

Training Neural Networks II

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 February 27, 2023



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unless otherwise stated

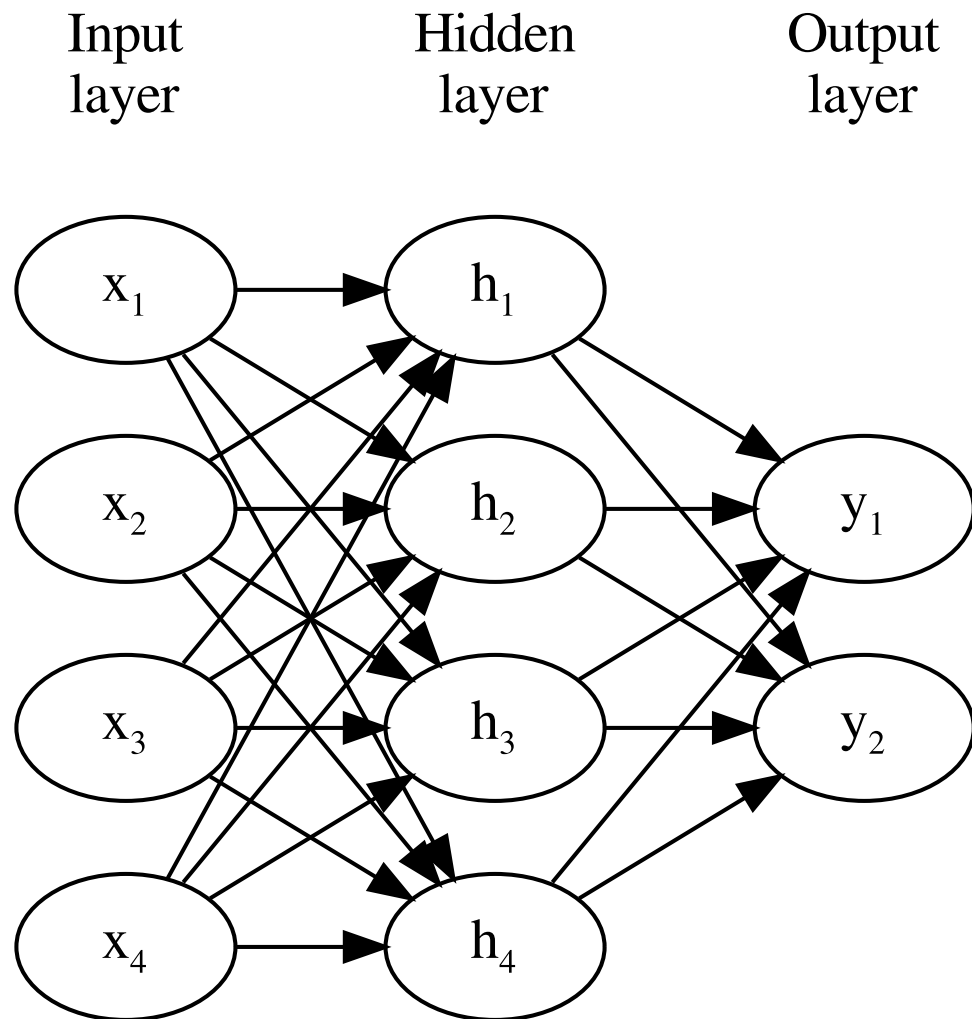
Let us have a dataset with training, validation, and test sets, each containing examples (\mathbf{x}, y) . Depending on y , consider one of the following output activation functions:

$$\begin{cases} \text{none} & \text{if } y \in \mathbb{R} \text{ and we assume variance is constant everywhere,} \\ \sigma & \text{if } y \text{ is a probability of an outcome,} \\ \text{softmax} & \text{if } y \text{ is a gold class index out of } K \text{ classes (or a full distribution).} \end{cases}$$

If $\mathbf{x} \in \mathbb{R}^D$, we can use a neural network with an input layer of size D , some number of hidden layers with nonlinear activations, and an output layer of size O (either 1 or the number of classes K) with the mentioned output function.

BTW, there are of course many functions, which could be used as output activations instead of σ and softmax; however, σ and softmax are almost universally used. One of the reason is that they can be derived using the maximum-entropy principle from a set of conditions, see the [Machine Learning for Greenhorns \(NPFL129\) lecture 5 slides](#). Additionally, they are the inverses of [canonical link functions](#) of the Bernoulli and categorical distributions, respectively.

Putting It All Together – Single-Hidden-Layer MLP



We have

$$h_i = f^{(1)} \left(\sum_j x_j W_{j,i}^{(1)} + b_i^{(1)} \right)$$

where

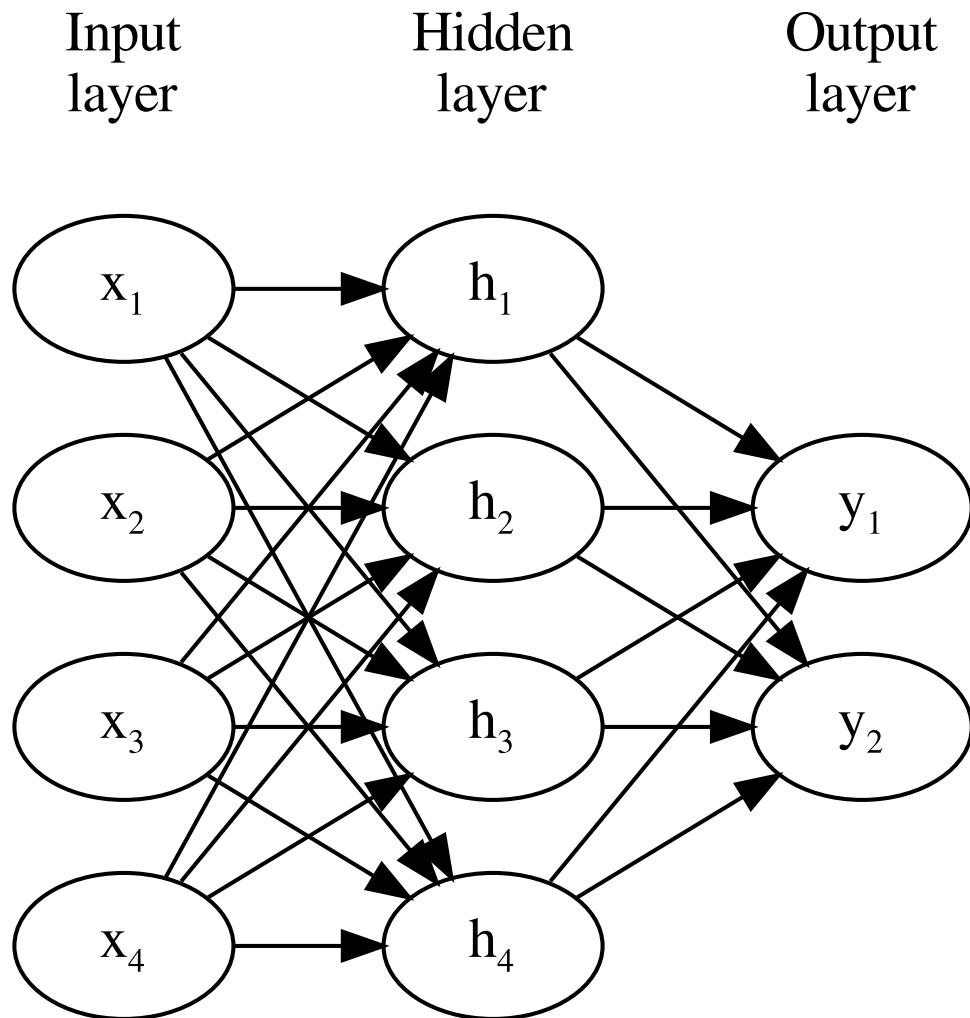
- $\mathbf{W}^{(1)} \in \mathbb{R}^{D \times H}$ is a matrix of **weights**,
- $\mathbf{b}^{(1)} \in \mathbb{R}^H$ is a vector of **biases**,
- $f^{(1)}$ is an activation function.

The weight matrix is also called a **kernel**.

The biases define general behaviour in case of zero/very small input.

Transformations of type $\mathbf{x}^T \mathbf{W}^{(1)} + \mathbf{b}$ are called **affine** instead of *linear*.

Putting It All Together – Single-Hidden-Layer MLP



Similarly

$$o_i = f^{(2)} \left(\sum_j h_j W_{j,i}^{(2)} + b_i^{(2)} \right)$$

with

- $\mathbf{W}^{(2)} \in \mathbb{R}^{H \times O}$ another matrix of weights,
- $\mathbf{b}^{(2)} \in \mathbb{R}^O$ another vector of biases,
- $f^{(2)}$ being an output activation function.

Altogether, the $\mathbf{W}^{(1)}$, $\mathbf{W}^{(2)}$, $\mathbf{b}^{(1)}$, and $\mathbf{b}^{(2)}$ form the **parameters** of the model, which we denote as a vector $\boldsymbol{\theta}$ in the model description and machine learning algorithms.

In our case, the parameters have a total size of $D \times H + H \times O + H + O$.

To train the network, we repeatedly sample m training examples and perform an SGD (or any of its adaptive variants), updating the parameters to minimize the loss derived by MSE

$$E(\boldsymbol{\theta}) = \mathbb{E}_{(\mathbf{x}, y) \sim \hat{p}_{\text{data}}} L(f(\mathbf{x}; \boldsymbol{\theta}), y):$$

$$\theta_i \leftarrow \theta_i - \alpha \frac{\partial E(\boldsymbol{\theta})}{\partial \theta_i}, \text{ or in vector notation, } \boldsymbol{\theta} \leftarrow \boldsymbol{\theta} - \alpha \nabla_{\boldsymbol{\theta}} E(\boldsymbol{\theta}).$$

We set the hyperparameters (size of the hidden layer, hidden layer activation function, learning rate, ...) using performance on the validation set and evaluate generalization error on the test set.

Putting It All Together – Batches

- We always process data in **batches**, i.e., matrices whose rows are the batch examples.
- We represent the network in a vectorized way (tensorized would be more accurate).

Instead of $H_{b,i} = f^{(1)} \left(\sum_j X_{b,j} W_{j,i}^{(1)} + b_i^{(1)} \right)$, we compute

$$\mathbf{H} = f^{(1)} \left(\mathbf{XW}^{(1)} + \mathbf{b}^{(1)} \right),$$

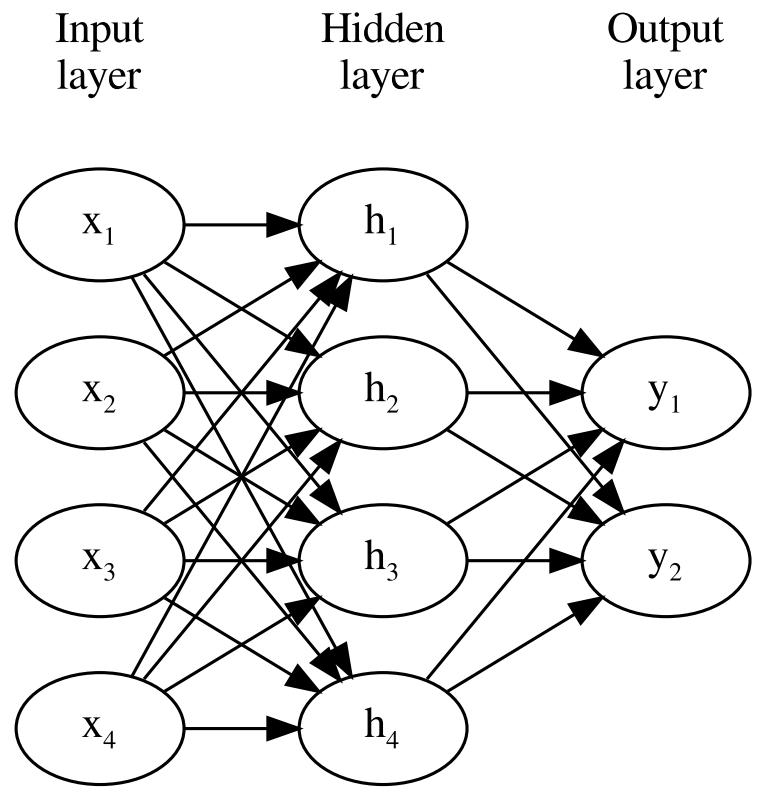
$$\mathbf{O} = f^{(2)} \left(\mathbf{HW}^{(2)} + \mathbf{b}^{(2)} \right) = f^{(2)} \left(f^{(1)} \left(\mathbf{XW}^{(1)} + \mathbf{b}^{(1)} \right) \mathbf{W}^{(2)} + \mathbf{b}^{(2)} \right).$$

