \\ \frac{\partial}{\partial A_{1m}} f & \cdots & \frac{\partial}{\partial A_{mn}} f \end{bmatrix}$$

- Trace:

$$\text{tr}A = \sum_{i=1}^n A_{ii} , \quad \text{tr}a = a , \quad \text{tr}ABC = \text{tr}CAB = \text{tr}BCA$$

- Some fact of matrix derivatives (without proof)

$$\nabla_A \text{tr}AB = B^T , \quad \nabla_A \text{tr}ABA^T C = CAB + C^T AB^T , \quad \nabla_A |A| = |A| (A^{-1})^T$$

Comments on the normal equation

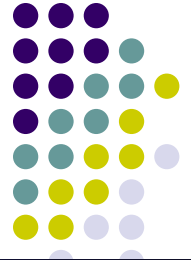


- In most situations of practical interest, the number of data points N is larger than the dimensionality k of the input space and the matrix \mathbf{X} is of full column rank. If this condition holds, then it is easy to verify that $X^T X$ is necessarily invertible.
- The assumption that $X^T X$ is invertible implies that it is positive definite, thus at the critical point we have found is a minimum.
- What if \mathbf{X} has less than full column rank? \rightarrow regularization (later).



Direct and Iterative methods

- Direct methods: we can achieve the solution in a single step by solving the normal equation
 - Using Gaussian elimination or QR decomposition, we converge in a finite number of steps
 - It can be infeasible when data are streaming in in real time, or of very large amount
- Iterative methods: stochastic or steepest gradient
 - Converging in a limiting sense
 - But more attractive in large practical problems
 - Caution is needed for deciding the learning rate α



Convergence rate

- **Theorem:** the steepest descent equation algorithm converge to the minimum of the cost characterized by normal equation:

$$\theta^{(\infty)} = (X^T X)^{-1} X^T y$$

If

$$0 < \alpha < 2/\lambda_{\max}[X^T X]$$

- A formal analysis of LMS need more math-mussels; in practice, one can use a small α , or gradually decrease α .



A Summary:

- LMS update rule

$$\theta_j^{t+1} = \theta_j^t + \alpha (y_n - \mathbf{x}_n^T \boldsymbol{\theta}^t) x_{n,i}$$

- Pros: on-line, low per-step cost, fast convergence and perhaps less prone to local optimum
- Cons: convergence to optimum not always guaranteed

- Steepest descent

$$\boldsymbol{\theta}^{t+1} = \boldsymbol{\theta}^t + \alpha \sum_{i=1}^n (y_n - \mathbf{x}_n^T \boldsymbol{\theta}^t) \mathbf{x}_n$$

- Pros: easy to implement, conceptually clean, guaranteed convergence
- Cons: batch, often slow converging

- Normal equations

$$\boldsymbol{\theta}^* = \left(X^T X \right)^{-1} X^T \bar{\mathbf{y}}$$

- Pros: a single-shot algorithm! Easiest to implement.
- Cons: need to compute pseudo-inverse $(X^T X)^{-1}$, expensive, numerical issues (e.g., matrix is singular ..), although there are ways to get around this ...



Geometric Interpretation of LMS

- The predictions on the training data are:

$$\hat{\bar{y}} = X\theta^* = X(X^T X)^{-1} X^T \bar{y}$$

- Note that

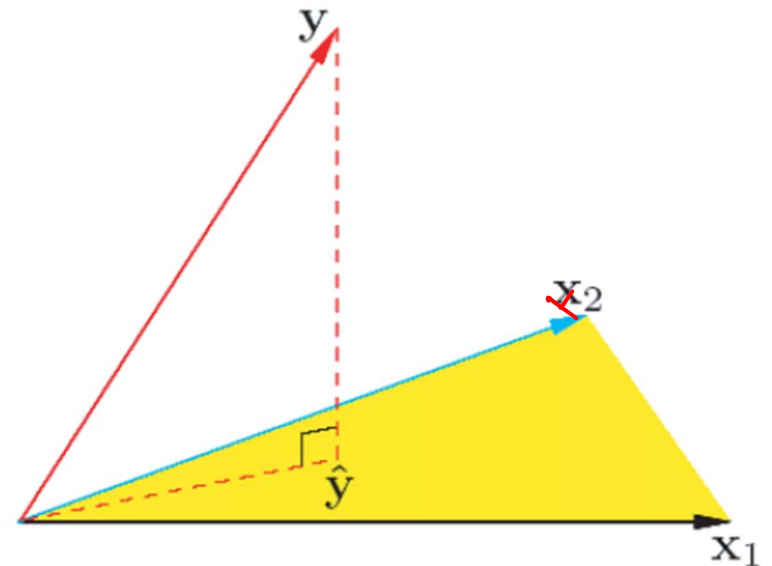
$$\hat{\bar{y}} - \bar{y} = (X(X^T X)^{-1} X^T - I)\bar{y}$$

and

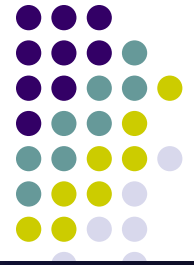
$$\begin{aligned} X^T(\hat{\bar{y}} - \bar{y}) &= X^T(X(X^T X)^{-1} X^T - I)\bar{y} \\ &= (X^T X(X^T X)^{-1} X^T - X^T)\bar{y} \\ &= 0 \quad !! \end{aligned}$$

$\hat{\bar{y}}$ is the orthogonal projection of \bar{y} into the space spanned by the column of X

$$\bar{y} = \begin{bmatrix} y_1 \\ y_2 \\ \vdots \\ y_n \end{bmatrix} \quad \mathbf{X} = \begin{bmatrix} \text{---} & \mathbf{x}_1 & \text{---} \\ \text{---} & \mathbf{x}_2 & \text{---} \\ \vdots & \vdots & \vdots \\ \text{---} & \mathbf{x}_n & \text{---} \end{bmatrix}$$



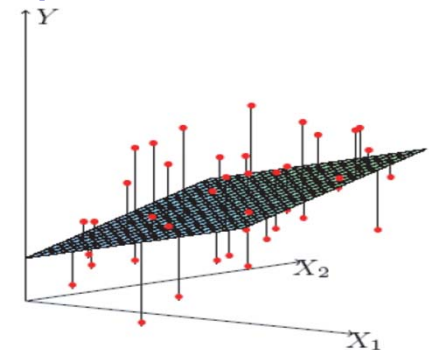
Probabilistic Interpretation of LMS



- Let us assume that the target variable and the inputs are related by the equation:

$$y_i = \theta^T \mathbf{x}_i + \varepsilon_i$$

where ε is an error term of unmodeled effects or random noise



- Now assume that ε follows a Gaussian $N(0, \sigma)$, then we have:

$$p(y_i | x_i; \theta) = \frac{1}{\sqrt{2\pi\sigma}} \exp\left(-\frac{(y_i - \theta^T \mathbf{x}_i)^2}{2\sigma^2}\right)$$

- By independence assumption:

$$L(\theta) = \prod_{i=1}^n p(y_i | x_i; \theta) = \left(\frac{1}{\sqrt{2\pi\sigma}}\right)^n \exp\left(-\frac{\sum_{i=1}^n (y_i - \theta^T \mathbf{x}_i)^2}{2\sigma^2}\right)$$

Probabilistic Interpretation of LMS, cont.



- Hence the log-likelihood is:

$$l(\theta) = n \log \frac{1}{\sqrt{2\pi}\sigma} - \frac{1}{\sigma^2} \frac{1}{2} \sum_{i=1}^n (y_i - \theta^T \mathbf{x}_i)^2$$

- Do you recognize the last term?

Yes it is:
$$J(\theta) = \frac{1}{2} \sum_{i=1}^n (\mathbf{x}_i^T \theta - y_i)^2$$

- Thus under independence assumption, LMS is equivalent to MLE of θ !

