

# Introduction to HPC Applications, Systems, Programming Models & Machine Learning & Data Analytics – Part Two

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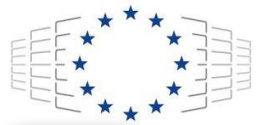
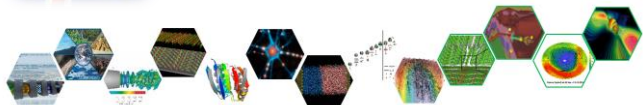
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# Introduction to HPC Applications, Systems, Programming Models – Part One

Dr. – Ing.  
Bernd Mohr



@ Supercomputing 2017, Denver, USA:  
Bernd Mohr first EU Chair of SC series

***Importance of our Community Building:***

***Attending & Participating at our community conferences like SC @USA or ISC @ Europe is crucial for learning, networking, sharing and building your career over time – world-wide experts become mentors & friends!***

# Outline – Part Two

- All HPC scripts will be available with ISC Online Material after the event
- All source & data free to use

## Machine Learning Fundamentals

- Learning Methods Overview, Prerequisites, Classification Application & Linear Perceptron Model
- Training & Testing Process using different Datasets, Food Inspection Classification Application Example
- Linear Regression Model & Logistic Regression Model

## Artificial Neural Network (ANN) Basics

- Handwritten Character Recognition MNIST Dataset & Understanding Multi-Class Classification Approach
- Limits of the Perceptron Learning Model, Multi-Output Perceptron Model & ANNs with Backpropagation
- Observe Growth of Trainable Parameter & Understanding Overfitting

## Convolutional Neural Network (CNN) Basics

- Moving from Shallow Learning to Deep Learning, MNIST Application Example with CNNs in Keras
- Understanding Feature Maps in CNN Architecture, Hyperparameter Complexity & Adam Optimizer
- Understanding Accuracy Improvements & Limits

## Selected Parallel & Scalable Machine & Deep Learning Techniques

- piSVM MPI Implementation & Remote Sensing Applications
- Computing Footprint in Training, Testing & Validation Methods, Distributed Training
- Parallel & Scalable HPDBSCAN for Data Clustering & Quantum Machine Learning, Hyperparameter tuning





# More Information: Full HPC Spring 2023 University Course



Prof Dr - Ing Morris Riedel

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## Selected Testimonials from our community:



'... I have found your wonderful HPC lectures on YouTube, and I am finding them very helpful for learning the supercomputing tools necessary for my research...'

- student in computational nuclear physics from USA



neeraj kumar · 1 year ago

Awesome work for HPC students....thanks



纵横10分钟 - 10 Minutes Across the World · 3 months ago 5 subscribers

Thanks so much for the lecture - One of the best courses. Very helpful!



Andy Gill @\_andy\_gill · Nov 15, 2021

Highly recommend this excellent introduction to state of the art HPC programming by @MorrisRiedel. Enough details to actually be useful, explained clearly using many examples. Thank you for posting it!

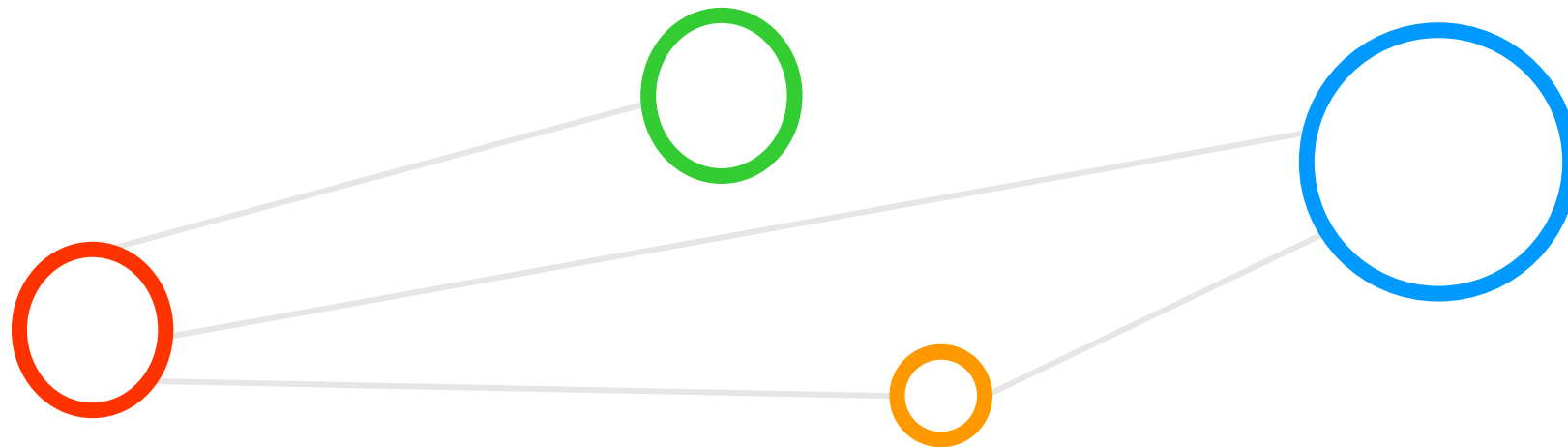


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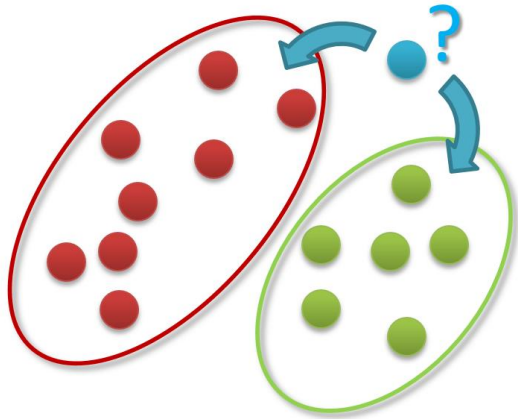


# Machine Learning Fundamentals



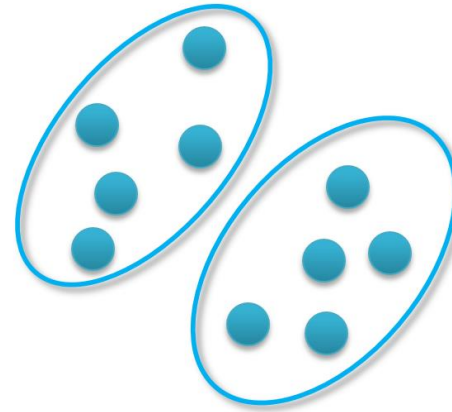
# Machine Learning Models – Short Overview & Introduction to Classification

## Classification



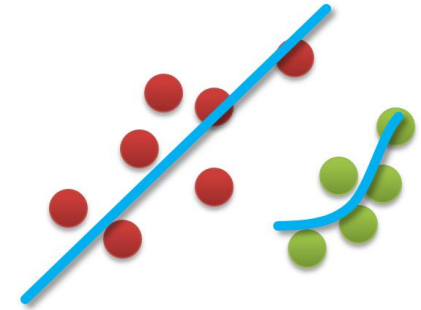
- Groups of data exist
- New data classified to existing groups

## Clustering



- No groups of data exist
- Create groups from data close to each other

## Regression



- Identify a line with a certain slope describing the data

▪ Machine learning methods can be roughly categorized in classification, clustering, or regression augmented with various techniques for data exploration, selection, or reduction – despite the momentum of deep learning, traditional machine learning algorithms are still widely relevant today

[1] [www.big-data.tips](http://www.big-data.tips), 'Data Classification'



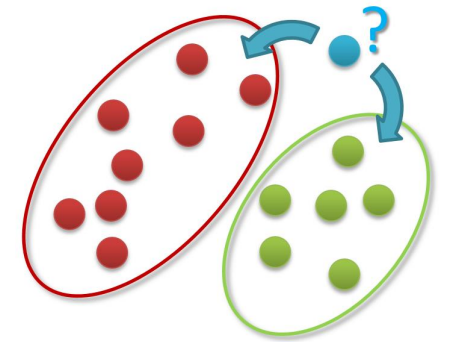
# Classification Machine Learning Model – Supervised Learning Example

- Each observation of the predictor measurement(s) has **an associated response measurement**:
  - Input  $\mathbf{x} = x_1, \dots, x_d$
  - Output  $y_i, i = 1, \dots, n$
  - Data  $(\mathbf{x}_1, y_1), \dots, (\mathbf{x}_N, y_N)$
  - (the output guides the learning process as a ‘supervisor’)
- Goal: Fit a model that relates the response to the predictors
  - **Prediction**: Aims of accurately predicting the response for future observations
  - **Inference**: Aims to better understanding the relationship between the response and the predictors

- Supervised learning approaches fits a model that related the response to the predictors
- Supervised learning approaches are used in classification algorithms such as SVMs
- Supervised learning works with data = [input, correct output]



## Classification



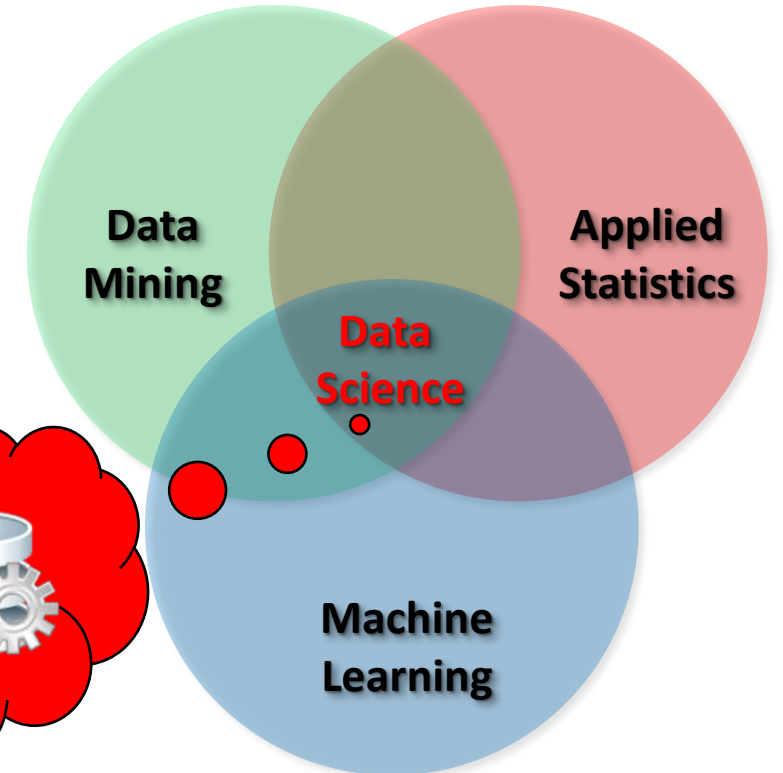
$(\mathbf{x}_1, y_1), \dots, (\mathbf{x}_N, y_N)$

# Machine Learning Prerequisites & Computing Challenges – Revisited

1. Some pattern exists
2. No exact mathematical formula
3. Data exists

- Idea ‘Learning from Big Data’
  - Shared with a wide variety of other disciplines
  - E.g. signal processing, big data data mining, etc.

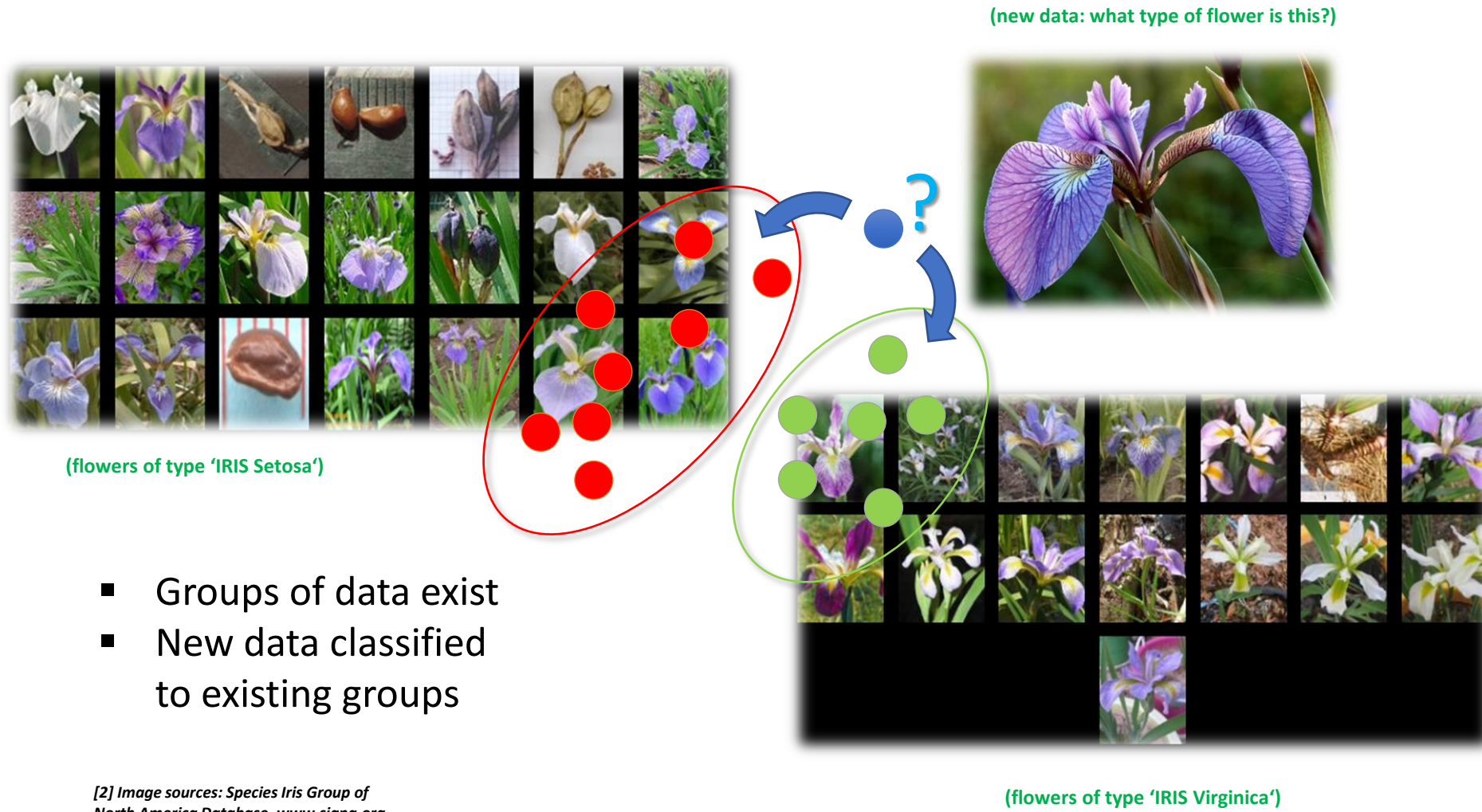
- Challenges
  - Data is often complex
  - Requires ‘Big Data analytics’
  - Learning from data requires processing time → Clouds or High Performance Computing



- Machine learning is a very broad subject and goes from very abstract theory to extreme practice ('rules of thumb')
- Training machine learning models needs processing time (clouds or high performance computing)
- While data analysis is more describing the process of analysis in the data, the term data analytics also includes and the necessary scalable or parallel infrastructure to perform analysis of 'big data'



# Simple Application Example: Classification of a Flower



[2] Image sources: Species Iris Group of North America Database, [www.signa.org](http://www.signa.org)

# The Learning Problem in the Example

(flowers of type 'IRIS Setosa')



(flowers of type 'IRIS Virginica')



[2] Image sources: Species Iris Group of North America Database, [www.signa.org](http://www.signa.org)

Learning problem: A prediction task

- Determine whether a new Iris flower sample is a “Setosa” or “Virginica”
- Binary (two class) classification problem
- What attributes about the data help?



(what type of flower is this?)



# Feasibility of Machine Learning in this Example

## 1. Some pattern exists:

- Believe in a 'pattern with 'petal length' & 'petal width' somehow influence the type

## 2. No exact mathematical formula

- To the best of our knowledge there is no precise formula for this problem

## 3. Data exists

- Data collection from UCI Dataset „Iris“
- 150 labelled samples (aka 'data points')
- Balanced: 50 samples / class

[3] UCI Machine Learning  
Repository Iris Dataset



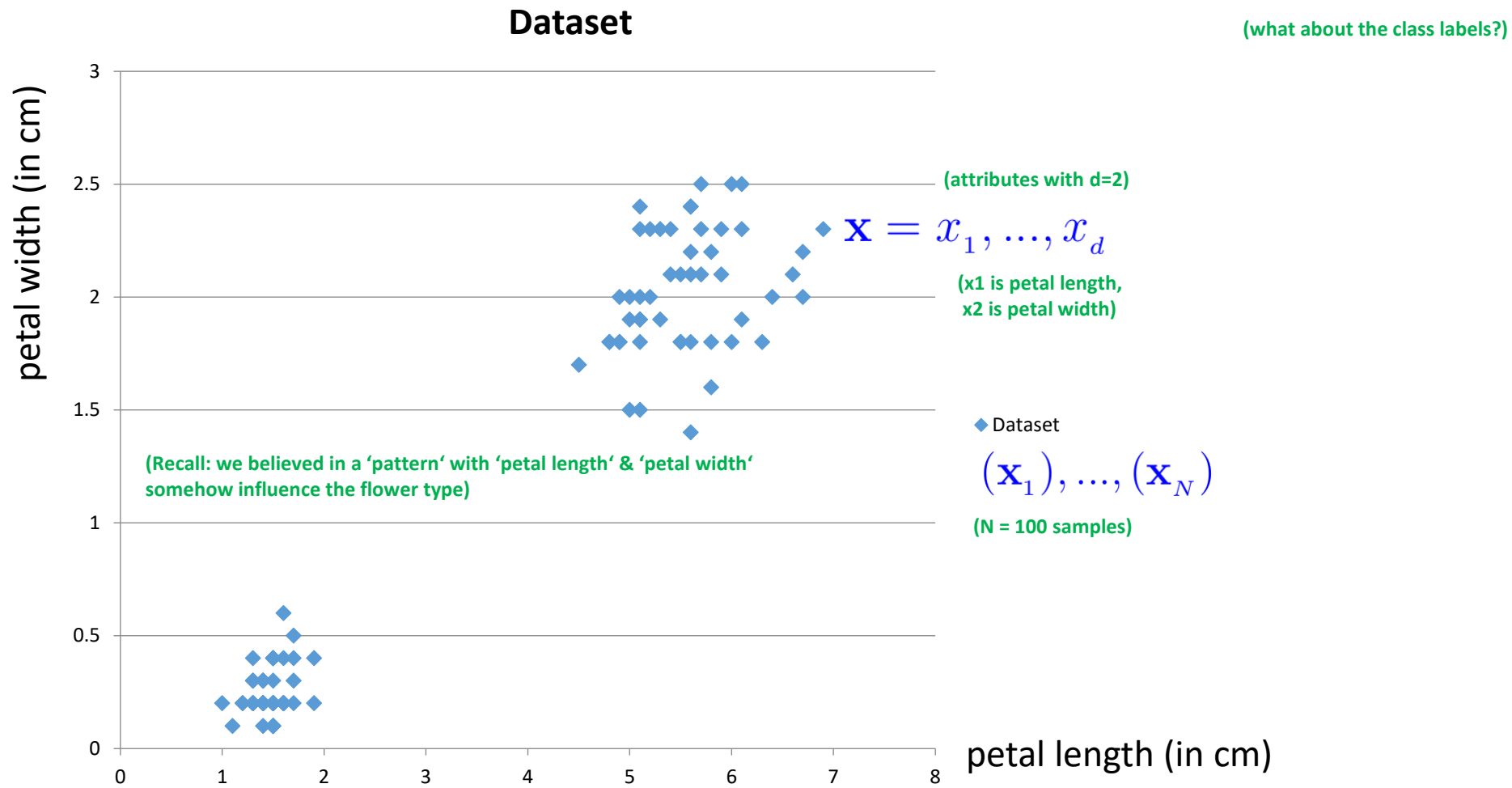
[4] Image source: Wikipedia, Sepal

(four data attributes for each  
sample in the dataset)

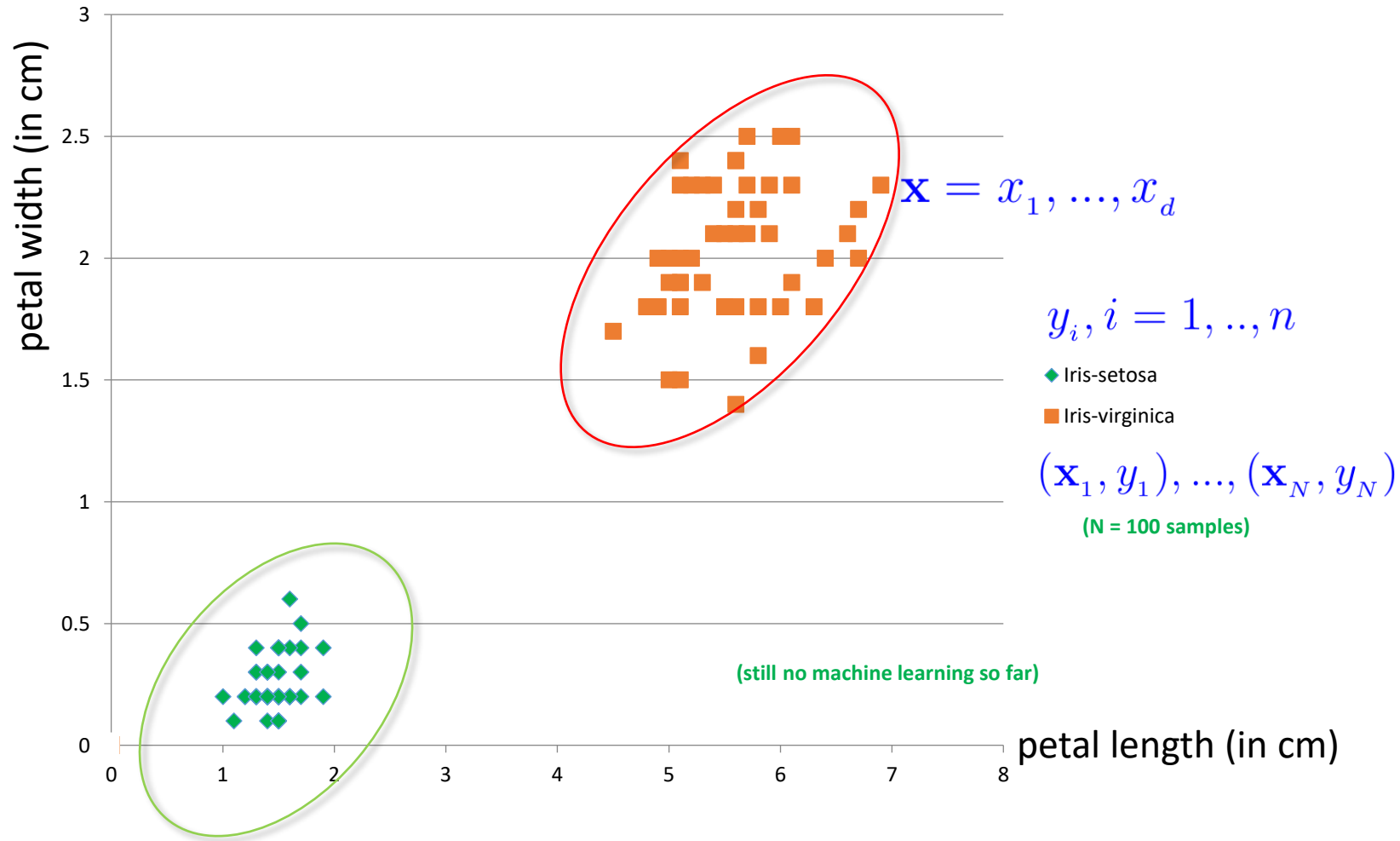
(one class label for each  
sample in the dataset)

- sepal length in cm
- sepal width in cm
- petal length in cm
- petal width in cm
- class: Iris Setosa, or Iris Versicolour, or Iris Virginica

# Check Preparation Phase: Plotting the Data (Two Classes)

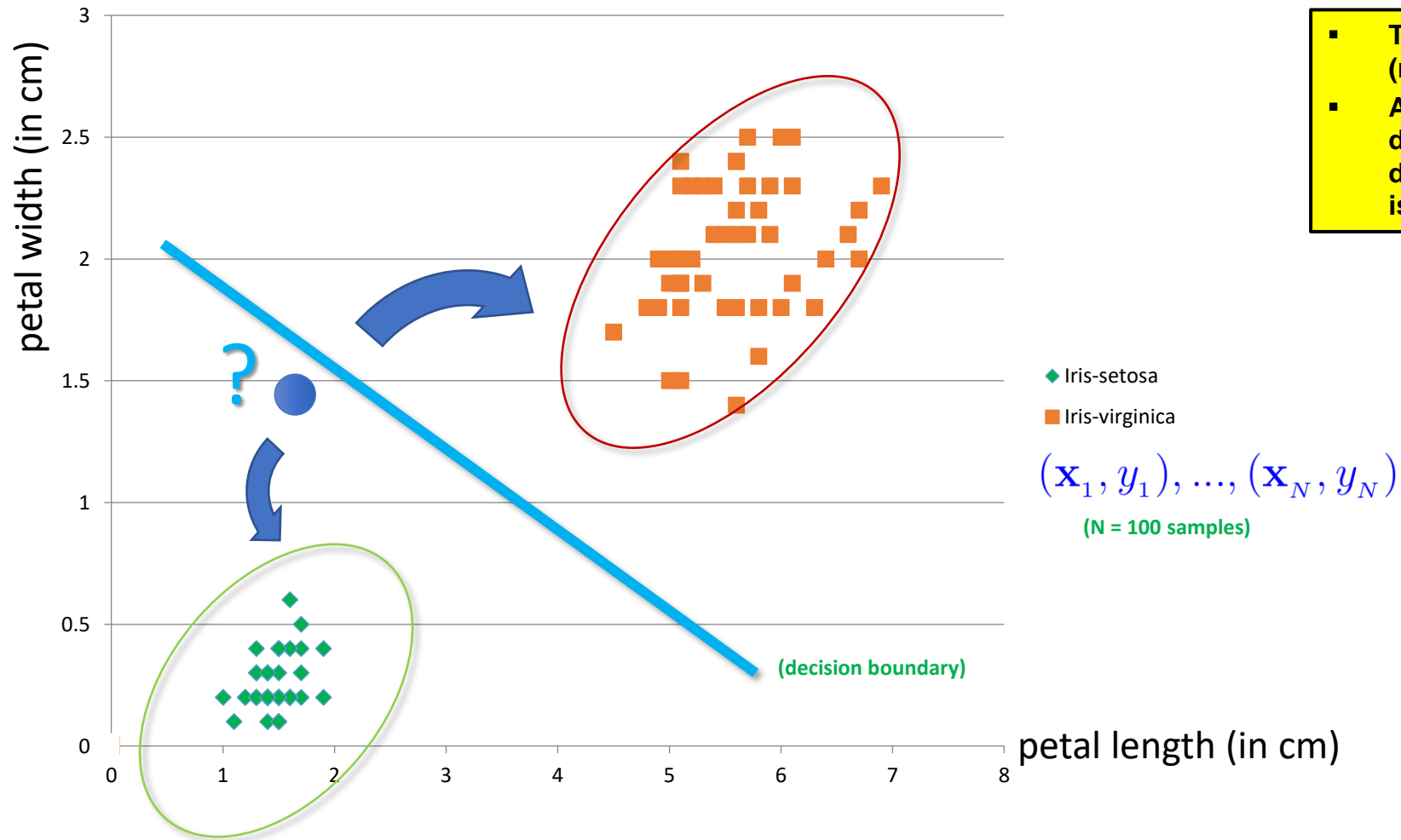


# Check Preparation Phase: Class Labels





# Linearly Seperable Data & Linear Decision Boundary



- The data is linearly seperable (rarely in practice)
- A line becomes a decision boundary to determine if a new data point is class red/green

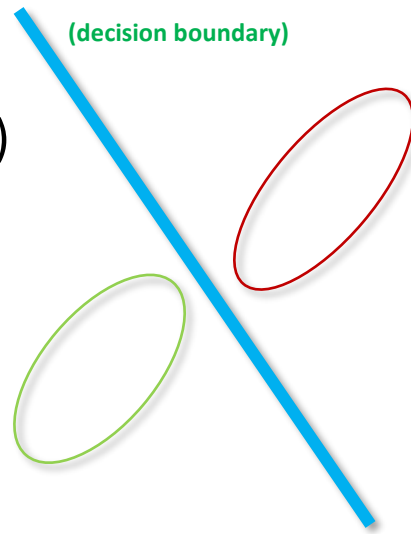
# Separating Line & Mathematical Notation

- Data exploration results
  - A line can be crafted between the classes since linearly separable data
  - All the data points representing Iris-setosa will be below the line
  - All the data points representing Iris-virginica will be above the line

- More formal mathematical notation

- Input:  $\mathbf{x} = x_1, \dots, x_d$  (attributes of flowers)

- Output:  
class +1 (Iris-virginica)  
or class -1 (Iris-setosa)



Iris-virginica if  $\sum_{i=1}^d w_i x_i > threshold$

Iris-setosa if  $\sum_{i=1}^d w_i x_i < threshold$

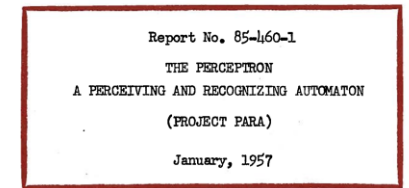
( $w_i$  and threshold are still unknown to us)

$$sign \left( \left( \sum_{i=1}^d w_i x_i \right) - threshold \right) \quad \text{(compact notation)}$$

# A Simple Linear Learning Model – The Perceptron

- Human analogy in learning
  - Human brain consists of nerve cells called **neurons**
  - Human brain learns by changing the **strength of neuron connections ( $w_i$ )** upon **repeated stimulation** by the same impulse (aka a **'training phase'**)
  - Training a perceptron model means adapting the weights  $w_i$
  - Done **until they fit input-output relationships** of the given **'training data'**

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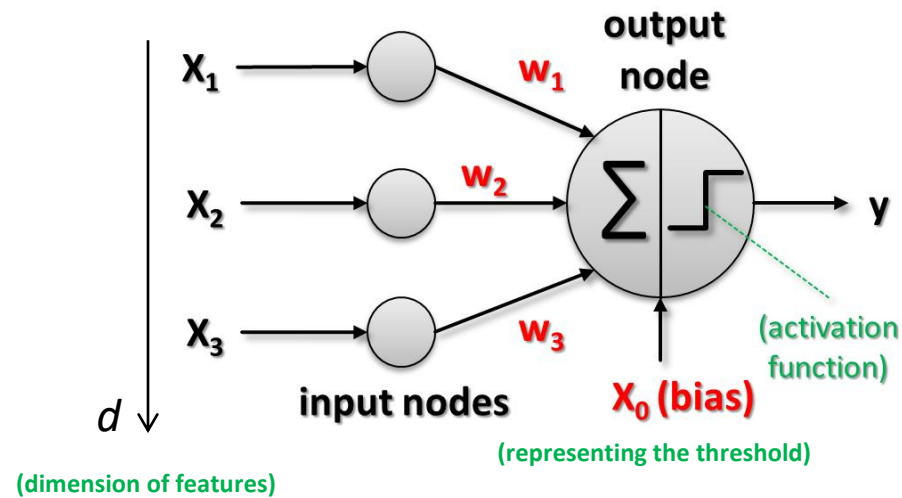
[5] F. Rosenblatt, 1957

$$\text{sign} \left( \underbrace{\left( \sum_{i=1}^d w_i x_i \right)}_{\text{(the signal)}} - \text{threshold} \right)$$

(activation function, +1 or -1)

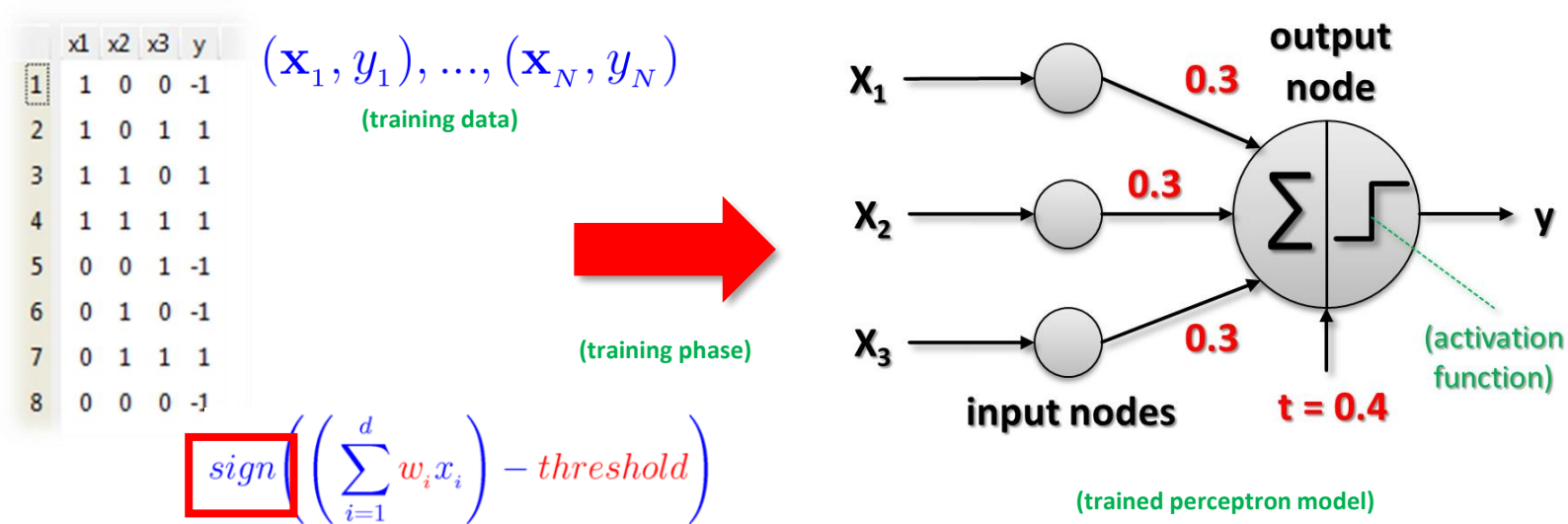
(training data)  $(\mathbf{x}_1, y_1), \dots, (\mathbf{x}_N, y_N)$

(modelled as bias term)





# Perceptron – Example of a Boolean Function



- Output node interpretation
  - More than just the weighted sum of the inputs – threshold (aka bias)
  - Activation function **sign (weighted sum): takes sign of the resulting sum**

$$y = 1, \text{ if } 0.3x_1 + 0.3x_2 + 0.3x_3 - 0.4 > 0$$

(e.g. consider sample #3, sum is positive (0.2) → +1)

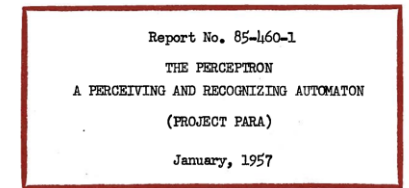
$$y = -1, \text{ if } 0.3x_1 + 0.3x_2 + 0.3x_3 - 0.4 < 0$$

(e.g. consider sample #6, sum is negative (-0.1) → -1)

# Summary Perceptron & Hypothesis Set $h(\mathbf{x})$

- When: Solving a **linear classification** problem
  - Goal: learn a simple value (+1/-1) above/below a certain threshold
  - Class label renamed: **Iris-setosa = -1** and **Iris-virginica = +1**
- Input:  $\mathbf{X} = x_1, \dots, x_d$  (attributes in one dataset)
- Linear formula (take attributes and give them different weights – think of ‘impact of the attribute’)
  - All learned formulas are **different hypothesis** for the given problem

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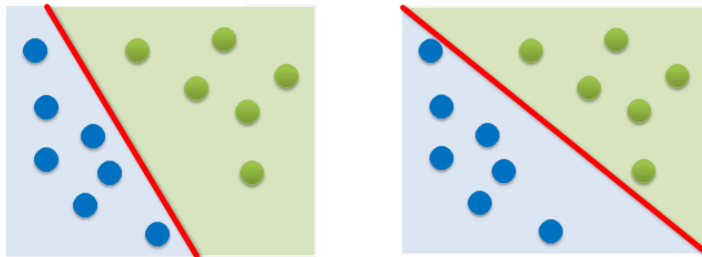


[5] F. Rosenblatt, 1957

$$h(\mathbf{x}) = \text{sign} \left( \left( \sum_{i=1}^d w_i x_i \right) - \text{threshold} \right); h \in \mathcal{H}$$

(parameters that define one hypothesis vs. another)

(each green space and blue space are regions of the same class label determined by sign function)



(red parameters correspond to the redline in graphics)

(but question remains: how do we actually learn  $w_i$  and threshold?)