The derivatives

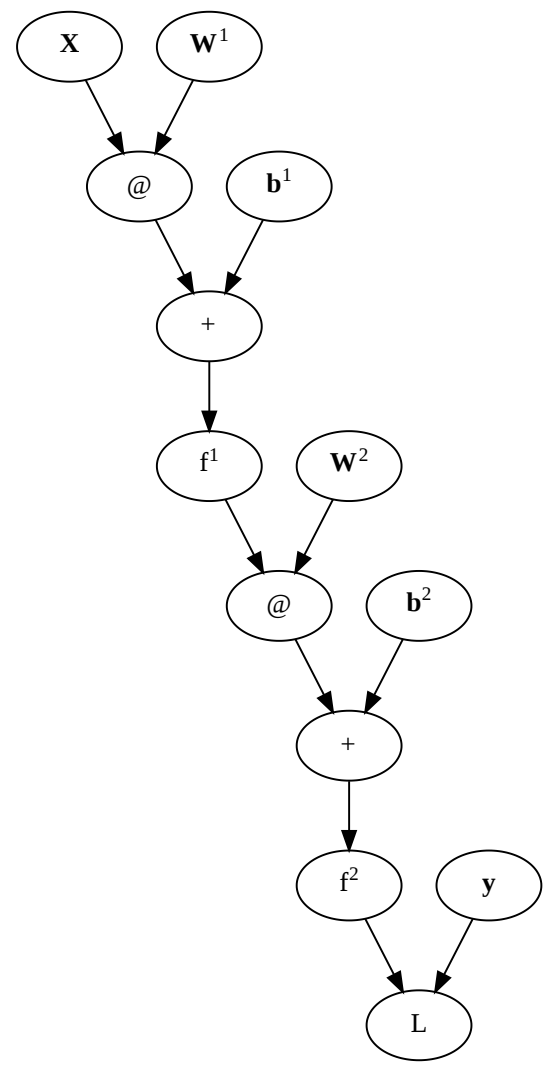
$$\frac{\partial f^{(1)} \left(\mathbf{XW}^{(1)} + \mathbf{b}^{(1)} \right)}{\partial \mathbf{X}}, \frac{\partial f^{(1)} \left(\mathbf{XW}^{(1)} + \mathbf{b}^{(1)} \right)}{\partial \mathbf{W}^{(1)}}, \dots$$

are then batches of matrices (called **Jacobians**) or even higher-dimensional tensors.

Putting It All Together – Computation Graph



→



Designing and training a neural network is not a one-shot action, but instead an iterative procedure.

- When choosing hyperparameters, it is important to verify that the model does not underfit and does not overfit.
- Underfitting can be checked by trying increasing model capacity or training longer, and observing whether the training performance increases.
- Overfitting can be tested by observing train/dev difference, or by trying stronger regularization and observing whether the development performance improves.

Regarding hyperparameters:

- We need to set the number of training epochs so that development performance stops increasing during training (usually later than when the training performance plateaus).
- Generally, we want to use large enough batch size, but such a one which does not slow us down too much (GPUs sometimes allow larger batches without slowing down training). However, because larger batch size implies less noise in the gradient, small batch size sometimes work as regularization (especially for vanilla SGD algorithm).

	Classical ('90s)	Deep Learning
Architecture	:::	::::::::::: CNN, RNN, Transformer, VAE, GAN, ...
Activation func.	\tanh, σ	\tanh , ReLU, LReLU, GELU, Swish (SiLU), SwiGLU, ...
Output function	none, σ	none, σ , softmax
Loss function	MSE	NLL (or cross-entropy or KL-divergence)
Optimization	SGD, momentum	SGD (+ momentum), RMSProp, Adam, SGDW, AdamW, ...
Regularization	L^2, L^1	L^2 , Dropout, Label smoothing, BatchNorm, LayerNorm, MixUp, WeightStandardization, ...

During training and evaluation, we use two kinds of error functions:

- **loss** is a *differentiable* function used during training,
 - NLL, MSE, Huber loss, Hinge, ...
- **metric** is any (and very often non-differentiable) function used during evaluation,
 - any loss, accuracy, F-score, BLEU, ...
 - possibly even human evaluation.

In TensorFlow, the losses and metrics are available in `tf.losses` and `tf.metrics` (aliases for `tf.keras.losses` and `tf.keras.metrics`).

The `tf.losses` offer two sets of APIs. The newer API ones are loss classes like

```
tf.losses.MeanSquaredError(
    reduction=tf.losses.Reduction.AUTO, name='mean_squared_error'
)
```

The created objects are subclasses of `tf.losses.Loss` and can be always called with three arguments:

```
__call__(y_true, y_pred, sample_weight=None)
```

which returns the loss of the given data, *reduced* using the specified reduction. If `sample_weight` is given, it is used to weight (multiply) the individual batch example losses before reduction.

- `tf.losses.Reduction.SUM_OVER_BATCH_SIZE`, which is the default of `.AUTO`;
- `tf.losses.Reduction.SUM`;
- `tf.losses.Reduction.NONE`.

The cross-entropy losses need to specify also the distribution in question:

- `tf.losses.BinaryCrossentropy`: the gold and predicted distributions are Bernoulli distributions (i.e., a single probability);
- `tf.losses.CategoricalCrossentropy`: the gold and predicted distributions are categorical distributions;
- `tf.losses.SparseCategoricalCrossentropy`: a special case, where the gold distribution is one-hot distribution (i.e., a single correct class), which is represented as the gold *class index*; therefore, it has one less dimension than the predicted distribution.

These losses expect probabilities on input, but offer `from_logits` argument, which can be used to indicate that logits are used instead of probabilities.

Old losses API

In addition to the loss objects, `tf.losses` offers methods like `tf.losses.mean_squared_error`, which process two arguments `y_true` and `y_pred` and do not reduce the batch example losses.

There are two important differences between metrics and losses.

1. metrics may be non-differentiable;
2. metrics **aggregate** results over multiple batches.

The metric objects are subclasses of `tf.metrics.Metric` and offer the following methods:

- `update_state(y_true, y_pred, sample_weight=None)` updates the value of the metric and stores it;
- `result()` returns the current value of the metric;
- `reset_states()` clears the stored state of the metric.

The most common pattern is using the provided method

```
__call__(y_true, y_pred, sample_weight=None)
```

which is a combination of `update_state` followed by a `result()`.

Apart from analogues of the losses

- `tf.metrics.MeanSquaredError`
- `tf.metrics.BinaryCrossentropy`
- `tf.metrics.CategoricalCrossentropy`
- `tf.metrics.SparseCategoricalCrossentropy`

the `tf.metrics` module provides

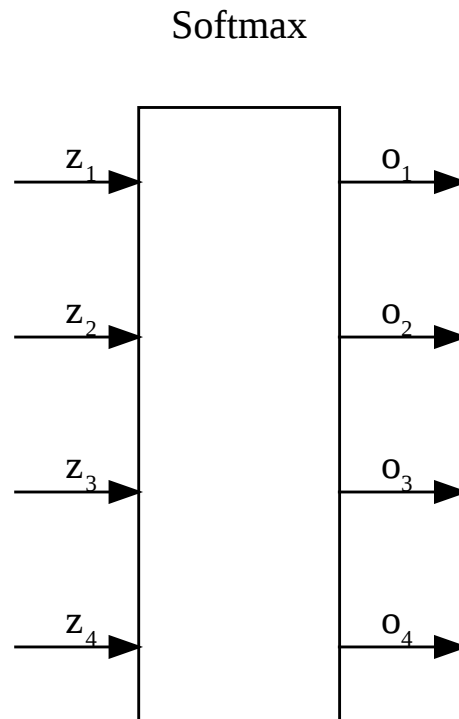
- `tf.metrics.Mean` computing averaged mean;
- `tf.metrics.Accuracy` returning accuracy, which is an average number of examples where the prediction is equal to the gold value;
- `tf.metrics.BinaryAccuracy` returning accuracy of predicting a Bernoulli distribution (the gold value is 0/1, the prediction is a probability);
- `tf.metrics.CategoricalAccuracy` returning accuracy of predicting a Categorical distribution (the argmaxes of gold and predicted distributions are equal);
- `tf.metrics.SparseCategoricalAccuracy` is again a special case of `CategoricalAccuracy`, where the gold distribution is represented as the gold class *index*.

Given the MSE loss of

$$L = (f(\mathbf{x}; \boldsymbol{\theta}) - y)^2,$$

the derivative with respect to the model output is simply:

$$\frac{\partial L}{\partial f(\mathbf{x}; \boldsymbol{\theta})} = 2(f(\mathbf{x}; \boldsymbol{\theta}) - y).$$



Let us have a softmax output layer with

$$o_i = \frac{e^{z_i}}{\sum_j e^{z_j}}.$$

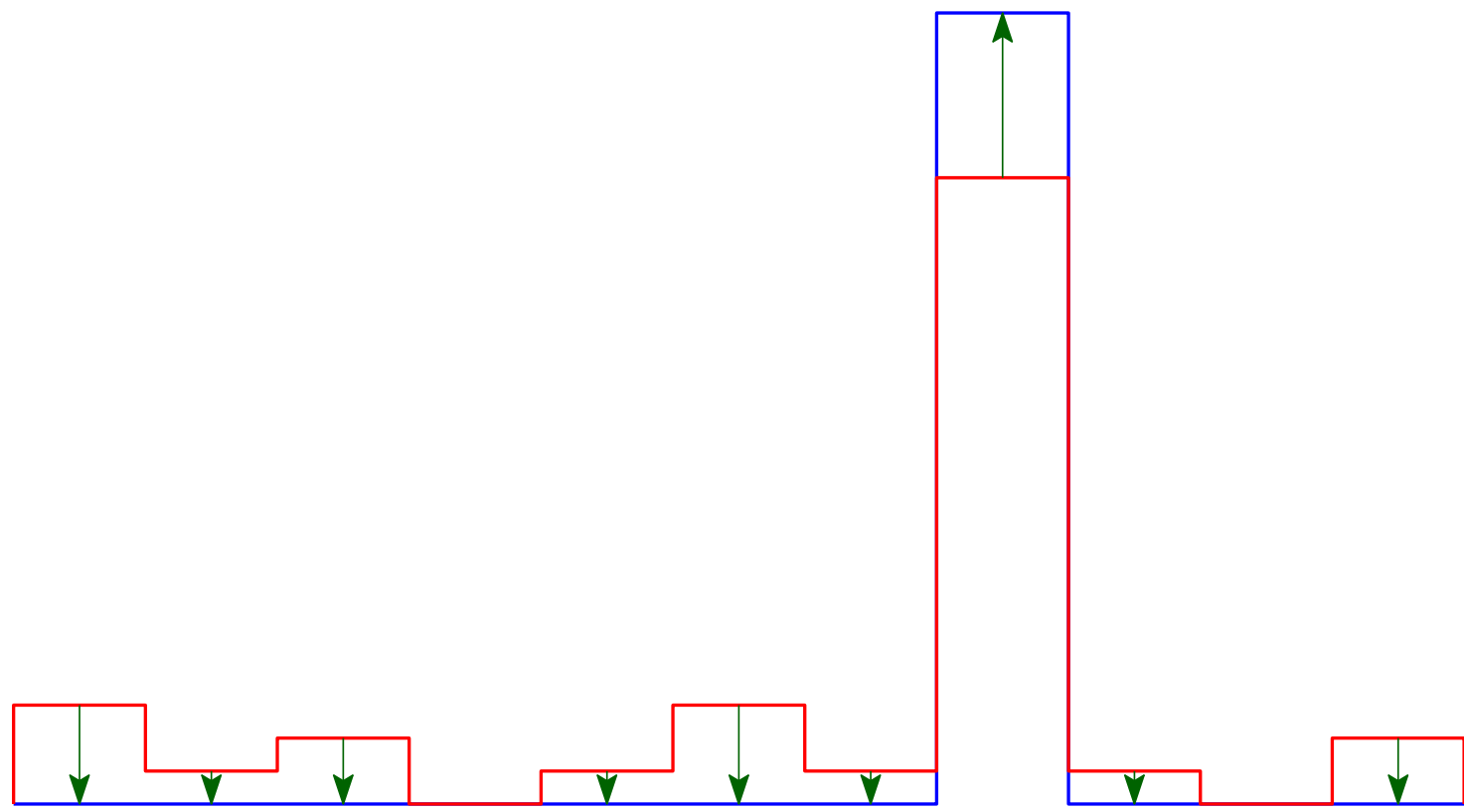
Consider now the MLE estimation. The loss for gold class index *gold* is then

$$L(\text{softmax}(\mathbf{z}), \text{gold}) = -\log o_{\text{gold}}.$$

The derivation of the loss with respect to \mathbf{z} is then

$$\begin{aligned}\frac{\partial L}{\partial z_i} &= \frac{\partial}{\partial z_i} \left[-\log \frac{e^{z_{\text{gold}}}}{\sum_j e^{z_j}} \right] = -\frac{\partial z_{\text{gold}}}{\partial z_i} + \frac{\partial \log(\sum_j e^{z_j})}{\partial z_i} \\ &= -[\text{gold} = i] + \frac{1}{\sum_j e^{z_j}} e^{z_i} \\ &= -[\text{gold} = i] + o_i.\end{aligned}$$

Therefore, $\frac{\partial L}{\partial \mathbf{z}} = \mathbf{o} - \mathbf{1}_{\text{gold}}$, where $\mathbf{1}_{\text{gold}}$ is the one-hot encoding (a vector with 1 at the index *gold* and 0 everywhere else).



