Deep feed-forward models

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- About deep feed-forward models
- Loss function and output layers
- Activation functions in hidden layers
- Universal approximation: depth matters
- Backprop: efficient computation of the loss gradient

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Deep feed-forward network

- $\cdot\,$ the simplest formulation of a deep model
- network: consists of a number of interconnected processing elements
- implemented as a sequence of fully connected layers i.e. affine transformations with non-linear activation

Goal:

- approximate the desired function $y = f^*(x)$ with a parametric model $\hat{y} = f(x, \Theta)$.
- $\cdot\,$ the model f maps the input ${\bf x}$ into predictive output ${\hat {\bf y}}$
- \cdot we jointly learn the parameters Θ from end to end

About deep feed-forward models

Details:

- the function $y = f^*(x)$ corresponds to the exact relationship between input x and output y
- our model $\hat{y} = f(x, \Theta)$ approximates the exact function
- we want to find the set of parameters Θ^* that provides the "best" approximation
- problem: we do not know what the function f*(x) looks like in most x; we only know f*(x) in a finite training set {(x_i, y_i)}^N_{i=1}.
- hence, we care about the ability to generalize
- the choice of the model will depend on the data (cf. no free lunch theorem)

Basic properties of deep models:

- information flows from input to output (feed-forward), there are no loops
- can be represented as a composition of simpler functions: $f(\mathbf{x}, \mathbf{\Theta}) = o(f_L(f_{L-1}(\cdots(f_1(\mathbf{x}, \mathbf{\Theta}_1)), \cdots), \mathbf{\Theta}_{L-1}), \mathbf{\Theta}_L)),$
- we refer to f_i as layers
- each layer includes exactly one non-linear activation
- the model depth (*L*): the number of layers f_i

About deep feed-forward models

We express the model in terms of auxiliary variables $\{h_l\}$:

 $h_1 = f_1(\boldsymbol{x}, \boldsymbol{\Theta}_1)$

$$h_{L-1} = f_{L-1}(h_{L-2}, \Theta_{L-1})$$
$$h_L = f_L(h_{L-1}, \Theta_L)$$
$$f(\mathbf{x}, \Theta) = o(h_L)$$

We denote the auxiliary variables as hidden or latent features

Layer width (D_l) : dimension of its feature vector, $\boldsymbol{h}^l \in \mathbb{R}^{D_l}$.

The supervision involves only the input **x** and the output **y**: model has a freedom to arrange hidden features in a manner that ensures the best approaximation. The basic form: sequence of fully-connected layers

• each f_i models an elementary non-linear transformation: parametric afine mapping with non-linear activation σ :

$$\mathbf{f}_k(\mathbf{h}_{k-1}) = \boldsymbol{\sigma}(\mathbf{W}_k\mathbf{h}_{k-1} + \mathbf{b}_k)$$

- We must get to know the fully-connected layers well:
 - basis for more complex layers (eg. convolutional)
 - building blocks of more complex architectures (eg. attention)

Other names:

- (feed-forward, deep) **fully-connected model** (with affine transformations)
- multi-layer perceptron (MLP)
- (feed-forward) (artificial) neural network

Fully connected layer



$$f(\mathbf{x}; \mathbf{W}, \mathbf{b}) = \boldsymbol{\sigma}(\mathbf{W} \cdot \mathbf{x} + \mathbf{b})$$
$$\boldsymbol{\sigma}(\mathbf{s})_i = \boldsymbol{\sigma}(\mathbf{s}_i)$$

Problem: determine the structure, equations and total number of parameters of a fully connected model for 2D data. if we know that the layer widths are: 5, 10, 5, 2..

Fully connected model

Problem: determine the structure, equations and total number of parameters of a fully connected model for 2D data. if we know that the layer widths are: 5, 10, 5, 2..



Relation to artificial neural networks:

- artificial neural networks study machine learning algorithms that are inspired by early models of the human brain
- on the other hand, deep learning is concerned with good generalization on real data

Question: how deep should a fully connected model be?

Seductive idea: L=1!

$$f(\mathbf{x}, \mathbf{\Theta} = (\mathbf{w}, b)) = \boldsymbol{\sigma}(\mathbf{w}^{\top}\mathbf{x} + b)$$

- strength: usual loss functions lead to convex optimization
- strength: guaranteed convergence
- weakness: our world is non-linear.