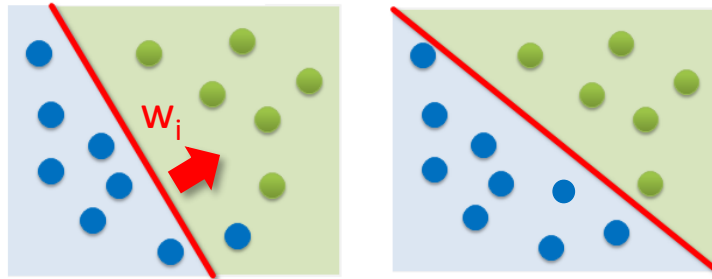
# Perceptron Learning Algorithm – Understanding Vector W

- When: If we believe there is a **linear pattern** to be detected
  - Assumption: **Linearly seperable data** (lets the algorithm converge)
  - Decision boundary: perpendicular vector  $\mathbf{w}_i$  fixes orientation of the line

$$\mathbf{w}^T \mathbf{x} = 0$$

$$\mathbf{w} \cdot \mathbf{x} = 0$$

(points on the decision boundary satisfy this equation)



$$h(\mathbf{x}) = \text{sign}(\mathbf{w}^T \mathbf{x})$$

(vector notation, using T = transpose)

$$\mathbf{w}_i = (w_{i1}, w_{i2}, \dots, w_{id})$$

$$\mathbf{w}_i^T = \begin{bmatrix} w_{i1} \\ w_{i2} \\ \dots \\ w_{id} \end{bmatrix}$$

$$\mathbf{x}_i = (x_{i1}, x_{i2}, \dots, x_{id})$$

$$h(\mathbf{x}) = \text{sign} \left( \left( \sum_{i=1}^d w_i x_i \right) + w_0 \right); w_0 = -\text{threshold}$$

$$h(\mathbf{x}) = \text{sign} \left( \left( \sum_{i=0}^d w_i x_i \right) \right); x_0 = 1$$

$$h(\mathbf{x}) = \text{sign}(\mathbf{w} \cdot \mathbf{x})$$

(equivalent dotproduct notation)

[6] Rosenblatt, 1958

(all notations are equivalent and result is a scalar from which we derive the sign)



# Perceptron Learning Algorithm – Learning Step

- Iterative Method using (labelled) training data  $(\mathbf{x}_1, y_1), \dots, (\mathbf{x}_N, y_N)$

(one point at a time is picked)

- Pick one misclassified training point where:

$$\text{sign}(\mathbf{w}^T \mathbf{x}_n) \neq y_n$$

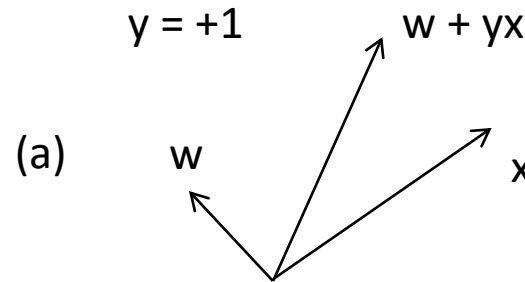
- Update the weight vector:

$$\mathbf{w} \leftarrow \mathbf{w} + y_n \mathbf{x}_n$$

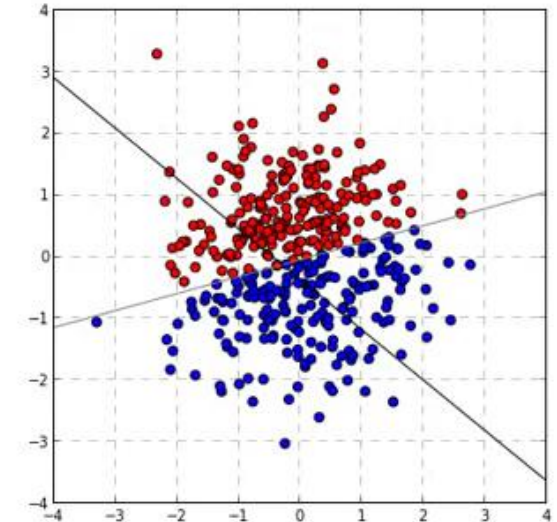
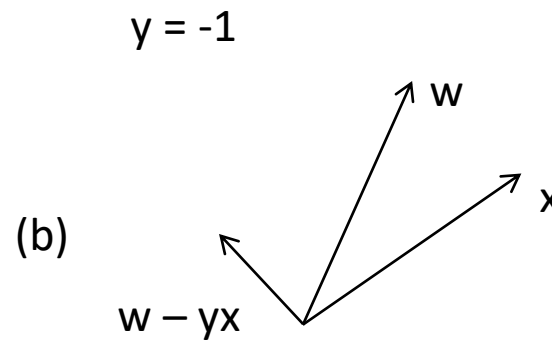
( $y_n$  is either +1 or -1)

- Terminates when there are no misclassified points

(converges only with linearly separable data)

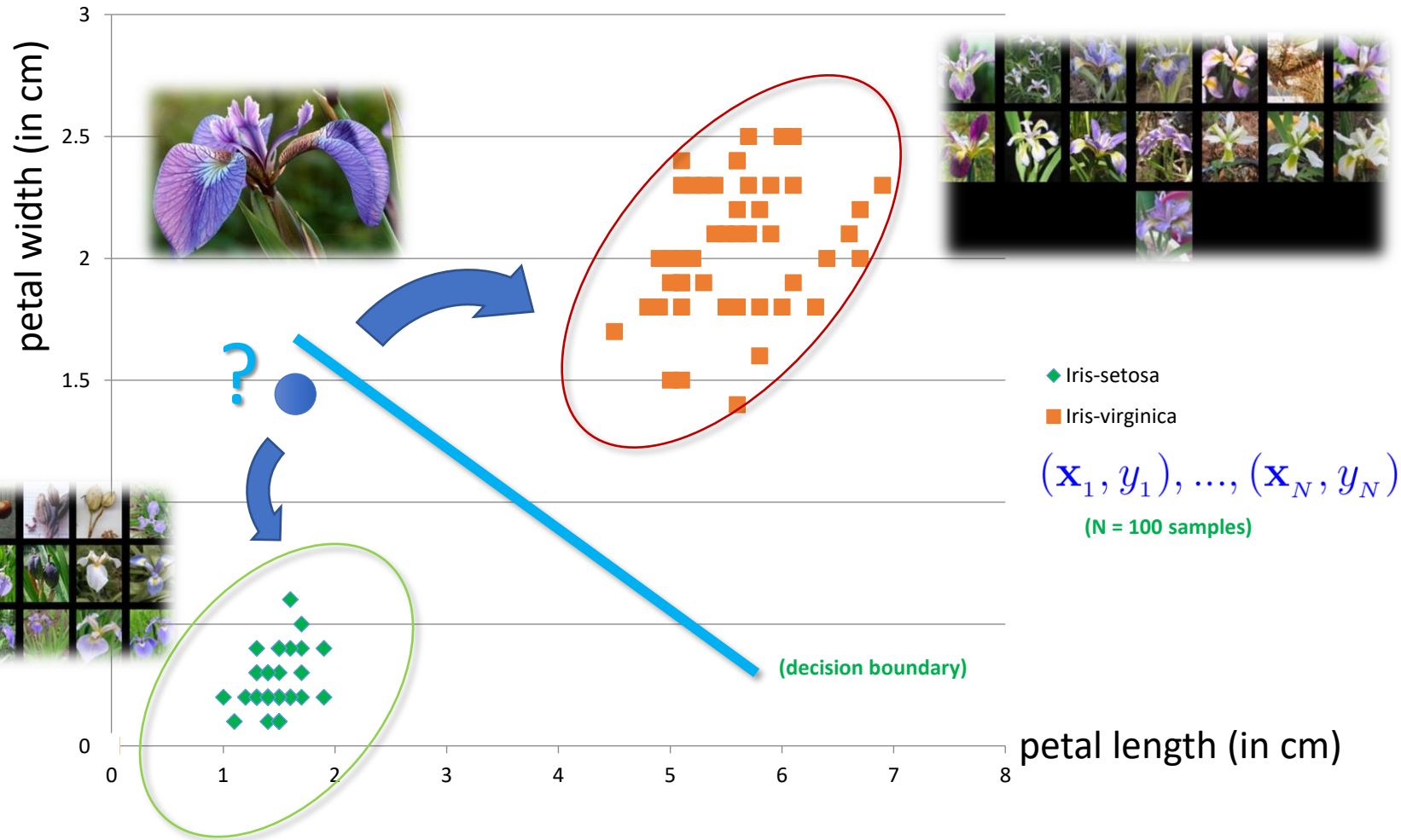


(a) adding a vector or  
(b) subtracting a vector



[7] Perceptron Visualization

# Predicting Task: Obtain Class of a new Flower 'Data Point'



# Food Inspection in Chicago: Advanced Application Example

## 1. Some pattern exists:

- Believe in a pattern with 'quality violations in checking restaurants' will somehow influence if food inspection pass or fail (binary classification)

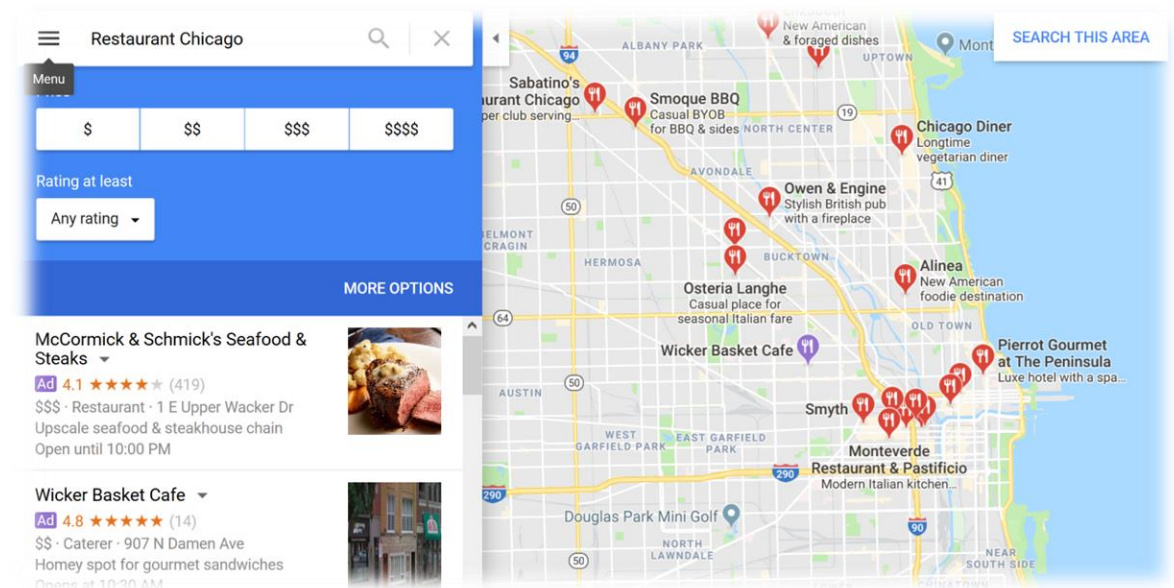
## 2. No exact mathematical formula

- To the best of our knowledge there is no precise formula for this problem

## 3. Data exists

- Data collection from City of Chicago

- The goal of the advanced machine learning application with food inspection of restaurants in the City of Chicago is to predict the outcome of food inspection of new Chicago restaurants given some of existing violations of older restaurants already obtained in Chicago
- A key question is if the new restaurant will pass or fail the inspection just based on violations





# Logistic Regression Using Non-Linear Activation Function

## ■ Linear Classification

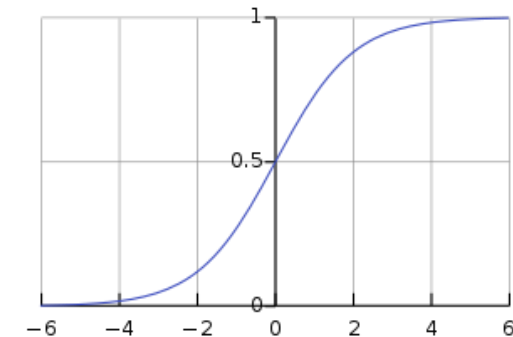
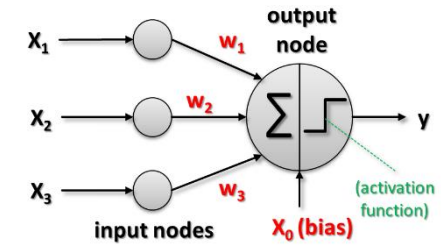
- Simple binary classification (linearly separable)
- Linear combination of the inputs  $x_i$  with weights  $w_i$

## ■ Linear Regression

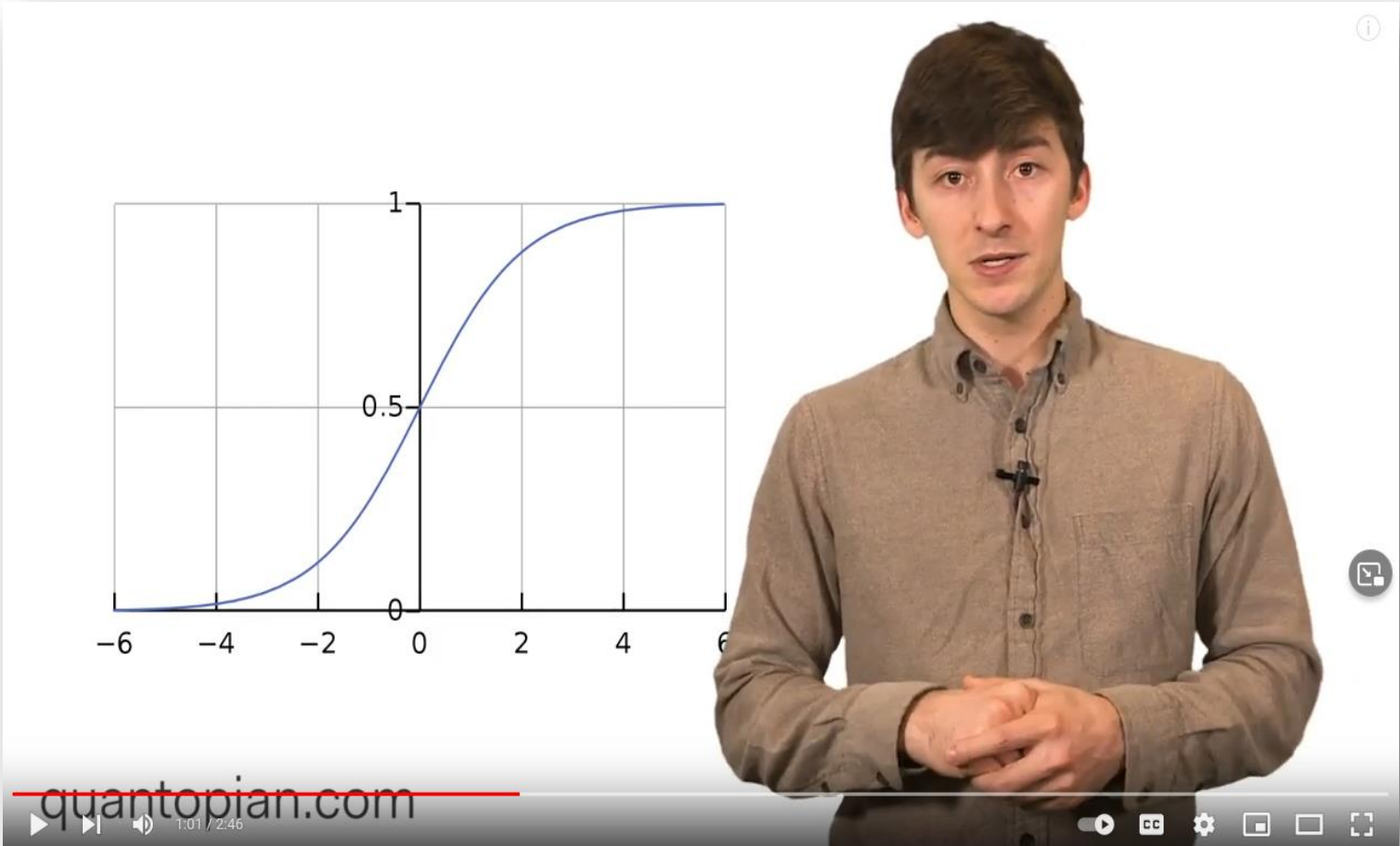
- Real value with the activation being the identity function
- E.g. how much sales given marketing money spend on TV advertising

## ■ Logistic Regression

- Model/error measure/learning algorithm is different
- Captures non-linear data dependencies using the so-called Sigmoid function
- Key idea is to bring values between 0 and 1 to estimate a probability
- (candidate model for pass/fail in our application)



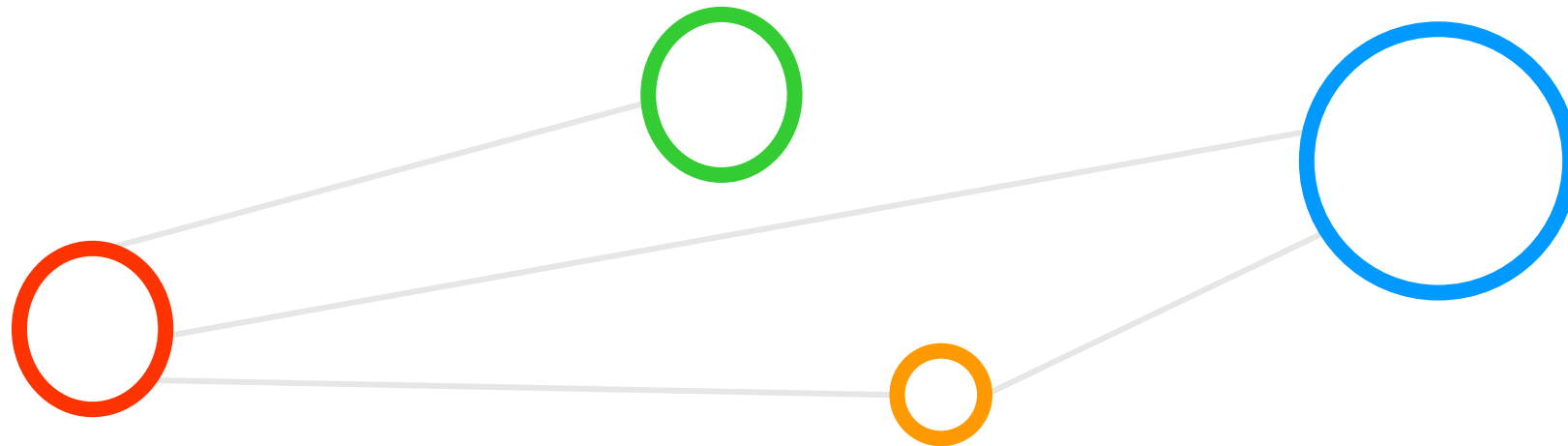
# [Video for further Studies] Logistic Regression Shortly Explained



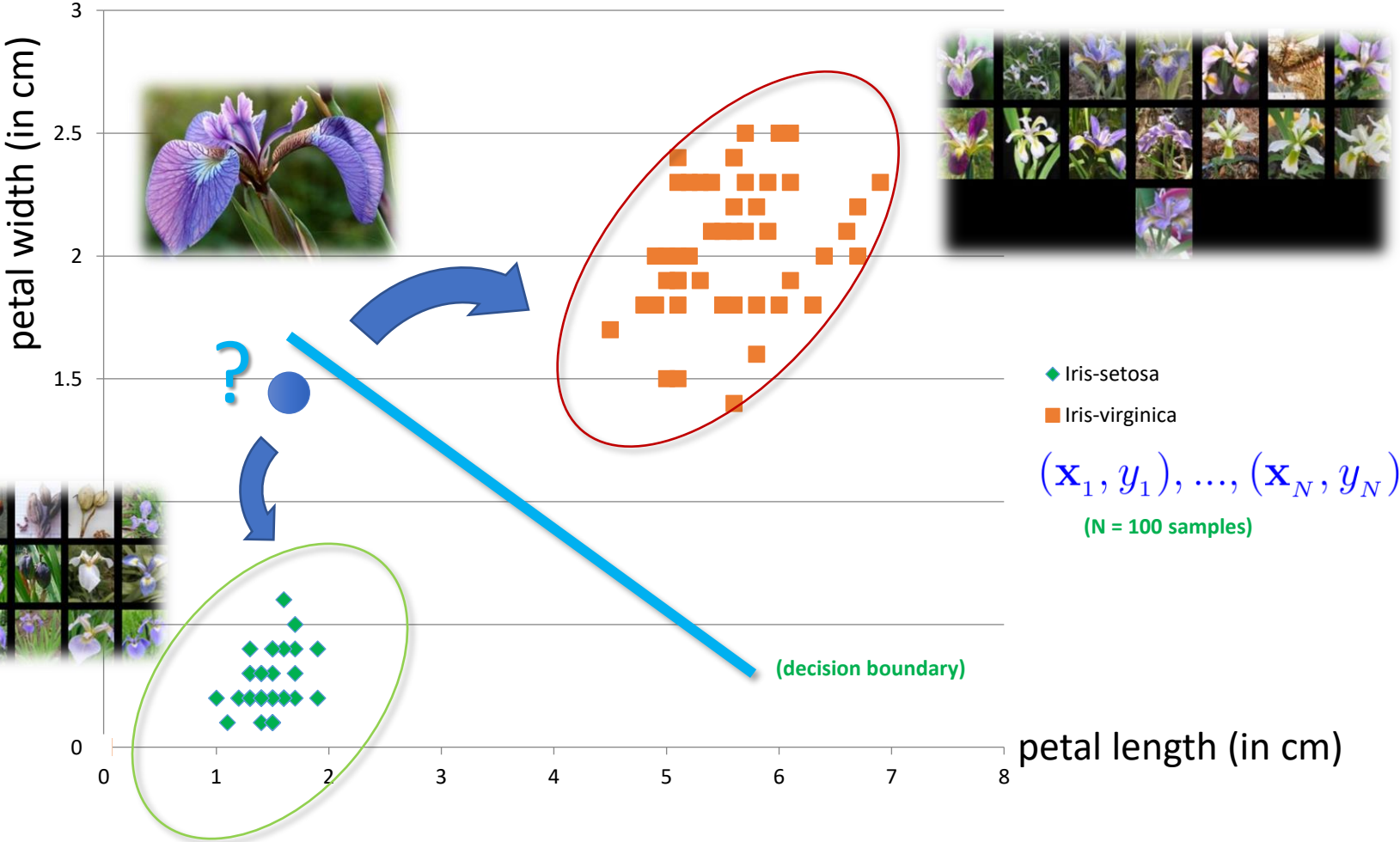
The video player displays a man in a brown shirt speaking. To his left is a graph of a sigmoid function, which is an S-shaped curve. The x-axis ranges from -6 to 6 with major ticks every 2 units. The y-axis ranges from 0 to 1 with major ticks at 0, 0.5, and 1. The curve passes through the point (0, 0.5) and approaches 0 as x goes to negative infinity and 1 as x goes to positive infinity. The video player interface includes a play button, a progress bar showing 1:01 / 2:46, and various control icons like volume, closed captions, settings, and full screen. The watermark 'quantopian.com' is visible in the bottom left of the video frame.

[8] YouTube video, Logistic Regression

# Artificial Neural Network (ANN) Basics



# Predicting Task: Obtain Class of a new Flower 'Data Point' – Revisited



[2] Image sources: Species Iris Group of North America Database, [www.signa.org](http://www.signa.org)



# Perceptron Model – General Mathematical Notation for one Neuron

non-linear activation function

linear combination of input data

$$\hat{y} = g \left( 1 * w_0 + \sum_{i=1}^m x_i * w_i \right)$$

Output

Bias

Sum

→

$$\hat{y} = g \left( w_0 + X^T \mathbf{w} \right)$$

- Simplify the perceptron learning model formula with techniques from linear algebra for mathematical convenience

Constants

$$X = \begin{bmatrix} x_1 \\ \vdots \\ x_m \end{bmatrix} \quad \mathbf{w} = \begin{bmatrix} w_1 \\ \vdots \\ w_m \end{bmatrix}$$

Input Data

Trainable Weights

# Multi-Class Classification: Handwritten Character Recognition MNIST Dataset

- Metadata
  - Not very challenging dataset, but **good for benchmarks & tutorials**
- When working with the dataset
  - Dataset is **not in any standard image format** like jpg, bmp, or gif (i.e. file format not known to a graphics viewer)
  - Data samples are stored in a simple **file format that is designed for storing vectors and multidimensional matrices** (i.e. **numpy arrays**)
  - The pixels of the handwritten digit images are organized row-wise with **pixel values ranging from 0 (white background) to 255 (black foreground)**
  - Images contain grey levels as a result of an anti-aliasing technique used by the normalization algorithm that generated this dataset

- Handwritten Character Recognition MNIST dataset is a subset of a larger dataset from US National Institute of Standards (NIST)
- MNIST handwritten digits includes corresponding labels with values 0-9 and is therefore a labeled dataset
- MNIST digits have been size-normalized to 28 \* 28 pixels & are centered in a fixed-size image for direct processing
- Two separate files for training & test: 60000 training samples (~47 MB) & 10000 test samples (~7.8 MB)

(10 class classification problem)



(downloads data into ~home/.keras/datasets as NPZ file format of numpy that provides storage of array data using gzip compression)



```
import numpy as np
from keras.datasets import mnist
```

```
# download and shuffled as training and testing set
(X_train, y_train), (X_test, y_test) = mnist.load_data()
```

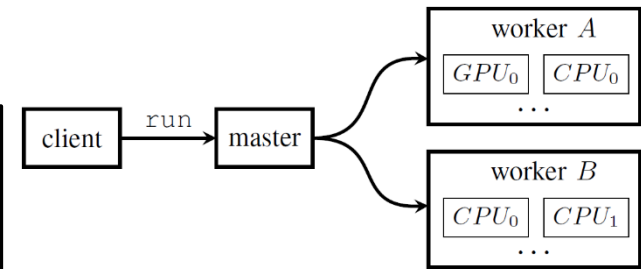


# Deep Learning Frameworks using GPUs also good for Artificial Neural Networks

## TensorFlow

- One of the most popular deep learning frameworks available today
- Execution on **multi-core CPUs or many-core GPUs**

- **Tensorflow is an open source library for deep learning models using a flow graph approach**
- **Tensorflow nodes model mathematical operations and graph edges between the nodes are so-called tensors (also known as multi-dimensional arrays)**
- **The Tensorflow tool supports the use of CPUs and GPUs (much more faster than CPUs)**
- **Tensorflow work with the high-level deep learning tool Keras in order to create models fast**
- **New versions of Tensorflow have Keras shipped with it as well & many further tools**



TensorFlow

[9] Tensorflow  
Web page



[10] Keras  
Web page

## Keras

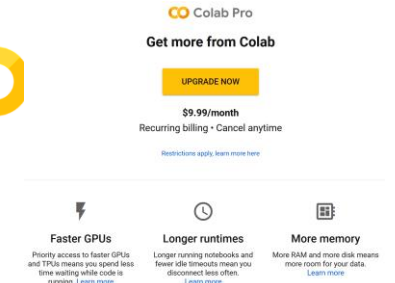
- Often used in combination with low-level frameworks like Tensorflow

- **Keras is a high-level deep learning library implemented in Python that works on top of existing other rather low-level deep learning frameworks like Tensorflow, CNTK, or Theano**
- **Created deep learning models with Keras run seamlessly on CPU and GPU via low-level deep learning frameworks**
- **The key idea behind the Keras tool is to enable faster experimentation with deep networks**
- **Keras is part of the more recent TensorFlow distributions**



# Using Google Colaboratory Cloud Infrastructure for Deep Learning with GPUs

- Google Colaboratory (free & pro version for 9.99 \$ / month)
  - 'Colab' notebooks are Jupyter notebooks that run in the Google cloud
  - Possible to run Apache Spark via PySpark Jupyter notebooks in Colab (cf. Lecture 3)
  - Possible to train Deep Learning networks via GPUs & Jupyter notebooks in Colab
  - Highly integrated with other Google services (e.g., Google Drive for data)
  - Access to vendor-specific Tensor Processing Units (TPUs)



[11] Google Colaboratory

(for international students:  
watch out – it uses the browser  
language automatically)

```
[ ] # initialize the optimizer and model
model = CNN.build(input_shape=INPUT_SHAPE, classes=NB_CLASSES)
model.compile(loss="categorical_crossentropy", optimizer=OPTIMIZER,
metrics=["accuracy"])

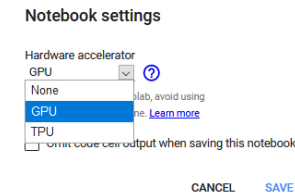
-----
TypeError                                 Traceback (most recent call last)
<ipython-input-11-13ca928a46c2> in <module>()
      1 # initialize the optimizer and model
----> 2 model = CNN.build(input_shape=INPUT_SHAPE, classes=NB_CLASSES)
      3 model.compile(loss="categorical_crossentropy", optimizer=OPTIMIZER,
      4 metrics=["accuracy"])

-----
5 frames
/usr/local/lib/python3.6/dist-packages/tensorflow/python/keras/utils/generic_utils.py in validate_kwargs(kwargs, allowed_kwargs, error_message)
    776 for kwarg in kwargs:
    777     if kwarg not in allowed_kwargs:
--> 778         raise TypeError(error_message, kwarg)
    779
    780

TypeError: ('Keyword argument not understood:', 'border_mode')
```

(Clouds also face this update problem)

■ The portability of deep learning codes is hindered by the frequent updates of the different APIs of deep learning frameworks like Keras, Tensorflow, etc. (cf. different AWS EC2 AMI versions)



(tutorials & codes need updates)

- Update Oct/2016: Updated for Keras 1.1.0, TensorFlow 0.10.0 and scikit-learn v0.18.
- Update Mar/2017: Updated for Keras 2.0.2, TensorFlow 1.0.1 and Theano 0.9.0.
- Update Sep/2019: Updated for Keras 2.2.5 API.