Gold distribution

Model distribution

Loss derivative with respect to the softmax inputs.

In the previous case, the gold distribution was *sparse*, with only one target probability being 1. In the case of general gold distribution \mathbf{g} , we have

$$L(\text{softmax}(\mathbf{z}), \mathbf{g}) = - \sum_i g_i \log o_i.$$

Repeating the previous procedure for each target probability, we obtain

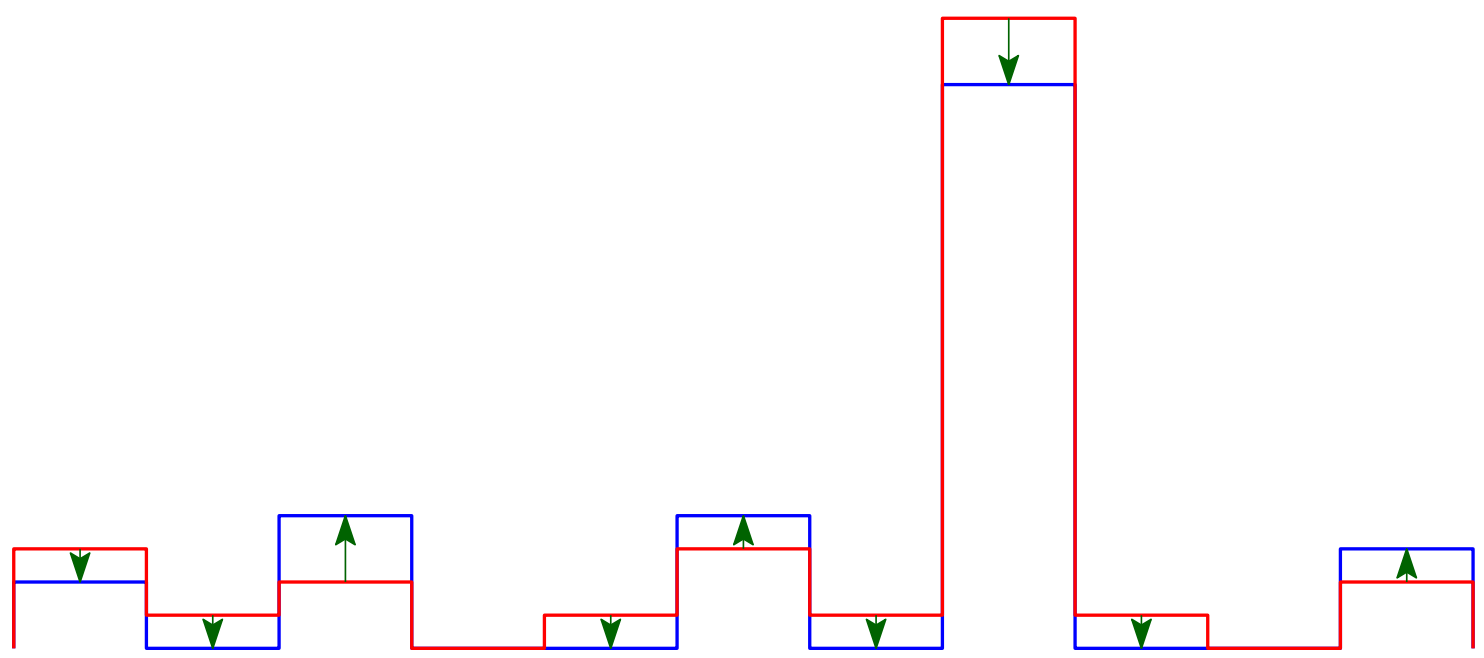
$$\frac{\partial L}{\partial \mathbf{z}} = \mathbf{o} - \mathbf{g}.$$

Sigmoid

Analogously, for $o = \sigma(z)$ we get $\frac{\partial L}{\partial z} = o - g$, where g is the target gold probability.

The result follows automatically from the fact that σ can be computed using softmax as

$$\text{softmax}([0 \ x])_1 = \frac{e^x}{e^x + e^0} = \frac{1}{1 + e^{-x}} = \sigma(x).$$



Gold distribution

Model distribution

Loss derivative with respect to the softmax inputs.

As already mentioned, regularization is any change in the machine learning algorithm that is designed to reduce generalization error but not necessarily its training error.

Regularization is usually needed only if training error and generalization error are different. That is often not the case if we process each training example only once. Generally the more training data, the better generalization performance without any explicit regularization.

- Early stopping
- L^2 , L^1 regularization
- Dataset augmentation
- Ensembling
- Dropout
- Label smoothing

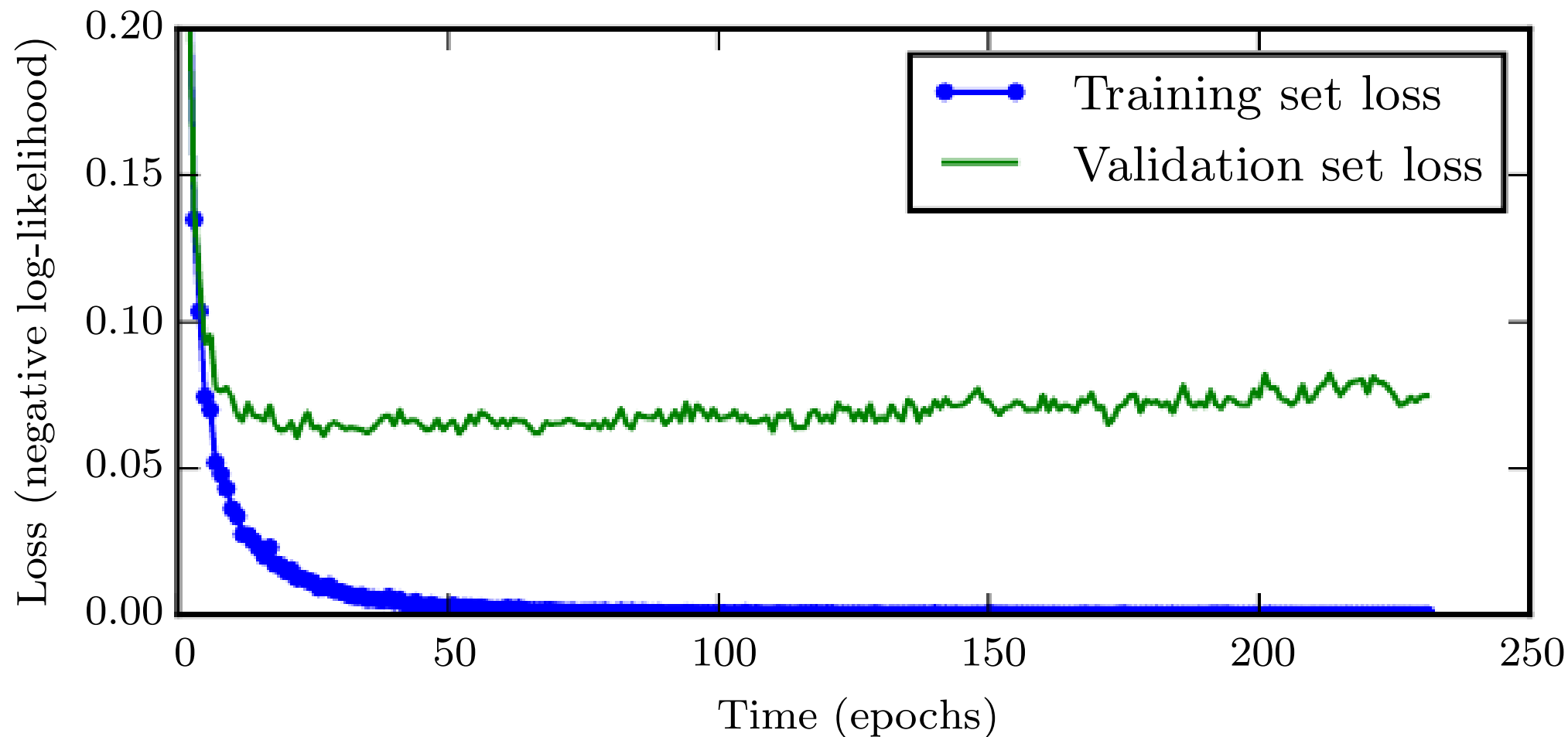


Figure 7.3 of "Deep Learning" book, <https://www.deeplearningbook.org>

L^2 -regularization is one of the oldest regularization techniques, which tries to prefer “simpler” models by endorsing models with **smaller weights**.

Concretely, **L^2 -regularization** (also called **Tikhonov regularization** or **weight decay**) penalizes models with large weights by utilizing the following error function:

$$\tilde{E}(\boldsymbol{\theta}; \mathbb{X}) = E(\boldsymbol{\theta}; \mathbb{X}) + \frac{\lambda}{2} \|\boldsymbol{\theta}\|_2^2$$

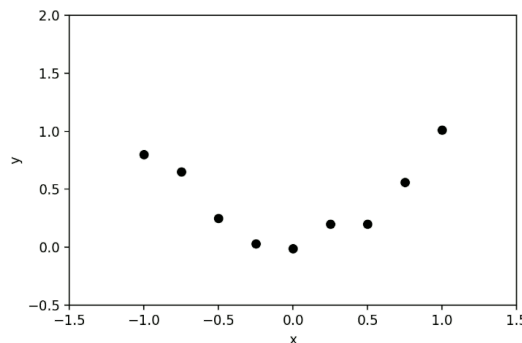
for a suitable (usually very small) λ .

Note that the L^2 -regularization is usually not applied to the *bias*, only to the “proper” weights, because we cannot really overfit via the bias.

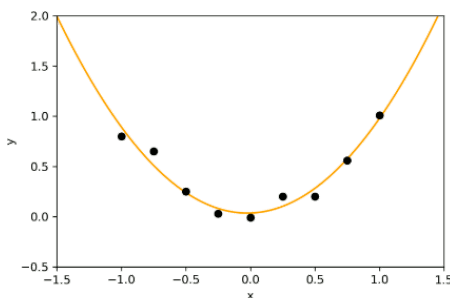
L2 Regularization

One way to look at L^2 -regularization is that it promotes smaller changes of the model (the Jacobian of a single layer with respect to the inputs depends on the weight matrix, because $\frac{\partial x^T \mathbf{W}}{\partial x} = \mathbf{W}$).