Case study: predicting gene expression



The genetic picture

causal SNPs

CGTTTCACTGTACAATTT

a univariate phenotype:

i.e., the expression intensity of
a gene



Association Mapping as Regression

	Phenotype (BMI)	Genotype
Individual 1	2.5	C T C T
		C A C T
Individual 2	4.8	G A G A
		C T C T
⋮		
Individual N	4.7	G T C T
		G T G T

Benign SNPs Causal SNP



Association Mapping as Regression

	Phenotype (BMI)	Genotype
Individual 1	2.5	.. 0 1 .. 0 0 . . .
Individual 2	4.8	.. 1 1 .. 1 1 . . .
⋮		
Individual N	4.7	.. 2 2 .. 1 0 . . .



y_i

=

$$\sum_{j=1}^J x_{ij} \beta_j$$

SNPs with large $|\beta_j|$ are relevant

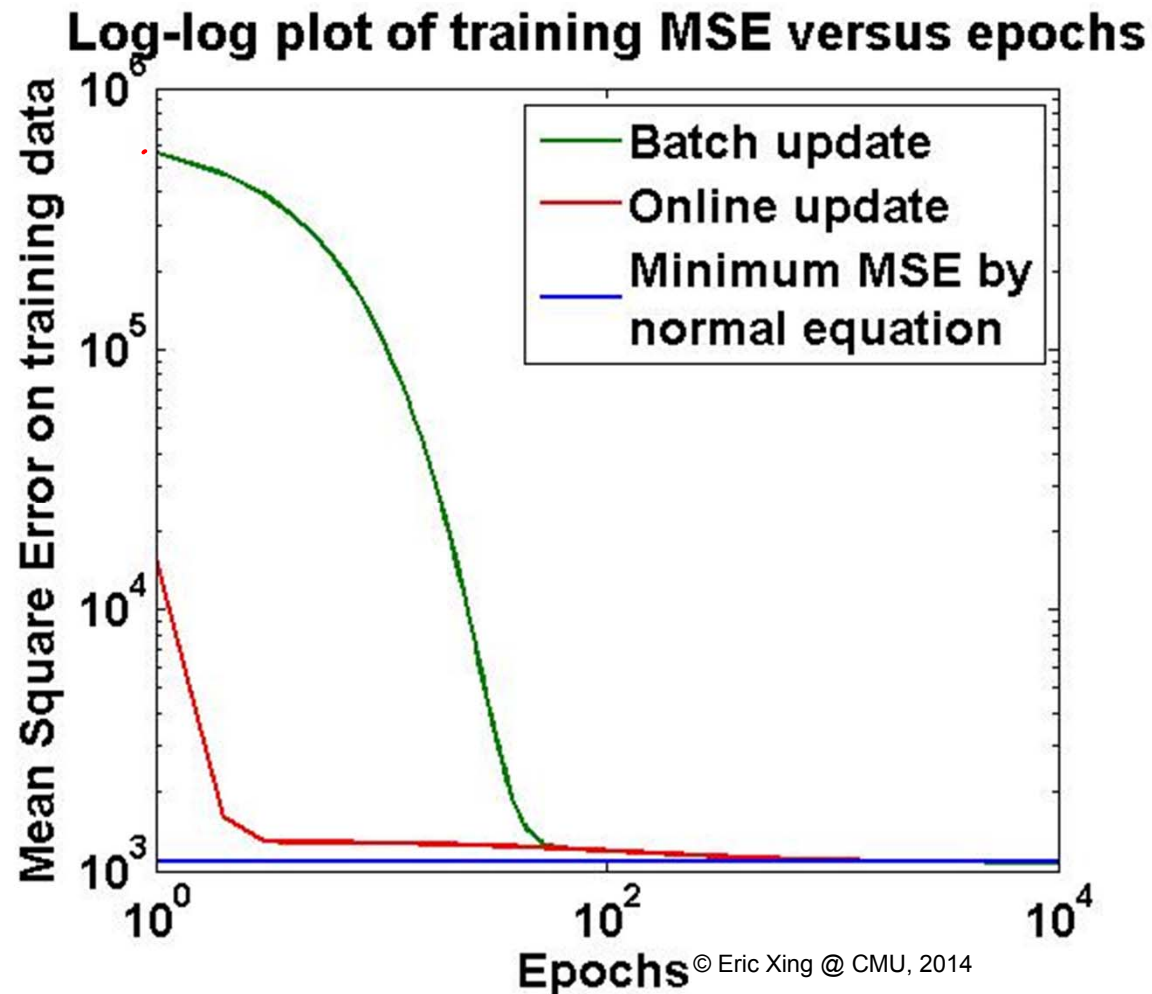
Experimental setup



- Asthama dataset
 - 543 individuals, genotyped at 34 SNPs
 - Diploid data was transformed into 0/1 (for homozygotes) or 2 (for heterozygotes)
 - $X=543 \times 34$ matrix
 - Y =Phenotype variable (continuous)
- A single phenotype was used for regression
- Implementation details
 - Iterative methods: Batch update and online update implemented.
 - For both methods, step size α is chosen to be a small fixed value (10^{-6}). This choice is based on the data used for experiments.
 - Both methods are only run to a maximum of 2000 epochs or until the change in training MSE is less than 10^{-4}

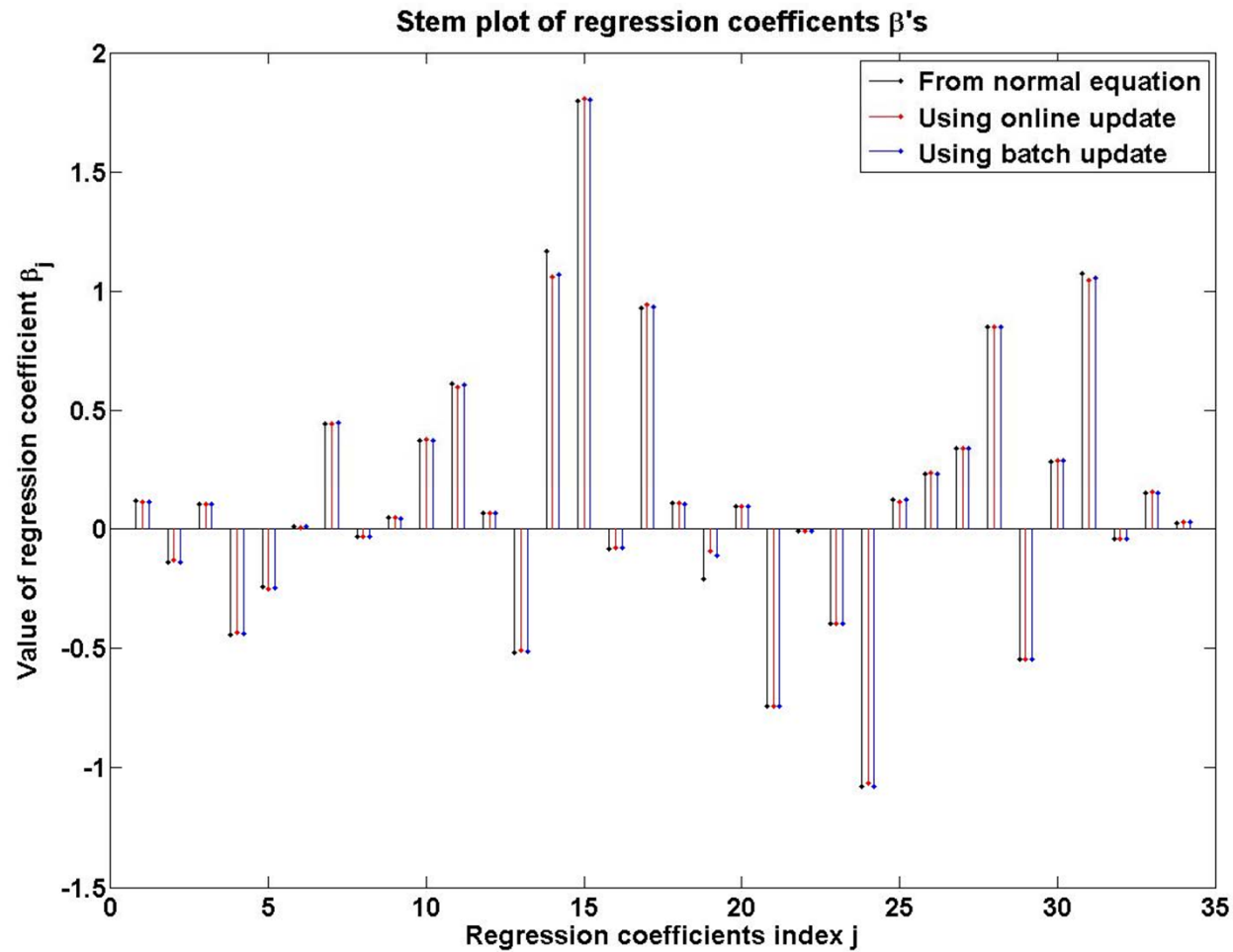


Convergence Curves



- For the batch method, the training MSE is initially large due to uninformed initialization
- In the online update, N updates for every epoch reduces MSE to a much smaller value.

The Learned Coefficients



Multivariate Regression for Trait Association Analysis



Trait

Genotype

Association Strength

2.1

=

T
G
A
A
C
C
A
T
G
A
A
G
T
A

x

?

y

=

X

x

β

Multivariate Regression for Trait Association Analysis



Trait

Genotype

Association Strength

2.1

=

TGAACCATGAAGTA

x



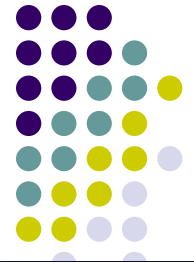
$$\beta^* = \arg \min_{\beta} (\mathbf{y} - \mathbf{X}\beta)^T (\mathbf{y} - \mathbf{X}\beta)$$

Many non-zero associations:
Which SNPs are truly significant?

