

## An introduction to artificial neural networks

Coryn A.L. Bailer-Jones

*Max-Planck-Institut für Astronomie, Königstuhl 17, 69117 Heidelberg, Germany*  
*email: calj@mpia-hd.mpg.de*

Ranjan Gupta

*IUCAA, Post Bag 4, Ganeshkhind, Pune-411007, India*  
*email: rag@iucaa.ernet.in*

Harinder P. Singh

*Department of Physics, Sri Venkateswara College, Benito Juarez Road, Dhoola Kuan*  
*New Delhi-110021, India*  
*email: hps@ttdsvc.ernet.in*

**Abstract.** Artificial neural networks are algorithms which have been developed to tackle a range of computational problems. These range from modelling brain function to making predictions of time-dependent phenomena to solving hard (NP-complete) problems. In this introduction we describe a single, yet very important, type of network known as a feedforward network. This network is a mathematical model which can be trained to learn an arbitrarily complex relationship between a data and a parameter domain, so can be used to solve interpolation and classification problems. We discuss the structure, training and interpretation of these networks, and their implementation, taking the classification of stellar spectra as an example.

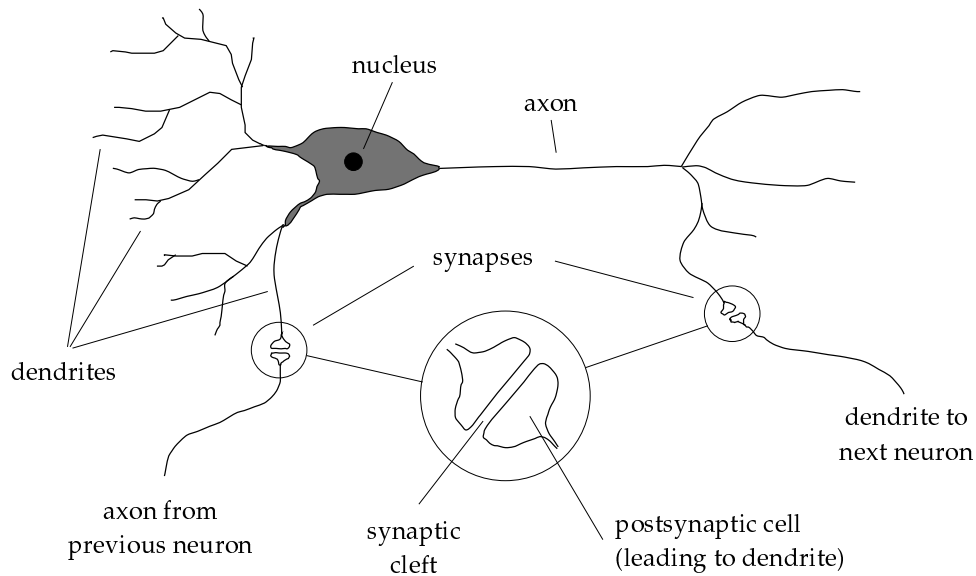
*Key words:* neural networks, nonlinear models, classification, interpolation, data modelling

### 1. Biological neural networks

Artificial neural networks were originally introduced as very simplified models of brain function, so it is initially instructive to consider the broad analogies between artificial

neural networks and biological ones. Of course, the latter are considerably more complex, and many artificial neural network models today bear very little resemblance to biological ones. For a history of the development of artificial neural networks, see, for example, the introduction of Hertz, Krogh and Palmer (1991).

The human brain consists of billions of interconnected neurons. These are cells which have specialized membranes which allow the transmission of signals to neighbouring neurons. Fig. 1 shows the structure of a neuron. A single *axon* extends from the basic cell body. This is the output from the neuron, and typically divides into many sub-branches before terminating at a *synapse*. Electrical pulses are transmitted along the axon to the synapses by the transfer of  $\text{Na}^+$  ions. The arrival of a voltage pulse stimulates the release of neurotransmitting chemicals across the synaptic cleft towards the postsynaptic cell, which is the receiving part of the next neuron. This postsynaptic cell passes the signal via the *dendrite* to the main part of the neuron body. The inputs from the different dendrites are then combined to produce an output signal which is passed along the axon, and so the process continues. However, a signal is only produced in the axon if there are enough inputs of sufficient strength to overcome some threshold value, and then the output is some nonlinear function of the input stimuli.



