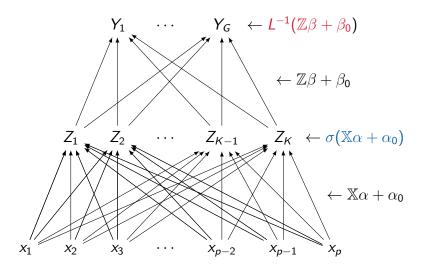
NEURAL NETWORKS AND DEEP LEARNING 2

-Introduction to Data Science-

Lecturer: Darren Homrighausen, PhD

HIERARCHICAL VIEW



RECALL: Single hidden layer neural network. Note the similarity to latent factor models

NEURAL NETWORKS: GENERAL FORM

Generalizing to multi-layer neural networks:

(I'm eliminating the bias term for simplicity) $\overline{}$

$$0 \text{ Layer} := \sigma(\alpha_{\text{lowest}}^{\top} X)$$

$$1 \text{ Layer} := \sigma(\alpha_{\text{lowest}+1}^{\top} (0 \text{ Layer}))$$

$$\vdots$$

$$Top \text{ Layer} := \sigma(\alpha_{\text{Top}}^{\top} (\text{Top - 1 Layer}))$$

$$L(\mu_{g}(X)) = \beta_{g0} + \beta_{g}^{\top} (\text{Top Layer}) \quad (g=1,...G)$$

This looks like iterated matrix multiplications

•
$$\mathbb{Z}_1 = \sigma(\mathbb{X}\alpha_1)$$

:
• $\mathbb{Z}_{L-1} = \sigma(\mathbb{Z}_{L-2}\alpha_{L-1})$
• $\mathbb{Z}_L = \sigma(\mathbb{Z}_{L-1}\alpha_L)$
• $L^{-1}(\beta^\top \mathbb{Z}_L)$

 $(\alpha_l \in \mathbb{R}^{K_{l-1} \times K_l}, \text{ and } K_0 = p))$

NEURAL NETWORKS: GENERAL FORM

Some comments on adding layers:

- It has been shown that one hidden layer is sufficient to approximate any piecewise continuous function (However, this may take a huge number of hidden units (i.e. K >> 1))
- By including multiple layers, we can have fewer hidden units per layer. Also, we can encode (in)dependencies that can speed computations

Back to nonparametric regression

Nonparametric regression

Suppose $Y \in \mathbb{R}$ and we are trying to nonparametrically fit the regression function

$$\mathbb{E}Y|X=f_*(X)$$

A common approach (particularly when p is small) is to specify a fixed basis of functions, $(\phi_k)_{k=1}^K$ (here K is a tuning parameter)

An example of this approach is splines

Nonparametric regression

REMINDER: Nonparametric regression looks like:

1. Write

$$f_*(X) = \sum_{k=1}^K \beta_k \phi_k(X) = \beta^\top \Phi(X)$$

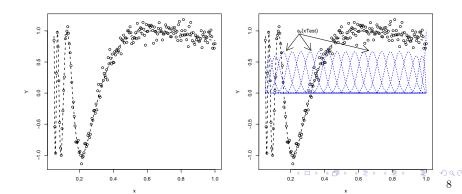
(Again, this is a feature map)

2. Estimate β with least squares (Often a basis function ϕ_k for larger k are rougher \Rightarrow choosing K controls smoothness)

EXAMPLE: ϕ_k could be the k^{th} b-spline basis function

BACK TO THE DOPPLER FUNCTION

```
x = seq(.05,1,length=200)
Y = sin(1/x) + rnorm(100,0,.1)
plot(x,Y)
xTest = seq(.05,1,length=1000)
lines(xTest,sin(1/xTest),col='black',lwd=2,lty=2)
```



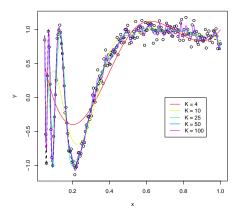
Nonparametric regression: Example

(Code for second plot on previous slide)

```
require(splines)
X = bs(x,df=20)
plot(x,Y)
lines(xTest, sin(1/xTest), col='black', lwd=2, lty=2)
matlines(x=x,X,lty=2,type='1',col='blue')
label = bquote(phi[k](xTest))
text(x=.3,y=1,label)
arrows (x0=.25, y0=.95, x1=.175, y1=max(X[,df/2])+0.02)
arrows (x0=.25,y0=.95,x1=.32,y1=max(X[,df/2])+0.02)
arrows (x0=.25, y0=.95, x1=.66, y1=max(X[,df/2])+0.02)
```