If a single layer is not an option, what solutions remain?

• Solution: use a nonlinear solution Φ in order to map the data into linearly separable **features**:

$$f(\mathbf{x}, \Phi, \mathbf{\Theta} = (\mathbf{w}, b)) = \Phi(\mathbf{x})^{\top}\mathbf{w} + b$$

- Three dominant ways to construct Φ :
 - · design a generic function Φ (suitable for all algorithms)
 - · hand-craft an algortihm-specific function Φ ,
 - · learn the function on the data $\Phi(x|\Theta_{\Phi})$.

Example: kernel functions

 eg. RBF function k(x, ·) implicitly maps the data into the infinite-dimensional feature vector Φ(x)

Problem: such functions assume local smoothness

 unfortunately, local smoothness is not good enough when dim(x)=10⁵



Hand-crafted features

Examples: SIFT descriptor (vision), word normalization (language), MFCC descriptors (speech).

Problem: requires domain knowledge, time-consuming process

Problem: (today we know) limited generalization power



The only approach left: learn the function $\Phi(x|\Theta_{\Phi})$, by optimizing parameters Θ_{Φ}

We can try to learn the layers separately: first learn the features Θ_{Φ} (eg. unsupervised), and only then learn the classifier w, b

- that would work better than linear model
- but nobody succeeded by learning more than two layers that way

Only one approach remains: learn a deep model from end to end:

· joint learning $\boldsymbol{\Theta} = (\boldsymbol{w}, b) \cup \boldsymbol{\Theta}_{\boldsymbol{\Phi}}$

Deep learning (end-to-end)

Advantages with respect to generic and hand-crafted mappings:

- we specify a *class* of functions $\Phi(x|\Theta_{\Phi})$ except of a specific function $\Phi(x)$
- \cdot class of functions is determined by the model structure
- we can have arbitrarily many layers (there is a sweet spot in practice)

Disadvantage with respect to generic and hand-crafted design:

- the optimization problem is no longer convex
- global convergence is not guaranteed

However, it turns out that the non-convex loss does not pose a problem in practice

Deep models are best suited when the data is generated by a composition of factors, for example, the face consists of a mouth, eyes, nose.....

If such factors exist, factored recognition can ensure efficient representation in the input space.

Some works suggest exponential efficiency with respect to approaches that rely on local smoothness prior [montufar14nips]

• shallow models, prototypes (k-NN), kernel functions

Deep learning (end-to-end)

Problem: learn a function that maps 2D points into RGB color

• these are the training data



Deep learning (end-to-end)

Problem: learn a function that maps 2D points into RGB color

- these are the training data
- intuitively unclear how to generalize



Problem: learn a function that maps 2D points into RGB color

- these are the training data
- intuitively unclear how to generalize
- the problem becomes much easier if we express the model as a sum of two independent 1D functions

$$f(x,y) = (f_R(x) + f_B(y))/2$$



- Consider the following function of two binary variables: $f^*(\mathbf{x}) = (x_0 \land \overline{x_1}) \lor (\overline{x_0} \land x_1).$
- Let us try to learn a linear model to approximate f^* : $f(\mathbf{x}, \mathbf{\Theta} = (\mathbf{w}, b)) = \mathbf{w}^\top \mathbf{x} + b$
- We are looking for $\Theta^* = (w^*, b^*)$ such that mean square error of the predictions becomes minimal:

$$J_{\text{MSE}}(\mathbf{Y}, f(\mathbf{X}, \boldsymbol{\Theta})) = \frac{1}{4} \sum_{i=1}^{4} (f(\mathbf{x}_i, \boldsymbol{\Theta}) - y_i)^2$$

• Later we shall see that such loss is not a good choice for classification problems, but here it is convenient because we can get the solution in a closed form.