**Figure 1.** Structure of a single neuron in the brain. Information is passed into the neuron from other neurons via the dendrites to the left. Given appropriate input stimuli, the cell sends an output signal along the axon to the synapses, where the signal is transmitted to other neurons.

The human brain consists of about  $10^{11}$  neurons, and each neuron has between a few and a few thousand synapses on its dendrites, giving a total of about  $10^{14}$  synapses (connections) in the brain. The ‘strength’ of the synaptic connection between neurons

can be chemically altered by the brain in response to favourable and unfavourable stimuli, in such a way as to adapt the organism to function optimally within its environment. The synapses are therefore believed to be the key to learning in biological systems.

## 2. Artificial neural networks

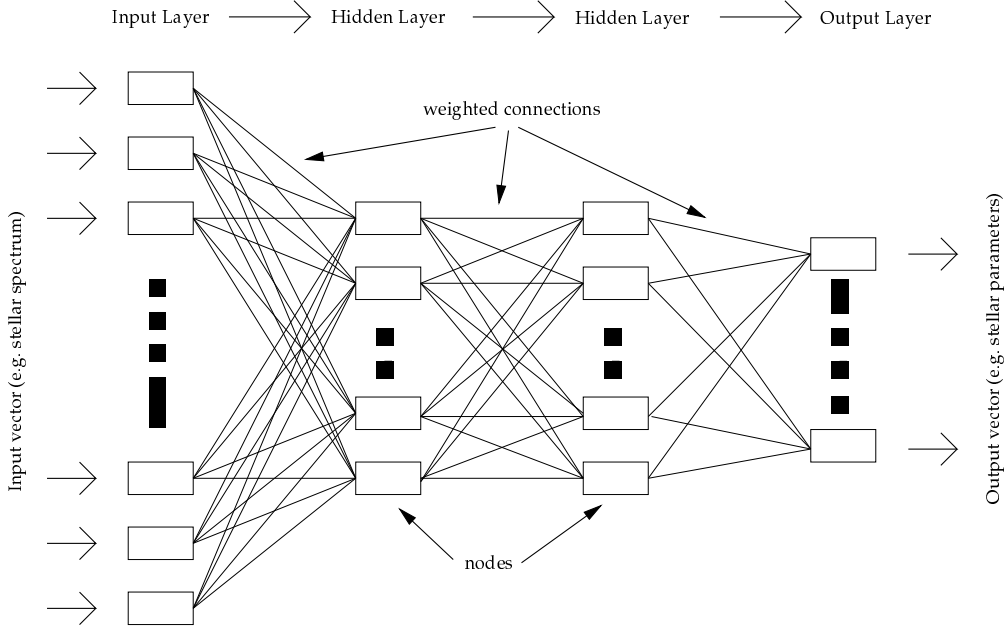
“Artificial neural networks” is a relatively loose term referring to mathematical models which have some kind of distributed architecture, that is, consist of processing nodes (analogous to neurons) with multiple connections (analogous to dendrites and axons). These connections generally have adaptable parameters which modify the signals which pass along them. There are numerous types of artificial neural networks for addressing many different types of problems, such as modelling memory, performing pattern recognition, and predicting the evolution of dynamical systems. Most networks therefore perform some kind of data modelling, and they may be split into two broad classes: *supervised* and *unsupervised*. The former refers to networks which attempt to learn the relationship between a data and a parameter domain, while the latter refers to networks used to find “natural” groupings with a data set independently of external constraints. What they have in common is the idea of learning about a problem through relationships intrinsically present in data, rather than through a set of predetermined rules. An introduction to several types of neural networks (but by no means all) is given by Hertz, Krogh and Palmer (1991). A less mathematical overview is provided by Beale & Jackson (1990).