Sparsity



- One common assumption to make **sparsity**.
- **Makes biological sense:** each phenotype is likely to be associated with a small number of SNPs, rather than all the SNPs.
- **Makes statistical sense:** Learning is now feasible in high dimensions with small sample size



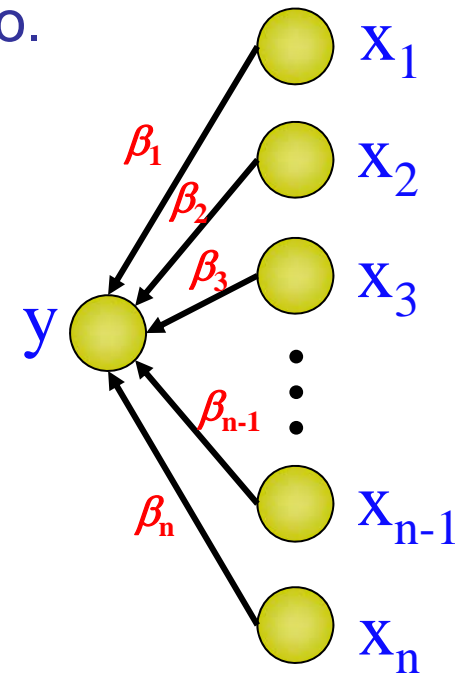
Sparsity: In a mathematical sense

- Consider least squares linear regression problem:
- Sparsity means most of the beta's are zero.

$$\hat{\beta} = \operatorname{argmin}_{\beta} \|\mathbf{Y} - \mathbf{X}\beta\|^2$$

subject to:

$$\sum_{j=1}^p \mathbb{I}[|\beta_j| > 0] \leq C$$



- But this is not convex!!! Many local optima, computationally intractable.

L1 Regularization (LASSO)

(Tibshirani, 1996)



- A convex relaxation.

Constrained Form

$$\hat{\beta} = \operatorname{argmin}_{\beta} \|\mathbf{Y} - \mathbf{X}\beta\|^2$$

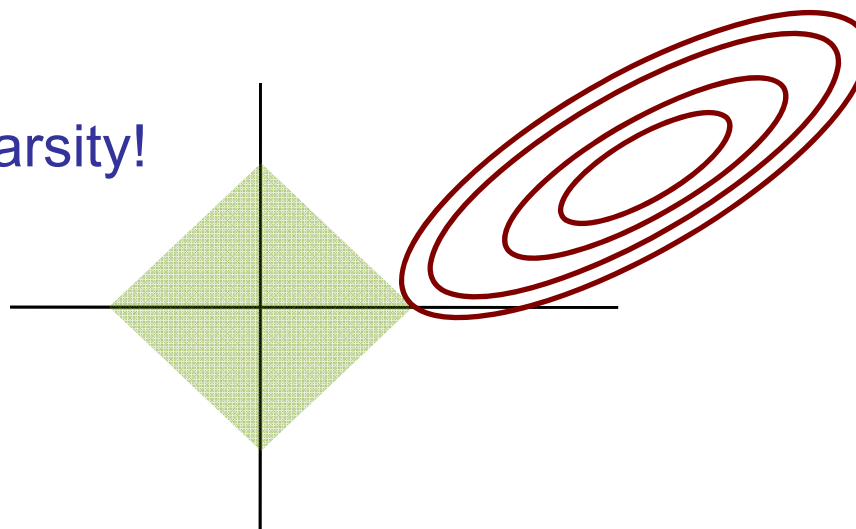
subject to:

$$\sum_{j=1}^p |\beta_j| \leq C$$

Lagrangian Form

$$\hat{\beta} = \operatorname{argmin}_{\beta} \|\mathbf{Y} - \mathbf{X}\beta\|^2 + \lambda \|\beta\|_1$$

- Still enforces sparsity!





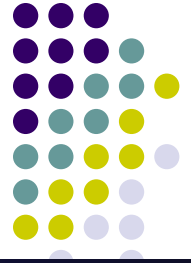
Theoretical Guarantees

- Assumptions
 - **Dependency Condition:** Relevant Covariates are not overly dependent
 - **Incoherence Condition:** Large number of irrelevant covariates cannot be too correlated with relevant covariates
 - **Strong concentration bounds:** Sample quantities converge to expected values quickly

If these assumptions are met, LASSO will asymptotically recover correct subset of covariates that are relevant.

Consistent Structure Recovery

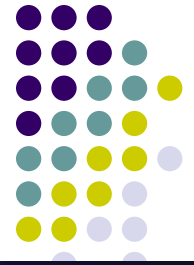
[Zhao and Yu 2006]



Theorem 4 (Gaussian Noise). Assume ε_i^n are i.i.d. Gaussian random variables. Under conditions (5), (6), (7) and (8), if there exists $0 \leq c_3 < c_2 - c_1$ for which $p_n = O(e^{n^{c_3}})$ then strong Irrepresentable Condition implies that Lasso has strong sign consistency. In particular, for $\lambda_n \propto n^{\frac{1+c_4}{2}}$ with $c_3 < c_4 < c_2 - c_1$,

$$P(\hat{\beta}^n(\lambda_n) =_s \beta^n) \geq 1 - o(e^{-n^{c_3}}) \rightarrow 1 \text{ as } n \rightarrow \infty.$$

Lasso for Reducing False Positives



Trait

Genotype

Association Strength

2.1

=

T
G
A
A
C
C
A
T
G
A
A
G
T
A

x



Lasso
Penalty
for sparsity

$$\beta^* = \arg \min_{\beta} (\mathbf{y} - \mathbf{X}\beta)^T (\mathbf{y} - \mathbf{X}\beta) + \lambda \sum_{j=1}^J |\beta_j|$$

Many zero associations (**sparse** results),
but what if there are multiple related traits?



Ridge Regression vs Lasso

$$\min_{\beta} (\mathbf{X}\beta - \mathbf{Y})^T (\mathbf{X}\beta - \mathbf{Y}) + \lambda \text{pen}(\beta) = \min_{\beta} J(\beta) + \lambda \text{pen}(\beta)$$

Ridge Regression:

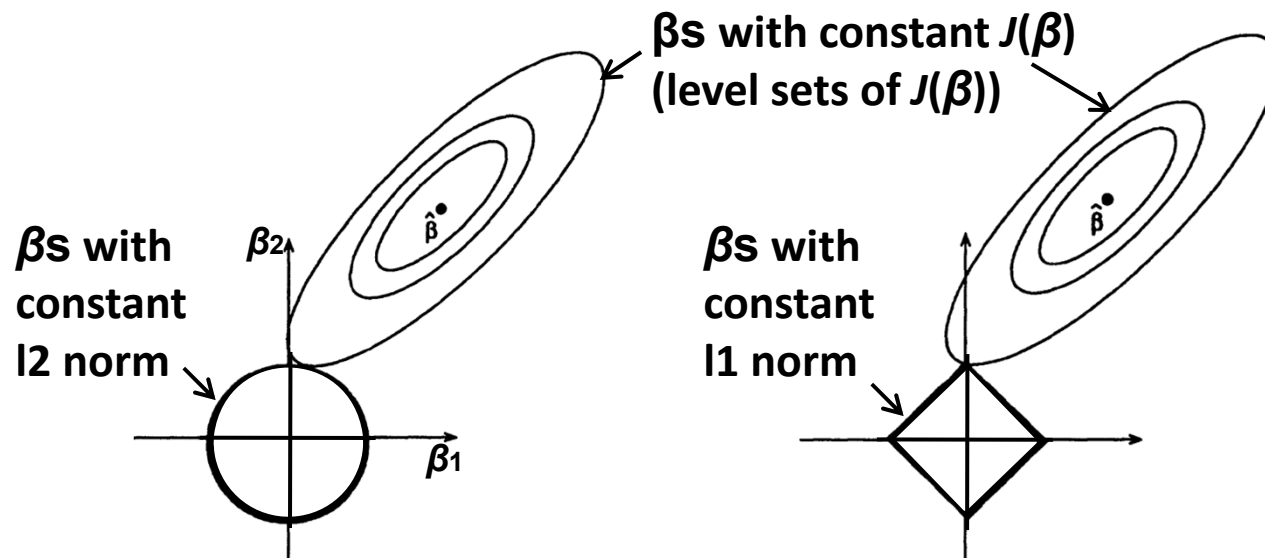
$$\text{pen}(\beta) = \|\beta\|_2^2$$

Lasso:

$$\text{pen}(\beta) = \|\beta\|_1$$

HOT

!