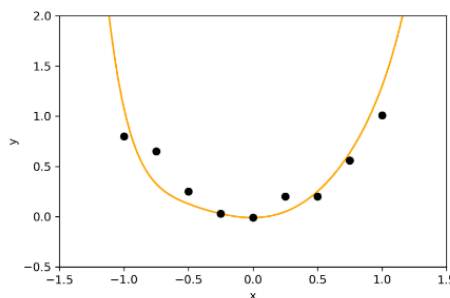
Considering the data points on the right, we present mean squared errors and L^2 norms of the weights for three linear regression models:



https://miro.medium.com/max/2880/1*0-fsK9RkqL3rogo2SnZPCg.png

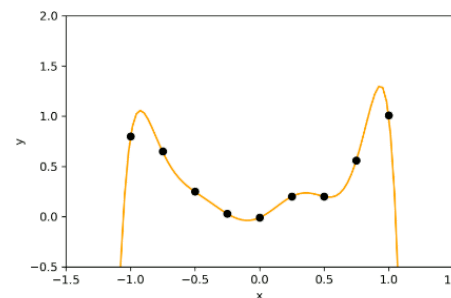


(a) #params = 3
MSE = 0.006
L2 norm = 0.90
L1 norm = 0.98



(b) #params = 9
MSE = 0.035
L2 norm = 1.06
L1 norm = 2.32

https://miro.medium.com/max/2880/1*DVfYChNDMNIS_7CVq2PhSQ.png



(c) #params = 9
MSE = 0
L2 norm = 32.69
L1 norm = 70.03

Figure a: $\hat{y} = 0.04 + 0.04x + 0.9x^2$

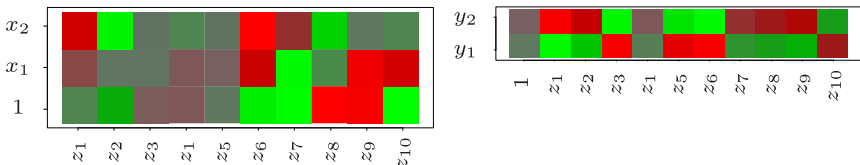
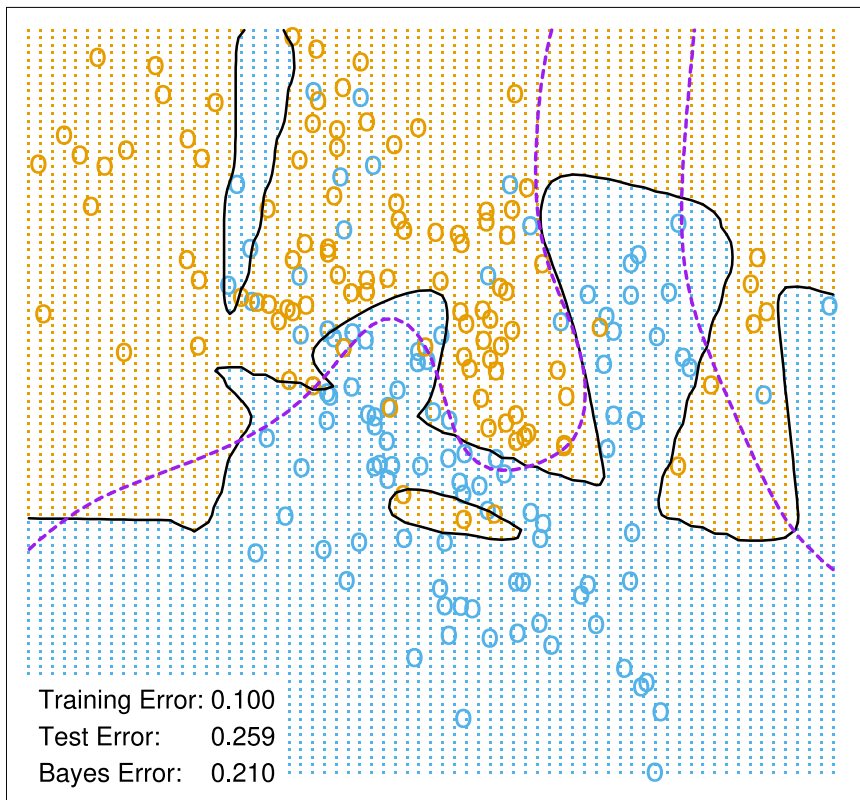
Figure b: $\hat{y} = -0.01 + 0.01x + 0.8x^2 + 0.5x^3 - 0.1x^4 - 0.1x^5 + 0.3x^6 - 0.3x^7 + 0.2x^8$

Figure c: $\hat{y} = -0.01 + 0.57x + 2.67x^2 - 4.08x^3 - 12.25x^4 + 7.41x^5 + 24.87x^6 - 3.79x^7 - 14.38x^8$

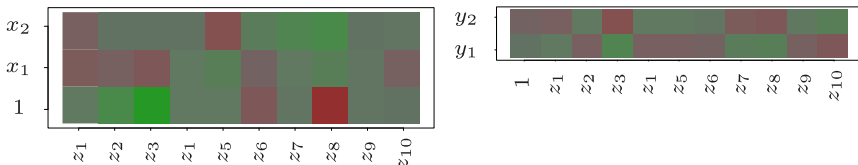
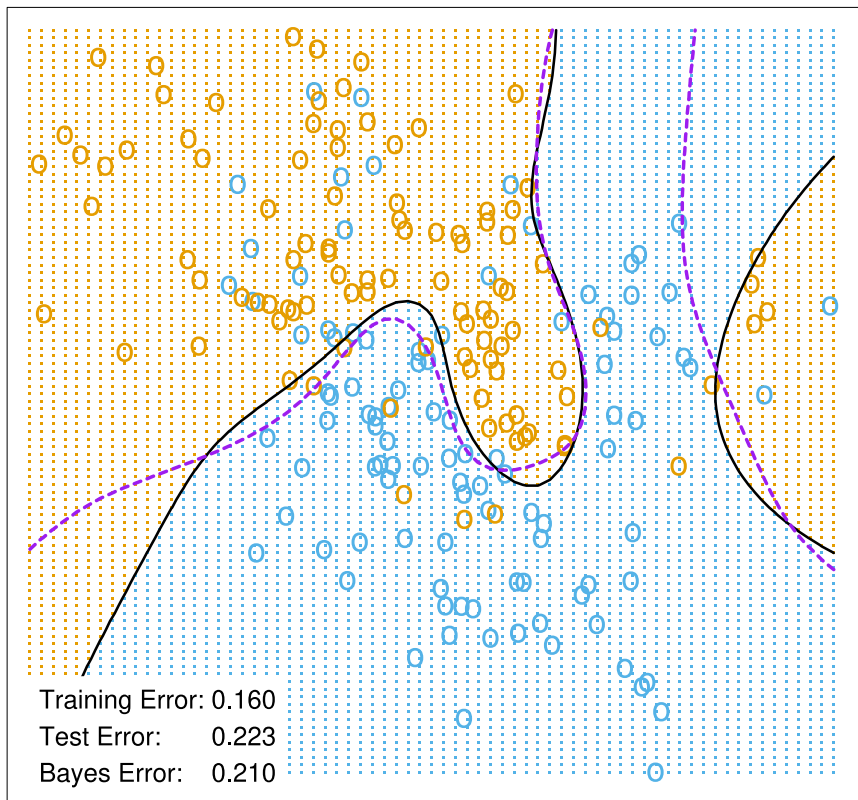
https://miro.medium.com/max/2880/1*UoIRIKXikCz7SFsPfSZrYQ.png

L2 Regularization

Neural Network - 10 Units, No Weight Decay



Neural Network - 10 Units, Weight Decay=0.02



Figures 11.4, 11.5 of "The Elements of Statistical Learning: Data Mining, Inference, and Prediction", <https://hastie.su.domains/ElemStatLearn/>

Another way to arrive at L^2 regularization is to utilize Bayesian inference.

With MLE we have

$$\boldsymbol{\theta}_{\text{MLE}} = \arg \max_{\boldsymbol{\theta}} p(\mathbb{X}; \boldsymbol{\theta}).$$

Instead, we may want to maximize **maximum a posteriori (MAP)** point estimate:

$$\boldsymbol{\theta}_{\text{MAP}} = \arg \max_{\boldsymbol{\theta}} p(\boldsymbol{\theta} | \mathbb{X}).$$

Using Bayes' theorem stating that

$$p(\boldsymbol{\theta} | \mathbb{X}) = \frac{p(\mathbb{X} | \boldsymbol{\theta}) p(\boldsymbol{\theta})}{p(\mathbb{X})},$$

we can rewrite the MAP estimate to

$$\boldsymbol{\theta}_{\text{MAP}} = \arg \max_{\boldsymbol{\theta}} p(\mathbb{X} | \boldsymbol{\theta}) p(\boldsymbol{\theta}).$$

The $p(\boldsymbol{\theta})$ are prior probabilities of the parameter values (our *preference*).

A common choice of the preference is the *small weights preference*, where the mean is assumed to be zero, and the variance is assumed to be σ^2 . Given that we have no further information, we employ the maximum entropy principle, which results in $p(\theta_i) = \mathcal{N}(\theta_i; 0, \sigma^2)$, so that $p(\boldsymbol{\theta}) = \prod_i \mathcal{N}(\theta_i; 0, \sigma^2) = \mathcal{N}(\boldsymbol{\theta}; \mathbf{0}, \sigma^2 \mathbf{I})$. Then

$$\begin{aligned}\boldsymbol{\theta}_{\text{MAP}} &= \arg \max_{\boldsymbol{\theta}} p(\mathbb{X}; \boldsymbol{\theta}) p(\boldsymbol{\theta}) \\ &= \arg \max_{\boldsymbol{\theta}} \prod_{i=1}^m p(\mathbf{x}^{(i)}; \boldsymbol{\theta}) p(\boldsymbol{\theta}) \\ &= \arg \min_{\boldsymbol{\theta}} \sum_{i=1}^m \left(-\log p(\mathbf{x}^{(i)}; \boldsymbol{\theta}) - \log p(\boldsymbol{\theta}) \right).\end{aligned}$$

By substituting the probability of the Gaussian prior, we get

$$\boldsymbol{\theta}_{\text{MAP}} = \arg \min_{\boldsymbol{\theta}} \sum_{i=1}^m \left(-\log p(\mathbf{x}^{(i)}; \boldsymbol{\theta}) + \frac{|\boldsymbol{\theta}|}{2} \log(2\pi\sigma^2) + \frac{\|\boldsymbol{\theta}\|_2^2}{2\sigma^2} \right).$$

The resulting parameter update during SGD with L^2 -regularization is

$$\theta_i \leftarrow \theta_i - \alpha \frac{\partial E}{\partial \theta_i} - \alpha \lambda \theta_i, \text{ or in vector notation, } \boldsymbol{\theta} \leftarrow \boldsymbol{\theta} - \alpha \nabla_{\boldsymbol{\theta}} E(\boldsymbol{\theta}) - \alpha \lambda \boldsymbol{\theta}.$$

This update can be rewritten to

$$\theta_i \leftarrow \theta_i (1 - \alpha \lambda) - \alpha \frac{\partial E}{\partial \theta_i}, \text{ or in vector notation, } \boldsymbol{\theta} \leftarrow \boldsymbol{\theta} (1 - \alpha \lambda) - \alpha \nabla_{\boldsymbol{\theta}} E(\boldsymbol{\theta}).$$

Terminologically, the update of weights in these two formulas is called *weight decay*, because the weights are multiplied by a factor $1 - \alpha \lambda < 1$, while adding the L^2 -norm of the parameters to the loss is called *L^2 -regularization*.

For SGD, they are equivalent – but once you add momentum or normalization by the estimated second moment (RMSProp, Adam), weight decay and L^2 -regularization are different.

L2 Regularization – AdamW

It has taken more than three years to realize that using Adam with L^2 -regularization does not work well. At the end of 2017, **AdamW** was proposed, which is Adam with weight decay.

