Machine Learning (COMP-652 and ECSE-608)

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Class web page: http://www.cs.mcgill.ca/~dprecup/courses/ml.html

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Outline

- Administrative issues
- What is machine learning?
- Types of machine learning
- Linear hypotheses
- Error functions
- Overfitting

Administrative issues

- Class materials:
 - No required textbook, but several textbooks available
 - Required or recommended readings (from books or research papers) posted on the class web page
 - Class notes: posted on the web page
- Prerequisites:
 - Knowledge of a programming language
 - Knowledge of probability/statistics, calculus and linear algebra; general facility with math
 - Some AI background is recommended *but not required*

Evaluation

- Four homework assignments (40%)
- Midterm examination (30%)
- Project (30%)
- Participation to class discussions (up to 1% extra credit)

What is learning?

- H. Simon: Any process by which a system improves its performance
- M. Minsky: Learning is making useful changes in our minds
- R. Michalsky: Learning is constructing or modifying representations of what is being experienced
- L. Valiant: Learning is the process of knowledge acquisition in the absence of explicit programming

Why study machine learning?

Engineering reasons:

- Easier to build a learning system than to hand-code a working program! E.g.:
 - Robot that learns a map of the environment by exploring
 - Programs that learn to play games by playing against themselves
- Improving on existing programs, e.g.
 - Instruction scheduling and register allocation in compilers
 - Combinatorial optimization problems
- Solving tasks that require a system to be adaptive, e.g.
 - Speech and handwriting recognition
 - "Intelligent" user interfaces

Why study machine learning?

Scientific reasons:

- Discover knowledge and patterns in highly dimensional, complex data
 - Sky surveys
 - High-energy physics data
 - Sequence analysis in bioinformatics
 - Social network analysis
 - Ecosystem analysis
- Understanding animal and human learning
 - How do we learn language?
 - How do we recognize faces?
- Creating real Al!

"If an expert system-brilliantly designed, engineered and implementedcannot learn not to repeat its mistakes, it is not as intelligent as a worm or a sea anemone or a kitten." (Oliver Selfridge).

Very brief history

- Studied ever since computers were invented (e.g. Samuel's checkers player)
- Very active in 1960s (neural networks)
- Died down in the 1970s
- Revival in early 1980s (decision trees, backpropagation, temporaldifference learning) - coined as "machine learning"
- Exploded since the 1990s
- Now: very active research field, several yearly conferences (e.g., ICML, NIPS), major journals (e.g., Machine Learning, Journal of Machine Learning Research), rapidly growing number of researchers
- The time is right to study in the field!
 - Lots of recent progress in algorithms and theory
 - Flood of data to be analyzed
 - Computational power is available
 - Growing demand for industrial applications

What are good machine learning tasks?

- There is no human expert
 - E.g., DNA analysis
- Humans can perform the task but cannot explain how E.g., character recognition
- Desired function changes frequently
 - E.g., predicting stock prices based on recent trading data
- Each user needs a customized function
 - E.g., news filtering

Important application areas

- Bioinformatics: sequence alignment, analyzing microarray data, information integration, ...
- Computer vision: object recognition, tracking, segmentation, active vision, ...
- Robotics: state estimation, map building, decision making
- Graphics: building realistic simulations
- Speech: recognition, speaker identification
- Financial analysis: option pricing, portfolio allocation
- E-commerce: automated trading agents, data mining, spam, ...
- Medicine: diagnosis, treatment, drug design,...
- Computer games: building adaptive opponents
- Multimedia: retrieval across diverse databases

Kinds of learning

Based on the information available:

- Supervised learning
- Reinforcement learning
- Unsupervised learning

Based on the role of the learner

- Passive learning
- Active learning

Passive and active learning

- Traditionally, learning algorithms have been *passive learners*, which take a given batch of data and process it to produce a hypothesis or model
 Data → Learner → Model
- Active learners are instead allowed to query the environment
 - Ask questions
 - Perform experiments
- Open issues: how to query the environment optimally? how to account for the cost of queries?

