



Deep Learning

You are here

Outline

Introduction to ML & Supervised ML

Introduction

Regression, Regularization

Classification, Logistic Regression

Bias/Variance trade-off

Supervised ML strikes back

Support Vector Machines
Gaussian Processes

Nearest Neighbors

Ensemble Methods: random forests

Gradient Boosting

Unsupervised ML

Clustering: KMeans, DBScan, GMM, Agglomerative Clustering

Anomaly Detection

Dimensionality Reduction:

- linear: PCA, NMF, ICA
- manifold learning: LLE, IsoMap, t-SNE

Self-Organizing Maps

Deep Learning

D.

Basics of NN: computation graphs

Training a NN: forth- and back-propagation

Optimization Algorithms
Transfer Learning

ResNets

Bayesian NN, Probabilistic BNN

Autoencoders and VAE

Deep Learning, The Revenge

Reinforcement Learning

Convolutional Neural Networks

Inception Module and MobileNet

Generative Adversarial Networks

(hints on) Recurrent Neural Networks

Transformers

Supervised Learning

Data: (x, y)

 $x \rightarrow y$

x is the data, with associated labels y

Goal:

learn a function that maps

"This thing is a dog"

Unsupervised Learning

Data: x there are no labels, only data x

Goal:

learn underlying structure of x



"These two things look alike"

Reinforcement Learning

Data: state-action pairs

Goal:

learn a policy π maximizing future rewards



"Cuddling this thing will make you happy"

Gradient descent is the process of moving towards the minimum of a cost function J, as fast as possible, in order to find the best-fit parameters w and b of your model.

The outline is quite simple:

- 1. start with some (random or not) initial values for w and 6
- 2. keep changing w and b in order to reduce J(w,b)
- 3. until you settle at or near the global minimum (pay attention to local minima)

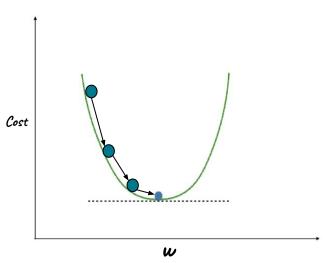
The weights are updated through an iterative process in this way:

$$w:=w-lpha\,\mathrm{d}J(w)/\mathrm{d}w$$

(the same for b)

with α being the **learning rate**, a fundamental hyperparameter especially in Deep Learning. Choosing the best learning rate is fundamental for efficiency and reaching the minimum if α is too small \rightarrow too slow, lots of steps to arrive to minimum if α is too large \rightarrow risk of missing the minimum and jumping over (overshoot)

Typical values for α range between 0.01 and 1.



Recap: batch gradient descent

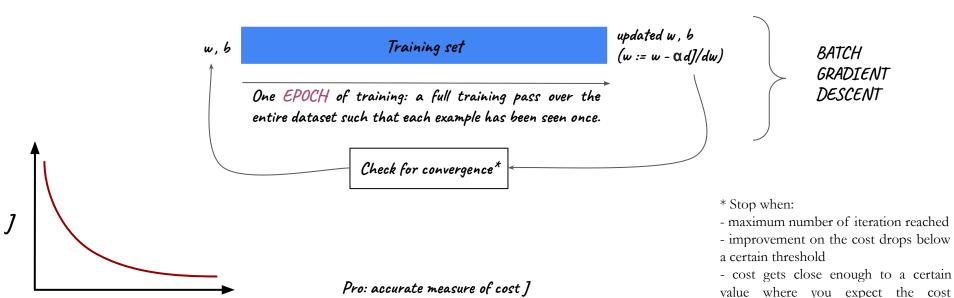
minimum (rare)

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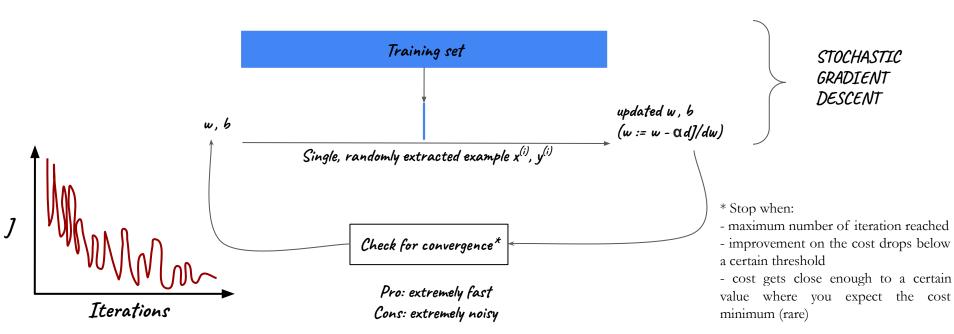
Cons: extremely slow for large datasets

Recap: stochastic gradient descent

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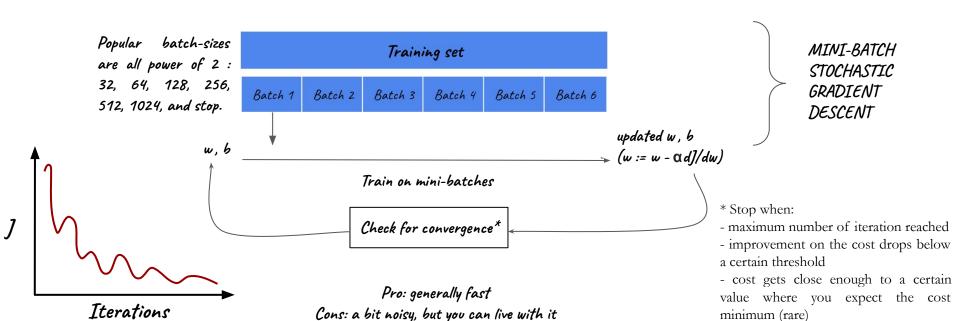


Recap: mini-batch gradient descent

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Galaxy size

Logistic regression is the centerpiece of classification problems. It originates from the sigmoid (or logistic) function:

$$g(z) = \frac{1}{1+e^{-z}}$$

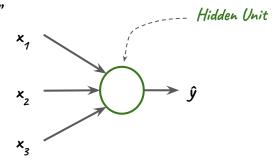
$$g(z)$$

where $f_{w,b}$ gives the probability that κ belongs to a certain class (e.g. if the galaxy is elliptical once given a galaxy size in kpc). With logistic regression you train on the training set, finding the optimal values for the weights w, b. The logistic loss function (*logloss*) evaluated on a single training example is:

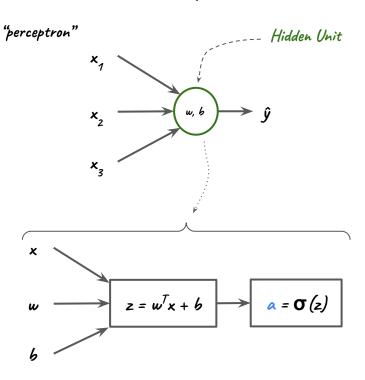
$$\mathcal{L}(f_{ec{w},b}(x^{(i)},y^{(i)})) = -y^{(i)}\log(f_{ec{w},b}(x^{(i)})) - (1-y^{(i)})\log(1-f_{ec{w},b}(x^{(i)}))$$

To understand what it actually does, let's look at a computation graph, with a simple three features input example.

"perceptron"



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The meaning of that circle symbol is:

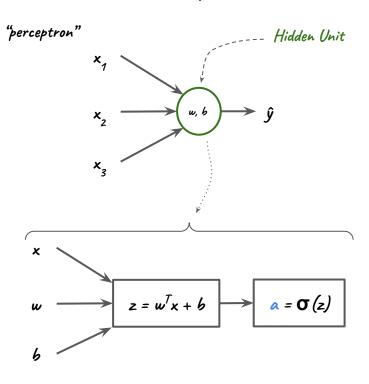
- take in input the features \boldsymbol{x}
- 2) combine the features with the hidden unit weights and bias

$$z = w^T x + b$$

3) compute the α value (activation) applying a function (here $\sigma(z)$)

This is a lone hidden unit, in this case a is equal to the output value ŷ.

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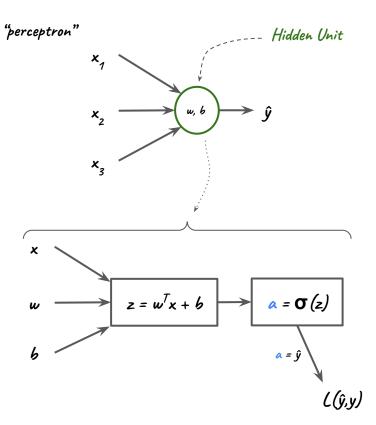
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Let's say that the function that we apply is, e.g., the *sigmoid* function that we saw in the first lesson, and that I reported in the previous slides.

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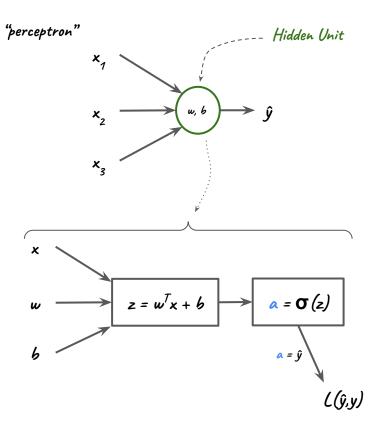
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So, I take the input features, I compute $\alpha = \sigma(w^T x + b)$, that activation value is actually the output \hat{y} , a predicted label, that I can compare with the real value associated to that set of features x, measuring a loss $\ell(\hat{y}, y)$.

A computation graph which is actually logistic regression

The core element of a neural network is the hidden unit (previously known as neuron).