Adam with L^2 -regularization, AdamW

- $\mathbf{s} \leftarrow \mathbf{0}, \mathbf{r} \leftarrow \mathbf{0}, t \leftarrow 0$
- Repeat until stopping criterion is met:
 - Sample a minibatch of m training examples $(\mathbf{x}^{(i)}, y^{(i)})$
 - $\mathbf{g} \leftarrow \frac{1}{m} \sum_i \nabla_{\boldsymbol{\theta}} (L(f(\mathbf{x}^{(i)}; \boldsymbol{\theta}), y^{(i)}) + \frac{\lambda}{2} \|\boldsymbol{\theta}\|^2)$
 - $t \leftarrow t + 1$
 - $\mathbf{s} \leftarrow \beta_1 \mathbf{s} + (1 - \beta_1) \mathbf{g}$
 - $\mathbf{r} \leftarrow \beta_2 \mathbf{r} + (1 - \beta_2) \mathbf{g}^2$
 - $\hat{\mathbf{s}} \leftarrow \mathbf{s} / (1 - \beta_1^t), \hat{\mathbf{r}} \leftarrow \mathbf{r} / (1 - \beta_2^t)$
 - $\boldsymbol{\theta} \leftarrow \boldsymbol{\theta} - \frac{\alpha_t}{\sqrt{\hat{\mathbf{r}} + \epsilon}} \hat{\mathbf{s}} - \alpha_t \lambda \boldsymbol{\theta}$

$$\boldsymbol{\theta} \leftarrow \boldsymbol{\theta} - \frac{\alpha_t}{\sqrt{\hat{\mathbf{r}} + \varepsilon}} \hat{\mathbf{s}} - \alpha_t \lambda \boldsymbol{\theta}$$

In some variants of the algorithm (notably in the original AdamW paper), the authors proposed not to use the learning rate in the weight decay (to decouple the influence of the learning rate on the weight decay).

However, this would mean that if you utilize learning rate decay, you would need to apply it manually also on the weight decay. So currently, both the implementation of `tf.optimizers.experimental.AdamW` (it will move to `tf.optimizers.AdamW` in TF 2.12) and `torch.optim.AdamW` multiplies the (possibly decaying) learning rate and the (constant) weight decay in the update.

Similar to L^2 -regularization, but could prefer low L^1 metric of parameters. We could therefore minimize

$$\tilde{E}(\boldsymbol{\theta}; \mathbb{X}) = E(\boldsymbol{\theta}; \mathbb{X}) + \lambda \|\boldsymbol{\theta}\|_1.$$

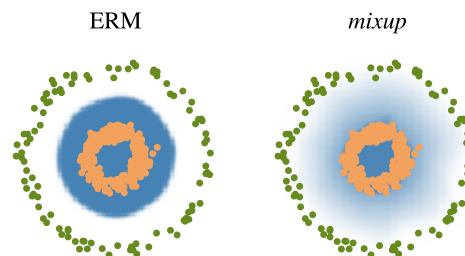
The corresponding SGD update is then

$$\theta_i \leftarrow \theta_i - \alpha \frac{\partial E J}{\partial \theta_i} - \min(\alpha \lambda, |\theta_i|) \text{sign}(\theta_i).$$

Empirically, L^1 -regularization does not work well with deep neural networks and is essentially never used, as far as I know.

For some data, it is cheap to generate slightly modified examples.

- Image processing: translations, horizontal flips, scaling, rotations, color adjustments, ...
 - Mixup (appeared in 2017)



(b) Effect of *mixup* on a toy problem.

Figure 1b of "mixup: Beyond Empirical Risk Minimization", <https://arxiv.org/abs/1710.09412>

- Speech recognition: noise, frequency change, ...
- More difficult for discrete domains like text.

Ensembling (also called **model averaging** or in some contexts *bagging*) is a general technique for reducing generalization error by combining several models. The models are usually combined by averaging their outputs (either distributions or output values in case of a regression).

The main idea behind ensembling is that if models have uncorrelated (independent) errors, then by averaging model outputs the errors will cancel out. If we denote the prediction of the i^{th} model on a training example (\mathbf{x}, y) as $y_i(\mathbf{x}) = y + \varepsilon_i(\mathbf{x})$, so that $\varepsilon_i(\mathbf{x})$ is the model error on example \mathbf{x} , the mean square error of the model is $\mathbb{E}[(y_i(\mathbf{x}) - y)^2] = \mathbb{E}[\varepsilon_i^2(\mathbf{x})]$.

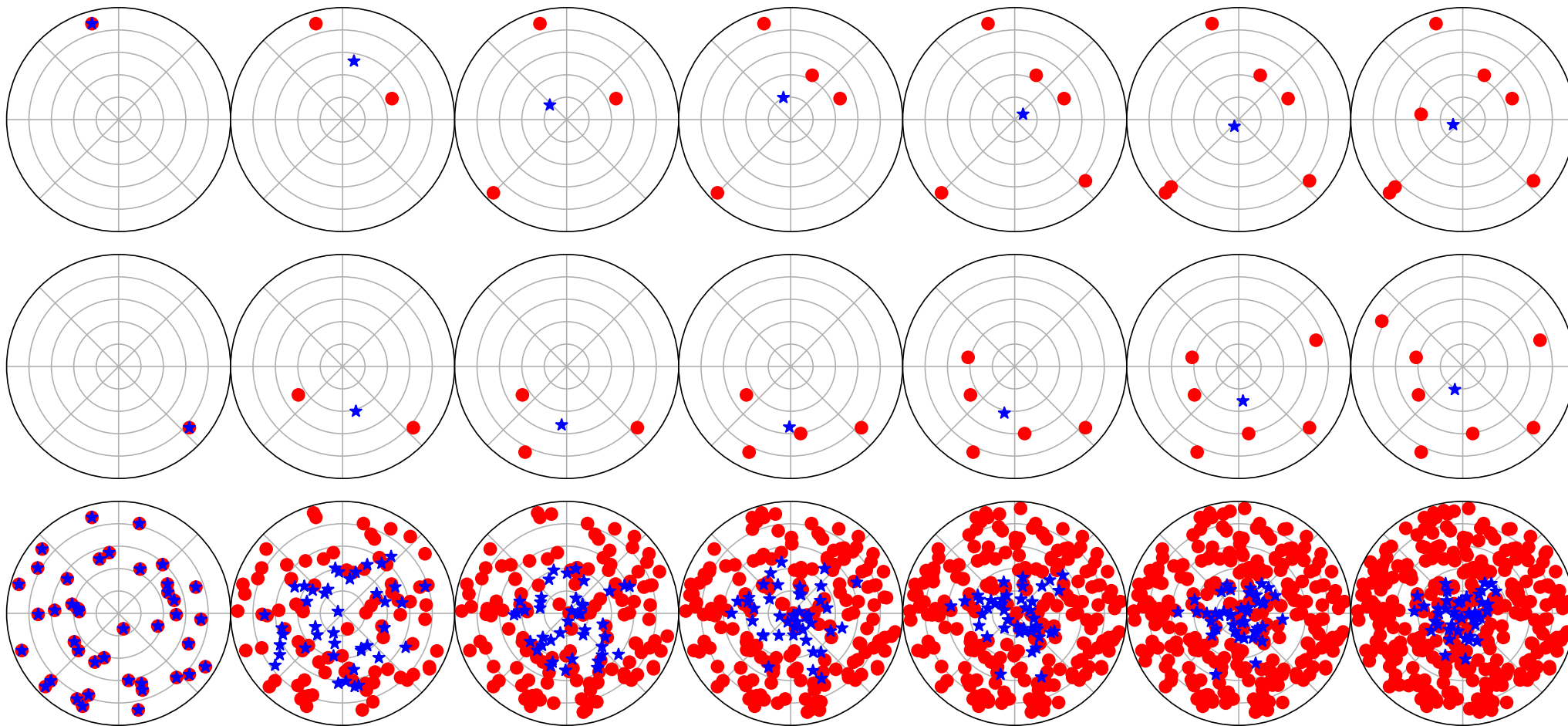
Because for uncorrelated identically distributed random values \mathbf{x}_i we have

$$\text{Var}\left(\sum \mathbf{x}_i\right) = \sum \text{Var}(\mathbf{x}_i), \quad \text{Var}(a \cdot \mathbf{x}) = a^2 \text{Var}(\mathbf{x}),$$

we get that $\text{Var}\left(\frac{1}{n} \sum_i \varepsilon_i\right) = \frac{1}{n} \left(\sum_i \frac{1}{n} \text{Var}(\varepsilon_i)\right)$, so the errors should decrease with the increasing number of models.

Regularization – Ensembling Visualization

Consider ensembling predictions generated uniformly on a planar disc:



Regularization – Ensembling

There are many possibilities how to train the models to ensemble:

- For neural network models, training models with independent random initialization is usually enough, given that the loss has many local minima, so the models tend to be quite independent just when using different random initialization.
- Algorithms with convex loss functions usually converge to the same optimum independent of randomization. In that case, we can use **bagging** (bootstrap aggregation), where we generate different training data for each model by sampling with replacement.
- Average models from last hours/days of training.

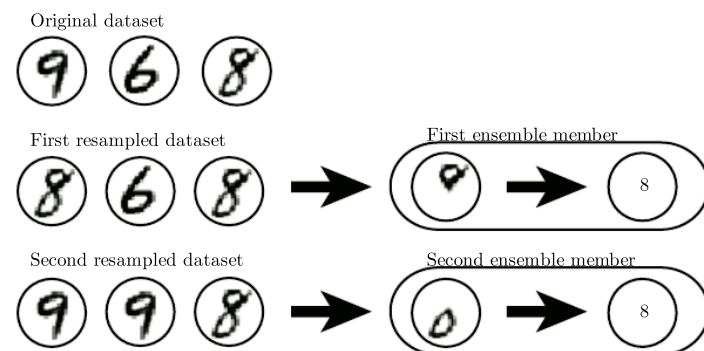


Figure 7.5 of "Deep Learning" book,
<https://www.deeplearningbook.org>

However, ensembling usually has high performance requirements.

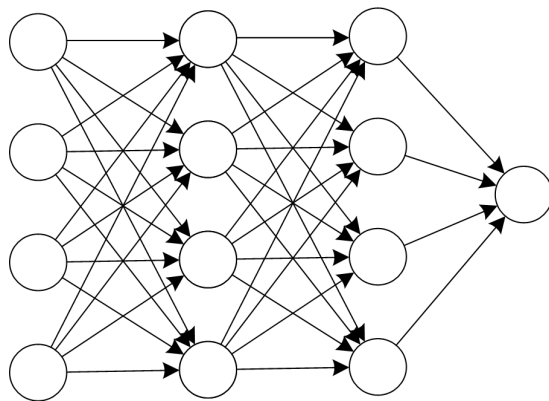
Regularization – Dropout

How to design good universal features?

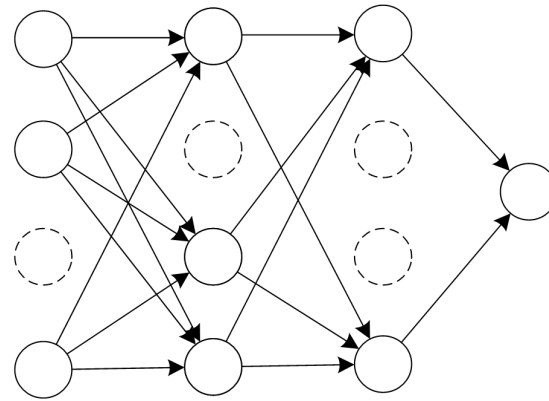
- In reproduction, evolution is achieved using gene swapping. The genes must not be just good with combination with other genes, they need to be universally good.

Idea of **dropout** by (Srivastava et al., 2014), in preprint since 2012.

When applying dropout to a layer, we drop each neuron independently with a probability of p (usually called **dropout rate**). To the rest of the network, the dropped neurons have value of zero.



(a) Standard Neural Network



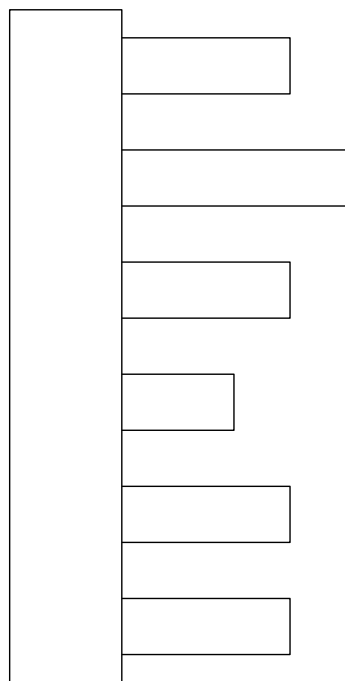
(b) Network after Dropout

Figure 4 of "Multiple Instance Fuzzy Inference Neural Networks" by Amine B. Khalifa et al.