**Lasso (l_1 penalty) results in sparse solutions – vector with more zero coordinates
Good for high-dimensional problems – don't have to store all coordinates!**



Bayesian Interpretation

- Treat the distribution parameters θ also as a *random variable*
- The *a posteriori* distribution of θ after seeing the data is:

$$p(\theta | D) = \frac{p(D | \theta)p(\theta)}{p(D)} = \frac{p(D | \theta)p(\theta)}{\int p(D | \theta)p(\theta)d\theta}$$

This is Bayes Rule

$$\text{posterior} = \frac{\text{likelihood} \times \text{prior}}{\text{marginal likelihood}}$$

Bayes, Thomas (1763) An essay towards solving a problem in the doctrine of chances. *Philosophical Transactions of the Royal Society of London*, 53:370-418



The prior $p(\cdot)$ encodes our prior knowledge about the domain

Regularized Least Squares and MAP



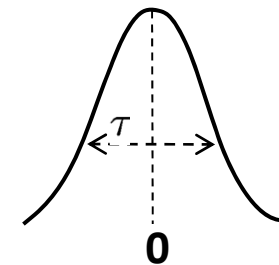
What if $(X^T X)$ is not invertible ?

$$\hat{\beta}_{\text{MAP}} = \arg \max_{\beta} \underbrace{\log p(\{(X_i, Y_i)\}_{i=1}^n | \beta, \sigma^2)}_{\text{log likelihood}} + \underbrace{\log p(\beta)}_{\text{log prior}}$$

I) Gaussian Prior

$$\beta \sim \mathcal{N}(0, \tau^2 \mathbf{I})$$

$$p(\beta) \propto e^{-\beta^T \beta / 2\tau^2}$$



$$\hat{\beta}_{\text{MAP}} = \arg \min_{\beta} \sum_{i=1}^n (Y_i - X_i \beta)^2 + \lambda \|\beta\|_2^2$$

Ridge Regression

Closed form: HW

constant(σ^2, τ^2)

Prior belief that β is Gaussian with zero-mean biases solution to “small” β

Regularized Least Squares and MAP



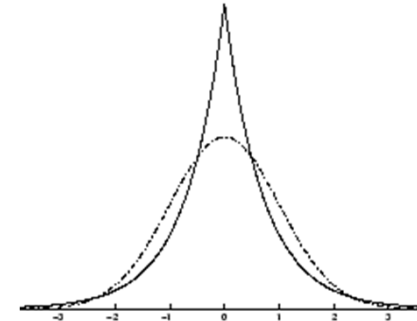
What if $(X^T X)$ is not invertible ?

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II) Laplace Prior

$$\beta_i \stackrel{iid}{\sim} \text{Laplace}(0, t)$$

$$p(\beta_i) \propto e^{-|\beta_i|/t}$$



$$\hat{\beta}_{\text{MAP}} = \arg \min_{\beta} \sum_{i=1}^n (Y_i - X_i \beta)^2 + \lambda \|\beta\|_1$$

Lasso

↓
constant(σ^2, t)

Closed form: HW

Prior belief that β is Laplace with zero-mean biases solution to “small” β



Take home message

- Gradient descent
 - On-line
 - Batch
- Normal equations
- Geometric interpretation of LMS
- Probabilistic interpretation of LMS, and equivalence of LMS and MLE under certain assumption (what?)
- Sparsity:
 - Approach: ridge vs. lasso regression
 - Interpretation: regularized regression versus Bayesian regression
 - Algorithm: convex optimization (we did not discuss this)
- LR does not mean fitting linear relations, but linear combination or basis functions (that can be non-linear)
- Weighting points by importance versus by fitness

After class material ...



Advanced Material: Beyond basic LR



- LR with non-linear basis functions
- Locally weighted linear regression
- Regression trees and Multilinear Interpolation

**We will discuss this in next class after we set the state right!
(if we've got time 😊)**

LR with non-linear basis functions



- LR does not mean we can only deal with linear relationships
- We are free to design (non-linear) features under LR

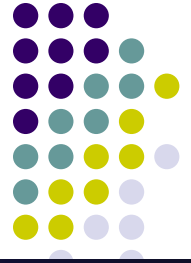
$$y = \theta_0 + \sum_{j=1}^m \theta_j \phi_j(x) = \theta^T \phi(x)$$

where the $\phi_j(x)$ are fixed basis functions (and we define $\phi_0(x) = 1$).

- Example: polynomial regression:

$$\phi(x) := [1, x, x^2, x^3]$$

- We will be concerned with estimating (distributions over) the weights θ and choosing the model order M .



Basis functions

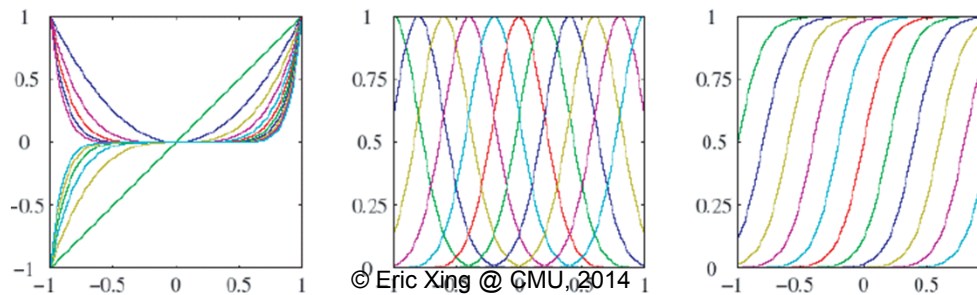
- There are many basis functions, e.g.:

- Polynomial $\phi_j(x) = x^{j-1}$

- Radial basis functions $\phi_j(x) = \exp\left(-\frac{(x - \mu_j)^2}{2s^2}\right)$

- Sigmoidal $\phi_j(x) = \sigma\left(\frac{x - \mu_j}{s}\right)$

- Splines, Fourier, Wavelets, etc



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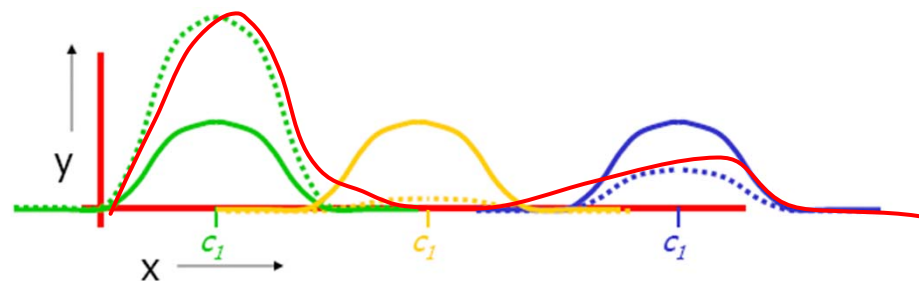
1D and 2D RBFs

- 1D RBF



$$y^{est} = \beta_1 \phi_1(x) + \beta_2 \phi_2(x) + \beta_3 \phi_3(x)$$

- After fit:



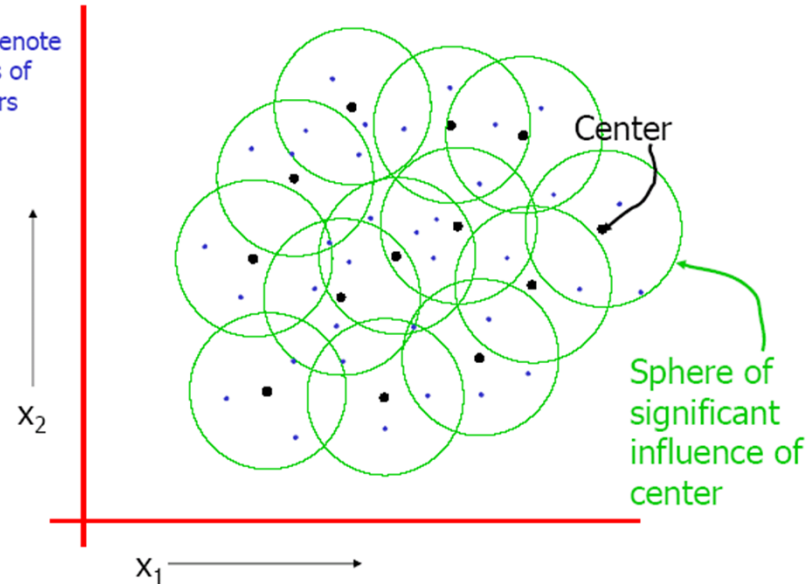
$$y^{est} = 2\phi_1(x) + 0.05\phi_2(x) + 0.5\phi_3(x)$$