■ Google Colaboratory offers 'Colab' notebooks that are implemented with Jupyter notebooks that in turn run in the Google cloud and are highly integrated with other Google cloud services such as Google Drive thus making 'Colab' notebooks easy to set up, access, and share with others

[12] Machine Learning Mastery MNIST Tutorial



# Data Inspection using Keras Dataset MNIST with Visualization in Jupyter

The screenshot shows a Jupyter Notebook titled "Preprocessing" with the following code cells:

```
In [1]: import numpy as np
from keras.datasets import mnist

Using TensorFlow backend.

WARNING:tensorflow:From /home/ec2-user/anaconda3/envs/tensorflow_p27/lib/python2.7/site-packages/tensorflow_core/_init_.py:1473: The name tf.estimator.inputs is deprecated. Please use tf.compat.v1.estimator.inputs instead.

In [2]: # download and shuffled as training and testing set
(X_train, y_train), (X_test, y_test) = mnist.load_data()

In [3]: # function to explore one hand-written character
def character_show(character):
    for y in character:
        row = ""
        for x in y:
            row += '{0: <4}'.format(x)
        print(row)

In [4]: # view first 10 hand-written characters
for i in range(0,50):
    character_show(X_test[i])
    print("\n")
    print("Label:")
    print(y_test[i])
    print("\n")
```

The output of the notebook shows a grid of 50 rows of 28x28 grayscale pixel values for the first 10 handwritten digits. The output is partially obscured by a redacted area at the bottom.



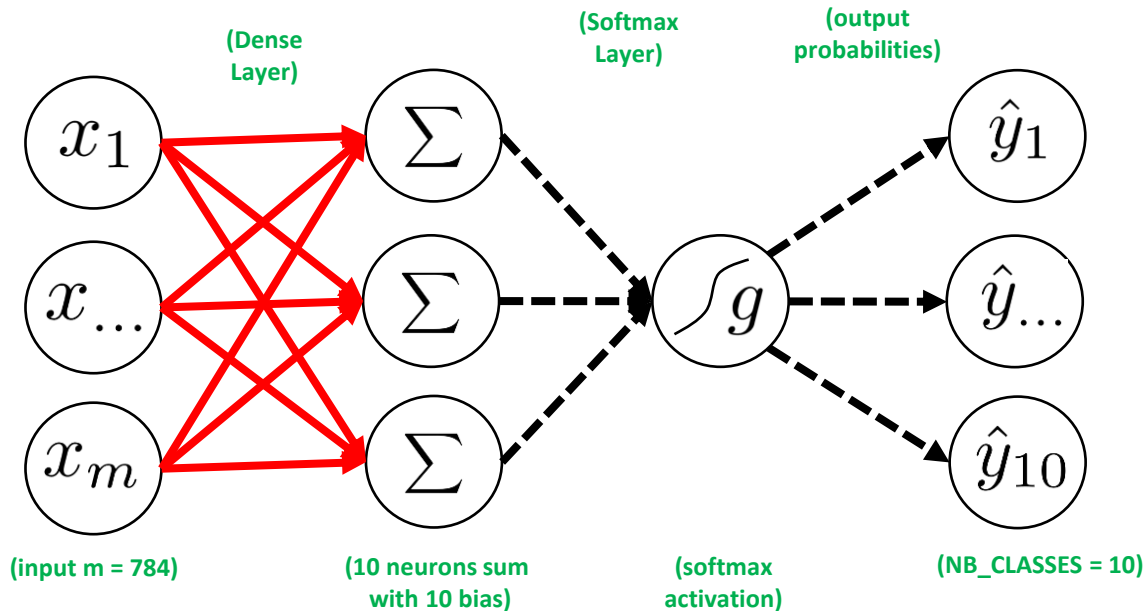




# MNIST Dataset & Multi Output Perceptron Model

- 10 Class Classification Problem

- Use 10 Perceptrons for 10 outputs with softmax activation function (enables probabilities for 10 classes)



```
from keras.models import Sequential
from keras.layers.core import Dense, Activation
```

```
# model Keras sequential
model = Sequential()
```

```
# add fully connected layer - input with output
model.add(Dense(NB_CLASSES, input_shape=(RESHAPED,)))
```

```
# add activation function layer to get class probabilities
model.add(Activation('softmax'))
```

```
# printout a summary of the model to understand model complexity
model.summary()
```

(parameters =  $784 * 10 + 10$  bias = 7850)

Layer (type)	Output Shape	Param #
dense_1 (Dense)	(None, 10)	7850
activation_1 (Activation)	(None, 10)	0
Total params: 7,850		
Trainable params: 7,850		
Non-trainable params: 0		

- Note that the output units are independent among each other in contrast to neural networks with one hidden layer
- The output of softmax gives class probabilities
- The non-linear Activation function 'softmax' represents a generalization of the sigmoid function – it squashes an n-dimensional vector of arbitrary real values into a n-dimensional vector of real values in the range of 0 and 1 – here it aggregates 10 answers provided by the Dense layer with 10 neurons

# MNIST Dataset & Compile Multi Output Perceptron Model

- **Compile** the model

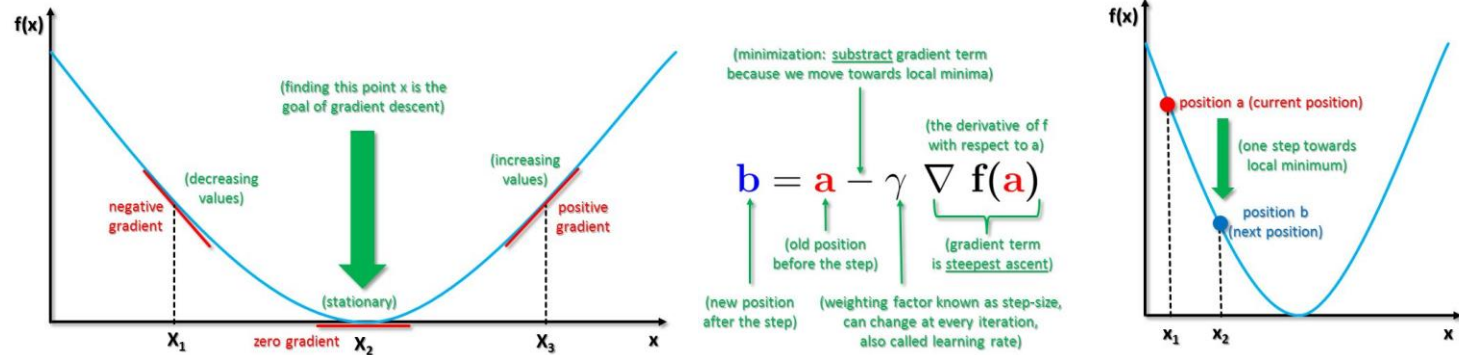
- **Optimizer** as algorithm used to update weights while training the model
- Specify **loss function** (i.e. objective function) that is used by the optimizer to navigate the space of weights
- (note: **process of optimization is also called loss minimization**)
- Indicate **metric** for model evaluation (e.g., accuracy)

- Specify **loss function**

- Compare prediction vs. given class label
- E.g. **categorical\_crossentropy**



```
from keras.optimizers import SGD
OPTIMIZER = SGD() # optimization technique
```



- **Compile the model to be executed by the Keras backend (e.g. TensorFlow)**
- **Optimizer Gradient Descent (GD) uses all the training samples available for a step within a iteration**
- **Optimizer Stochastic Gradient Descent (SGD) converges faster: only one training samples used per iteration**
- **Loss function is a multi-class logarithmic loss: target is  $t_{i,j}$  and prediction is  $p_{i,j}$**
- **Categorical crossentropy is suitable for multiclass label predictions (default with softmax)**



```
# specify loss, optimizer and metric
model.compile(loss='categorical_crossentropy', optimizer=OPTIMIZER, metrics=['accuracy'])
```

$$L_i = -\sum_j t_{i,j} \log(p_{i,j})$$

[14] Big Data Tips, Gradient Descent

# Full Script: MNIST Dataset – Model Parameters & Data Normalization

```
import numpy as np
from keras.datasets import mnist
from keras.models import Sequential
from keras.layers.core import Dense, Activation
from keras.optimizers import SGD
from keras.utils import np_utils
```

```
# parameter setup
NB_EPOCH = 20
BATCH_SIZE = 128
NB_CLASSES = 10 # number of outputs = number of digits
OPTIMIZER = SGD() # optimization technique
VERBOSE = 1
```

```
# download and shuffled as training and testing set
(X_train, y_train), (X_test, y_test) = mnist.load_data()

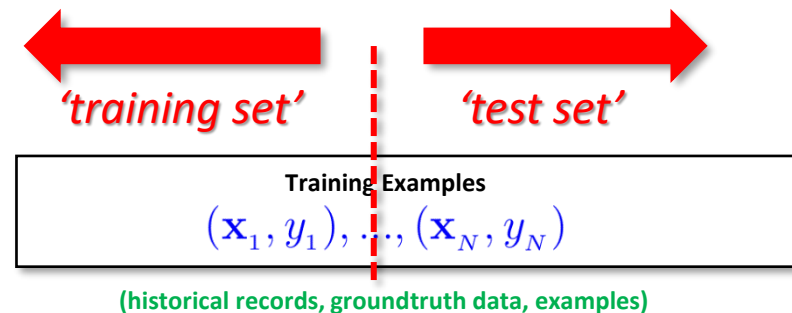
# X_train is 60000 rows of 28x28 values --> reshaped in 60000 x 784
RESHAPED = 784
X_train = X_train.reshape(60000, RESHAPED)
X_test = X_test.reshape(10000, RESHAPED)
X_train = X_train.astype('float32')
X_test = X_test.astype('float32')
```

```
# normalize
X_train /= 255
X_test /= 255
```

```
# output number of samples
print(X_train.shape[0], 'train samples')
print(X_test.shape[0], 'test samples')
```

- **NB\_CLASSES: 10 Class Problem**
- **NB\_EPOCH: number of times the model is exposed to the overall training set – at each iteration the optimizer adjusts the weights so that the objective function is minimized – increasing leads to better accuracy, but also to overfitting**
- **BATCH\_SIZE: number of training instances taken into account before the optimizer performs a weight update to the whole model**
- **OPTIMIZER: Stochastic Gradient Descent ('SGD')**

- **Data load shuffled between training and testing set in files**
- **Data preparation, e.g. X\_train is 60000 samples / rows of 28 x 28 pixel values that are reshaped in 60000 x 784 including type specification (i.e. float32)**
- **Data normalization: divide by 255 – the max intensity value to obtain values in range [0,1]; a usual technique, e.g. model training smoother & avoids data structure problems**



# Full Script: MNIST Dataset – Fitting a Multi Output Perceptron Model

(full script continued from previous slide)

```
# convert class label vectors using one hot encoding
Y_train = np_utils.to_categorical(y_train, NB_CLASSES)
Y_test = np_utils.to_categorical(y_test, NB_CLASSES)
```

```
# model Keras sequential
model = Sequential()
```

```
# add fully connected layer - input with output
model.add(Dense(NB_CLASSES, input_shape=(RESHAPED,)))
```

```
# add activation function layer to get class probabilities
model.add(Activation('softmax'))
```

```
# printout a summary of the model to understand model complexity
model.summary()
```

```
# specify loss, optimizer and metric
model.compile(loss='categorical_crossentropy', optimizer=OPTIMIZER, metrics=['accuracy'])
```

```
# model training
history = model.fit(X_train, Y_train, batch_size=BATCH_SIZE, epochs=NB_EPOCH, verbose=VERBOSE)
```

```
# model evaluation
score = model.evaluate(X_test, Y_test, verbose=VERBOSE)
print("Test score:", score[0])
print('Test accuracy:', score[1])
```

- The Sequential() Keras model is a linear pipeline (aka ‘a stack’) of various neural network layers including Activation functions of different types (e.g. softmax)
- Dense() represents a fully connected layer used in ANNs that means that each neuron in a layer is connected to all neurons located in the previous layer

- The non-linear activation function ‘softmax’ is a generalization of the sigmoid function – it squashes an n-dimensional vector of arbitrary real values into a n-dimensional vector of real values in the range of 0 and 1 – here it aggregates 10 answers provided by the Dense layer with 10 neurons

- Loss function is a multi-class logarithmic loss: target is  $t_{i,j}$  and the prediction is  $p_{i,j}$

$$L_i = -\sum_j t_{i,j} \log(p_{i,j})$$

- Train the model (‘fit’) using selected batch & epoch sizes on training & test data



# Running a Simple ANN with no hidden layers – Multi-Output-Perceptron

```

127.0.0.1:8888/notebooks/ANN_0_Hidden.ipynb
jupyter ANN_0_Hidden Last Checkpoint: 27 minutes ago (unsaved changes)
Environment (conda_tensorflow_p27)
File Edit View Insert Cell Kernel Widgets Help Trusted
+ %> Run Code

In [1]: from __future__ import print_function
import numpy as np
from keras.datasets import mnist
from keras.models import Sequential
from keras.layers.core import Dense, Activation
from keras.optimizers import SGD
from keras.utils import np_utils
np.random.seed(1671) # for reproducibility

Using TensorFlow backend.

WARNING:tensorflow:From /home/ec2-user/anaconda3/envs/tensorflow_p27/lib/python2.7/site-packages/tensorflow_core/__init__.py:1473: The name tf.estimator.inputs is deprecated. Please use tf.compat.v1.estimator.inputs instead.

In [2]: # parameter setup
NB_EPOCH = 200
BATCH_SIZE = 128
VERBOSE = 1
NB_CLASSES = 10 # number of outputs = number of digits
OPTIMIZER = SGD() # optimization technique
N_HIDDEN = 128
VALIDATION_SPLIT=0.2 # 20% of TRAIN is reserved for VALIDATION

In [3]: # download and shuffled as training and testing set
(X_train, y_train), (X_test, y_test) = mnist.load_data()
# X_train is 60000 rows of 28x28 values --> reshaped in 60000 x 784
RESHAPED = 784
X_train = X_train.reshape(60000, RESHAPED)
X_test = X_test.reshape(10000, RESHAPED)
X_train = X_train.astype('float32')
X_test = X_test.astype('float32')
# normalize
X_train /= 255
X_test /= 255
print(X_train.shape[0], 'train samples')
print(X_test.shape[0], 'test samples')
# convert class vectors to binary class matrices
Y_train = np_utils.to_categorical(y_train, NB_CLASSES)
Y_test = np_utils.to_categorical(y_test, NB_CLASSES)

60000 train samples
10000 test samples
    
```

```

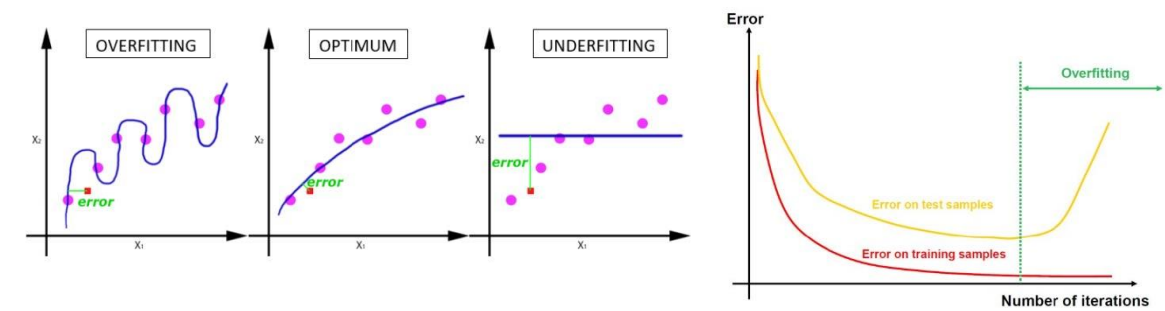
Epoch 199/200
48000/48000 [=====] - 1s 14us/step - loss: 0.2762 - acc: 0.9229 - val_loss: 0.2757 - val_acc: 0.9241
Epoch 200/200
48000/48000 [=====] - 1s 14us/step - loss: 0.2761 - acc: 0.9230 - val_loss: 0.2756 - val_acc: 0.9241

Out[6]: 'Mon, 26 Oct 2020 23:12:30 +0000'

In [7]: # model evaluation
score = model.evaluate(X_test, Y_test, verbose=VERBOSE)
print("Test score:", score[0])
print('Test accuracy:', score[1])

10000/10000 [=====] - 0s 22us/step
Test score: 0.2773858518630266
Test accuracy: 0.9227
    
```

- Note that the outcome of the training process is the result of optimization techniques like SGD that tend to vary 'a bit'
- Note that the outcome of the training process can be dependent on the length of training increasing accuracy to a certain point when overfitting starts
- Overfitting can be controlled with validation and regularization techniques that belong to advanced machine learning methods to be studied in full university machine learning course in detail



# MNIST Dataset – A Multi Output Perceptron Model – Output & Evaluation

```

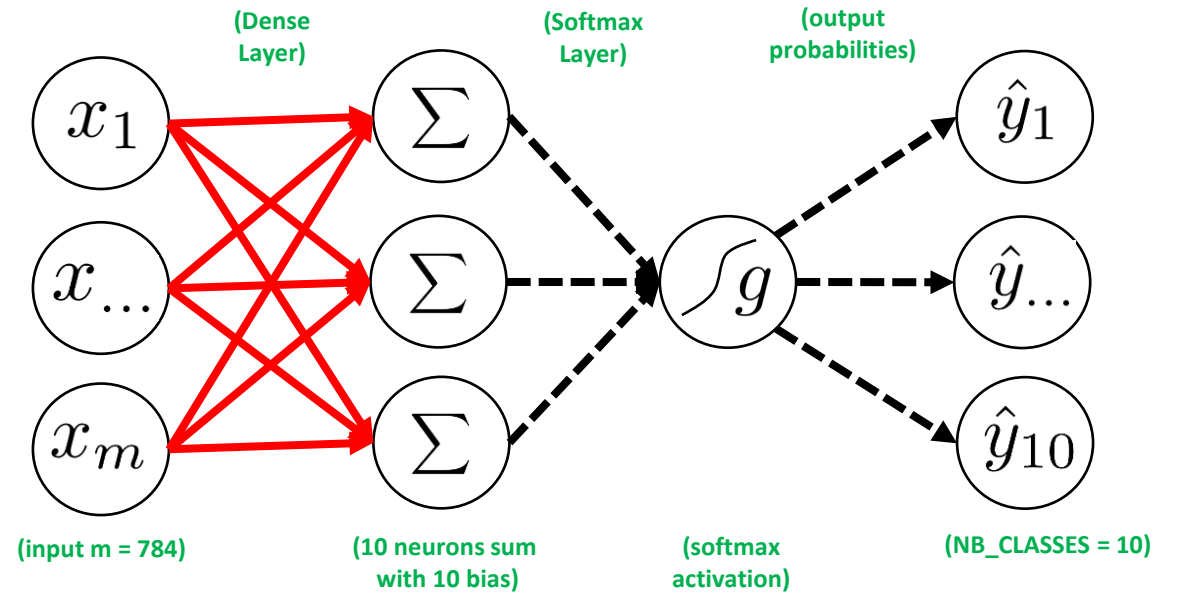
Epoch 7/20
60000/60000 [=====] - 2s 26us/step - loss: 0.4419 - acc: 0.8838
Epoch 8/20
60000/60000 [=====] - 2s 26us/step - loss: 0.4271 - acc: 0.8866
Epoch 9/20
60000/60000 [=====] - 2s 25us/step - loss: 0.4151 - acc: 0.8888
Epoch 10/20
60000/60000 [=====] - 2s 26us/step - loss: 0.4052 - acc: 0.8910
Epoch 11/20
60000/60000 [=====] - 2s 26us/step - loss: 0.3968 - acc: 0.8924
Epoch 12/20
60000/60000 [=====] - 2s 25us/step - loss: 0.3896 - acc: 0.8944
Epoch 13/20
60000/60000 [=====] - 2s 26us/step - loss: 0.3832 - acc: 0.8956
Epoch 14/20
60000/60000 [=====] - 2s 25us/step - loss: 0.3777 - acc: 0.8969
Epoch 15/20
60000/60000 [=====] - 2s 25us/step - loss: 0.3727 - acc: 0.8982
Epoch 16/20
60000/60000 [=====] - 1s 24us/step - loss: 0.3682 - acc: 0.8989
Epoch 17/20
60000/60000 [=====] - 1s 25us/step - loss: 0.3641 - acc: 0.9001
Epoch 18/20
60000/60000 [=====] - 1s 25us/step - loss: 0.3604 - acc: 0.9007
Epoch 19/20
60000/60000 [=====] - 2s 25us/step - loss: 0.3570 - acc: 0.9016
Epoch 20/20
60000/60000 [=====] - 1s 24us/step - loss: 0.3538 - acc: 0.9023
    
```

```

# model evaluation
score = model.evaluate(X_test, Y_test, verbose=VERBOSE)
print("Test score:", score[0])
print('Test accuracy:', score[1])
    
```

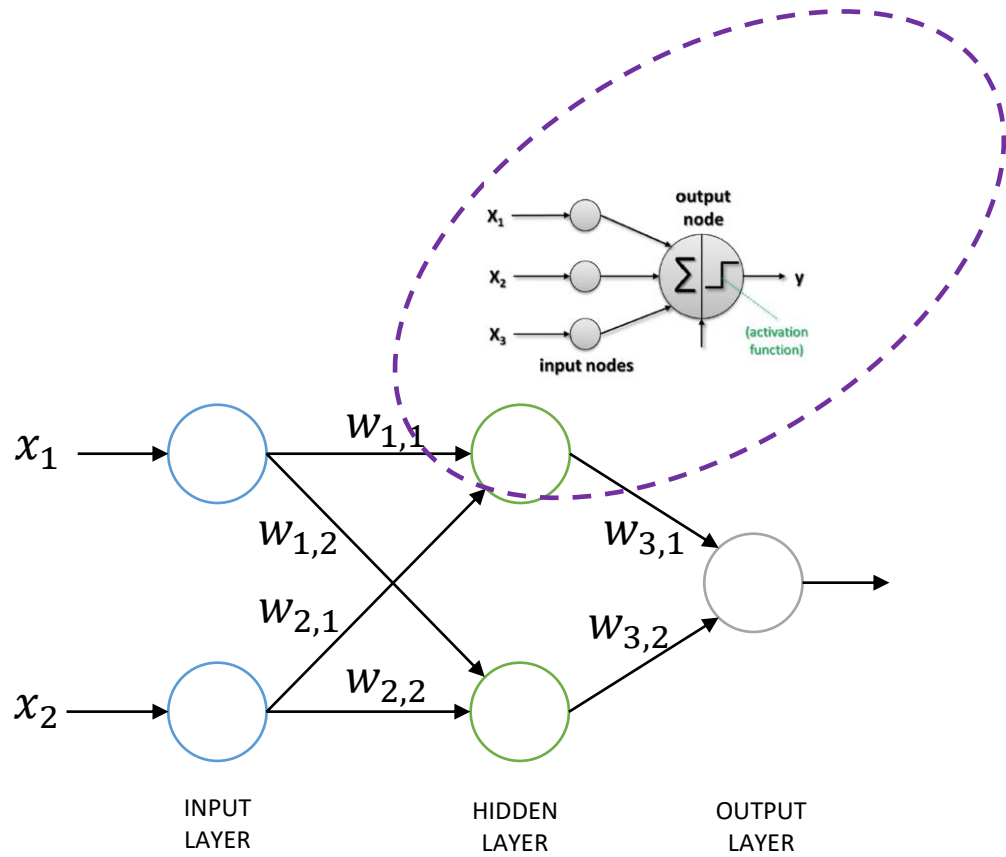
```

10000/10000 [=====] - 0s 41us/step
Test score: 0.33423959468007086
Test accuracy: 0.9101
    
```

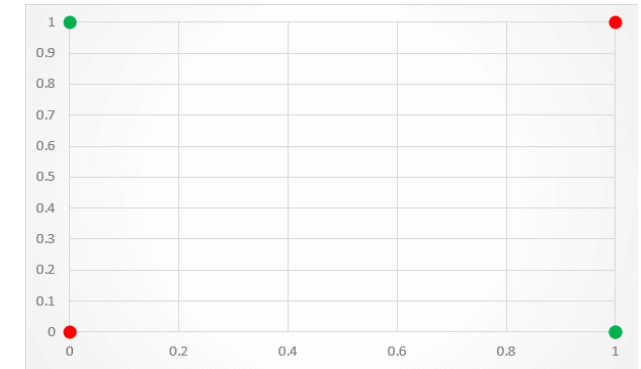


- How to improve the model design by extending the neural network topology?
- Which layers are required?
- Think about input layer need to match the data – what data we had?
- Maybe hidden layers?
- How many hidden layers?
- What activation function for which layer (e.g. maybe ReLU)?
- Think Dense layer – Keras?
- Think about final Activation as Softmax → output probability

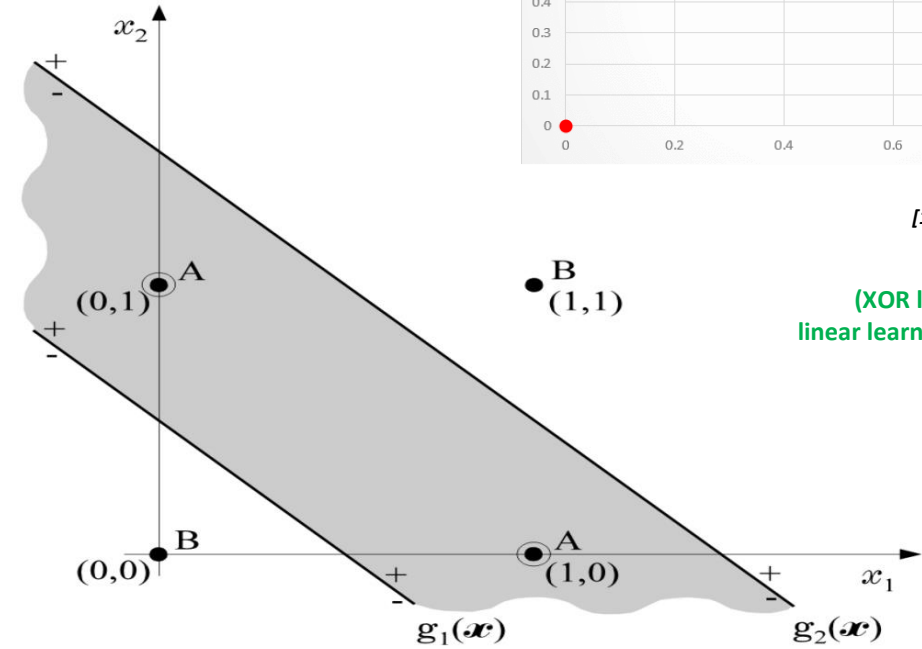
# From Limits of Linear Perceptron Model to Multi Layer Perceptrons (MLP)



Input 1	Input 2	Output
0	0	0
0	1	1
1	1	0
1	0	1



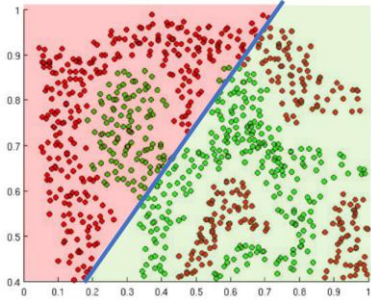
[15] The XOR Problem



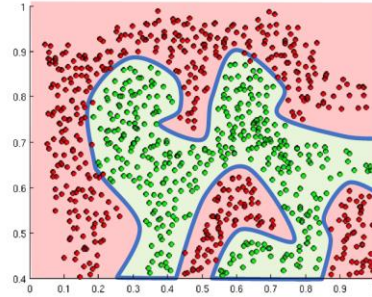
(XOR limit of linear learning models)

[16] Multilayer Perceptron

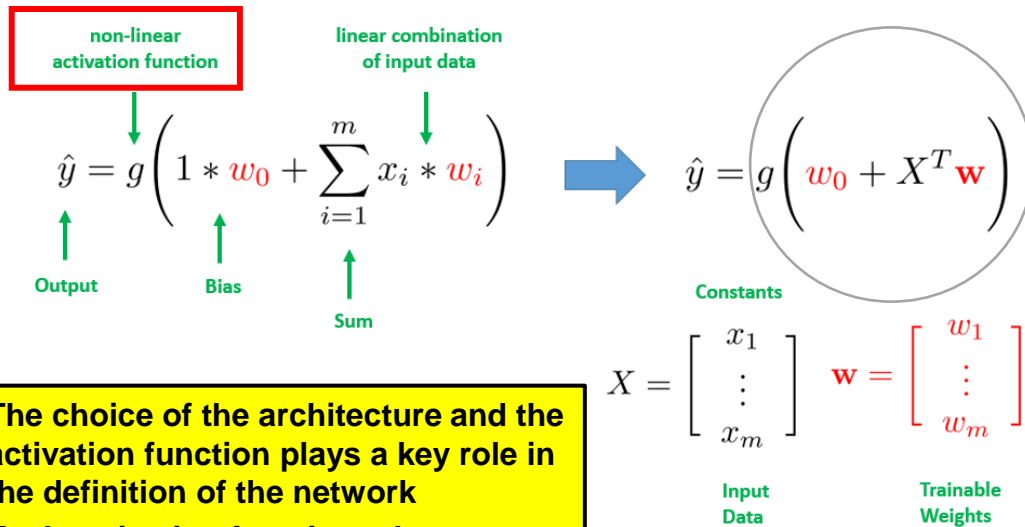
# Introduce Non-Linearities – The Role of Activation Functions in ANNs



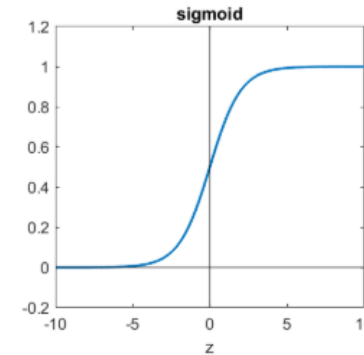
Linear Activation functions produce linear decisions no matter the network size



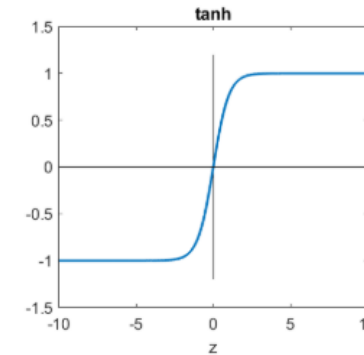
Non-linearities allow us to approximate arbitrarily complex functions



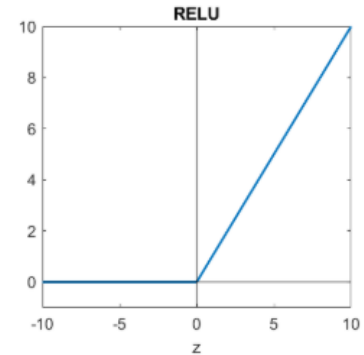
- The choice of the architecture and the activation function plays a key role in the definition of the network
- Each activation function takes a single number and performs a certain fixed mathematical operation on it