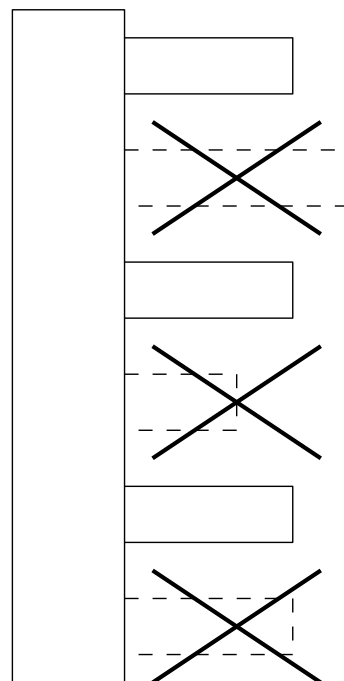
Regularization – Dropout

Dropout is performed only when training, during inference no nodes are dropped. However, in that case we need to **scale the activations down** by a factor of $1 - p$ to account for more neurons than usual.

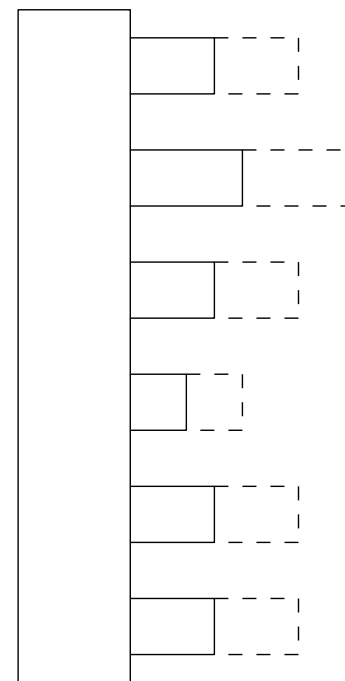
Neuron Activations



Training



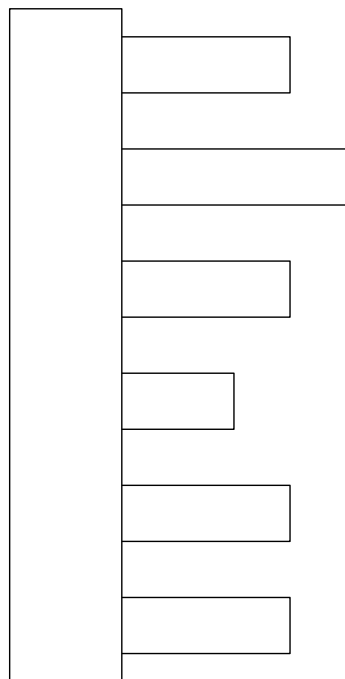
Inference



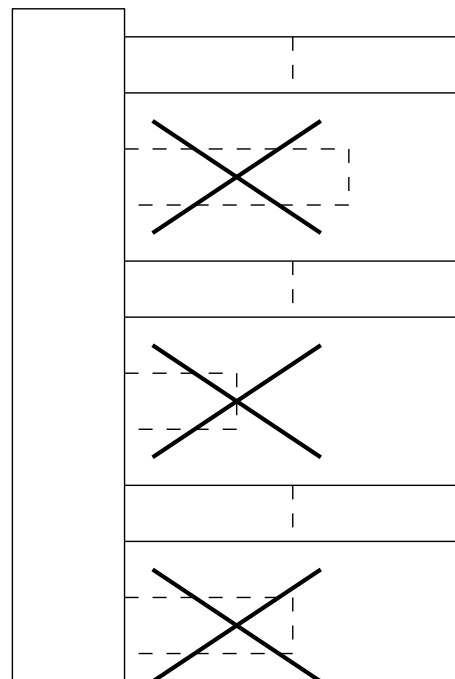
Regularization – Dropout

In practice, the dropout is implemented by instead **scaling the activations up** during training by a factor of $1/(1 - p)$ and then **doing nothing** during inference.

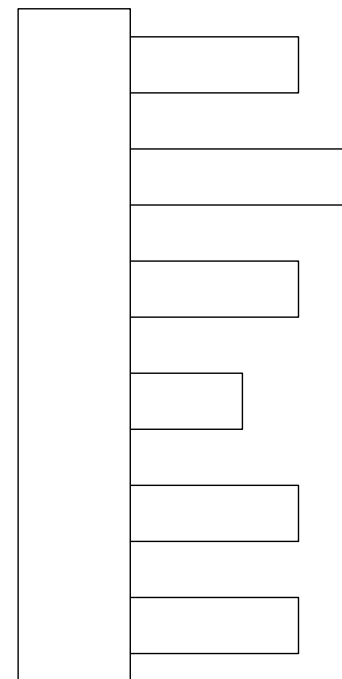
Neuron Activations



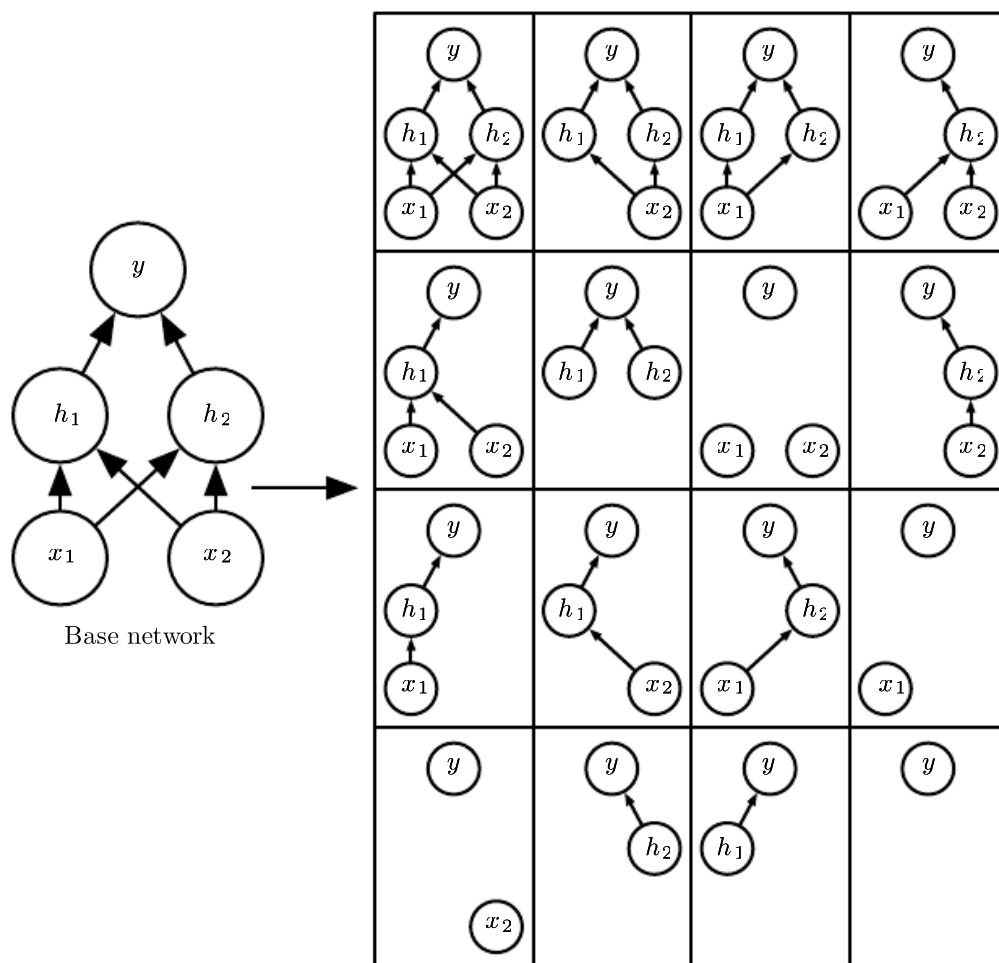
Training



Inference



Regularization – Dropout as Ensembling



Ensemble of subnetworks

Figure 7.6 of "Deep Learning" book, <https://www.deeplearningbook.org>

We can understand dropout as a layer obtaining inputs \mathbf{x} and multiplying them element-wise by a vector of Bernoulli random variables \mathbf{z} , where each z_i is 0 with a probability p :

$$\text{dropout}(\mathbf{x}|\mathbf{z}) = \mathbf{x} \odot \mathbf{z}.$$

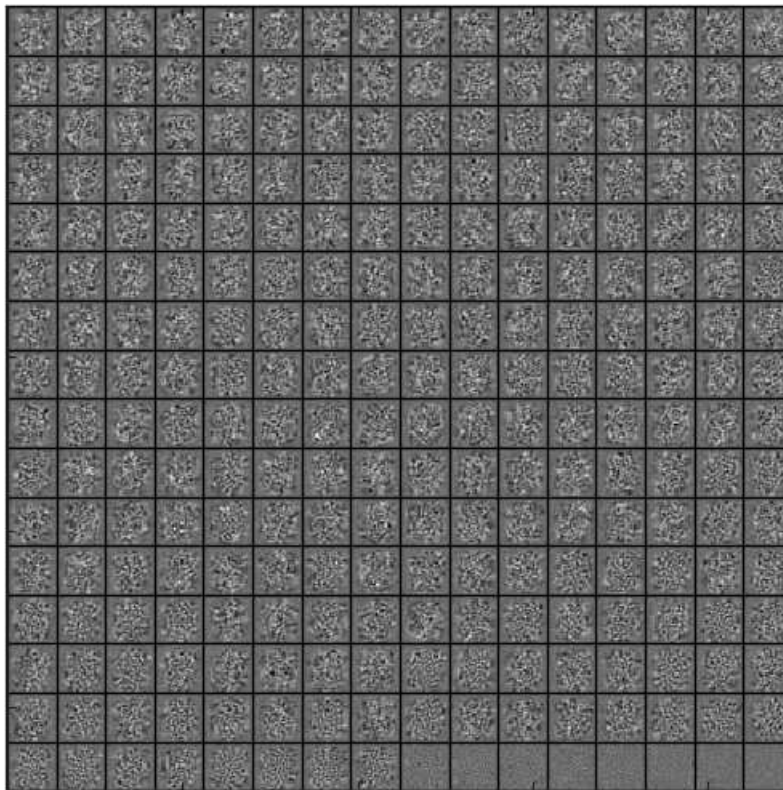
- During training, we sample \mathbf{z} randomly.
- During inference, we compute an expectation over all \mathbf{z} :

$$\begin{aligned} \mathbb{E}_{\mathbf{z}} [\mathbf{x} \odot \mathbf{z}] &= p \cdot \mathbf{x} \odot \mathbf{0} + (1 - p) \cdot \mathbf{x} \odot \mathbf{1} \\ &= (1 - p) \cdot \mathbf{x}. \end{aligned}$$

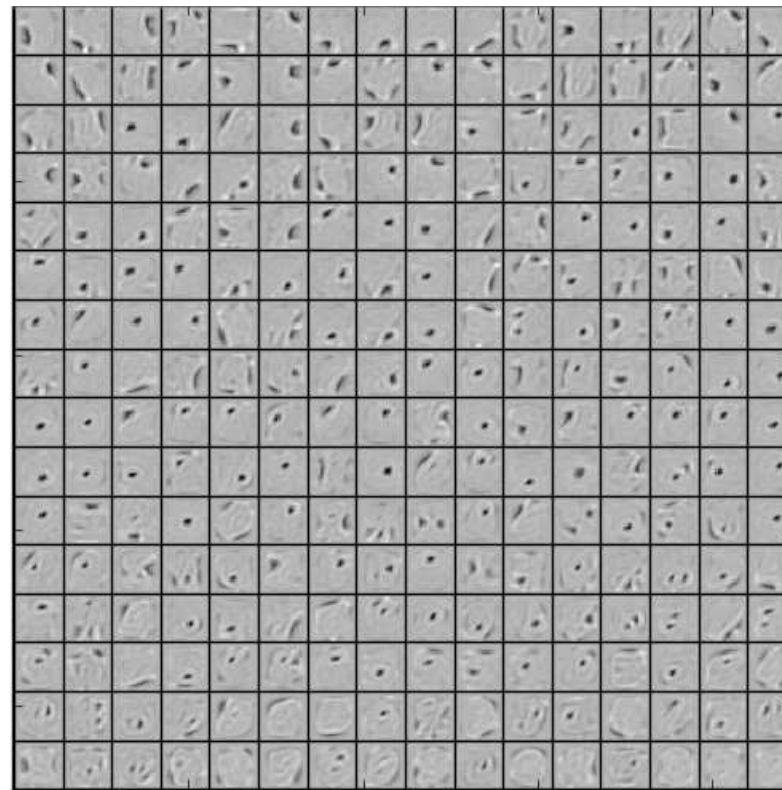
- In order for the inference to be an identity, we can use $\text{dropout}(\mathbf{x}|\mathbf{z}) = \frac{1}{1-p} \cdot \mathbf{x} \odot \mathbf{z}$.

```
def dropout(inputs, rate=0.5, training=False):  
    def do_inference():  
        return inputs  
  
    def do_train():  
        random_noise = tf.random.uniform(tf.shape(inputs))  
        mask = tf.cast(random_noise >= rate, tf.float32)  
        return inputs * mask / (1 - rate)  
  
    if training:  
        return do_train()  
    else:  
        return do_inference()
```


Regularization – Dropout Effect



(a) Without dropout



(b) Dropout with $p = 0.5$.

