
Learning, Part 2

Zero Error Linear Decision Boundary

Linear and Nonlinear

Linear Boundary
1

Linear Boundary 2

Nonlinear Boundary
1

Nonlinear Boundary
2

Neural Networks

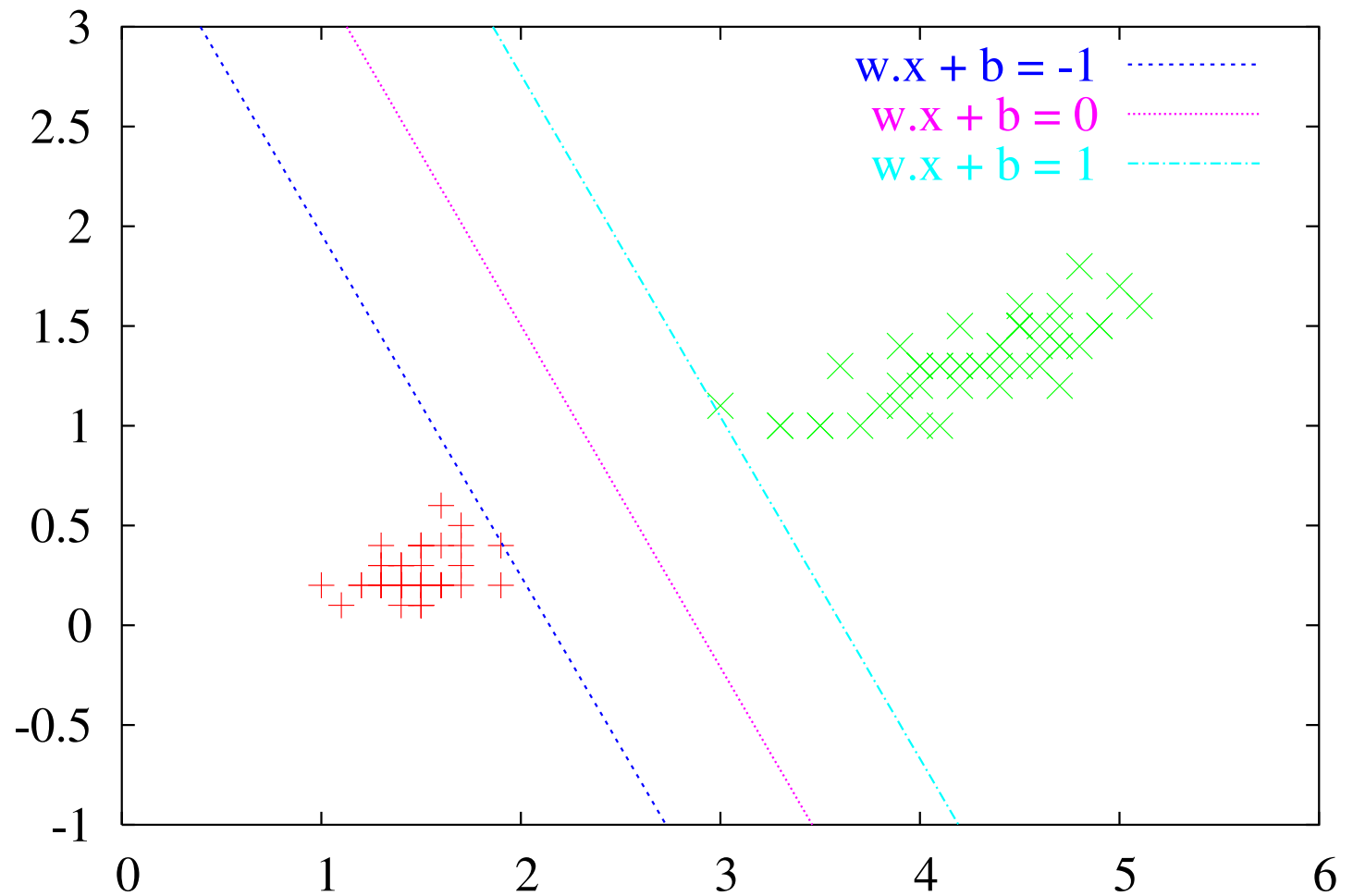
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Low Error Linear Decision Boundary