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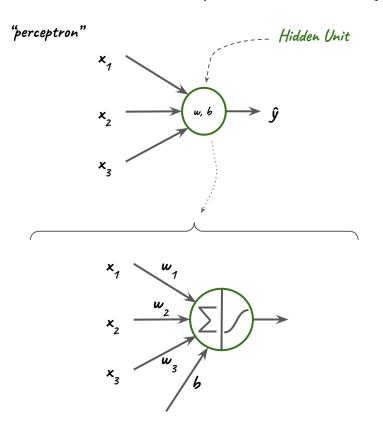
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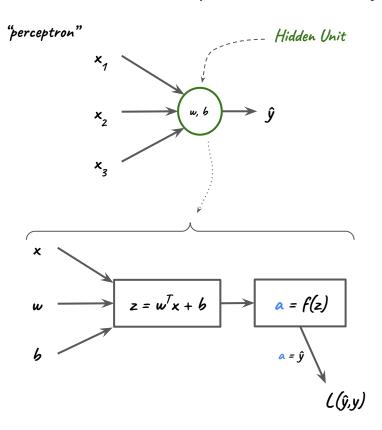
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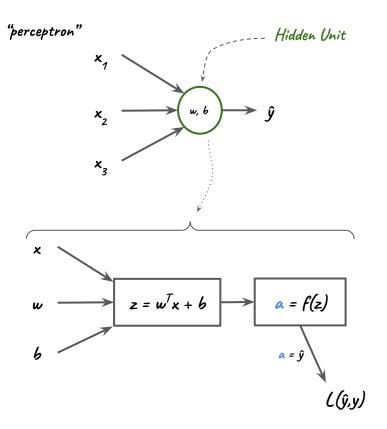
$$y = f(z) = z$$

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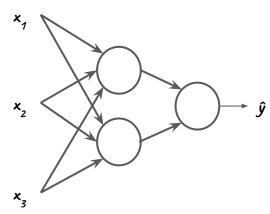
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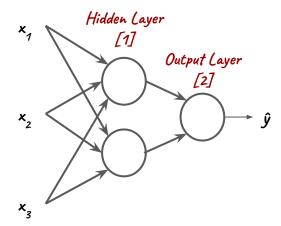
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That is linear regression.



Input Layer



Here we have a two-layers neural network.

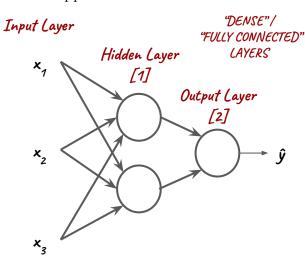
The layer with the input features is called the *input layer*.

The final layer, the one that outputs the prediction \hat{y} , is called the *output layer*.

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Notice how the *input layer* features enters **ALL** the hidden units of the first layer, and how **ALL** the outputs of the first layer enters the output layer.

Everything is "fully-connected" in this Neural Network.



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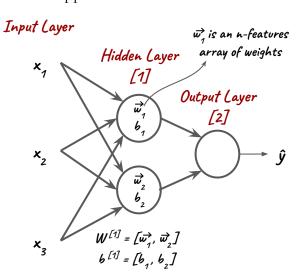
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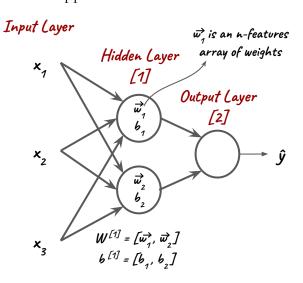
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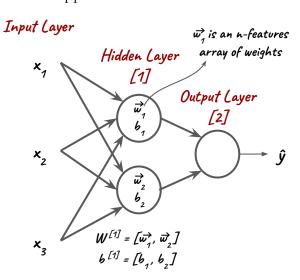
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8 parameters + 3 parameters = 11 parameters



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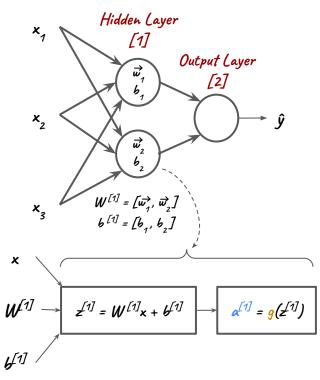
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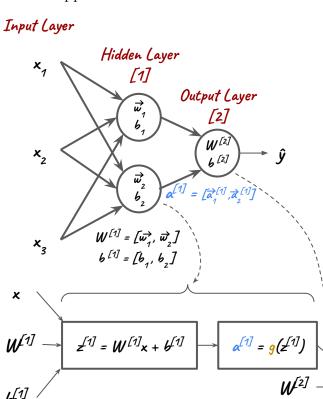
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W2]



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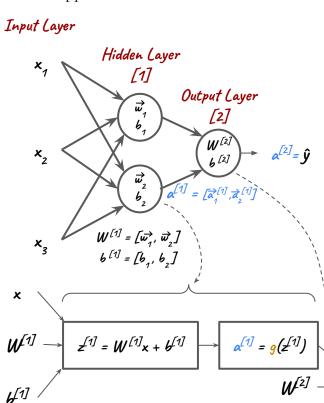
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Then it inputs in the **second (output) layer** the **activation** $a^{[1]}$ calculated in the previous layer, evaluating the second layer activation $a^{[2]}$ with another activation function $g(z^{[2]})$, which in principle could be different from the one used in the first layer.

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 $\alpha^{[2]} = g(z^{[2]})$

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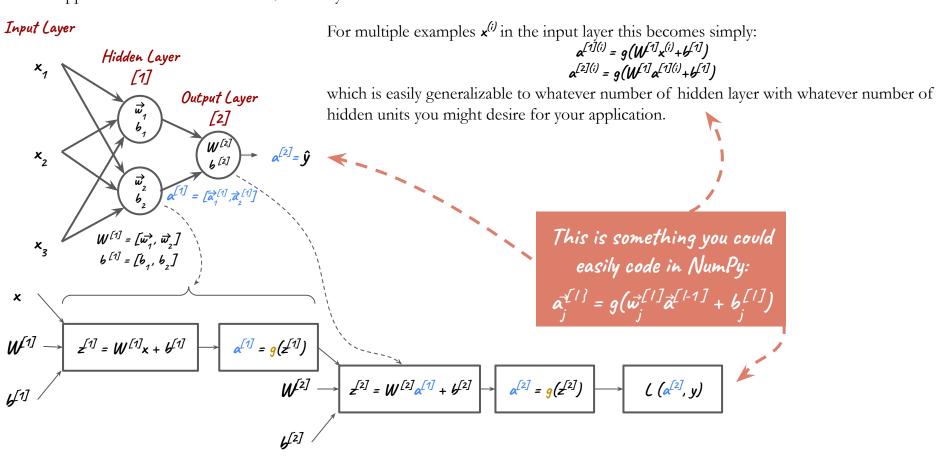
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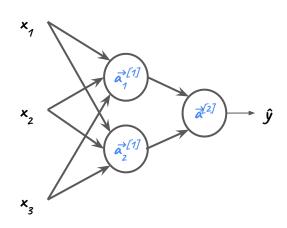
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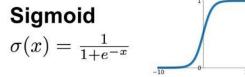
(a[2], y)

This is the output layer, so: $\alpha^{(2)} = \hat{y}$

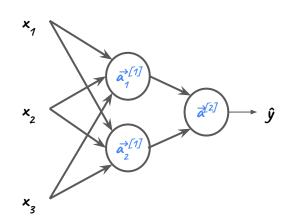
and we can compute a loss function between the predicted and observed labels.

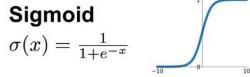




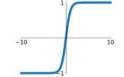


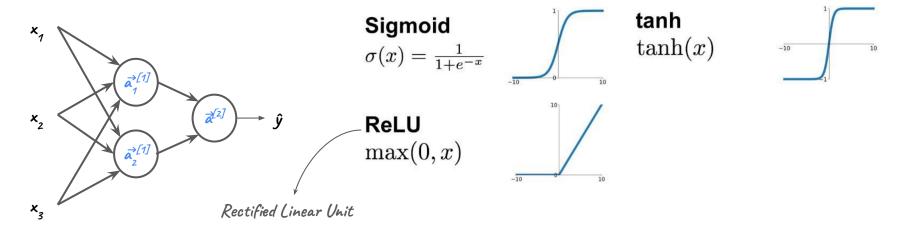
There are plenty to choose, but usually everybody uses the same three or four.

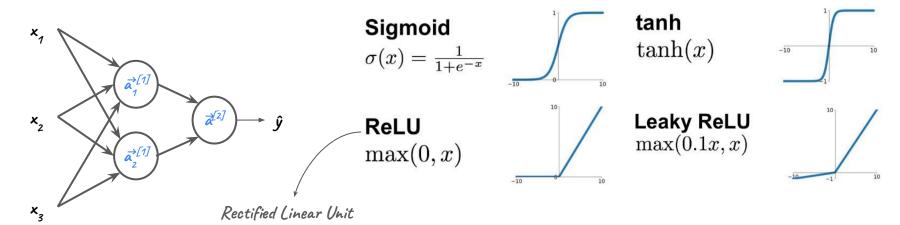


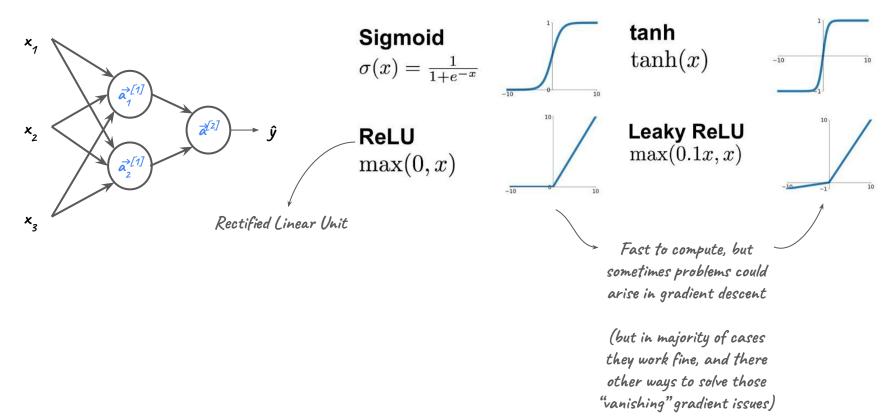


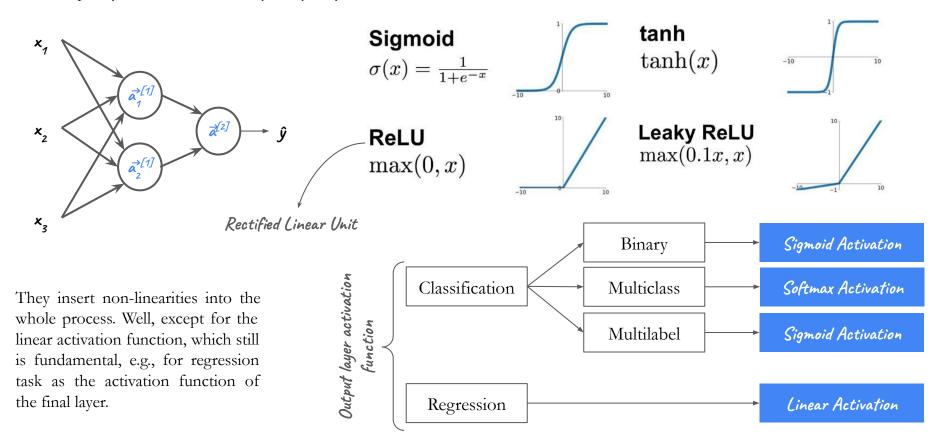
tanh tanh(x)











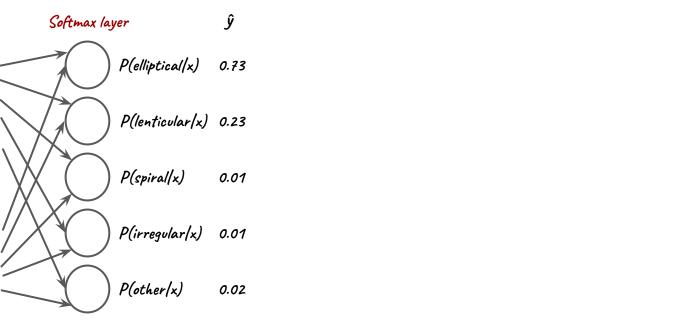
We saw in the first lecture that a possible solution to problems with multiple classes is to use a **one-vs-all** (a.k.a. **one-vs-rest**) approach, or a **one-vs-one** algorithm, where each class probability is evaluated against all the others combined or taken individually.