Notation:

$$\mathbf{w}' = \begin{bmatrix} w_1, w_2, b \end{bmatrix}^{\top}, \quad \mathbf{X}' = \begin{bmatrix} x_{11} = 0, x_{12} = 0, 1\\ x_{21} = 0, x_{22} = 1, 1\\ x_{31} = 1, x_{32} = 0, 1\\ x_{41} = 1, x_{42} = 1, 1 \end{bmatrix}, \quad \mathbf{y} = \begin{bmatrix} y_1 = 0\\ y_2 = 1\\ y_3 = 1\\ y_4 = 0 \end{bmatrix}$$

Let us express the loss in a convenient form:

$$J_{MSE}(y, X', w') = \frac{1}{N} \|X'w' - y\|_{2}^{2} = \frac{q^{\top}q}{N}, q = X'w' - y$$

Now we can determine the gradient by chaining rule:

$$\nabla_{w'} J_{\text{MSE}}(y, \mathbf{X}', w')^{\top} = \frac{\partial J}{\partial w} = \frac{\partial J}{\partial q} \frac{\partial q}{\partial w}$$
$$= \frac{2 \cdot q^{\top}}{N} \cdot \mathbf{X}' = \frac{2}{N} (\mathbf{X}' w' - y)^{\top} \mathbf{X}'$$

We are looking for the minimum of the function J:

$$abla_{w'}J = \mathbf{0} \rightarrow w' = \left(X'^{\top}X'\right)^{-1}X'^{\top}y$$

Solution: **w**^{*} = **0**, *b*^{*} = 0.5 (??!)

Conclusion: linear model has insufficient capacity to solve the XOR problem.

- Minski and Papert published this in their 1969 book: Perceptrons: An Introduction to Computational Geometry
- this was viewed as a limitation of all learning approaches and contributed to the first AI winter (1974-1980)
- backprop was invented in 1970 by Seppo Linnainmaa...

- Let us introduce an additional non-linear layer
 - $\cdot\,$ it has to be such if we wish a non-linear composite model
- The hinge function is a default non-linear function today: $g(x) = \text{ReLU}(x) = \max(0, x).$
- Non-linearity affects each vector element separately: $g(x)_i = g(x_i)$
- Now we can formulate our composite modeli:

$$f(\mathbf{x}, \mathbf{\Theta}) = \mathbf{w}_2^\top \mathbf{h} + b_2,$$
$$\mathbf{h} = g(\mathbf{W}_1^\top \mathbf{x} + b_1)$$

- *h*: vector of (learned) hidden features
- *W*₁, *b*₁: learned parameters for mapping data to hidden features

Solution:
$$W_1 = \begin{bmatrix} 1 & 1 \\ 1 & 1 \end{bmatrix}$$
, $b_1 = \begin{bmatrix} 0 \\ -1 \end{bmatrix}$, $w_2 = \begin{bmatrix} 1 \\ -2 \end{bmatrix}$, $b_2 = 0$.

$$f(x,y) = 1 \cdot \max(x+y,0) - 2 \cdot \max(x+y-1,0) + 0$$



Solution:
$$W_1 = \begin{bmatrix} 1 & 1 \\ 1 & 1 \end{bmatrix}, b_1 = \begin{bmatrix} 0 \\ -1 \end{bmatrix}, W_2 = \begin{bmatrix} 1 \\ -2 \end{bmatrix}, b_2 = 0.$$

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$$f(x,y) = 1 \cdot \max(x+y,0) - 2 \cdot \max(x+y-1,0) + 0$$



- We have shown the solution without going into details of how to find it
- parameters of deep models are most often determined through gradient optimization of the loss.
- \cdot The presented solution is the global minimum of the loss
 - in the general case, the gradient descent will lead to some local minimum (if we train to convergence), since the loss will typically be non-convex
 - · choice of the local minimum will depend on initialization

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Learning by minimizing empirical risk

• Learning corresponds to finding Θ^* that minimizes the empirical risk:

$$J(\Theta|\mathbf{X},\mathbf{Y}) = \frac{1}{N} \sum_{i=1}^{N} \ell(\mathbf{y}_i, f(\mathbf{x}_i, \Theta)) + \lambda \Omega(\Theta)$$
$$\Theta^* = \arg\min_{\Theta} J(\Theta|\mathbf{X}, \mathbf{Y})$$