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Linear Boundary 2

Nonlinear Boundary 1

Nonlinear Boundary 2

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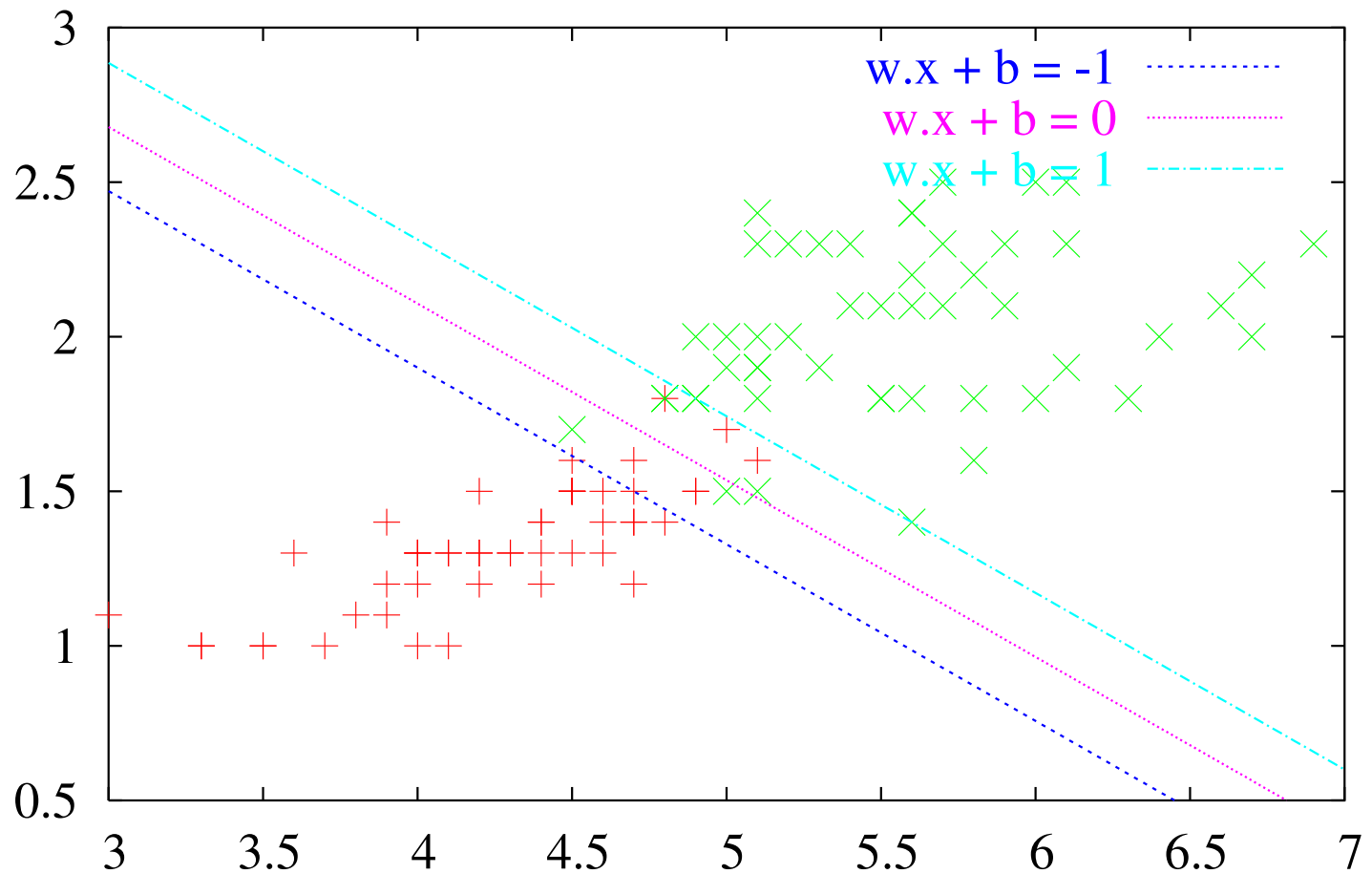
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Nonlinear Decision Boundary

