

# Introduction to machine learning and neural networks

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## Introduction and applications

Demonstration of overfitting in regression / first steps with R

Classifying images of digits

# Machine learning intro: image classification example

ML is all about learning predictive functions  $f(x) \approx y$ , where

- ▶ Inputs/features  $x$  can be easily computed using traditional algorithms, e.g. matrix of pixel intensities in an image.
- ▶ Outputs/labels  $y$  are what we want to predict, easy to get by asking a human, but hard to compute using traditional algorithms, e.g. image class.
- ▶ Input  $x$  = image of digit, output  $y \in \{0, 1, \dots, 9\}$ ,
  - this is a classification problem with 10 classes.



$$f(\text{0}) = 0$$



$$f(\text{1}) = 1$$

- ▶ Traditional/unsupervised algorithm: I give you a pixel intensity matrix  $x \in \mathbb{R}^{16 \times 16}$ , you code a function  $f$  that returns one of the 10 possible digits. Q: how to do that?

## Supervised machine learning algorithms

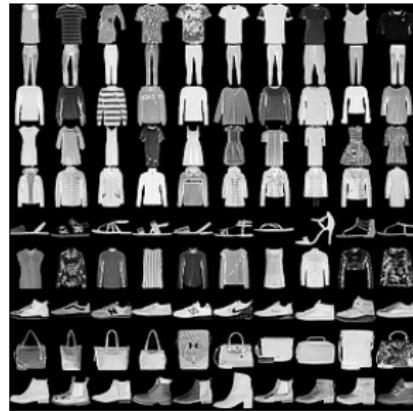
I give you a training data set with paired inputs/outputs, e.g.

0	1	2	3	4	5	6	7	8	9
0	1	2	3	4	5	6	7	8	9
0	1	2	3	4	5	6	7	8	9
0	1	2	3	4	5	6	7	8	9
0	1	2	3	4	5	6	7	8	9
0	1	2	3	4	5	6	7	8	9
0	1	2	3	4	5	6	7	8	9
0	1	2	3	4	5	6	7	8	9
0	1	2	3	4	5	6	7	8	9
0	1	2	3	4	5	6	7	8	9

Your job is to code an algorithm that learns the function  $f$  from the training data. (you don't code  $f$ )

Source: [github.com/cazala/mnist](https://github.com/cazala/mnist)

# Advantages of supervised machine learning

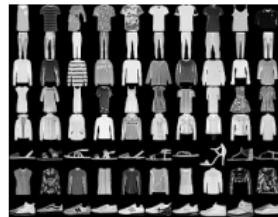


- ▶ Input  $x \in \mathbb{R}^{16 \times 16}$ , output  $y \in \{0, 1, \dots, 9\}$  types the same!
- ▶ Can use same learning algorithm regardless of pattern.
- ▶ Pattern encoded in the labels (not the algorithm).
- ▶ Useful if there are many un-labeled data, but few labeled data (or getting labels is long/costly).
- ▶ State-of-the-art accuracy (if there is enough training data).

# Learning two different functions

Say LEARN is a learning algorithm you have coded.

0	1	2	3	4	5	6	7	8	9
0	1	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>	<b>6</b>	<b>7</b>	<b>8</b>	<b>9</b>
0	1	2	3	4	5	6	7	8	9
0	1	2	3	4	5	6	7	8	9
0	1	2	3	4	5	6	7	8	9
0	1	2	3	4	5	6	7	8	9
0	1	2	3	4	5	6	7	8	9



$\text{LEARN}([0 \ 1 \ 2 \ 3 \ 4 \ 5 \ 6 \ 7 \ 8 \ 9]) \rightarrow f$ ,  $\text{LEARN}([\text{grid of images}]) \rightarrow g$



- ▶ Then we would expect  $f(\text{circle}) = 0$ ,  $f(\text{bar}) = 1$



- ▶  $g(\text{boot}) = \text{shoe}/0$ ,  $g(\text{pants}) = \text{pants}/1$



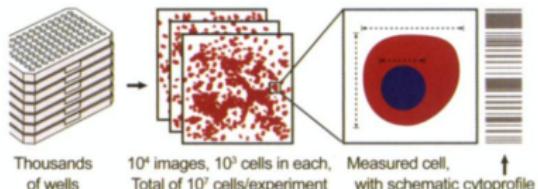
- ▶ Q: what happens if you do  $f(\text{boot})$ , or  $g(\text{circle})$ ?

# Machine learning for cell image classification (CellProfiler)

Jones *et al.* PNAS 2009. Scoring diverse cellular morphologies in image-based screens with iterative feedback and machine learning.

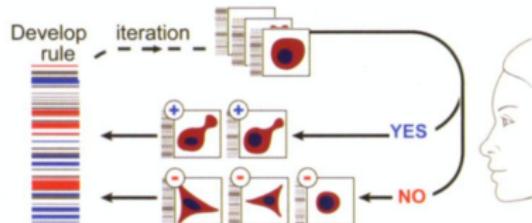
## A Automated Cell Image Processing

Cytoprofile of 500+ features measured for each cell



## B Iterative Machine Learning

System presents cells to biologist for scoring, in batches



- ▶ Input  $x$  = image of cell,
- ▶ Output  $y \in \{\text{yes, no}\}$  (binary classification),

- ▶  $f(\text{ } ) = \text{yes,}$
- ▶  $f(\text{ } ) = \text{no.}$

# Machine learning for image segmentation (LabelMe)

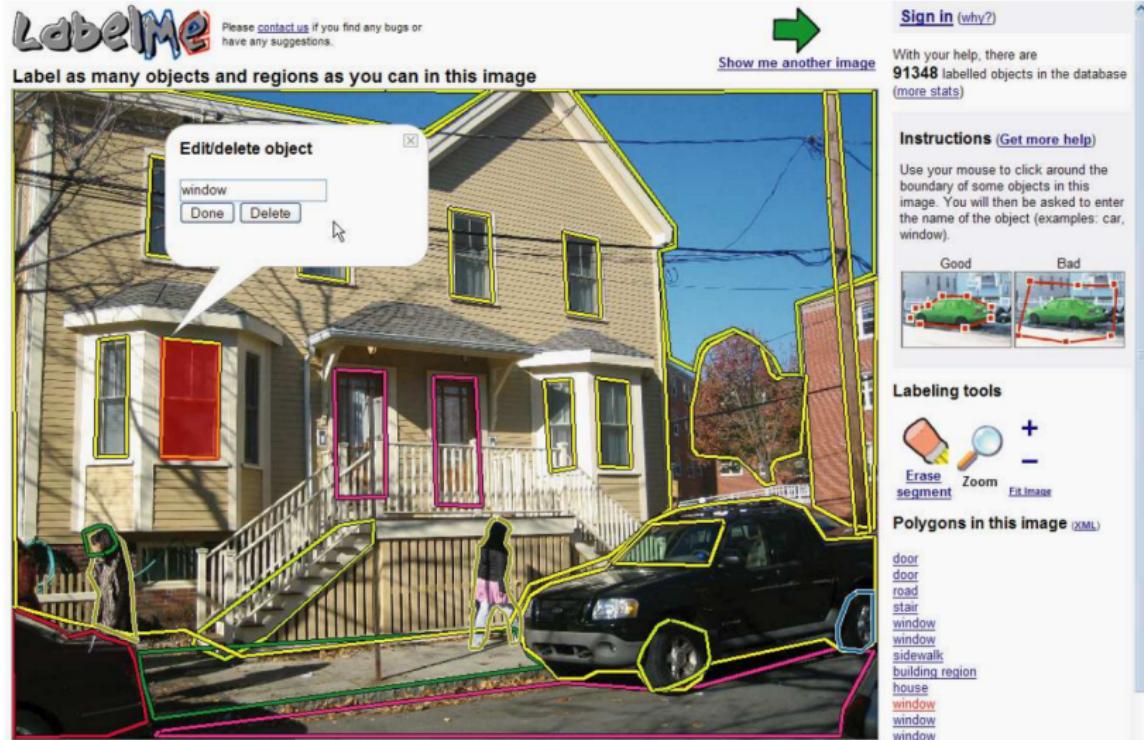


Figure 1. A screenshot of the labeling tool in use. The user is shown an image along with possibly one or more existing annotations, which are drawn on the image. The user has the option of annotating a new object by clicking along the boundary of the desired object and labeling it. Labels can be added to the image with the "Edit/delete object" dialog.



# Machine learning for image segmentation

Paper cup



Rock



Statue



Chair



Russell *et al.* 2007.

Q: What are the types/dimensions of  $x, y, f$  in this example?

# Machine learning for spam filtering (Gmail)

A few of your incoming messages has been suspended ➔ Spam

De Vito Raffael... Jan 11, 2019, 5:11 AM (3 days ago) ★ ↗  
to no\_reply@micorosoft.net

Why is this message in spam? It is similar to messages that were identified as spam in the past.

Report not spam

Re: Assay Standards call, September 5th

Myrto Kostadima ko... Tue, Sep 5, 2017, 1:45 AM ↗  
to Cath, assay\_standards@lhec-intranet.org

Please, find attached a short presentation on the ChIP-seq analysis pipeline.

Thanks,  
Myrto

Your Messages Has Been Suspended By Microsoft Outlook Team

A few of your incoming messages has been suspended by Microsoft verification Team because your email box account has not been properly verify. Do [verify](#) now to receive your suspended messages now.

On 01/09/2017 20:09, Cath Ennis wrote:

Hello all

We have a call scheduled for Tuesday September 5<sup>th</sup> at 6am Pacific time. I've attached the minutes from the last call (also available on the IHEC intranet)

Want:  $f(\text{email message}) \in \{0, 1\}$  – binary classification, spam=1 or not=0.

# Machine learning for translation (google books)

LE COMTE  
DE  
COUNT OF MONTE-CRISTO.  
MONTE-CRISTO  
BY  
ALEXANDRE DUMAS

ABRIDGED AND ANNOTATED  
BY  
EDGAR EWING BRANDON, A.M.  
*Professor of French in Miami University*

Twenty Illustrations,  
DRAWN ON WOOD BY M. VALENTIN,  
AND EXECUTED BY THE MOST EMINENT ENGLISH ENGRAVERS,  
UNDER THE SUPERINTENDENCE OF MR. CHARLES HEATH.



NEW YORK  
HENRY HOLT AND COMPANY  
1900

IN TWO VOLUMES.  
VOL. II.

LONDON:  
CHAPMAN AND HALL, 186 STRAND.

1846.

## Machine learning for translation (google translate)

— Fernand! lui cria-t-il, de mes noms, je n'aurais besoin de t'en dire qu'un seul pour te foudroyer; mais ce nom, tu le devines, n'est-ce pas? ou plutôt tu te le rappelles? car, malgré tous mes chagrins, toutes mes tortures, je te montre aujourd'hui un visage que le bonheur de la vengeance rajeunit, un visage que tu dois avoir vu bien souvent dans tes rêves depuis ton mariage... avec Mercédès, ma fiancée!

Le général, la tête renversée en arrière, les mains étendues, le regard fixe, dévora en silence ce terrible spectacle; puis, allant chercher la muraille comme point d'appui, il s'y glissa lentement jusqu'à la porte par laquelle il sortit à reculons, en laissant échapper ce seul cri lugubre, lamentable, déchirant:

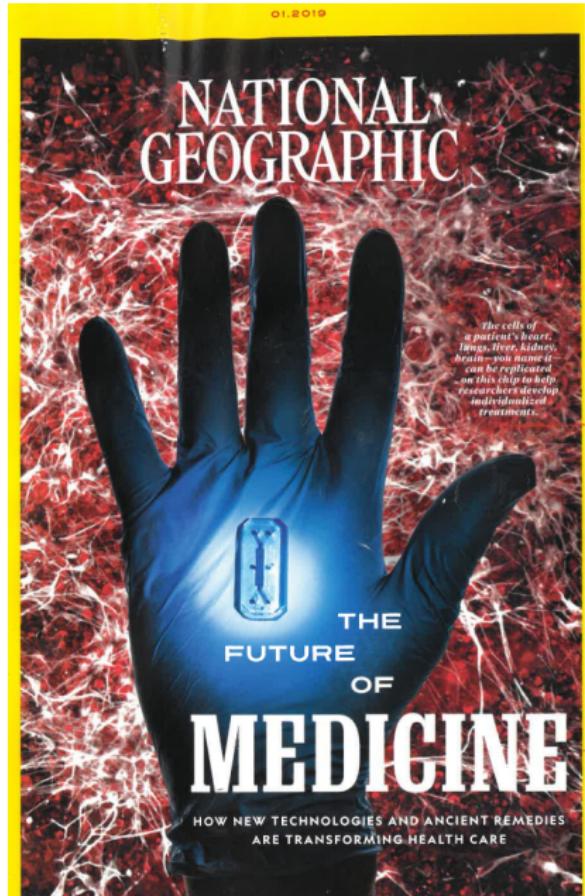
Want:  $f($  — Edmond Dantès!  $) =$

“Fernand,” cried he, “of my hundred names I need only tell you one to overwhelm you! But you guess it now; do you not?—or, rather, you remember it? For, notwithstanding all my sorrows and my tortures, I shew you to-day a face which the happiness of revenge makes young again—a face you must often have seen in your dreams since your marriage with Mercédès, my betrothed!”

The general, with his head thrown back, hands extended, gaze fixed, looked silently at this dreadful apparition; then seeking the wall to support him, he glided along close to it until he reached the door, through which he went out backwards, uttering this single mournful, lamentable, distressing cry,—

“Edmond Dantès!”

# Machine learning for medical diagnosis



f( ) =

Thanks to artificial intelligence and machine learning, diagnostic tools can be trained to read tissue samples and radiologic scans. Google researchers fed more than a quarter-million patients' retinal scans into algorithms that recognize patterns—and the technology "learned" to spot which patterns predict a patient has high blood pressure or is at increased risk for heart attack or stroke. In some comparisons, digital tools produced more accurate analyses than did human pathologists, dermatolo-



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## Machine learning algorithms in practice

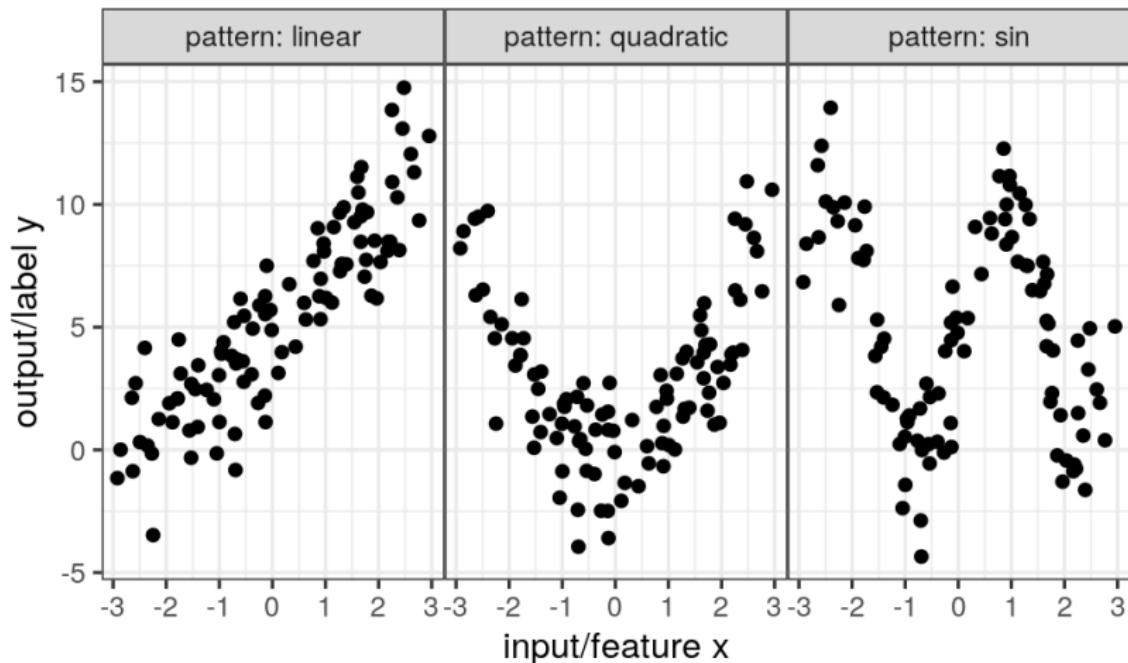
To learn a good prediction function for any of these problems (and any other problems you encounter in your work) you should

- ▶ First gather a data set that you will use to train/test the machine learning algorithms, typically a CSV file with rows for observations and columns for features.
- ▶ Use K-fold Cross-Validation to repeatedly divide the observations into train/test sets. Use the train set to learn machine learning model parameters, and use the test set to estimate prediction error.
- ▶ Try a variety of different learning algorithms (e.g. linear models, random forests, boosting, neural networks, support vector machines, ...), because you don't know which one will result in most accurate predictions for any given data set.
- ▶ Each learning algorithm has different hyper-parameters which control complexity/regularization/overfitting, and typically must be chosen by the user (you) by dividing the train data into subtrain/validation sets. Use the hyper-parameter which is best with respect to the held-out validation set.

