## Feature Maps

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Slides based on Lecture 4d from David Rosenberg's course material.

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# The Input Space ${\mathfrak X}$

- ullet Our general learning theory setup: no assumptions about  ${\mathcal X}$
- But  $\mathfrak{X} = \mathsf{R}^d$  for the specific methods we've developed:
  - Ridge regression
  - Lasso regression
  - Support Vector Machines
- Our hypothesis space for these was all affine functions on  $R^d$ :

$$\mathcal{F} = \left\{ x \mapsto w^T x + b \mid w \in \mathbb{R}^d, b \in \mathbb{R} \right\}.$$

• What if we want to do prediction on inputs not natively in R<sup>d</sup>?

# The Input Space $\mathfrak X$

- Often want to use inputs not natively in R<sup>d</sup>:
  - Text documents
  - Image files
  - Sound recordings
  - DNA sequences

$$\phi(\kappa) = \begin{bmatrix} \phi_1(\kappa) \\ \vdots \\ \phi_d(\kappa) \end{bmatrix}$$

- But everything in a computer is a sequence of numbers
  - The *i*th entry of each sequence should have the same "meaning"
  - All the sequences should have the same length

#### Feature Extraction

#### Definition

Mapping an input from  $\mathfrak{X}$  to a vector in  $\mathbb{R}^d$  is called **feature extraction** or **featurization**.

## **Raw Input**

Feature Vector

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$$\mathcal{X} \xrightarrow{x} \overset{\text{Feature}}{\Longrightarrow} \frac{\phi(x)}{\text{Extraction}}$$

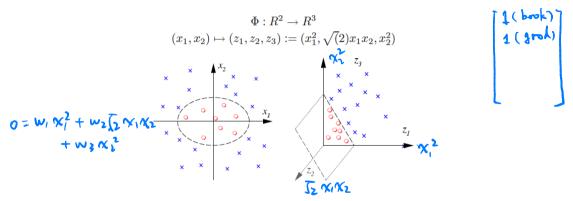
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# Linear Models with Explicit Feature Map

- Input space: X (no assumptions)
- Introduce feature map  $\phi: \mathcal{X} \to \mathbb{R}^d$
- The feature map maps into the feature space  $R^d$ .
- Hypothesis space of affine functions on feature space:

$$\mathcal{F} = \left\{ x \mapsto w^T \phi(x) + b \mid w \in \mathbb{R}^d, b \in \mathbb{R} \right\}.$$

# Geometric Example: Two class problem, nonlinear boundary



- With identity feature map  $\phi(x) = (x_1, x_2)$  and linear models, can't separate regions
- With appropriate featurization  $\phi(x) = (x_1, x_2, x_1^2 + x_2^2)$ , becomes linearly separable .
- Video: http://youtu.be/3liCbRZPrZA

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## Expressivity of Hypothesis Space

- For linear models, to grow the hypothesis spaces, we must add features.
- Sometimes we say a larger hypothesis is more expressive.
  - (can fit more relationships between input and action)
- Many ways to create new features.

Handling Nonlinearity with Linear Methods

## Example Task: Predicting Health

- General Philosophy: Extract every feature that might be relevant
- Features for medical diagnosis
  - height
  - weight
  - body temperature
  - blood pressure
  - etc...

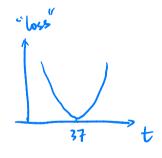
#### Feature Issues for Linear Predictors

- For linear predictors, it's important how features are added
  - The relation between a feature and the label may not be linear
  - There may be complex dependence among features
- Three types of nonlinearities can cause problems:
  - Non-monotonicity
  - Saturation
  - Interactions between features

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## Non-monotonicity: The Issue

- Feature Map:  $\phi(x) = [1, temperature(x)]$
- Action: Predict health score  $y \in R$  (positive is good)
- Hypothesis Space  $\mathcal{F}=\{affine functions of temperature\}$
- Issue:
  - Health is not an affine function of temperature.
  - Affine function can either say
    - Very high is bad and very low is good, or
    - Very low is bad and very high is good,
    - But here, both extremes are bad.



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From Percy Liang's "Lecture 3" slides from Stanford's CS221, Autumn 2014.