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Linear Boundary 2

Nonlinear
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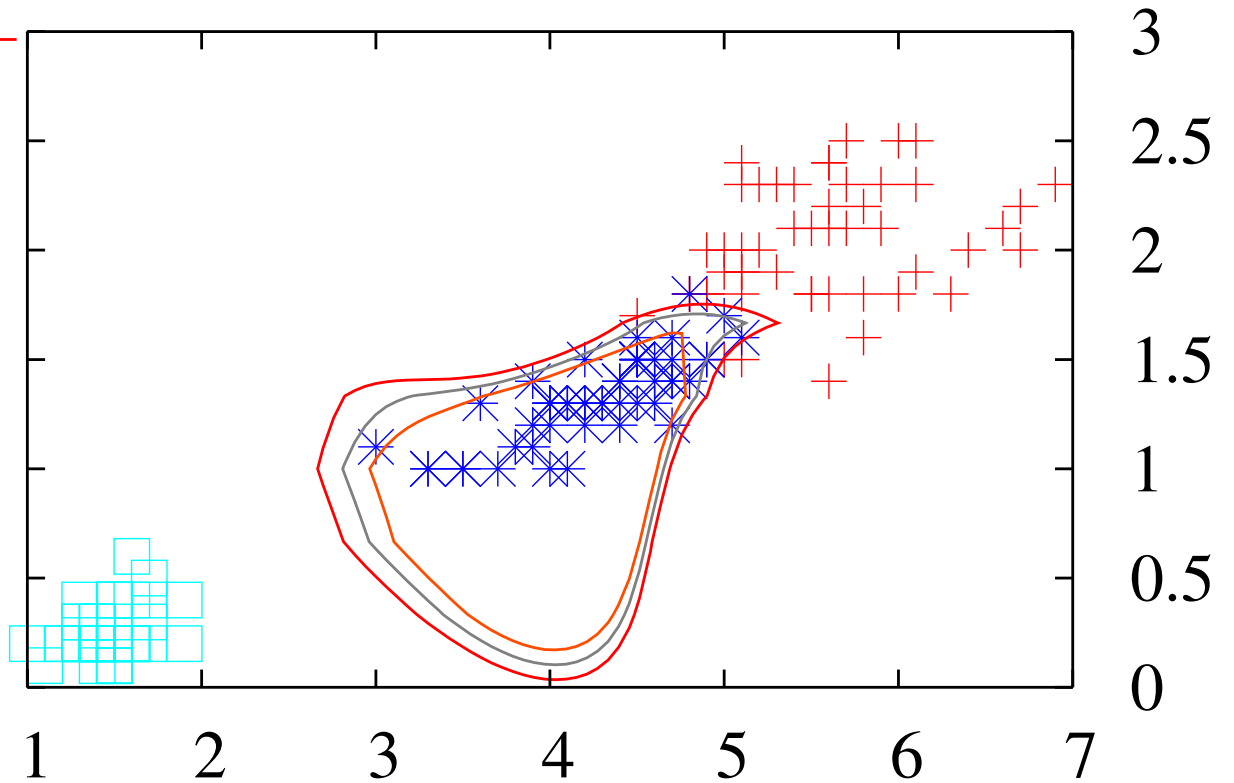
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1 ———
0 ———
-1 ———



Zoom on Nonlinear Decision Boundary

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Linear Boundary 2

Nonlinear Boundary

1

▷ Nonlinear

Boundary 2

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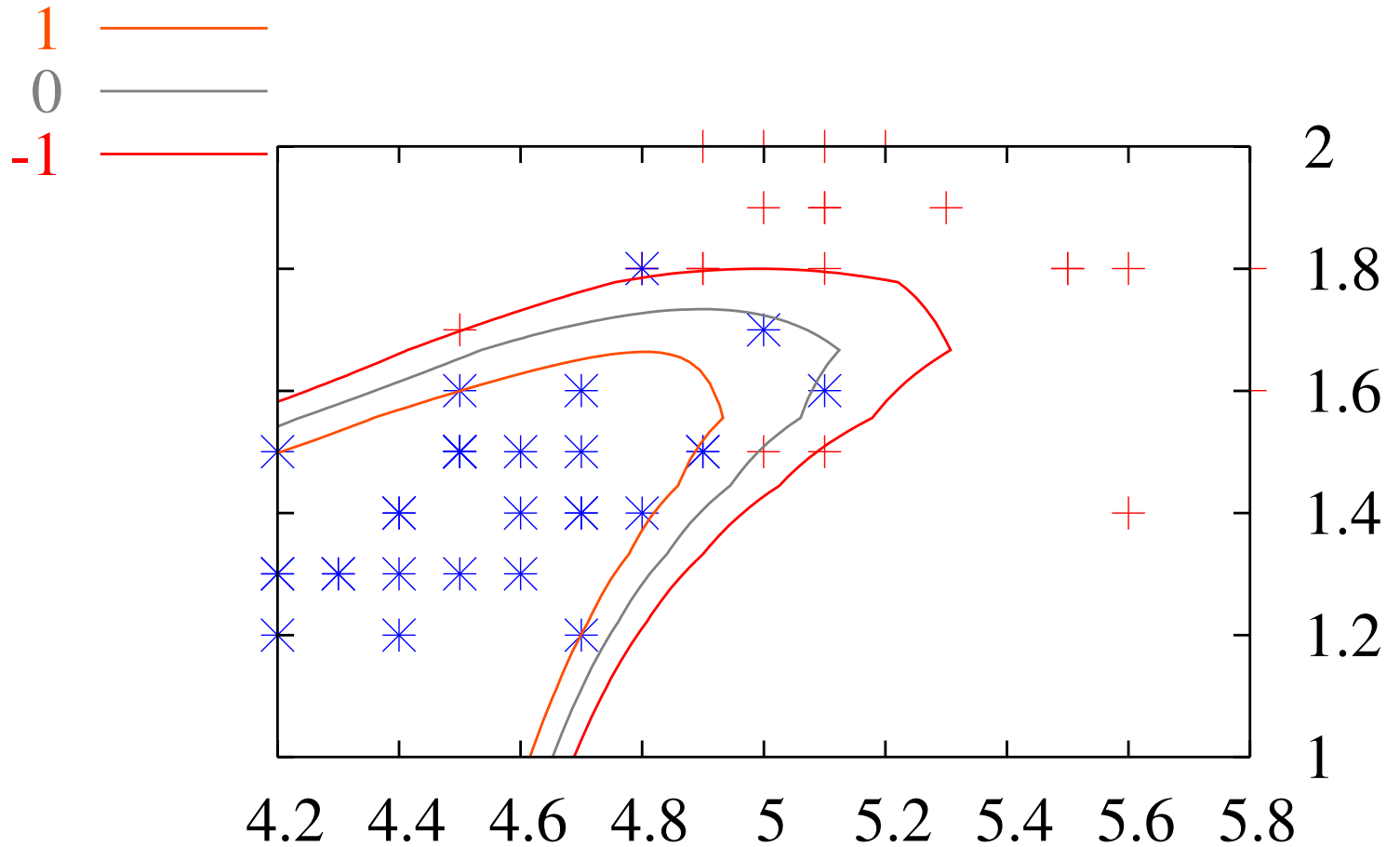
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Motivation for Neural Networks