## Goal of this section: demonstrate overfitting

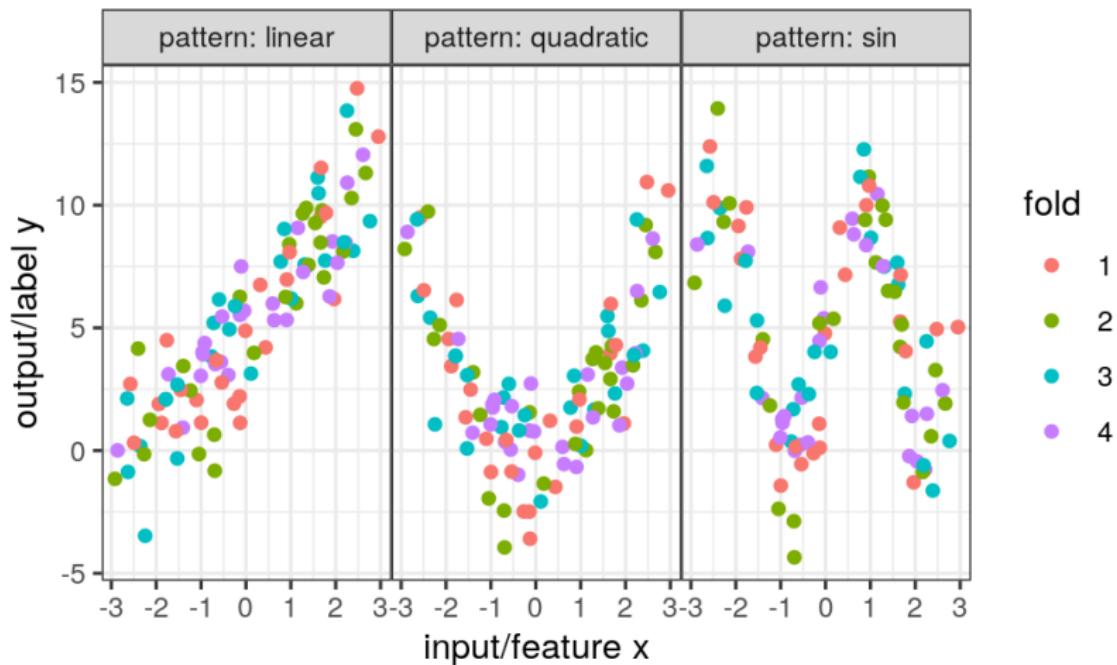
- ▶ The goal of supervised machine learning is to get accurate predictions on new/unseen/held-out test data.
- ▶ Any machine learning algorithm is prone to overfit, which means providing better predictions on the train/subtrain set than on a held-out validation/test set. (BAD)
- ▶ To learn a model which does NOT overfit (GOOD), you need to first divide your train set into subtrain/validation sets.
- ▶ Code for figures in this section: <https://github.com/tdhock/2020-yiqi-summer-school/blob/master/figure-overfitting.R>

## Three different data sets/patterns



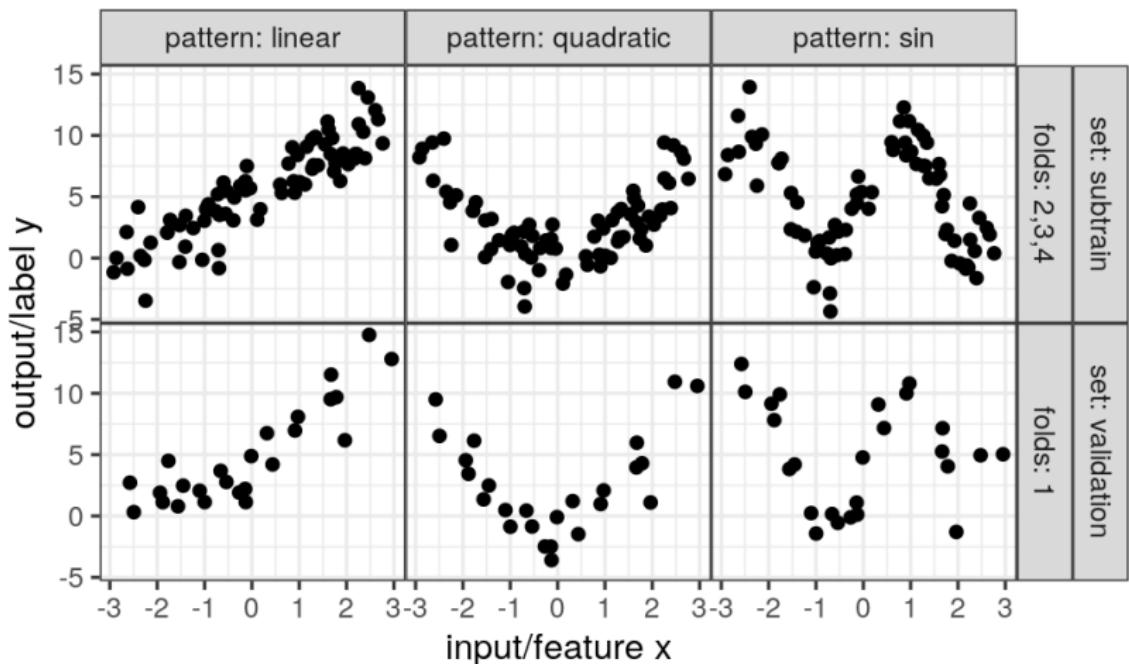
- ▶ We illustrate this using a single input/feature  $x \in \mathbb{R}$ .
- ▶ We use a regression problem with outputs  $y \in \mathbb{R}$ .
- ▶ Goal is to learn a function  $f(x) \in \mathbb{R}$ .

# Illustration of 4-fold cross-validation



Randomly assign each observation a fold ID from 1 to 4.

## Illustration of subtrain/validation split



- ▶ For validation fold 1, all observations with that fold ID are considered the validation set.
- ▶ All other observations are considered the subtrain set.

## How to do this in practice? Use R with an IDE

- ▶ Machine learning model training/evaluation is easy to code in programming languages which support vectors/matrices.
- ▶ I recommend using R because it has so many useful packages (each one for a different machine learning model/algorithm).
- ▶ Download R at <https://cloud.r-project.org/>
- ▶ If you already use Emacs (classic text editor), I recommend using R inside Emacs Speaks Statistics, which is a great Interactive Development Environment (IDE) for R  
<http://ess.r-project.org/> Also see my screencasts about ESS, <https://www.youtube.com/playlist?list=PLwc48KSH3D1Onsed66FPLywMSIQmAhUYJ>
- ▶ Otherwise, I recommend using RStudio, which is another popular IDE for R  
<https://rstudio.com/products/rstudio/download>

## CSV data tables for machine learning

- ▶ One row for each observation.
- ▶ One column for the output/label/y (in regression the label is a real number, in classification the label is a class/category).
- ▶ The other columns should be inputs/features/X that will be used to predict the corresponding output/label/y.

Example: [https://raw.githubusercontent.com/tdhock/2020-yiqi-summer-school/master/data\\_linear.csv](https://raw.githubusercontent.com/tdhock/2020-yiqi-summer-school/master/data_linear.csv)

	x	y
1:	-1.40694802	0.933196336
2:	-0.76725660	3.832773444
3:	0.43712018	4.202983135
4:	2.44924674	13.089055084
...		

## Download CSV data and get file names into R

Download a copy of the github repo to get these data sets

<https://github.com/tdhock/2020-yiqi-summer-school/archive/master.zip>

Set working directory to the downloaded repo:

```
> setwd("~/teaching/2020-yiqi-summer-school")
> getwd()
[1] "/home/tdhock/teaching/2020-yiqi-summer-school"
```

Get a character vector of CSV data file names:

```
> csv.vec <- Sys.glob("data_*.csv")
> csv.vec
[1] "data_linear.csv"      "data_quadratic.csv"
[3] "data_sin.csv"
```

## Read each CSV data file into R

```
sim.data.list <- list()
for(csv in csv.vec){
  pattern <- gsub("data_|.csv", "", csv)
  sim.dt <- data.table::fread(csv)
  set.seed(1) # for reproducibility.
  fold.vec <- sample(rep(1:4, l=nrow(sim.dt)))
  sim.data.list[[pattern]] <- data.table::data.table(
    pattern, sim.dt, fold=factor(fold.vec))
} # combine into a single data table:
sim.data <- do.call(rbind, sim.data.list)
```

- ▶ gsub removes prefix/suffix from CSV file name.
- ▶ data.table::fread gets CSV data from file into R.
- ▶ set.seed used to get consistent/reproducible results from the pseudo-random number generator (sample).
- ▶ data.table::fun means use fun from data.table package, install it via `install.packages("data.table")`

## Result of reading CSV data into R

Result is:

```
> sim.data
   pattern          x          y fold
1: linear -1.4069480  0.9331963    4
2: linear -0.7672566  3.8327734    3
3: linear  0.4371202  4.2029831    1
4: linear  2.4492467 13.0890551    2
5: linear -1.7899084  2.0791987    3
---
296:     sin  1.7838530  4.0502991    1
297:     sin -0.2683533 -0.1097264    1
298:     sin -0.5394955 -0.5539398    1
299:     sin  1.8652215 -0.2262517    4
300:     sin  0.6295997  8.8124249    4
```

## Assign each observation to subtrain/validation set

```
validation.fold <- 1
sim.data[, set := ifelse(
  fold==validation.fold, "validation", "subtrain")]
> sim.data
```

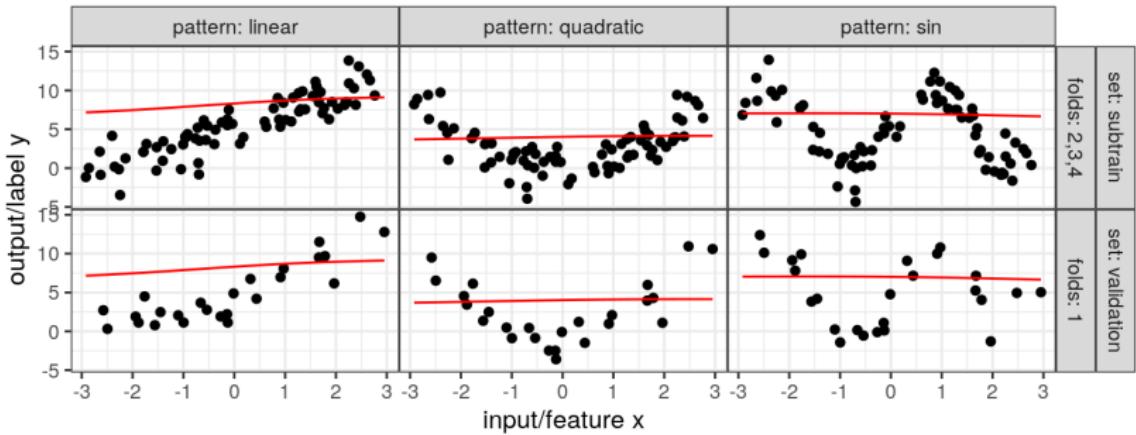
	pattern	x	y	fold	set
1:	linear	-1.4069480	0.9331963	4	subtrain
2:	linear	-0.7672566	3.8327734	3	subtrain
3:	linear	0.4371202	4.2029831	1	validation
4:	linear	2.4492467	13.0890551	2	subtrain
5:	linear	-1.7899084	2.0791987	3	subtrain
---					
296:	sin	1.7838530	4.0502991	1	validation
297:	sin	-0.2683533	-0.1097264	1	validation
298:	sin	-0.5394955	-0.5539398	1	validation
299:	sin	1.8652215	-0.2262517	4	subtrain
300:	sin	0.6295997	8.8124249	4	subtrain

## Neural network with one hidden layer, 20 hidden units

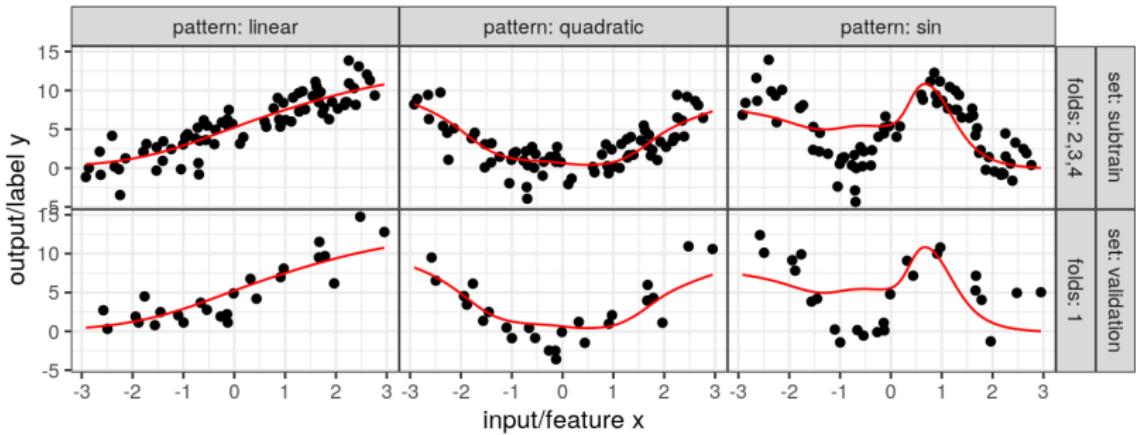
Use for loops to fit different neural network models for each data set and number of iterations.

```
maxit.values <- 10^seq(0, 4)
pattern.values <- c("linear", "quadratic", "sin")
for(i in maxit.values)for(p in pattern.values){
  pattern.data <- sim.data[pattern==p]
  fit <- nnet::nnet(
    y ~ x,
    pattern.data[set=="subtrain"],
    size=20,      #hidden units
    linout=TRUE,  #for regression
    maxit=i)     #max number of iterations
  ...
}
```