In this article we introduce what is one of the most important types of supervised neural networks, called a *feedforward multilayer perceptron*. The term *perceptron* is historical, and refers to the function performed by the nodes. *Feedforward* means that there is a definite input and output, and a flow of data in one direction. This is in contrast to *recurrent neural networks* in which data flows in a loop: these are important when time plays a relevant role in the problem (see, e.g., Bailer-Jones 1998a). A comprehensive treatment of feedforward networks is provided by Bishop (1996), where more details on many of the themes discussed below can be found.

Fig. 2 shows a feedforward neural network with several inputs, two hidden layers, and several outputs. Each node in the input layer holds a value,  $x_i$ . In our example application, the input vector,  $(x_1, x_2, \dots, x_i, \dots)$ , is a stellar spectrum, and the output vector,  $(y_1, y_2, \dots, y_i, \dots)$ , is a set of stellar parameters, e.g.  $T_{\text{eff}}$ ,  $\log g$  and  $[M/H]$ . Each of the input nodes connects to every node in the next layer of nodes, the first ‘hidden’ layer, and each of these connections has a weight,  $w_{i,j}$ , associated with it. A node in the hidden layer forms a weighted sum of its inputs, and passes this through a *nonlinear transfer function*, such that the output from the  $j^{\text{th}}$  hidden node is

$$p_j = \tanh \left( \sum_i w_{i,j} x_i \right) . \quad (1)$$

These values are passed to a second hidden layer which performs a similar processing, the output from that layer being the vector  $\mathbf{q}$



**Figure 2.** Feedforward artificial neural network architecture. This particular example shows two hidden layers. Not explicitly shown are the bias nodes: each layer has an extra node that holds a constant value (and has no inputs) and provides an offset to the subsequent layers. These are necessary for the network to model functions properly.

$$q_k = \tanh \left( \sum_j w_{j,k} p_j \right) . \quad (2)$$

The output layer then performs a simple sum of its inputs, so that the network output,  $\mathbf{y}$ , is

$$y_l = \sum_k w_{k,l} q_k . \quad (3)$$

The tanh function in the hidden layers provides the nonlinear capability of the network. Other nonlinear functions are possible; the sigmoidal function ( $1/(1 + \exp[-\sum wx])$ ) is often used. Both functions map an infinite possible input range onto a finite output range,  $-1$  to  $+1$  in the case of tanh. This imitates the transfer function of neurons.

### 3. Network training

The weights,  $\mathbf{w}$ , appearing in equations 1-3 are the free parameters of the network. Clearly, in order for the network to yield appropriate outputs for given inputs, the weights must be set to suitable values. This is done by *training* the network on a set of input vectors for which the ideal outputs (or *targets*) are already known. This is *supervised learning*. The idea is that the network will encapsulate the relationship between, in our example, a wide range of types of stellar spectra and their associated physical parameters. The network does not come up with a new classification system, but rather will *interpolate* the training data to give a *generalization* of the relationship between spectra and their classifications. We can then use this trained network to obtain reliable classifications for unclassified spectra.

Supervised learning proceeds by minimizing an error function with respect to all of the network weights. The error function typically used is the sum-of-squares error, which for a single input vector,  $n$ , is

$$e^{\{n\}} = \frac{1}{2} \sum_l \beta_l (y_l^{\{n\}} - T_l^{\{n\}})^2 \quad , \quad (4)$$

where  $T_l$  is the target output value for the  $l^{\text{th}}$  output node. The  $\beta_l$  terms allow us to assign different weights to different outputs, and thereby give more priority to determining certain outputs correctly.

