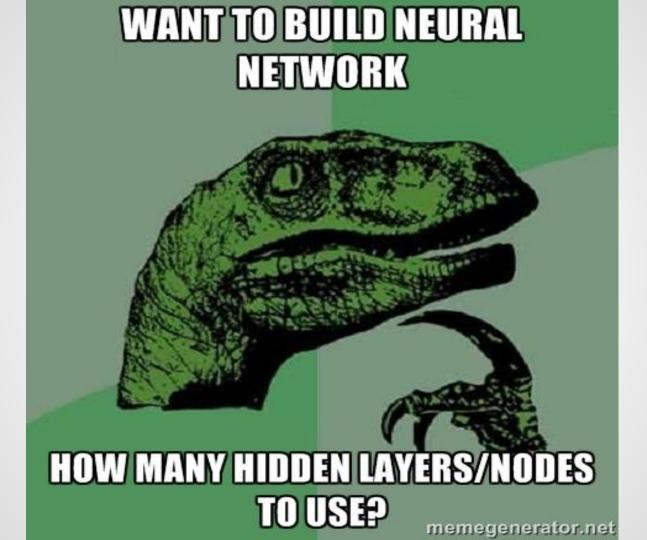
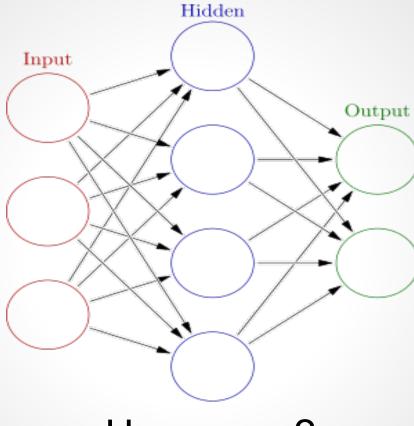
# How many hidden layers and nodes?

D.Stathakis, JRC European Commission 2009

Group 21
Aditya Kumar Akash 120050046
Deependra Patel 120050032
Nishant Kumar Singh 120050043



# **Training**



How many?

#### **Motivation**

- 2 decades, no exact solution until today
- Traditionally, it has been based on trial and error, heuristics, pruning and constructive methods
- None of them gives optimal or at least near-optimal solution
- This paper describes a genetic algorithm which aims to optimize performance while minimizing network complexity

# Theoretical bounds of optimal topology

In single hidden layer, number of neurons can be as high as number of training samples

The purpose of using a second hidden layer is to drastically reduce the total required number of hidden node

Huang (2003) proved that in the two-hidden-layer case, with m output neurons, the number of hidden nodes that are enough to learn N samples with negligibly small error is given by -

$$2\sqrt{(m+2)N}$$

Specifically, he suggests that the sufficient number of hidden nodes in the first layer is √(m+2)N + 2√N/(m+2) and in the second is m√N/(m+2)

# **CURRENT METHODS 1. Trial and error**

This is the most primitive path, and it will yield severely suboptimal structures.

## 2. Heuristic Search

#### Objective

To devise a formula that estimates the number of nodes in the hidden layers as a function of the number of input and output nodes

- Could be used to estimate a single exact topology or a range of topologies that should be searched.
- These heuristics are used prior to applying trial and error

eg: grapheme to phoneme asst., we take (n+m)/2

## 3. Exhaustive search

- Search through all possible topologies
- Very slow process
- Testing each topology takes lots of time
- Infeasible

# 4. Pruning and constructive algorithms

Aim at devising an efficient network structure by incrementally adding/removing links

Hessian matrix:-

$$h_{ij} = \frac{\partial^2 E}{\partial u_i \partial u_j}$$

## Optimal Brain Damage

- Progressively removes weights that causes least increase in training error
- To simplify computation, it assumes the hessian matrix of network is diagonal

## Optimal Brain Surgeon

- > It makes no such assumption and takes more time
- Does not demand retraining after pruning the weights
- Outperforms the rest

#### PROPOSED METHOD

- Uses genetic algorithm
- A novel fitness function is introduced to evaluate each solution
- This function aims at concurrently maximizing classification accuracy and at the same time minimizing network complexity