- The loss $\ell(y, \hat{y})$ reflects our ``disappointment'' due to model prediction \hat{y} being different than the desired value *y*.
- The regularizer $\Omega(\Theta)$ penalizes parameter vectors that correspond to mappings that we assume unlikely

$$\ell_{01}(\boldsymbol{y}, \hat{\boldsymbol{y}}) = \begin{cases} 0, & ext{ako } \boldsymbol{y} = \hat{\boldsymbol{y}} \\ 1, & ext{inače} \end{cases}$$

• This loss is not differentiable, so that minimizing $J(X, Y, \Theta)$ requires combinatorial optimization

Probabilistic loss

- Suppose we allow probabilistic predictions in the form of a distribution: $P(\hat{Y}|x; \Theta)$
- Then, a principled loss can be formulated as negative log-likelihood:

$$\ell_{\mathrm{MLE}}(y, \hat{Y}) = -\log P(\hat{Y} = y | x; \Theta)$$

• we can formulate regression by predicting a normal (Gaussian) distribution (simplest case - unit covariance)

$$p(\hat{\mathbf{Y}} = \mathbf{y} | \mathbf{x}; \mathbf{\Theta}) = \mathcal{N}(\mathbf{y} | \mu = f(\mathbf{x}, \mathbf{\Theta}), \Sigma = \mathbf{I})$$

• we can formulate classification by predicting a categorical posterior distribution (generalized Bernoulli)

$$P(\hat{Y}|\mathbf{x};\boldsymbol{\Theta}) = \sigma(f(\mathbf{x},\boldsymbol{\Theta}))$$

• we can even formulate deterministic prediction by plugging in the Dirac δ -distribution (this leads to 0 – 1 loss)

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Negative log-likelihood

- Maximum likelihood estimation is versatile:
 - no need for model-specific loss formulations
 - the only requirement is probabilistic output: $p(\hat{Y}|\mathbf{x}; \boldsymbol{\Theta})$
 - the loss function is $\ell_{\text{MLE}}(y, \hat{Y}) = -\log p(\hat{Y} = y | \mathbf{x}; \boldsymbol{\Theta})$
- Gaussian predictions with unit covariance lead to mean square error:

$$\ell_{MSE}(\mathbf{y}, \hat{\mathbf{Y}}) = (\mathbf{y} - f(\mathbf{x}, \boldsymbol{\Theta}))^2$$

Categorical predictions lead to multinomial logistic loss:

$$\ell_{\mathsf{MLL}}(y, \hat{Y}) = -\log \sigma_y(f(\mathbf{x}, \Theta))$$

• All these variants of negative log-likelihood are differentiable.

 \Rightarrow can be learned with gradient descent
Negative log-likelihood vs cross-entropy

Sometimes we wish to treat the labels as random variables

- eg. use smooth labels instead of one-hot ones (a form of regularization)
- eg. produce the labels with another probabilistic model (distillation, semi-supervised learning)

In this case, the loss can be expressed as cross entropy:

$$\ell_{\rm CE}(Y,\hat{Y}) = -\sum_{y} p(Y=y) \log P(\hat{Y}=y|\mathbf{x};\boldsymbol{\Theta})$$

The following statements are easily shown (homework):

- cross entropy is related to KL divergence, a measure of "distance" between two distributions
- negative log-likelihood is a special case of cross entropy

Categoric predictions

A categoric model M must meet the following constraints:

- $M_i(\mathbf{x}, \mathbf{\Theta}) \in [0, 1] \ \forall i,$
- $\sum_{i}^{C} M_i(\mathbf{x}, \mathbf{\Theta}) = 1$

Typically, we ensure this through softmax activation:

$$P(\hat{Y}|\mathbf{x}; \mathbf{\Theta}) = M(\mathbf{x}, \mathbf{\Theta}) = \text{softmax}(f(\mathbf{x}, \mathbf{\Theta}))$$

Learning with softmax enforces unnormalized log-posteriors in the the last layer features $z = f(x, \Theta)$ (also known as logits):

$$z_i = \log \operatorname{const} + \log P(\hat{y} = i | \mathbf{x}; \boldsymbol{\Theta})$$
.