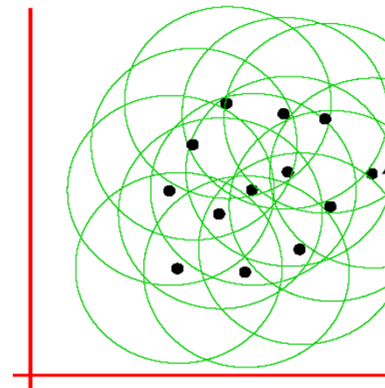
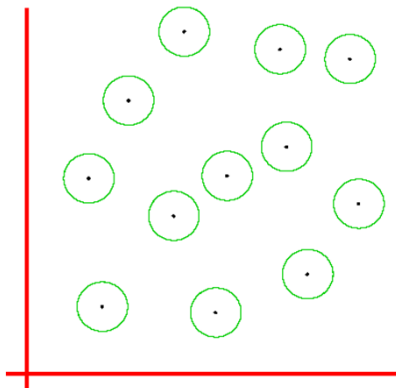
Good and Bad RBFs

- A good 2D RBF

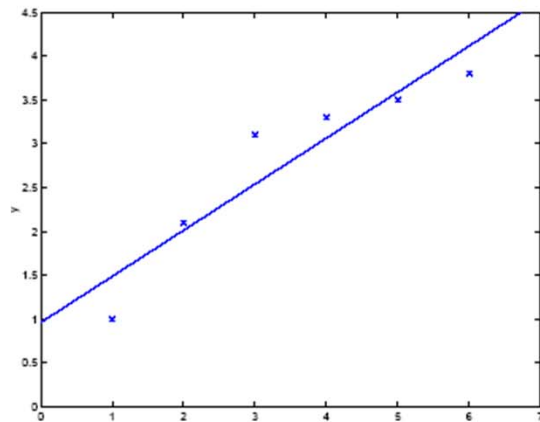
Blue dots denote coordinates of input vectors



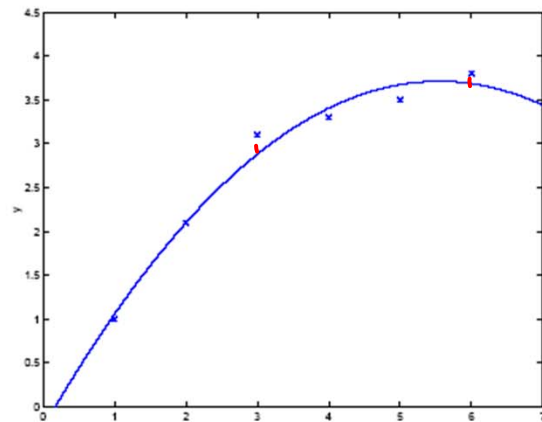
- Two bad 2D RBFs



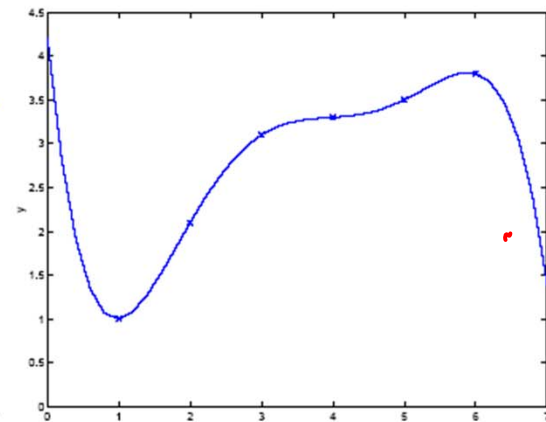
Overfitting and underfitting



$$y = \theta_0 + \theta_1 x$$



$$y = \theta_0 + \theta_1 x + \theta_2 x^2$$

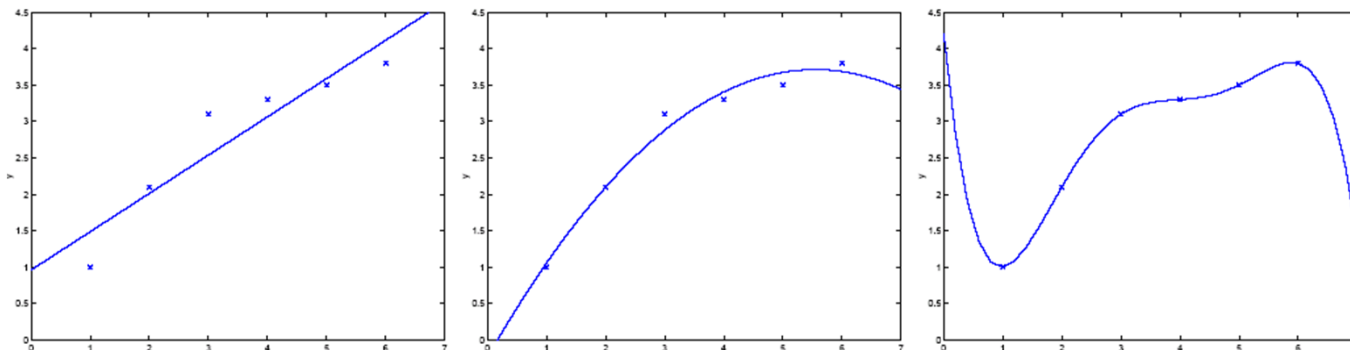


$$y = \sum_{j=0}^5 \theta_j x^j$$



Bias and variance

- We define the bias of a model to be the expected generalization error even if we were to fit it to a very (say, infinitely) large training set.
- By fitting "spurious" patterns in the training set, we might again obtain a model with large generalization error. In this case, we say the model has large variance.



Locally weighted linear regression



- The algorithm:

Instead of minimizing

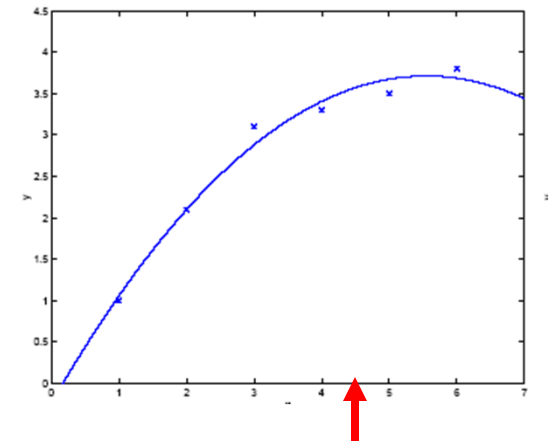
$$J(\theta) = \frac{1}{2} \sum_{i=1}^n (\mathbf{x}_i^T \theta - y_i)^2$$

now we fit θ to minimize

$$J(\theta) = \frac{1}{2} \sum_{i=1}^n w_i (\mathbf{x}_i^T \theta - y_i)^2$$

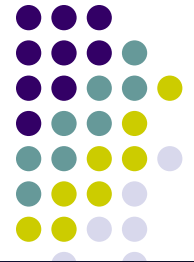
Where do w_i 's come from?

$$w_i = \exp\left(-\frac{(\mathbf{x}_i - \mathbf{x})^2}{2\tau^2}\right)$$



- where \mathbf{x} is the query point for which we'd like to know its corresponding y

→ Essentially we put higher weights on (errors on) training examples that are close to the query point (than those that are further away from the query)



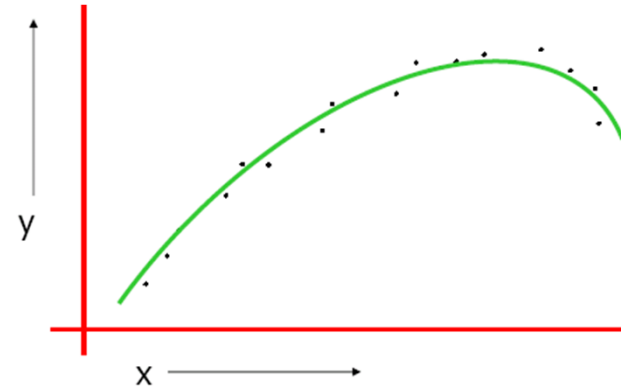
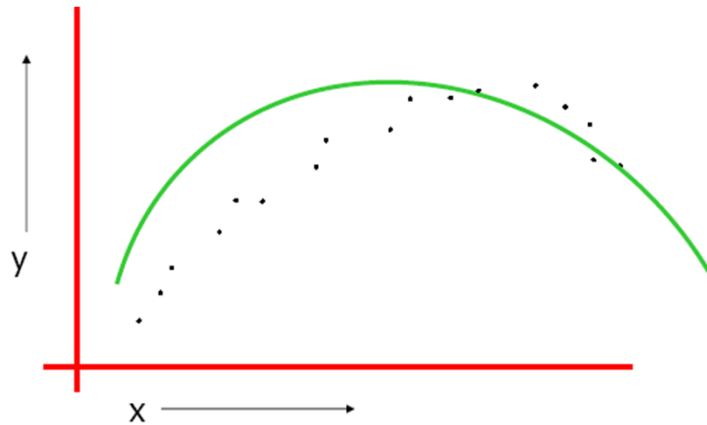
Parametric vs. non-parametric

- Locally weighted linear regression is the second example we are running into of a **non-parametric** algorithm. (what is the first?)
- The (unweighted) linear regression algorithm that we saw earlier is known as a **parametric** learning algorithm
 - because it has a fixed, finite number of parameters (the θ), which are fit to the data;
 - Once we've fit the θ and stored them away, we no longer need to keep the training data around to make future predictions.
 - In contrast, to make predictions using locally weighted linear regression, we need to keep the entire training set around.
- The term "**non-parametric**" (roughly) refers to the fact that the amount of stuff we need to keep in order to represent the hypothesis grows linearly with the size of the training set.