## Introduction to genetic algorithm

- Inspired from natural selection
- Involves inheritance, selection, crossover and mutation
- Algorithm is started with a set of solutions (represented by chromosomes) called population
- Solutions from one population are taken and are used to form a new population

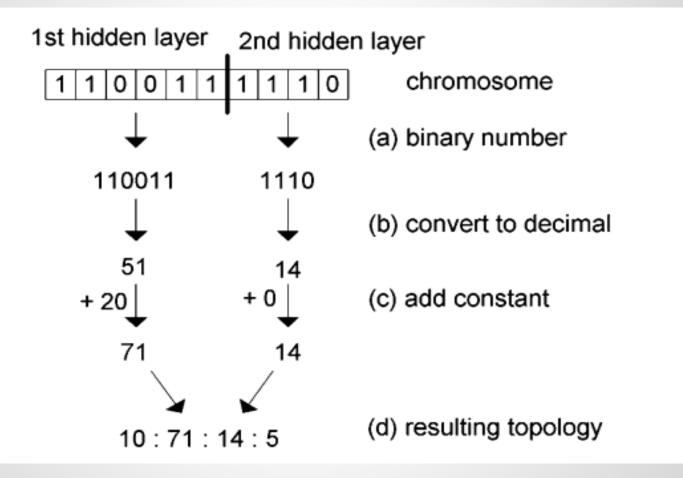
#### Continued...

- New population is better than the old one
- Solutions which are selected to form new solutions(offsprings) are selected according to their fitness
- The more suitable they are, more the chances they have to reproduce
- This process is repeated until either the maximum number of generations has been produced or a fitness level has been reached

## Symbiosis of genetic algorithm and neural networks

- Topologies with upto two hidden layers are searched
- The number of nodes of the two hidden layers is coded into binary chromosome
- The length I of the chromosome is 10 bits
- First six bits are reserved for first layer and next four for the second layer

#### Continued...



#### Continued...

- Convert binary numbers corresponding to each hidden layer into decimal numbers
- A constant is then added to each of those decimals
- For the second layer, constant is kept as 0 to account for the question - 'How many layers?'. Possibility of network with one hidden layer

#### Continued..

- The number of bits required is decided by the two equations shown earlier
- In our example with 10 inputs and 5 outputs, topologies between 10:[20-83]:[0-15]:5 are searched
- The network is built based on the number of nodes dictated by the genetic algorithm
- It is then trained and its performance is evaluated according to the fitness function

## Parameters of the Genetic Algorithm

Two point crossover is performed with a probability of 0.75. Example:

Parent 1: 1100|010|100

Parent 2: 0101|001|011

After crossover:

Offspring 1: 1100|001|100

Offspring 2: 0101|010|011

Tournament selection is performed with tournament size = 4

#### **Tournament selection**

- A few individuals chosen at random from the population
- The winner of each tournament is selected with probability p
- Second best selected with probability p\*(1-p), third p\*((1-p)^2) and so on
- Many such tournaments are performed

#### Parameters of GA continued...

- Uniform mutation is performed with mutation probability of 0.01
- Mutation important for genetic diversity
- Arbitrary bit in the genetic sequence changed from its original state
- Fitness scaling is performed to maintain even selection pressure throughout the GA

#### Parameters of GA continued...

The formulas are set according to Goldberg (let n denote the population size and I the length of chromosome)

- $\rightarrow$  n =  $\ell \log^2_{10}(l)$
- $\rightarrow$  max function evaluations =  $l \log_{10} l$
- $\rightarrow$  max generations =  $\ell^2 \log_{10}^2(l) / n$

## **Fitness Function**

 Novel Fitness function is used in the current genetic algorithm to rank the population

 Assumption Made : Compactness an additional benefit and hence the fitness function incorporates it

## Formula for Novel Fitness fn

$$f = e + s \frac{c - c_{\min}}{c_{\max} - c_{\min}}.$$

- e overall verification error for current topology
- c complexity factor, measured as number of weights
- c\_min, c\_max depend on the given chromosome length
- s accuracy sacrifice %