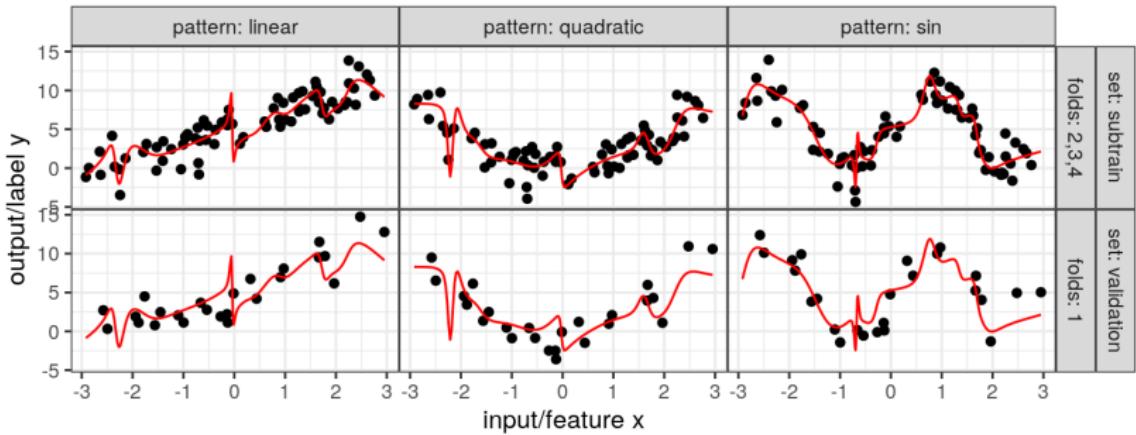
Neural network, 20 hidden units, 1 gradient descent iterations



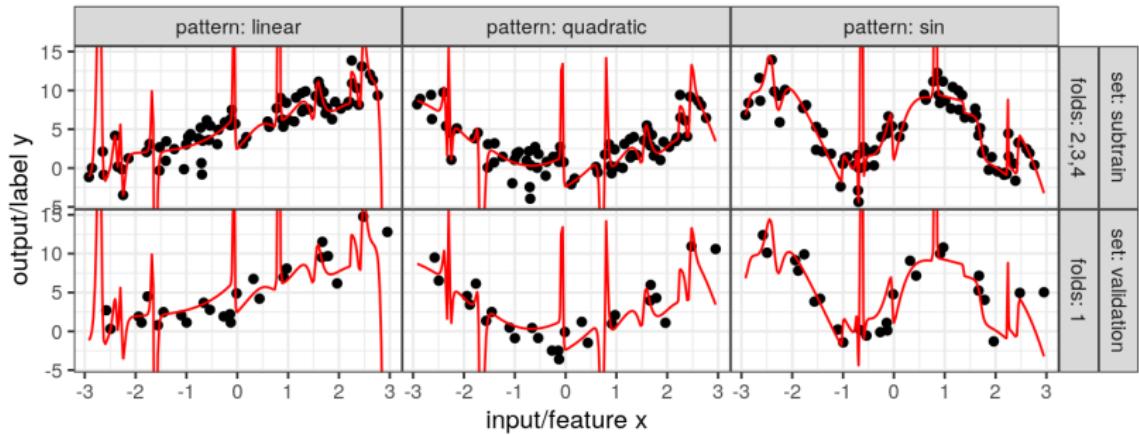
Neural network, 20 hidden units, 10 gradient descent iterations



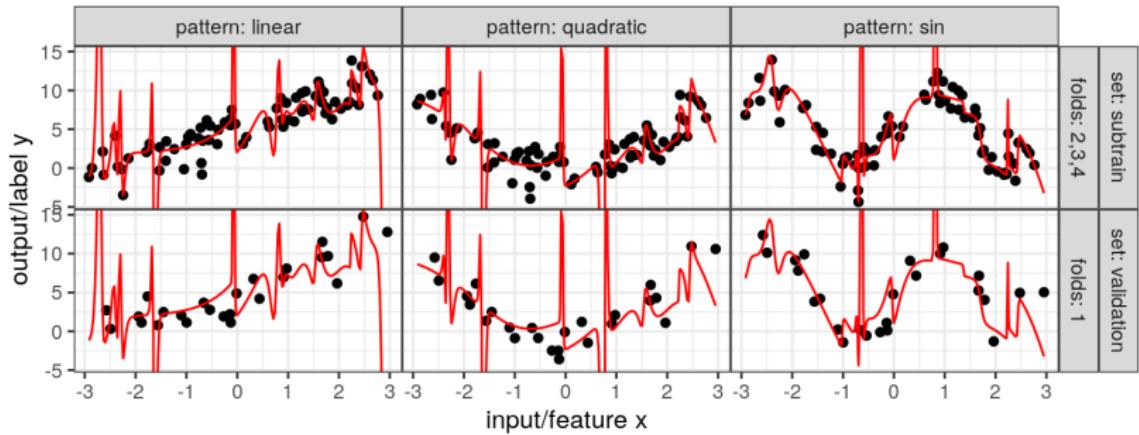
Neural network, 20 hidden units, 100 gradient descent iterations



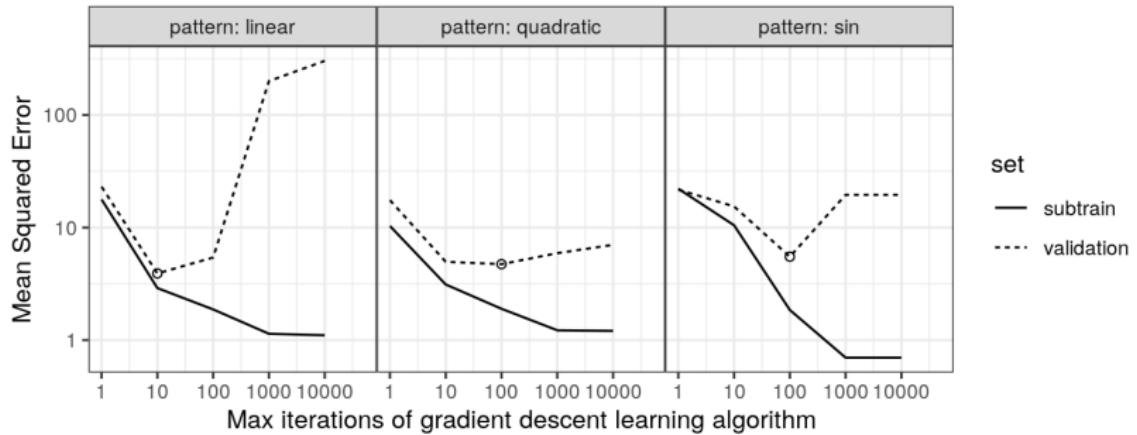
Neural network, 20 hidden units, 1000 gradient descent iterations



Neural network, 20 hidden units, 10000 gradient descent iterations



## Neural network, 20 hidden units



# Overfitting

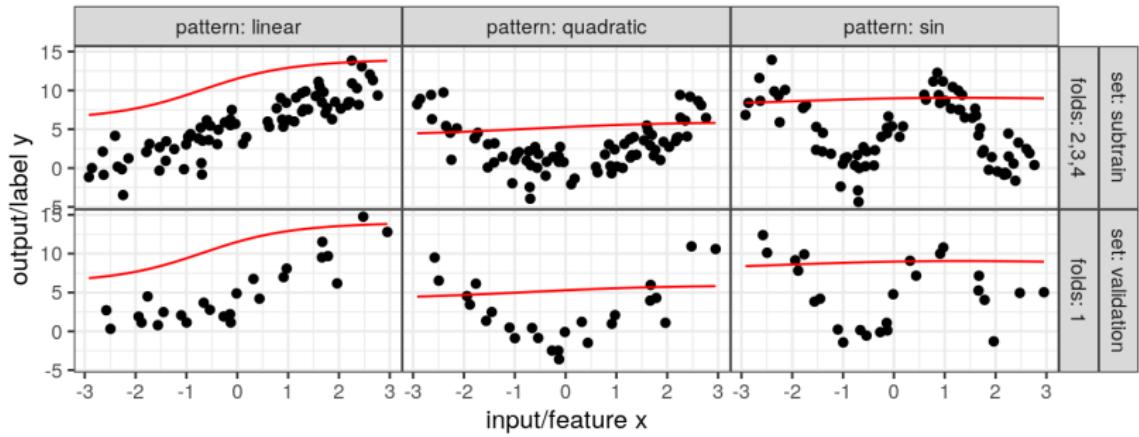
- ▶ Happens when subtrain error/loss decreases but validation error increases (as a function of some hyper-parameter)
- ▶ Here the hyper-parameter is the number of iterations of gradient descent, and overfitting starts after 10–100 iterations.
- ▶ To maximize prediction accuracy you need to choose a hyper-parameter with minimal validation error/loss.
- ▶ This optimal hyper-parameter will depend on the data set.
- ▶ In this case that means choosing 10 iterations for the linear data set, and 100 iterations for the quadratic/sin data sets.

## Computing substrain/validation loss curves

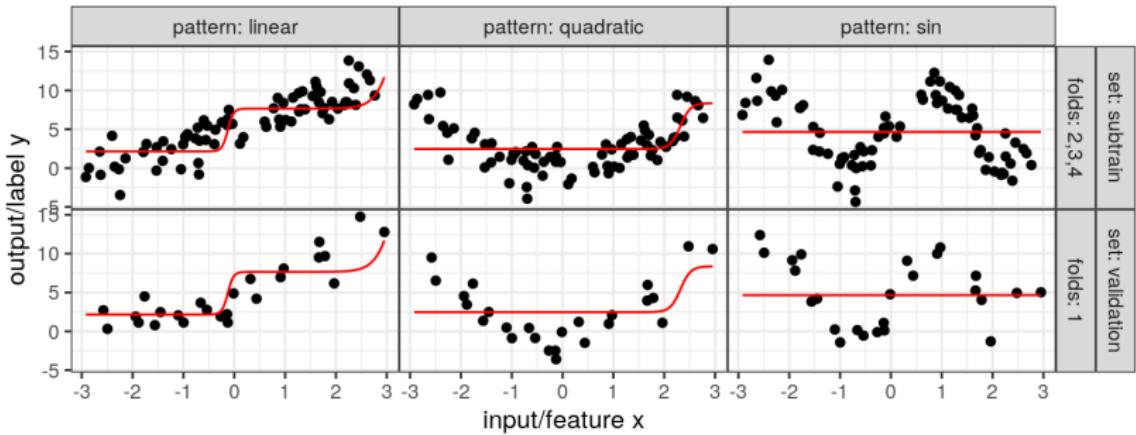
```
...
pattern.data[, pred.y := as.numeric(predict(
    fit, pattern.data))]
loss.dt.list[[paste(pattern, maxit)]] <- data.table(
    pattern=p, maxit=i, pattern.data[, .(
        mse=mean((pred.y-y)^2)), by=set])
}
loss.dt <- do.call(rbind, loss.dt.list)
```

	pattern	set	maxit	mse
1:	linear	substrain	1	17.7785881
2:	linear	validation	1	23.0575528
...				
29:	sin	substrain	10000	0.6970455
30:	sin	validation	10000	19.5051785

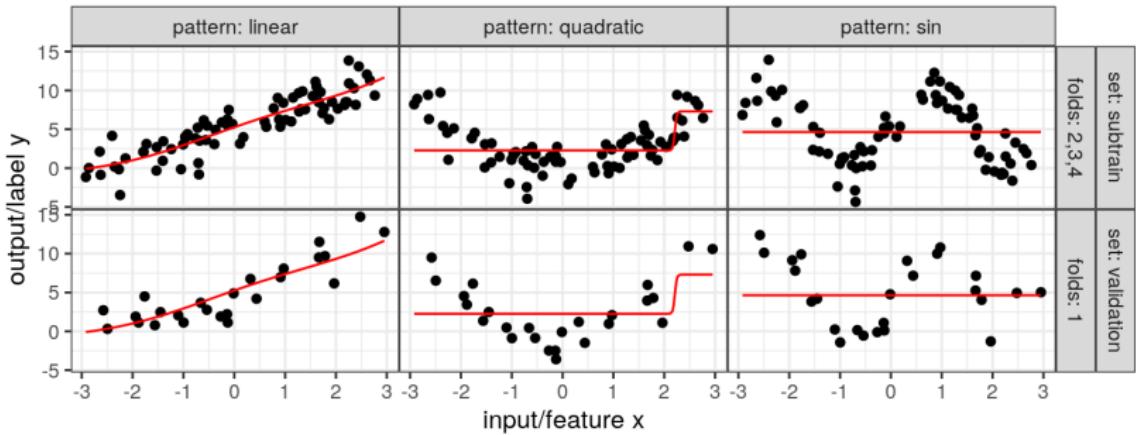
## Neural network, 2 hidden units, 1 gradient descent iterations



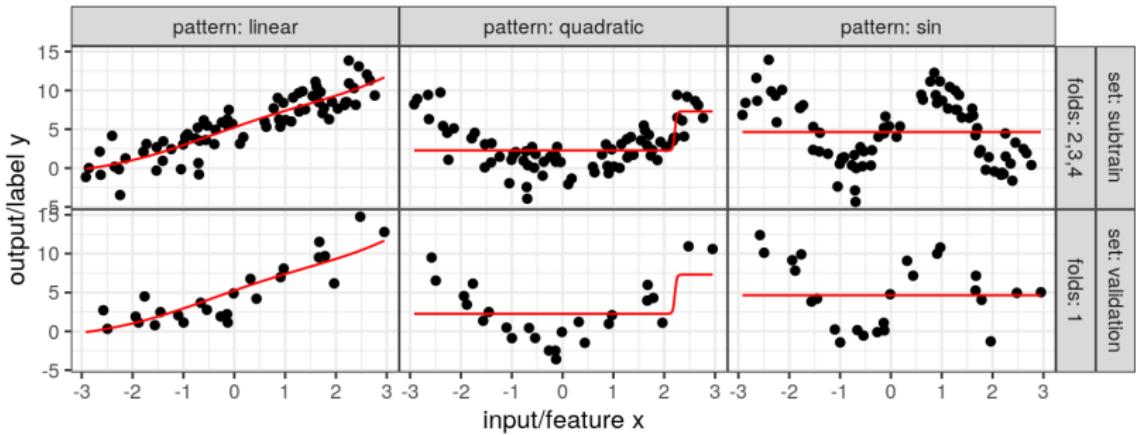
Neural network, 2 hidden units, 10 gradient descent iterations



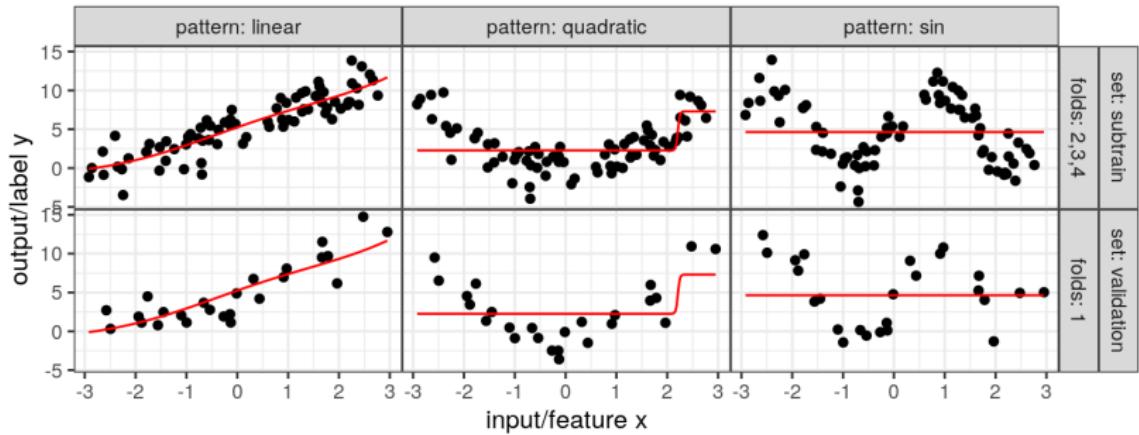
Neural network, 2 hidden units, 100 gradient descent iterations