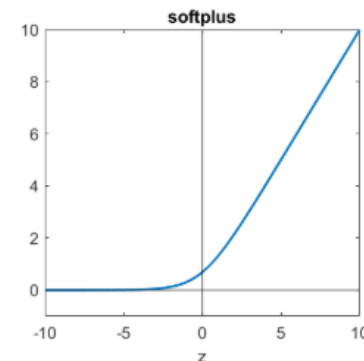
$$g(z) = \frac{1}{1 + e^{-z}}$$



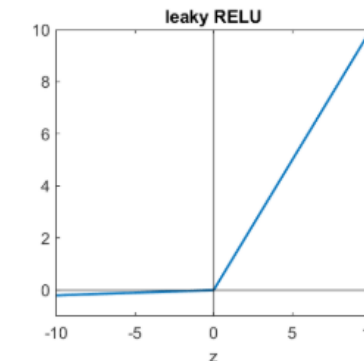
$$g(z) = \tanh z$$



$$g(z) = \max(z, 0)$$

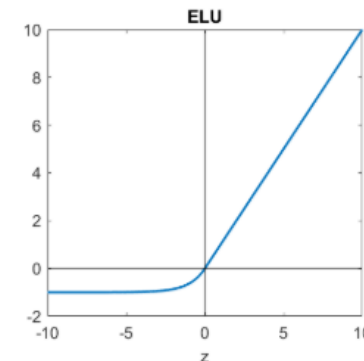


$$g(z) = \log(1 + e^z)$$



$$g(z) = \max(z, z\alpha)$$

$$0 < \alpha < 1$$

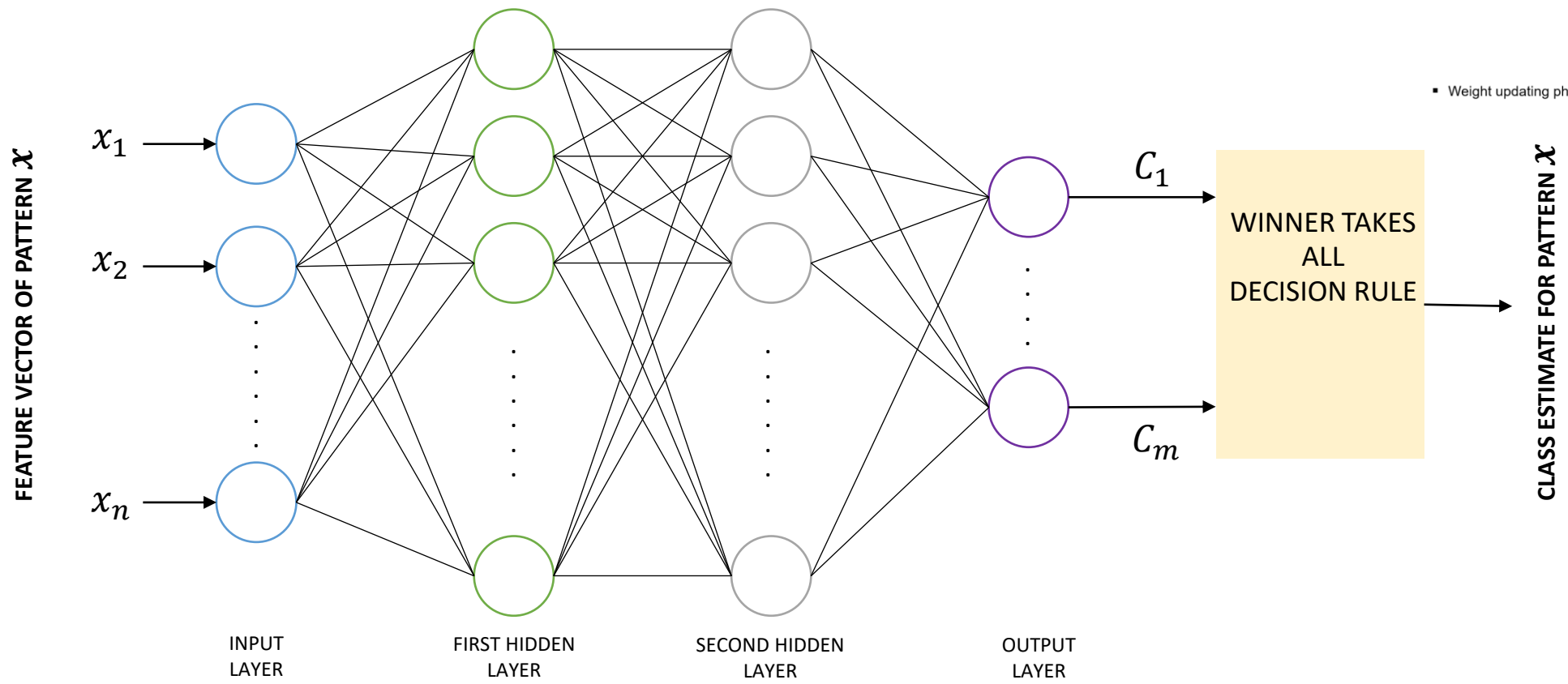
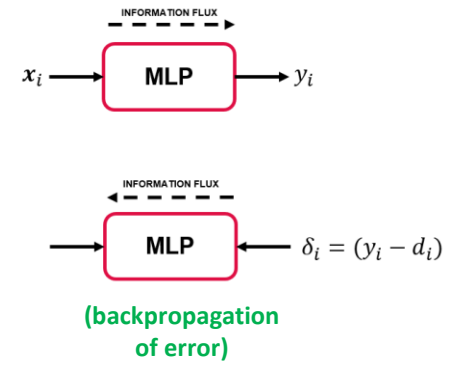


[17] Introduction to Deep Learning [18] Understanding the Neural Network

# Artificial Neural Network (ANN) Basic Network Topology & Learning Algorithm

- Forward interconnection of several layers of perceptrons create an Artificial Neural Network (ANNs)
- Multi Layer Perceptrons (MLPs) can be used as universal approximators
- In classification problems, they allow modeling nonlinear discriminant functions
- Interconnecting neurons aims at increasing the capability of modeling complex input-output relationships

- Forward propagation phase
- Backward propagation phase
- Weight updating phase





# Training an ANN with Backpropagation performing Weight Updates

1. Initialize weights randomly  $\sim \mathcal{N}(0, \sigma^2)$
2. Loop until convergence
3. Compute gradient  $\frac{\partial \mathcal{L}(W)}{\partial W}$  (it explains how the loss changes with respect to each of the weights) (compute-intensive)
4. Update weights  $W := W - \eta \frac{\partial \mathcal{L}(W)}{\partial W}$  (in the opposite direction of the gradient)
5. Return weights (Learning rate) (goal: minimize the 'Loss' using an optimization problem)

- The Learning Rate determines the adjustment magnitude & how much do you trust the computed gradient
- Computing the gradients using optimization is the most computational part when there is a high number of weights

$$\mathcal{L} = \frac{1}{N} \sum_{i=1}^N (y_i - d_i)^2$$

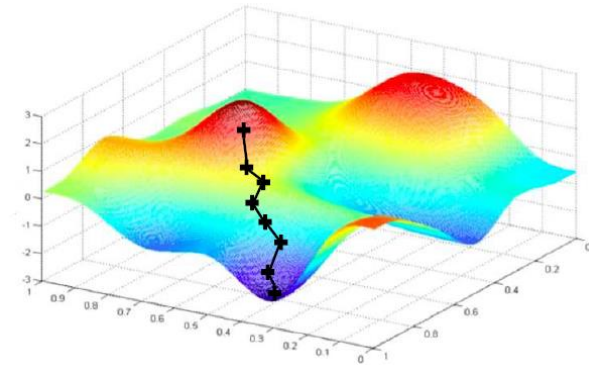
TOTAL NUMBER OF TRAINING SAMPLES

OUTPUT VALUE OBTAINED BY THE MLP FOR THE i-th SAMPLE

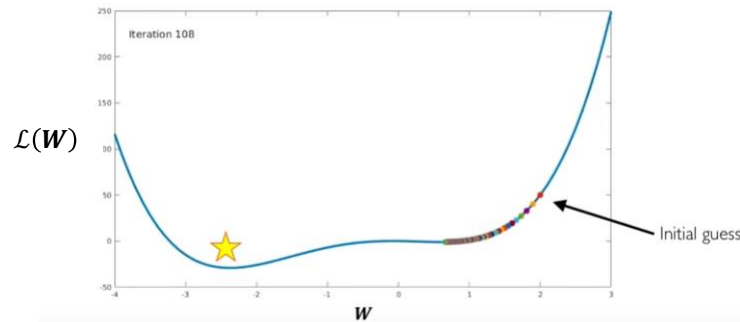
DESIRED OUTPUT (TARGET) VALUE FOR THE i-th SAMPLE

$$W^* = \arg \min \sum_{i=1}^N \mathcal{L}(f(x_i; W), d_i)$$

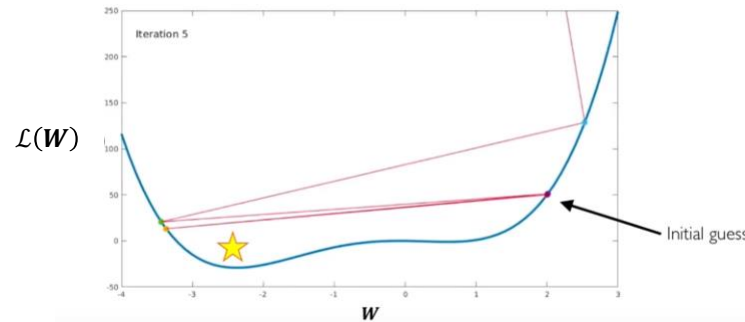
$$W^* = \arg \min_W \mathcal{L}(W)$$



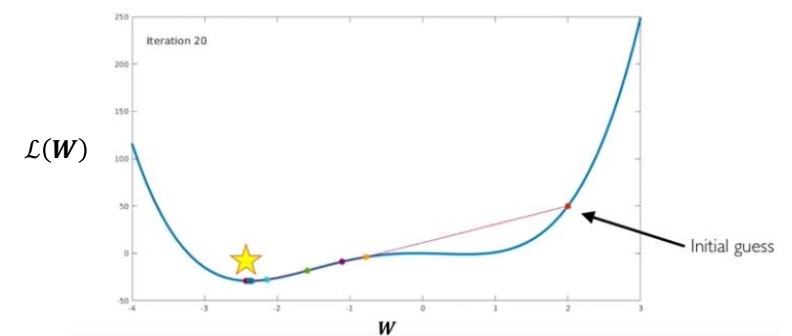
[17] Introduction to Deep Learning



**Small learning rate** converges slowly and gets stuck in false local minima



**Large learning rates** overshoot, become unstable and diverge



**Stable learning rates** converge smoothly and avoid local minima

# MNIST Dataset – Add Two Hidden Layers for Artificial Neural Network (ANN)

- All parameter value remain the same as before
- We add N\_HIDDEN as parameter in order to set 128 neurons in one hidden layer – this number is a hyperparameter that is not directly defined and needs to be find with parameter search

```
# parameter setup
NB_EPOCH = 20
BATCH_SIZE = 128
NB_CLASSES = 10 # number of outputs = number of digits
OPTIMIZER = SGD() # optimization technique
VERBOSE = 1
N_HIDDEN = 128 # number of neurons in one hidden layer
```

```
# model Keras sequential
model = Sequential()
```

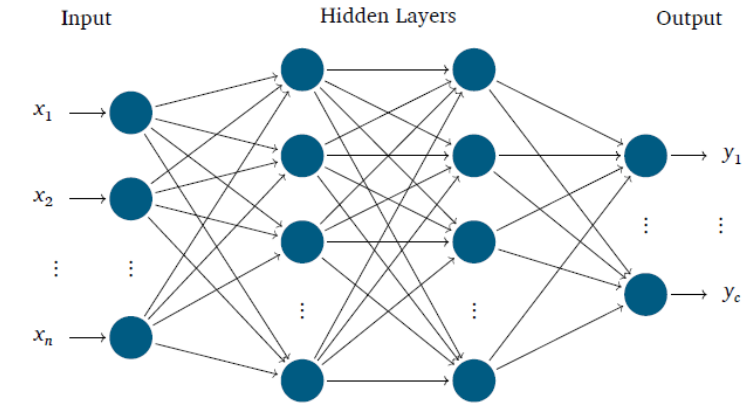
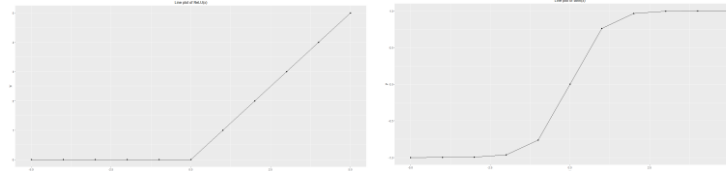
```
# modeling step
# 2 hidden layers each N_HIDDEN neurons
model.add(Dense(N_HIDDEN, input_shape=(RESHAPED,)))
model.add(Activation('relu'))
model.add(Dense(N_HIDDEN))
model.add(Activation('relu'))
model.add(Dense(NB_CLASSES))
```

```
# add activation function layer to get class probabilities
model.add(Activation('softmax'))
```

[19] big-data.tips,  
'Relu Neural Network'

[20] big-data.tips,  
'tanh'

(activation functions ReLU & Tanh)



```
model.add(Dense(N_HIDDEN))
model.add(Activation('relu'))
```

```
model.add(Dense(N_HIDDEN))
model.add(Activation('tanh'))
```

- The non-linear Activation function 'relu' represents a so-called Rectified Linear Unit (ReLU) that only recently became very popular because it generates good experimental results in ANNs and more recent deep learning models – it just returns 0 for negative values and grows linearly for only positive values
- A hidden layer in an ANN can be represented by a fully connected Dense layer in Keras by just specifying the number of hidden neurons in the hidden layer

# Running a Simple ANN with two hidden layers (200 Epochs: very long learning)

```
127.0.0.1:8888/notebooks/ANN_2_Hidden.ipynb
jupyter ANN_2_Hidden Last Checkpoint: 30 minutes ago (unsaved changes)
Environment (conda_tensorflow_p27)

In [1]: from __future__ import print_function
import numpy as np
from keras.datasets import mnist
from keras.models import Sequential
from keras.layers.core import Dense, Activation
from keras.optimizers import SGD
from keras.utils import np_utils
np.random.seed(1671) # for reproducibility

Using TensorFlow backend.

WARNING:tensorflow:From /home/ec2-user/anaconda3/envs/tensorflow_p27/lib/python2.7/site-packages/tensorflow_core/_init_.py:1473: The name tf.estimator.inputs is deprecated. Please use tf.compat.v1.estimator.inputs instead.

In [2]: # parameter setup
NB_EPOCH = 200
BATCH_SIZE = 128
VERBOSE = 1
NB_CLASSES = 10 # number of outputs = number of digits
OPTIMIZER = SGD() # optimization technique
N_HIDDEN = 128
VALIDATION_SPLIT=0.2 # 20% of TRAIN is reserved for VALIDATION

In [3]: # download and shuffled as training and testing set
(X_train, y_train), (X_test, y_test) = mnist.load_data()
# X_train is 60000 rows of 28x28 values --> reshaped in 60000 x 784
RESHAPED = 784
X_train = X_train.reshape(60000, RESHAPED)
X_test = X_test.reshape(10000, RESHAPED)
X_train = X_train.astype('float32')
X_test = X_test.astype('float32')
# normalize
X_train /= 255
X_test /= 255
print(X_train.shape[0], 'train samples')
print(X_test.shape[0], 'test samples')
# convert class vectors to binary class matrices
Y_train = np_utils.to_categorical(y_train, NB_CLASSES)
Y_test = np_utils.to_categorical(y_test, NB_CLASSES)

60000 train samples
10000 test samples
```

```
Epoch 196/200
48000/48000 [=====] - 1s 26us/step - loss: 0.0160 - acc: 0.9978 - val_loss: 0.0884 - val_acc: 0.97
50
Epoch 197/200
48000/48000 [=====] - 1s 26us/step - loss: 0.0158 - acc: 0.9979 - val_loss: 0.0888 - val_acc: 0.97
46
Epoch 198/200
48000/48000 [=====] - 1s 26us/step - loss: 0.0157 - acc: 0.9978 - val_loss: 0.0891 - val_acc: 0.97
48
Epoch 199/200
48000/48000 [=====] - 1s 26us/step - loss: 0.0155 - acc: 0.9979 - val_loss: 0.0890 - val_acc: 0.97
49
Epoch 200/200
48000/48000 [=====] - 1s 26us/step - loss: 0.0154 - acc: 0.9979 - val_loss: 0.0892 - val_acc: 0.97
47

Out[6]: 'Mon, 26 Oct 2020 23:19:28 +0000'

In [7]: # model evaluation
score = model.evaluate(X_test, Y_test, verbose=VERBOSE)
print("Test score:", score[0])
print("Test accuracy:", score[1])

10000/10000 [=====] - 0s 27us/step
Test score: 0.07618757467148826
Test accuracy: 0.9764
```

# MNIST Dataset – ANN Model Parameters & Output Evaluation (20 Epochs)

```
Epoch 7/20
60000/60000 [=====] - 1s 18us/step - loss: 0.2743 - acc: 0.9223
Epoch 8/20
60000/60000 [=====] - 1s 18us/step - loss: 0.2601 - acc: 0.9266
Epoch 9/20
60000/60000 [=====] - 1s 18us/step - loss: 0.2477 - acc: 0.9301
Epoch 10/20
60000/60000 [=====] - 1s 18us/step - loss: 0.2365 - acc: 0.9329
Epoch 11/20
60000/60000 [=====] - 1s 18us/step - loss: 0.2264 - acc: 0.9356
Epoch 12/20
60000/60000 [=====] - 1s 18us/step - loss: 0.2175 - acc: 0.9386
Epoch 13/20
60000/60000 [=====] - 1s 18us/step - loss: 0.2092 - acc: 0.9412
Epoch 14/20
60000/60000 [=====] - 1s 18us/step - loss: 0.2013 - acc: 0.9432
Epoch 15/20
60000/60000 [=====] - 1s 18us/step - loss: 0.1942 - acc: 0.9454
Epoch 16/20
60000/60000 [=====] - 1s 18us/step - loss: 0.1876 - acc: 0.9472
Epoch 17/20
60000/60000 [=====] - 1s 18us/step - loss: 0.1813 - acc: 0.9487
Epoch 18/20
60000/60000 [=====] - 1s 18us/step - loss: 0.1754 - acc: 0.9502
Epoch 19/20
60000/60000 [=====] - 1s 18us/step - loss: 0.1700 - acc: 0.9522
Epoch 20/20
60000/60000 [=====] - 1s 18us/step - loss: 0.1647 - acc: 0.9536
```

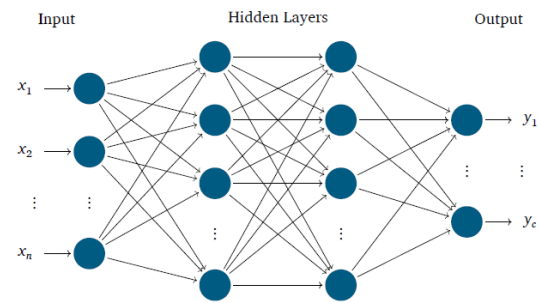
```
# model evaluation
score = model.evaluate(X_test, Y_test, verbose=VERBOSE)
print("Test score:", score[0])
print('Test accuracy:', score[1])
```

```
10000/10000 [=====] - 0s 33us/step
Test score: 0.16286438911408185
Test accuracy: 0.9514
```

- ✓ **Multi Output Perceptron: ~91,01% (20 Epochs)**
- ✓ **ANN 2 Hidden Layers: ~95,14 % (20 Epochs)**



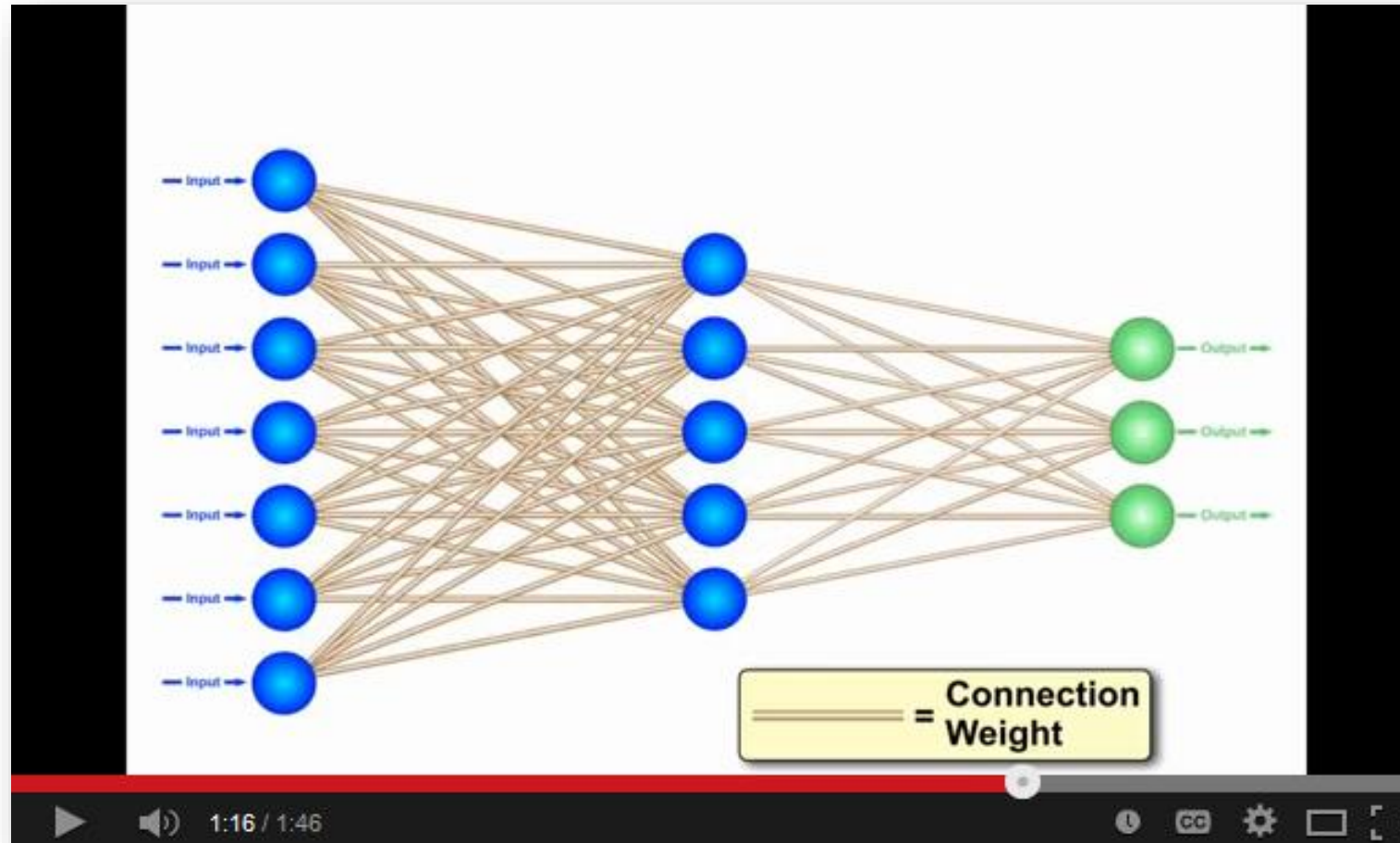
```
# printout a summary of the model to understand model complexity
model.summary()
```



Layer (type)	Output Shape	Param #
dense_1 (Dense)	(None, 128)	100480
activation_1 (Activation)	(None, 128)	0
dense_2 (Dense)	(None, 128)	16512
activation_2 (Activation)	(None, 128)	0
dense_3 (Dense)	(None, 10)	1290
activation_3 (Activation)	(None, 10)	0
Total params: 118,282		
Trainable params: 118,282		
Non-trainable params: 0		

- **Dense Layer connects every neuron in this dense layer to the next dense layer with each of its neuron also called a fully connected network element with weights as trainable parameters**
- **Choosing a model with different layers is a model selection that directly also influences the number of parameters (e.g. add Dense layer from Keras means new weights)**
- **Adding a layer with these new weights means much more computational complexity since each of the weights must be trained in each epoch (depending on #neurons in layer)**

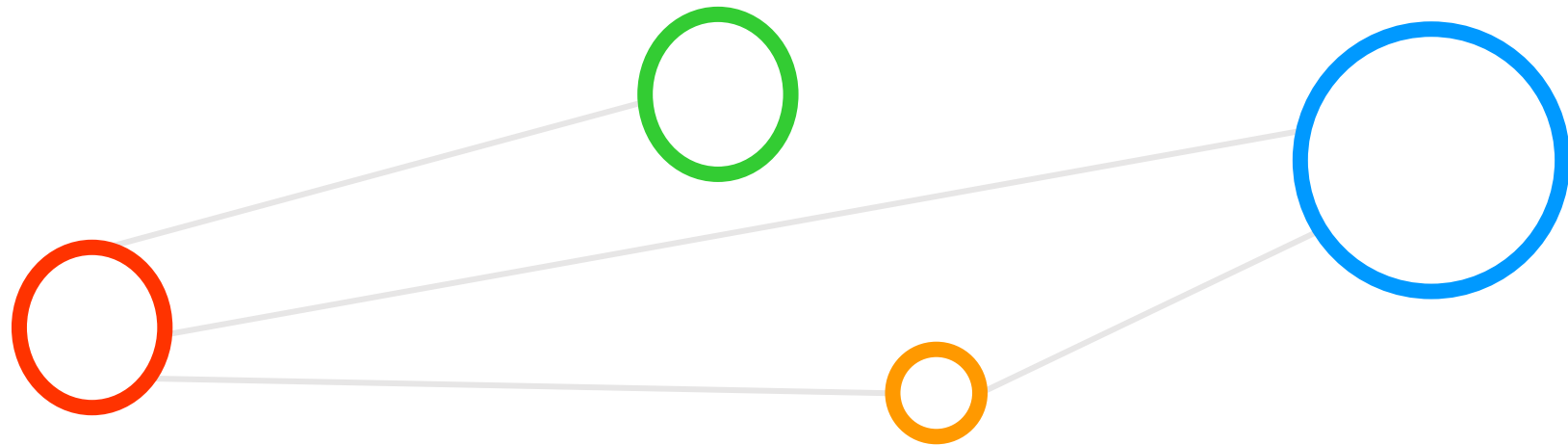
# [Video for further Studies] Neural Networks Summary



[21] YouTube Video, Neural Networks – A Simple Explanation

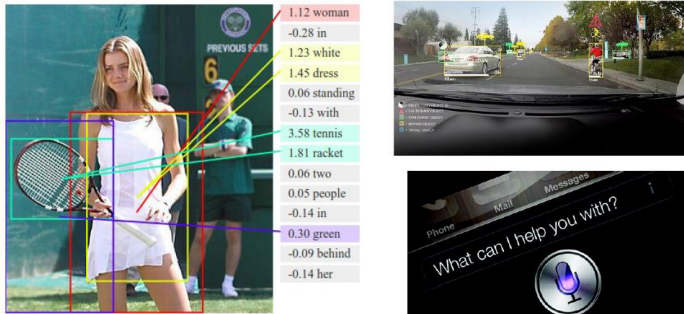


# Convolutional Neural Network (CNN) Basics

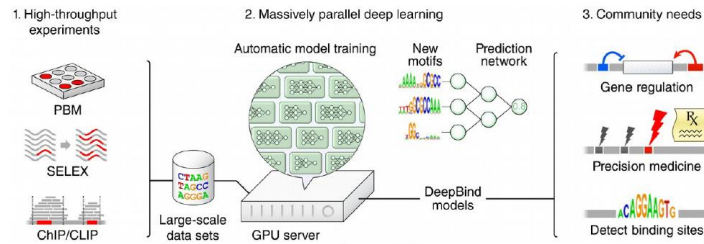


# Impact of Deep Learning in Various Application Domains

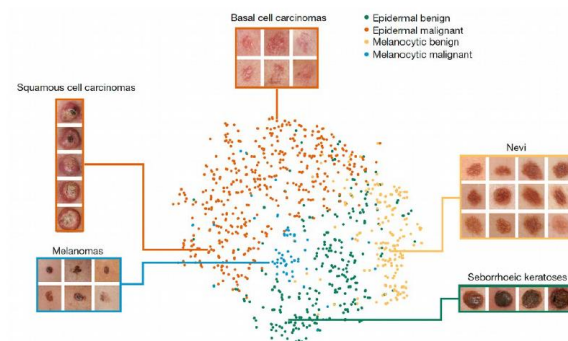
## Vision, Natural Speech



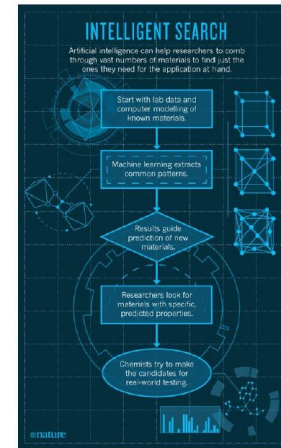
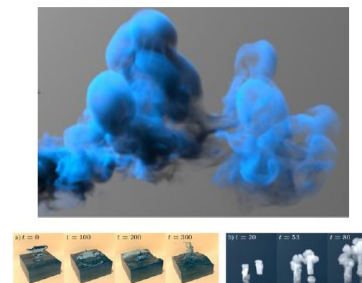
## Omics (Genomics, Proteomics, Metabolomics)



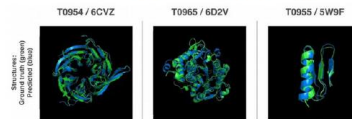
## Medical Imaging and Diagnostics



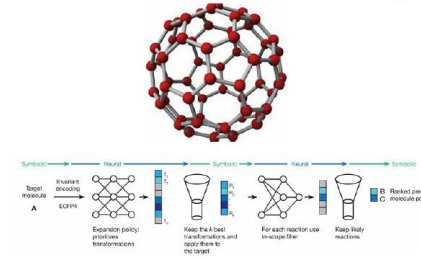
## Fluid, Gas Dynamics



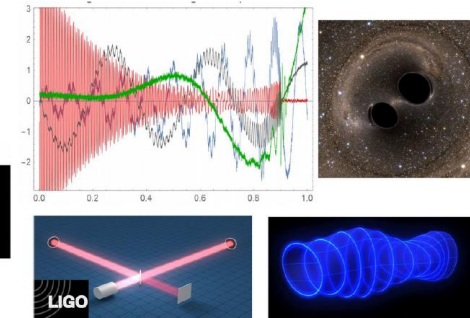
## Protein Folding



## Material Science, Chemistry



## Gravitational Waves Detection



## Games, Control Optimization



# Complex Relationships: ML & DL vs. HPC/Clouds & Big Data

Turing Award 2018: Recognition of **Deep Neural Networks** as critical component of computing

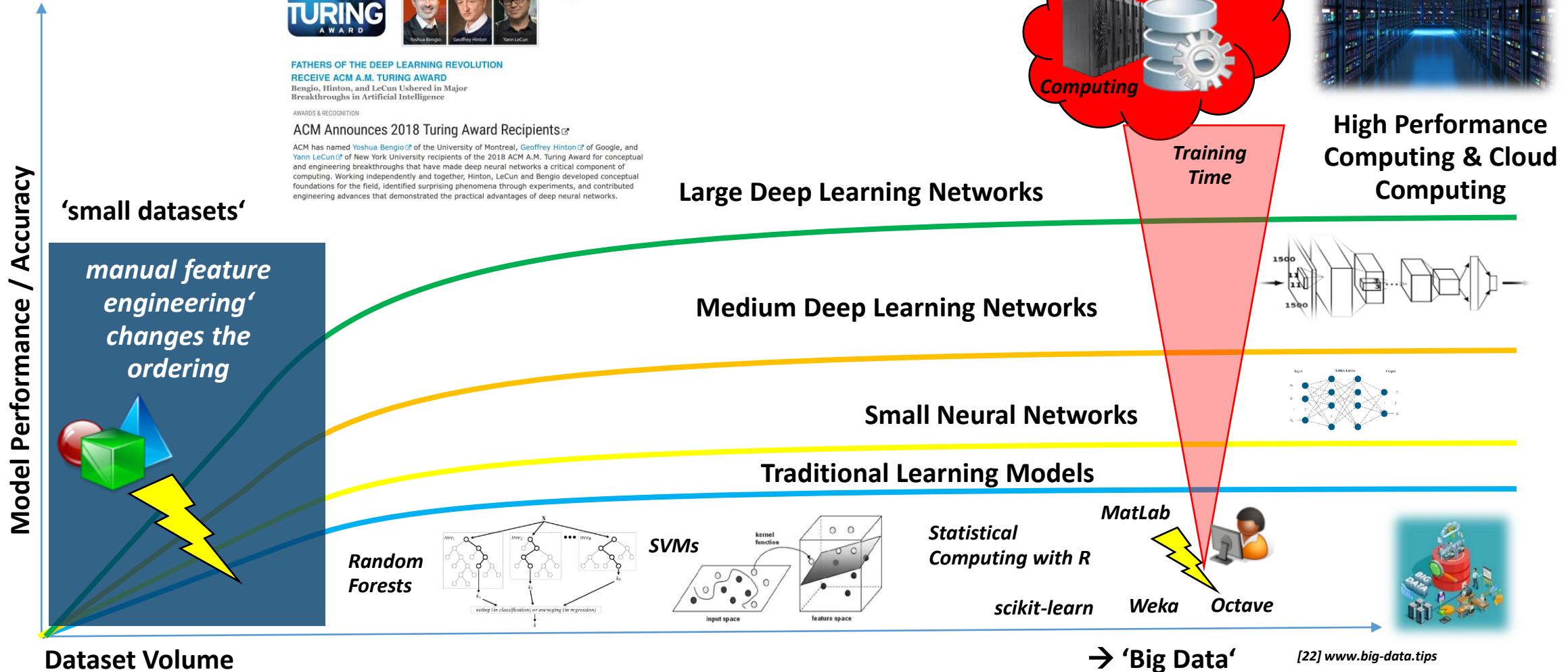


**FATHERS OF THE DEEP LEARNING REVOLUTION RECEIVE ACM A.M. TURING AWARD**  
Bengio, Hinton, and LeCun ushered in major breakthroughs in Artificial Intelligence

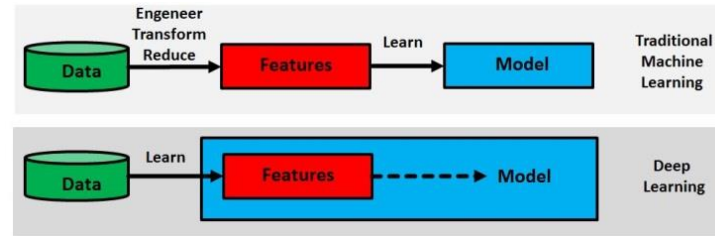
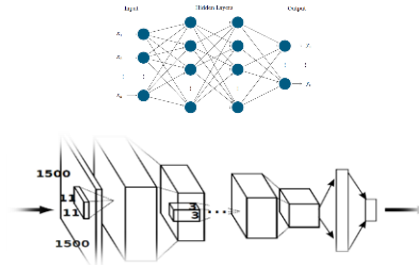
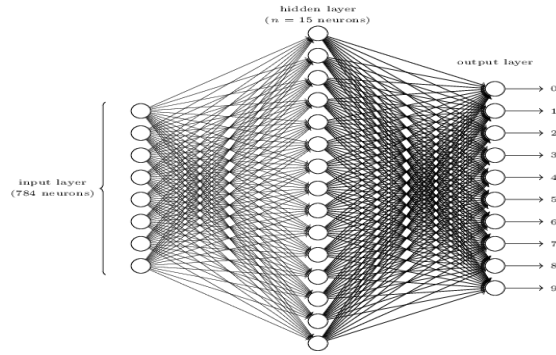
AWARDS & RECOGNITION

ACM Announces 2018 Turing Award Recipients

ACM has named [Yoshua Bengio](#) of the University of Montreal, [Geoffrey Hinton](#) of Google, and [Yann LeCun](#) of New York University recipients of the 2018 ACM A.M. Turing Award for conceptual and engineering breakthroughs that have made deep neural networks a critical component of computing. Working independently and together, Hinton, LeCun and Bengio developed conceptual foundations for the field, identified surprising phenomena through experiments, and contributed engineering advances that demonstrated the practical advantages of deep neural networks.