## **Novel Fitness fn conti.**

```
Example - 1100111110
```

- 1st layer, Max Nodes = (2^6 1) + 20 = 83, Min = 20
- 2nd layer, Max Nodes = (2^4 1) + 0 = 15, Min = 0
- Highest allowed structure = 10 : 83 : 15 : 5
- Lowest allowed structure = 10 : 20 : 5

#### Thus,

$$C_{max} = 10*83 + 83*15 + 15*5 = 2150$$
  
 $C_{min} = 10*20 + 20*5 = 300$ 

#### **Parameters of Fitness Function**

- C\_min, C\_max used for normalization to smoothly incorporate the fitness function, Ideally C\_min = 1, C\_max = inf.
- 's' Denotes the sacrifice in accuracy for compactness, s = 1 denotes 1% sacrifice, s = 0 denoted compactness is no more an objective
- With s = 0 we exert on pressure on algorithm and hence the more accurate individuals prevail
- For easy classification s is relaxed while for difficult ones s is kept low

## **Properties of Fitness fn**

- Only user defined parameter is 's', intuitive meaning, Hence applicable in other areas
- Logarithmic scaling of complexity ?? No need here since rank selection is adopted. Relieves from setting the curvature of logarithmic function used in case of proportional functions
- Wide scaling <u>Highest scaled take over the population gene</u> <u>pool too quickly</u> thus preventing algorithm from searching other areas
- On the other hand, if the <u>scaled values vary little</u>, all individuals have approximately the same chance of reproduction and the <u>search progresses very slowly</u>.

## **Experimental Results**

Experimental data refers to **Lefkas Island**.

#### Input Characteristics:

- 7 LANDSAT Bands
- Elevation
- Slope
- Aspect

#### **Output** Classes:

<u>5 landuse classes</u>: artificial surfaces, agricultural areas, forest and semi-natural areas, wetlands, and water bodies

# **Method Comparison**

#### Heuristics

Method name	Reference	Range	Topology	Mean Max	Min	$\sigma$
Kanellopoulos– Wilkinson rule also Hush rule	Kanellopoulos and Wilkinson (1997)	Low	10:20:5	70.54 71.87	68.48	0.69
	Hush (1989); Kanellopoulos and Wilkinson (1997)	Medium High		73.42 74.98 7 73.91 75.21 7		

Obs: Larger topology estimate better, stated another way heuristics tend to underestimate the complexity of the network,

None suggest use of second hidden layer

# **Pruning**

Method name	Reference	Range	Topology	Mean	Max	Min	$\sigma$
Optimal Brain Surgeon (OBD)	LeCun <i>et al.</i> (1990)			_	75.48	_	-
Optimal Brain Damage (OBS)	Hassibi and Stork (1993)			_	75.26	-	-

The maximum topology proposed among all heuristics, a 10:40:5 structure, is adopted as the starting point.

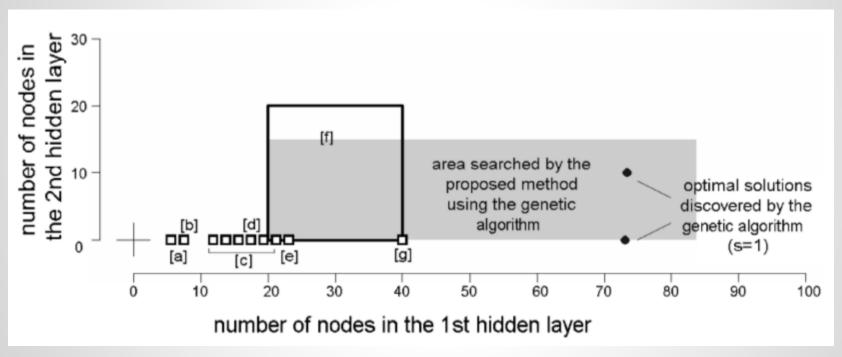
The computation resources required for both algorithms are high, with Optimal Brain Surgeon being more demanding.