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## Non-monotonicity: Solution 1

Transform the input:

$$\phi(x) = \left[1, \{\text{temperature}(x) - 37\}^2\right],$$

where 37 is "normal" temperature in Celsius.

- Ok, but requires manually-specified domain knowledge
  - Do we really need that?
  - What does  $w^T \phi(x)$  look like?

$$w^{T}\phi(x) = w_{1} + w_{2}(t-3)^{2}$$

$$= w'_{1} + w_{2}t^{2} + w_{3}t$$

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### Non-monotonicity: Solution 2

• Think less, put in more:

$$\phi(x) = \left[1, \text{temperature}(x), \{\text{temperature}(x)\}^2\right].$$

More expressive than Solution 1.

#### General Rule

Features should be simple building blocks that can be pieced together.

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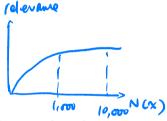
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### Saturation: The Issue

- Setting: Find products relevant to user's query
- Input: Product x
- Action: Score the relevance of x to user's query
- Feature Map:

$$\phi(x) = [1, N(x)],$$

where N(x) = number of people who bought x.



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• We expect a monotonic relationship between N(x) and relevance, but also expect diminishing return.

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### Saturation: Solve with nonlinear transform

• Smooth nonlinear transformation:

$$\phi(x) = [1, \log\{1 + N(x)\}]$$

- ullet log  $(\cdot)$  good for values with large dynamic ranges
- Discretization (a discontinuous transformation):

$$\phi(x) = (1(0 \leqslant N(x) < 10), 1(10 \leqslant N(x) < 100), \ldots)$$

Small buckets allow quite flexible relationship

From Percy Liang's "Lecture 3" slides from Stanford's CS221, Autumn 2014.

#### Interactions: The Issue

- Input: Patient information x
- Action: Health score  $y \in R$  (higher is better)
- Feature Map

$$\phi(x) = [\mathsf{height}(x), \mathsf{weight}(x)]$$

- Issue: It's the weight relative to the height that's important.
- Impossible to get with these features and a linear classifier.
- Need some interaction between height and weight.

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# Interactions: Approach 1

- Google "ideal weight from height"
- J. D. Robinson's "ideal weight" formula (for a male):

$$weight(kg) = 52 + 1.9 [height(in) - 60]$$

Make score square deviation between height(h) and ideal weight(w)

$$f(x) = (52 + 1.9 [h(x) - 60] - w(x))^{2}$$

$$f(x) = (52 + 1.9 [h(x) - 60] - w(x))^2$$
 • WolframAlpha for complicated Mathematics:

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$$f(x) = 3.61h(x)^2 - 3.8h(x)w(x) - 235.6h(x) + w(x)^2 + 124w(x) + 3844$$

From Percy Liang's "Lecture 3" slides from Stanford's CS221. Autumn 2014.

## Interactions: Approach 2

Just include all second order features:

$$\phi(x) = \left[1, h(x), w(x), h(x)^2, w(x)^2, \underbrace{h(x)w(x)}_{\text{cross term}}\right]$$

More flexible, no Google, no WolframAlpha.

#### General Principle

Simpler building blocks replace a single "smart" feature.

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Interaction terms are useful building blocks to model non-linearities in features.

- Suppose we start with  $x = (1, x_1, \dots, x_d) \in \mathbb{R}^{d+1} = \mathcal{X}$ .
- Consider adding all monomials of degree M:  $x_1^{p_1} \cdots x_d^{p_d}$ , with  $p_1 + \cdots + p_d = M$ .
  - Monomials with degree 2 in 2D space:  $x_1^2$ ,  $x_2^2$ ,  $x_1x_2$
- How many features will we end up with?  $\binom{M+d-1}{M}$  ("stars and bars")
- This leads to extremely large data matrices
  - For d = 40 and M = 8, we get 314457495 features.

# Big Feature Spaces

Very large feature spaces have two potential issues:

- Overfitting
- Memory and computational costs

#### Solutions:

- Overfitting we handle with regularization.
- Kernel methods can help with memory and computational costs when we go to high (or infinite) dimensional spaces.