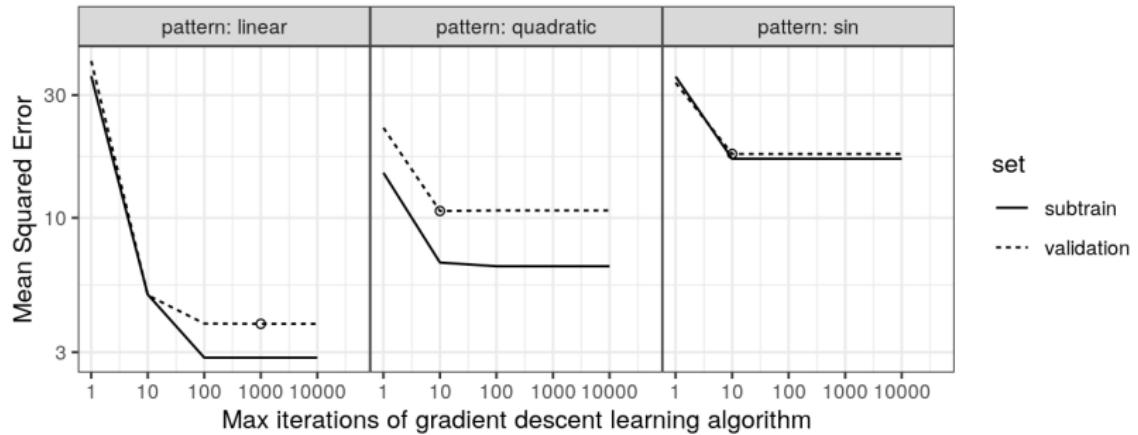
Neural network, 2 hidden units, 1000 gradient descent iterations



Neural network, 2 hidden units, 10000 gradient descent iterations



## Neural network, 2 hidden units



## How does the neural network predict/learn?

$$\text{hidden units: } h_j = \sigma(w_j x + b_j)$$

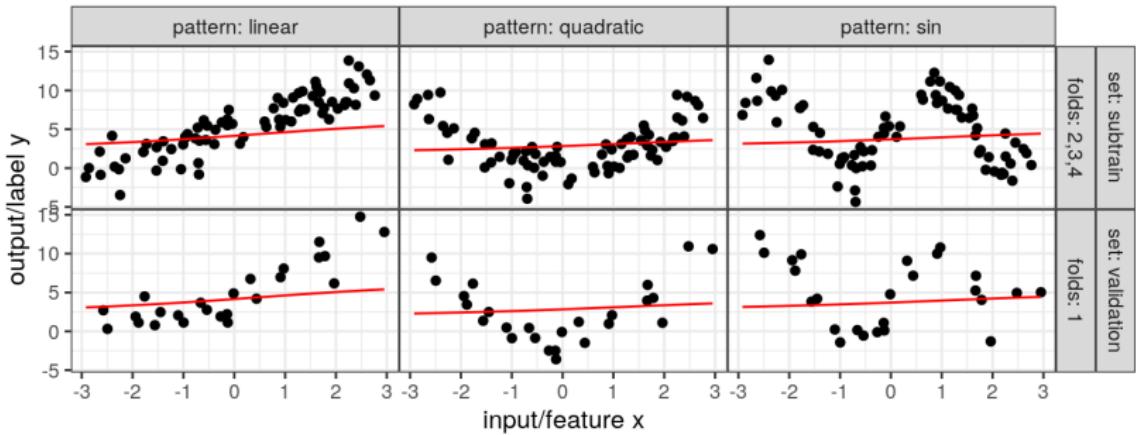
$$\text{predicted output: } o = \hat{y} = \sum_{j=1}^u v_j h_j + \beta$$

- ▶  $x \in \mathbb{R}$  is the input/feature (i1 in R notation below).
- ▶  $\sigma : \mathbb{R} \rightarrow [0, 1]$  is the sigmoid activation function,  
 $\sigma(z) = (1 + e^{-z})^{-1}$ .
- ▶  $h_j \in [0, 1]$  are  $u$  hidden units, e.g.  $u \in \{20, 2\}$  (h1,h2).
- ▶  $w_j, b_j, v_j, \beta \in \mathbb{R}$  are the weights which are learned using gradient descent to minimize the squared error, e.g. for  $u = 2$ :

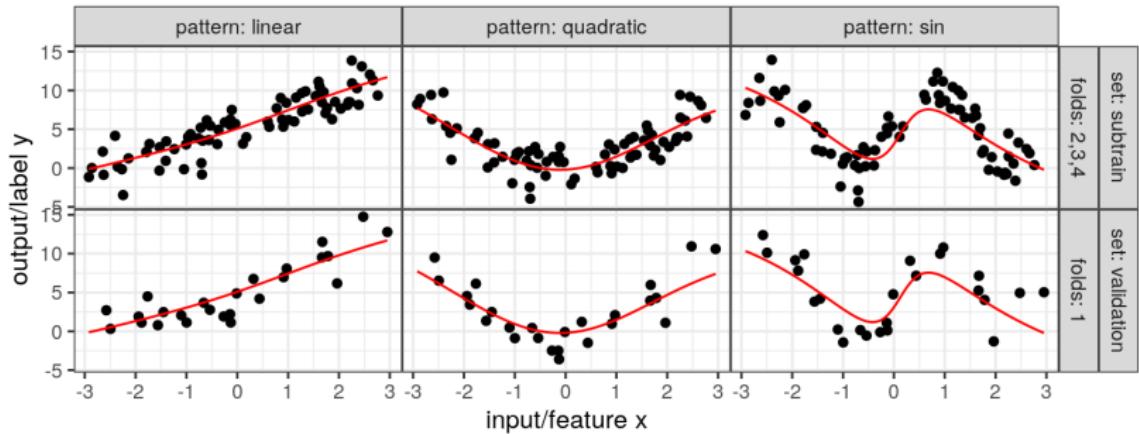
```
> coef(fit)
```

b->h1	i1->h1	b->h2	i1->h2
-40.770433	-2.636863	-20.719333	2.629485
b->o	h1->o	h2->o	
4.643852	18.973225	28.348342	

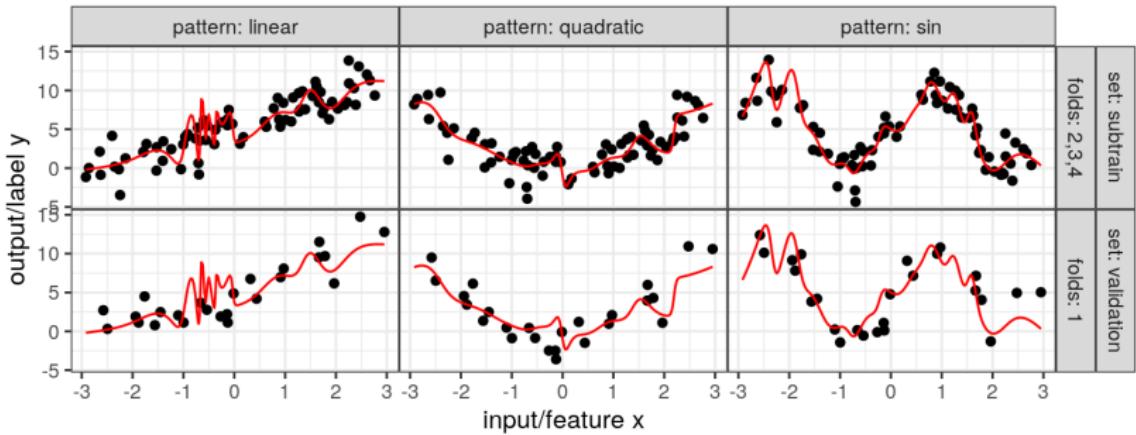
Neural network, 200 hidden units, 1 gradient descent iterations



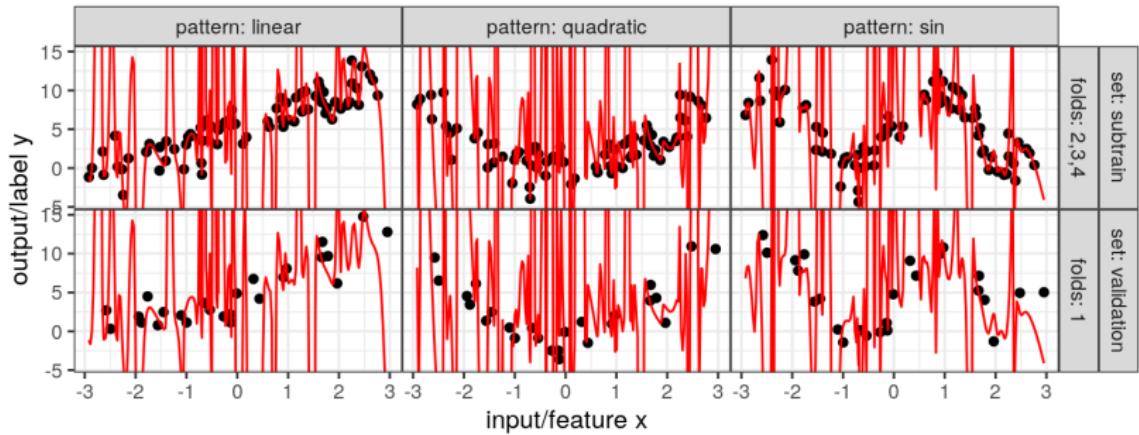
Neural network, 200 hidden units, 10 gradient descent iterations



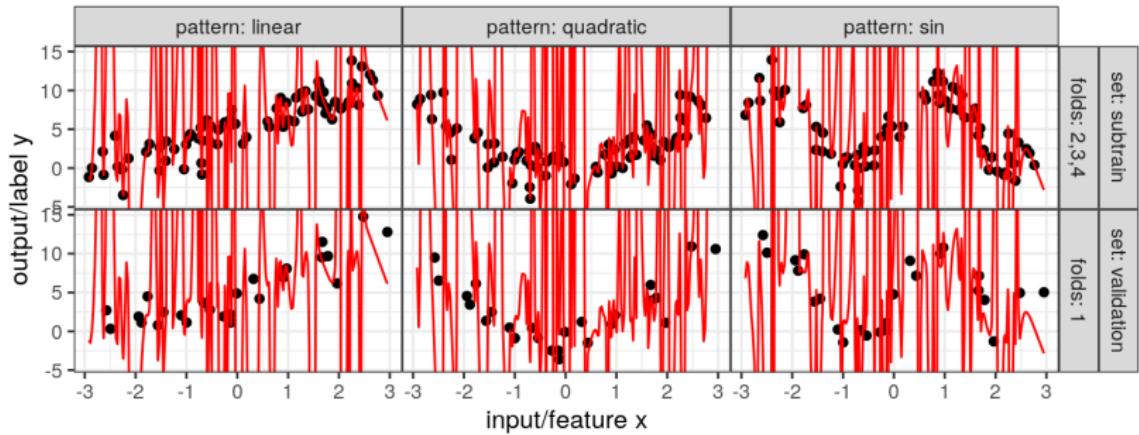
Neural network, 200 hidden units, 100 gradient descent iterations



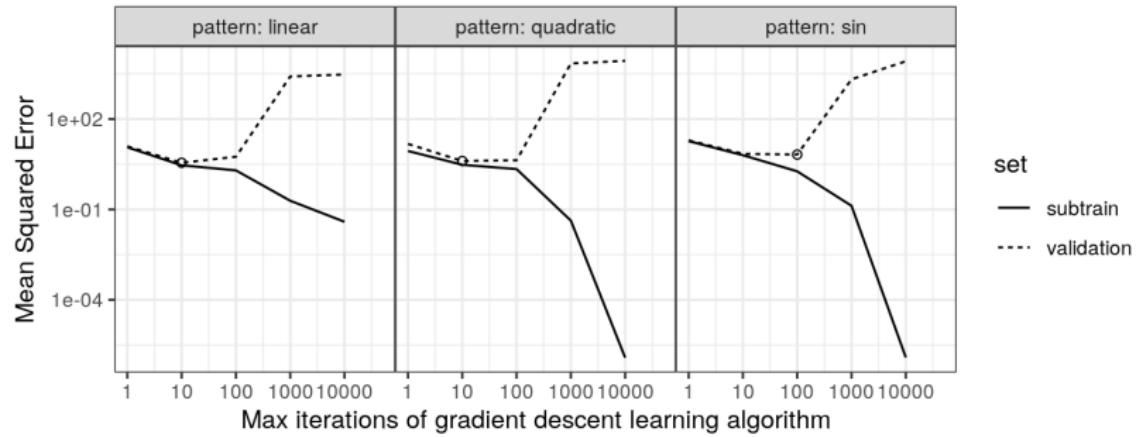
Neural network, 200 hidden units, 1000 gradient descent iterations