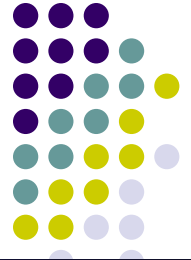


Robust Regression

- The best fit from a quadratic regression
- But this is probably better ...

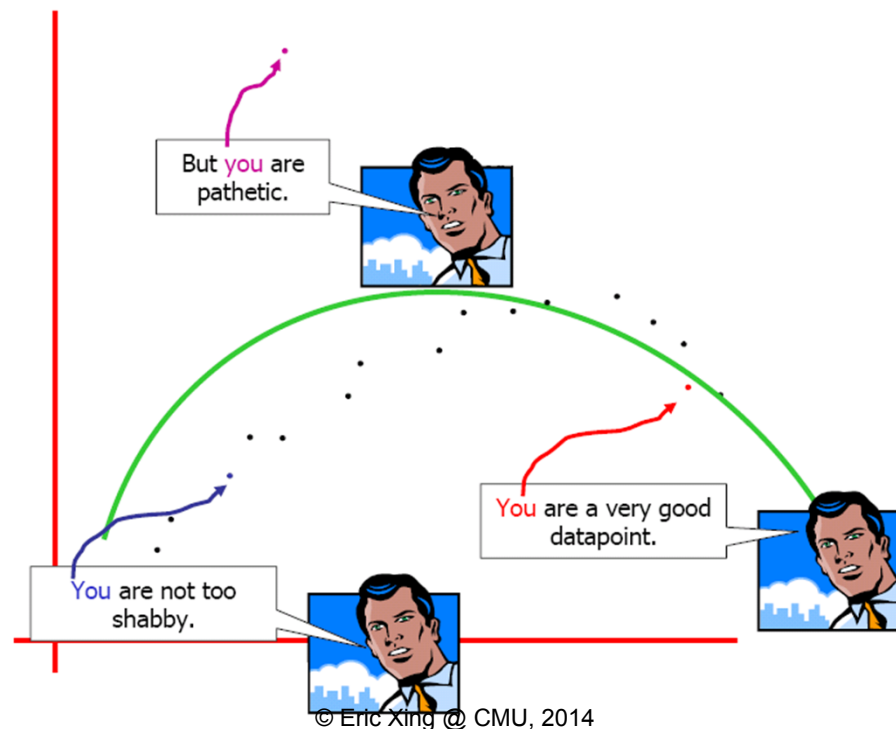


How can we do this?



LOESS-based Robust Regression

- Remember what we do in "locally weighted linear regression"?
→ we "score" each point for its impotence
- Now we score each point according to its "fitness"

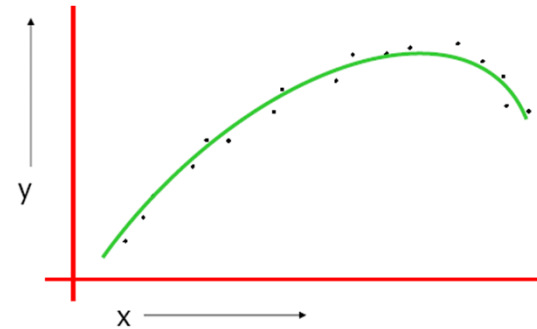


Robust regression



- For $k = 1$ to $R \dots$
 - Let (x_k, y_k) be the k th datapoint
 - Let y_k^{est} be predicted value of y_k
 - Let w_k be a weight for data point k that is large if the data point fits well and small if it fits badly:

$$w_k = \phi\left((y_k - y_k^{\text{est}})^2\right)$$



- Then redo the regression using weighted data points.
- Repeat whole thing until converged!

Robust regression—probabilistic interpretation



- What regular regression does:

Assume y_k was originally generated using the following recipe:

$$y_k = \theta^T \mathbf{x}_k + \mathcal{N}(0, \sigma^2)$$

Computational task is to find the Maximum Likelihood estimation of θ

Robust regression—probabilistic interpretation



- What LOESS robust regression does:

Assume y_k was originally generated using the following recipe:

with probability p : $y_k = \theta^T \mathbf{x}_k + \mathcal{N}(0, \sigma^2)$

but otherwise $y_k \sim \mathcal{N}(\mu, \sigma_{\text{huge}}^2)$

Computational task is to find the Maximum Likelihood estimates of θ , p , μ and σ_{huge} .

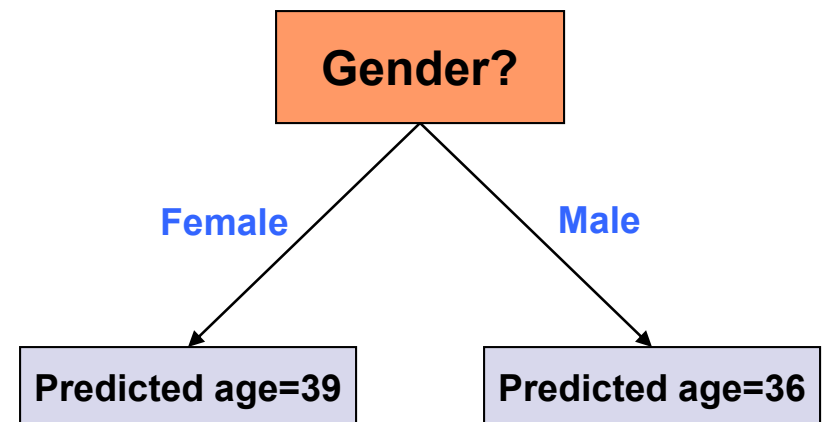
- The algorithm you saw with iterative **reweighting/refitting** does this computation for us. Later you will find that it is an instance of the famous **E.M.** algorithm

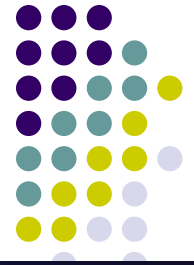


Regression Tree

- Decision tree for regression

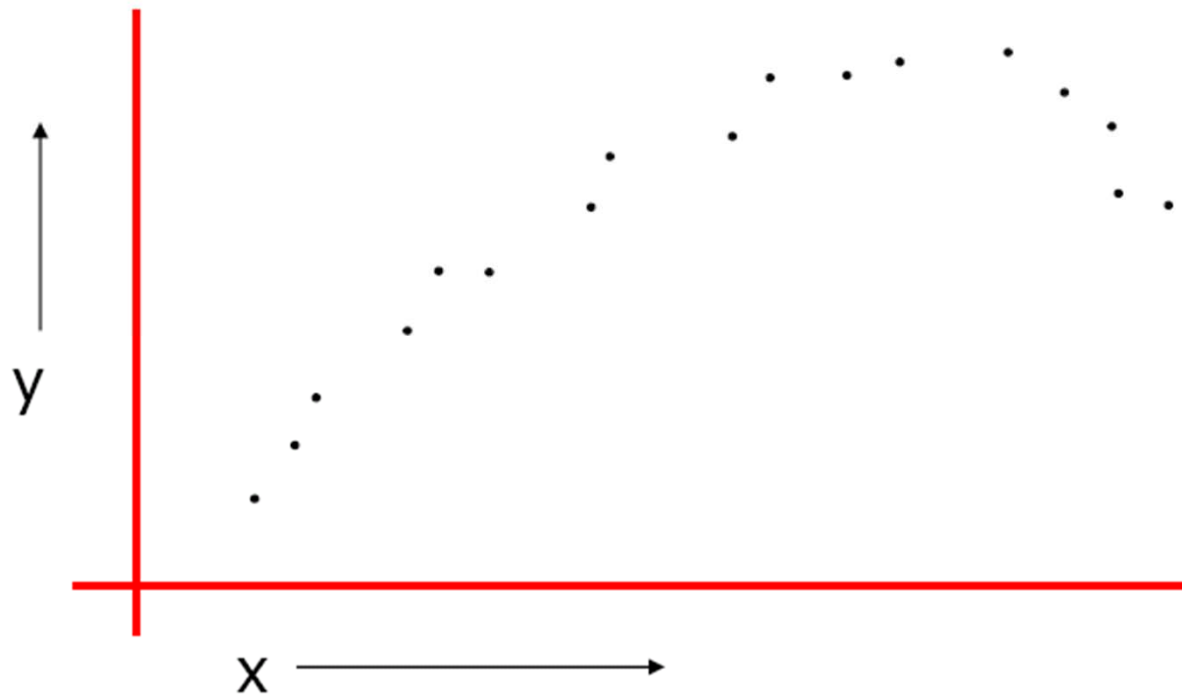
Gender	Rich?	Num. Children	# travel per yr.	Age
F	No	2	5	38
M	No	0	2	25
M	Yes	1	0	72
:	:	:	:	:

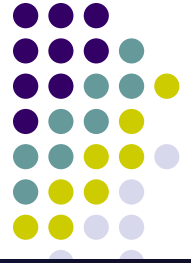




A conceptual picture

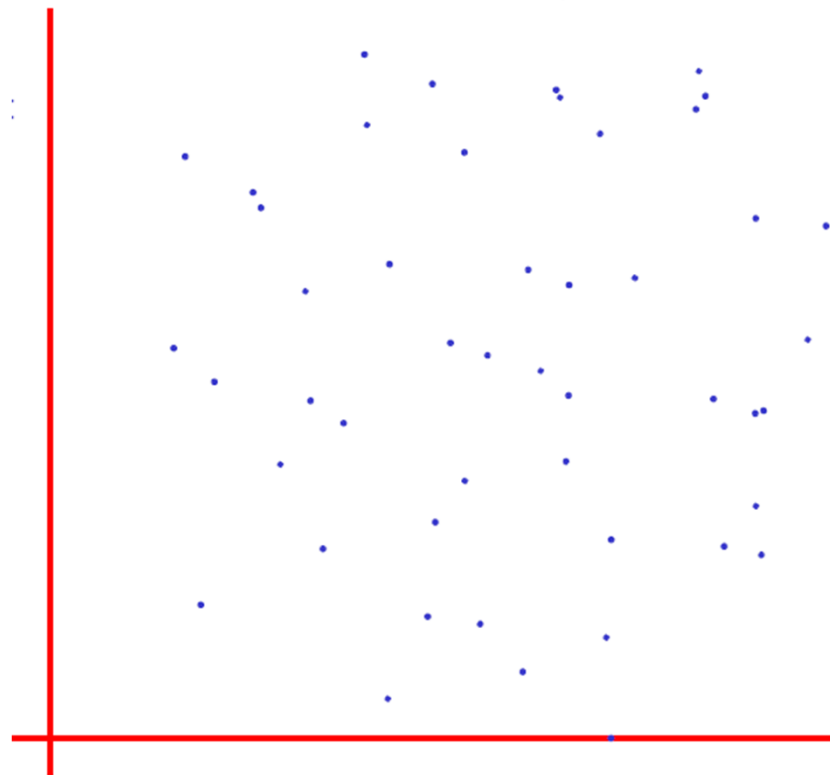
- Assuming regular regression trees, can you sketch a graph of the fitted function $y^*(x)$ over this diagram?





How about this one?

- Multilinear Interpolation



- We wanted to create a continuous and piecewise linear fit to the data

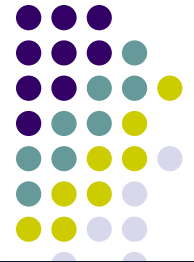


Take home message

- Gradient descent
 - On-line
 - Batch
- Normal equations
- Geometric interpretation of LMS
- Probabilistic interpretation of LMS, and equivalence of LMS and MLE under certain assumption (what?)
- Sparsity:
 - Approach: ridge vs. lasso regression
 - Interpretation: regularized regression versus Bayesian regression
 - Algorithm: convex optimization (we did not discuss this)
- LR does not mean fitting linear relations, but linear combination or basis functions (that can be non-linear)
- Weighting points by importance versus by fitness

Appendix





Parameter Learning from *iid* Data

- Goal: estimate distribution parameters θ from a dataset of N independent, identically distributed (*iid*), fully observed, training cases

$$D = \{x_1, \dots, x_N\}$$

- Maximum likelihood estimation (MLE)
 1. One of the most common estimators
 2. With iid and full-observability assumption, write $L(\theta)$ as the likelihood of the data:

$$\begin{aligned} L(\theta) &= P(x_1, x_2, \dots, x_N; \theta) \\ &= P(x_1; \theta) P(x_2; \theta), \dots, P(x_N; \theta) \\ &= \prod_{i=1}^N P(x_i; \theta) \end{aligned}$$

3. pick the setting of parameters most likely to have generated the data we saw:

$$\theta^* = \arg \max_{\theta} L(\theta) = \arg \max_{\theta} \log L(\theta)$$



Example: Bernoulli model

- Data:
 - We observed N *iid* coin tossing: $D=\{1, 0, 1, \dots, 0\}$
- Representation:



Binary r.v:

$$x_n = \{0,1\}$$

- Model:
$$P(x) = \begin{cases} 1-\theta & \text{for } x=0 \\ \theta & \text{for } x=1 \end{cases} \Rightarrow P(x) = \theta^x (1-\theta)^{1-x}$$

- How to write the likelihood of a single observation x_i ?

$$P(x_i) = \theta^{x_i} (1-\theta)^{1-x_i}$$

- The likelihood of dataset $D=\{x_1, \dots, x_N\}$:

$$P(x_1, x_2, \dots, x_N | \theta) = \prod_{i=1}^N P(x_i | \theta) = \prod_{i=1}^N (\theta^{x_i} (1-\theta)^{1-x_i}) = \theta^{\sum_{i=1}^N x_i} (1-\theta)^{\sum_{i=1}^N 1-x_i} = \theta^{\text{\#head}} (1-\theta)^{\text{\#tails}}$$



Maximum Likelihood Estimation

- Objective function:

$$\ell(\theta; D) = \log P(D | \theta) = \log \theta^{n_h} (1 - \theta)^{n_t} = n_h \log \theta + (N - n_h) \log(1 - \theta)$$

- We need to maximize this w.r.t. θ
- Take derivatives wrt θ

$$\frac{\partial \ell}{\partial \theta} = \frac{n_h}{\theta} - \frac{N - n_h}{1 - \theta} = 0 \quad \Rightarrow \quad \hat{\theta}_{MLE} = \frac{n_h}{N} \quad \text{or} \quad \hat{\theta}_{MLE} = \frac{1}{N} \sum_i x_i$$

Frequency as sample mean

- Sufficient statistics

- The counts, n_h , where $n_k = \sum_i x_i$, are **sufficient statistics** of data D



Overfitting

- Recall that for Bernoulli Distribution, we have

$$\hat{\theta}_{ML}^{\text{head}} = \frac{n^{\text{head}}}{n^{\text{head}} + n^{\text{tail}}}$$

- What if we tossed too few times so that we saw zero head?

We have $\hat{\theta}_{ML}^{\text{head}} = 0$, and we will predict that the probability of seeing a head next is zero!!!

- The rescue: *"smoothing"*

- Where n' is known as the pseudo- (imaginary) count

$$\hat{\theta}_{ML}^{\text{head}} = \frac{n^{\text{head}} + n'}{n^{\text{head}} + n^{\text{tail}} + n'}$$

- But can we make this more formal?



Bayesian Parameter Estimation

- Treat the distribution parameters θ also as a *random variable*
- The *a posteriori* distribution of θ after seeing the data is:

$$p(\theta | D) = \frac{p(D | \theta)p(\theta)}{p(D)} = \frac{p(D | \theta)p(\theta)}{\int p(D | \theta)p(\theta)d\theta}$$

This is Bayes Rule

$$\text{posterior} = \frac{\text{likelihood} \times \text{prior}}{\text{marginal likelihood}}$$

Bayes, Thomas (1763) An essay towards solving a problem in the doctrine of chances. *Philosophical Transactions of the Royal Society of London*, 53:370-418



The prior $p(\cdot)$ encodes our prior knowledge about the domain

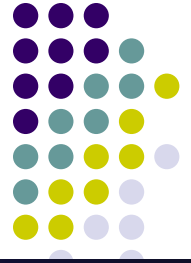
Frequentist Parameter Estimation



Two people with different priors $p(\theta)$ will end up with different estimates $p(\theta|D)$.

- Frequentists dislike this “subjectivity”.
- Frequentists think of the parameter as a **fixed, unknown constant**, not a random variable.
- Hence they have to come up with different “objective” **estimators** (ways of computing from data), instead of using Bayes’ rule.
 - These estimators have different properties, such as being “unbiased”, “minimum variance”, etc.
 - The **maximum likelihood estimator**, is one such estimator.

Discussion



θ or $p(\theta)$, this is the problem!