Example: A data set

Cell Nuclei of Fine Needle Aspirate



- Cell samples were taken from tumors in breast cancer patients before surgery, and imaged
- Tumors were excised
- Patients were followed to determine whether or not the cancer recurred, and how long until recurrence or disease free

Data (continued)

- Thirty real-valued variables per tumor.
- Two variables that can be predicted:
 - Outcome (R=recurrence, N=non-recurrence)
 - Time (until recurrence, for R, time healthy, for N).

tumor size	texture	perimeter	 outcome	time
18.02	27.6	117.5	N	31
17.99	10.38	122.8	N	61
20.29	14.34	135.1	R	27

. . .

Terminology							
tumor size	texture	perimeter		outcome	time		
18.02	27.6	117.5		N	31		
17.99	10.38	122.8		N	61		
20.29	14.34	135.1		R	27		

- Columns are called *input variables* or *features* or *attributes*
- The outcome and time (which we are trying to predict) are called *output* variables or targets
- A row in the table is called *training example* or *instance*
- The whole table is called *(training) data set*.
- The problem of predicting the recurrence is called *(binary)* classification
- The problem of predicting the time is called *regression*

tumor size	texture	perimeter	 outcome	time
18.02	27.6	117.5	Ν	31
17.99	10.38	122.8	Ν	61
20.29	14.34	135.1	R	27

More formally

- A training example *i* has the form: $\langle x_{i,1}, \ldots x_{i,n}, y_i \rangle$ where *n* is the number of attributes (30 in our case).
- We will use the notation \mathbf{x}_i to denote the column vector with elements $x_{i,1}, \ldots x_{i,n}$.
- The training set D consists of m training examples
- We denote the $m \times n$ matrix of attributes by X and the size-m column vector of outputs from the data set by y.

Supervised learning problem

- \bullet Let ${\mathcal X}$ denote the space of input values
- \bullet Let ${\mathcal Y}$ denote the space of output values
- Given a data set $D \subset \mathcal{X} \times \mathcal{Y}$, find a function:

 $h: \mathcal{X} \to \mathcal{Y}$

such that $h(\mathbf{x})$ is a "good predictor" for the value of y.

- *h* is called a *hypothesis*
- Problems are categorized by the type of output domain
 - If $\mathcal{Y} = \mathbb{R}$, this problem is called *regression*
 - If \mathcal{Y} is a categorical variable (i.e., part of a finite discrete set), the problem is called *classification*
 - If \mathcal{Y} is a more complex structure (eg graph) the problem is called *structured prediction*

Steps to solving a supervised learning problem

- 1. Decide what the input-output pairs are.
- 2. Decide how to encode inputs and outputs.

This defines the input space \mathcal{X} , and the output space \mathcal{Y} .

(We will discuss this in detail later)

3. Choose a class of hypotheses/representations $\ensuremath{\mathcal{H}}$.

4. ...

Example: What hypothesis class should we pick?

y

2.49

0.83

-0.25

3.10

0.87

0.02

-0.12

1.81

-0.83

0.43



Linear hypothesis

• Suppose y was a linear function of \mathbf{x} :

$$h_{\mathbf{w}}(\mathbf{x}) = w_0 + w_1 x_1 (+ \cdots)$$

- w_i are called *parameters* or *weights*
- To simplify notation, we can add an attribute $x_0 = 1$ to the other n attributes (also called *bias term* or *intercept term*):

$$h_{\mathbf{w}}(\mathbf{x}) = \sum_{i=0}^{n} w_i x_i = \mathbf{w}^T \mathbf{x}$$

where \mathbf{w} and \mathbf{x} are vectors of size n + 1.

How should we pick w?

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Error minimization!

- Intuitively, w should make the predictions of h_w close to the true values y on the data we have
- Hence, we will define an *error function* or *cost function* to measure how much our prediction differs from the "true" answer
- $\bullet\,$ We will pick ${\bf w}$ such that the error function is minimized

How should we choose the error function?

Least mean squares (LMS)

- Main idea: try to make $h_{\mathbf{w}}(\mathbf{x})$ close to y on the examples in the training set
- We define a *sum-of-squares* error function

$$J(\mathbf{w}) = \frac{1}{2} \sum_{i=1}^{m} (h_{\mathbf{w}}(\mathbf{x}_i) - y_i)^2$$

(the 1/2 is just for convenience)

• We will choose \mathbf{w} such as to minimize $J(\mathbf{w})$

Steps to solving a supervised learning problem

- 1. Decide what the input-output pairs are.
- 2. Decide how to encode inputs and outputs.