Nonparametric regression: Example

```
require(splines)
X = bs(x,df=K)
Yhat = predict(lm(Y~.,data=X))
```



Nonparametric regression

The weaknesses of this approach are:

- The basis, ϕ_k , is fixed and independent of the supervisor (Just the coefficients β are estimated)
- If p is large, then nonparametrics doesn't work well at all (See previous discussion on curse of dimensionality)
- If the basis doesn't 'agree' with f_* , then K will have to be large to capture the structure
- What if parts of f_* have substantially different structure? (Such is the case in this example..)

An alternative would be to have the data tell us what kind of basis to use

(We have done a version of this twice in the class: kernel methods and boosting)

NEURAL NETWORKS: EXAMPLE

Let's return to the example

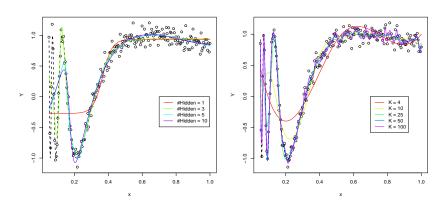
We can fit the data with a single layer NN with varying #'s of hidden units K

A notable difference with B-splines is that 'wiggliness' doesn't necessarily increase with ${\it K}$

Some specifics:

- I used the R package neuralnet
 (This uses the resilient backpropagation version of the gradient descent)
- I regularized via a stopping criterion ($||\partial \ell||_{\infty} < 0.01$)
- I did 3 replications
 (This means I did three starting values and then averaged the results)

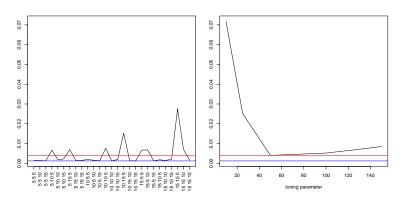
NEURAL NETWORKS: EXAMPLE



Single layer NN (left) vs. B-splines (right)

NEURAL NETWORKS: RISK

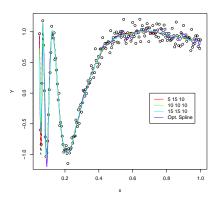
What's the estimation equality? $MSE = \mathbb{E}(\hat{f}(X) - f_*(X))^2$



3 layer¹ NN vs. B-splines

¹The numbers mean (#(layer 1) #(layer 2) #(layer 3))

NEURAL NETWORKS: EXAMPLE



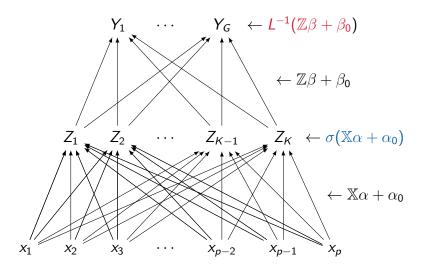
Optimal NNs vs. Optimal B-spline fit

NEURAL NETWORKS: CODE FOR EXAMPLE

```
trainingdata = cbind(x, Y)
colnames(trainingdata) = c("Input", "Output")
testdata
              = xTest
require("neuralnet")
        = c(10,5,15)
K
nRep
          = 3
nn.out
           = neuralnet(Output~Input,trainingdata,
                       hidden=K, threshold=0.01,
                       rep=nRep)
nn.results = matrix(0,nrow=length(testdata),ncol=nRep)
for(reps in 1:nRep){
  pred.obj = compute(nn.out, testdata,rep=reps)
  nn.results[,reps] = pred.obj$net.result
Yhat = apply(nn.results,1,mean)
```

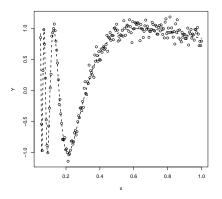
Hierarchical view

HIERARCHICAL VIEW

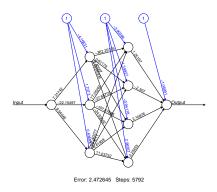


RECALL: Single hidden layer neural network. Note the similarity to latent factor models イロト イ団ト イミト イミト

STILL USING THE DOPPLER FUNCTION...