Figure 7: Features learned on MNIST with one hidden layer autoencoders having 256 rectified linear units.

Figure 7 of "Dropout: A Simple Way to Prevent Neural Networks from Overfitting", <http://jmlr.org/papers/volume15/srivastava14a/srivastava14a.pdf>

Problem with softmax MLE loss is that it is *never satisfied*, always pushing the gold label probability higher (but it saturates near 1).

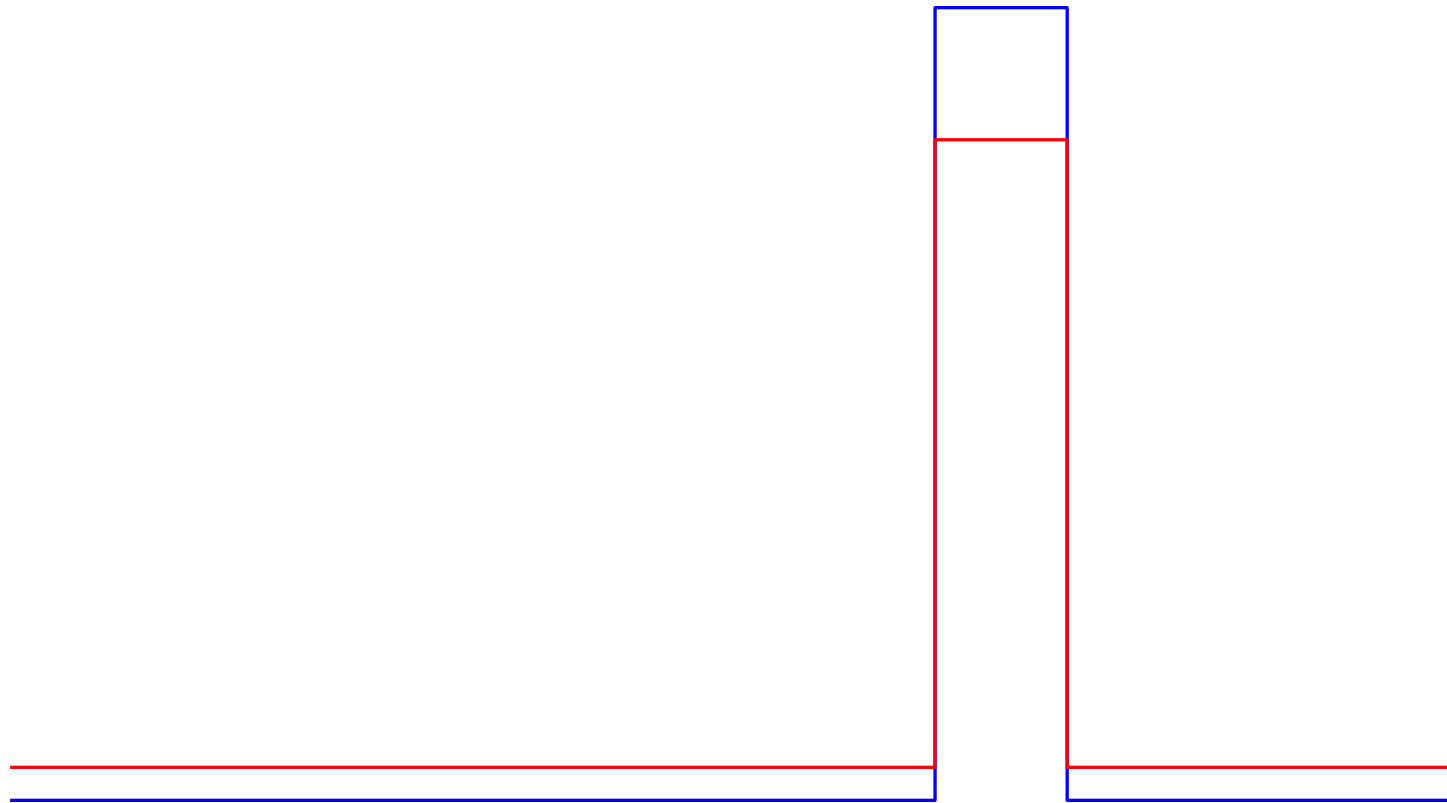
This behaviour can be responsible for overfitting, because the network is always commanded to respond more strongly to the training examples, not respecting similarity of different training examples.

Ideally, we would like a full (non-sparse) categorical distribution of classes for training examples, but that is usually not available.

We can at least use a simple smoothing technique, called *label smoothing*, which allocates some small probability volume α uniformly for all possible classes.

The target distribution is then

$$(1 - \alpha)\mathbf{1}_{gold} + \alpha \frac{1}{\text{number of classes}}.$$



Gold distribution

Smoothed distribution

When you need to regularize (your model is overfitting), then a good default strategy is to:

- use data augmentation if possible;
- use dropout on all hidden dense layers (not on the output layer), good default dropout rate is 0.5 (or use 0.3-0.1 if the model is underfitting);
- use weight decay (AdamW) for convolutional networks;
- use label smoothing (start with 0.1);
- if you require best performance and have a lot of resources, also perform ensembling.

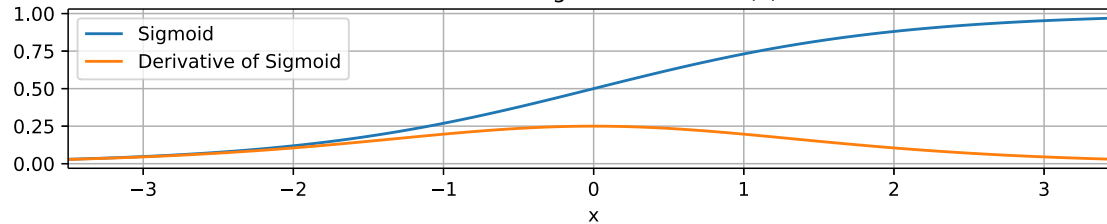
The training process might or might not converge. Even if it does, it might converge slowly or quickly.

A major issue of convergence of deep networks is to make sure that the gradient with respect to all parameters is reasonable at all times, i.e., it does not decrease or increase too much with depth or in different batches.

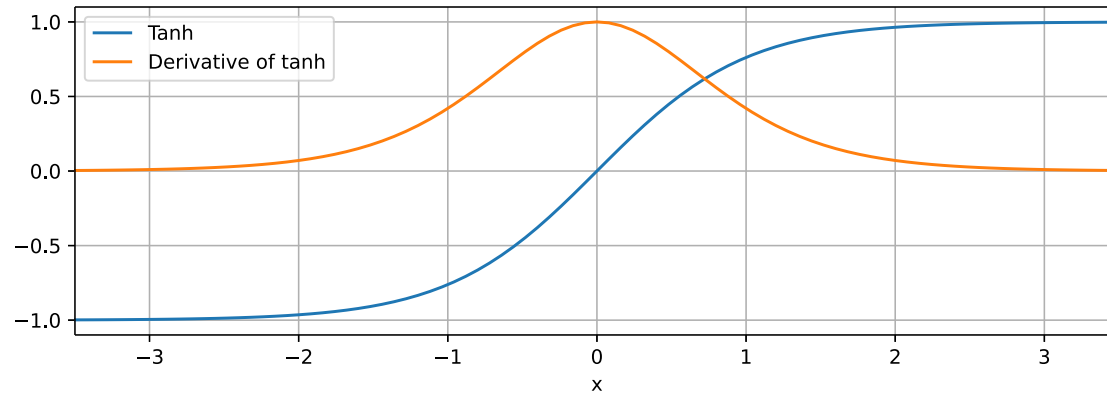
There are *many* factors influencing the gradient, convergence and its speed, we now mention three of them:

- saturating nonlinearities,
- parameter initialization strategies,
- gradient clipping.

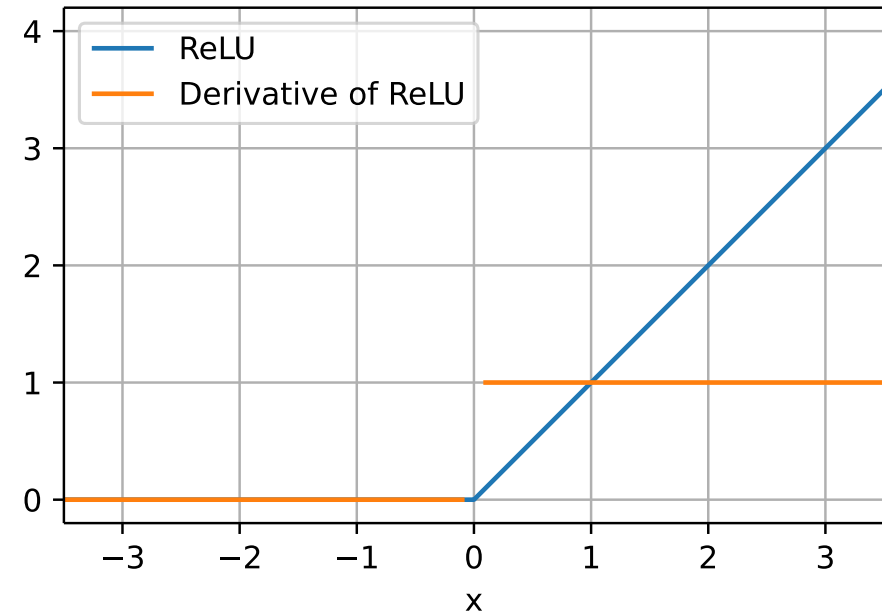
Plot of the Sigmoid Function $\sigma(x)$



Plot of the Tanh Function



Plot of the ReLU Function



Neural networks usually need random initialization to *break symmetry*.

- Biases are usually initialized to a constant value, usually 0.
- Weights are usually initialized to small random values, either with uniform or normal distribution.
 - The scale matters for deep networks!
 - Originally, people used $U \left[-\frac{1}{\sqrt{n}}, \frac{1}{\sqrt{n}} \right]$ distribution.
 - Xavier Glorot and Yoshua Bengio, 2010: *Understanding the difficulty of training deep feedforward neural networks*.