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- Neural networks are inspired by neurons and their connections in the brain.
- An artificial neuron, called a *unit*, has inputs and an output.
- The output can be connected to other units.
- Typically, the output of a unit is computed by a linear function of its inputs passed through an activation function.
- Learning is adjusting weights to reduce error.
- Advantage: can learn non-linear functions.
- Disadvantage: more parameters and no guarantee of optimality.

Feedforward Neural Networks

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- *A feed-forward* neural network is the most common type. Its units are organized as a directed acyclic graph.
- *Input units* simply output the values of the input features.
- *Hidden units* input the values of other units and produce outputs for other units.
- *Output units* produce predictions of output features.
- One standard model is an initial “layer” of input units, which feed into a single layer of hidden units, which feed into the output units.

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Procedure $NNOutput(e, H, O)$

Inputs $e = \mathbf{x}$: the inputs of an example e

H : hidden units with activation function f_H
and weights \mathbf{w}_{H_j} for each hidden unit H_j

O : output unit with activation function f_O
and weights \mathbf{w}_O

for each hidden unit H_j

$$h_j \leftarrow f_H(\mathbf{w}_{H_j} \cdot \mathbf{x})$$

$$o \leftarrow f_O(\mathbf{w}_O \cdot \mathbf{h})$$

return o

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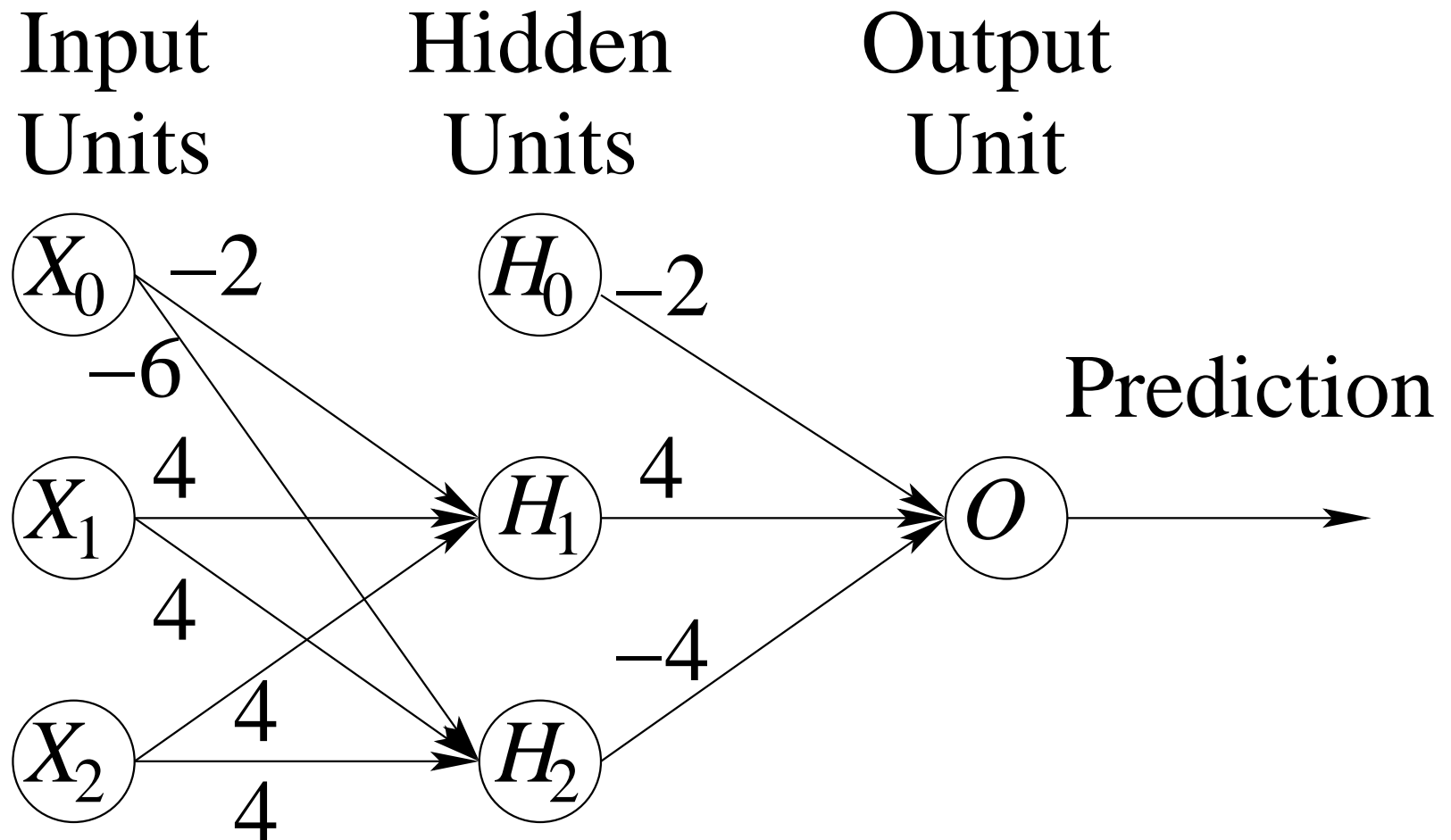
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Suppose $f(z) = 1/(1 + e^{-z})$ is the activation function. X_0 and H_0 always output 1.