Proof:

softmax
$$(z)_y = \frac{\exp(z_y)}{\sum_j \exp(z_j)} = P(\hat{Y} = y | x; \Theta)$$
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Classification with softmax

Let us apply the negative log-likelihood to the softmax: $\ell(y, \hat{Y}) = -\log \operatorname{softmax}(z)_y = \log \sum_j \exp(z_j) - z_y \approx \max_j z_j - z_y$

We draw the following intuitive conclusions:

- when the model is correct $(\max_j z_j = z_y)$ the loss is ≈ 0 .
- when the model is incorrect $(\max_j z_j \neq z_y)$, the loss is mostly affected by the strongest incorrect prediction.
- such behavior is very similar to 0-1 loss: negative log-likelihood is an upper bound of the 0-1 loss.

The following relation is easily shown (homework):

$$\frac{d\ell_{\mathsf{MLE}}(y,\mathsf{softmax}(z))}{dz_i} = \mathsf{softmax}(z)_i - \llbracket y = i \rrbracket$$

Softmax properties

Invariance to the addition of a constant: $softmax(z) = softmax(z + c) = softmax(z - \max_{j} z_{j})$

A better name (which did not catch on): softargmax

"Real softmax" would correspond to log-sum-exp:

$$LSE(\mathbf{z}) = \log \sum_{i} e^{z_i} = \max(\mathbf{z}) + \log \sum_{i} e^{z_i - \max(\mathbf{z})}$$

Softmax-weighted average corresponds to scalar product of softmax and input:

softmax-mean(z, q) = softmax $(z)^{\top}x$

Although the softmax output is C-dimensional, there are only C – 1 degrees of freedom (output is a distribution)

Consequently, we can fix one input (eg. to 0) without reducing the generality.

Often there is no difference between the two variants

• in these cases we choose C-dimensional inputs for simplicity

Binary classification: sigmoid-activated outputs

If C = 2 then:

$$P(\hat{y} = 1 | \mathbf{x}) = \text{softmax}(z)_1 = \frac{\exp(z_1)}{\exp(z_0) + \exp(z_1)}$$
$$= \frac{1}{1 + \exp(z_0 - z_1)}$$

If we set $z_0 := 0$, we get:

$$P(\hat{y}=1|\mathbf{x})=\sigma(z_1)$$

- \Rightarrow soft-max generalizes the sigmoid activation for C > 2
- ⇒ categorical distribution generalizes Bernoulli distribution for C>2.

The following relation is easily shown (homework):

$$\frac{d\ell_{CE}(y,\sigma(z))}{dz} = \sigma(z) - y$$
³⁶

Why prefer negative log-likelihood to mean-square error for classification?

$$\ell_{\mathsf{MSE}}(y,\sigma(z)) = (y-\sigma(z))^2$$

Let us observe the ℓ_{MSE} gradient with respect to the logits: $\frac{\partial \ell_{MSE}(y, \sigma(z))}{\partial z} = 2(\sigma(z) - y)(1 - \sigma(z))\sigma(z)$

When the sigmoid saturates (z >> 0 ili $z \ll 0$), the loss gradient is small:

- this holds regardless of whether $\sigma(z)$ is close to y or not
- the model can not learn from such examples.

MSE as classification loss??

Main weakness: MSE ignores the intrinsic constraints of probabilistic distribution

Suppose we have data \mathbf{x}_1 i \mathbf{x}_2 that belong to the class y=2: $\mathbf{y}_1^{OH} = \mathbf{y}_2^{OH} = [0,0,1]$

Moreover, suppose we get the following predictions: $P(Y|x_1) = [0.8, 0, 0.2], P(Y|x_2) = [0.4, 0.4, 0.2]$

Equally wrong predictions lead to different losses:

 $L_{MSE}(\mathbf{x}_1, \mathbf{Y}_1^{OH}) = 1.28, L_{MSE}(\mathbf{x}_2, \mathbf{Y}_2^{OH}) = 0.96$

In the classification context, MLE outperforms MSE.

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We had solved the problem by inserting a non-linear layer

• it had to be non-linear, otherwise the composite problem would be (again) linear.

The hinge function (ReLU) is the default non-linearity today:

$$g(x) = \operatorname{ReLU}(x) = \max(0, x)$$

Non-linearities activate each dimension separately:

$$g(\mathbf{x})_i = g(\mathbf{x}_i)$$

ReLU activation

The hinge function (rectified linear unit):

 $g(x) = \operatorname{ReLU}(x) = \max(0, x)$

Advantages:

.