This defines the input space \mathcal{X} , and the output space \mathcal{Y} .

- 3. Choose a class of hypotheses/representations $\ensuremath{\mathcal{H}}$.
- 4. Choose an error function (cost function) to define the best hypothesis
- 5. Choose an algorithm for searching efficiently through the space of hypotheses.

Notation reminder

- Consider a function $f(u_1, u_2, \ldots, u_n) : \mathbb{R}^n \mapsto \mathbb{R}$ (for us, this will usually be an error function)
- The *partial derivative* w.r.t. u_i is denoted:

$$\frac{\partial}{\partial u_i} f(u_1, u_2, \dots, u_n) : \mathbb{R}^n \mapsto \mathbb{R}$$

The partial derivative is the derivative along the u_i axis, keeping all other variables fixed.

The gradient ∇f(u₁, u₂, ..., u_n) : ℝⁿ → ℝⁿ is a function which outputs a vector containing the partial derivatives.
That is:

$$\nabla f = \left\langle \frac{\partial}{\partial u_1} f, \frac{\partial}{\partial u_2} f, \dots, \frac{\partial}{\partial u_n} f \right\rangle$$

A bit of algebra

$$\begin{aligned} \frac{\partial}{\partial w_j} J(\mathbf{w}) &= \frac{\partial}{\partial w_j} \frac{1}{2} \sum_{i=1}^m (h_{\mathbf{w}}(\mathbf{x}_i) - y_i)^2 \\ &= \frac{1}{2} \cdot 2 \sum_{i=1}^m (h_{\mathbf{w}}(\mathbf{x}_i) - y_i) \frac{\partial}{\partial w_j} (h_{\mathbf{w}}(\mathbf{x}_i) - y_i) \\ &= \sum_{i=1}^m (h_{\mathbf{w}}(\mathbf{x}_i) - y_i) \frac{\partial}{\partial w_j} \left(\sum_{l=0}^n w_l x_{i,l} - y_i \right) \\ &= \sum_{i=1}^m (h_{\mathbf{w}}(\mathbf{x}_i) - y_i) x_{i,j} \end{aligned}$$

Setting all these partial derivatives to 0, we get a linear system with (n+1) equations and (n+1) unknowns.

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The solution

• Recalling some multivariate calculus:

$$\nabla_{\mathbf{w}} J = \nabla_{\mathbf{w}} \frac{1}{2} (\mathbf{X}\mathbf{w} - \mathbf{y})^T (\mathbf{X}\mathbf{w} - \mathbf{y})$$

=
$$\nabla_{\mathbf{w}} \frac{1}{2} (\mathbf{w}^T \mathbf{X}^T \mathbf{X} \mathbf{w} - \mathbf{y}^T \mathbf{X} \mathbf{w} - \mathbf{w}^T \mathbf{X}^T \mathbf{y} + \mathbf{y}^T \mathbf{y})$$

=
$$\mathbf{X}^T \mathbf{X} \mathbf{w} - \mathbf{X}^T \mathbf{y}$$

• Setting gradient equal to zero:

$$\mathbf{X}^{T}\mathbf{X}\mathbf{w} - \mathbf{X}^{T}\mathbf{y} = 0$$

$$\Rightarrow \mathbf{X}^{T}\mathbf{X}\mathbf{w} = \mathbf{X}^{T}\mathbf{y}$$

$$\Rightarrow \mathbf{w} = (\mathbf{X}^{T}\mathbf{X})^{-1}\mathbf{X}^{T}\mathbf{y}$$

• The inverse exists if the columns of ${f X}$ are linearly independent.

Example: Data and best linear hypothesis y = 1.60x + 1.05



Linear regression summary

- The optimal solution (minimizing sum-squared-error) can be computed in polynomial time in the size of the data set.
- The solution is $\mathbf{w} = (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{y}$, where \mathbf{X} is the data matrix augmented with a column of ones, and \mathbf{y} is the column vector of target outputs.
- A very rare case in which an analytical, exact solution is possible

Coming back to mean-squared error function...

- Good intuitive feel (small errors are ignored, large errors are penalized)
- Nice math (closed-form solution, unique global optimum)
- Geometric interpretation



• Any other interpretation?