Y	X_0, X_1, X_2	H_1	H_2	O
0	1, 0, 0	$f(-2) \approx 0.12$	$f(-6) \approx 0.00$	$f(-1.53) \approx 0.18$
1	1, 0, 1	$f(2) \approx 0.88$	$f(-2) \approx 0.12$	$f(1.05) \approx 0.74$
1	1, 1, 0	$f(2) \approx 0.88$	$f(-2) \approx 0.12$	$f(1.05) \approx 0.74$
0	1, 1, 1	$f(6) \approx 1.00$	$f(2) \approx 0.88$	$f(-1.53) \approx 0.18$

For example for $X_0, X_1, X_2 = 1, 0, 1$, then

$$H_1 = f(-2 * 1 + 4 * 0 + 4 * 1) = f(2) \approx 0.88$$

$$H_2 = f(-6 * 1 + 4 * 0 + 4 * 1) = f(-2) \approx 0.12$$

$$O \approx f(-2 + 4 * 0.88 - 4 * 0.12) \approx f(1.05) \approx 0.74$$

0.74 is closer to 1 than 0, so predict 1, which = Y .

Learning by Backpropagation

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Procedure $NNLearn(e, H, O, \mathbf{h}, o, \eta)$

Inputs $e = (\mathbf{x}, y)$: inputs and output of example e

H : hidden units with activation function f_H
and weights \mathbf{w}_{H_j} for each hidden unit H_j

O : output unit with activation function f_O
and weights \mathbf{w}_O

\mathbf{h}, o : outputs of hidden and output units

η : learning rate

$\delta_O \leftarrow$ derivative of $\mathbf{w}_O \cdot \mathbf{h}$ wrt error

$\mathbf{w}_O \leftarrow \mathbf{w}_O - \eta \delta_O \mathbf{h}$

for each hidden unit H_j

$\delta_{H_j} \leftarrow \delta_O * \text{derivative of } \mathbf{w}_{H_j} \cdot \mathbf{x} \text{ wrt } \mathbf{w}_O \cdot \mathbf{h}$

$\mathbf{w}_{H_j} \leftarrow \mathbf{w}_{H_j} - \eta \delta_{H_j} \mathbf{x}$

Comments on Backpropagation

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- If squared error and $f(z) = 1/(1 + e^{-z})$
 $\delta_O \leftarrow o(1 - o)(o - y)$
 $\delta_{H_j} \leftarrow h_j(1 - h_j)(w_{O,j})\delta_O$
- *NNLearn* needs to be applied for many epochs (passes over the examples).
- The weights need to be initialized to small random values.
- Multiple runs might be needed to find good weights (and hyperparameters such as number of hidden units).
- More sophisticated gradient descent algorithms are much faster than backpropagation.