Neural network, 200 hidden units, 10000 gradient descent iterations



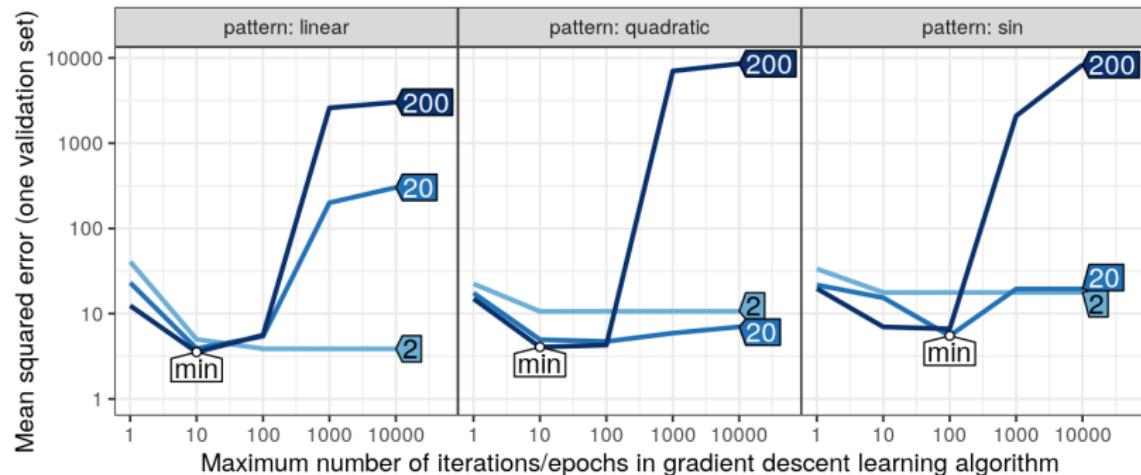
## Neural network, 200 hidden units



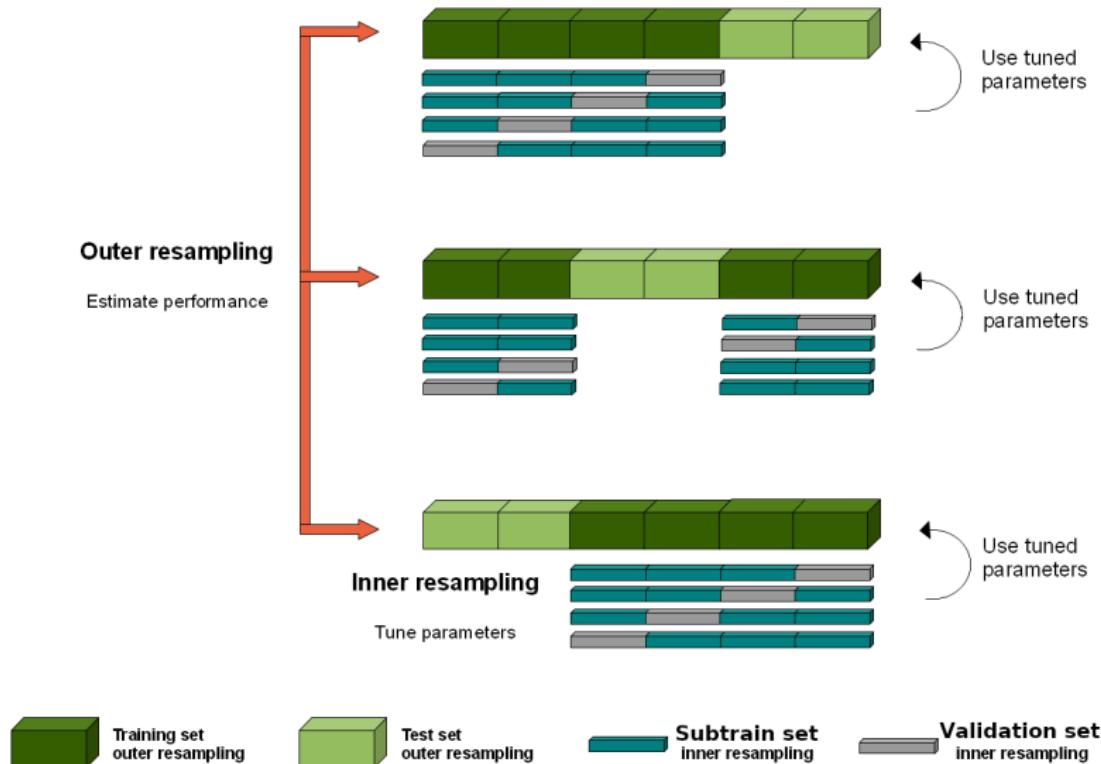
## Comments on other model sizes

- ▶ Model with 2 hidden units is less flexible: does not fit the subtrain data as well, and does not overfit as much.
- ▶ Model with 200 hidden units is more flexible: fits subtrain data even better, and overfits even more.
- ▶ Therefore model size / number of hidden units is also a hyper-parameter that can be used to control overfitting.
- ▶ R code is the same, just use `size=2` or `200` (or even better, I wrote a for loop over size values).
- ▶ Conclusion: you need to use a held-out validation set to choose the best values of hyper-parameters (e.g. number of iterations, hidden units).

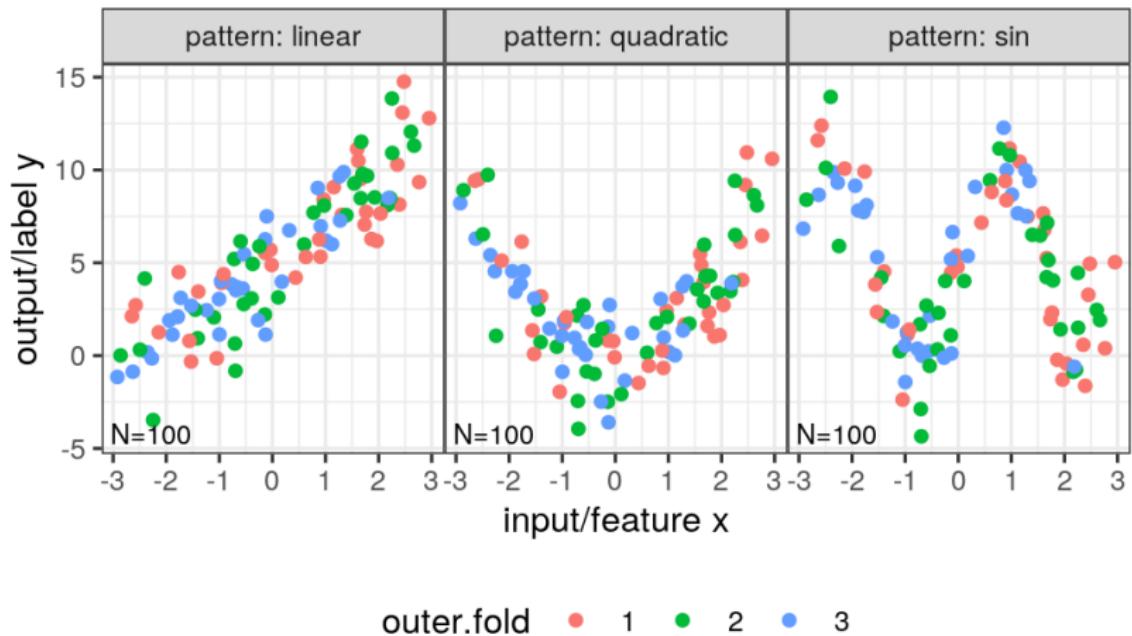
# Minimizing over both hyper-parameters



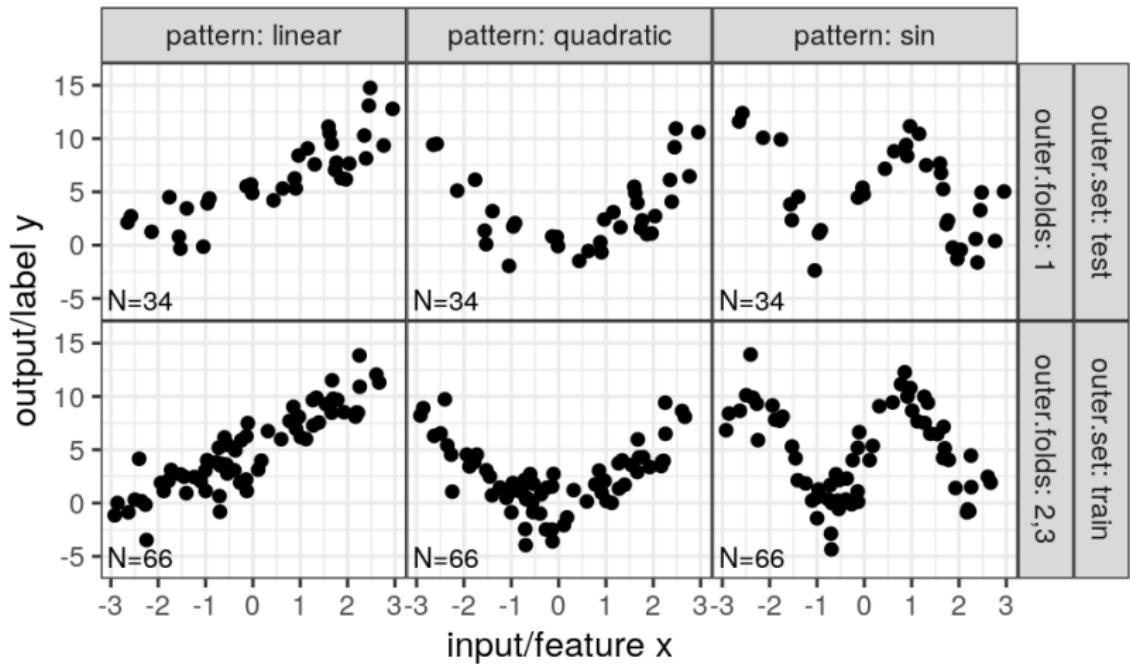
# Inner and outer cross-validation



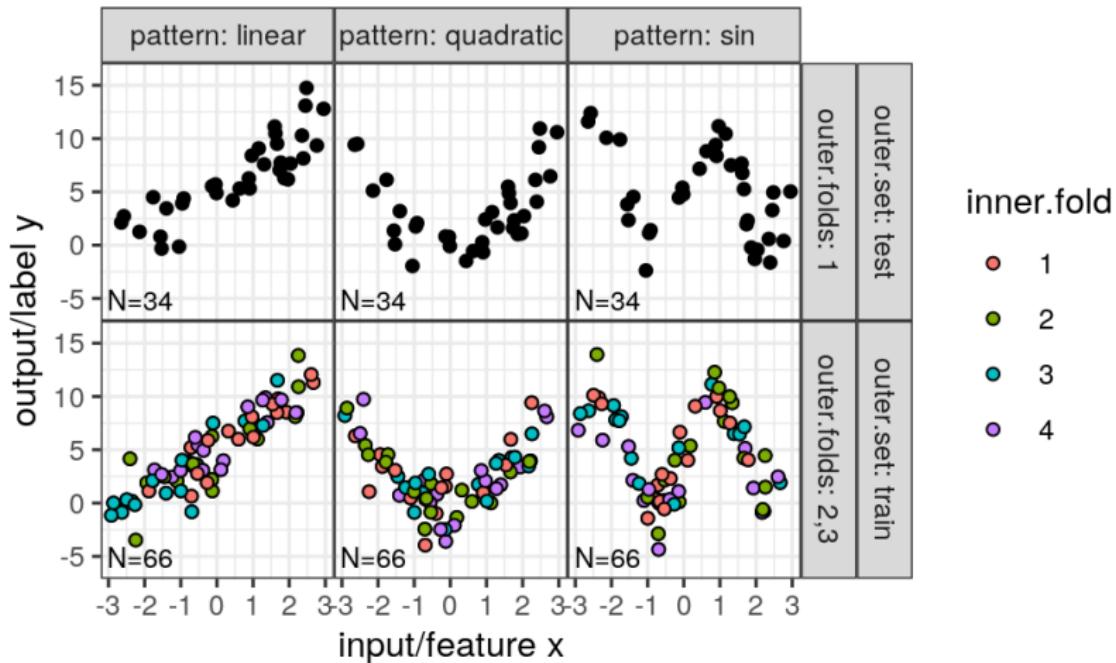
## Outer cross-validation



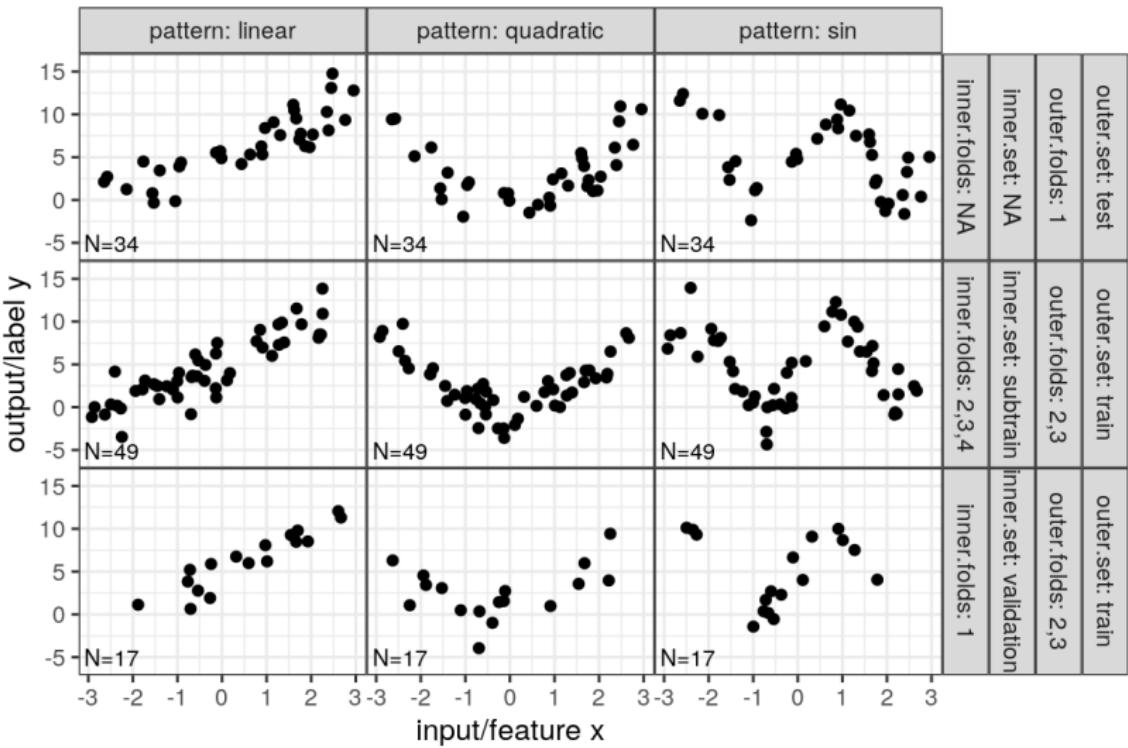
## Fix one outer fold = one train/test split