HIERARCHICAL FROM EXAMPLE



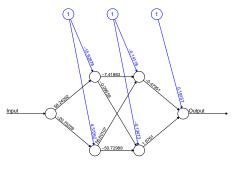
This is a directed acyclic graph (DAG)

One of the main curses/benefits of neural networks is the ability to localize

This makes neural networks very customizable, but commits the data analyst to intensively examining the data

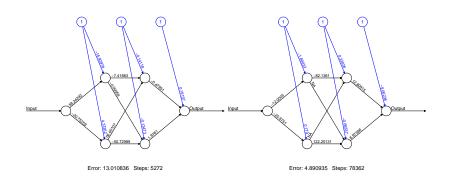
Suppose we are using 1 input and we want to restrict the implicit DAG

That is, we might want to constrain some of the weights to 0



Error: 13.010836 Steps: 5272

Unconstrained neural network



Not-constrained vs. constrained

Detour: Scores

SCORES

A frequently used term in statistics is scores

EXAMPLE A: In PCA we form $\mathbb{X} - \overline{\mathbb{X}} = UDV^{\top}$ and Z = UD are called the (PCA) scores

(I'm using the notation Z = UD to connect it to the neural network notation)

Although it might not look like it, the scores are fitted values

EXAMPLE B: In OLS, we estimate $\hat{\beta}$ via least squares and form the fitted values

$$\hat{Y} = X\hat{\beta}$$

EXAMPLE A: We can recover the PCA scores via

$$(\mathbb{X} - \overline{\mathbb{X}})V = UD \underbrace{V^{\top}V} = UD$$

identity

(Note that PCA isn't applicable in the Doppler problem.. why?)

SCORES

We can get the same sort of information as the PCA scores from neural networks

$$\hat{A}_1 = \begin{bmatrix} \hat{\alpha}_1 0 & \hat{\alpha}_2 0 \\ \hat{\alpha}_1 & \hat{\alpha}_2 \end{bmatrix} = \begin{bmatrix} -15.93 & 4.37 \\ 58.24 & -20.75 \end{bmatrix}$$

Augment $X \in \mathbb{R}^{n \times 1}$ with intercept column:

$$\mathbb{X} = \begin{bmatrix} 1 & X_1 \\ 1 & X_2 \\ & \vdots \\ 1 & X_n \end{bmatrix}$$

 1^{st} layer's scores: $\mathbb{Z}_1 = \sigma(\mathbb{X}\hat{A}_1) \in \mathbb{R}^{n \times 2}$ 2^{nd} layer's scores: $\mathbb{Z}_2 = \sigma(\mathbb{Z}_1\hat{A}_2) \in \mathbb{R}^{n \times 2}$ Input -7.41983 -7.41983 -9.0.72989 -9.0.72989

Error: 13.010836 Steps: 5272

Scores: First Layer

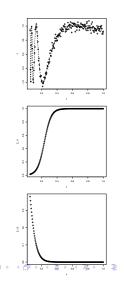
We can get scores from neural networks as well

$$\hat{\mathcal{A}}_1 = \begin{bmatrix} \hat{\alpha}_1 0 & \hat{\alpha}_2 0 \\ \hat{\alpha}_1 & \hat{\alpha}_2 \end{bmatrix} = \begin{bmatrix} -15.93 & 4.37 \\ 58.24 & -20.75 \end{bmatrix}$$

Augment $X \in \mathbb{R}^{n \times 1}$ with intercept column:

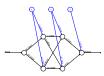
$$\mathbb{X} = \begin{bmatrix} 1 & X_1 \\ 1 & X_2 \\ & \vdots \\ 1 & X_n \end{bmatrix}$$

 1^{st} layer's scores: $\mathbb{Z}_1 = \sigma(\mathbb{X}\hat{A}_1) \in \mathbb{R}^{n \times 2}$



