Neural Network Weight Selection Using Genetic Algorithms

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Neural Networks

Neural networks generally consist of five components.

- A directed graph known as the network topology whose nodes represent the neurodes (or processing elements) and whose arcs represent the connections.
- A state variable associated with each neurode.
- A real-valued weight associated with each connection.
- A real-valued bias associated with each neurode.
- A transfer function f for each neurode such that the state of the neurode is $f(\omega_i x_i \beta)$.

Genetic algorithms require five components.

- A way of encoding solutions to the problem on chromosomes.
- An evaluation function which returns a rating for each chromosome given to it.
- A way of initializing the population of chromosomes.
- Operators that may be applied to parents when they reproduce to alter their genetic composition. Standard operators are mutation and crossover.
- Parameter settings for the algorithm, the operators, and so forth.

(19.3, 0.05, -1.2, 345, 2.0, 7.7, 68.0)

 $\stackrel{Mutation}{\longrightarrow} (19.3, 0.05, -1.2, 345, 2.0, 8.2, 68.0)$

Figure 1: Mutation Operation

(19.3, 0.05, -1.2, 345, 2.0, 7.7, 68.0), (17.6, 0.05, -1.2, 250, 3.0, 7.7, 70.0)

 $\stackrel{Crossover}{\longrightarrow} (17.6, 0.05, -1.2, 345, 3.0, 7.7, 68.0)$

Figure 2: Crossover Operation

Given these five components, a genetic algorithm operates according to the following steps.

- Initialize the population using the initialization procedure, and evaluate each member of the initial population.
- Reproduce until a stopping criterion is met.

Reproduction consists of iterations of the following steps.

- Choose one or more parents to reproduce. Selection is stochastic, but the individuals with the highest evaluations are favored in the selection.
- Choose a genetic operator and apply it to the parents.
- Evaluate the children and accumulate them into a generation. After accumulating enough individuals, insert them into the population, replacing the worst current members of the population.

Pattern Classification Problem

A Problem with k features and M classes:

Given a set of training examples, selects the most likely class for any instance not in the training set.

An **instance** is represented by a k-dimensional feature vector.

An **example** is represented by a instance and the class that the instance belongs to.

Real Examples

- Handwritten character recognition
- Speech recognition
- Blood cell classification
- . . .

Neural Networks

- Sigmoid Feed-forward Neural Network
- Weighted Probabilistic Neural Networks (WPNN)

Sigmoid Feed-forward Networks

A feed-forward network

- A directed graph without cycles.
- A multi-layer network Neurones in each layer (except output layer) are completely connected to the forward layer.
- Each neuron in the network is a sigmoid unit







Application for Pattern Classification (Cont.)

- Representation:
 - Input representation: $\langle x_1, x_2, \ldots, x_k \rangle$
 - Class representation: $<1,0,\ldots,0>$ for class 1
- Evaluation Function The sum of squared errors
- Training Algorithms
 - Backpropagation Algorithm
 - Genetic Algorithm

Weighted Probabilistic Neural Network (WPNN)

- WPNN is a pattern classification algorithm which falls into the broad class of "nearest-neighbor-like" algorithms.
- Likelihood Function: "distance" \rightarrow probability

Note: Distance is between an instance to all examples of a particular class, not just one example.

WPNN Likelihood Function

"Let the exemplars from class *i* be the *k*-vector \vec{x}_j^i for $j = 1, ..., N_i$. Then the likelihood function for class *i* is

$$L_i(\vec{x}) = \frac{1}{N_i(2\pi)^{k/2}(det\Sigma)^{1/2}} \sum_{j=1}^{N_i} e^{-(\vec{x} - \vec{x}_j^i)^T \Sigma^{-1}(\vec{x} - \vec{x}_j^i)}$$

 $L_i(\vec{x})$ describes the probability that the instance x is of class i." Note: the class likelihood functions are sums of identical anisotropic Gaussians centered at the examples divided by N_i .

WPNN Conditional Probability

The value of the likelihood function for a particular instance may fail to classify the instance. For example, $L_1(\vec{x}) = 0.3, L_j(\vec{x}) = 0.01$ for $j = 2, 3, \ldots, M$.