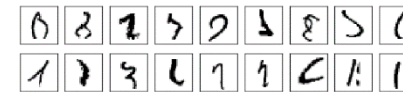
# Innovative Deep Learning Techniques – Revisited



[23] M. Riedel, 'Deep Learning - Using a Convolutional Neural Network', Invited YouTube Lecture, six lectures, University of Ghent, 2017

[24] M. Riedel et al., 'Introduction to Deep Learning Models', JSC Tutorial, three days, JSC, 2019

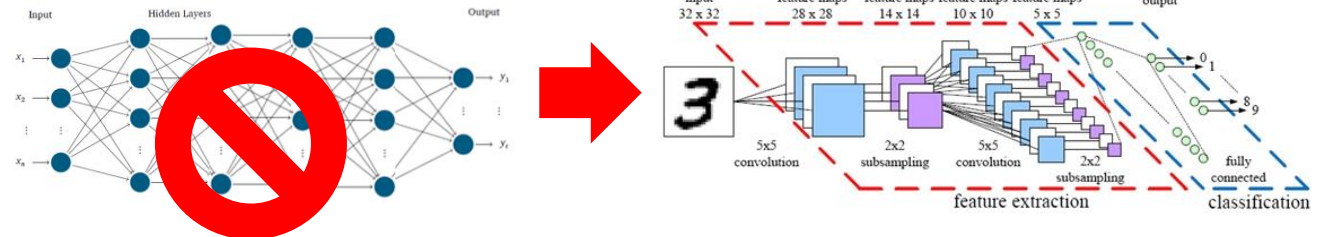
■ Innovation via specific layers and architecture types is the success of Convolutional Neural Networks (CNNs)



[27] A. Rosebrock



[25] H. Lee et al., 'Convolutional Deep Belief Networks for Scalable Unsupervised Learning of Hierarchical Representations'



[26] Neural Network 3D Simulation

www.cybercontrols.org

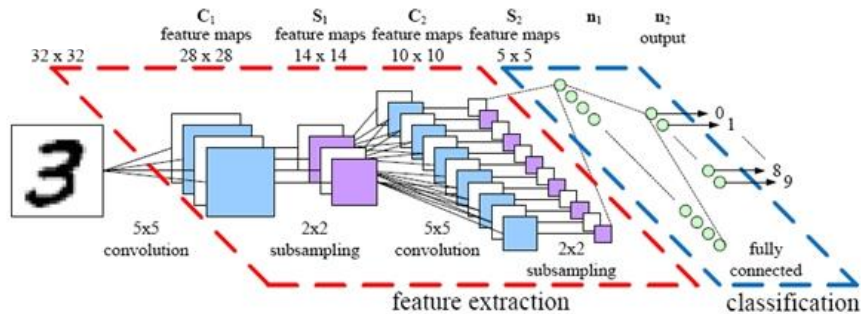


# CNNs – Basic Principles & Local Receptive Fields

- Convolutional Neural Networks (CNNs/ConvNets) implement a connectivity pattern between neurons inspired by the animal visual cortex and use several types of layers (convolution, pooling)
- CNN key principles are local receptive fields, shared weights, and pooling (or down/sub-sampling)
- CNNs are optimized to take advantage of the spatial structure of the data

## Simple application example

- MNIST database written characters
- Use CNN architecture with different layers
- Goal: automatic classification of characters



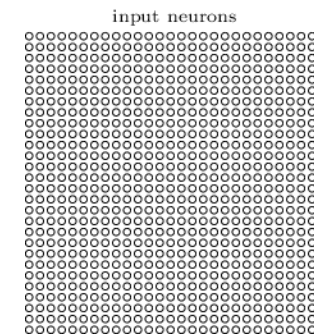
[27] A. Rosebrock



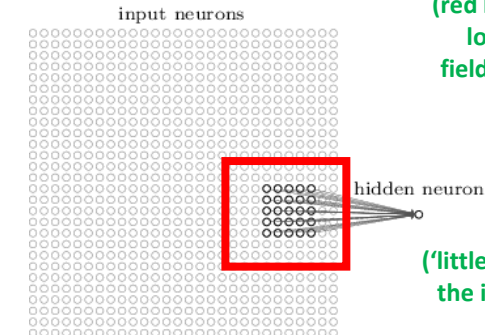
[28] M. Nielsen

## MNIST dataset example

- 28 \* 28 pixels modeled as square of neurons in a convolutional net
- Values correspond to the 28 \* 28 pixel intensities as inputs



(28 \* 28 pixel image)



(5 \* 5 local connectivity)

(red box indicate the local receptive field for the hidden neuron)

('little window' on the input pixels)



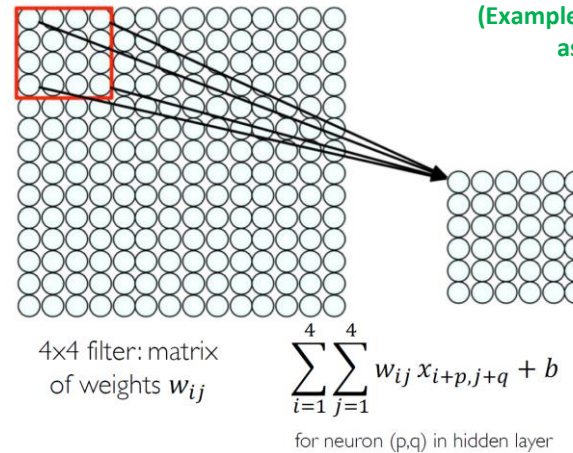
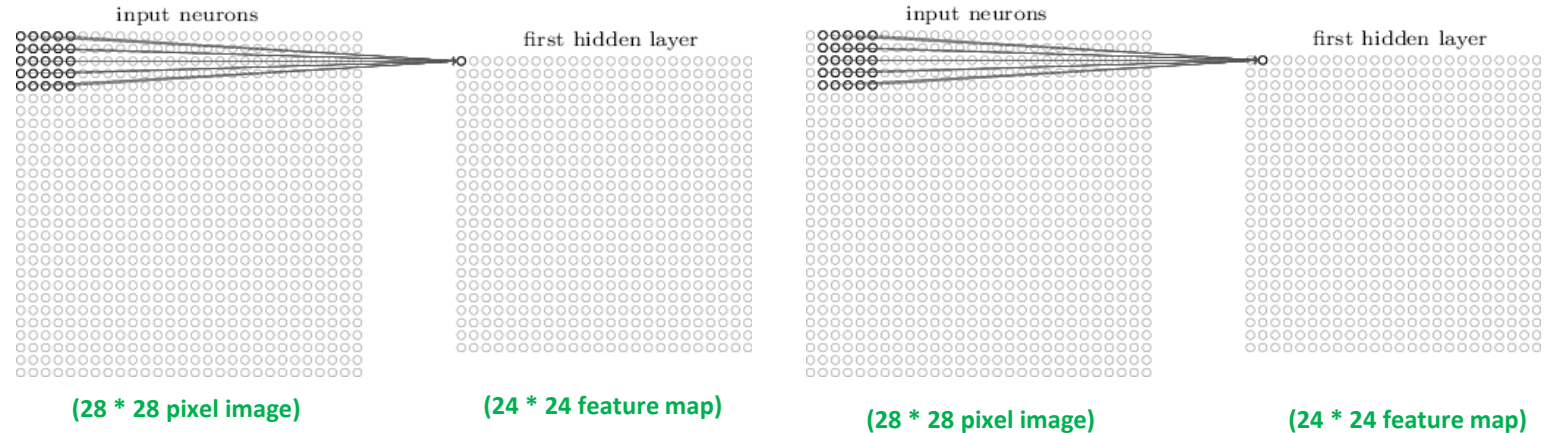
# CNNs – Principle Sliding with Convolutions & Feature Maps

- MNIST database example

- Apply stride length = 1
- Creates 'feature map' of 24 \* 24 neurons (hidden layer)

- Role of Convolutions & Filter

- Valid convolution does not exceed the input's boundary
- Same convolution adds a so called 'padding' to maintain the input's dimension for each convolutional layer
- Feature maps reflect where in the input a part of local features were activated by the applied filter



(Example: apply a set of weights as filter to extract local features)

[17] Introduction to Deep Learning

1 <sub>x1</sub>	1 <sub>x0</sub>	1 <sub>x1</sub>	0	0
0 <sub>x0</sub>	1 <sub>x1</sub>	1 <sub>x0</sub>	1	0
0 <sub>x1</sub>	0 <sub>x0</sub>	1 <sub>x1</sub>	1	1
0	0	1	1	0
0	1	1	0	0

Image

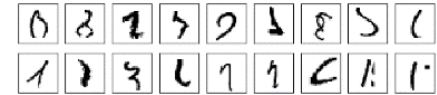
4		

Convolved Feature

# CNNs – Understanding Application Example MNIST - Summary

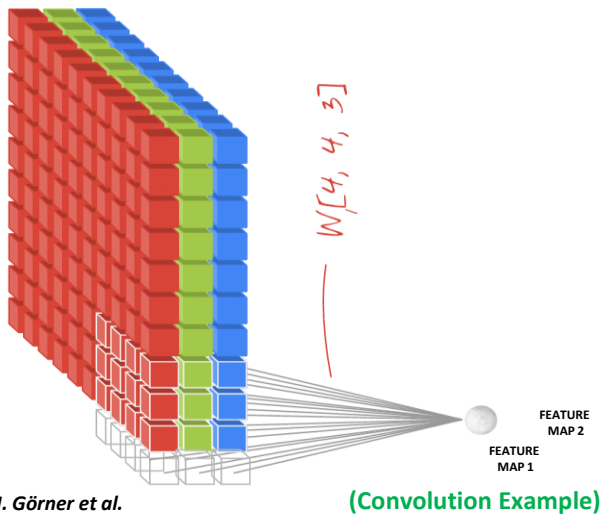
- MNIST database example

- Pooling Layer & Apply ‘fully connected layer (flatten)’ : layer connects every neuron from the max-pooling outcome layer to every neuron of the 10 out neurons

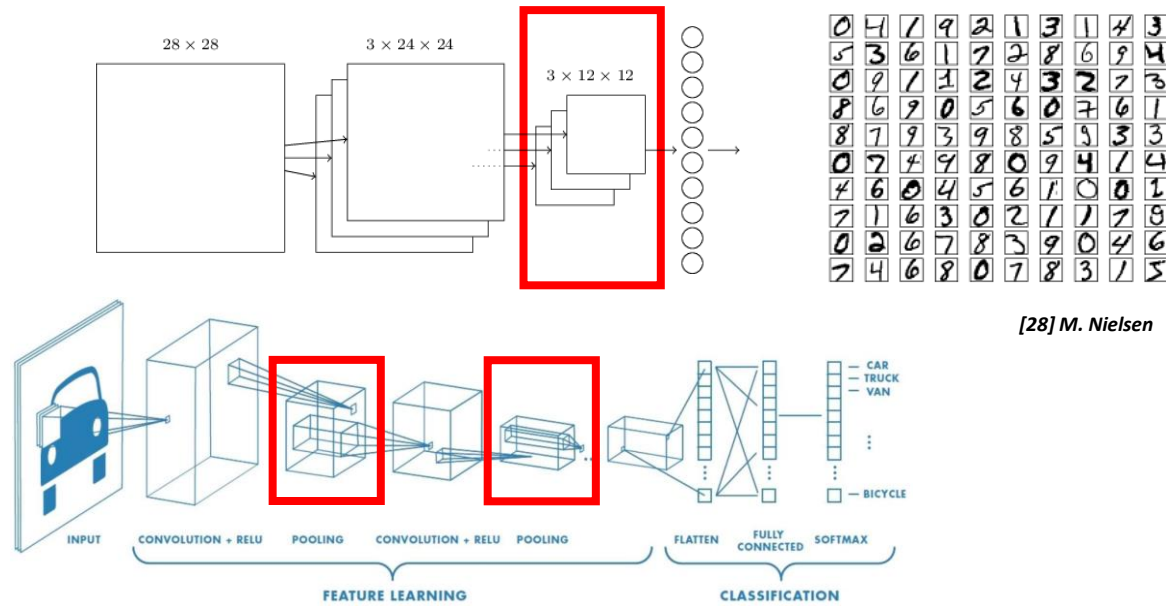


(another indicator that even with cutting edge technology machine learning never achieves 100% performance)

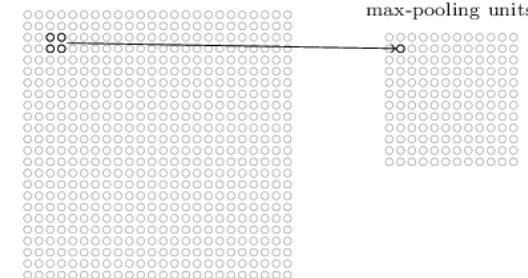
(Pooling layers: simplify the information in the output from the Convolution, e.g. max pooling)



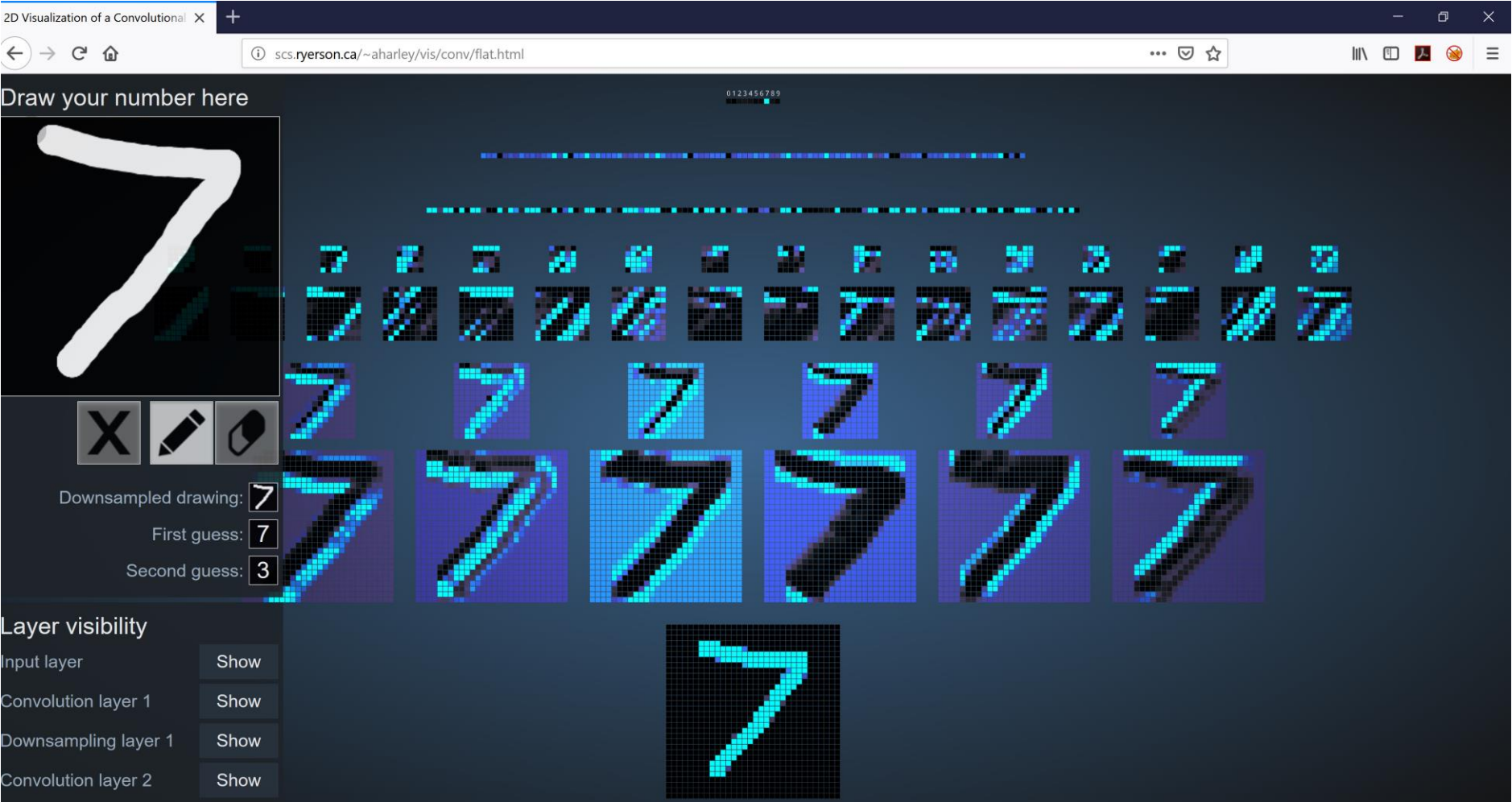
[29] M. Görner et al.



hidden neurons (output from feature map)



# Understanding Feature Maps & Convolutions Summary – Online Web Tool



[30] Harley, A.W., An Interactive Node-Link Visualization of Convolutional Neural Networks



# MNIST Dataset – Convolutional Neural Network (CNN) Model

```

from keras import backend as K
from keras.models import Sequential
from keras.layers.convolutional import Conv2D
from keras.layers.convolutional import MaxPooling2D
from keras.layers.core import Activation
from keras.layers.core import Flatten
from keras.layers.core import Dense
from keras.datasets import mnist
from keras.utils import np_utils
from keras.optimizers import SGD, RMSprop, Adam
import numpy as np
import matplotlib.pyplot as plt

```

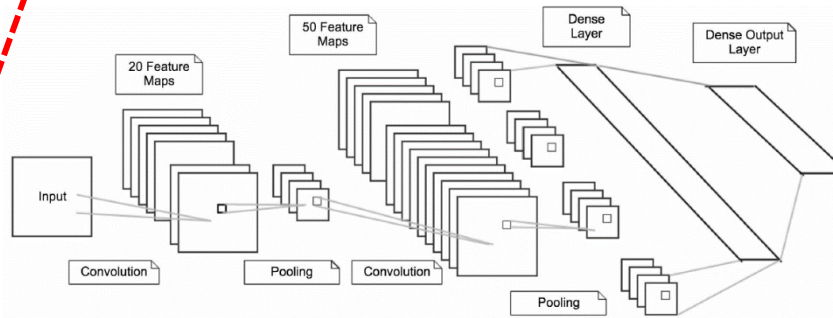
```

#define the CNN model
class CNN:
    @staticmethod
    def build(input_shape, classes):
        model = Sequential()
        # CONV => RELU => POOL
        model.add(Conv2D(20, kernel_size=5, padding="same",
            input_shape=input_shape))
        model.add(Activation("relu"))
        model.add(MaxPooling2D(pool_size=(2, 2), strides=(2, 2)))
        # CONV => RELU => POOL
        model.add(Conv2D(50, kernel_size=5, border_mode="same"))
        model.add(Activation("relu"))
        model.add(MaxPooling2D(pool_size=(2, 2), strides=(2, 2)))
        # Flatten => RELU layers
        model.add(Flatten())
        model.add(Dense(500))
        model.add(Activation("relu"))
        # a softmax classifier
        model.add(Dense(classes))
        model.add(Activation("softmax"))
        return model

```

- Increasing the number of filters learned to 50 in the next layer from 20 in the first layer
- Increasing the number of filters in deeper layers is a common technique in deep learning architecture modeling
- Flattening the output as input for a Dense layer (fully connected layer)
- Fully connected / Dense layer responsible with softmax activation for classification based on learned filters and features

[31] A. Gulli et al.



```

# initialize the optimizer and model
model = CNN.build(input_shape=INPUT_SHAPE, classes=NB_CLASSES)
model.compile(loss="categorical_crossentropy", optimizer=OPTIMIZER,
    metrics=["accuracy"])

```

```

# printout a summary of the model to understand model complexity
model.summary()

```

Layer (type)	Output Shape	Param #
conv2d_1 (Conv2D)	(None, 20, 28, 28)	520
activation_1 (Activation)	(None, 20, 28, 28)	0
max_pooling2d_1 (MaxPooling2D)	(None, 20, 14, 14)	0
conv2d_2 (Conv2D)	(None, 50, 14, 14)	25050
activation_2 (Activation)	(None, 50, 14, 14)	0
max_pooling2d_2 (MaxPooling2D)	(None, 50, 7, 7)	0
flatten_1 (Flatten)	(None, 2450)	0
dense_1 (Dense)	(None, 500)	1225500
activation_3 (Activation)	(None, 500)	0
dense_2 (Dense)	(None, 10)	5010
activation_4 (Activation)	(None, 10)	0

```

Total params: 1,256,080
Trainable params: 1,256,080
Non-trainable params: 0

```

# MNIST Dataset – Model Parameters & 2D Input Data

```
# parameter setup
NB_EPOCH = 20
BATCH_SIZE = 128
VERBOSE = 1
OPTIMIZER = Adam()
VALIDATION_SPLIT=0.2
IMG_ROWS, IMG_COLS = 28, 28 # input image dimensions
NB_CLASSES = 10 # number of outputs = number of digits
INPUT_SHAPE = (1, IMG_ROWS, IMG_COLS)
```

```
# data: shuffled and split between train and test sets
(X_train, y_train), (X_test, y_test) = mnist.load_data()
K.set_image_dim_ordering("th")
# consider them as float and normalize
X_train = X_train.astype('float32')
X_test = X_test.astype('float32')
X_train /= 255
X_test /= 255
```

```
# we need a 60K x [1 x 28 x 28] shape as input to the CONVNET
X_train = X_train[:, np.newaxis, :, :]
X_test = X_test[:, np.newaxis, :, :]
print(X_train.shape[0], 'train samples')
print(X_test.shape[0], 'test samples')
# convert class vectors to binary class matrices
y_train = np_utils.to_categorical(y_train, NB_CLASSES)
y_test = np_utils.to_categorical(y_test, NB_CLASSES)
```

- **OPTIMIZER:** Adam - advanced optimization technique that includes the concept of a momentum (a certain velocity component) in addition to the acceleration component of Stochastic Gradient Descent (SGD)
- Adam computes individual adaptive learning rates for different parameters from estimates of first and second moments of the gradients
- Adam enables faster convergence at the cost of more computation and is currently recommended as the default algorithm to use (or SGD + Nesterov Momentum)

[32] D. Kingma et al., 'Adam: A Method for Stochastic Optimization'

- Compared to the Multi-Output Perceptron and Artificial Neural Networks (ANN) model, the input dataset remains as 2d matrice with 1 x 28 x 28 per image, including also the class vectors that are converted to binary class matrices



# MNIST Dataset – CNN Model Output & Evaluation

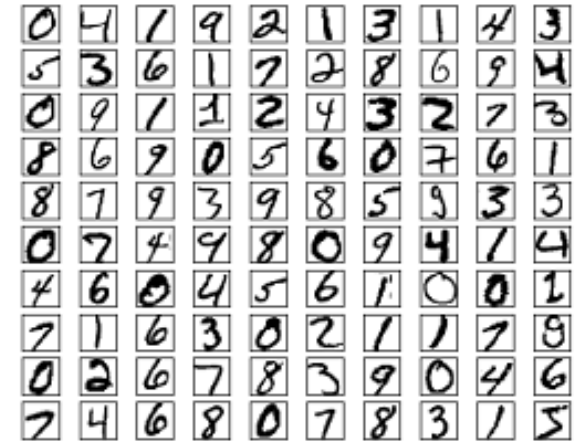
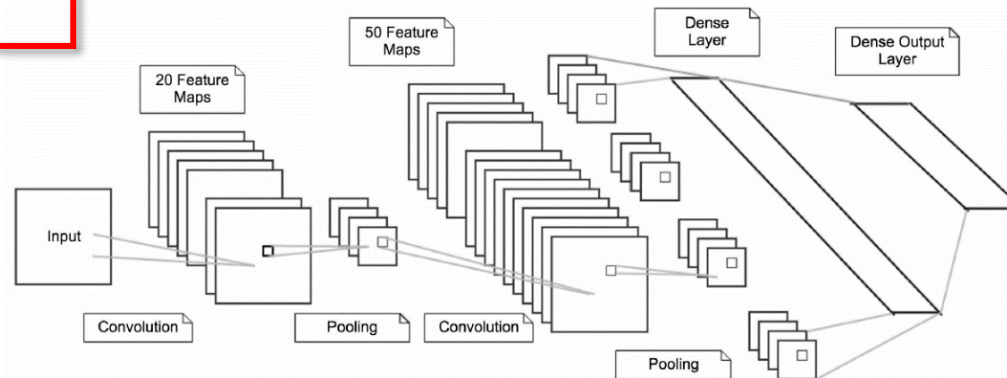
```
Epoch 14/20
48000/48000 [=====] - 4s 88us/step - loss: 0.0065 - acc: 0.9980 - val_loss: 0.0346 - val_acc: 0.9921
Epoch 15/20
48000/48000 [=====] - 4s 89us/step - loss: 0.0030 - acc: 0.9990 - val_loss: 0.0418 - val_acc: 0.9903
Epoch 16/20
48000/48000 [=====] - 4s 88us/step - loss: 0.0057 - acc: 0.9980 - val_loss: 0.0470 - val_acc: 0.9910
Epoch 17/20
48000/48000 [=====] - 4s 88us/step - loss: 0.0043 - acc: 0.9985 - val_loss: 0.0440 - val_acc: 0.9906
Epoch 18/20
48000/48000 [=====] - 4s 88us/step - loss: 0.0046 - acc: 0.9985 - val_loss: 0.0474 - val_acc: 0.9891
Epoch 19/20
48000/48000 [=====] - 4s 88us/step - loss: 0.0047 - acc: 0.9986 - val_loss: 0.0353 - val_acc: 0.9928
Epoch 20/20
48000/48000 [=====] - 4s 88us/step - loss: 3.4055e-04 - acc: 1.0000 - val_loss: 0.0374 - val_acc: 0.9927
```

```
# model evaluation
score = model.evaluate(X_test, y_test, verbose=VERBOSE)
print("Test score:", score[0])
print('Test accuracy:', score[1])
```

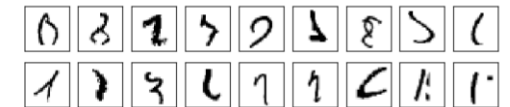
```
10000/10000 [=====] - 1s 70us/step
Test score: 0.0303058747581508
Test accuracy: 0.9936
```

- ✓ **Multi Output Perceptron:**  
~91,01% (20 Epochs)
- ✓ **ANN 2 Hidden Layers:**  
~95,14 % (20 Epochs)
- ✓ **CNN Deep Learning Model:**  
~99,36 % (20 Epochs)

[31] A. Gulli et al.

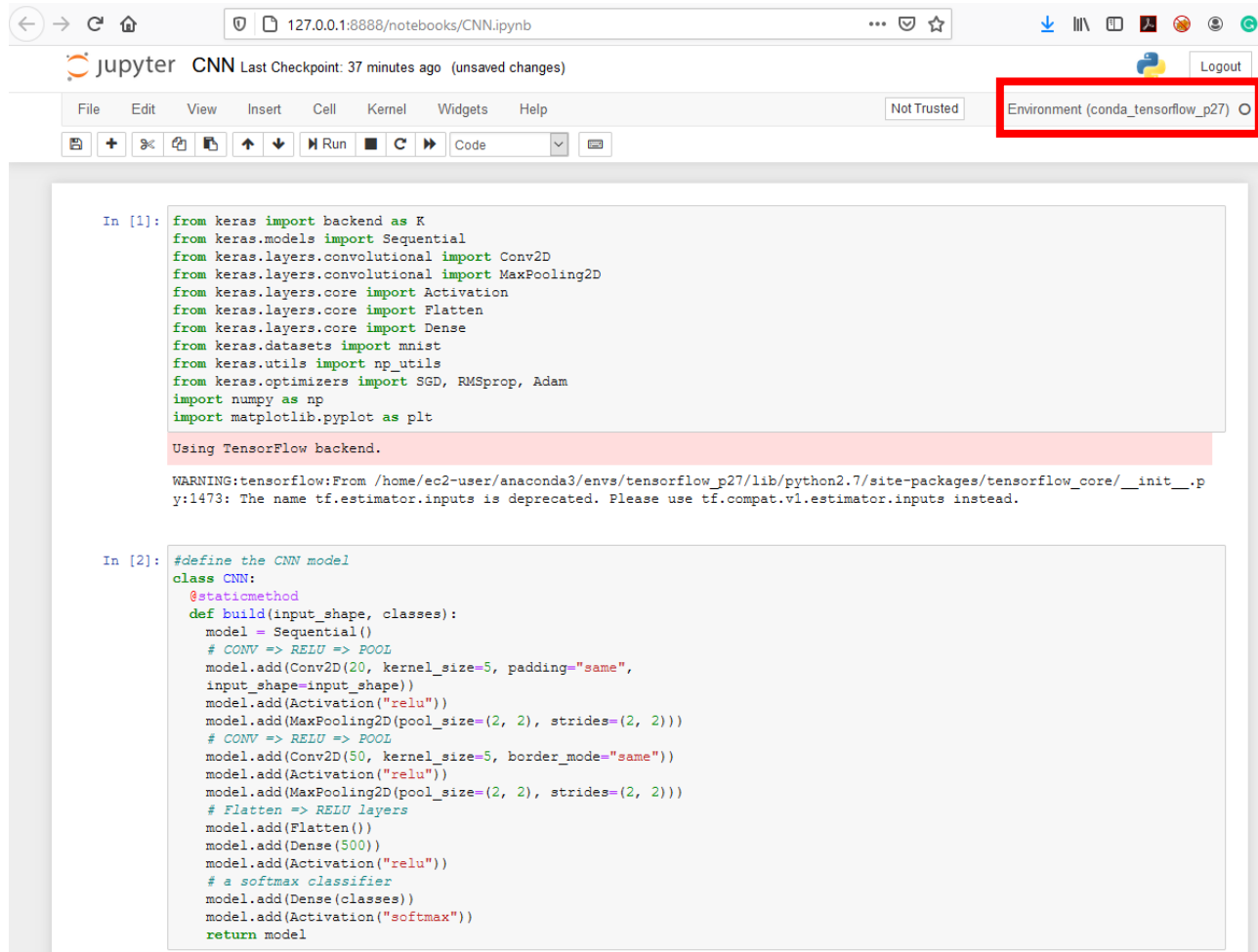


Why not 100%



some samples even for a human unrecognizable

# Running a Deep Learning Model with Convolutional Neural Network (CNN)



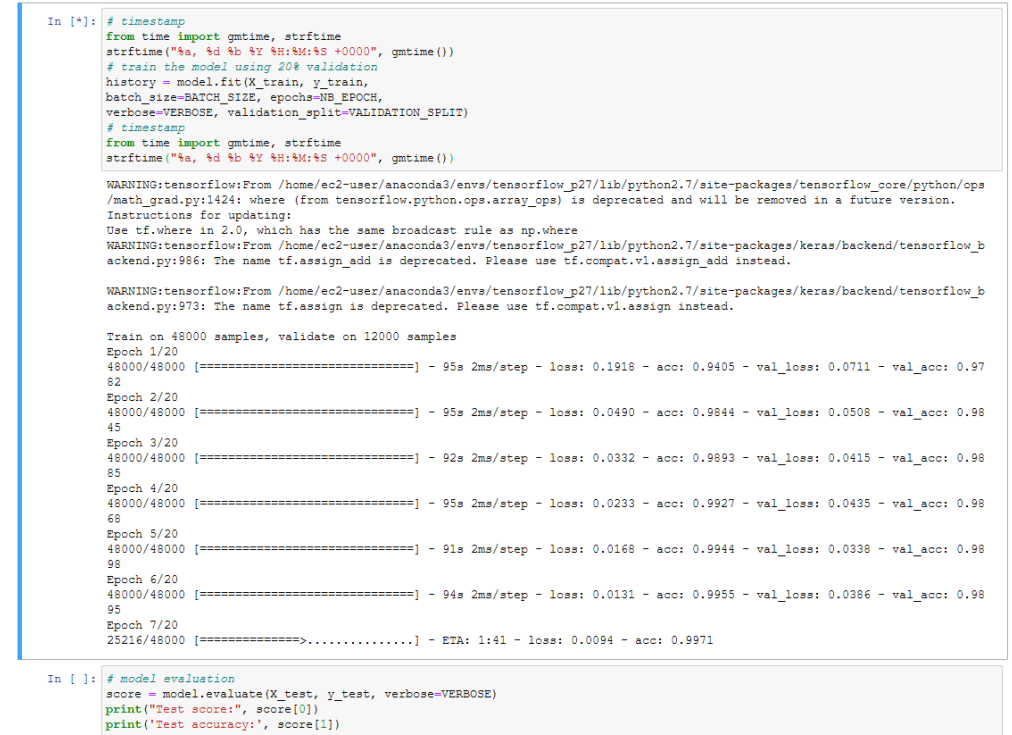
The screenshot shows a Jupyter Notebook titled 'CNN' with a 'conda\_tensorflow\_p27' environment. The code in the first cell imports necessary libraries like keras, tensorflow, and numpy. The second cell defines a CNN class with a build method that constructs a neural network with two convolutional layers, pooling, and a softmax classifier.

```
In [1]: from keras import backend as K
from keras.models import Sequential
from keras.layers.convolutional import Conv2D
from keras.layers.convolutional import MaxPooling2D
from keras.layers.core import Activation
from keras.layers.core import Flatten
from keras.layers.core import Dense
from keras.datasets import mnist
from keras.utils import np_utils
from keras.optimizers import SGD, RMSprop, Adam
import numpy as np
import matplotlib.pyplot as plt

Using TensorFlow backend.

WARNING:tensorflow:From /home/ec2-user/anaconda3/envs/tensorflow_p27/lib/python2.7/site-packages/tensorflow_core/__init__.py:1473: The name tf.estimator.inputs is deprecated. Please use tf.compat.v1.estimator.inputs instead.

In [2]: #define the CNN model
class CNN:
    @staticmethod
    def build(input_shape, classes):
        model = Sequential()
        # CONV -> RELU -> POOL
        model.add(Conv2D(20, kernel_size=5, padding="same",
            input_shape=input_shape))
        model.add(Activation("relu"))
        model.add(MaxPooling2D(pool_size=(2, 2), strides=(2, 2)))
        # CONV -> RELU -> POOL
        model.add(Conv2D(50, kernel_size=5, border_mode="same"))
        model.add(Activation("relu"))
        model.add(MaxPooling2D(pool_size=(2, 2), strides=(2, 2)))
        # Flatten -> RELU layers
        model.add(Flatten())
        model.add(Dense(500))
        model.add(Activation("relu"))
        # a softmax classifier
        model.add(Dense(classes))
        model.add(Activation("softmax"))
        return model
```