The authors theoretically and experimentally show that a suitable way to initialize a $\mathbb{R}^{n \times m}$ matrix is

$$U \left[-\sqrt{\frac{6}{m+n}}, \sqrt{\frac{6}{m+n}} \right].$$

Convergence – Parameter Initialization

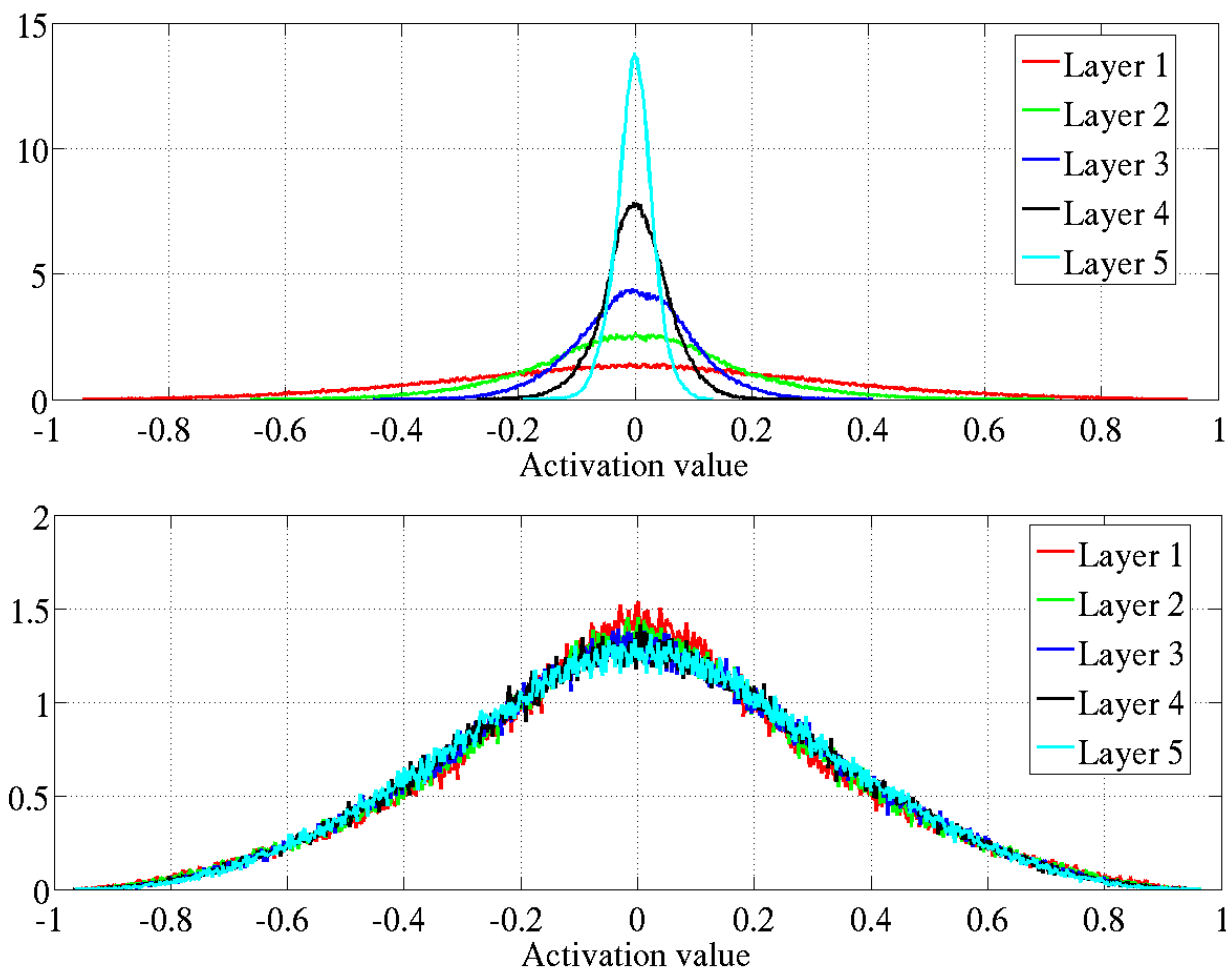


Figure 6 of "Understanding the difficulty of training deep feedforward neural networks", <http://proceedings.mlr.press/v9/glorot10a/glorot10a.pdf>

Convergence – Parameter Initialization

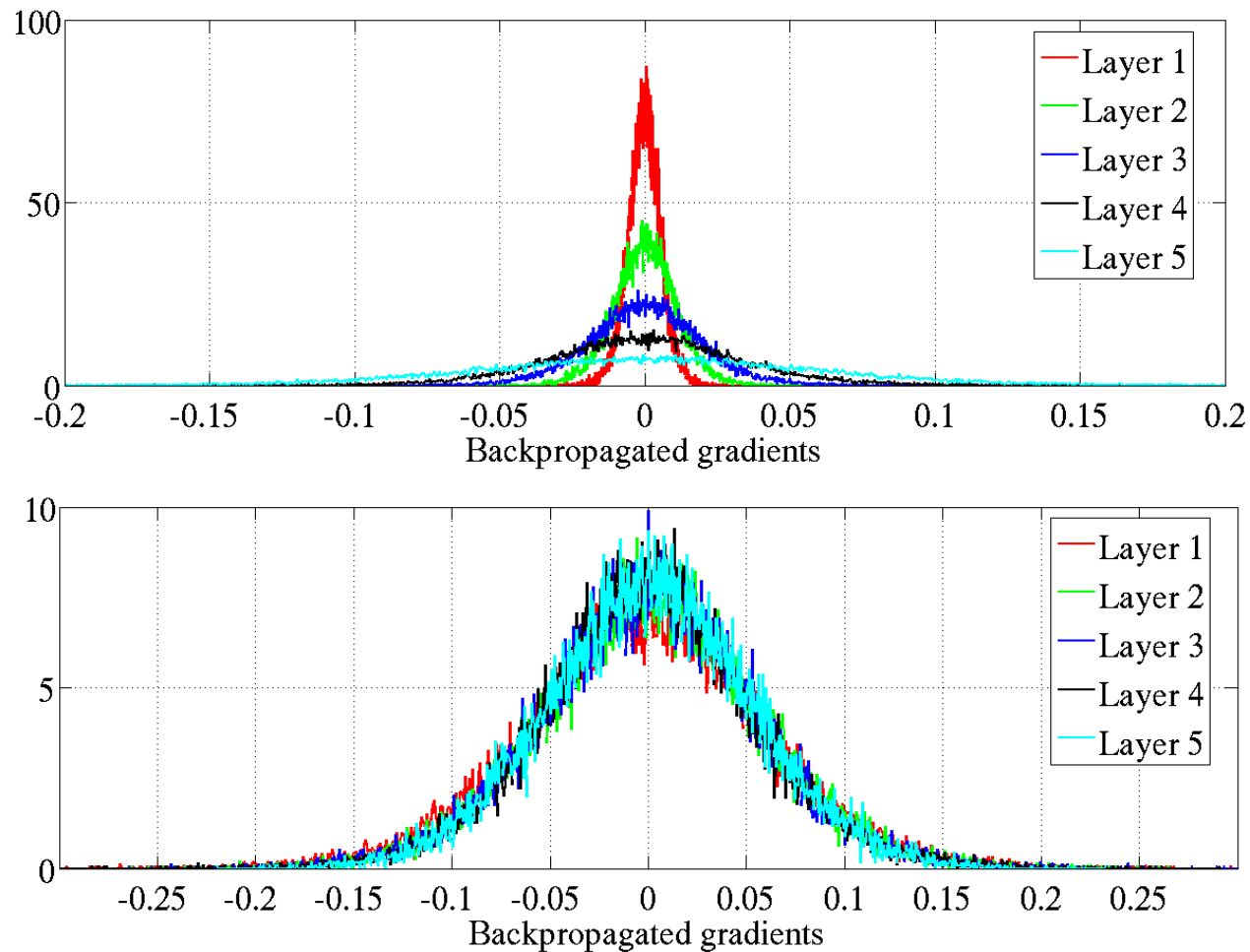


Figure 7 of "Understanding the difficulty of training deep feedforward neural networks", <http://proceedings.mlr.press/v9/glorot10a/glorot10a.pdf>

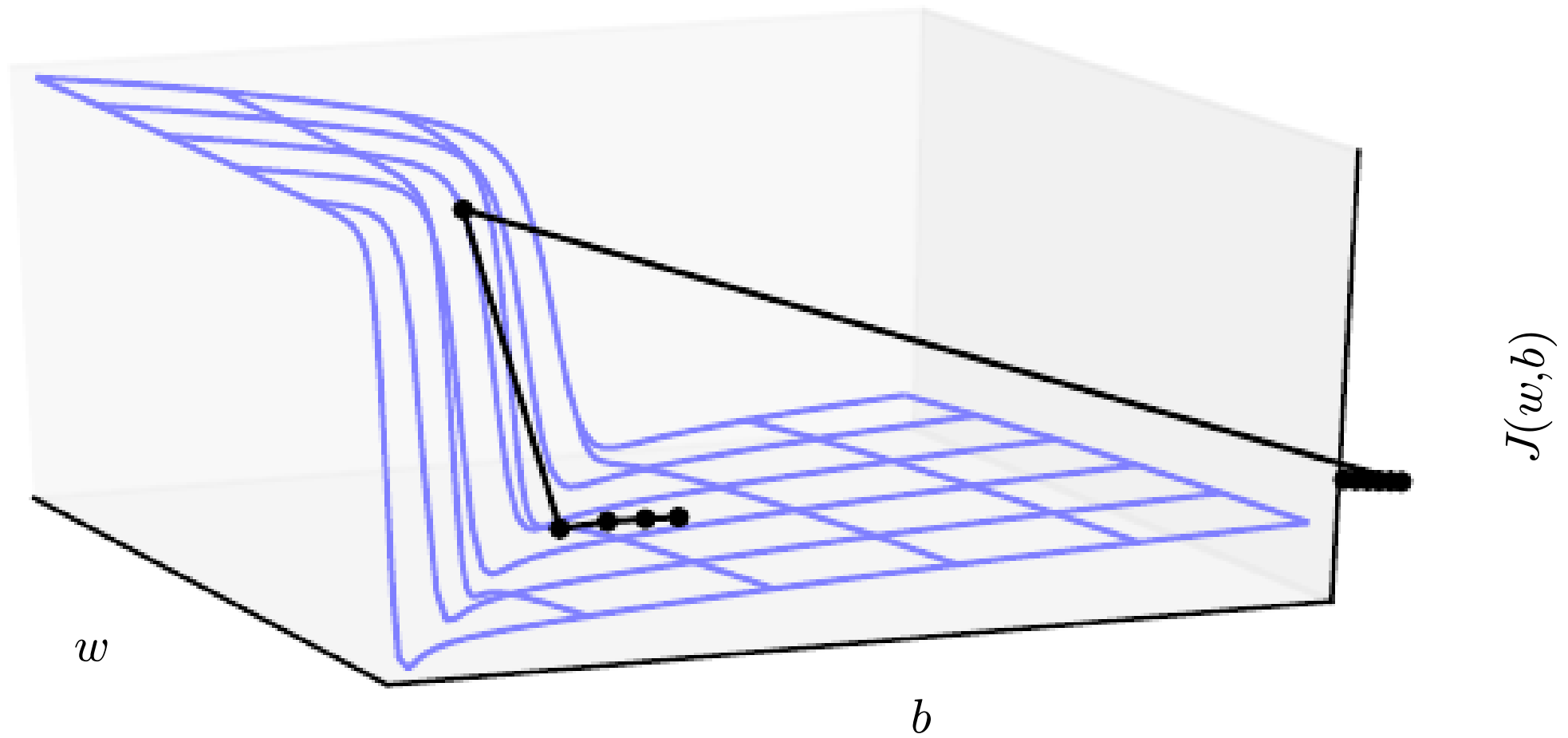


Figure 8.3 of "Deep Learning" book, <https://www.deeplearningbook.org>

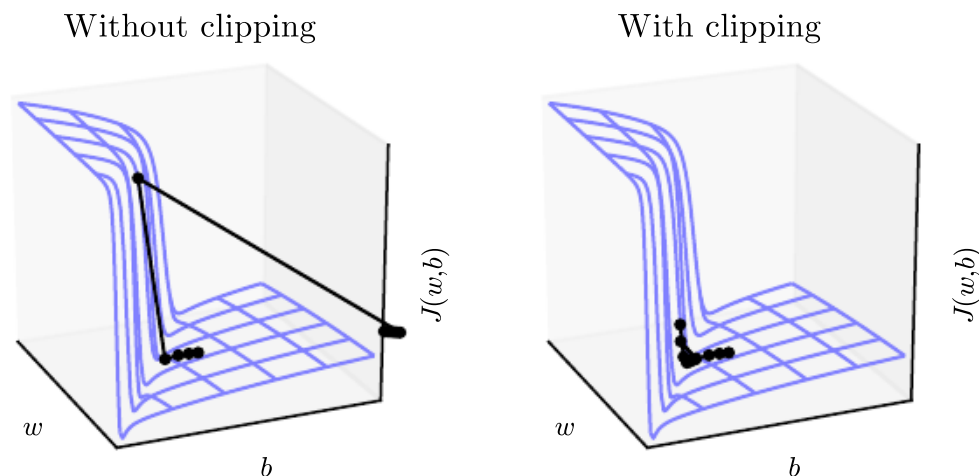


Figure 10.17 of "Deep Learning" book, <https://www.deeplearningbook.org>

Using a given maximum norm, we may *clip* the gradient.

$$\mathbf{g} \leftarrow \begin{cases} \mathbf{g} & \text{if } \|\mathbf{g}\| \leq c, \\ c \frac{\mathbf{g}}{\|\mathbf{g}\|} & \text{if } \|\mathbf{g}\| > c. \end{cases}$$

Clipping can be performed per weight (parameter `clipvalue` of `tf.optimizers.Optimizer`), **per variable** (`clipnorm`) or for the gradient as a whole (`global_clipnorm`).