- in the active state it admits both the signal (forward pass) and the gradients (backward pass)
- allows to propagate gradients according to output activations

"Shortcoming" 1: the gradient is undefined at x = 0

• implementations use either the left (0) or the right (1) subgradient.

ReLU activation

Shortcoming 2: in the non-active state the hinge function stops both the signal and the gradient, but:

- \cdot there exist bijective generalizations
 - Leaky ReLU: $g(x, \alpha) = \max(0, x) + \alpha \min(0, x)$.
 - Soft Plus: $g(x) = \log(1 + e^x)$.
- batch normalization ensures that ReLU inputs have zero mean and unit variance
 - ⇒ in each learning iteration we have 50% active activations for each feature

Shortcoming 3: the outputs have non-zero means

- this problem is again solved by batch normalization
- $\cdot\,$ models that do not use batchnorm prefer GELU activation

sigmoid-like activations

Sigmoid: $\sigma(x) = (1 + \exp(-x))^{-1}$

- supresses the gradijent when saturated
 - the learning stops due to vanishing gradients
- mostly avoided in modern architectures
 - they are sometimes used in specialized roles (LSTMs, flows)

hyperbolic tangent: $tanh(x) = \frac{exp(2x)-1}{exp(2x)+1}$

- similar to the sigmoid, but better due to resembling identitety around x = 0
 - it ensures zero-mean outputs (for zero-mean inputs)
 - simple backprop in case of small inputs
- it still suppresses the gradients when saturated
 - relationship between tanh and σ :

$$\tanh(x) = 2\sigma(2x) - 1$$

Exp.-linear function:

$$\mathsf{ELU}(x;\alpha) = \lambda \cdot \begin{cases} \alpha(e^x - 1); \text{ for } x < 0\\ x; \text{ for } x \ge 0 \end{cases}$$

Gaussian Error Linear Unit:

GELU (x) =
$$x \cdot \Phi(x) = x \cdot \frac{1}{2} \left[1 + erf(x/\sqrt{2}) \right]$$

Advantages: does not saturate; zero mean and unit variance

Disadvantages: complex, non-bijective (requires caching)

Other contenders:

- maxout (see the book...)
- any non-linear function may work fine (even cosine...)

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Theorem: a fully connected model with at least one hidden layer with non-polynomial activation can approximate any finite-dimensional Borel measureable function with arbitrary small error, if the model has enough hidden dimensions.

- each continuous fuction defined on a bounded closed subset of \mathbb{R}^n is Borel measureable.
- no need to adjust activations: it suffices that we have one hidden layer.

The theorem only guarantees sufficient capacity:

- if the function *f** were known, then we could approximate it arbitrarily well
- however, the function is not known: instead we only have training data $(\mathcal{X}, \mathcal{Y})$.

The theorem says nothing about whether some algorithm can learn an f^* that generalizes well.

• the theorem only states that a sufficiently powerful model can overfit to training data

The theorem does not specify the hidden dimensionality that ensures a given approximation error, but there is an upper bound

• in the worst case, we need exponentially many hidden features:

 $\dim(\mathbf{h}) \sim O(a^{\dim(\mathbf{x})})$

- each of these features corresponds to the input configuration that requires a distinctive output
 - intuition: we require $O(2^n)$ minterms to learn an arbitrary logical function of n variables

Universal approximation theorem

Problem: design a two-level model that uses affine and Relu mappings to approximate the following function:



Rješenje:

- h10 = np.maximum(X-0,0)
- h11 = np.maximum(X-1,0)
- h12 = np.maximum(X-2,0)
- h13 = np.maximum(X-3,0)
- h14 = np.maximum(X-6,0)
- h21 = 1*h10 1*h11 + 2*h12 3*h13 + 1*h14

Literature: http://neuralnetworksanddeeplearning.com/chap4.html

Deep models may require fewer hidden activations

• some functions can be very efficiently represented with composite mappings.