Neural Networks offers a way out of training #Classes(#Classes-1)/2 models, similar to what **OvA** does, but all hardcoded within the network as the softmax output layer.

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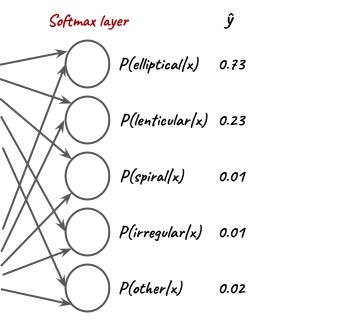
The *softmax layer* is a layer where the number of hidden units is equal to the number of *C classes*; the hidden units in the *softmax* layer will give back the probability that an example belongs to that particular class, so in the De Vaucoulers galaxy classification example:



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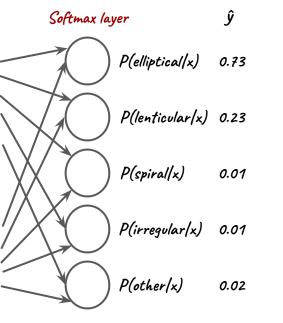
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which naturally outputs values between 0 and $1 \rightarrow \mathbf{probabilities}$

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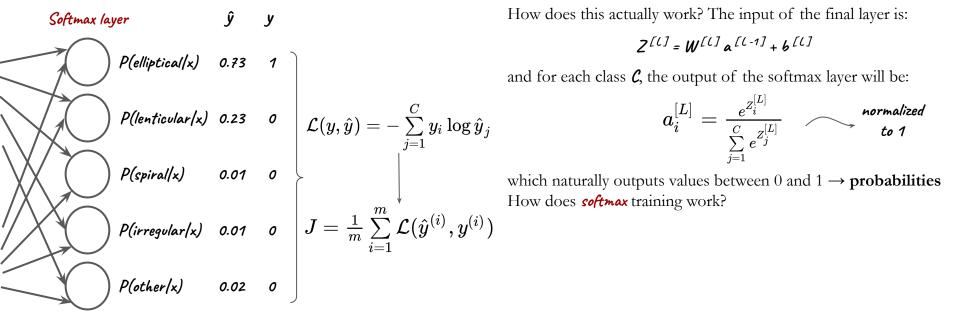
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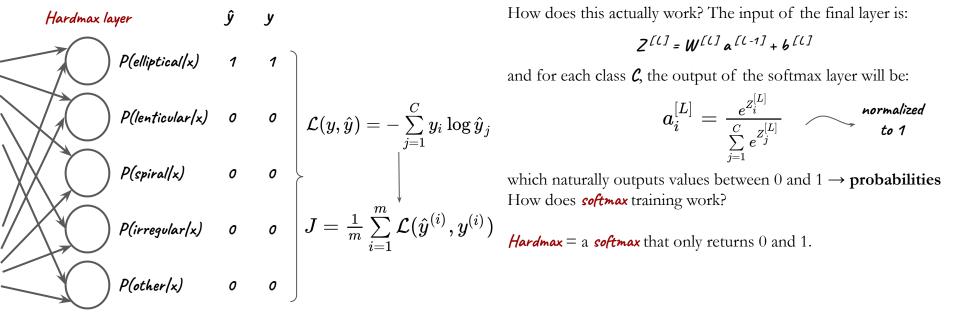
The *softmax layer* is a layer where the number of hidden units is equal to the number of *C classes*, the hidden units in the *softmax* layer will give back the probability that an example belongs to that particular class, so in the De Vaucoulers galaxy classification example:



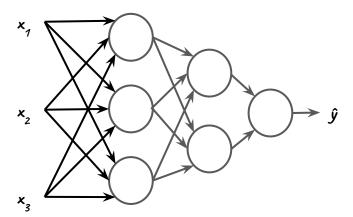
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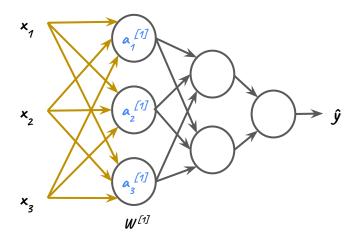
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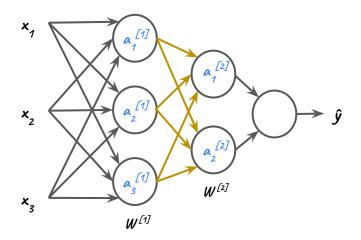
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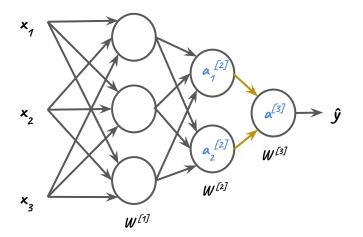
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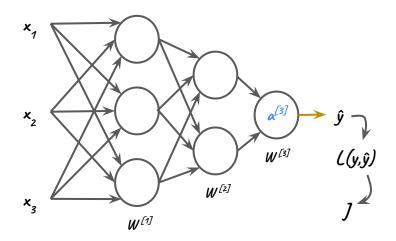


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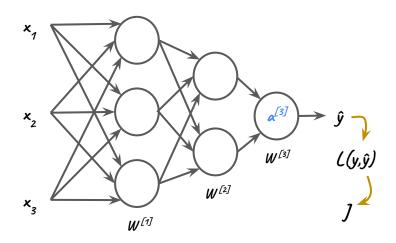
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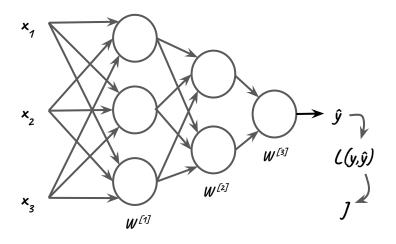
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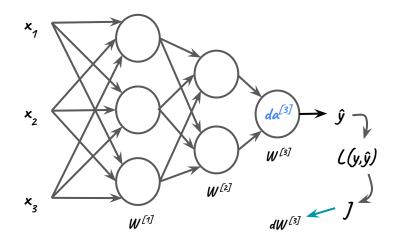
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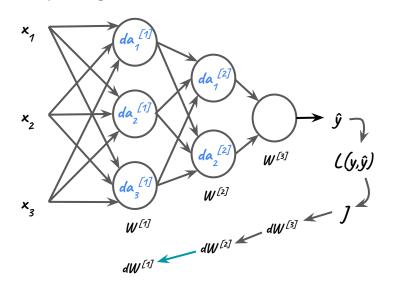
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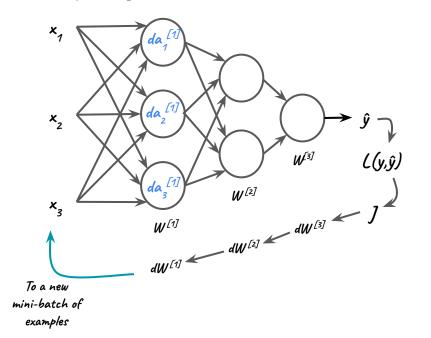
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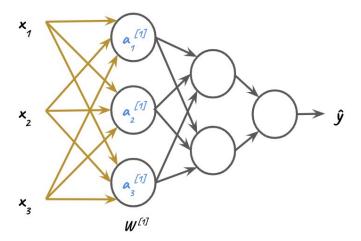
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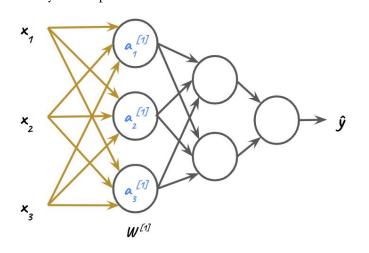
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Is the process of passing through the *hidden layers* backwards, updating each time the weights W^{IJ} of every *hidden layer* with the new values computed during gradient descent by multiplying the gradient of the cost with the learning rate.

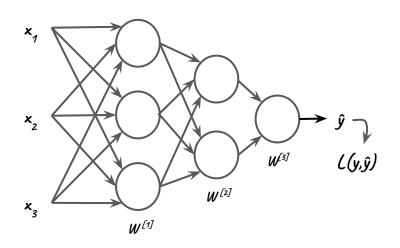
DEEP LEARNING

Perhaps you might wonder why it is called "deep learning".

The reason is that once you start to stack layers on layers, you'll notice how the first layers *learn simple structures* from your data, and the deeper you go with the model, the more *complex things* it is able to learn and reproduce.

As we saw in the first day, it is important to keep a balance between the need to fit to the training data and the need have a model as generalizable as possible, avoiding overfitting. This is achieved by keeping the model as simple as possible with *regularization*.

We saw that *regularization* forces the model weights to be close to (ℓ_2) or exactly zero (ℓ_1) , under the assumption that small values for the model weight \rightarrow simpler model, and it actually works fine.



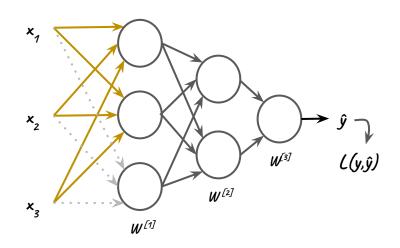
 ℓ_2 and ℓ_1 also apply for Neural Networks, being additional terms to add to the cost function J.

Early stopping, surprisingly, works also fine for Neural Networks, with the necessary caveat that you must always know what you're doing, so don't try this at home.