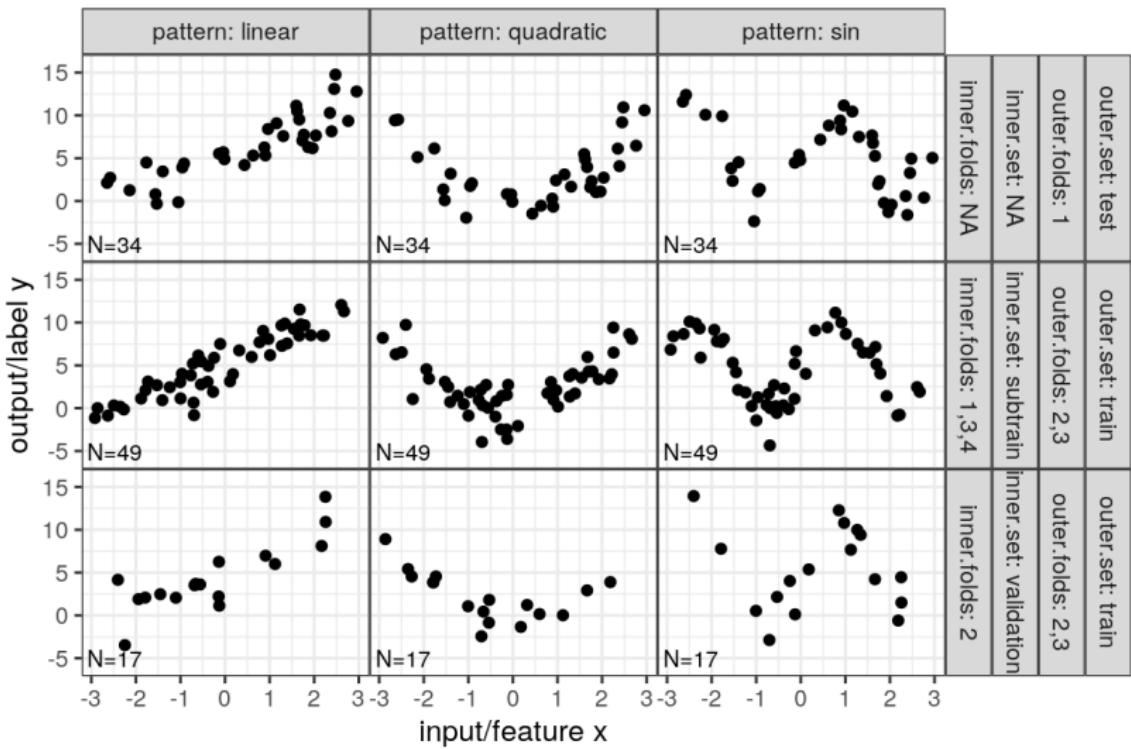
## Assign inner folds for one train set



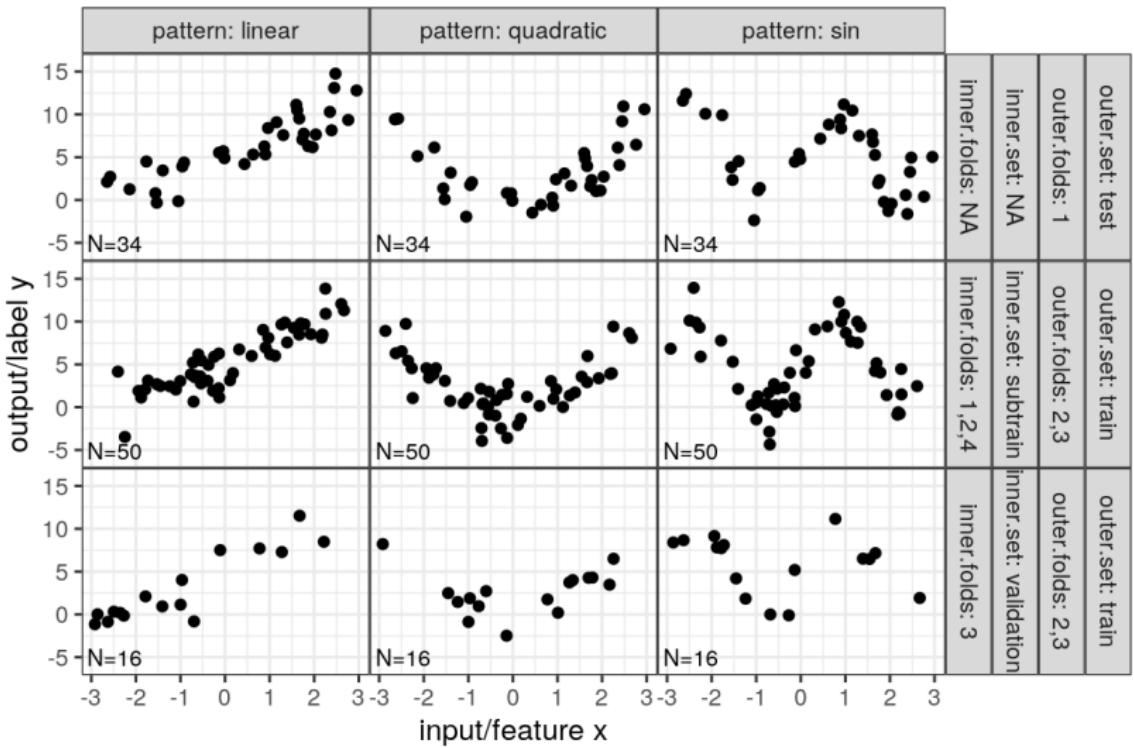
# Fix one inner fold = one subtrain/validation split



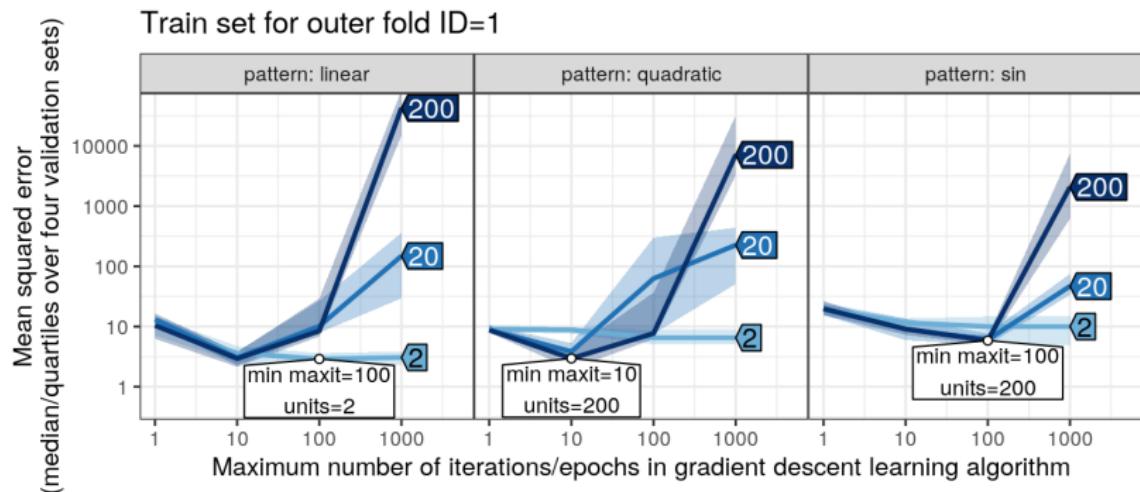
## Another subtrain/validation split



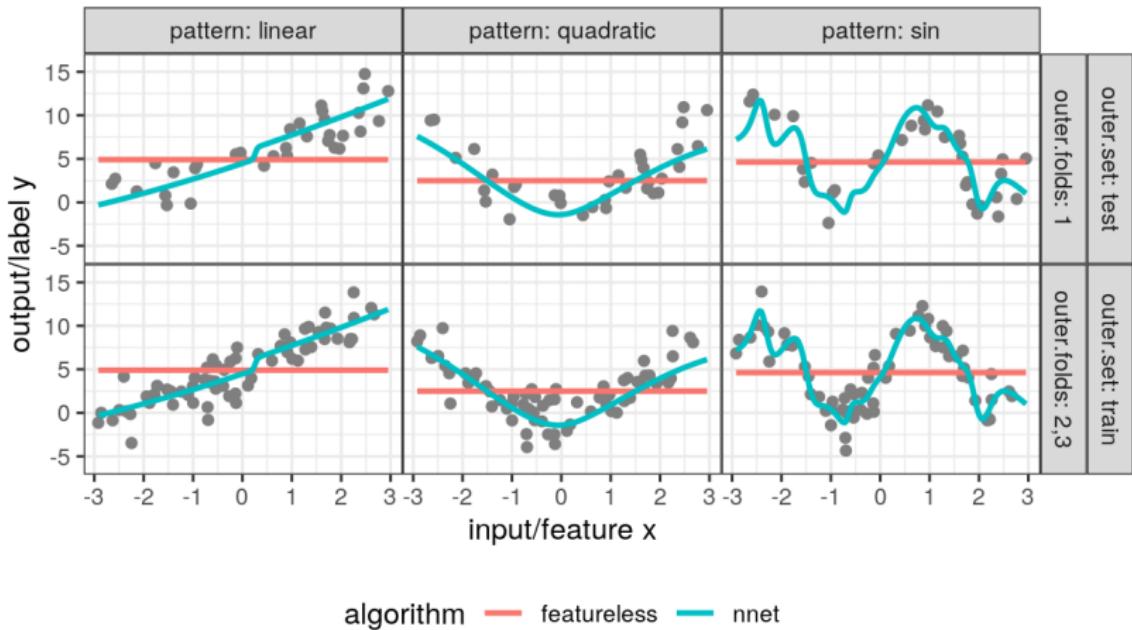
## Another subtrain/validation split



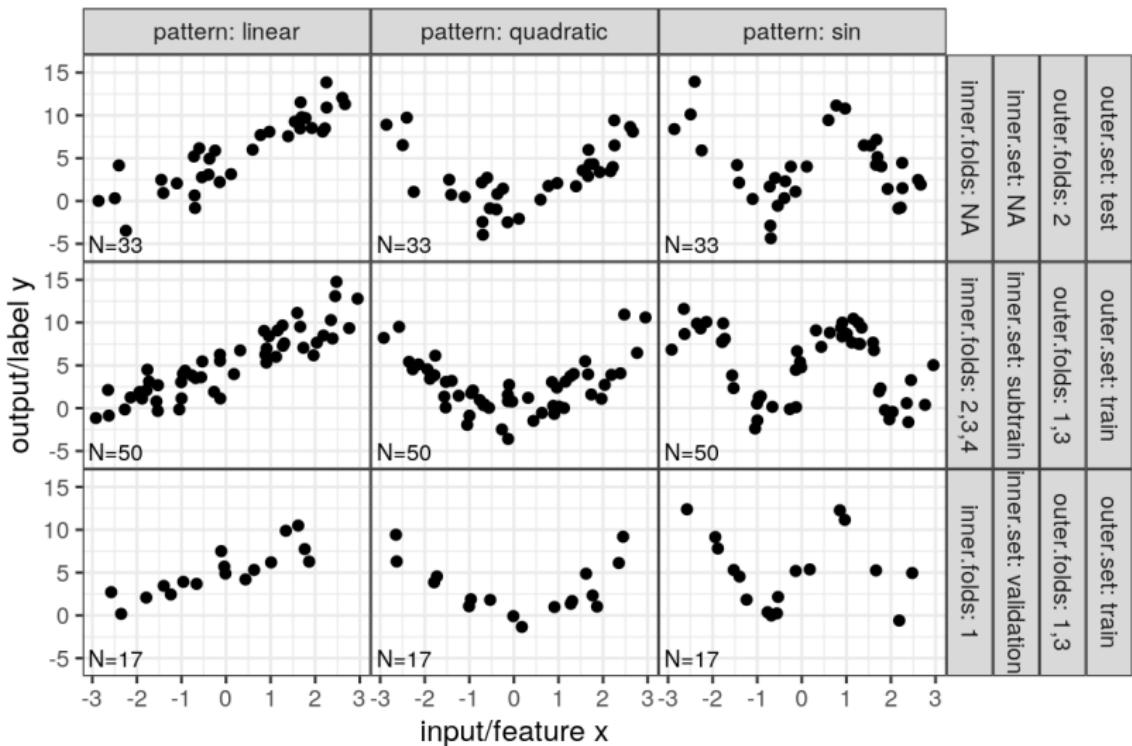
# Average validation error over four splits



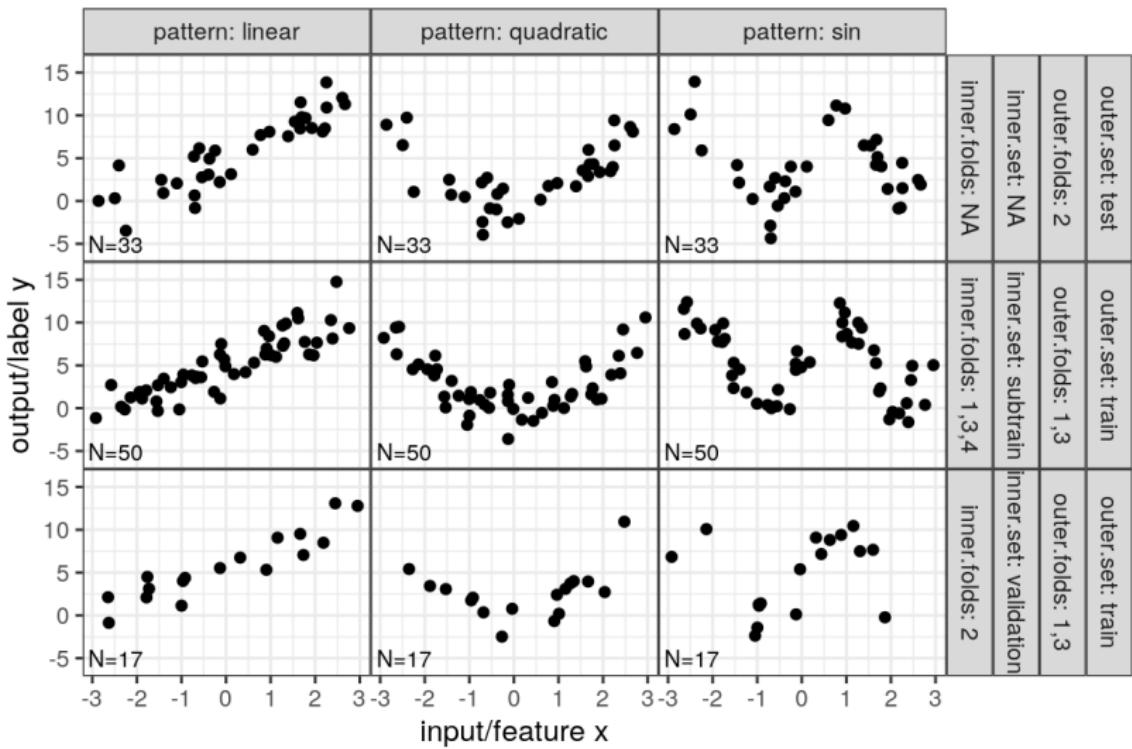
# Final models fit to entire train set using best hyper-parameters



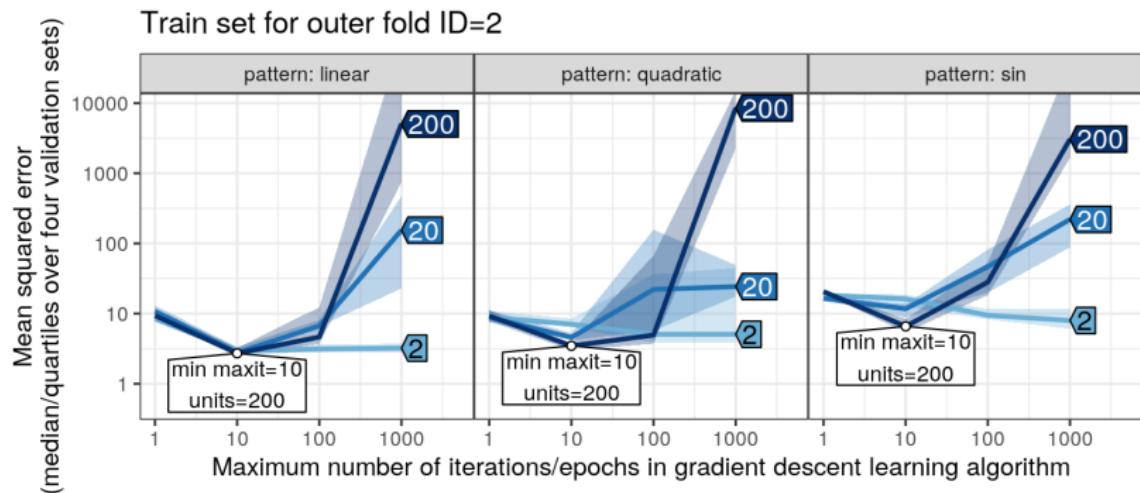
# Fix one inner fold = one subtrain/validation split