The conditional probability for class i:

$$P_i(\vec{x}) = L_i(\vec{x}) / \sum_{j=1}^M L_j(\vec{x})$$

WPNN Feature Weights

- Training WPNN consists of selecting the entries of matrix Σ .
- Matrix Σ is restricted to be a diagonal matrix.
- The inverse of each entry in Σ is a weight on the corresponding feature.

WPNN Implementation

It is called a "neural network" because its implementation is natural mapped onto a two-layer feed-forward network:

- k neurons in the input layer
- *M* neurons with standard normal Gaussian transfer functions in the output layer
- kN neurons in the single hidden layers, each with a linear function ax b.

 $k{:}$ number of features, $M{:}$ number of classes $N{:}$ number of total examples

WPNN Implementation (Cont.)

The ((i-1)k+j)th hidden neuron (i < N, j < k):

- 1. is connected to the *jth* input neuron with weight a = 1.0
- 2. has bias b equal to the value of jth feature of the ith example
- 3. is connected to the *nth* output neuron, where *n* is the class of the *ith* exemplar, with weight w_j , where w_j is the selected feature weight for the *jth* feature.

An Example

Two classes, two features, and three examples:

No.	Instance	Class
1	< 1, 2 >	1
2	< 2, 5 >	2
3	< 3, 4 >	1



Survey of Hybrid Systems

• Supportive combinations

Supportive combinations typically involve using one of these methods to prepare data for consumption by the other. For example, using a genetic algorithm to select features for use by neural network classifiers.

• Collaborative combinations

Collaborative combinations typically involve using the genetic algorithm to determine the neural network weights or the topology or learning algorithm.

Supportive Combinations

- Using NN to assist GA
- Using GA to assist NN
 - Data preparation
 - Evolving network parameters and learning rules
 - Using GA to explain and analyze NN
- Using GA and NN independently

Collaborative Combinations

GA to select weight

Two basic differences between different approaches: architectures (feedforward sigmoidal, WPNN, cascade-correlation, recurrent sigmoidal, recurrent linear threshold, feedforward with step functions and feedback with step functions) and difference in GA itself.

Collaborative Combinations

- GA to specify NN topology
 - A genotype representation must be devised and an attendant mapping from genotype to phenotype must be provided.
 - There must be a protocol for exposing the phenotype to the task environment.
 - There must be a learning method to fine tune the network function.
 - There must be a fitness measure.
 - There must be method for generating new genotypes.
- GA to learn the NN learning algorithm

Reasons to apply GA to NN

- Finding the global optima
- For recurrent networks
- For networks with discontinuous functions
- GA can optimize any combination of weights, biases, topology and transfer functions
- The ability to use arbitrary evaluation function

Disadvantage and Solutions

- Excessive computing time
- Solutions:
 - Using specialized NN hardware
 - Using the best local learning algorithm
 - Using parallel implementations of GA
 - By finding the best division of labor between the local and evolutionary learning paradigms to make the best possible use of training time

GA Formulation for Training Sigmoid FFNN

- Problem Formulation
- Initialization
- Genetic Operators
- Fitness Evaluation
- Genetic Parameters

Problem Formulation

- Individuals are the NN's themselves;
- no string encoding
- Topology is fixed, weights are evolved

Population Initialization

• Selection of initial weights



Plot of $e^{-|x|}$

Fitness Evaluation

• Sum of Squared Error

$$E(\vec{\omega}) \equiv 1/2 \sum_{d \in D} \sum_{k \in outputs} (t_{kd} - o_{kd})^2$$

Genetic Operators - Sigmoid FFNN

- Unbiased-Mutate-Weights Select from probability distribution
- Biased-Mutate-Weights Add in previous weight
- Mutate-Nodes Mutatation grouped by node (schema preservation)
- Crossover-Weights Uniform-crossover vs. point-crossover
- Crossover-Nodes Crossover grouped by node (schema preservation)
- Crossover-Features Competing conventions
- Mutate-Weakest-Nodes Non-random selection of least contributive
- Hillclimb One step in direction of gradient

Parameter Values

- Population-Size: 50
- Generation-Size: 1
- Parent-Scalar: [0.89, 0.93]

First, Comparing the three mutations resulted in a clear order

- 1. MUTATE-NODES
- 2. BIASED-MUTATE-WEIGHTS
- 3. UNBIASED-MUTATE-WEIGHTS



Second, a comparing of the three crossovers produced no clear winner.