- A *support vector machine* (SVM) assigns a weight α_i to each training example (\mathbf{x}_i, y_i) (\mathbf{x}_i is a vector of the values of the input features, and y_i is either -1 or 1).
- A [simplified] SVM computes a value on \mathbf{x} by summing over all examples:

$$h(\mathbf{x}) = \sum_i \alpha_i y_i k(\mathbf{x}, \mathbf{x}_i)$$

where k is a *kernel function*.

- Learning is by optimizing the error function:

$$\begin{aligned} &\text{minimize } \|h\|^2/2 + C \sum_i \max(0, 1 - y_i h(\mathbf{x}_i)) \\ &\text{subject to } 0 \leq \alpha_i \leq C \end{aligned}$$

where $\|h\|$ is the size of h in kernel space

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□ The effect of a kernel function is to expand the number of features for linear classification.

□ E.g., the kernel function $k(\mathbf{x}_i, \mathbf{x}_j) = (\mathbf{x}_i \cdot \mathbf{x}_j)^2$ corresponds to a dot product of all the quadratic combinations of input features. If $\mathbf{x}_i = (1, a, b)$ and $\mathbf{x}_j = (1, c, d)$, then

$$\begin{aligned}(\mathbf{x}_i \cdot \mathbf{x}_j)^2 &= (1 + ac + bd)^2 \\ &= 1 + 2ac + 2bd + a^2c^2 + 2acbd + b^2d^2 = \\ &(1, \sqrt{2}a, \sqrt{2}b, a^2, \sqrt{2}ab, b^2) \cdot (1, \sqrt{2}c, \sqrt{2}d, c^2, \sqrt{2}cd, d^2)\end{aligned}$$

□ The gaussian kernel $k(\mathbf{x}_i, \mathbf{x}_j) = \exp(-\gamma \|\mathbf{x}_i - \mathbf{x}_j\|^2)$ is a common choice. [γ is an extra parameter to choose.]

Slow SVM Learning Algorithm

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Procedure $SVM\text{Learner}(X, Y, E, k, C)$

Inputs X, Y : input features and target feature

E : set of training examples

k : kernel function

C : “soft margin” constant

initialize all example weights α_i to zero

repeat until termination

for each example $e = (\mathbf{x}_i, y_i) \in E$

$$\hat{y} \leftarrow \sum_j \alpha_j y_j k(\mathbf{x}_i, \mathbf{x}_j)$$

if $y_i \hat{y} < 1$ and $\alpha_i < C$ increase α_i

if $y_i \hat{y} > 1$ and $\alpha_i > 0$ decrease α_i

return α

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- Each example with $\alpha_i > 0$ is called a *support vector*.
- The solution will have $\alpha_i = 0$ if $y_i \hat{y} > 1$ and $\alpha_i = C$ if $y_i \hat{y} < 1$ with inbetween values for $y_i \hat{y} = 1$.
- An SVM has a global minimum with no local minima, though actual algorithms only get very close.
- Multiple runs might be needed to find the best kernel function and soft margin constant.
- More sophisticated SVM algorithms are much faster.

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- The k -nearest neighbor algorithm classifies a test ex. by finding the k closest training exs., returning the most common class.
- Suppose 10% noise (best possible test error is 10%). With enough training exs., a test ex. will agree with its nearest neighbor with prob. $(.9)(.9) + (.1)(.1) = .82$ (both not noisy or both noisy) and disagree with prob. $(.9)(.1) + (.1)(.9) = .18$.
- 1-NN converges to less than twice optimal error (3-NN to less than 32% higher).
- Problems: distance measure, number of exs. needed, find NNs efficiently.

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Run Continued

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- There are many algorithms for learning a single hypothesis.
- *Ensemble learning* will learn many hypotheses: run different algorithms or run the same algorithm on different training sets.
- *Bagging* runs a learning algorithm on repeated subsamples of the training set.
- If there are n examples, then a subsample of n examples is generated by sampling with replacement.
- On a test example, each hypothesis casts 1 vote for the class it predicts.

Boosting

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- In *boosting*, the hypotheses are learned in sequence.
- Both hypotheses and examples have weights with different purposes.
- After each hypothesis is learned, its weight is based on its error rate, and the weights of the training examples (initially all equal) are also modified.
- On a test example, when each hypothesis predicts a class, its weight is the size of its vote. The ensemble predicts the class with the highest vote.