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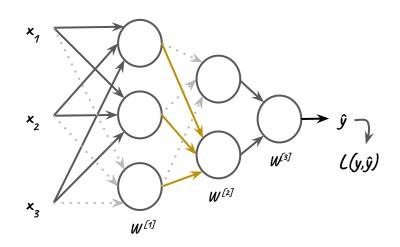
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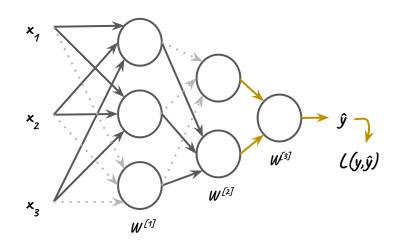
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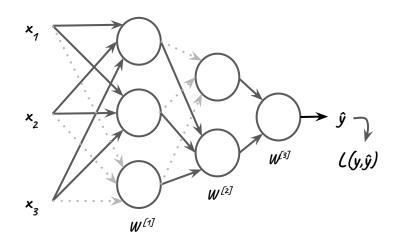
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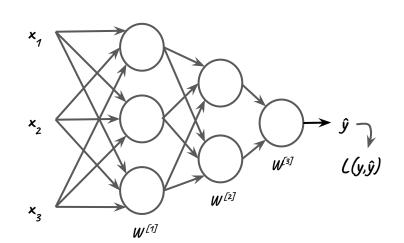
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The concept is: for each training step, do not update ALL the weights in the layers, but turn a random fraction (e.g. 20%) off, and update all the others. This works by forcing the Neural Network to spread the weights over the hidden units, shrinking the weights exactly as L, would do.

Vanishing/Exploding Gradients

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A common problem found when training <u>**DEEP**</u> NN is the one of *vanishing/exploding gradients*, meaning that the gradient might become exponentially high or low, thus halting the whole training process.

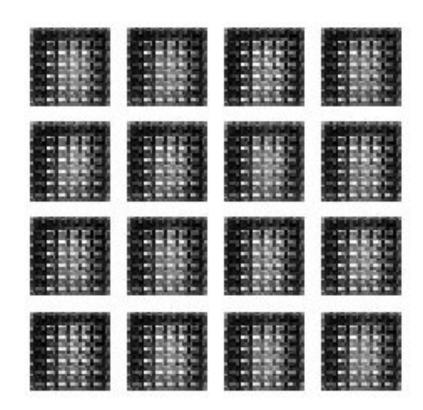
The problem has been recently solved (2014, but the idea dates back to 1961) with *skip connections* and *residual blocks* (*ResNet*, more on later). Another common practice to reduce the chance of *vanishing/exploding gradients* is weight initialization:

$$W^{[I]} = [random\ initialization] * np.sqrt(1/n^{[I-1]}) * Xavier\ initialization" the number of features entering the layer * (np.sqrt(2/n^{[I-1]}) works better with RelU * (np.sqrt(2/(n^{[I-1]}+n^{[I]})) Bengio & co. initialization$$



Tensorflow (by Google) is an open source library to build, train, evaluate, deploy in production, in general work in a ML environment.

```
import tensorflow as tf
mnist = tf.keras.datasets.mnist
(x_train, y_train),(x_test, y_test) = mnist.load_data()
x_{train}, x_{test} = x_{train} / 255.0, x_{test} / 255.0
model = tf.keras.models.Sequential([
  tf.keras.layers.Flatten(input_shape=(28, 28)),
  tf.keras.layers.Dense(128, activation='relu'),
  tf.keras.layers.Dropout(0.2),
  tf.keras.layers.Dense(10, activation='softmax')
])
model.compile(optimizer='adam',
              loss='sparse_categorical_crossentropy',
              metrics=['accuracy'])
model.fit(x_train, y_train, epochs=5)
model.evaluate(x_test, y_test)
```

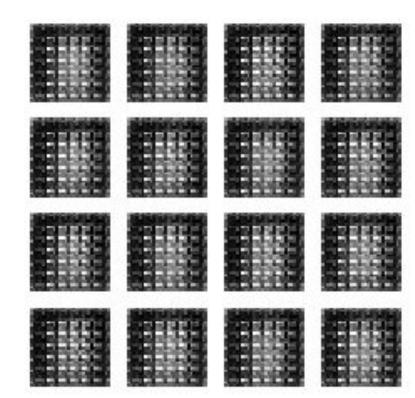






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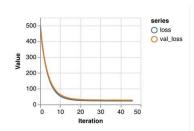


STATUS

Starting training process...

Baseline loss (meanSquaredError) is 85.58

TRAINING PROGRESS



Epoch 47 of 200 completed.

Top 5 weights by magnitude

School drop-out rate -3.9945 Number of rooms per house 2.6450

Distance to commute -2.4197 School class size -1.6939

Distance to highway 1.4261

Train Neural Network Reg hidden layer)

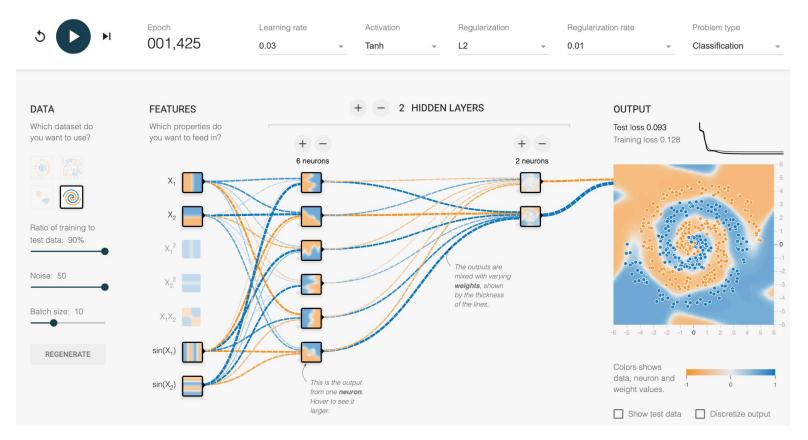
Train Neural Network Regressor (2



Tensorflow Playground						
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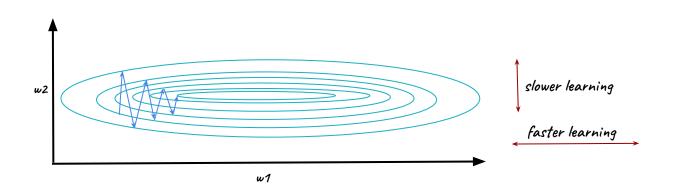


Optimization algorithms

Right now we only saw one possible optimization algorithm, that is the algorithm used to perform gradient descent and reach the minimum of the cost function *J*. It is the simplest possible form of gradient descent:

$$w := w - \alpha \, \mathrm{d}J(w)/\mathrm{d}w$$

which is fine, but as you can imagine there are other possible techniques to perform gradient descent *faster* and *safer* (meaning, avoiding local minima and vanishing/exploding gradients).



Optimization algorithms - Momentum

 β controls the number of past

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Gradient descent with Momentum: evaluates an exponentially weighted average of the gradients, and use that gradient to update the weights This smooths out the steps of gradient descent.

For each training iteration t.

- 1) compute **dw** on current batch/mini-batch
 - 2) compute $V_{dw} = \beta V_{dw} + (1 \beta) dw$
 - 3) update the weights $w := w \alpha V_{dw}$

slower learning

faster learning

https://distill.pub/2017/momentum/, for an overly detailed description of Momentum, with lots of visualizations

the same for b

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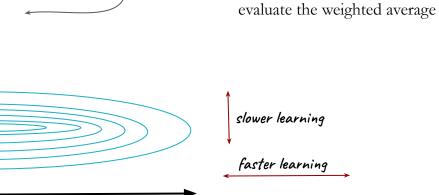
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w2

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the same for b

Two hyperparameters:
- α - β (usually 0.9)

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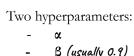
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Root Mean Square prop.: same concept as momentum, you want to speed up learning for certain weights and slow down learning for other weights, but with some small tweaks to the algorithm

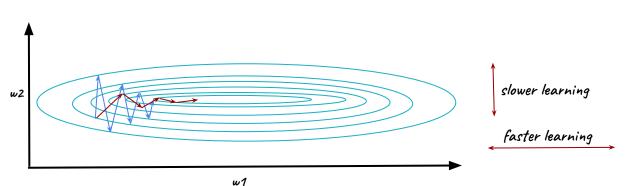
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To avoid dividing per zero, a small η (e.g. 10^{-8}) is added to the code



in this case you can even try a high value for α without risking overshooting



the same for b

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Adam (which stands for Adaptive Moment estimator) is one of those rare case of an algorithm that works well literally everywhere and for every application, and as such you will always see it used as *the* optimization algorithm of a NN. It puts together *Momentum* and *RMSProp*.

Adam starts by initializing $S_{du} = 0$ and $V_{du} = 0$

For each training iteration t.

- compute dw on current mini-batch
 - compute:

$$V_{dw} = \beta_M V_{dw} + (1 - \beta_M) dw$$

$$S_{dw} = \beta_R S_{dw} + (1 - \beta_R) dw^2$$

compute bias correction: 3)

$$V^{corr}_{dw} = V_{dw} / (1 - \beta^t_{M})$$

$$S^{corr}_{dw} = S_{dw} / (1 - \beta^t_{R})$$

update the weights $w := w - \alpha V_{dw}^* (1/\sqrt{S_{dw}})$

$$V_{dw} = \beta_M V_{dw} + (1 - \beta_M) dw$$

$$S_{dw} = \beta_R S_{dw} + (1 - \beta_R) dw^2$$

$$\beta_R \longrightarrow 0.9$$

$$\beta_R \longrightarrow 0.999$$

$$Adam Authors$$

Optimization algorithms - Learning rate decay

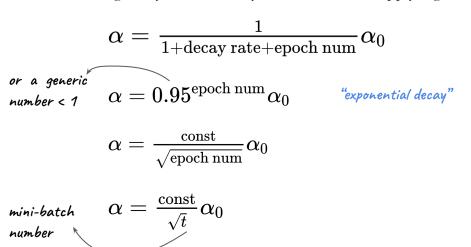
Right now we only saw one possible optimization algorithm, that is the algorithm used to perform gradient descent and reach the minimum of the cost function *J.* It is the simplest possible form of gradient descent:

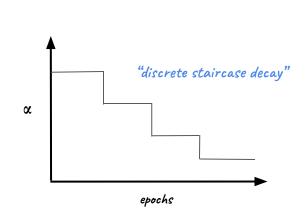
$$w := w - \alpha \, \mathrm{d}J(w)/\mathrm{d}w$$

which is fine, but as you can imagine there are other possible techniques to perform gradient descent *faster* and *safer* (meaning, avoiding local minima and vanishing/exploding gradients).

Learning rate decay means slowing reduce a while training with time. It's not an optimization algorithm on its own, learning rate decay can be easily attached to Momentum, RMSProp or Adam.