Third, when MUTATE-WEAKEST-NODE was added to a mutation and crossover operator, it improved performance only at the beginning. The performance decreased after a certain small amount of time.



Finally, a generic training algorithm with operators MUTATE-NODES and CROSSOVER-NODES outperformed backpropagation on our problem.



Backpropagation Algorithm is more computationally efficient than Genetic Algorithm.

Five Components of the genetic algorithm with WPNN:

Representation

- a "logarithmic" representation for large dynamic range with proportional resolution;
- Map: $n \to B^{n-k_0}$

2. Evaluation Function

- "leaving-one-out" technique: a special form of cross-validation
- Performance function:

$$E = \sum_{i=1}^{M} \sum_{j=1}^{N_i} \left\{ \left[1 - \tilde{P}_i(\vec{x}_j^i) \right]^2 + \sum_{q \neq i} \left[\tilde{P}_q(\vec{x}_j^i) \right]^2 \right\}$$





For $k = 1, 2, \dots, k$ Err(k) = 0

1. Randomly select a training data point and hide its class label



For $k = 1, 2, \dots, k$ Err(k) = 0

- 1. Randomly select a training data point and hide its class label
- 2. Using the remaining data and given k to predict the class label for the left data point



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from the true label Repeat the procedure until all training examples are tested.



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Repeat the procedure until all training examples are tested.

Choose the k whose $\operatorname{Err}(k)$ is minimal

Empirical observations and theoretical arguments show WPNN works best when only a small fraction of the exemplars contribute significantly. So we reject a particular, for any exemplar x_i^i , if

$$\sum_{\vec{x} \neq \vec{x}_j} e^{-(\vec{x} - \vec{x}_j^i)^T \Sigma^{-1} (\vec{x} - \vec{x}_j^i)} > \left(\sum_{i=1}^M N_i\right) / P$$

Where P = 4

3. Initialization Procedure

WPNN: integers chosen randomly in [1, K].

Where K depends on the desired range and resolution for weight

4. Genetic Operators

WPNN: standard GA mutation and uniform crossover

Uniform crossover example:

Individual	Genotype					
А	abcdef	E				
В	qwerty	J				
Offspring	qbedti	f				

The gene at locus j, where $0 \le j < \text{string length}$, from both parents have equal selected for the new offspring.

Why uniform?

- A real-valued representation is used and no particular
- ordering to the feature weights.

5. Parameter Values

- Population-Size: 1600
- Generation-Size:
 - 1. "steady-state" approach. With the large population a steady-state GA is used, which means Generation-Size is small relative to Population-Size.
 - 2. for using a single CPU, Generation-size = 1.
- Parent-Scalar: the smaller the Parent-Scalar, the faster the converge.

Parent-Scalar = 0.9

Sample run of a steady-state GA

- 1. Initial population(randomly generated)
 - (a) (1001010) eval = 3
 - (b) (0100110) eval = 3
 - (c) (1101011) eval = 5
 - (d) (0110101) eval = 4
- 2. New children, from crossover or 3rd and 4th:
 - (a) (1101101) eval = 5
 - (b) (0110011) eval = 4

Experimental Results — WPNN

WPNN was a new and untested algorithm, the experiments with WPNN centered on the overall performance rather than on the training algorithm.

4 data set designed are used to illustrate both the advantages and shortcomings of WPNN.

First Data Set:

- 1. It is a training set that is generated during an effort to classify simulated sonar signals.
- 2. 10 features
- 3. 5 classes
- 4. 516 total exemplars

Second Data Set: Same as first data set except:

5 more features are added (which were random numbers uniformly distributed between 0 and 1 hence contained no information relevant to classification)

Third Data Set: Same as first data set except:

10 irrelevent features are added. (total of 20 features)

Fourth Data Set:

- Has 20 features just like the third data set.
- Pair each of the true feature with one of the irrelevant features.
- Mixing up the relevant features with the irrelevant features with via linear combinations.

•
$$0.5(f_i + g_i)$$
 and $0.5(f_i - g_i + 1)$

Dataset	1	2	3	4
Backpropagation	11	16	20	13
PNN	9	94	109	29
WPNN	10	11	11	25