Example Boosting Algorithm

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Procedure *AdaBoost*(E, A, t)

Inputs E, A : examples and learning algorithm

n : number of hypotheses to generate

initialize example weights w to $1/\text{number of exs.}$

for i from 1 to n

$h[i] \leftarrow A(E, w)$

$\epsilon \leftarrow$ sum weights of exs. missed by $h[i]$

for j from 1 to number of examples

if $h[i]$ is correct on example j

then $w[j] \leftarrow 0.5 * w[j] / (1 - \epsilon)$

else $w[j] \leftarrow 0.5 * w[j] / \epsilon$

weight of $h[i] \leftarrow \log((1 - \epsilon) / \epsilon)$

return $h[1 \dots n]$ and their weights

Example Run of AdaBoost

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Using the 14 examples as a training set:

- Example weights are initialized to $1/14$.
- The hypothesis $\text{windy} = \text{false} \leftrightarrow \text{class} = \text{pos}$ is wrong on 5 of the 14 examples.
- The weights of the correctly classified examples are multiplied by $14/18$, and incorrectly classified by $14/10$. An example will have a weight of $1/18$ or $1/10$.
- This hypothesis has a weight of $\log(9/5)$.
- Note that after weight updating, the sum of the correctly classified examples equals the sum of the incorrectly classified examples.

Example Run of AdaBoost, Continued

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- The next hypothesis must be different from the previous one to have error less than $1/2$.
- Now the hypothesis outlook = overcast \leftrightarrow class = pos has an error rate of $29/90$.
- The weights of the correctly classified examples are multiplied times $90/122$, and the incorrectly classified times $90/58$.
- This hypothesis has a weight of $\log(61/29)$.

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Regularization

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- *Overfitting* is when the learner uses regularities that appear in the training set, but do not appear in the test set.
- The learning algorithm might fit noise in the training data (outliers, random fluctuations).
- The learning algorithm might fit to unique values in the training data (id numbers).
- The learning algorithm might “memorize” the training examples (more parameters than examples).
- This can be observed when running the algorithm longer does better on the training set, but does worse on the test set.

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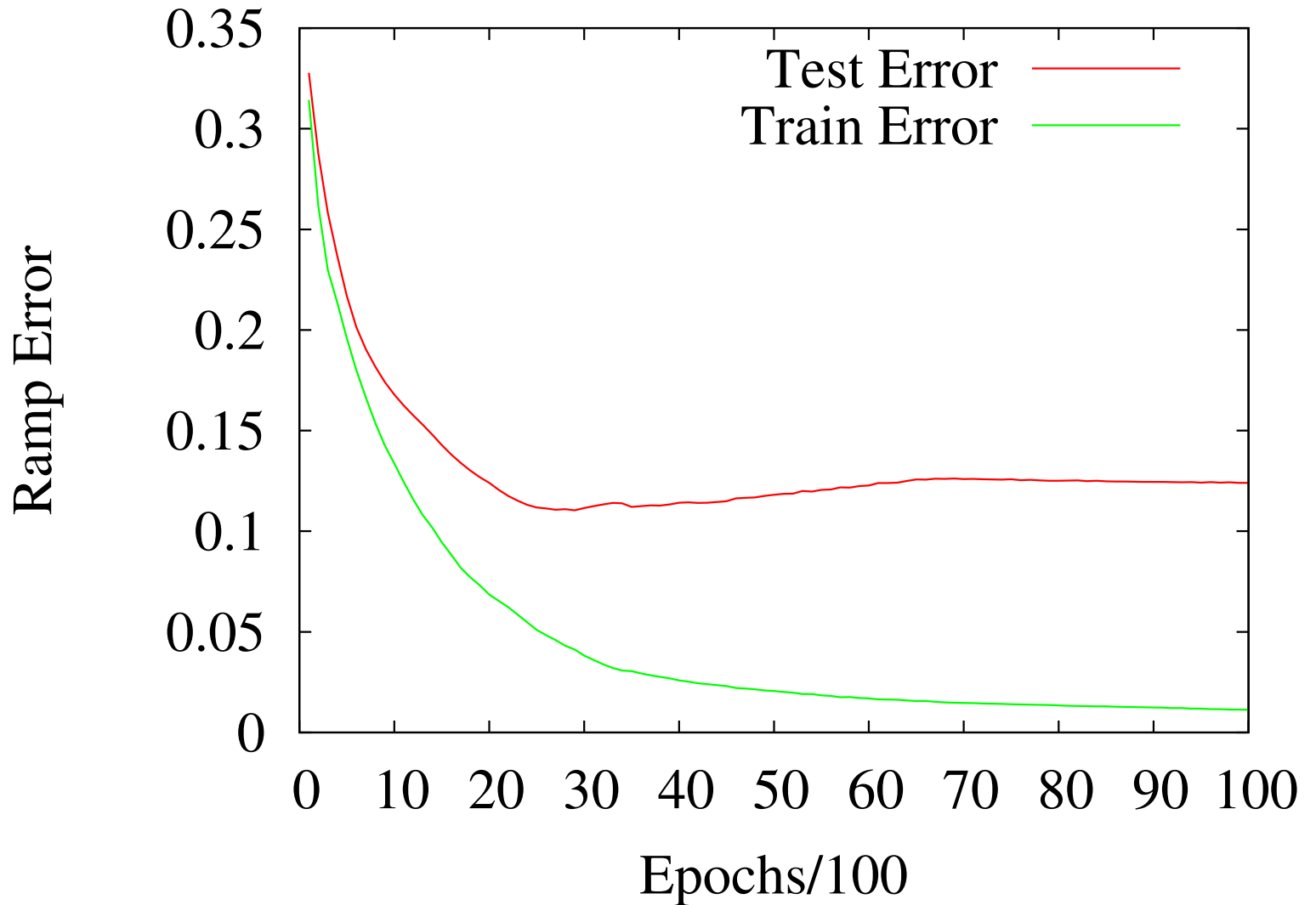
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- *A validation set* is one way to avoid overfitting.
- The original data is divided into a training set, a validation set and a test set (more about test set later).
- As the learning algorithm runs on the training set, keep track of its error on the validation set. Use the algorithm settings that minimize validation error.