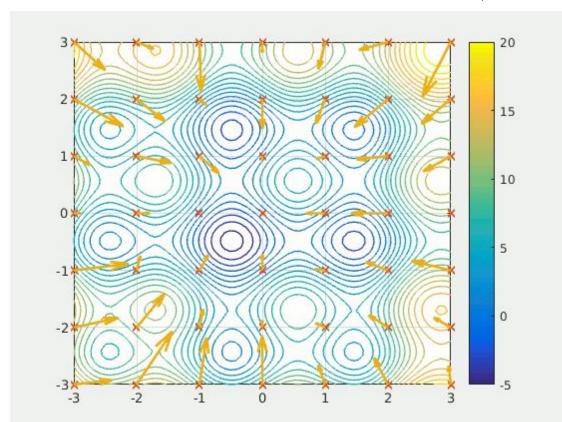
There isn't a single way to do \(\alpha\)-decay, various methods apply, e.g.:





Optimization algorithms - Particle Swarm Optimization

Particle Swarm Optimization is, for once, an idea taken from another field (Animal Social Scie... Studies) that actually applies on this one.



The idea of *PSO* is to emulate the social behaviour of birds and fishes, by initializing a *set* of candidate solutions to search for an optima.

So, it is not just a single particle searching for the minimum of the cost *J*, but a set (*swarm*) of particles.

The idea is similar to that of walkers in MonteCarlo sampling, with particles talking with each other and sharing their knowledges about the parameter space they're sampling.

Particles are scattered around the search-space, and they move around it to find the position of the optima. Their movements are affected by:

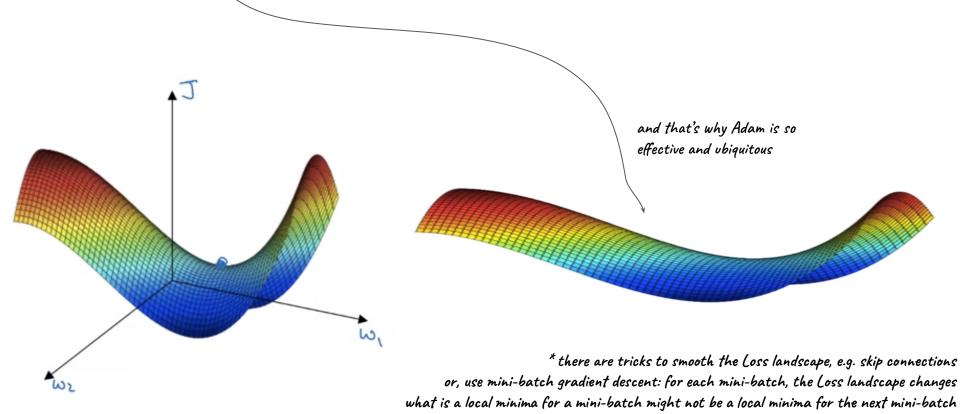
- (1) their cognitive desire to search individually
- (2) the collective action of the group or its neighbors.

Kennedy and Eberhart, "Particle Swarm Optimization," Proceedings of the IEEE International Joint Conference on Neural Networks, 1995, pp. 1942-1948.

Optimization algorithms - Saddle, not minima

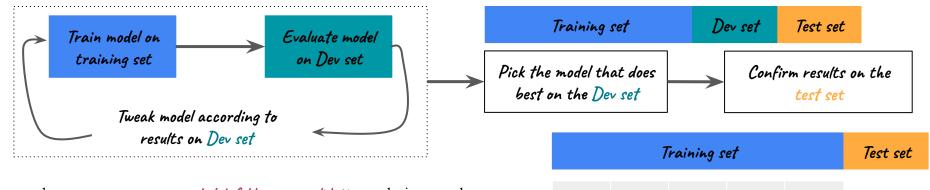
However, keep in mind that in Deep Learning, the thing you should be worried the most is not getting stuck in local minimum, which almost never happen *, but on saddle points.

Especially extremely elongated saddle points, that create a local plateau where your gradient descent might vanish.



Hyperparameter Optimization

As we've already seen in the previous lectures, a typical ML algorithm has at least one major hyperparameter that needs optimal tuning to lead to the best possible performances. That's the reason why the typical workflow in ML splits the sets between train/Dev/test:



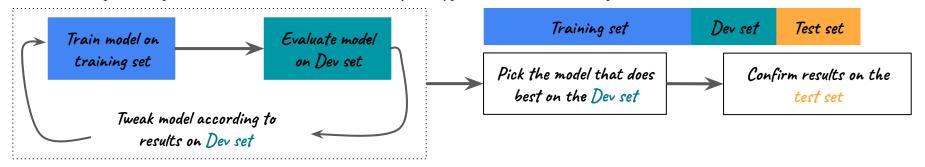
... and you can use, e.g., nested k-folds cross-validation technique to be as most general as possible \rightarrow

Training set					lest sel
Fold 1	Fold 2	Fold 3	Fold 4	Fold 5	
Fold 1	Fold 2	Fold 3	Fold 4	Fold 5	Split 1
Fold 1	Fold 2	Fold 3	Fold 4	Fold 5	Split 2
Fold 1	Fold 2	Fold 3	Fold 4	Fold 5	Split 3
Fold 1	Fold 2	Fold 3	Fold 4	Fold 5	Split 4
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		_			_

Final evaluation ightarrow

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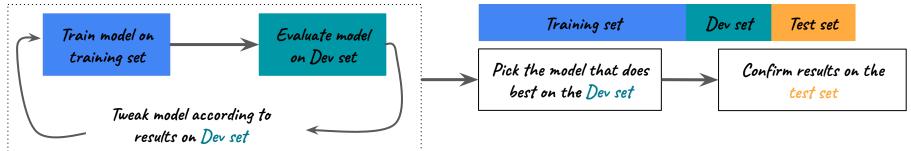


But this does not answer a question: when you have more than one important hyperparameter, how do you *actually* look for the best possible values?

scikit-learn has its own big module for model selection:

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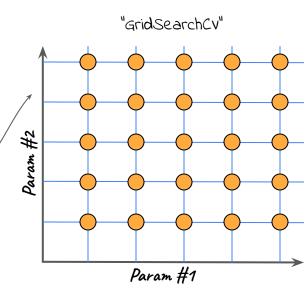


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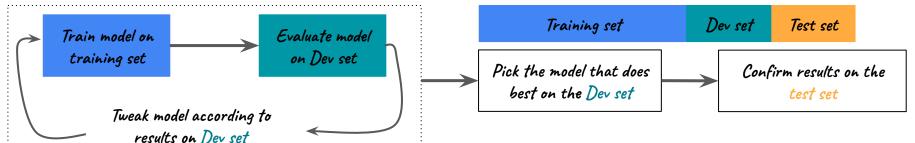
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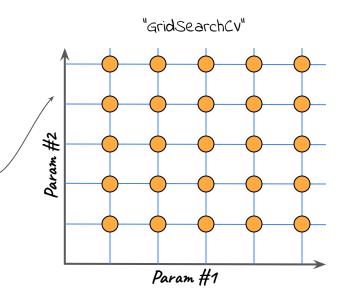
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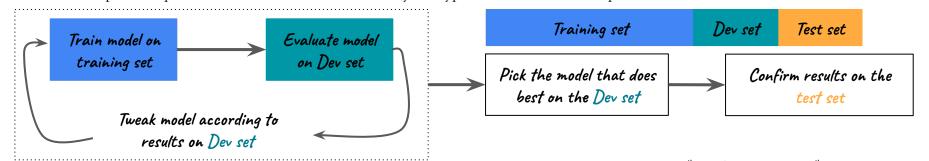
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If the number of hyperparameters is relatively small, GridSearchCV is computationally feasible and actually used.



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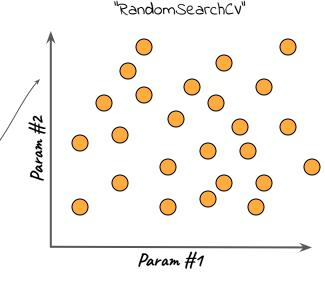


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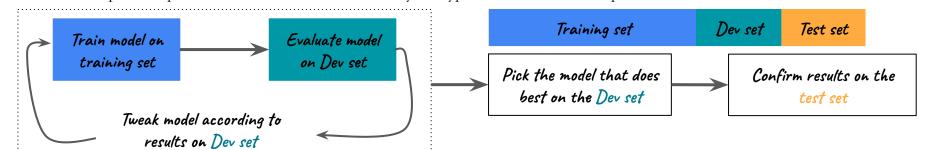
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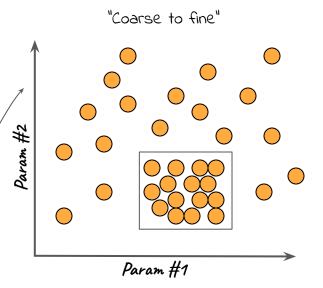
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Batch Normalization

Batch normalization makes the hyperaparameters search easier, and the whole network more robust.

It is an extension of feature scaling, applied to the $\alpha^{[l]}$ values (well, actually to the z's that enter activation functions) in any hidden layer, so as to train the weights $W^{[l]}$, $U^{[l]}$ faster.

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So, for a generic intermediate layer, batch normalization would compute the Z-score with values $z^{(1)} \dots z^{(n)} \to Z^{(i)}_{norm}$, but then instead of feeding the hidden units directly $Z^{(i)}_{norm}$, it gives them a slightly changed version:

$${\hat Z}^{(i)} = \gamma Z_{
m norm}^{(i)} + eta$$

with β and γ learnable parameters of the model, allowing to give the normalized values whatever the mean or range you want it to be (e.g. for sigmoid activation functions).

β parameter

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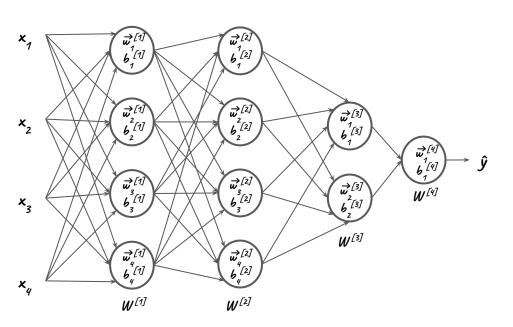
Batch normalization makes the weights more robust, acting in a sense like a regularization scheme, with three great advantages:

- faster training: although each iteration will be slower because of the extra normalization calculation during forth- and back-propagation, it should converge much more quickly
- allows to set a higher learning rate a, thereby increasing the speed of training.

whatever the mean or range you want it to be (e.g. for sigmoid activation functions).

optimal weight initialization: batch normalization reduces the sensitivity to the initial starting weights.