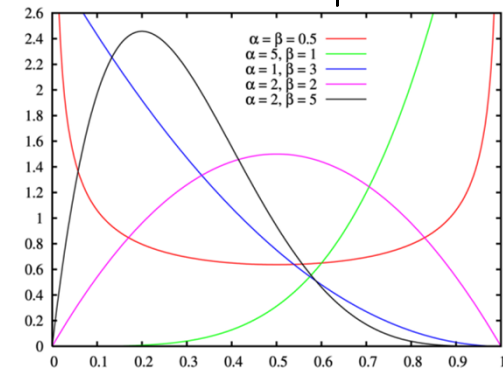
Bayesian estimation for Bernoulli



- Beta distribution:

$$P(\theta; \alpha, \beta) = \frac{\Gamma(\alpha + \beta)}{\Gamma(\alpha)\Gamma(\beta)} \theta^{\alpha-1} (1-\theta)^{\beta-1} = B(\alpha, \beta) \theta^{\alpha-1} (1-\theta)^{\beta-1}$$

- When x is discrete $\Gamma(x+1) = x\Gamma(x) = x!$

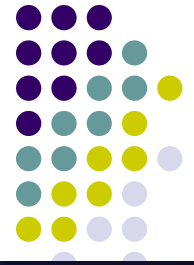


- Posterior distribution of θ :

$$P(\theta | x_1, \dots, x_N) = \frac{p(x_1, \dots, x_N | \theta) p(\theta)}{p(x_1, \dots, x_N)} \propto \theta^{n_h} (1-\theta)^{n_t} \times \theta^{\alpha-1} (1-\theta)^{\beta-1} = \theta^{n_h+\alpha-1} (1-\theta)^{n_t+\beta-1}$$

- Notice the isomorphism of the posterior to the prior,
- such a prior is called a **conjugate prior**
- α and β are hyperparameters (parameters of the prior) and correspond to the number of “virtual” heads/tails (pseudo counts)

Bayesian estimation for Bernoulli, con'd



- Posterior distribution of θ :

$$P(\theta | x_1, \dots, x_N) = \frac{p(x_1, \dots, x_N | \theta) p(\theta)}{p(x_1, \dots, x_N)} \propto \theta^{n_h} (1 - \theta)^{n_t} \times \theta^{\alpha-1} (1 - \theta)^{\beta-1} = \theta^{n_h + \alpha - 1} (1 - \theta)^{n_t + \beta - 1}$$

- Maximum *a posteriori* (MAP) estimation:

$$\theta_{MAP} = \arg \max_{\theta} \log P(\theta | x_1, \dots, x_N)$$

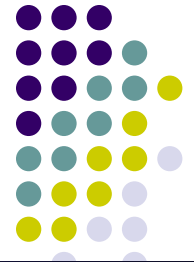
- Posterior mean estimation:

$$\theta_{Bayes} = \int \theta p(\theta | D) d\theta = C \int \theta \times \theta^{n_h + \alpha - 1} (1 - \theta)^{n_t + \beta - 1} d\theta = \frac{n_h + \alpha}{N + \alpha + \beta}$$

**Data parameters
can be understood
as pseudo-counts**

- Prior strength: $A = \alpha + \beta$

- A can be interpreted as the size of an imaginary data set from which we obtain the **pseudo-counts**



Effect of Prior Strength

- Suppose we have a uniform prior ($\alpha=\beta=1/2$), and we observe $\vec{n} = (n_h = 2, n_t = 8)$
- Weak prior $A = 2$. Posterior prediction:

$$p(x = h | n_h = 2, n_t = 8, \vec{\alpha} = \vec{\alpha}' \times 2) = \frac{1+2}{2+10} = 0.25$$

- Strong prior $A = 20$. Posterior prediction:

$$p(x = h | n_h = 2, n_t = 8, \vec{\alpha} = \vec{\alpha}' \times 20) = \frac{10+2}{20+10} = 0.40$$

- However, if we have enough data, it washes away the prior. e.g., $\vec{n} = (n_h = 200, n_t = 800)$. Then the estimates under weak and strong prior are $\frac{1+200}{2+1000}$ and $\frac{10+200}{20+1000}$, respectively, both of which are close to 0.2



Example 2: Gaussian density

- Data:
 - We observed N *iid* real samples:
 $D = \{-0.1, 10, 1, -5.2, \dots, 3\}$

- Model: $P(x) = (2\pi\sigma^2)^{-1/2} \exp\left\{-\frac{(x - \mu)^2}{2\sigma^2}\right\}$

- Log likelihood:

$$\ell(\theta; D) = \log P(D | \theta) = -\frac{N}{2} \log(2\pi\sigma^2) - \frac{1}{2} \sum_{n=1}^N \frac{(x_n - \mu)^2}{\sigma^2}$$

- MLE: take derivative and set to zero:

$$\frac{\partial \ell}{\partial \mu} = (1/\sigma^2) \sum_n (x_n - \mu)$$

$$\frac{\partial \ell}{\partial \sigma^2} = -\frac{N}{2\sigma^2} + \frac{1}{2\sigma^4} \sum_n (x_n - \mu)^2$$



$$\mu_{MLE} = \frac{1}{N} \sum_n (x_n)$$

$$\sigma_{MLE}^2 = \frac{1}{N} \sum_n (x_n - \mu_{ML})^2$$



MLE for a multivariate-Gaussian

- It can be shown that the MLE for μ and Σ is

$$\mu_{MLE} = \frac{1}{N} \sum_n (x_n)$$

$$\Sigma_{MLE} = \frac{1}{N} \sum_n (x_n - \mu_{ML})(x_n - \mu_{ML})^T = \frac{1}{N} S$$

where the scatter matrix is

$$S = \sum_n (x_n - \mu_{ML})(x_n - \mu_{ML})^T = \left(\sum_n x_n x_n^T \right) - N \mu_{ML} \mu_{ML}^T$$

- The sufficient statistics are $\sum_n x_n$ and $\sum_n x_n x_n^T$.
- Note that $X^T X = \sum_n x_n x_n^T$ may not be full rank (eg. if $N < D$), in which case Σ_{ML} is not invertible

$$x_n = \begin{pmatrix} x_n^1 \\ x_n^2 \\ \vdots \\ x_n^K \end{pmatrix}$$

$$X = \begin{pmatrix} \text{---} x_1^T \text{---} \\ \text{---} x_2^T \text{---} \\ \vdots \\ \text{---} x_N^T \text{---} \end{pmatrix}$$



Bayesian estimation

- Normal Prior:

$$P(\mu) = (2\pi\sigma_0^2)^{-1/2} \exp\left\{-\frac{(\mu - \mu_0)^2}{2\sigma_0^2}\right\}$$

- Joint probability:

$$P(x, \mu) = (2\pi\sigma^2)^{-N/2} \exp\left\{-\frac{1}{2\sigma^2} \sum_{n=1}^N (x_n - \mu)^2\right\} \\ \times (2\pi\sigma_0^2)^{-1/2} \exp\left\{-\frac{(\mu - \mu_0)^2}{2\sigma_0^2}\right\}$$

- Posterior:

$$P(\mu | \mathbf{x}) = (2\pi\tilde{\sigma}^2)^{-1/2} \exp\left\{-\frac{(\mu - \tilde{\mu})^2}{2\tilde{\sigma}^2}\right\}$$

where $\tilde{\mu} = \frac{N/\sigma^2}{N/\sigma^2 + 1/\sigma_0^2} \bar{x} + \frac{1/\sigma_0^2}{N/\sigma^2 + 1/\sigma_0^2} \mu_0$, and $\tilde{\sigma}^2 = \left(\frac{N}{\sigma^2} + \frac{1}{\sigma_0^2}\right)^{-1}$



Bayesian estimation: unknown μ , known σ

$$\mu_N = \frac{N/\sigma^2}{N/\sigma^2 + 1/\sigma_0^2} \bar{x} + \frac{1/\sigma_0^2}{N/\sigma^2 + 1/\sigma_0^2} \mu_0, \quad \tilde{\sigma}^2 = \left(\frac{N}{\sigma^2} + \frac{1}{\sigma_0^2} \right)^{-1}$$

- The posterior mean is a convex combination of the prior and the MLE, with weights proportional to the relative noise levels.
- The precision of the posterior $1/\sigma_N^2$ is the precision of the prior $1/\sigma_0^2$ plus one contribution of data precision $1/\sigma^2$ for each observed data point.

- Sequentially updating the mean

- $\mu_* = 0.8$ (unknown), $(\sigma^2)_* = 0.1$ (known)

- Effect of single data point

$$\mu_1 = \mu_0 + (x - \mu_0) \frac{\sigma_0^2}{\sigma^2 + \sigma_0^2} = x - (x - \mu_0) \frac{\sigma_0^2}{\sigma^2 + \sigma_0^2}$$

- Uninformative (vague/ flat) prior, $\sigma_0^2 \rightarrow \infty$

$$\mu_N \rightarrow \mu_0$$

