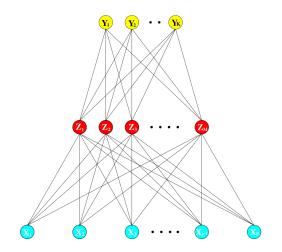
Intro to Neural Networks

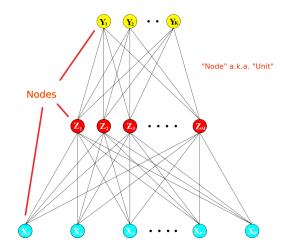
Matthew S. Shotwell, Ph.D.

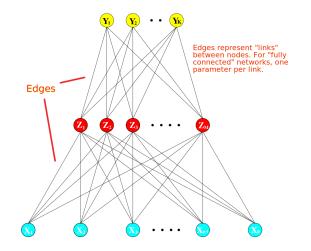
Department of Biostatistics Vanderbilt University School of Medicine Nashville, TN, USA

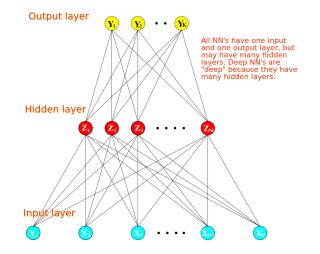
April 8, 2020

・ロト ・母ト ・ヨト ・ヨト ・ヨー うへで









- ▶ NN in previous diagram is for a K class problem
- output layer has K nodes, one for each class
- NN for regression would have just one output node

<ロト < 回 ト < 三 ト < 三 ト 三 三 のへで</p>

Model formula for NN in previous figure; K class problem:

- ▶ input layer: P features, $x_1 \cdots, x_P$
- hidden layer: for each hidden node $m = 1, \dots, M$

$$z_m = \sigma(\alpha_{0m} + \alpha_{1m}x_1 + \dots + \alpha_{Pm}x_P)$$

• output layer: for each output node $k = 1, \ldots, K$

$$t_k = s(\beta_{0k} + \beta_{1k}z_1 + \dots + \beta_{Mk}z_M)$$

< ロ > < 回 > < 三 > < 三 > < 三 > < 三 > < ○ < ○</p>

σ is an "activation" function
 s is a "link" function, e.g., logit, or softmax function

Model formula for NN in previous figure; K class problem:

- ▶ input layer: P features, $x_1 \cdots, x_P$
- hidden layer: for each hidden node $m = 1, \dots, M$

$$z_m = \sigma(\alpha_{0m} + \alpha_{1m}x_1 + \dots + \alpha_{Pm}x_P)$$

• output layer: for each output node $k = 1, \ldots, K$

$$t_k = s(\beta_{0k} + \beta_{1k}z_1 + \dots + \beta_{Mk}z_M)$$

layers are "fully connected" to lower layerall nodes in lower layer contribute to each node of layer above

Model formula for NN in previous figure; K class problem:

- input layer: P features, $x_1 \cdots, x_P$
- ▶ hidden layer: for each hidden node m = 1, ..., M

$$z_m = \sigma(\alpha_{0m} + \alpha_{1m}x_1 + \dots + \alpha_{Pm}x_P)$$

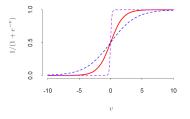
• output layer: for each output node $k = 1, \ldots, K$

$$t_k = s(\beta_{0k} + \beta_{1k}z_1 + \dots + \beta_{Mk}z_M)$$

- α_{0m} and β_{0k} called "bias" parameters
- # params: hidden $M \times (P+1)$; output $K \times (M+1)$

σ Activation function

 σ is an "activation function" designed to mimic the behavior of neurons in propagating signals in the (human) brain. The activation function makes the model nonlinear (in the parameters)



▶ sigmoid -
$$\sigma(x) = \frac{1}{1+e^{-x}}$$

- ► ReLU $\sigma(x) = \max(0, x)$
- ReLU "rectified linear unit"
- ReLU faster training vs sigmoid, may perform better

\boldsymbol{s} Link function

The link function transforms the output nodes so that they make sense for the type of probelm: classification or regression.