Deep models can be exponentially more efficient than their shallow counterparts

Consider learning the n-way XOR:

- shallow model requires $O(2^n)$ hidden activations
- a suitable deep model requires O(n) hidden activations

Deep models and mapping efficiency

ReLU-activated models define piecewise linear functions over regions of the input space:

- the number of these regions is proportional to model flexibility (capacity)
- deep models have exponentially more regions than shallow models with the same number of activations



Figure 1: Binary classification using a shallow model with 20 hidden units (solid line) and a deep model with two layers of 10 units each (dashed line). The right panel shows a close-up of the left panel. Filled markers indicate errors made by the shallow model.

[montufar14nips]



Empirical results show that image classification models generalize better when we increase the depth

• x: model depth, y: classification accuracy

Increasing the model width leads to smaller improvements



Deep convolutional models for image classification generalize better than the shallow ones on SVHN sequence transcription

• x: number of parameters, y: classification accuracy

Increased depth introduces a bias that favors generalization

- $\cdot\,$ shallow models already overfit with $2\cdot 10^7$ parameters
- deep models generalize well even with $6 \cdot 10^7$ parameters 53

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Supervised learning

Forward pass —computes the model predictions $\hat{y} = f(x, \Theta)$ and the loss $J(y, \hat{y}) = J(y, f(x, \Theta))$

Backward pass —computes the loss gradient with respect to model parameters: $\nabla_{\Theta} J(\mathbf{y}, f(\mathbf{x}, \Theta)) = (\frac{\partial J(\mathbf{y}, f(\mathbf{x}, \Theta))}{\partial \Theta})^{\top}$

Optimization algorithm —typically a variant of the stochastic gradient descent:

- $\cdot \Theta' = \Theta \delta \cdot \nabla_{\Theta} J(\mathbf{y}, f(\mathbf{x}, \Theta))$
- more details about this some other time...

Backward propagation of errors (*backprop*); a simple and efficient approach to compute gradients of composite functions.

Derivatives of composite functions

Chain rule: a recipe to find derivatives of a composition of differentiable functions.

In the scalar case, y = g(x) and z = f(y) = f(g(x)), we get: $\frac{dz}{dx} = \frac{dz}{dy}\frac{dy}{dx} = \frac{df(y)}{dy}\frac{dg(x)}{dx}$

In the vector case, y = g(x) and z = f(y) = f(g(x)), we get:

$$\frac{\partial z}{\partial \mathbf{x}} = \frac{\partial z}{\partial \mathbf{y}} \frac{\partial \mathbf{y}}{\partial \mathbf{x}} \quad \text{or} \quad \nabla_{\mathbf{x}} z = \left(\frac{\partial \mathbf{y}}{\partial \mathbf{x}}\right)^{\top} \nabla_{\mathbf{y}} z.$$

• $\frac{\partial z}{\partial \mathbf{v}}$ and $\frac{\partial \mathbf{y}}{\partial \mathbf{x}}$ are Jacobians 1 × *n* and *n* × *m*;

•
$$\frac{\partial z}{\partial x_i} = \sum_j \frac{\partial z}{\partial y_j} \frac{\partial y_j}{\partial x_i}$$

We perform such steps to compute the gradients in each layer.

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Backprop: iterative application of the chain rule

Consider a model f_{Θ} that maps a datum **x** into predictions \hat{y} :

 $\hat{y} = f(\mathbf{x}, \boldsymbol{\Theta})$

The loss gradient with respect to parameters of the *l*-th layer is:

$$\frac{\partial \mathcal{L}(y,\hat{y})}{\partial \Theta^{l}} = \frac{\partial \mathcal{L}(y,\hat{y})}{\partial \hat{y}} \frac{\partial \hat{y}}{\partial h^{L}} \frac{\partial h^{L}}{\partial h^{L-1}} \cdots \frac{\partial h^{l}}{\partial \Theta^{l}}$$
$$= \frac{\partial \mathcal{L}(y,\hat{y})}{\partial h^{L}} \frac{\partial f^{L}(h^{L-1},\Theta^{L})}{\partial h^{L-1}} \cdots \frac{\partial f^{l+1}(h^{l},\Theta^{l+1})}{\partial h^{l}} \frac{\partial f^{l}(h^{l-1},\Theta^{l})}{\partial \Theta^{l}}$$

For each layer we must determine the partial derivative...