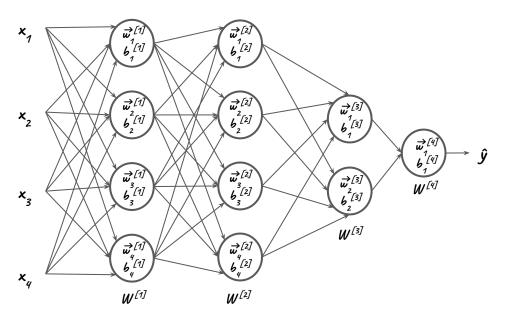
The power of Neural Networks does not only reside in their being low-bias machines that naturally adapts to GPUs with (lots) of libraries and softwares that takes the most out of the technique, but also in subtleties that makes them extremely versatile and able to adapt even to cases when you do not have that much training data in your hands. *Transfer Learning* is one of those.



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The idea is: transfer the knowledge obtained by the training other people made on another Neural Network to your case, where the knowledge, in this case, are the network weights.

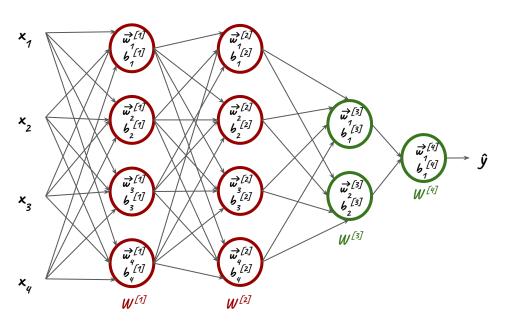
The application is even simpler: take the other model, freeze the weights of the first layers ("pre-training") and train the weights of the final layers (or even just the output layer) on your training set.



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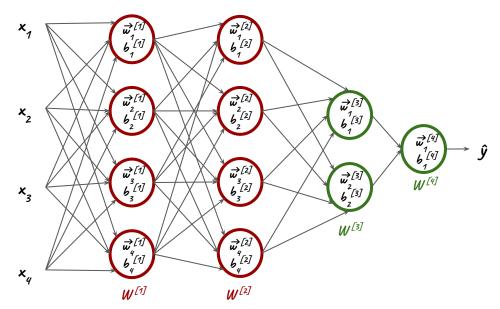
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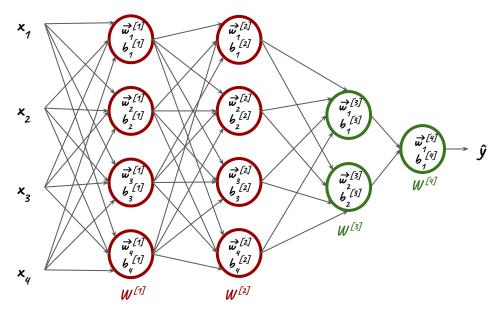
There are multiple reasons to use transfer learning.

Let's suppose you want to train a Convolutional Neural Network to recognize your face in pictures. There's no need to train from scratch a CNN finding billions of pictures of human beings just to recognize your face; instead, you could take the weights of the first layers of a typical CNN doing face-recognition, and train only the final softmax layer to recognize your face out of all the possible people, with just a reduced training set of your pictures.

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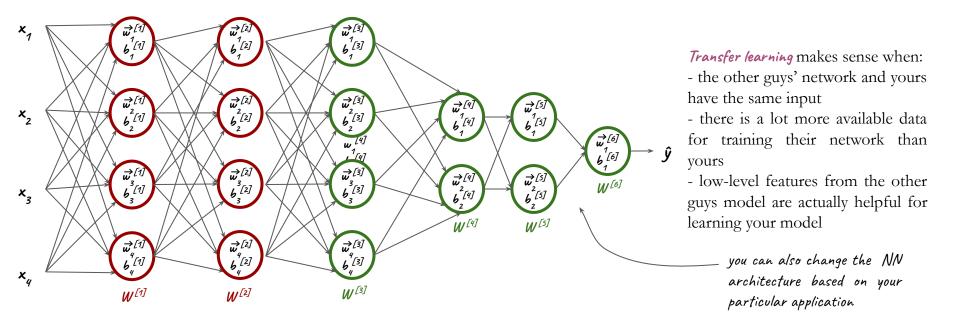
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Translate this into Astrophysics: you do not have to look for billions of pictures of galaxies to do morphological classification on your particular, specific case: there are already lots public models capable of at least recognize a galaxy and its feature out of noise, so take those models, and train the final layers only on your dataset.

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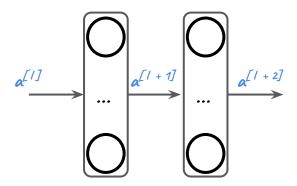
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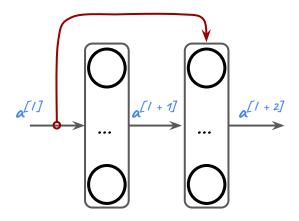
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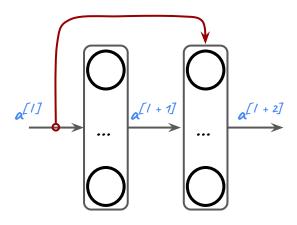
Skip connections take activations from a layer, and inputs it in a deeper layer, as in the example below:

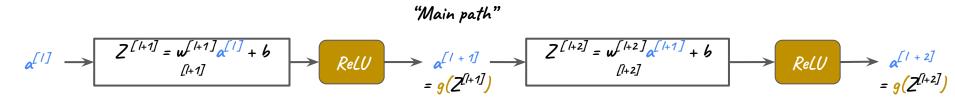


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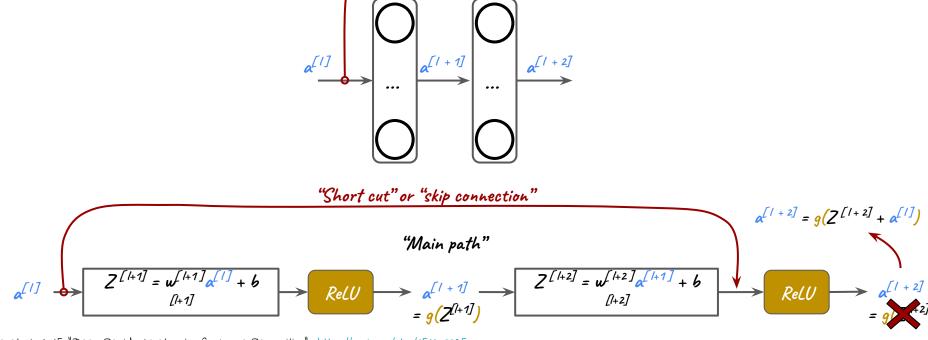




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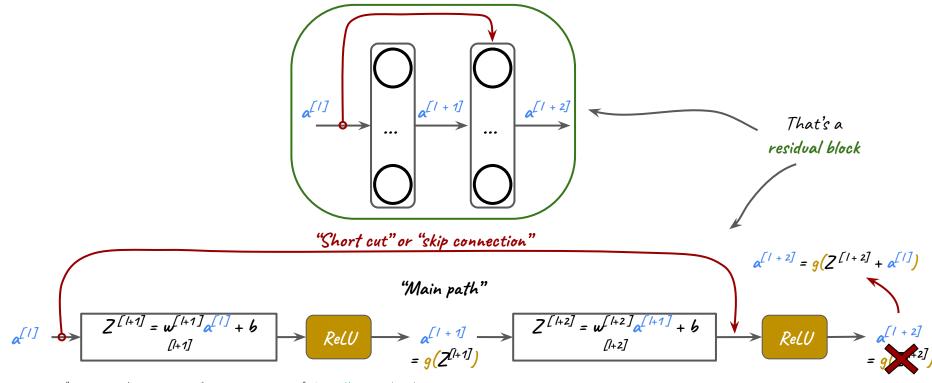
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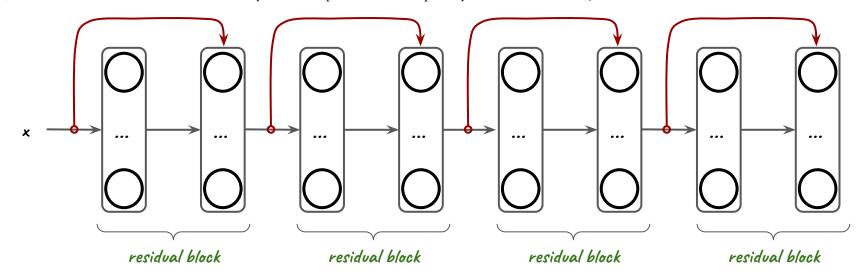
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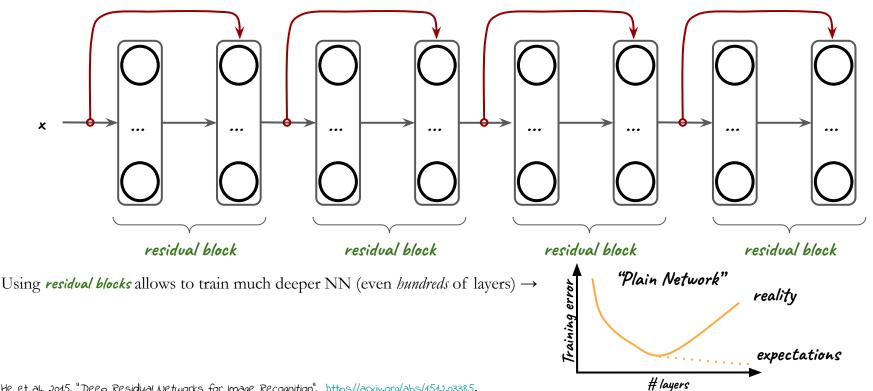
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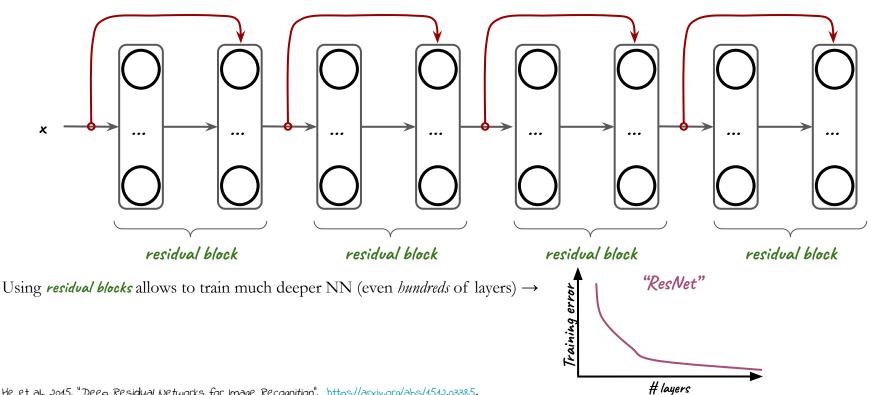
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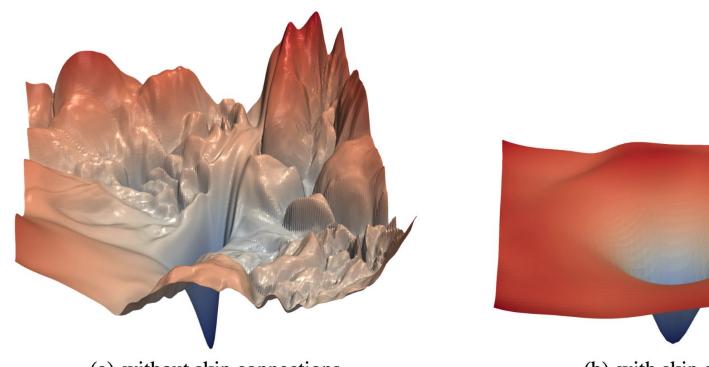
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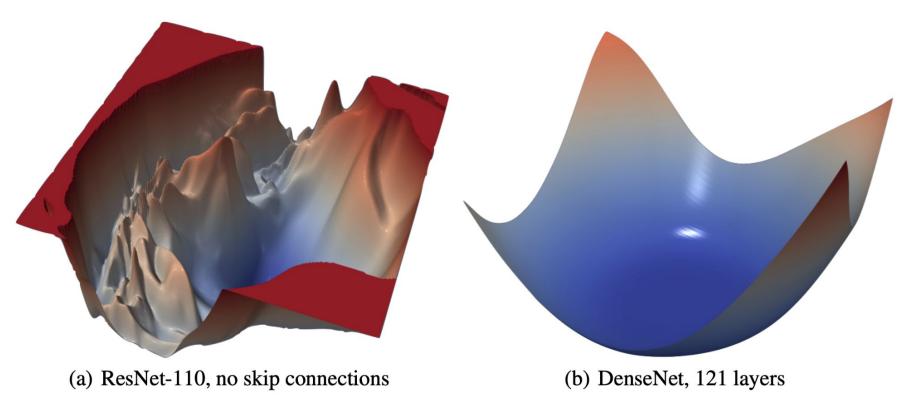