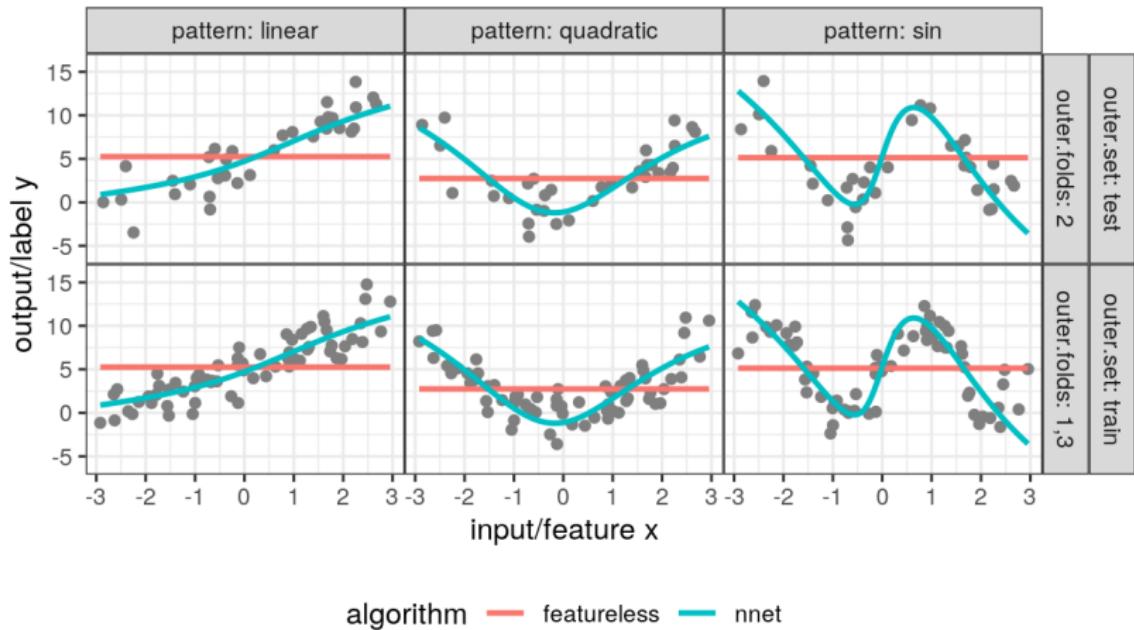
# Another subtrain/validation split



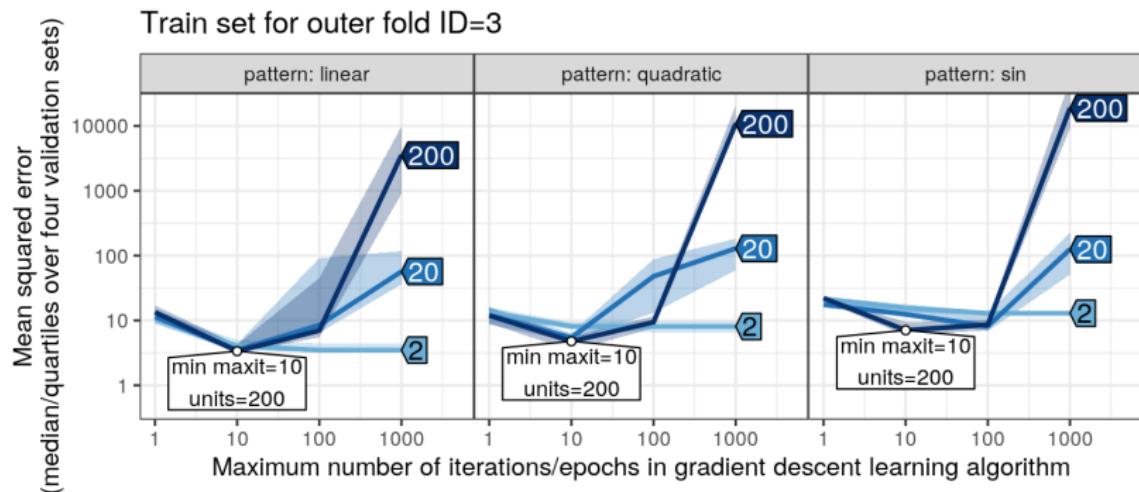
# Average validation error over four splits



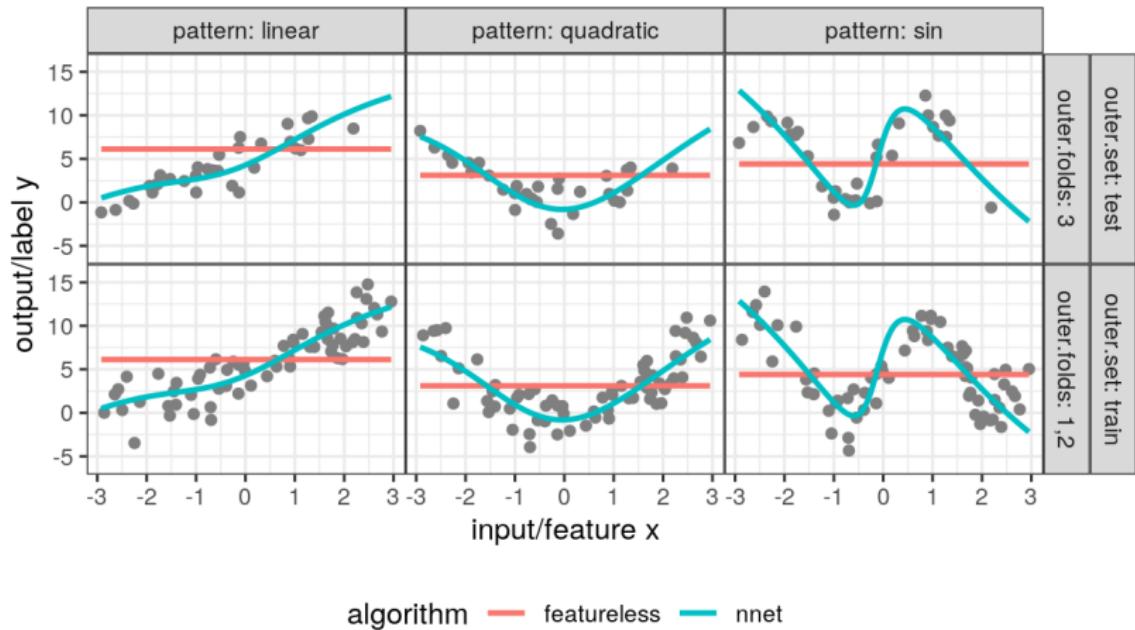
# Final models fit to entire train set using best hyper-parameters



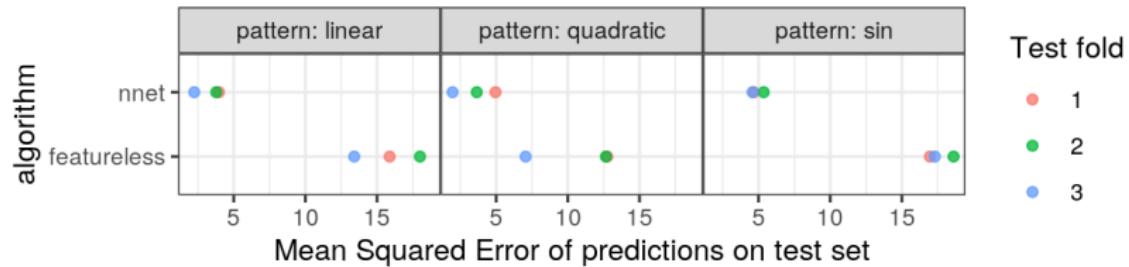
# Average validation error over four splits



# Final models fit to entire train set using best hyper-parameters



## Finally compare test error across all train/test splits



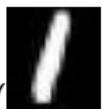
Introduction and applications

Demonstration of overfitting in regression / first steps with R

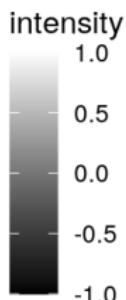
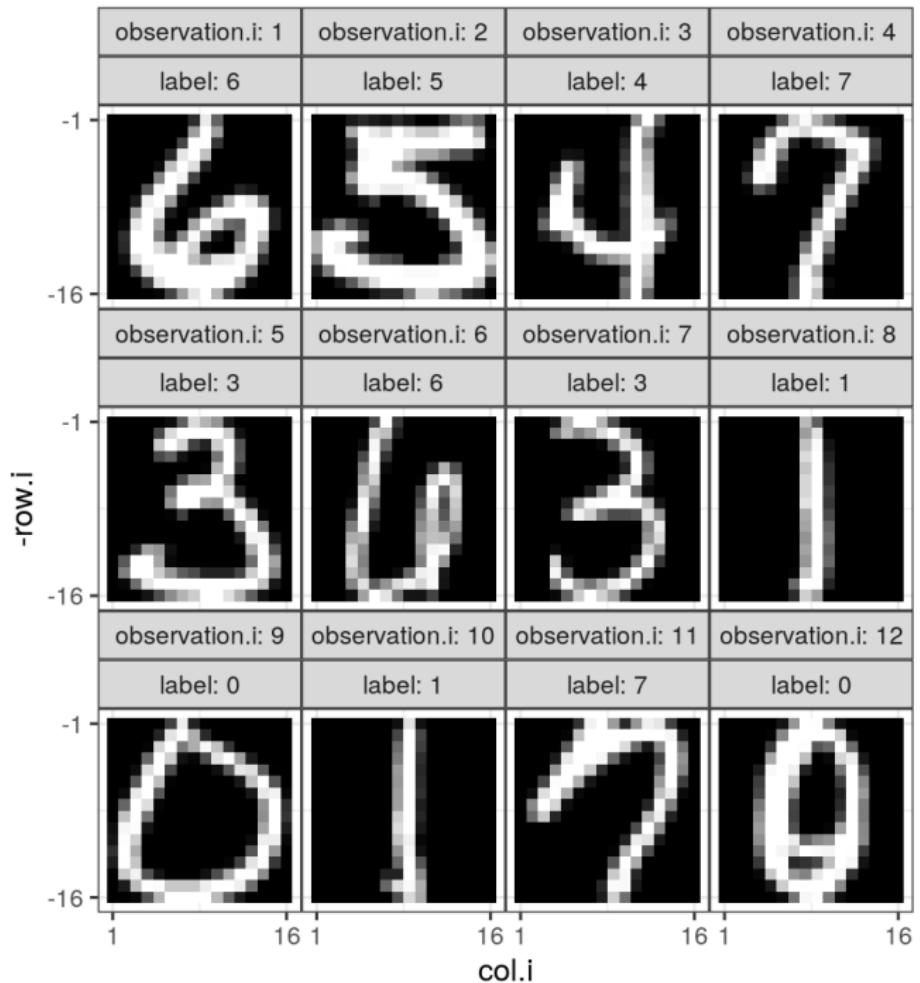
Classifying images of digits

# Image classification

- ▶ One of the most popular/successful applications of machine learning.
- ▶ Input: image file  $x \in \mathbb{R}^{h \times w \times c}$  where  $h$  is the height in pixels,  $w$  is the width,  $c$  is the number of channels, e.g. RGB image  $c = 3$  channels.
- ▶ In this tutorial we use images with  $h = w = 16$  pixels and  $c = 1$  channel (grayscale, smaller values are darker).
- ▶ Output: class/category  $y$  (from a finite set).
- ▶ In this tutorial there are ten image classes  $y \in \{0, 1, \dots, 9\}$ , one for each digit.



- ▶ Want to learn  $f$  such that  $f(\textcircled{0}) = 0$ ,  $f(\textcircled{1}) = 1$ , etc.
- ▶ Code for figures in this section: <https://github.com/tdhock/2020-yiqi-summer-school/blob/master/figure-validation-loss.R>



## Representation of digits in CSV

- ▶ Each image/observation is one row.
- ▶ First column is output/label/class to predict.
- ▶ Other 256 columns are inputs/features (pixel intensity values).

1:	6	-1	-1	...	-1.000	-1.000	-1
2:	5	-1	-1	...	-0.671	-0.828	-1
3:	4	-1	-1	...	-1.000	-1.000	-1
4:	7	-1	-1	...	-1.000	-1.000	-1
5:	3	-1	-1	...	-0.883	-1.000	-1
6:	6	-1	-1	...	-1.000	-1.000	-1
7:	3	-1	-1	...	-1.000	-1.000	-1
8:	1	-1	-1	...	-1.000	-1.000	-1
9:	0	-1	-1	...	-1.000	-1.000	-1
10:	1	-1	-1	...	-1.000	-1.000	-1
11:	7	-1	-1	...	-1.000	-1.000	-1
12:	0	-1	-1	...	-1.000	-1.000	-1

## Converting label column to matrix for neural network

This is a “one hot” encoding of the class labels.

```
zip.dt <- data.table::fread("zip.gz")
zip.y.mat <- keras::to_categorical(zip.dt$V1)
```

	0	1	2	3	4	5	6	7	8	9
[1,]	0	0	0	0	0	0	1	0	0	0
[2,]	0	0	0	0	0	1	0	0	0	0
[3,]	0	0	0	0	1	0	0	0	0	0
[4,]	0	0	0	0	0	0	0	1	0	0
[5,]	0	0	0	1	0	0	0	0	0	0
[6,]	0	0	0	0	0	0	1	0	0	0
[7,]	0	0	0	1	0	0	0	0	0	0
[8,]	0	1	0	0	0	0	0	0	0	0
[9,]	1	0	0	0	0	0	0	0	0	0
[10,]	0	1	0	0	0	0	0	0	0	0
[11,]	0	0	0	0	0	0	0	1	0	0
[12,]	1	0	0	0	0	0	0	0	0	0

## Conversion to array for input to neural network

Use array function with all columns except first as data.

```
zip.size <- 16  
zip.X.array <- array(  
  data = unlist(zip.dt[1:nrow(zip.dt), -1]),  
  dim = c(nrow(zip.dt), zip.size, zip.size, 1))
```

Need to specify dimensions of array:

- ▶ Observations: same as the number of rows in the CSV table.
- ▶ Pixels wide: 16.
- ▶ Pixels high: 16.
- ▶ Channels: 1 (greyscale image).

## Linear model R code

```
library(keras)
linear.model <- keras::keras_model_sequential() %>%
  keras::layer_flatten(
    input_shape = c(16, 16, 1)) %>%
  keras::layer_dense(
    units = 10,
    activation = 'softmax')
```

- ▶ First layer must specify shape of inputs (here 16x16x1).
- ▶ `layer_flatten` converts any shape to a single dimension of units (here 256).
- ▶ `layer_dense` uses all units in the previous layer to predict each unit in the layer.
- ▶ `units=10` because there are ten possible classes for an output.
- ▶ `activation='softmax'` is required for the last/output layer in multi-class classification problems.