A probabilistic assumption

- Assume y_i is a noisy target value, generated from a hypothesis $h_{\mathbf{w}}(\mathbf{x})$
- More specifically, assume that there exists \mathbf{w} such that:

$$y_i = h_{\mathbf{w}}(\mathbf{x}_i) + \epsilon_i$$

where ϵ_i is random variable (noise) drawn independently for each \mathbf{x}_i according to some Gaussian (normal) distribution with mean zero and variance σ .

 $\bullet\,$ How should we choose the parameter vector $\mathbf{w}?$

Bayes theorem in learning

Let h be a hypothesis and D be the set of training data. Using Bayes theorem, we have:

$$P(h|D) = \frac{P(D|h)P(h)}{P(D)},$$

where:

- P(h) is the prior probability of hypothesis h
- $P(D) = \int_h P(D|h)P(h)$ is the probability of training data D (normalization, independent of h)
- P(h|D) is the probability of h given D
- P(D|h) is the probability of D given h (*likelihood of the data*)

Choosing hypotheses

- What is the most probable hypothesis given the training data?
- Maximum a posteriori (MAP) hypothesis h_{MAP} :

$$h_{MAP} = \arg \max_{h \in \mathcal{H}} P(h|D)$$

= $\arg \max_{h \in \mathcal{H}} \frac{P(D|h)P(h)}{P(D)}$ (using Bayes theorem)
= $\arg \max_{h \in \mathcal{H}} P(D|h)P(h)$

Last step is because P(D) is independent of h (so constant for the maximization)

• This is the Bayesian answer (more in a minute)

Maximum likelihood estimation

$$h_{MAP} = \arg\max_{h \in \mathcal{H}} P(D|h)P(h)$$

• If we assume $P(h_i) = P(h_j)$ (all hypotheses are equally likely a priori) then we can further simplify, and choose the maximum likelihood (ML) hypothesis:

$$h_{ML} = \arg \max_{h \in \mathcal{H}} P(D|h) = \arg \max_{h \in \mathcal{H}} L(h)$$

- Standard assumption: the training examples are *independently identically distributed (i.i.d.)*
- This alows us to simplify P(D|h):

$$P(D|h) = \prod_{i=1}^{m} P(\langle \mathbf{x}_{i}, y_{i} \rangle | h) = \prod_{i=1}^{m} P(y_{i} | \mathbf{x}_{i}; h) P(\mathbf{x}_{i})$$

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The $\log\,trick$

• We want to maximize:

$$L(h) = \prod_{i=1}^{m} P(y_i | \mathbf{x}_i; h) P(\mathbf{x}_i)$$

This is a product, and products are hard to maximize!

• Instead, we will maximize $\log L(h)!$ (the log-likelihood function)

$$\log L(h) = \sum_{i=1}^{m} \log P(y_i | \mathbf{x}_i; h) + \sum_{i=1}^{m} \log P(\mathbf{x}_i)$$

• The second sum depends on D, but not on h, so it can be ignored in the search for a good hypothesis

Maximum likelihood for regression

• Adopt the assumption that:

$$y_i = h_{\mathbf{w}}(\mathbf{x}_i) + \epsilon_i,$$

where $\epsilon_i \sim \mathcal{N}(0, \sigma)$.

- The best hypothesis maximizes the likelihood of $y_i h_{\mathbf{w}}(\mathbf{x}_i) = \epsilon_i$
- Hence,

$$L(\mathbf{w}) = \prod_{i=1}^{m} \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{1}{2}\left(\frac{y_i - h_{\mathbf{w}}(\mathbf{x}_i)}{\sigma}\right)^2}$$

because the noise variables ϵ_i are from a Gaussian distribution

Applying the \log trick

$$\log L(\mathbf{w}) = \sum_{i=1}^{m} \log \left(\frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{1}{2} \frac{(y_i - h_{\mathbf{w}}(\mathbf{x}_i))^2}{\sigma^2}} \right)$$
$$= \sum_{i=1}^{m} \log \left(\frac{1}{\sqrt{2\pi\sigma^2}} \right) - \sum_{i=1}^{m} \frac{1}{2} \frac{(y_i - h_{\mathbf{w}}(\mathbf{x}_i))^2}{\sigma^2}$$