▶ regression - identity link - s(t) = t

classification - softmax link -

$$s(t_k) = \frac{e^{t_k}}{\sum_{l=1}^{K} e^{t_l}}$$

- ロ ト - 4 日 ト - 4 日 ト - 4 日 ト - 9 0 0

s(*t_k*) is a value between 0 and 1
 s(*t_k*) is *P*(*y_k* = 1|*x*) where *y_k* is target code for class *k*

- NN's often have large number of parameters: θ
- fit NN's by minimizing average loss in training data!
- Regression: $\overline{\operatorname{err}} = \sum_{i=1}^{N} (y_i f(x_i, \theta))^2$
- ► Classificatin: err = -∑^N_{i=1}∑^K_{k=1} y_{ik} log f_k(x_i, θ), where y_{ik} is the indicator for class k, and f_k gives the probability for class k. This is called the "cross-entropy" or "deviance"
- other loss functions can be used
- err minimized w.r.t. θ using a gradient descent algorithm called "back-propagation" or "backprop"

・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・
 ・

- backprop iterates two steps:
- forward step: fix θ and compute values of hidden and output nodes z_m and t_k, and apply link function to get predictions f(x_i) (probabilities for classification, or numerical prediction for regression)

▶ backward step: fix z_m , t_k , $f(x_i)$ and update θ using a gradient descent step

- ▶ usually don't want global minimum of err due to overfitting
- # of iters of backprop, learning rate (of gradient descent algorithm), and shrinkage penalties (weight decay, dropout) are tuning parameters

Shrinkage:

weight decay: modified objective

 $\overline{\mathrm{err}}(\theta) + \lambda J(\theta)$

where

$$J(\theta) = \sum_{k} \theta_k^2$$

like a ridge penalty, λ is tuning parameter

dropout: at each round of training, some of the hidden or input nodes are Z_m or X_m are ignored (assigned a value zero); ignored inputs are selected at random at each round of backprop; number of ignored features is tuning parameter

Simple NN in R: nnet.R

<ロト < 団ト < 三ト < 三ト < 三 ・ つへの</p>

Extending NN's

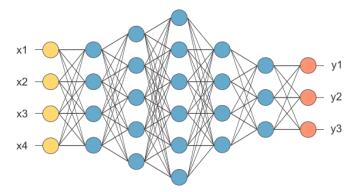
► The real power of NN's comes through various extensions:

▲ロト ▲ □ ト ▲ 三 ト ▲ 三 ト ○ ○ ○ ○ ○ ○

- Additional hidden layers
- Modifying connectivity between layers
- Processing between layers

Additional layers

More than one hidden layer:



- local connectivity: hidden units only receive input from a subset of "local" units in the layer below; not "fully" connected
- say there are 3 hidden nodes and 9 features

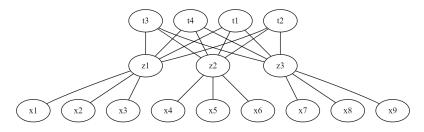
$$z_1 = \sigma(\alpha_{01} + \alpha_{11}x_1 + \alpha_{21}x_2 + \alpha_{31}x_3)$$

$$z_2 = \sigma(\alpha_{02} + \alpha_{12}x_4 + \alpha_{22}x_5 + \alpha_{32}x_6)$$

$$z_3 = \sigma(\alpha_{03} + \alpha_{13}x_7 + \alpha_{23}x_8 + \alpha_{33}x_9)$$

- each hidden node linked to just 3 of 9 features
- hidden layer just $3 \times 4 = 12$ parameters instead of 30
- output layer typically always fully connected

Local connectivity



▲□▶ ▲□▶ ▲ 三▶ ▲ 三▶ ● ○ ○ ○ ○

Iocal connectivity: there can be some "overlap": some hidden nodes can take input from some of the same featuresw

say there are 3 hidden nodes and 9 features

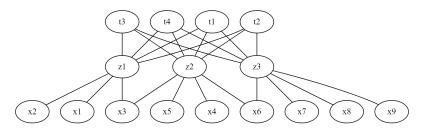
$$z_1 = \sigma(\alpha_{01} + \alpha_{11}x_1 + \alpha_{21}x_2 + \alpha_{31}x_3)$$

$$z_2 = \sigma(\alpha_{02} + \alpha_{12}x_3 + \alpha_{22}x_4 + \alpha_{32}x_5 + \alpha_{42}x_6)$$

$$z_3 = \sigma(\alpha_{03} + \alpha_{13}x_6 + \alpha_{23}x_7 + \alpha_{33}x_8 + \alpha_{43}x_9)$$

<ロト < 回 ト < 三 ト < 三 ト 三 三 のへで</p>

Local connectivity



- weight sharing: some hidden units share weights
- only makes sense in combination with local connectivity
- say there are 3 hidden nodes and 9 features

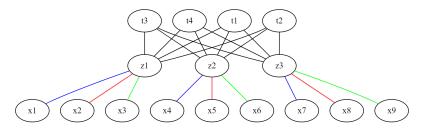
$$z_{1} = \sigma(\alpha_{01} + \alpha_{1}x_{1} + \alpha_{2}x_{2} + \alpha_{3}x_{3})$$

$$z_{2} = \sigma(\alpha_{02} + \alpha_{1}x_{4} + \alpha_{2}x_{5} + \alpha_{3}x_{6})$$

$$z_{3} = \sigma(\alpha_{03} + \alpha_{1}x_{7} + \alpha_{2}x_{8} + \alpha_{3}x_{9})$$