(a) without skip connections

(b) with skip connections

ResNet-56, with or without skip connections

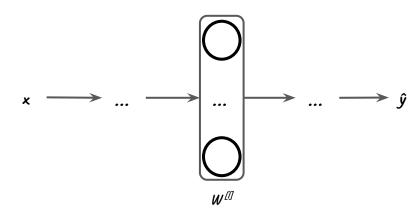
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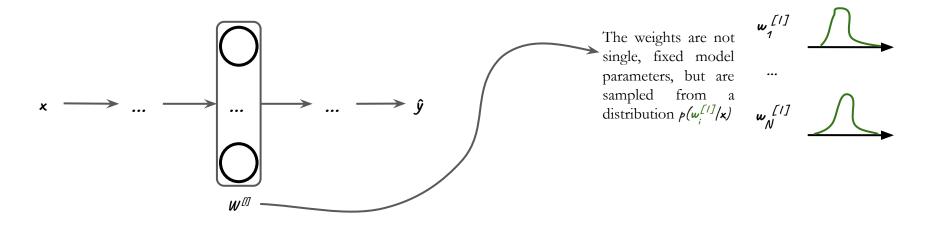
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For each new example you'll get a single *point estimate* prediction, with no information about the uncertainty of the model, nor the prediction \rightarrow enters *Bayesian Neural Networks*: instead of learning a single, specific set of *weights*, try to learn the whole *weight distributions*, encoding the model uncertainty, from which we can sample to produce (still, *a single*) output for a given input.



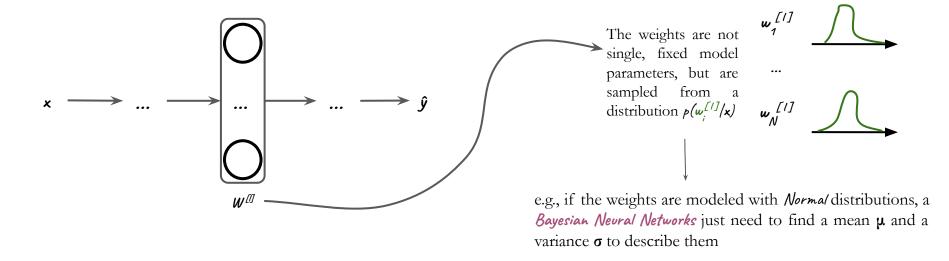
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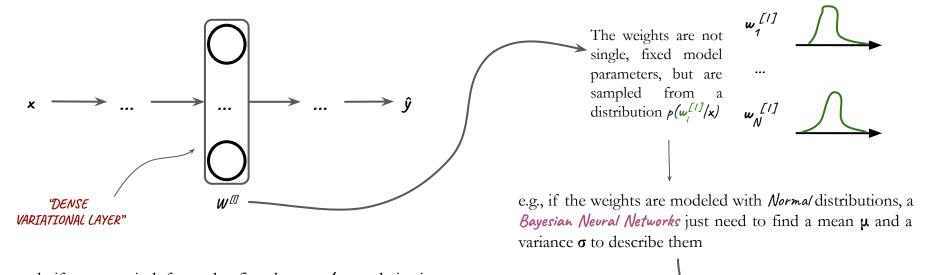
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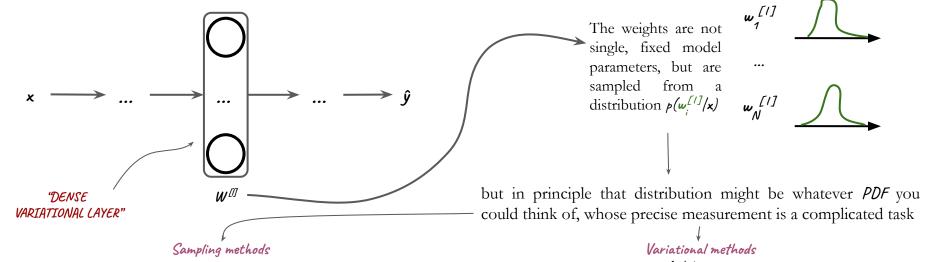


and, if you remind from the first lesson, ℓ_2 regularization forces the weights to be normally distributed with mean zero (and ℓ_4 actually forces some of those to be exactly zero)

https://keras.io/examples/keras_recipes/bayesian_neural_networks/

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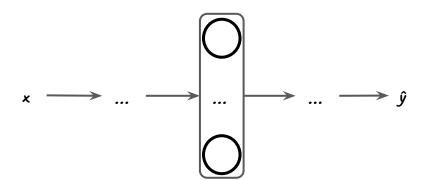
Approximately compute the distribution by generating a finite set of weights $(w_1 ... w_N)$ whose distribution matches p(w/x) in the limit of large N. The challenge is to relatively quickly produce a small number N of network samples that yield a decent approximation of the PDF.

Directly model the posterior p(w/x) using a parametrized distribution q_{ϕ} (w) called the approximate posterior, then iteratively improve the approximation by solving a suitable optimization problem (stochastic variational inference), e.g. minimizing the Kullback-Liebler divergence

Probabilistic Neural Networks

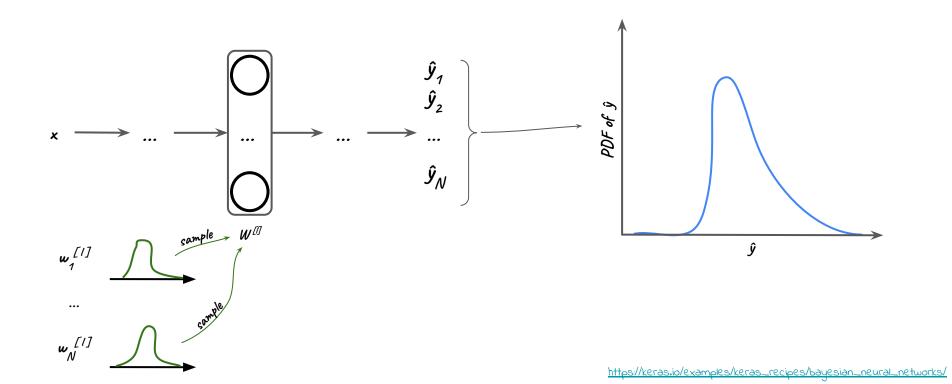
Once you have the weights distributions $\rho(\omega_i^{\lceil l \rceil}/x)$, the next step is to sample from the distribution each time to produce a *distribution* in output, instead of a single point estimate, for the predicted label \hat{y} .

In this way, the model will capture both the *epistemic* (model) and *aleatoric* (data) uncertainties, due to the stochastic process generating the *weights* distributions inevitably influenced by the irreducible noise in the data.

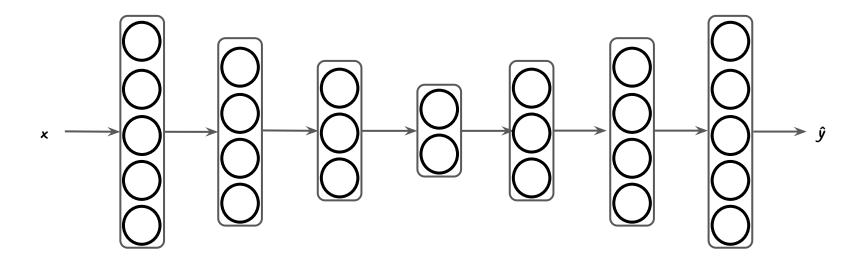


Once you have the weights distributions $\rho(\omega_i^{\lceil l \rceil}/x)$, the next step is to sample from the distribution each time to produce a *distribution* in output, instead of a single point estimate, for the predicted label \hat{y} .

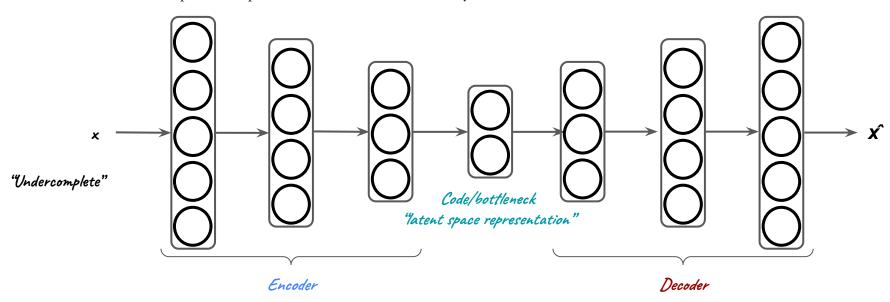
In this way, the model will capture both the *epistemic* (model) and *aleatoric* (data) uncertainties, due to the stochastic process generating the *weights* distributions inevitably influenced by the irreducible noise in the data.