The screenshot shows the output of the training process. It includes a timestamp, training progress for 7 epochs, and a final evaluation step. The output shows that the model's accuracy is approximately 99.71% after 7 epochs.

```
In [4]: # timestamp
from time import gmtime, strftime
strftime("%a, %d %b %Y %H:%M:%S +0000", gmtime())
# train the model using 20# validation
history = model.fit(X_train, y_train,
    batch_size=BATCH_SIZE, epochs=NB_EPOCH,
    verbose=VERBOSE, validation_split=VALIDATION_SPLIT)
# timestamp
from time import gmtime, strftime
strftime("%a, %d %b %Y %H:%M:%S +0000", gmtime())

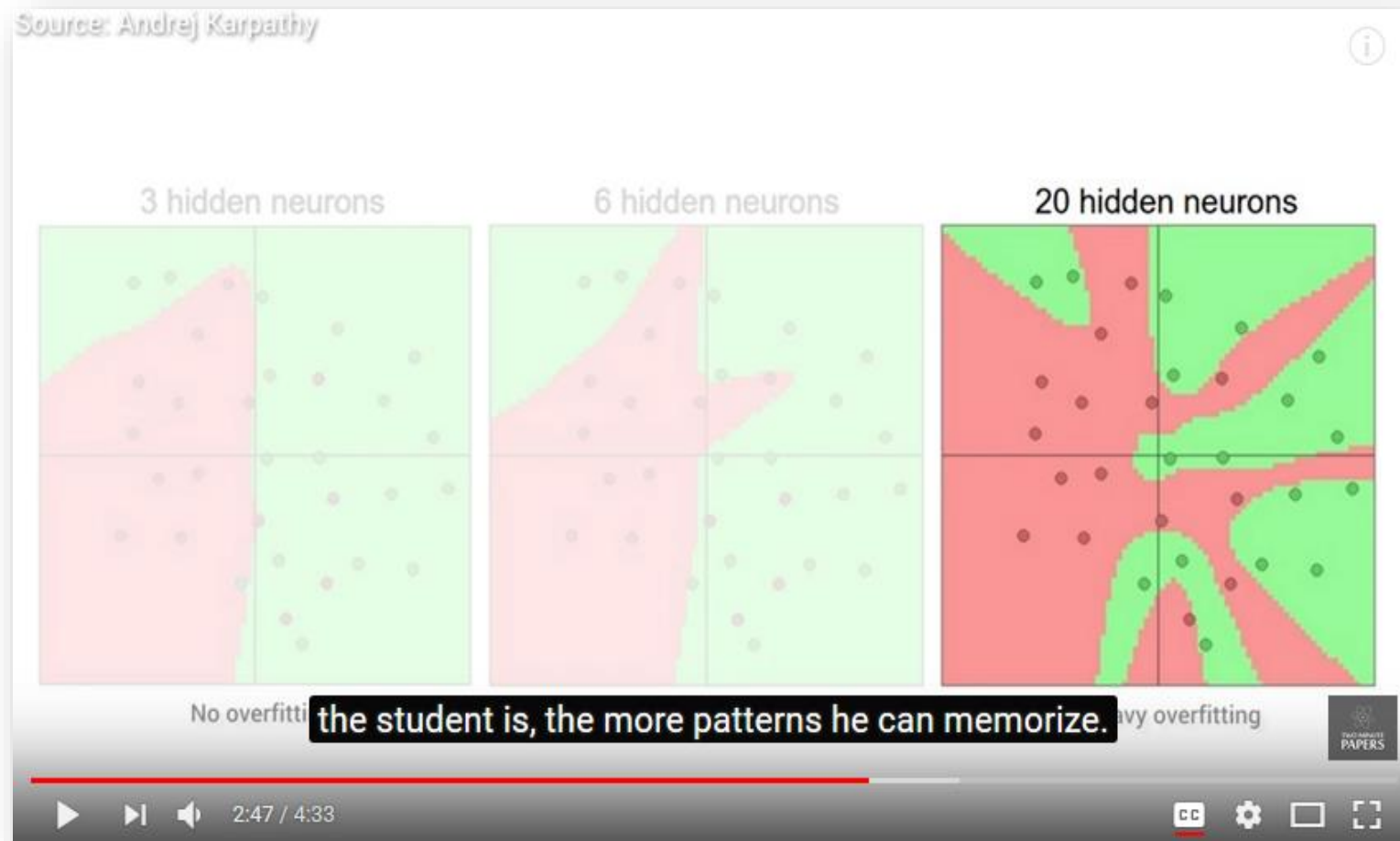
WARNING:tensorflow:From /home/ec2-user/anaconda3/envs/tensorflow_p27/lib/python2.7/site-packages/tensorflow_core/python/ops/math_grad.py:1424: where (from tensorflow.python.ops.array_ops) is deprecated and will be removed in a future version.
Instructions for updating:
Use tf.where in 2.0, which has the same broadcast rule as np.where
WARNING:tensorflow:From /home/ec2-user/anaconda3/envs/tensorflow_p27/lib/python2.7/site-packages/keras/backend/tensorflow_backend.py:986: The name tf.assign_add is deprecated. Please use tf.compat.v1.assign_add instead.
WARNING:tensorflow:From /home/ec2-user/anaconda3/envs/tensorflow_p27/lib/python2.7/site-packages/keras/backend/tensorflow_backend.py:973: The name tf.assign is deprecated. Please use tf.compat.v1.assign instead.

Train on 48000 samples, validate on 12000 samples
Epoch 1/20
48000/48000 [=====] - 95s 2ms/step - loss: 0.1918 - acc: 0.9405 - val_loss: 0.0711 - val_acc: 0.9782
Epoch 2/20
48000/48000 [=====] - 95s 2ms/step - loss: 0.0490 - acc: 0.9844 - val_loss: 0.0508 - val_acc: 0.9845
Epoch 3/20
48000/48000 [=====] - 92s 2ms/step - loss: 0.0332 - acc: 0.9893 - val_loss: 0.0415 - val_acc: 0.9885
Epoch 4/20
48000/48000 [=====] - 95s 2ms/step - loss: 0.0233 - acc: 0.9927 - val_loss: 0.0435 - val_acc: 0.9868
Epoch 5/20
48000/48000 [=====] - 91s 2ms/step - loss: 0.0168 - acc: 0.9944 - val_loss: 0.0338 - val_acc: 0.9898
Epoch 6/20
48000/48000 [=====] - 94s 2ms/step - loss: 0.0131 - acc: 0.9955 - val_loss: 0.0386 - val_acc: 0.9895
Epoch 7/20
25216/48000 [=====>.....] - ETA: 1:41 - loss: 0.0094 - acc: 0.9971

In [ ]: # model evaluation
score = model.evaluate(X_test, y_test, verbose=VERBOSE)
print("Test score:", score[0])
print("Test accuracy:", score[1])
```

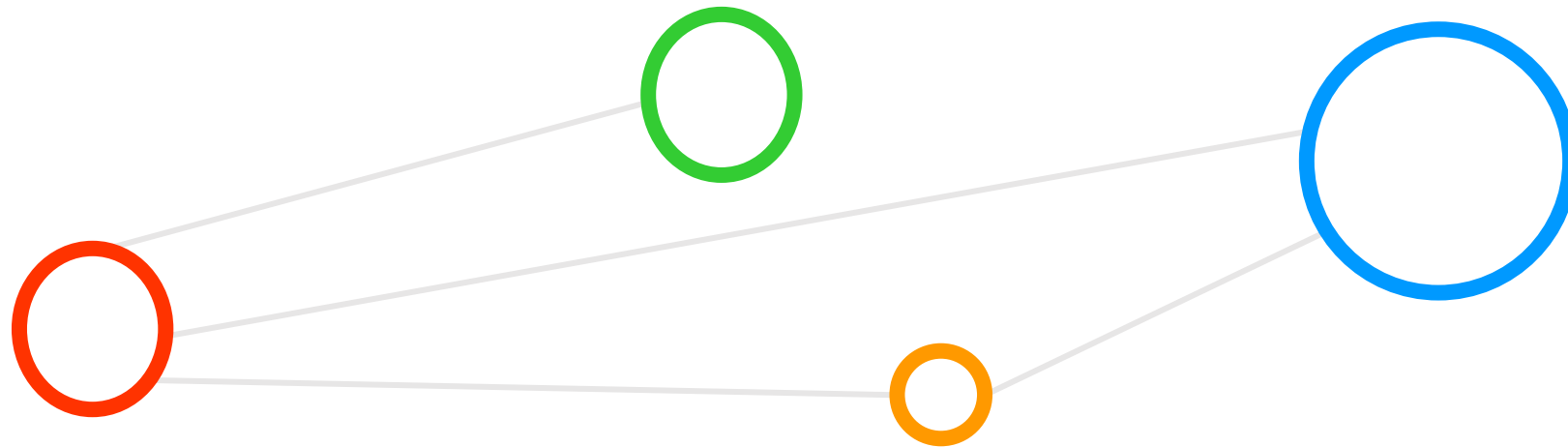
- Using Deep Learning Techniques such as Convolutional Neural Networks (CNNs) in clouds can lead to significant improvements in accuracy, but also to significant longer run-times than traditional Artificial Neural Networks (ANNs) and are thus much more costly in clouds
- Using CPU resources for deep learning techniques is usually not recommended

# [Video for further Studies] Overfitting in Deep Neural Networks



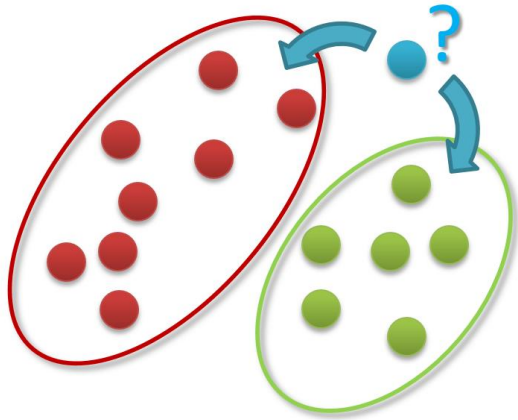
[33] YouTube Video, Overfitting and Regularization For Deep Learning

# Selected Parallel & Scalable Machine & Deep Learning Techniques



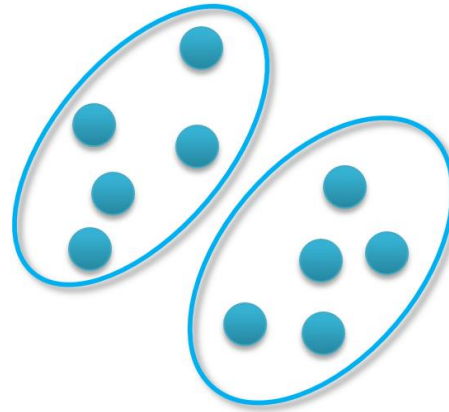
# Machine Learning Models – Understanding Parallel Benefits

## Classification



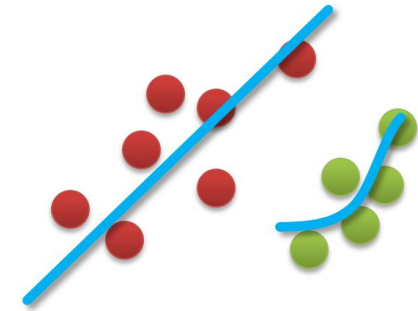
- Groups of data exist
- New data classified to existing groups

## Clustering



- No groups of data exist
- Create groups from data close to each other

## Regression



- Identify a line with a certain slope describing the data

▪ Machine learning methods can be roughly categorized in classification, clustering, or regression augmented with various techniques for data exploration, selection, or reduction – despite the momentum of deep learning, traditional machine learning algorithms are still widely relevant today

[1] [www.big-data.tips](http://www.big-data.tips), 'Data Classification'



# Terminologies & Different Dataset Elements & Processes – Overview

## Machine Learning Models

- Based on various algorithms that learn from existing data sets

## Labelled Dataset (samples)

- 'in-sample' data given to us:  $(\mathbf{x}_1, y_1), \dots, (\mathbf{x}_N, y_N)$

## Learning vs. Memorizing

- The goal is to create a system that works well 'out of sample'
- In other words we want to classify 'future data' (out of sample) correct

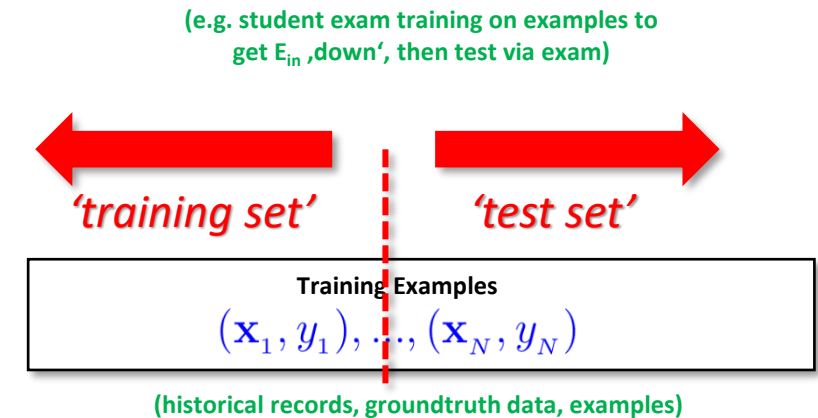
## Dataset Part One: Training set

- Used for training a machine learning algorithms
- Result after using a training set: a trained system

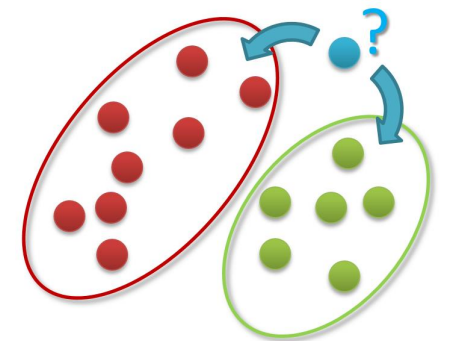
## Dataset Part Two: Test set

- Used for testing whether the trained system might work well
- Result after using a test set: accuracy of the trained model

(exact separation is rule of thumb, but different in each data analysis case: e.g., 10% training data, 90% test data)

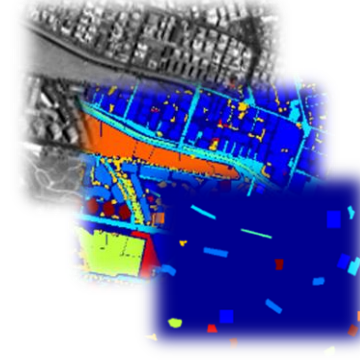


## Classification

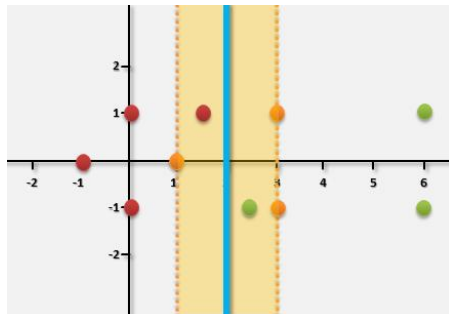
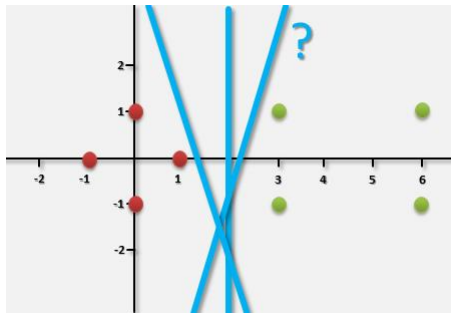


# Parallel and Scalable Machine Learning – Parallel Support Vector Machine (SVM)

- ‘Different kind’ of parallel algorithms
  - ‘learn from data’ instead of modelling/approximate reality with physics
  - Parallel algorithms often useful to reduce ‘overall time for data analysis’
- E.g. Parallel Support Vector Machines (SVMs) Technique
  - Data classification algorithm PiSVM using MPI to reduce ‘training time’
  - Example: classification of land cover masses from satellite image data



Class	Training	Test
Buildings	18126	163129
Blocks	10982	98834
Roads	16353	147176
Light Train	1606	14454
Vegetation	6962	62655
Trees	9088	81792
Bare Soil	8127	73144
Soil	1506	13551
Tower	4792	43124
Total	77542	697859



[35] C. Cortes & V. Vapnik, ‘Support Vector Networks’, *Machine Learning*, 1995

[36] [www.big-data.tips](http://www.big-data.tips), ‘SVM Train’

```
#!/bin/bash -x
#SBATCH--nodes=4
#SBATCH--ntasks=96
#SBATCH--ntasks-per-node=24
#SBATCH--output=mpi-out.%j
#SBATCH--error=mpi-err.%j
#SBATCH--time=04:00:00
#SBATCH--partition=batch
#SBATCH--mail-user=m.riedel@fz-juelich.de
#SBATCH--mail-type=ALL
#SBATCH--job-name=train-indianpines-4-96-24
#SBATCH--reservation=ml-hpc-2

### location executable
PISVM=/homea/hpclub/train001/tools/pisvm-1.2.1/pisvm-train

### location data
TRAINDATA=/homea/hpclub/train001/data/indianpines/indian_raw_training.el

### submit
srun $PISVM -D -o 1024 -q 512 -c 100 -g 8 -t 2 -m 1024 -s 0 $TRAINDATA

#!/bin/bash -x
#SBATCH--nodes=4
#SBATCH--ntasks=96
#SBATCH--ntasks-per-node=24
#SBATCH--output=mpi-out.%j
#SBATCH--error=mpi-err.%j
#SBATCH--time=04:00:00
#SBATCH--partition=batch
#SBATCH--mail-user=m.riedel@fz-juelich.de
#SBATCH--mail-type=ALL
#SBATCH--job-name=pred-indianpines-4-96-24
#SBATCH--reservation=ml-hpc-2

### location executable
PISVMPRED=/homea/hpclub/train001/tools/pisvm-1.2.1/pisvm-predict

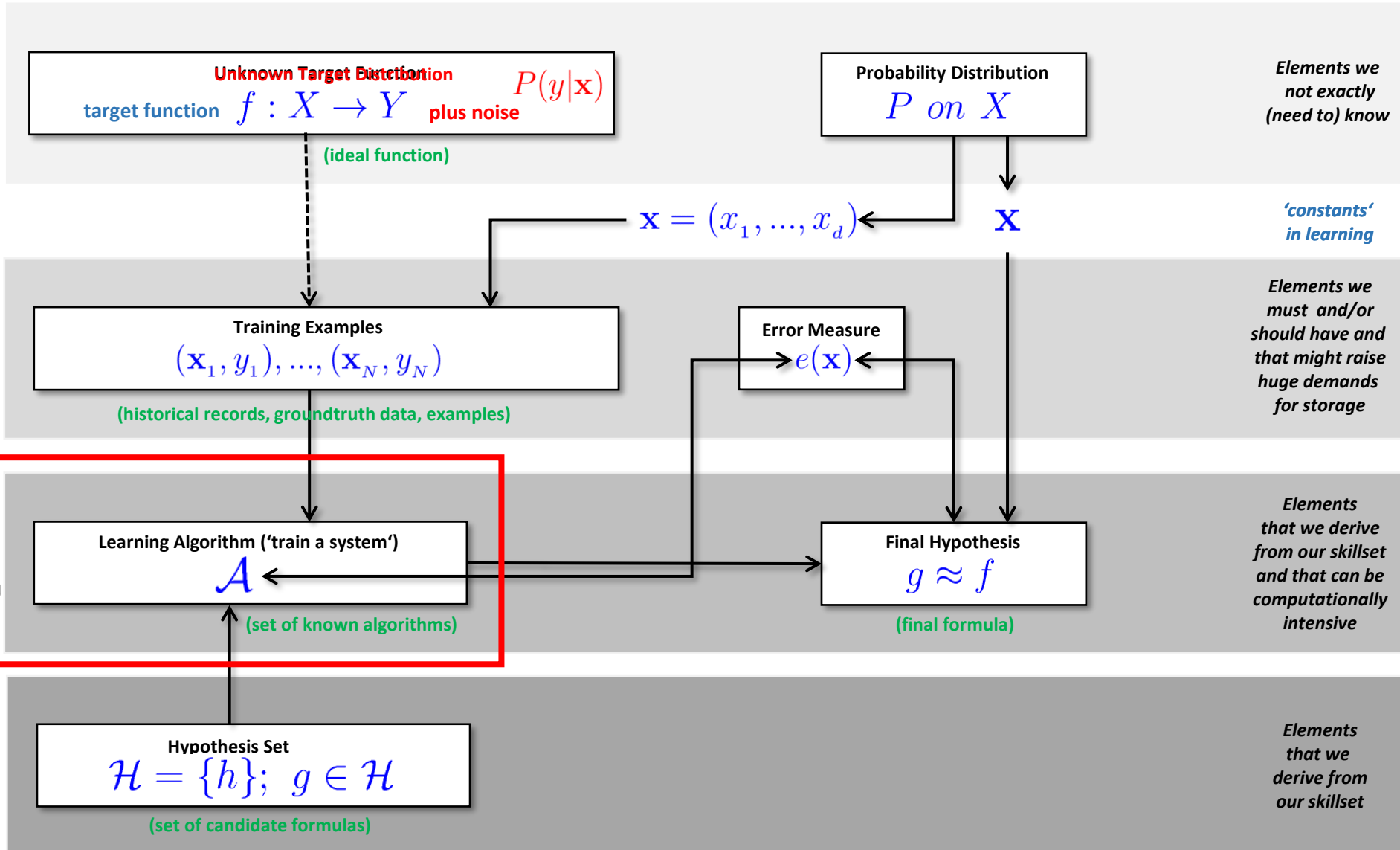
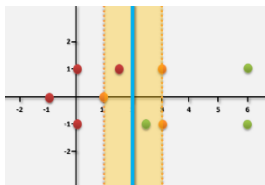
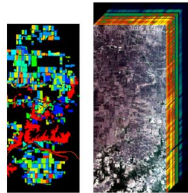
### location data
TESTDATA=/homea/hpclub/train001/data/indianpines/indian_raw_test.el

### trained model data
MODELDATA=/homea/hpclub/train001/tools/pisvm-1.2.1/indian_raw_training.el.model

### submit
srun $PISVMPRED $TESTDATA $MODELDATA results.txt
```

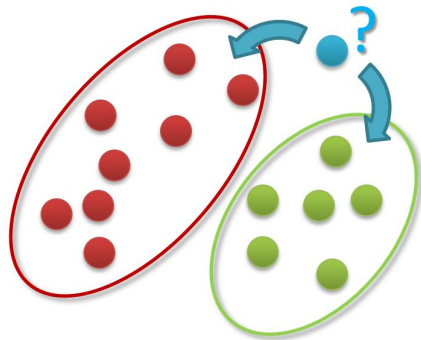
[34] G. Cavallaro & M. Riedel & J.A. Benediktsson et al., ‘On Understanding Big Data Impacts in Remotely Sensed Image Classification Using Support Vector Machine Methods’, *Journal of Applied Earth Observations and Remote Sensing*, 2015

# Overview Machine Learning Terminologies & Computing-intensive Processes

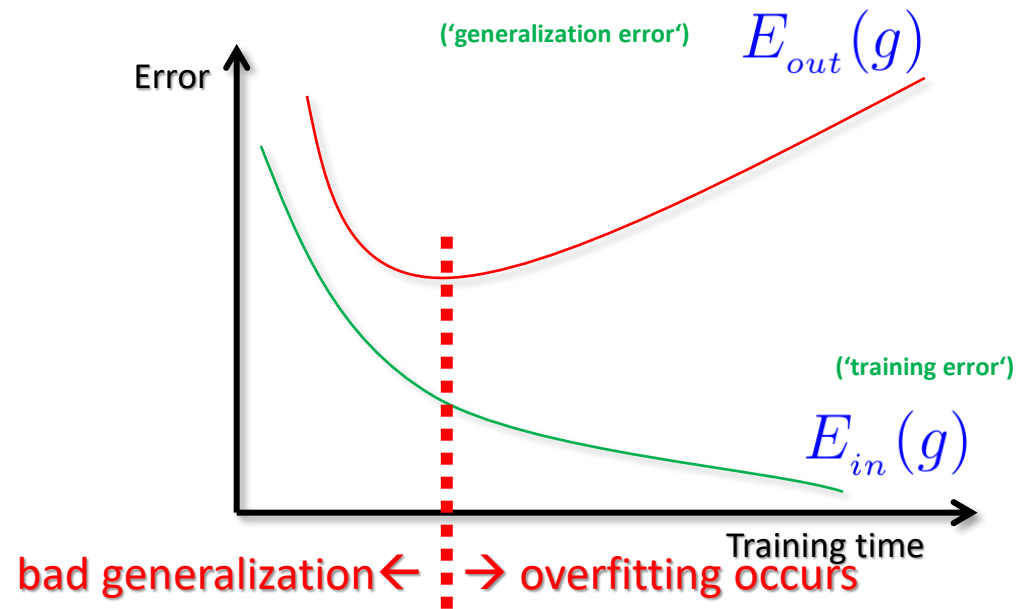


# Problem of Overfitting – Clarifying Terms

## Classification



$(\mathbf{x}_1, y_1), \dots, (\mathbf{x}_N, y_N)$



- A good model must have low training error ( $E_{in}$ ) and low generalization error ( $E_{out}$ )
- Model overfitting is if a model fits the data too well ( $E_{in}$ ) with a poorer generalization error ( $E_{out}$ ) than another model with a higher training error ( $E_{in}$ )
- The two general approaches to prevent overfitting are (1) validation and (2) regularization

[37] [www.big-data.tips](http://www.big-data.tips), 'Generalization in Machine Learning'

# Validation Technique – Proper Model Selection Process is Compute-Intensive

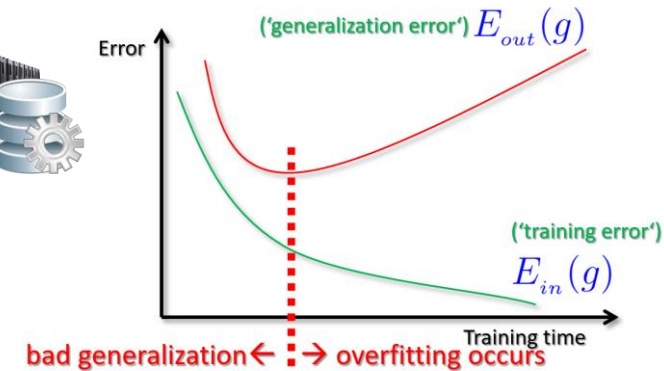
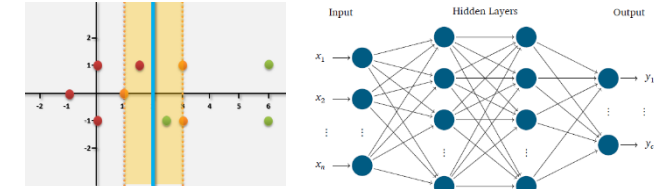
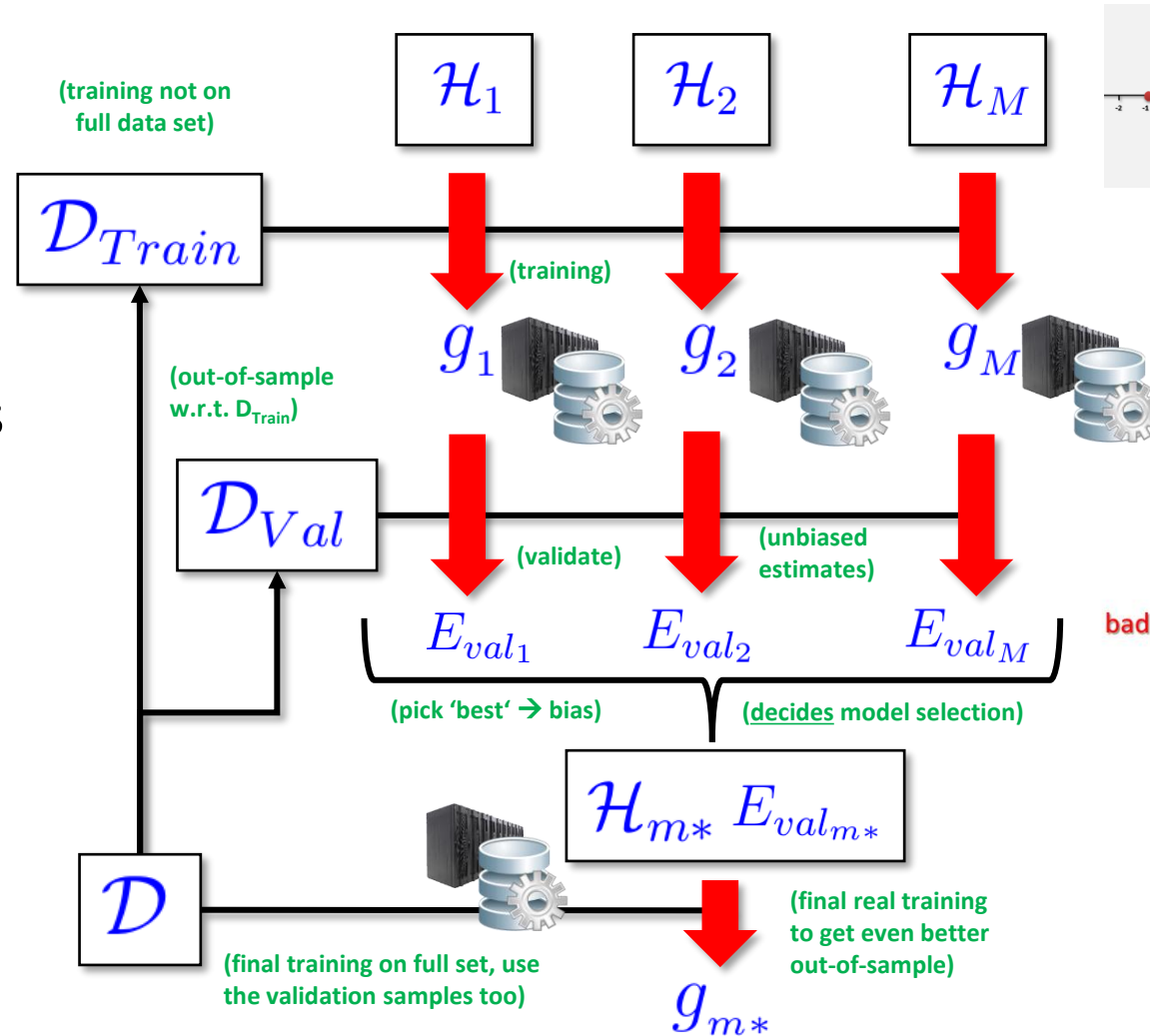
Hypothesis Set  
 $\mathcal{H} = \{h\}; g \in \mathcal{H}$