The most popular method for training the network is with the *backpropagation algorithm* (Rumelhart, Hinton and Williams 1986a,b), in which we determine the gradient of the error function with respect to each of the weights in the network. The network is first initialized by setting the weights to small random values. We then pass one of the training vectors (spectra) through the network and evaluate the outputs,  $y_l$ , as well as all the intermediate node values in the network (the vectors  $\mathbf{p}$  and  $\mathbf{q}$ ). Differentiating equation 4 with respect to a weight,  $w_{k,l}$ , in the final weights layer and substituting for  $y_l$  from equation 3 gives

$$\frac{\partial e}{\partial w_{k,l}} = \beta_l (y_l - T_l) q_k \quad , \quad (5)$$

where we have used the fact that the weights are causally independent of one another and of the node values. (We have dropped the  $n$  superscript to make things less crowded.) We can determine the gradient of  $e$  with respect to a weight,  $w_{j,k}$ , in the previous weights layer in the same way, but now  $q_k$  is a function of  $w_{j,k}$ ,

$$\frac{\partial e}{\partial w_{j,k}} = \sum_l \beta_l (y_l - T_l) \frac{\partial y_l}{\partial w_{j,k}} = \frac{\partial q_k}{\partial w_{j,k}} \sum_l \beta_l (y_l - T_l) w_{k,l} \quad . \quad (6)$$

The remaining partial derivative can then be evaluated from equation 2. Thus we see that the error gradient with respect to any weight in the network can be evaluated by

*propagating* the error dependency back through the network to that weight. This can be continued for any number of layers, up to the input layer, giving us the complete error gradient vector,  $\partial e/\partial \mathbf{w}$ , where  $\mathbf{w}$  is the set of all network weights.

Having evaluated the error gradient vector, there are a number of ways in which it can be used to train the network. The most common is the *gradient descent* process, in which we adjust the weight vector in the direction of the negative of the gradient vector, i.e.

$$\Delta \mathbf{w} = -\mu \frac{\partial e}{\partial \mathbf{w}} . \quad (7)$$

The factor  $\mu$  determines how large a step is made, and typically has a value between 0.1 and 1.0. The gradient is then recalculated using the new values of the weights, and over many iterations the weights should move towards a value which represents a small value of the error,  $e$ . For training to strictly converge (i.e. reach an exact minimum),  $\mu$  would have to decrease to zero during training, although in practice a sufficiently small error can often be achieved without this.

Of course, we are generally interested in getting the network to generalize its input–output mapping for a range of types of objects (e.g. different classes of stars), so we must correspondingly use a range of objects which are representative of the problem. We therefore apply the above training algorithm successively to each vector in a training sample. We may either update the weights after each vector is passed, or we can keep the weights fixed and update only after the errors have been calculated for all vectors in the training set. In this latter case, known as *batch* training, we update using the the average error,

$$E = \frac{1}{N} \sum_{n=1}^{n=N} e^{\{n\}} , \quad (8)$$

and the corresponding average error gradient vector.

The network training is a nonlinear minimization process in  $W$  dimensions, where  $W$  is the number of weights in the network. As  $W$  is typically large, this can lead to various complications. One of the most important is the problem of local minima. Clearly, training using the gradient descent technique will stop upon reaching the bottom of a minimum, but this may only be a *local* minimum, not the *global* minimum, and may correspond to a much larger error. To help avoid local minima, a *momentum term* is sometimes added to the weight update equation 7, in which a fraction,  $\nu$ , of the previous weight update is added to the current one. This provides some ability to “escape” from small local minima: the larger is  $\nu$  the larger are the minima which can be ignored, although possible at the expense of slower convergence. A brute force approach to avoiding local minima is to retrain the network several times from different initial random weights, and see whether each case converges to the same solution. If the majority do, this could be taken as the best solution. Alternatively, we could use all