# Genetic algorithm - accuracy over compactness

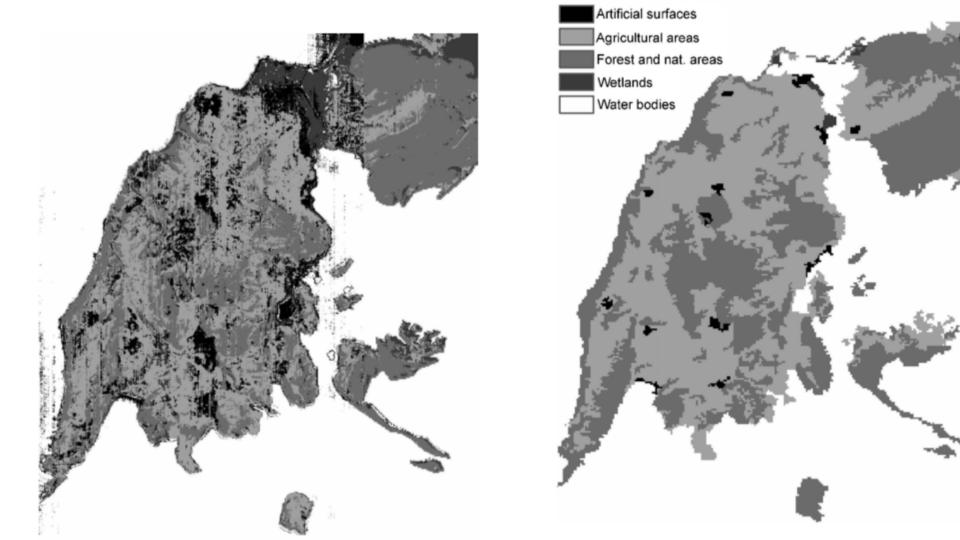
Method name	Reference	Range	Topology	Mean Max	Min	$\sigma$
Genetic algorithm $(s=1)$	Proposed method	10:[20-83]: [0-15]:5	10:73:10:5	70.53 78.34	19.96	15.29
			10:73:5	71.23 77.03	19.96	15.29
Genetic algorithm $(s=0)$		10:[20–83]: [0–15]:5	10:74:14:5	71.62 79.63	24.33	13.44

- For s = 1, Two complexity solutions evolve, One of them has only 1 hidden layer
- The accuracy difference for s = 0, s = 1 is not much since the classification problem is difficult, so here it makes less sense to use s > 1

# **Solution Space Searched**



The proposed method reveals several superior solutions



#### CONCLUSION

- The <u>heuristics tend to underestimate complexity</u>.
- On the number of hidden layers, when seeking to optimize accuracy the <u>use of a second layer is desirable</u>
- <u>Ideally searching with the genetic algorithm</u> should <u>cover for the</u> <u>theoretical bounds of the number of hidden nodes</u> per layer

# **Questions?**

# **Thank You**

#### References

- 1. <a href="http://www.tandfonline.com/doi/pdf/10.1080/01431160802549278">http://www.tandfonline.com/doi/pdf/10.1080/01431160802549278</a>
- http://en.wikipedia.org/wiki/Genetic\_algorithm
- 3. <a href="http://www.obitko.com/tutorials/genetic-algorithms/parameters.php">http://www.obitko.com/tutorials/genetic-algorithms/parameters.php</a>
- 4. <a href="http://www.obitko.com/tutorials/genetic-algorithms/ga-basic-description.php">http://www.obitko.com/tutorials/genetic-algorithms/ga-basic-description.php</a>
- 5. <a href="http://en.wikipedia.org/wiki/Tournament selection">http://en.wikipedia.org/wiki/Tournament selection</a>
- 6. <a href="http://en.wikipedia.org/wiki/Mutation\_%28genetic\_algorithm%29">http://en.wikipedia.org/wiki/Mutation\_%28genetic\_algorithm%29</a>
- 7. <a href="http://www.cse.unr.edu/~sushil/class/gas/notes/scaling/index.html">http://www.cse.unr.edu/~sushil/class/gas/notes/scaling/index.html</a>
- 8. <a href="http://www.nd.com/products/genetic/crossover.htm">http://www.nd.com/products/genetic/crossover.htm</a>
- 9. <a href="http://yann.lecun.com/exdb/publis/pdf/lecun-90b.pdf">http://yann.lecun.com/exdb/publis/pdf/lecun-90b.pdf</a>
- 10. <a href="http://en.wikipedia.org/wiki/Hessian matrix">http://en.wikipedia.org/wiki/Hessian matrix</a>