(set of candidate formulas across models)

- Many different models Use validation error to perform select decisions
- Careful consideration:
  - 'Picked means decided' hypothesis has already bias ( $\rightarrow$  contamination)
  - Using  $\mathcal{D}_{Val}$  M times

Final Hypothesis  
 $g_{m^*} \approx f$

(test this on unseen data good, but depends on availability in practice)



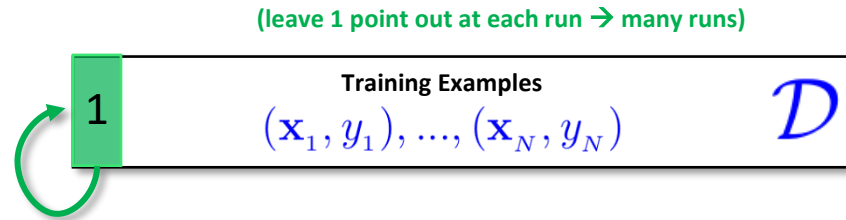
- Model selection is choosing (a) different types of models or (b) parameter values inside models
- Model selection takes advantage of the validation error in order to decide  $\rightarrow$  'pick the best'



# Validation Technique – Cross-Validation – Leave-more-out

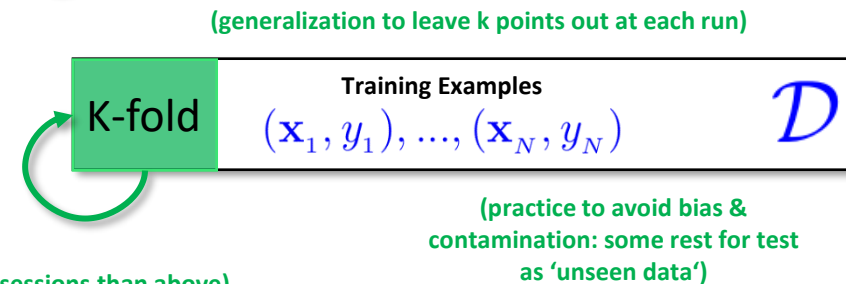
- Leave-one-out

- N training sessions on N – 1 points each time



- Leave-more-out

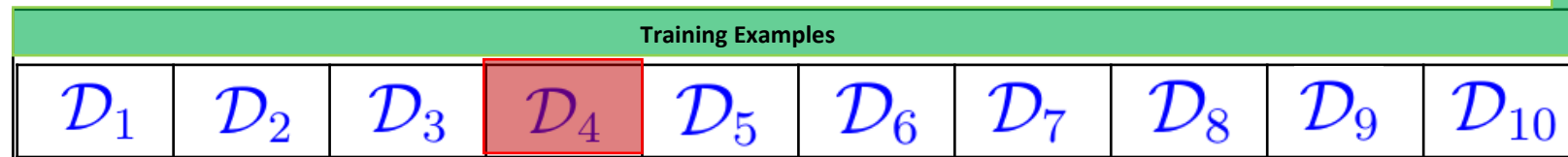
- Break data into number of folds
  - N/K training sessions on N – K points each time
  - Example: ‘10-fold cross-validation’ with K = N/10 multiple times (N/K)



(fewer training sessions than above)

(use 1/10 for validation, use 9/10 for training, then another 1/10 ... N/K times)

$D$   
(dataset)

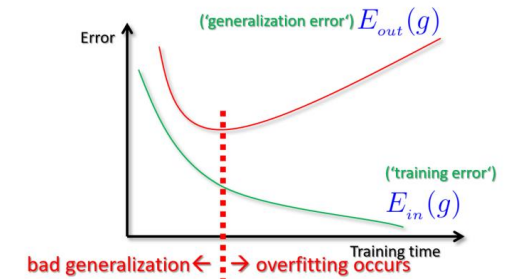


(involved in training now)

(involved in training now)

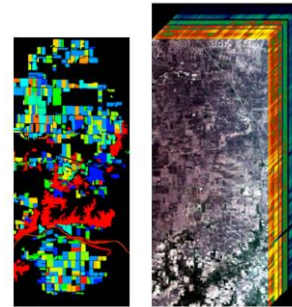
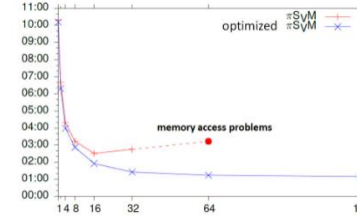
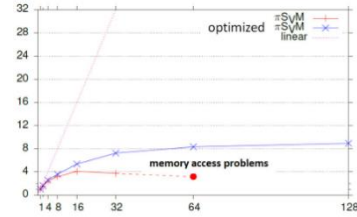
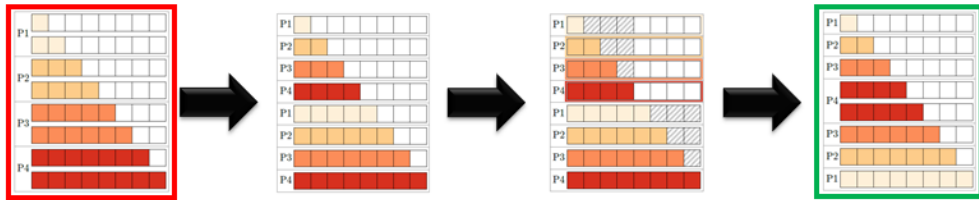
[38] [www.big-data.tips](http://www.big-data.tips), ‘Cross Validation’

- 10-fold cross validation is mostly applied in practical problems by setting K = N/10 for real data
- Having N/K training sessions on N – K points each leads to long runtimes (→ use parallelization)



# Parallel Support Vector Machine (SVM) – piSVM MPI Implementation & Impact

- Original piSVM 1.2 version (2011)
  - Open-source and based on libSVM library, C
  - Message Passing Interface (MPI)
  - New version appeared 2014-10 v. 1.3 (no major improvements)
  - Lack of 'big data' support (e.g. memory)
- Tuned scalable parallel piSVM tool 1.2.1
  - Highly scalable version maintained by Juelich
  - Based on original piSVM 1.2 tool
  - Optimizations: load balancing; MPI collectives



```
//Jeder Prozess hat l Werte an die Stelle Rang * l in p_cache_status geschrieben
for(int k = 0; k < p; ++k)
{
    //Jeder Prozess broadcastet sein Ergebnis zu allen anderen Prozessen
    MPI_Bcast(&p_cache_status[k * l], l, MPI_CHAR, k, comm);
}
// Using MPI_Allgather() instead

for(int i = 0; i < p; ++i) {
    if(rank == i) { //Der sendende Prozess kopiert in den Sendebereich
        for(int j = 0; j < lmn; ++j)
            G_buf[j] = G_n[j];
    }
    //Alle anderen Prozesse erhalten die Daten
    MPI_Bcast(G_buf, lmn, MPI_DOUBLE, i, comm);
    //Und addieren sie auf
    for(int j = 0; j < lmn; ++j)
        G[not_work_set[j]] += G_buf[j];
}
// Using MPI_Allreduce() instead
```

Scenario 'pre-processed data', 10xCV serial: accuracy (min)

$\gamma/C$	1	10	100	1000	10 000
2	48.90 (18.81)	65.01 (19.57)	73.21 (20.11)	75.55 (22.53)	74.42 (21.21)
4	57.53 (16.82)	70.74 (13.94)	75.94 (13.53)	76.04 (14.04)	74.06 (15.55)
8	64.18 (18.30)	74.45 (15.04)	<b>77.00</b> (14.41)	75.78 (14.65)	74.58 (14.92)
16	68.37 (23.21)	76.20 (21.88)	76.51 (20.69)	75.32 (19.60)	74.72 (19.66)
32	70.17 (34.45)	75.48 (34.76)	74.88 (34.05)	74.08 (34.03)	73.84 (38.78)

Scenario 'pre-processed data', 10xCV parallel: accuracy (min)

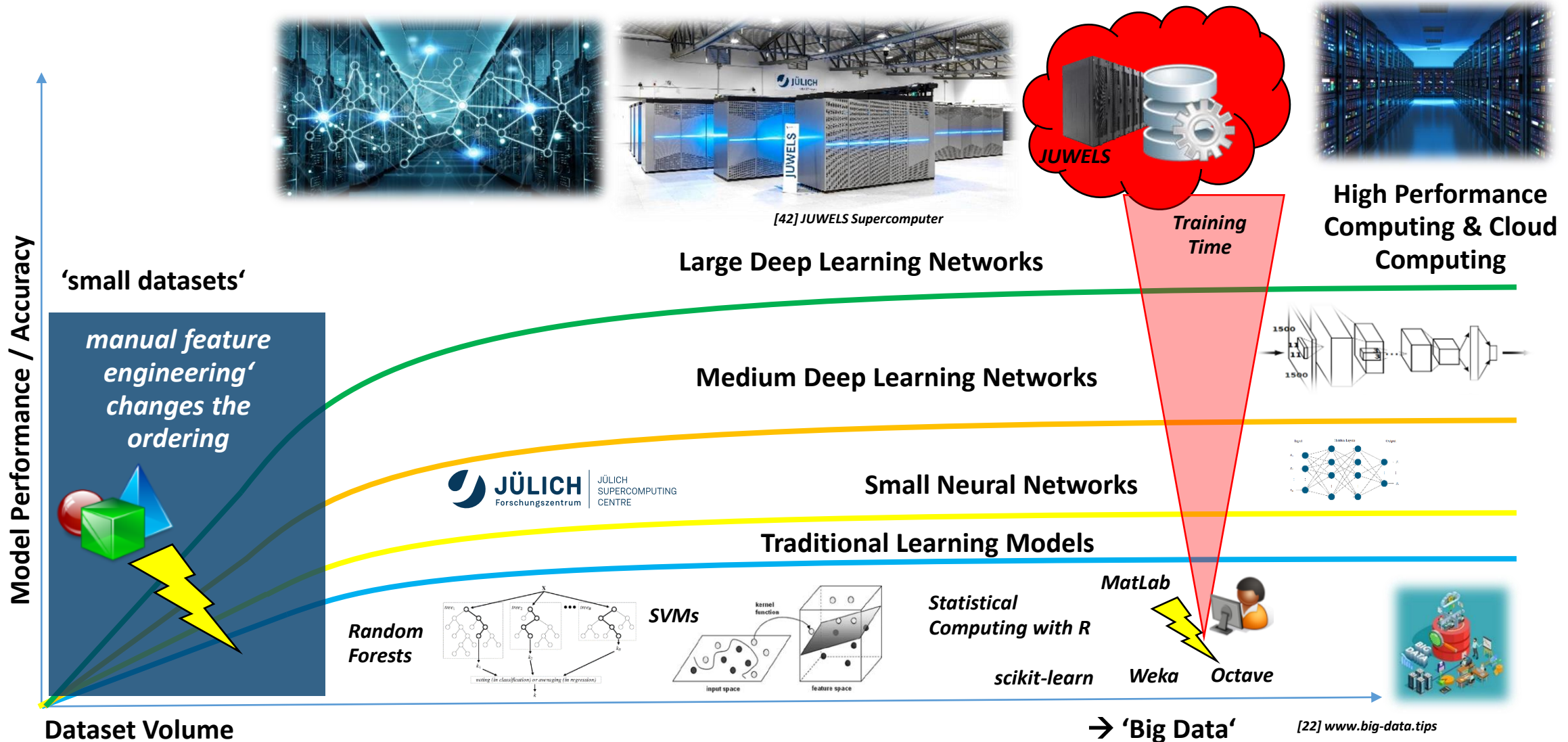
$\gamma/C$	1	10	100	1000	10 000
2	75.26 (1.02)	65.12 (1.03)	73.18 (1.33)	75.76 (2.35)	74.53 (4.40)
4	57.60 (1.03)	70.88 (1.02)	75.87 (1.03)	76.01 (1.33)	74.06 (2.35)
8	64.17 (1.02)	74.52 (1.03)	<b>77.02</b> (1.02)	75.79 (1.04)	74.42 (1.34)
16	68.57 (1.33)	76.07 (1.33)	76.40 (1.34)	75.26 (1.05)	74.53 (1.34)
32	70.21 (1.33)	75.38 (1.34)	74.69 (1.34)	73.91 (1.47)	73.73 (1.33)

First Result: best parameter set from 14.41 min to 1.02 min  
 Second Result: all parameter sets from ~9 hours to ~35 min

[39] piSVM on SourceForge, 2008

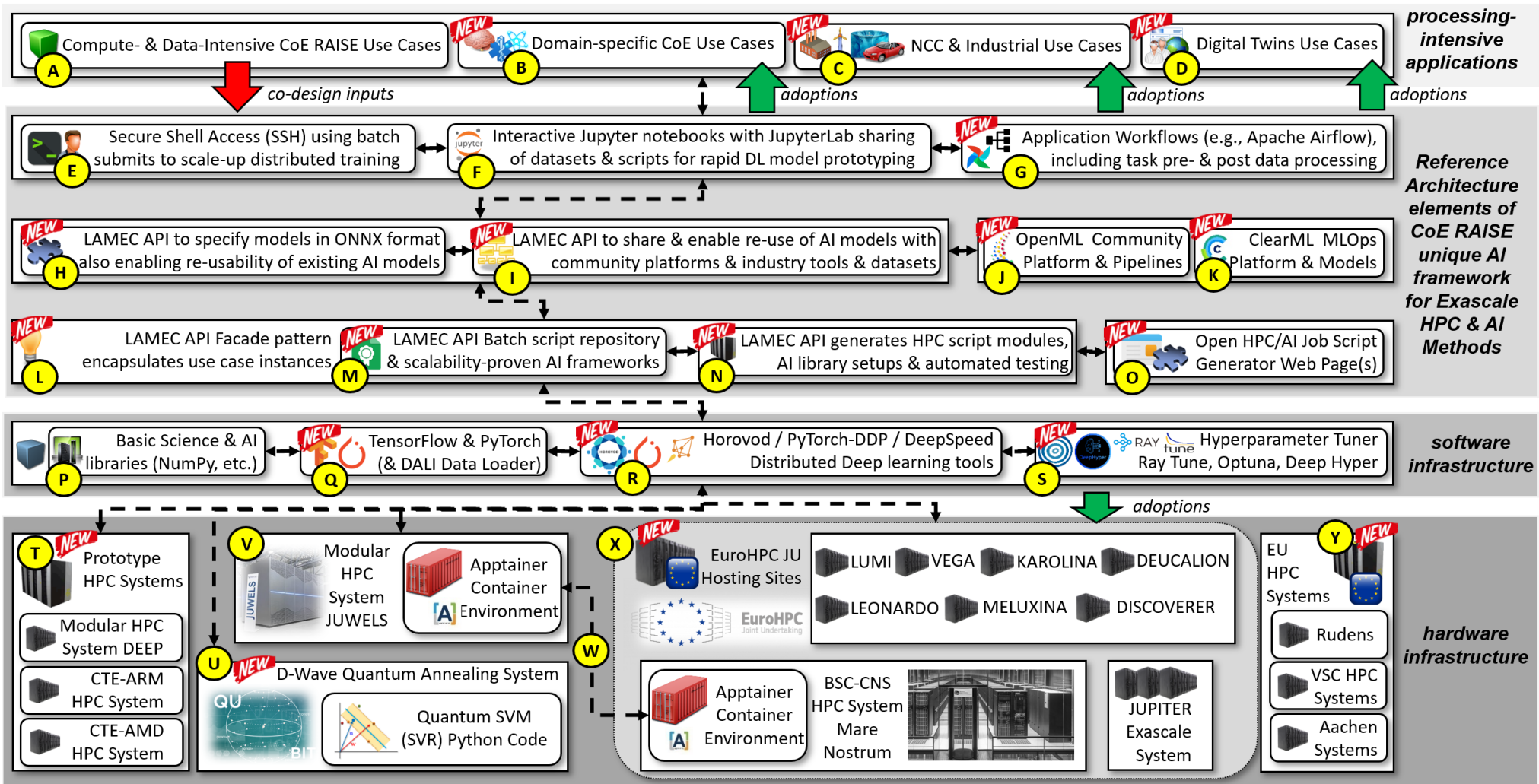
[34] G. Cavallaro & M. Riedel & J.A. Benediktsson et al., 'On Understanding Big Data Impacts in Remotely Sensed Image Classification Using Support Vector Machine Methods', Journal of Applied Earth Observations and Remote Sensing

# HPC Relationship to 'Big Data' in Machine & Deep Learning – Scalability





# HPC Relationship to 'Big Data' in Machine & Deep Learning – Complexity



[58] CoE RAISE Web Page



▪ Load Modules, Environments, and Containers (LAMEC)

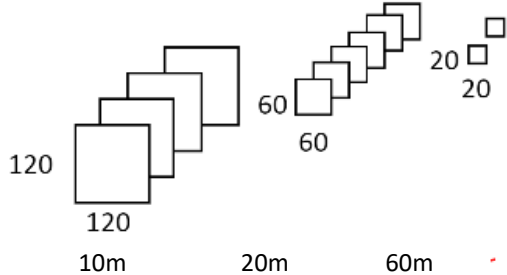
[57] M. Riedel, C. Barakat, S. Fritsch, M. Aach, J. Busch, A. Lintermann, A. Schuppert, S. Brynjólfsson, H. Neukirchen, M. Book: Enabling Hyperparameter-Tuning of AI Models for Healthcare Using the CoE RAISE Unique AI Framework for HPC, in proceedings of IEEE MIPRO 2023, Croatia

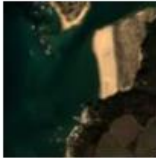


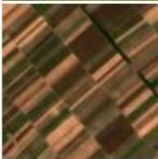


# Multispectral Remote Sensing Dataset Example used in Deep Learning

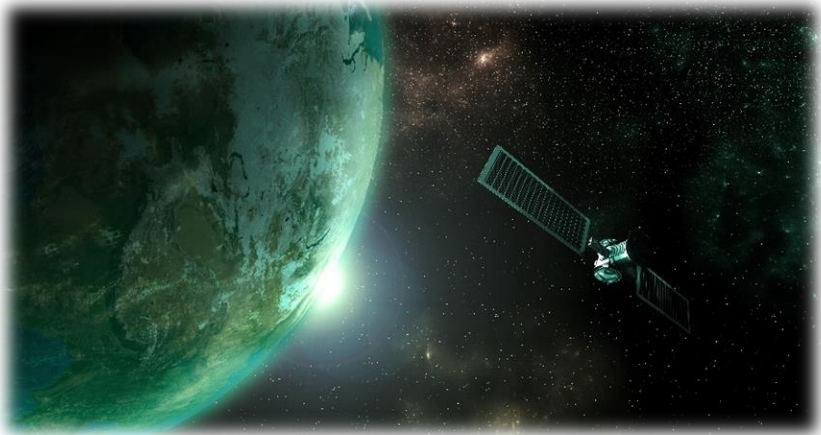
Datasets	Image type	Image per class	Scene classes	Annotation type	Total images	Spatial resolution (m)	Image sizes	Year
BigEarthNet	Satellite MS	328 to 217119	43	Multi label	590,326	10	120x120	2018
						20	60x60	
						60	20x20	

[40] G. Sumbul et al.

[41] Big Earth Net Dataset



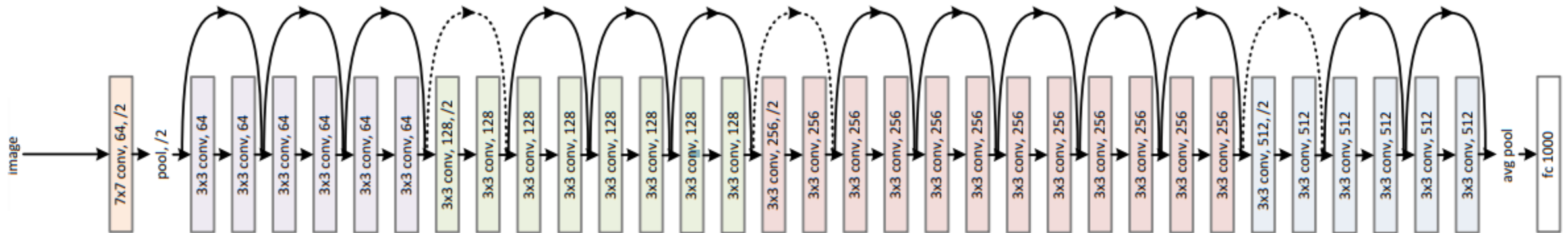
	permanently irrigated land, sclerophyllous vegetation, beaches, dunes, sands, estuaries, sea and ocean		non-irrigated arable land, fruit trees and berry plantations, agro-forestry areas, transitional woodland/shrub
	permanently irrigated land, vineyards, beaches, dunes, sands, water courses		non-irrigated arable land
	coniferous forest, mixed forest, water bodies		discontinuous urban fabric, non-irrigated arable land, land principally occupied by agriculture, broad-leaved forest





# Deep Learning via RESNET-50 Architecture – A Case for interconnecting GPUs

- Classification of land cover in scenes in Remote Sensing
  - Very suitable for parallelization via distributed training on multi GPUs

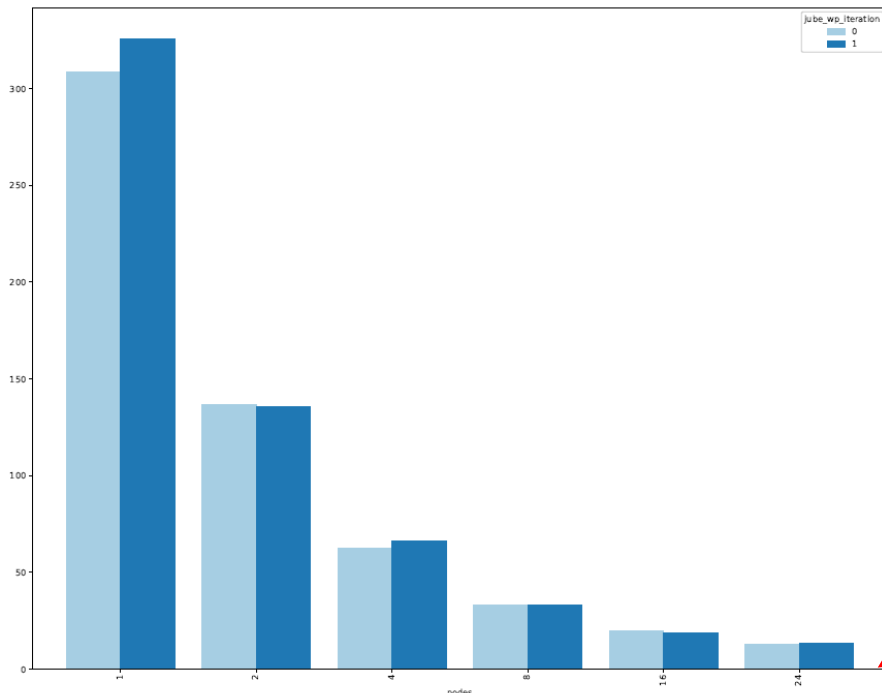


[45] RESNET

- RESNET-50 is a known neural network architecture that has established a strong baseline in terms of accuracy
- The computational complexity of training the RESNET-50 architecture relies in the fact that it has ~ 25.6 millions of trainable parameters
- RESNET-50 still represents a good trade-off between accuracy, depth and number of parameters
- The setup of RESNET-50 makes it very suitable for parallelization via distributed training on multi GPUs

# HPC Enables Faster Training of AI Models: Distributed Training (Benefit #1)

Time per epoch [sec]



A partition of the JUWELS system has 56 compute nodes, each with 4 NVIDIA V100 GPUs (equipped with 16 GB of memory)

24 nodes x 4 GPUs = 96 GPUs

- Horovod is a distributed training framework used in combination with low-level deep learning frameworks like Tensorflow
- Horovod uses MPI for inter-process communication, e.g., `MPI_Allreduce()`
- Distributed training using data parallelism approach means: (1) Gradients for different batches of data are calculated separately on each node; (2) But averaged across nodes to apply consistent updated to the deep learning model in each node

Other distributed training approaches possible with DeepSpeed

[46] DeepSpeed

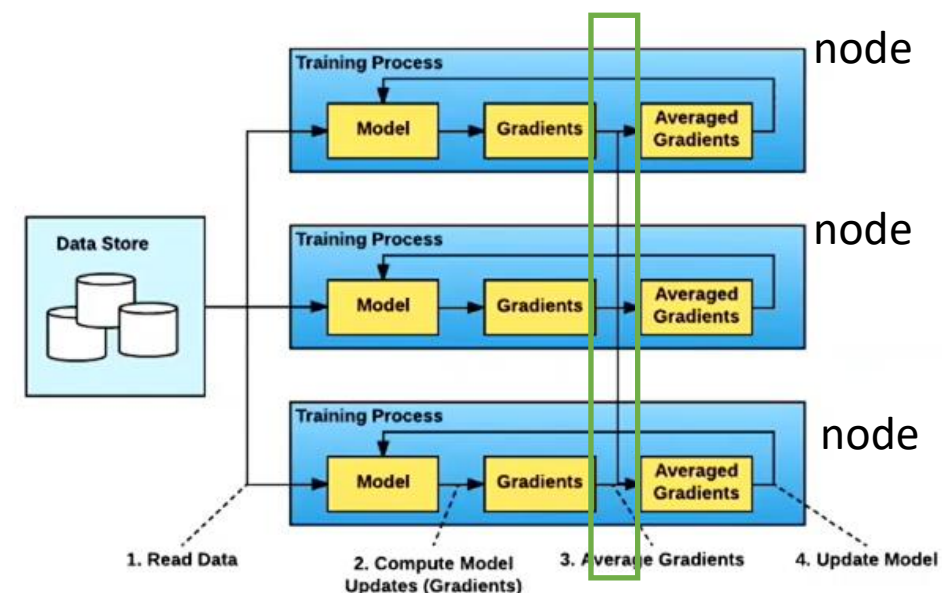
DeepSpeed is a deep learning optimization library that makes distributed training easy, efficient, and effective.

10x Larger Models

10x Faster Training

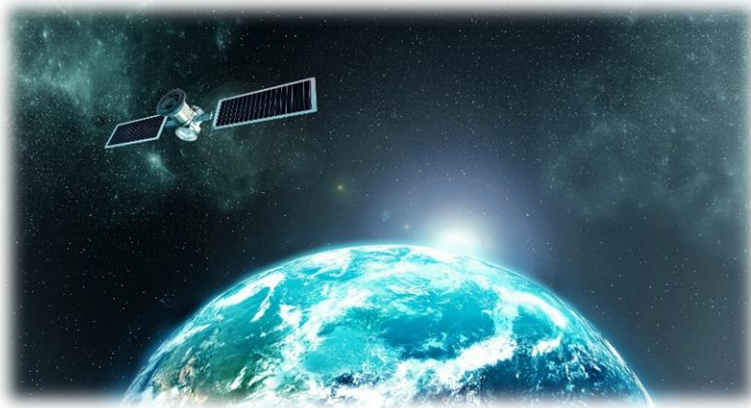
Minimal Code Change

Horovod distributed training via `MPI_Allreduce()`



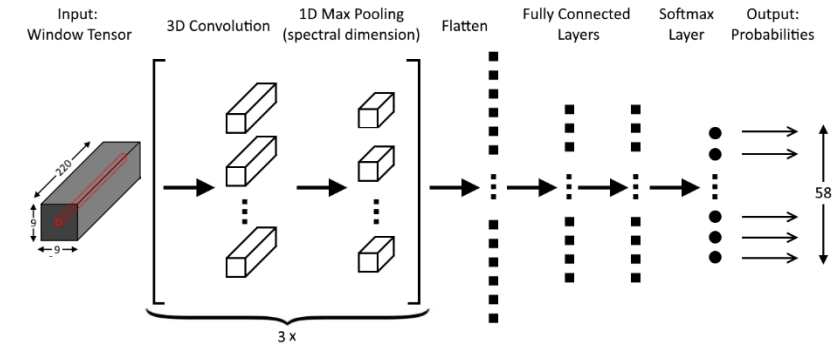
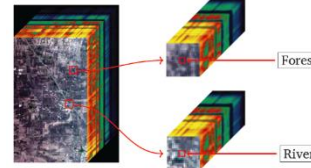
[47] Horovod

# Deep Learning Application Example – Using High Performance Computing



- Using Convolutional Neural Networks (CNNs) with hyperspectral remote sensing image data

[43] J. Lange and M. Riedel et al., IGARSS Conference, 2018

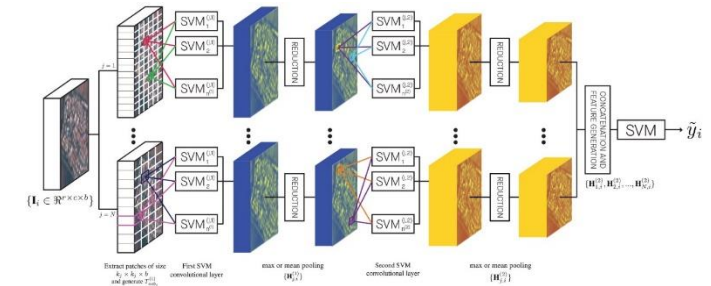


- Find Hyperparameters & joint 'new-old' modeling & transfer learning given rare labeled/annotated data in science (e.g. 36,000 vs. 14,197,122 images ImageNet)

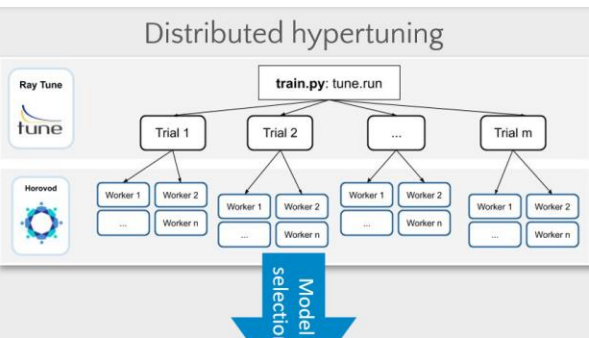
Feature	Representation / Value
Conv. Layer Filters	48, 32, 32
Conv. Layer Filter size	(3, 3, 5), (3, 3, 5), (3, 3, 5)
Dense Layer Neurons	128, 128
Optimizer	SGD
Loss Function	mean squared error
Activation Functions	ReLU
Training Epochs	600
Batch Size	50
Learning Rate	1
Learning Rate Decay	$5 \times 10^{-6}$

[44] G. Cavallaro, M. Riedel et al., IGARSS 2019

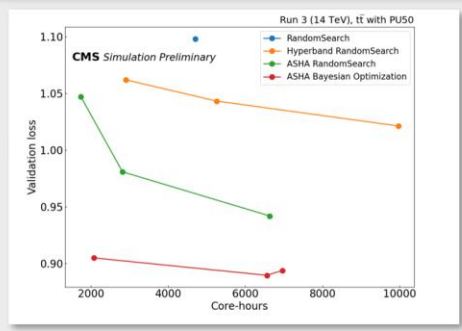
(challenges of big data & hyperparameter tuning)



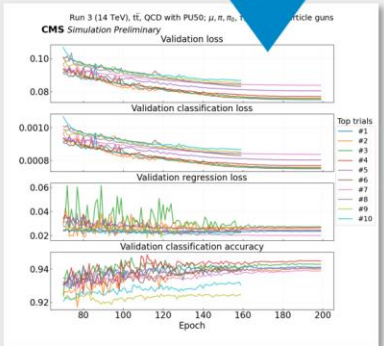
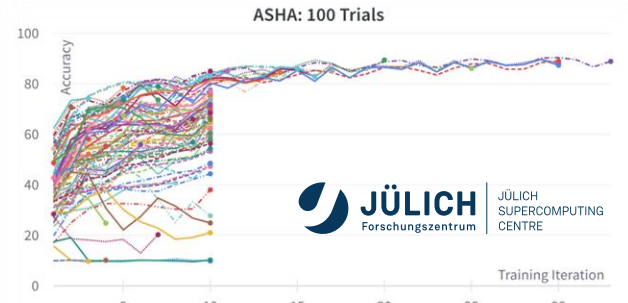
# HPC Enables Better AI Models: Hyperparameter Tuning (Benefit #2)



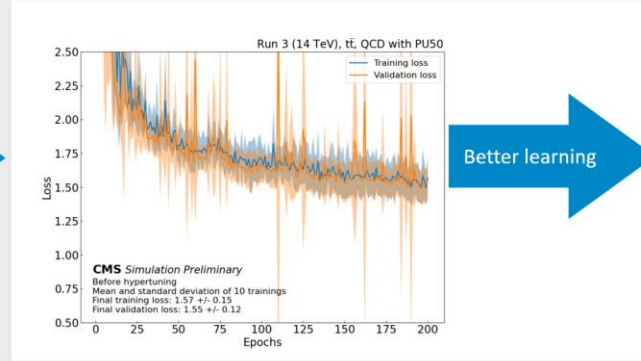
Hypertuning evaluation



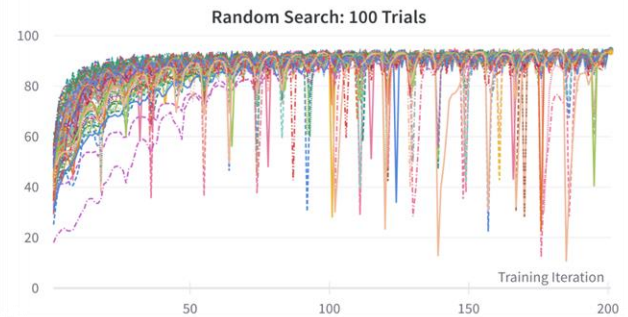
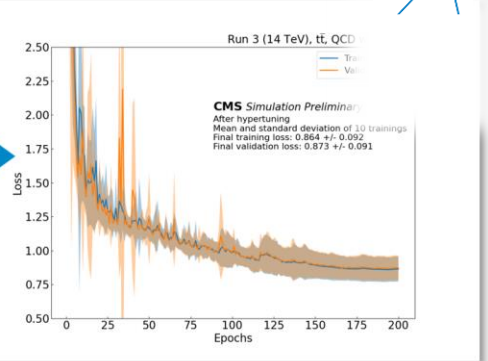
Mean validation loss decreased by ~44% giving a significant performance improvement



Assess learning variability



Better learning



MDPI

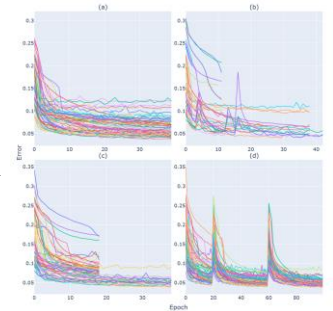
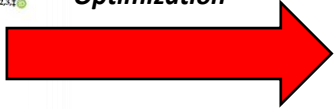
Article

### Developing an Artificial Intelligence-Based Representation of a Virtual Patient Model for Real-Time Diagnosis of Acute Respiratory Distress Syndrome

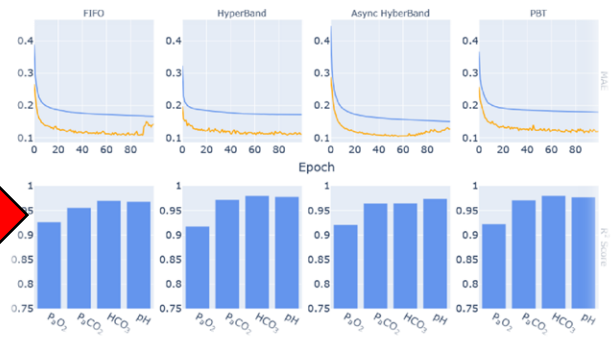
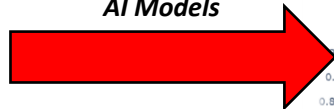
Chadi Barakat<sup>1,2,3,4,\*</sup>, Konstantin Sharafutdinov<sup>3,4</sup>, Josefine Busch<sup>1</sup>, Sina Saffaran<sup>5</sup>, Declan G. Bates<sup>6</sup>, Jonathan G. Hardman<sup>6</sup>, Andreas Schuppert<sup>3,4</sup>, Sigurður Brynjólfsson<sup>2</sup>, Sebastian Fritsch<sup>1,3,7,8</sup>, and Morris Riedel<sup>1,2,3,4</sup>

<sup>1</sup> Jülich Supercomputing Centre, Forschungszentrum Jülich, 52428 Jülich, Germany  
<sup>2</sup> School of Engineering and Natural Science, University of Iceland, 107 Reykjavík, Iceland  
<sup>3</sup> SMITH Consortium of the German Medical Informatics Initiative, 07747 Leipzig, Germany  
<sup>4</sup> Joint Research Centre for Computational Biomedicine, University Hospital RWTH Aachen, 52074 Aachen, Germany  
<sup>5</sup> School of Engineering, University of Warwick, CV4 7AL Coventry, UK  
<sup>6</sup> School of Medicine, University of Nottingham, NG7 2RD Nottingham, UK  
<sup>7</sup> Department of Intensive Care Medicine, University Hospital RWTH Aachen, 52074 Aachen, Germany  
<sup>8</sup> Correspondence: c.barakat@fz-juelich.de  
 \* Current address: Jülich Supercomputing Centre, Forschungszentrum Jülich, 52428 Jülich, Germany.  
 † These authors contributed equally to this work.