- ... with respect to parameters (if they exist) $\frac{\partial f^l(\mathbf{h}^{l-1}, \mathbf{\Theta}^l)}{\partial \mathbf{\Theta}^l}$,
- and with respect to the input $\frac{\partial f^l(h^{l-1}, \Theta^l)}{\partial h^{l-1}}$ (only if we have not computed all required gradients);
- problem: Θ^l can be a matrix (fully connected layer)
- problem: h^l i Θ^l can be a 4th-order tensor (conv. layer)

Gradients with respect to higher-order tensors (>1)

We *can* compute the gradients with respect to tensors in the same way as for vectors:

- we first determine the gradients for a vectorized tensor...
- ... and subsequently reshape them according to the tensor shape.

Assume $\mathbf{X} \in \mathbb{R}^{m_1} \times \mathbb{R}^{m_2} \times \cdots \mathbb{R}^{m_M}$, $\mathbf{Y} \in \mathbb{R}^{n_1} \times \mathbb{R}^{n_2} \times \cdots \mathbb{R}^{n_N}$

- Then $\frac{\partial \operatorname{vec}(Y)}{\partial \operatorname{vec}(X)}$ is a Jacobian with dimensions $(n_1n_2\cdots n_N) \times (m_1m_2\cdots m_M).$
- again, the backprop boils down to multiplication of Jacobians:

$$\frac{\partial z}{\partial \operatorname{vec}(X)} = \frac{\partial z}{\partial \operatorname{vec}(Y)} \frac{\partial \operatorname{vec}(Y)}{\partial \operatorname{vec}(X)}$$

Backprop for parameters of a fully connected layer

However, the default recipe is often inefficient due to ignoring the fine-grained structure of the particular layer.

We focus on the parameters of a fully connected layer.

• this is the default formulation:

$$\frac{\partial \mathcal{L}}{\partial \text{vec}(W_k)} = \frac{\partial \mathcal{L}}{\partial h_k} \cdot \frac{\partial h_k}{\partial \text{vec}(W_k)}$$

 \cdot This is the efficient recipe from the lab instructions:

$$\frac{\partial \mathcal{L}}{\partial \mathbf{W}_{k}} = \left(\frac{\partial \mathcal{L}}{\partial w_{kij}}\right)_{D_{k} \times D_{k-1}} = \left[\frac{\partial \mathcal{L}}{\partial \mathbf{h}_{k}}\right]^{\top} \cdot \mathbf{h}_{k-1}^{\top}$$

The following statements are easily shown (homework):

- \cdot the two formulations deliver the same gradients
- their complexities are $O(D_k^2 \cdot D_{k-1})$ and $O(D_k \cdot D_{k-1})$









Fully connected model: backprop (=dynamic programming)



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In order to compute the gradients automatically, we represent a deep model as a computational graph.

The roots of the graph represent inputs, labels, parameters and hyperparameters.

All other nodes represent differentiable functions.

All nodes have only one output (for simplicity):

- this restriction does not reduce the generality
- the output can be a scalar, vector, matrix or tensor.

Example: a classification model with two fully connected layers and L2 regularization

 $\cdot\,$ for the sake of simplicity, we omit the offsets b

Representation of a deep model with computational graph



PyTorch hello world



Main ingredient: reverse-mode automatic differentiation

Homework: solve $x^5 - x^4 - x = -1$ with gradient descent.

Let us go back to our computational graph



The corresponding code in PyTorch

```
import torch, torch.nn.functional as F
```

```
# roots: input, label, parameters, hiperparameter
x = torch.tensor([1.,1.])
y = torch.tensor(0.)
W1 = torch.tensor([[0.5,0], [0,1]], requires_grad=True)
W2 = torch.tensor([1.,0.], requires_grad=True)
lambda1 = torch.tensor(0.01)
```

```
# model
h1 = torch.relu(W1 @ x)
JMLE = F.binary_cross_entropy_with_logits(W2 @ h1, y)
J = JMLE + lambda1 * (W1.pow(2).sum() + W2.pow(2).sum())
```

```
# ask autograd to compute the gradients
J.backward()
print(W1.grad)
```