SCORES: SECOND LAYER

We can get scores from neural networks as well



$$\hat{A}_2 = \begin{bmatrix} \hat{\alpha}_1 0 & \hat{\alpha}_2 0 \\ \hat{\alpha}_1 & \hat{\alpha}_2 \end{bmatrix} = \begin{bmatrix} -0.14181 & -0.13473 \\ -7.41883 & 56.60107 \\ -0.09559 & -50.72989 \end{bmatrix}$$

Augment $X \in \mathbb{R}^{n \times 1}$ with intercept column:

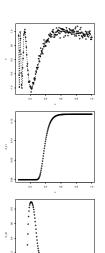
$$[1, \mathbb{Z}_1] \in \mathbb{R}^{n \times 3}$$

 1^{st} layer's scores: $\mathbb{Z}_1 = \sigma(\mathbb{X}\hat{A}_1) \in \mathbb{R}^{n imes 2}$

2nd layer's scores:

$$\mathbb{Z}_2 = \sigma([1, \mathbb{Z}_1] \hat{A}_2) \in \mathbb{R}^{n \times 2}$$

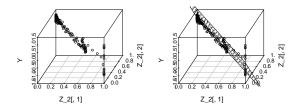
 $([1,\mathbb{Z}_1]$ indicates augmenting \mathbb{Z}_1 with an intercept column)

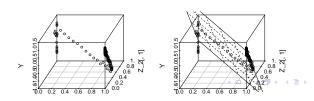


Back to localization

Scores of the second hidden layer vs. the supervisor

 $(2^{nd} \text{ layer's scores: } \mathbb{Z}_2 = \sigma([1,\mathbb{Z}_1]\hat{A}_2) \in \mathbb{R}^{n \times 2}$, the supervisor is Y (the noisy version of the doppler function)



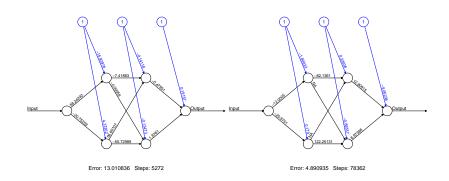


We can localize in neuralnet via the exclude parameter

To use it, do the following:

exclude is a $E \times 3$ matrix, with E the number of exclusions

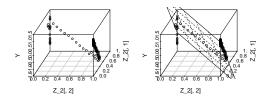
- first column stands for the layer
- the second column for the input neuron
- the third column for the output neuron

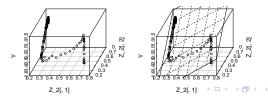


Not-constrained vs. constrained

Scores of the second hidden layer vs. the supervisor for constrained \overline{NN}

 $(2^{nd} \text{ layer's scores: } \mathbb{Z}_2 = \sigma([1,\mathbb{Z}_1]\hat{A}_2) \in \mathbb{R}^{n \times 2}$, the supervisor is Y (the noisy version of the doppler function)

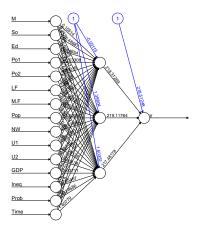




NEURAL NETWORKS: CRIME DATA

```
M
percentage of males aged 1424.
So
indicator variable for a Southern state.
F.d
mean years of schooling.
Po1
police expenditure in 1960.
LF
labour force participation rate.
M.F
number of males per 1000 females.
. . .
У
rate of crimes in a particular category per capita
```

NEURAL NETWORKS: CRIME DATA



NEURAL NETWORKS: CRIME DATA

We may want to constrain the neural network to have neurons specifically about

- Demographic variables
- Police expenditure
- Economics

This type of prior information can be encoded via exclude

(This is, in my opinion, when neural networks work well)

Tuning parameters

NEURAL NETWORKS: TUNING PARAMETERS

We can use a GIC method:

$$AIC = training error + 2\hat{d}f \hat{\sigma}^2$$

(This is reported by neuralnet, by setting likelihood = T)

Or via cross-validation

NEURAL NETWORKS: TUNING PARAMETERS

Unfortunately, neuralnet provides a somewhat bogus measure of AIC/BIC

Here is the relevant part of the code

```
if (likelihood) {
   synapse.count = length(weights) - length(exclude)
   aic = 2 * error + (2 * synapse.count)
   bic = 2 * error + log(nrow(response))*synapse.count
}
```

They use the number of parameters for the degrees of freedom!