Hyperparameter Optimization



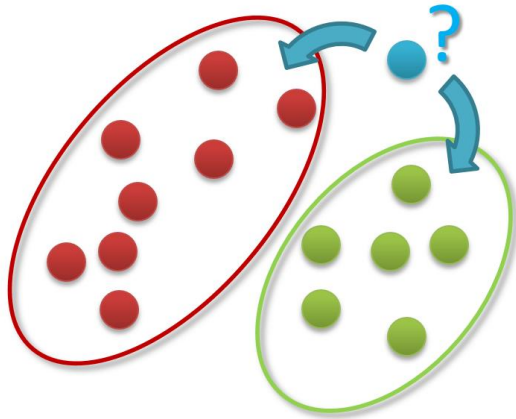
Train Better AI Models





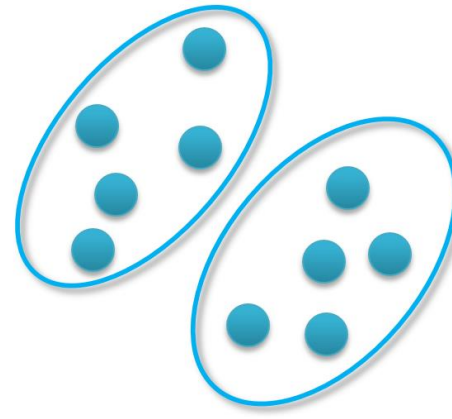
# Machine Learning Models – Short Overview & Introduction to Classification

## Classification



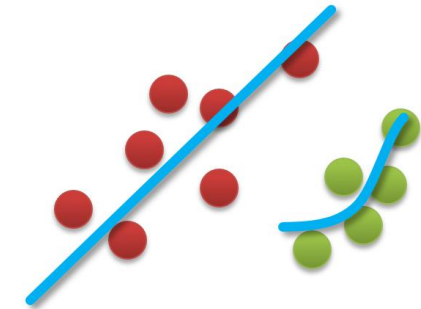
- Groups of data exist
- New data classified to existing groups

## Clustering



- No groups of data exist
- Create groups from data close to each other

## Regression



- Identify a line with a certain slope describing the data

▪ Machine learning methods can be roughly categorized in classification, clustering, or regression augmented with various techniques for data exploration, selection, or reduction – despite the momentum of deep learning, traditional machine learning algorithms are still widely relevant today



# Data Science Example: DBSCAN Clustering Algorithm

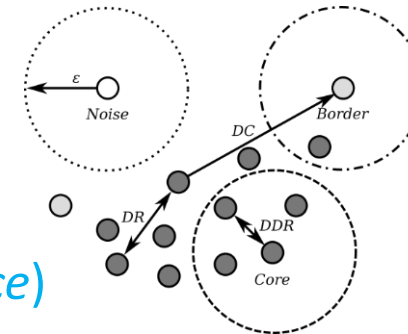
## ■ DBSCAN Algorithm

- Introduced 1996 and most cited clustering algorithm
- Groups number of similar points into clusters of data
- Similarity is defined by a distance measure (e.g. *euclidean distance*)

## ■ Distinct Algorithm Features

- Clusters a variable number of clusters (cf. K-Means Clustering with K clusters)
- Forms arbitrarily shaped clusters (no 'bow ties')
- Identifies inherently also outliers/noise

- Density-based spatial clustering of applications with noise (DBSCAN) is a data clustering algorithm that requires only two parameters and has no requirement to specify number of clusters
- Parameter Epsilon: Algorithm looks for a similar point within a given search radius Epsilon
- Parameter minPoints: Algorithm checks that cluster consist of a given minimum number of points



(MinPoints = 4)

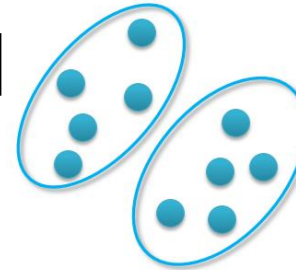
(DR = Density Reachable)

(DDR = Directly Density Reachable)

(DC = Density Connected)

[49] Ester et al.

Clustering



```
#!/bin/bash
#SBATCH --job-name=HPDBSCAN
#SBATCH -o HPDBSCAN-%j.out
#SBATCH -e HPDBSCAN-%j.err
#SBATCH --nodes=2
#SBATCH --ntasks=4
#SBATCH --ntasks-per-node=4
#SBATCH --time=00:20:00
#SBATCH --cpus-per-task=4
#SBATCH --reservation=ml-hpc-1

export OMP_NUM_THREADS=4

# location executable
HPDBSCAN=/homea/hpclub/train001/tools/hpdbscan/dbscan

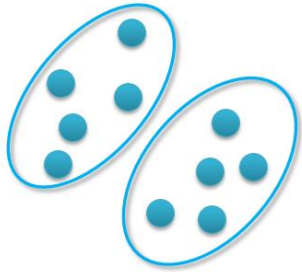
# your own copy of bremen small
BREMENSMALLDATA=/homea/hpclub/train001/bremenSmall.h5

# your own copy of bremen big
BREMENBIGDATA=/homea/hpclub/train001/bremen.h5

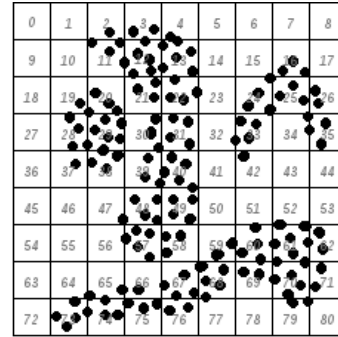
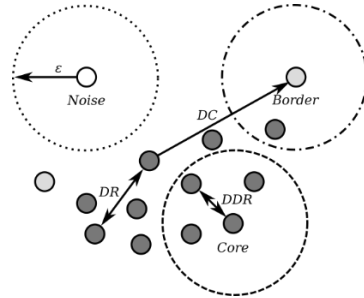
srun $HPDBSCAN -m 100 -e 300 -t 12 $BREMENSMALLDATA
```

[48] M. Goetz and M. Riedel et al,  
Proceedings IEEE Supercomputing Conference, 2015

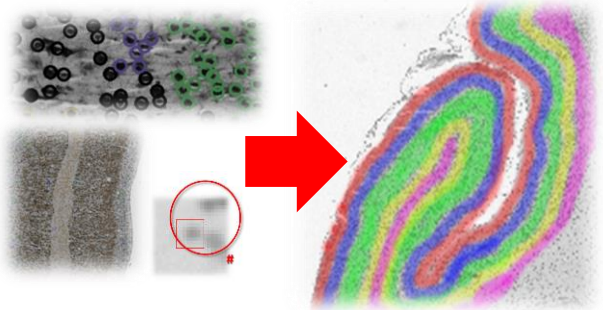
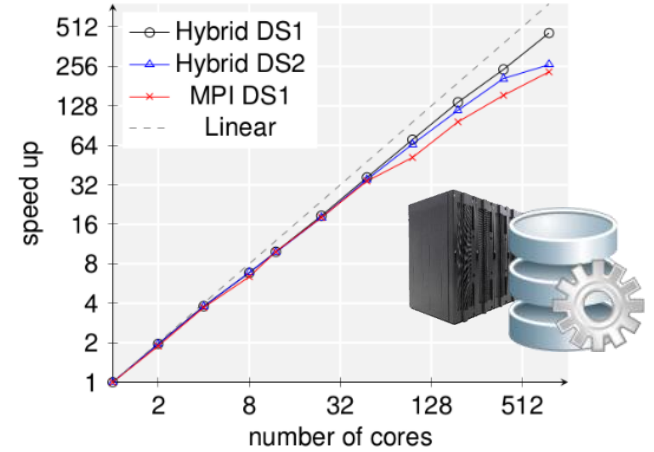
# Data Parallelism Example: Smart Domain Decomposition in Data Sciences



**Clustering**

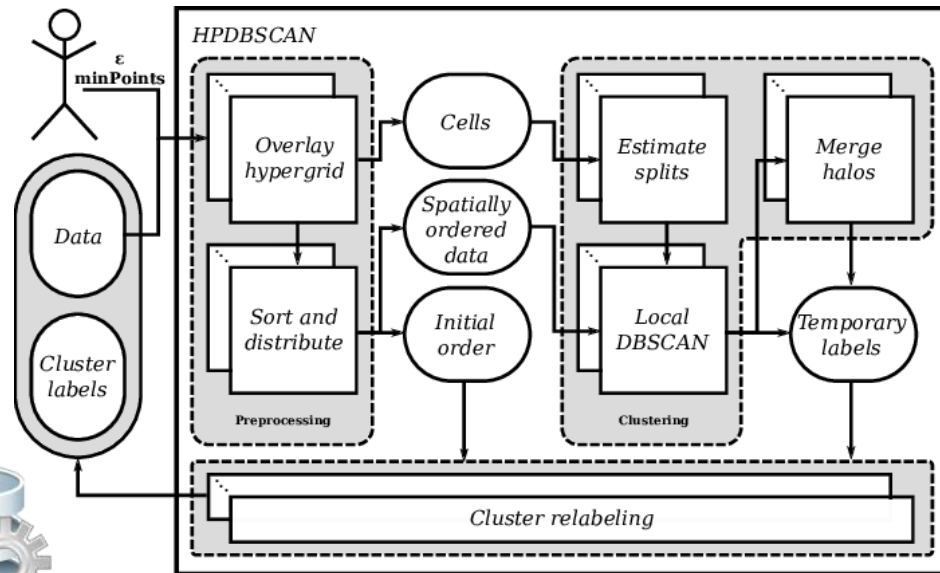


Overlaid spatial grid

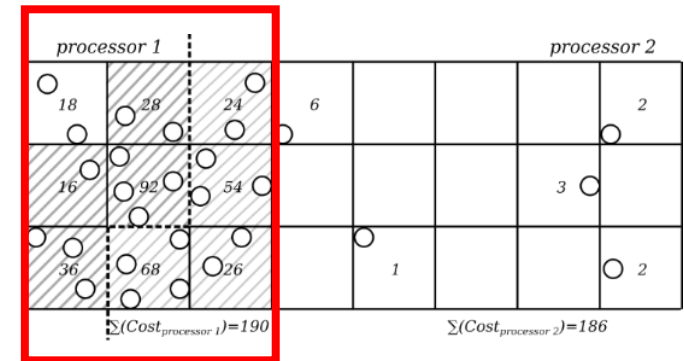


[48] M. Goetz and M. Riedel et al, Proceedings IEEE Supercomputing Conference, 2015

Algorithm can be used in many application domains



cluster merge across halo regions/layers



# HPDBSCAN Clustering – Using Parallel File Formats & File Systems

```
[morris@jotunn hpdbscan]$ more HPDBSCAN-199947.out
Calculating Cell Space...
  Computing Dimensions... [OK] in 0.054227
  Computing Cells... [OK] in 0.022971
  Sorting Points... [OK] in 0.133213
  Distributing Points... [OK] in 0.111046
DBSCAN...
  Local Scan... [OK] in 139.057375
  Merging Neighbors... [OK] in 0.010089
  Adjust Labels ... [OK] in 0.010476
  Rec. Init. Order ... [OK] in 0.556270
  Writing File ... [OK] in 0.170748
Result...
  21 Clusters
  2974394 Cluster Points
  25606 Noise Points
  2949094 Core Points
Took: 140.453795s
```

```
[morris@jotunn hpdbscan]$ h5dump -d /Clusters bremenSmall.h5
HDF5 "bremenSmall.h5" {
DATASET "/Clusters" {
  DATATYPE H5T_STD_I64LE
  DATASPACE SIMPLE { ( 3000000 ) / ( 3000000 ) }
  DATA {
(0): 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
(23): 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
(46): 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
(69): 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
(92): 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
(115): 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0, 0,
```

- The input data for the parallel & scalable HPDBSCAN clustering algorithm is a HDF5 file and all the processors read in parallel chunks of the data
- The HDF5 file before the execution of HPDBSCAN has 0 as Cluster Ids for its specific initialization



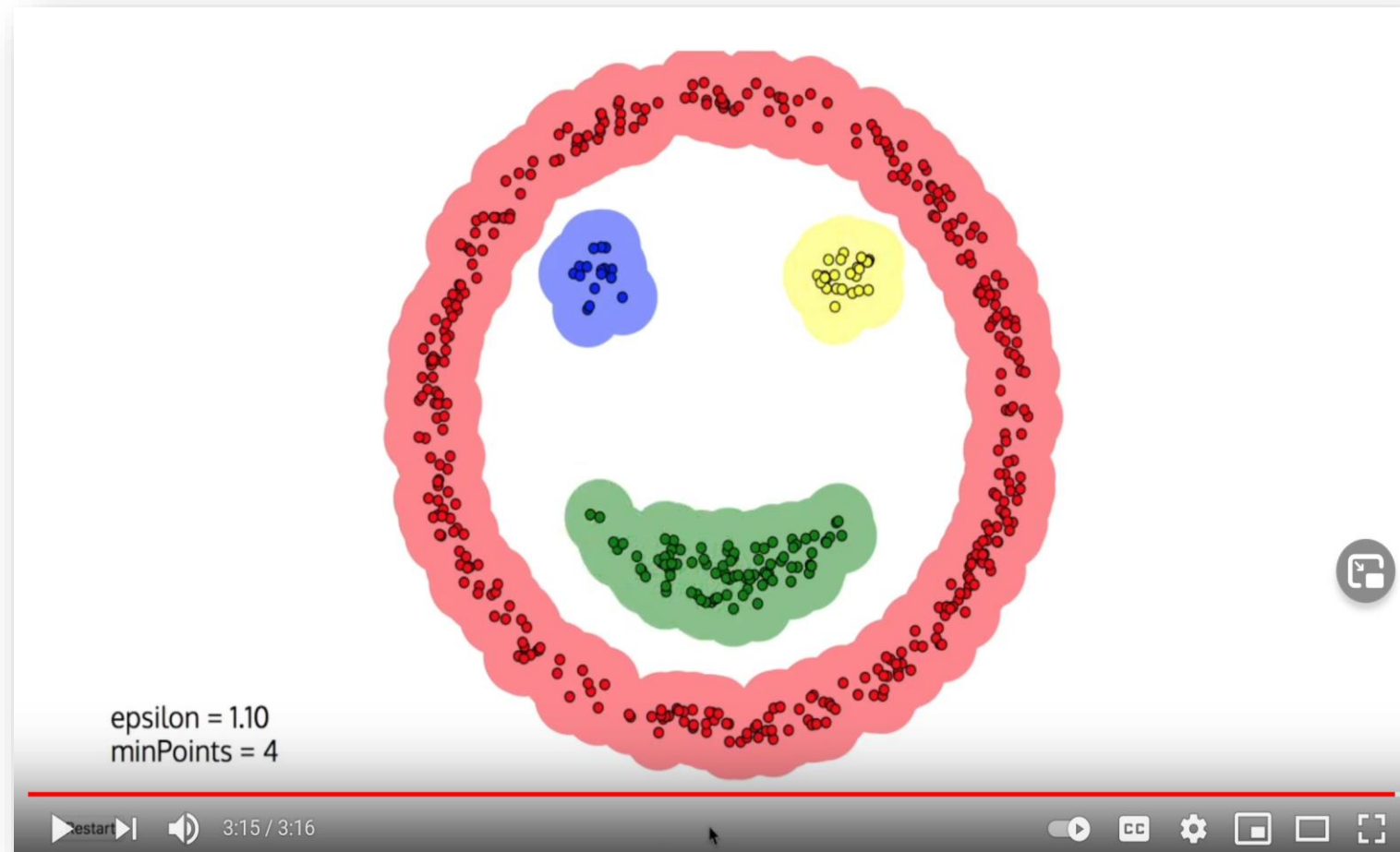
- The standard out of the HPDBSCAN parallel & scalable DBSCAN clustering algorithm is not the result of the DBSCAN clustering algorithm and only shows meta information such as the numbers of clusters found, noise, and running time

- The real outcome of the parallel & scalable HPDBSCAN algorithm is directly written into the HDF5 file assigning for each point cloud data element a specific cluster ID, or using minus numbers to indicate noise points (no real clusters)

```
[morris@jotunn hpdbscan]$ h5dump -d /Clusters bremenSmall.h5
HDF5 "bremenSmall.h5" {
DATASET "/Clusters" {
  DATATYPE H5T_STD_I64LE
  DATASPACE SIMPLE { ( 3000000 ) / ( 3000000 ) }
  DATA {
(0): -45205, -45205, -45205, -45205, -45205, -45205, -45205, -45205,
(8): -45205, -45205, -45205, -45205, -45205, -45205, -45205, -45205,
(17): -45205, -4108, -45205, -45205, -45205, -45205, -45205, -45205,
(25): -45205, -45205, -45205, -45205, -45205, -45205, -45205, -45205,
(33): -45205, -45205, -45205, -45205, -45205, -45205, -45205, -45205,
(41): -45205, -45205, -45205, -45205, -45205, -45205, -45205, -45205,
(49): -45205, -45205, -45205, -45205, -45205, -45205, -45205, -45205,
(57): -45205, -45205, -45205, -45205, -45205, -45205, -45205, -45205,
(65): -45205, -45205, -45205, -45205, -45205, -45205, -45205, -45205,
(73): -45205, -45205, -45205, -45205, -45205, -45205, -45205, -45205,
(81): -45205, -45205, -45205, -490063, -45205, -45205, -45205, -45205,
(89): -45205, -45205, -45205, -45205, -45205, -45205, -45205, 45205,
(97): -45205, -45205, -45205, -45205, -45205, -45205, -45205, -45205,
(105): -45205, -45205, -45205, -45205, -45205, -45205, -45205, -45205,
(113): -45205, -45205, -45205, -45205, -45205, -45205, -490063, -45205,
```

[48] M. Goetz and M. Riedel et al, Proceedings IEEE Supercomputing Conference, 2015

# [Video for further Studies] DBSCAN Algorithm Steps & Visualization Example



[50] YouTube video, DBSCAN Explanation and Visualization

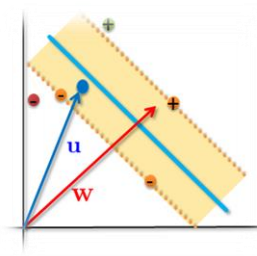
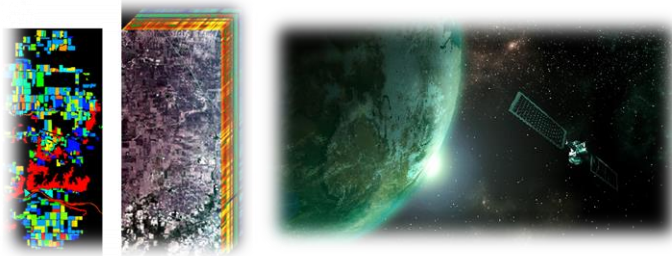
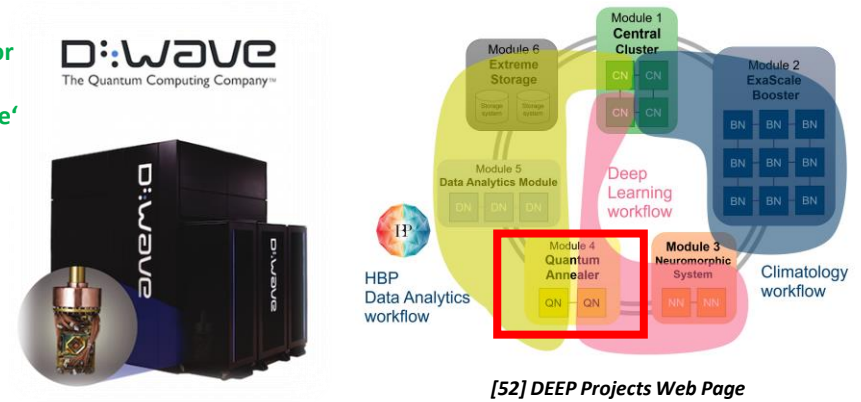


# Emerging Quantum Machine Learning – Initial Results on Quantum Annealing

- **Disruptive Technology** in the HPC Ecosystems
  - Different concept than traditional (super)computers
  - Information as '0' or '1' or both simultaneously

computing time for optimization  
'in a blink of an eye'

[51] D-Wave Systems YouTube Channel



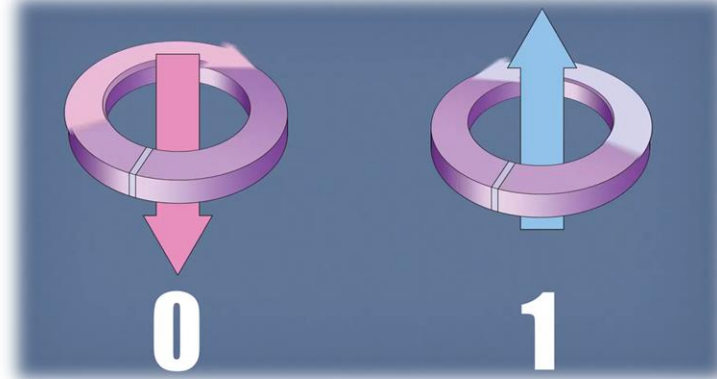
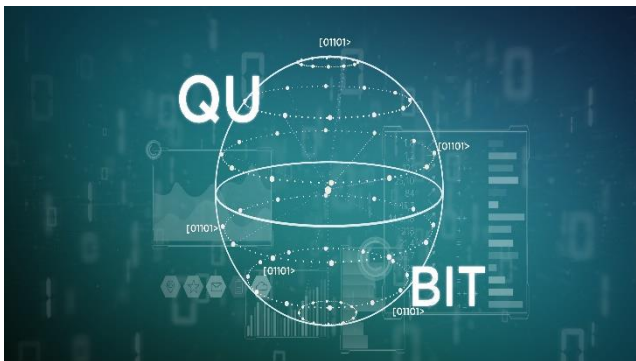
```
In [ ]: from quantum_SVM import *

# Hyperparameters
B=[2,3,5,10]
K=[2,3]
xi=[0,1,5]
gamma=[-1,0.125,0.25,0.5,1,2,4,8]
n_experiments=len(B)*len(K)*len(xi)*len(gamma)

hyperparameters=np.zeros(n_experiments,4), dtype=float)

path_data_key='input_datasets/calibration/'+id_dataset+'/'
data_key = id_dataset+'calibrain'
path_out='outputs/calibration/'+id_dataset+'/'

trainacc=np.zeros(10fold), dtype=float)
trainauc=np.zeros(10fold), dtype=float)
trainaup=np.zeros(10fold), dtype=float)
```



[53] M. Riedel, UTMessan 2020 YouTube Video



- [54] G. Cavallaro & M. Riedel et al., 'Approaching Remote Sensing Image Classification with ensembles of support vector machines on the D-Wave Quantum Annealer'
- [55] M. Riedel, G. Cavallaro, J.A. Bendiktsson, 'Practice and Experience in Using Parallel and Scalable Machine Learning in Remote Sensing from HPC Over Cloud to Quantum Computing'
- [56] Delilbasic, A., Cavallaro, G., Willsch, M., Melgani, F., Riedel, M., Michielsen, K., 'Quantum Support Vector Machine Algorithms for Remote Sensing Data Classification'



# More Information: Full HPC Spring 2023 University Course



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Research Group Leader, Juelich Supercomputing Centre, Forschungszentrum Juelich, Germany

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Prologue  
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LECTURE 0  
Prologue  
January 10, 2022  
Online Lecture

## Selected Testimonials from our community:



'... I have found your wonderful HPC lectures on YouTube, and I am finding them very helpful for learning the supercomputing tools necessary for my research...'

- student in computational nuclear physics from USA



neeraj kumar · 1 year ago

Awesome work for HPC students....thanks



纵横10分钟 - 10 Minutes Across the World · 3 months ago 5 subscribers

Thanks so much for the lecture - One of the best courses. Very helpful!



Andy Gill @\_andy\_gill · Nov 15, 2021

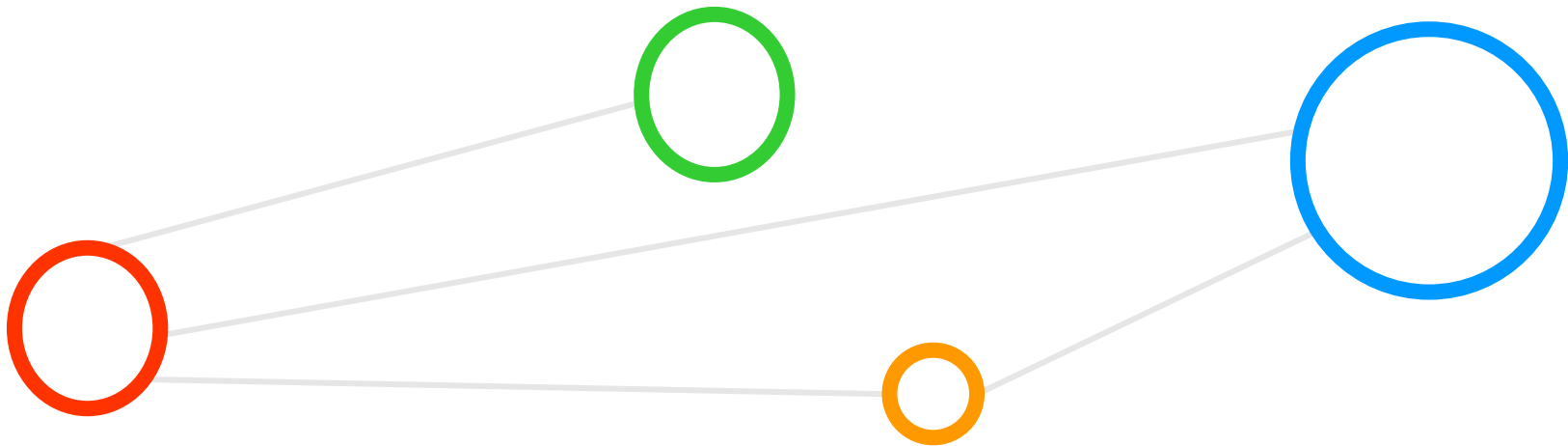
Highly recommend this excellent introduction to state of the art HPC programming by @MorrisRiedel. Enough details to actually be useful, explained clearly using many examples. Thank you for posting it!



youtube.com  
2021 High Performance Computing Course  
High Performance Computing Course - Advanced Scientific Computing 16 university lectures with ...

<https://www.youtube.com/channel/UCWC4VKHmL4NZgFfKoHtANKg>

# Lecture Bibliography



# Lecture Bibliography (1)

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<http://www.signa.org>
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<https://archive.ics.uci.edu/ml/datasets/Iris>
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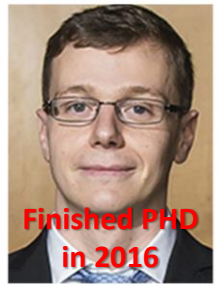
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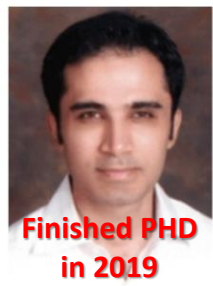
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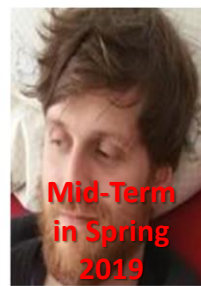
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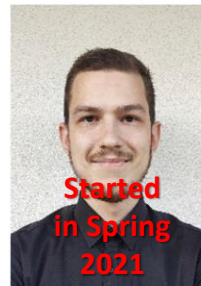
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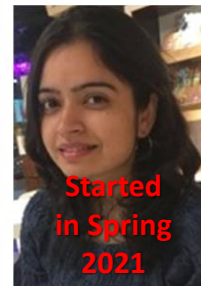
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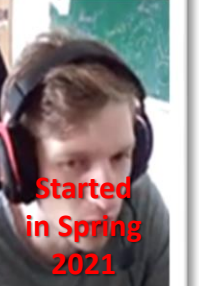
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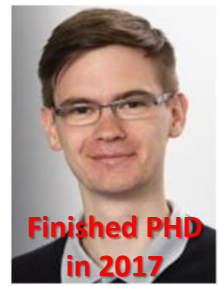
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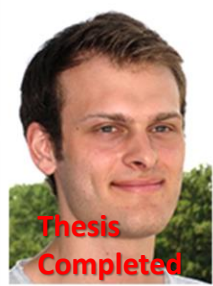
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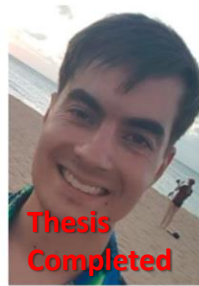
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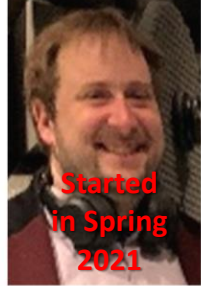
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