## Keras model compilation

```
linear.model %>% keras::compile(  
  loss = keras::loss_categorical_crossentropy,  
  optimizer = keras::optimizer_adadelta(),  
  metrics = c('accuracy'))
```

In compile you can specify

- ▶ a loss function, which is directly optimized/minimized in each iteration of the gradient descent learning algorithm.  
<https://keras.io/api/losses/>
- ▶ an optimizer, which is the version of gradient descent learning algorithm to use.  
<https://keras.io/api/optimizers/>
- ▶ an evaluation metric to monitor, not directly optimized via gradient descent, but usually more relevant/interpretable for the application (e.g. accuracy is the proportion of correctly predicted labels). <https://keras.io/api/metrics/>

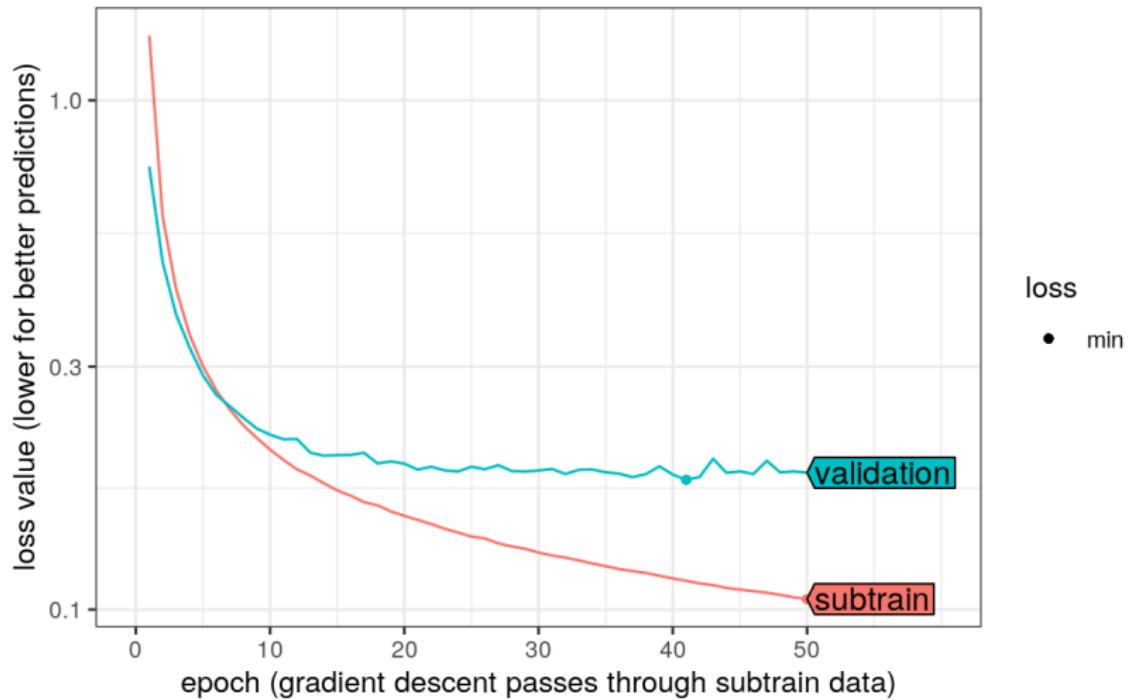
## Keras model fitting

```
linear.model %>% keras::fit(  
  zip.X.array, zip.y.mat,  
  epochs = 50,  
  validation_split = 0.2  
)
```

In fit you can specify

- ▶ Train data inputs `zip.X.array` and outputs `zip.y.mat` (required).
- ▶ Number of full passes of gradient descent through the subtrain data (`epochs`). In each epoch the gradient with respect to each subtrain observation is computed once.
- ▶ `validation_split=0.2` which means to use 80% subtrain (used for gradient descent parameter updates), 20% validation (used for hyper-parameter selection).

## Linear model

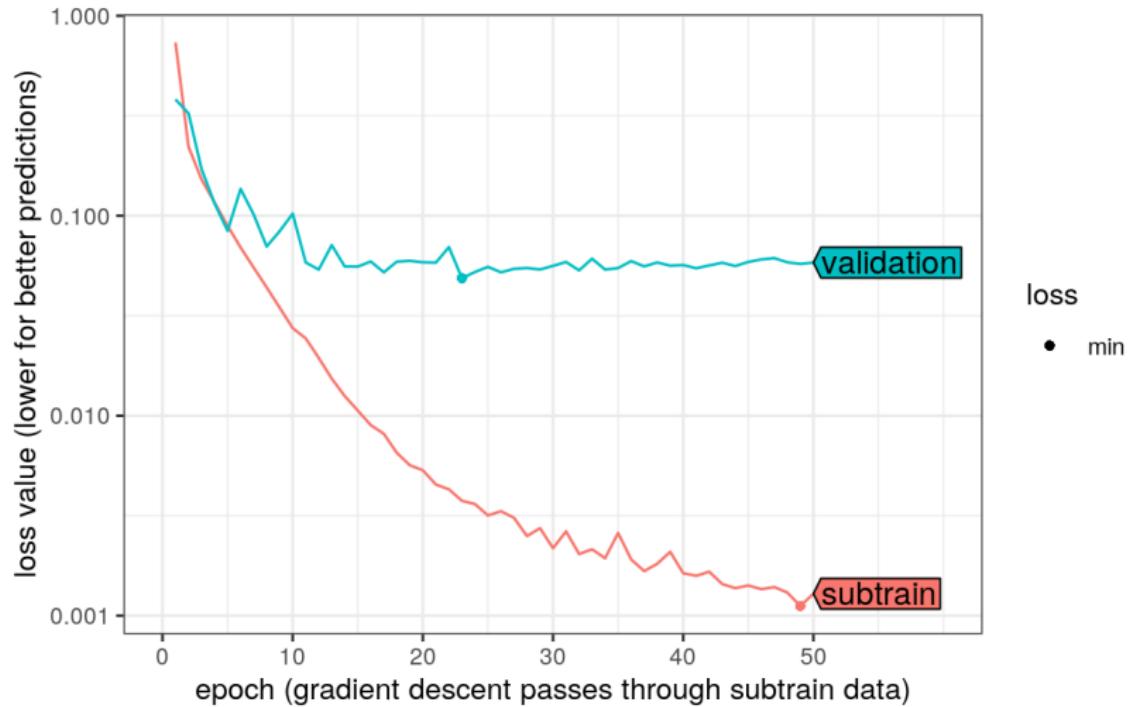


## Sparse (convolutional) model R code

```
library(keras)
conv.model <- keras_model_sequential() %>%
  layer_conv_2d(
    input_shape = dim(zip.X.array)[-1] ,
    filters = 20,
    kernel_size = c(3,3),
    activation = 'relu') %>%
  layer_max_pooling_2d(pool_size = c(2, 2)) %>%
  layer_flatten() %>%
  layer_dense(units = 100, activation = 'relu') %>%
  layer_dense(
    units = ncol(zip.y.mat),
    activation = 'softmax')
```

- ▶ Sparse: few inputs are used to predict each unit in `layer_conv_2d`.
- ▶ Exploits structure of image data to make learning easier/faster.

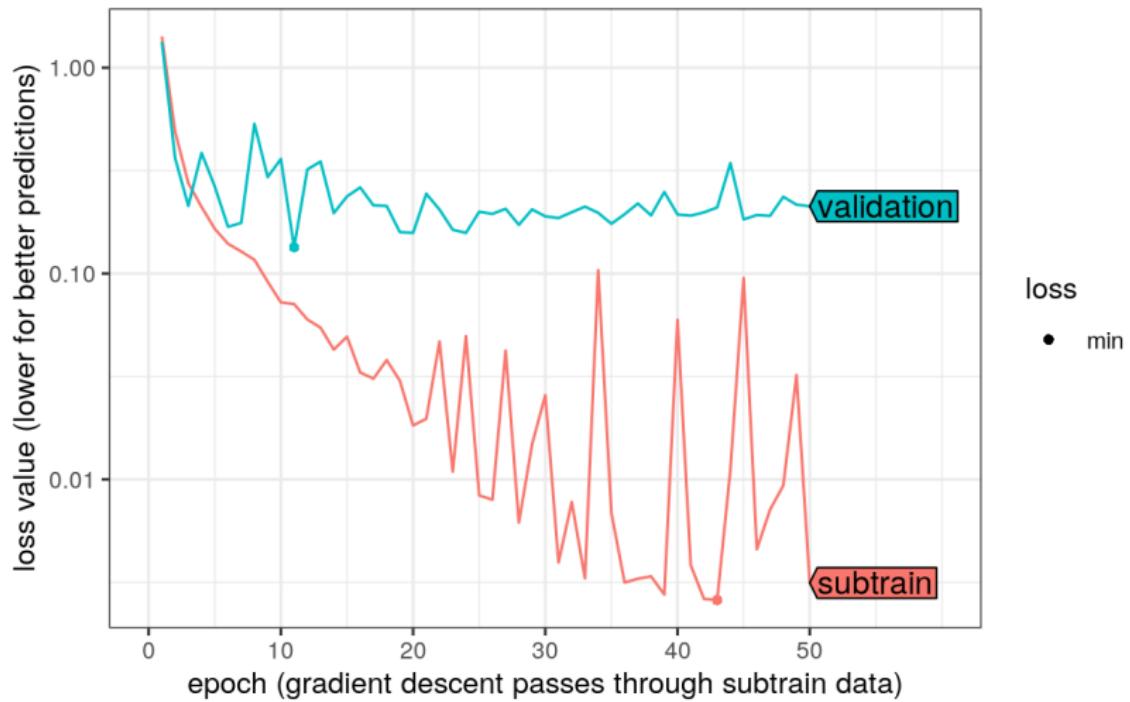
## Convolutional neural network

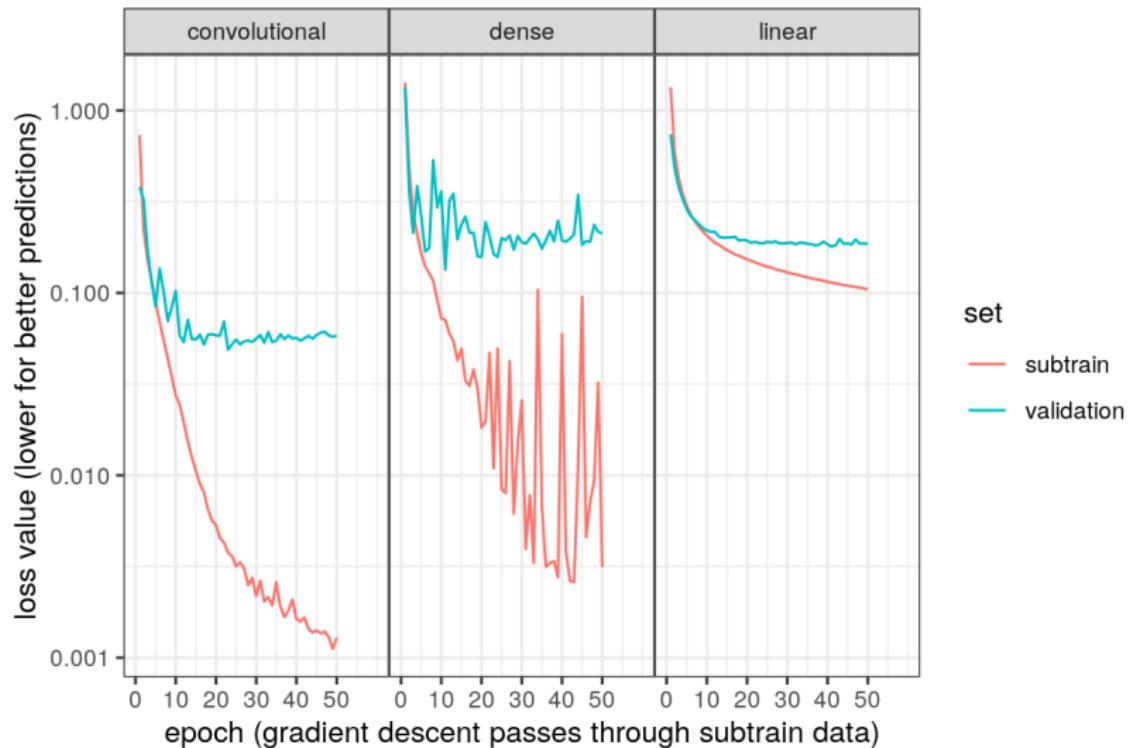


## Dense (fully connected) neural network R code

```
library(keras)
dense.model <- keras_model_sequential() %>%
  layer_flatten(
    input_shape = dim(zip.X.array)[-1]) %>%
  layer_dense(units = 100, activation = 'relu') %>%
  layer_dense(
    units = ncol(zip.y.mat),
    activation = 'softmax')
```

## Dense (fully connected) neural network with 8 hidden layers





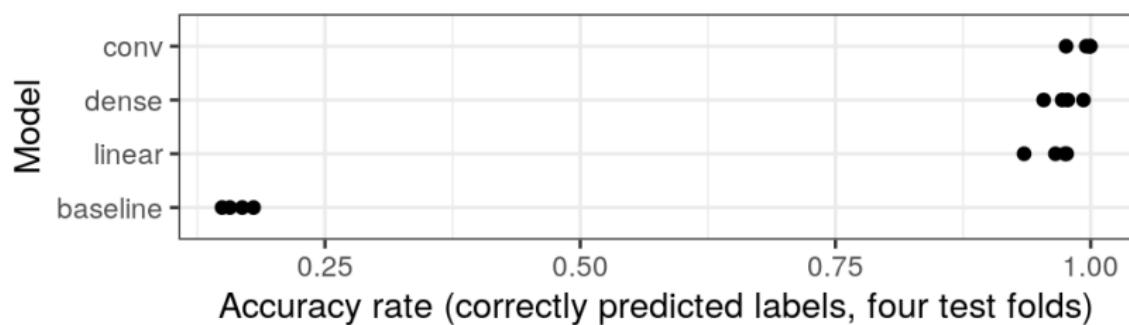
## 4-fold cross-validation for model evaluation

Does the convolutional model provide more accurate predictions on unseen test data? First randomly assign a fold ID to each image/observation, then for each test fold ID from 1 to 4:

- ▶ Hold out the images/observations with the test fold ID as a test set (not used at all for model training).
- ▶ Use the other images/observations to train all three models using `validation_split=0.2`.
- ▶ Plot the validation loss curve as a function of the number of epochs, and select the number of epochs which minimizes the validation loss (hyper-parameter learning).
- ▶ Re-train with `epochs`=the learned number of epochs and `validation_split=0` (all train data used for gradient descent parameter updates).
- ▶ Finally compute the prediction accuracy with respect to the held-out test set.

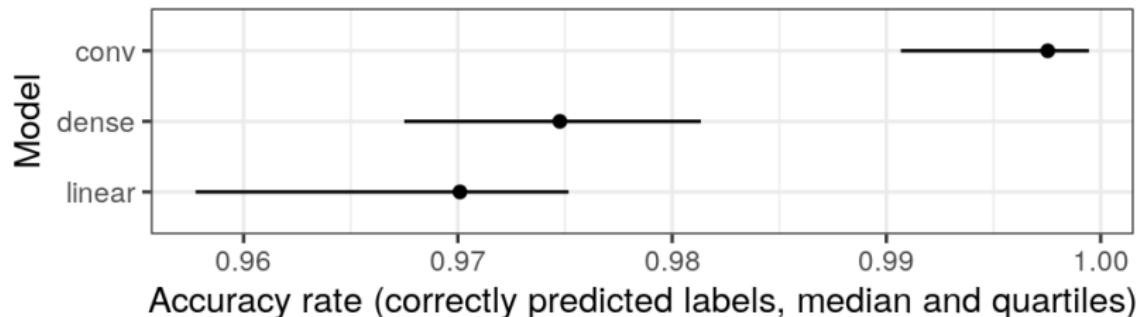
Plot accuracy values (or mean/sd) after learning/testing four times (one for each test fold).

## Accuracy rates for each test fold



- ▶ Always a good idea to compare with the trivial baseline model which always predicts the most frequent class in the train set. (ignoring all inputs/features)
- ▶ Here we see that the baseline is much less accurate than the three learned models, so they are clearly learning something non-trivial.
- ▶ Code for test accuracy figures: <https://github.com/tdhock/2020-yiqi-summer-school/blob/master/figure-test-accuracy.R>

## Zoom to learned models



- ▶ Dense neural network slightly more accurate than linear model, convolutional significantly more accurate than others.
- ▶ Conclusion: convolutional neural network should be preferred for most accurate predictions in these data.
- ▶ Maybe not the same conclusion in other data sets, with the same models. (always need to do cross-validation experiments to see which model is best in any given data set)
- ▶ Maybe other models/algorithms would be even more accurate in these data. (more/less layers, more/less units, completely different algorithm such as random forests, boosting, etc)