- typically each hidden node retains a distinct bias (α_{0m})
- hidden layer has just 3 + 3 = 6 parameters
- number of links \neq number of parameters (more links)

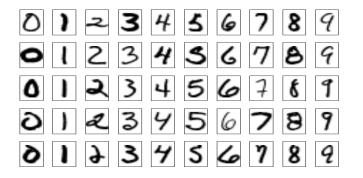
Local connectivity + weight sharing



▲□▶ ▲圖▶ ▲臣▶ ▲臣▶ 三臣 - のへ⊙

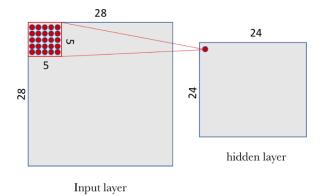
Example: zipcode data

- hand-written integers
- output: 10-class classification
- ▶ input: 16×16 B&W image



Example: zipcode data

- ▶ input: 16×16 B&W image
- when input is an array (2D in this case), typically the hidden units in a hidden layer are represented as an array too
- \blacktriangleright figure below shows local connectivity with 5×5 "kernel"



Local connectivity

- ▶ 3×3 kernel
- ► stride 1
- edge handling

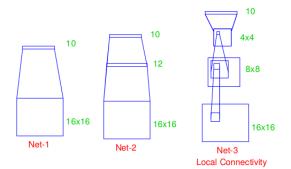
Local conn. w/weight sharing (convolution)

Convolution is type of shape detector or "feature map":

Example: zipcode data

- Net 1: no hidden layer, same as multinomial logistic regression, (256+1)10 = 2570 parameters
- Net 2: one hidden layer, 12 units, fully connected, (256+1)12 + (12+1)10 = 3214 parameters
- ▶ Net 3: two hidden layers, locally connected, 1226 parameters
- Net 4: two hidden layers, locally connected, weight sharing, 1132 parameters, 2266 links

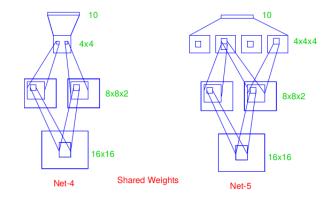
Local connectivity



・ロト・日本・日本・日本・日本・日本

Local connectivity and shared weights

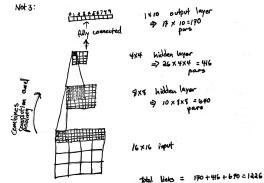
AKA: convolutional neural networks



 groups of hidden units form "shape detectors" or "feature maps"

more complex shape detectors near output layer

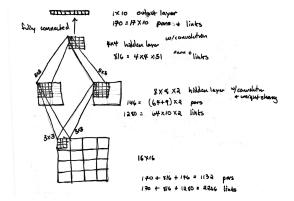
Net 3



total links = 170+446+6703742 total parametric (marging) = total links

<ロト < 目 > < 目 > < 目 > < 目 > < 目 > < 0 < 0</p>

Net 4



Performance on zipcode data

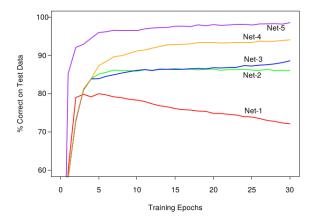
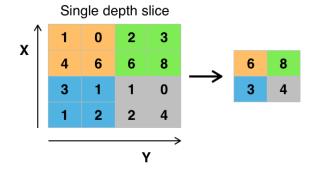


FIGURE 11.11. Test performance curves, as a function of the number of training epochs, for the five networks of Table 11.1 applied to the ZIP code data. (Le Cun, 1989)

<□ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ >

Processing between layers

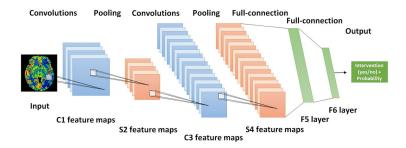
- Pooling/subsampling: down-sample data from a layer by summarizing of a group of units
- Max-pooling: summarize using maximum:



< □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □ > < □

Deep learning and deep neural networks

- Deep learning uses deep NNs
- Deep NNs are simply NNs with many layers, complex connectivity, and processing steps between layers:



Complex NNs in R

- ► No (good) native R libraries for complex NNs
- ▶ R can interface to good libraries, e.g., Keras, TensorFlow

See https://keras.rstudio.com/