Maximizing the right hand side is the same as minimizing:

$$\sum_{i=1}^{m} \frac{1}{2} \frac{(y_i - h_{\mathbf{w}}(\mathbf{x}_i))^2}{\sigma^2}$$

This is our old friend, the sum-squared-error function! (the constants that are independent of h can again be ignored)

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Maximum likelihood hypothesis for least-squares estimators

• Under the assumption that the training examples are i.i.d. and that we have *Gaussian target noise*, the maximum likelihood parameters w are those minimizing the sum squared error:

$$\mathbf{w}^* = \arg\min_{\mathbf{w}} \sum_{i=1}^m \left(y_i - h_{\mathbf{w}}(\mathbf{x}_i) \right)^2$$

- This makes explicit the hypothesis behind minimizing the sum-squared error
- If the noise is not normally distributed, maximizing the likelihood will not be the same as minimizing the sum-squared error
- In practice, different loss functions are used depending on the noise assumption

A graphical representation for the data generation process



- Circles represent (random) variables)
- Arrows represent dependencies between variables
- Some variables are observed, others need to be inferred because they are hidden (latent)
- New assumptions can be incorporated by making the model more complicated

Predicting recurrence time based on tumor size



Is linear regression enough?

- Linear regression is too simple for most realistic problems But it should be the first thing you try for real-valued outputs!
- Problems can also occur is $\mathbf{X}^T \mathbf{X}$ is not invertible.
- Two possible solutions:
 - 1. Transform the data
 - Add cross-terms, higher-order terms
 - More generally, apply a transformation of the inputs from \mathcal{X} to some other space \mathcal{X}' , then do linear regression in the transformed space
 - 2. Use a different hypothesis class (e.g. non-linear functions)
- Today we focus on the first approach

Polynomial fits

- Suppose we want to fit a higher-degree polynomial to the data. (E.g., $y = w_2 x^2 + w_1 x^1 + w_0$.)
- Suppose for now that there is a single input variable per training sample.
- How do we do it?

Answer: Polynomial regression

- Given data: $(x_1, y_1), (x_2, y_2), \dots, (x_m, y_m)$.
- Suppose we want a degree-*d* polynomial fit.
- $\bullet~$ Let ${\bf y}~$ be as before and let

$$\mathbf{X} = \begin{bmatrix} x_1^d & \dots & x_1^2 & x_1 & 1 \\ x_2^d & \dots & x_2^2 & x_2 & 1 \\ \vdots & \vdots & \vdots & \vdots & \vdots \\ x_m^d & \dots & x_m^2 & x_m & 1 \end{bmatrix}$$

• Solve the linear regression $\mathbf{X}\mathbf{w}\approx\mathbf{y}$.

Example of quadratic regression: Data matrices

$$\mathbf{X} = \begin{bmatrix} 0.75 & 0.86 & 1 \\ 0.01 & 0.09 & 1 \\ 0.73 & -0.85 & 1 \\ 0.76 & 0.87 & 1 \\ 0.19 & -0.44 & 1 \\ 0.18 & -0.43 & 1 \\ 1.22 & -1.10 & 1 \\ 0.16 & 0.40 & 1 \\ 0.93 & -0.96 & 1 \\ 0.03 & 0.17 & 1 \end{bmatrix} \quad \mathbf{y} = \begin{bmatrix} 2.49 \\ 0.83 \\ -0.25 \\ 3.10 \\ 0.87 \\ 0.02 \\ -0.12 \\ 1.81 \\ -0.83 \\ 0.43 \end{bmatrix}$$

$\mathbf{X}^T \mathbf{X}$

 $\mathbf{X}^T \mathbf{X} =$

$\begin{bmatrix} 0.75 \ 0.01 \ 0.73 \ 0.76 \ 0.19 \ 0.18 \ 1.22 \ 0.16 \ 0.93 \ 0.03 \\ 0.86 \ 0.09 \ -0.85 \ 0.87 \ -0.44 \ -0.43 \ -1.10 \ 0.40 \ -0.96 \ 0.17 \\ 1 \ 1 \ 1 \ 1 \ 1 \ 1 \ 1 \ 1 \ 1 \ 1$	$\begin{array}{c} 0.75\\ 0.01\\ 0.73\\ 0.76\\ 0.19\\ 0.18\\ 1.22\\ 0.16\\ 0.93\\ 0.03\\ \end{array}$	$\begin{array}{c} 0.86\\ 0.09\\ -0.85\\ 0.87\\ -0.44\\ -0.43\\ -1.10\\ 0.40\\ -0.96\\ 0.17\end{array}$	1 1 1 1 1 1 1 1 1	
$= \begin{bmatrix} 4.11 & -1.64 & 4.95 \\ -1.64 & 4.95 & -1.39 \\ 4.95 & -1.39 & 10 \end{bmatrix}$				