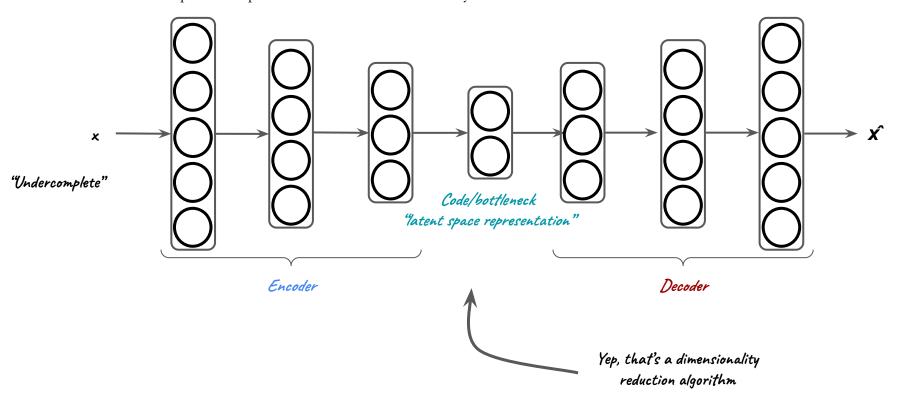
Take a look at this Neural Network. What do you think it is doing?



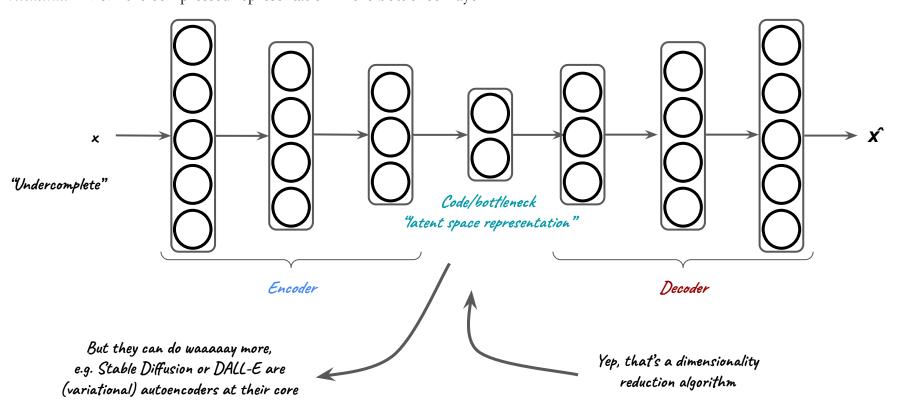
In doing so, it passes through a bottleneck, called the *code* or "latent space representation", which is a compressed representation of the input data. The first part is called the *encoder*, which learns how to compress x into the *code*, and the second part is the *decoder*, that learns how to reconstruct \hat{x} from the compressed representation in the bottleneck layer.



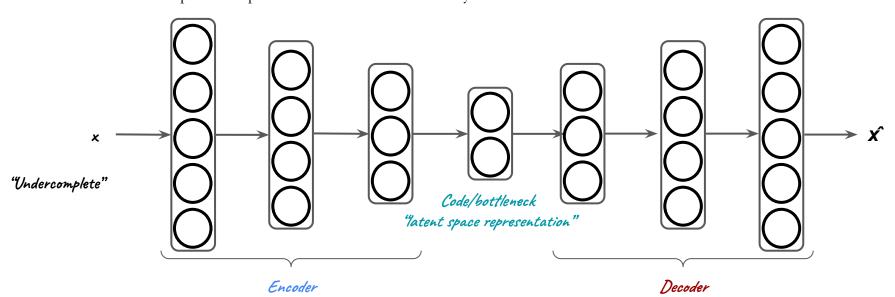
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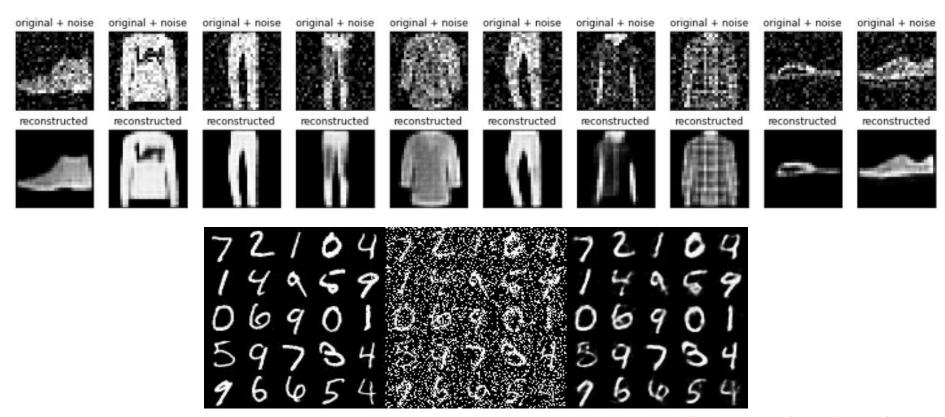


In practice, an autoencoder is trying to learn a function $f_{W,b}(x) = h$ which encodes the data information, and a function $g_{W,b}(h) = g_{W,b}(f_{W,b}(x)) = \hat{X}$ decoding the data from the latent space representation, while minimizing the reconstruction error.

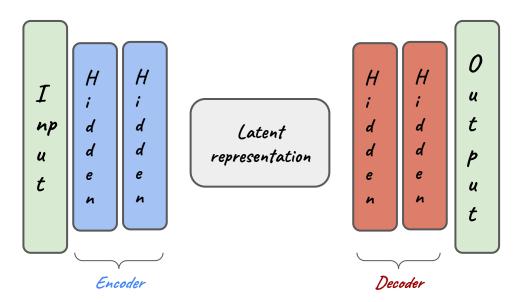
Of course, learning the identity function is useless, so these are designed to be unable to learn to copy perfectly. By placing constraints on the network, such as by limiting the number of hidden units, we can discover interesting structure about the data.

Autoencoders

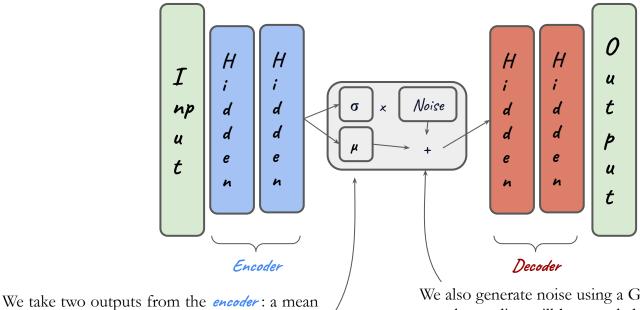
An autoencoder can be used to, e.g., doing dimensionality reduction and for visualization, for anomaly detection, but also for image denoising, or to generate new data that resembles the input data.



In *VAE* the latent representation will be probabilistic, so the later output is partially determined by chance, so *VAE* can create new instances of the data that look and feel like they came from the training set.



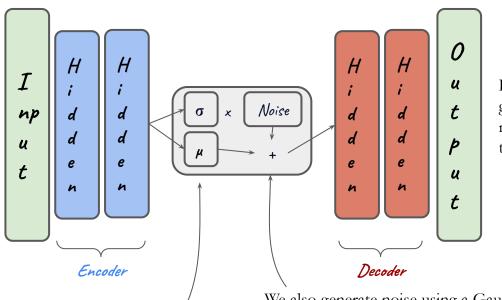
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encoding μ and a variance σ of the encoding

We also generate noise using a Gaussian distribution, so that the actual encoding will be sampled from the distribution using the mean encoding μ and the noise multiplied encoding variance σ .

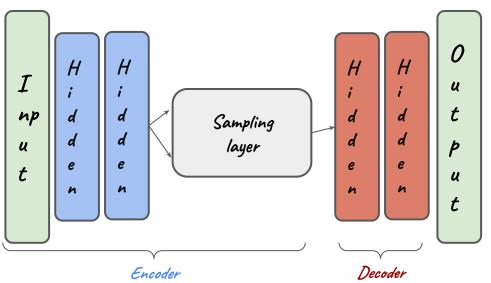
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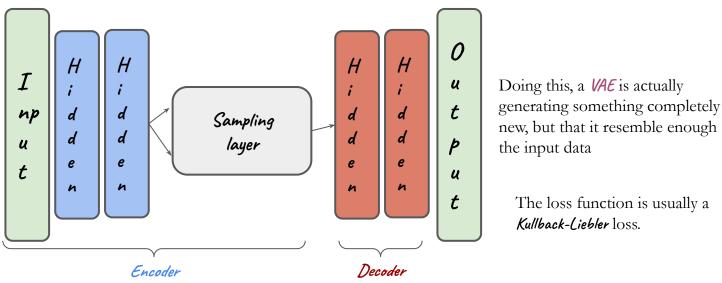


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The loss function is usually a

Kullback-Liebler loss.

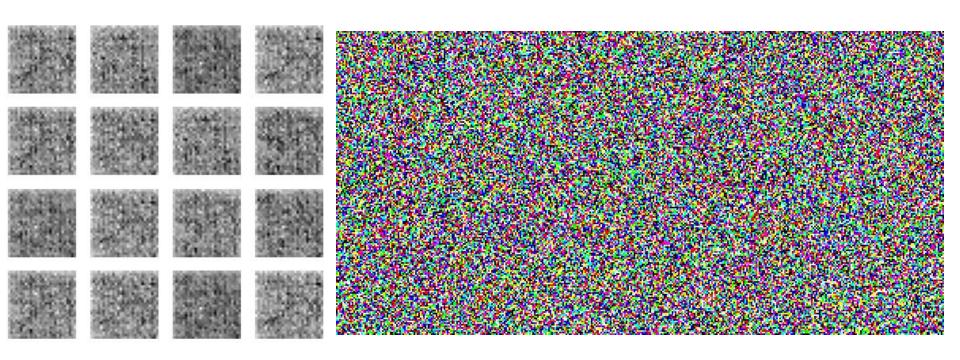
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Variational Autoencoders

Funny thing: until 2013 autoencoders were seen as the cute, didactic algorithm with absolute no practical application in real world. Now *Stable Diffusion* and *DALL-E* work on autoencoders, or better, on variational autoencoders.

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