k -Fold Cross-Validation

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- In k -fold cross-validation, the data is divided into a training set and a test set.
- The training set is evenly divided into k folds.
- Then k times: Train the learning algorithm on $k - 1$ folds and validate on the remaining fold.
- Average over the results.

Regularization

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- *Regularization* is a way to avoid overfitting.
- A learning model might have many parameters.
 - A NN with 10 hidden units has over 10 times more parameters than a linear model.
 - In natural language tasks, a parameter for every word (millions of words) or every word-word combination.
- If a linear model has as many or more parameters than examples, then the examples can be “memorized”.
- Regularization modifies the error function so there is a penalty for larger parameters.

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- Suppose we want to minimize hinge loss over a set of examples. That is, find \mathbf{w} minimizing:

$$\sum_i \max(0, 1 - y_i * (\mathbf{w} \cdot \mathbf{x}_i))$$

- Perhaps this formulation leads to overfitting. To regularize this, we can add a term that penalizes large weights.

$$\lambda(\mathbf{w} \cdot \mathbf{w})/2 + \sum_i \max(0, 1 - y_i(\mathbf{w} \cdot \mathbf{x}_i))$$

- The weight update in the *LinearLearn* algorithm becomes:

$$\mathbf{w} \leftarrow \mathbf{w}(1 - \eta\lambda/m) + \eta\delta\mathbf{x}$$

where m is the number of examples.

- Use validation to find λ .

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- Recall: the original data is divided into a training set, a validation set and a test set.
- Do not look at the test set.
- Do all the processing and analysis you want using the training and validation sets.
- Do not look at the test set.
- Select a hypothesis (or a few hypotheses) by training on the training and validation sets combined, using the best settings on the best algorithm(s).
- Evaluate hypothesis (or hypotheses) on the test set.

Can I Use the Test Set Again?

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- In theory, **no**. By reusing the test set, you are in effect using the test set for training (searching for an algorithm that does well on the test set).
- In practice, a test set can be reused once or twice without getting into too much trouble, probably, maybe.
- Using many different datasets (and so many different test sets) from a variety of domains can be used to empirically compare algorithms.
- Moral: Prepare a second test set (maybe a third or fourth) so you have one in reserve after a possible initial failure.

Error Rate and Comparing Algorithms

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Paired Difference

- What is the true error rate?
 - If ϵ is the test error rate, and m examples in the test set ($m > 100$), then a 95% confidence interval is:

$$\epsilon \pm 1.96 \sqrt{\frac{\epsilon(1-\epsilon)}{m-1}}$$

- Is algorithm A better than algorithm B?
 - Use the paired-difference test (next page assuming more than 100 examples in test set).

Paired-Difference Test

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Let h_1 and h_2 be the two hypotheses to compare.
For each $(\mathbf{x}_i, y_i) \in E$:

$$\text{let } d_i = \begin{cases} 1 & \text{if } h_1(x_i) = y_i \neq h_2(x_i) \\ 0 & \text{if } h_1(x_i) = h_2(x_i) \\ -1 & \text{if } h_1(x_i) \neq y_i = h_2(x_i) \end{cases}$$

Calculate:

$$u = \frac{\sum_{i=1}^m d_i}{m} \quad s^2 = \frac{\sum_{i=1}^m (d_i - u)^2}{m - 1} \quad z = \frac{u}{s/\sqrt{m}}$$

If $z \geq 1.96$, answer “ h_1 is better than h_2 ”.

If $z \leq -1.96$, answer “ h_1 is worse than h_2 ”.

The significance is 0.05.

If $|z| < 1.96$, then answer “I don’t know”.