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Solving for $\ensuremath{\mathbf{w}}$

$$\mathbf{w} = (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{X}^T \mathbf{y}$$

=
$$\begin{bmatrix} 4.11 & -1.64 & 4.95 \\ -1.64 & 4.95 & -1.39 \\ 4.95 & -1.39 & 10 \end{bmatrix}^{-1} \begin{bmatrix} 3.60 \\ 6.49 \\ 8.34 \end{bmatrix} = \begin{bmatrix} 0.68 \\ 1.74 \\ 0.73 \end{bmatrix}$$

So the best order-2 polynomial is $y = 0.68x^2 + 1.74x + 0.73$.

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Linear function approximation in general

• Given a set of examples $\langle \mathbf{x}_i, y_i \rangle_{i=1...m}$, we fit a hypothesis

$$h_{\mathbf{w}}(\mathbf{x}) = \sum_{k=0}^{K-1} w_k \phi_k(\mathbf{x}) = \mathbf{w}^T \phi(\mathbf{x})$$

where ϕ_k are called basis functions

• The best w is considered the one which minimizes the sum-squared error over the training data:

$$\sum_{i=1}^{m} (y_i - h_{\mathbf{w}}(\mathbf{x}_i))^2$$

• We can find the best w in closed form:

$$\mathbf{w} = (\mathbf{\Phi}^T \mathbf{\Phi})^{-1} \mathbf{\Phi}^T \mathbf{y}$$

or by other methods (e.g. gradient descent - as will be seen later)

Linear models in general

- By linear models, we mean that the hypothesis function $h_w(x)$ is a *linear function of the parameters* w
- This *does not mean the* $h_{\mathbf{w}}(\mathbf{x})$ *is a linear function of the input vector* \mathbf{x} (e.g., polynomial regression)
- In general

$$h_{\mathbf{w}}(\mathbf{x}) = \sum_{k=0}^{K-1} w_k \phi_k(\mathbf{x}) = \mathbf{w}^T \phi(\mathbf{x})$$

where ϕ_k are called *basis functions*

- Usually, we will assume that $\phi_0(\mathbf{x}) = 1, \forall \mathbf{x}$, to create a bias term
- The hypothesis can alternatively be written as:

$$h_{\mathbf{w}}(\mathbf{x}) = \mathbf{\Phi}\mathbf{w}$$

where Φ is a matrix with one row per instance; row j contains $\phi(\mathbf{x}_j)$.

• Basis functions are *fixed*

Example basis functions: Polynomials



"Global" functions: a small change in x may cause large change in the output of many basis functions

Example basis functions: Gaussians



- μ_k controls the position along the x-axis
- σ controls the width (activation radius)
- μ_k , σ fixed for now (later we discuss adjusting them)
- Usually thought as "local" functions: if σ is relatively small, a small change in x only causes a change in the output of a few basis functions (the ones with means close to x)

Example basis functions: Sigmoidal



- μ_k controls the position along the x-axis
- $\bullet \ s$ controls the slope
- μ_k , s fixed for now (later we discuss adjusting them)
- "Local" functions: a small change in x only causes a change in the output of a few basis (most others will stay close to 0 or 1)

Order-2 fit



Order-3 fit



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Order-4 fit



Order-5 fit



Is this a better fit to the data?

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Order-6 fit



Is this a better fit to the data?

Order-7 fit



Order-8 fit



Order-9 fit



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Overfitting

- A general, <u>HUGELY IMPORTANT</u> problem for all machine learning algorithms
- We can find a hypothesis that predicts perfectly the training data but *does not generalize* well to new data
- E.g., a lookup table!
- We are seeing an instance here: if we have a lot of parameters, the hypothesis "memorizes" the data points, but is wild everywhere else.
- Next time: defining overfitting formally, and finding ways to avoid it