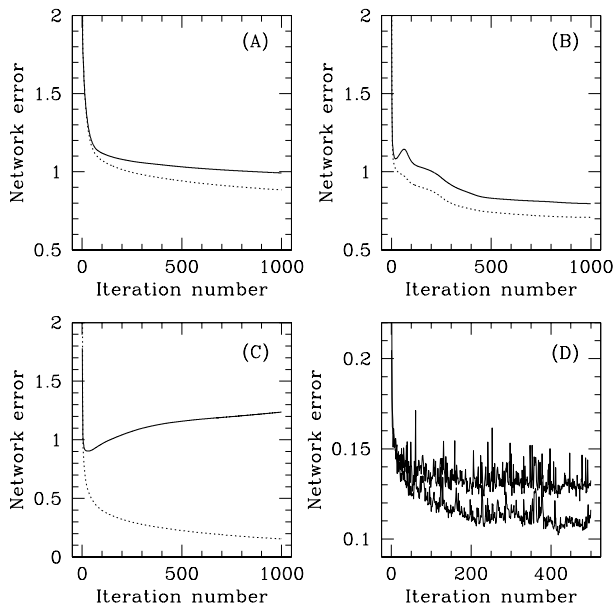
of these networks together in a *committee*, in which the outputs from the networks are averaged.

A common difficulty is knowing when to stop training, as true convergence in all but very simple (or fortuitous) cases is rarely achieved. Without searching the entire weights space, it is generally impossible to know whether the minimum reached is the global one or just a local one, and if local, how much better the global minimum is. Furthermore, the minimum may have a long, almost horizontal bottom (thinking analogously in three dimensions), so it will often be necessary to terminate training when the gradient falls below some small value. Setting the convergence gradient too small will slow convergence; too large and the search may terminate prematurely.

Fig. 3 shows several examples of how the error in equation 8 changes during training. The dotted line shows the error on the actual training data. However, we really want to test whether the network has generalized its mapping, rather than “memorized” the specific training data. The solid line therefore shows the error evaluated on a separate test data set, which is not used to modify the weights. Under certain circumstances it is possible for the network to fit the noise in the training set, in which case it is said to have *overtrained*. This can be seen in Fig. 3c, where the error on the test data starts to rise at some point, although the error on the training data continues to fall. Frequently, both errors continue to drop monotonically until levelling out (Fig. 3a), which is desirable. (A difference in the errors finally achieved may reflect differences in the training and test data sets.) Fig. 3b shows an example of where the error on the test data rises briefly, and then drops: we should not, therefore, cease training at the first sign of a rising test error. In certain cases the error can be very noisy (Fig. 3d), oscillating around a minimum value, probably due to  $\mu$  in equation 7 being held constant. Anecdotal evidence suggests that this is more likely to occur when multiple outputs are used, although it depends on the minimization algorithm used.

Gradient descent only uses knowledge of the local gradient, which will not, in general, point towards the global minimum, thus requiring many steps to reach this minimum. Numerous other optimization techniques exist to overcome this problem. One is the method of *conjugate gradients*, in which a more efficient gradient is defined using the second order derivatives (albeit implicitly) of the error with respect to the weights. Several other methods make direct use of the second derivatives (or *Hessian* matrix) to obtain more rapid convergence. We refer the interested reader to any number of books on optimization techniques for further details.

The mapping produced by a neural network is an interpolation of the training data. Whenever data are interpolated, there needs to be some kind of complexity control to prevent overfitting the data. For example, if we imagine fitting a curve through five noisy data points in a two-dimensional space, then – unless the data are co-linear – a fourth order polynomial model will give a better fit (a perfect one in fact) than a straight line. This is unlikely to be appropriate, though, as the data contain noise. Hence there is a trade off between obtaining a good fit and not fitting the noise, and a compromise can be achieved through the use of a *regularization* technique. A very basic approach is to split the training sample into two parts, and during training on one half, monitor the total error on the other half. As seen in Fig. 3, the test error *may* start to rise after some number of iterations, indicating that the network has begun to fit the noise (or *overfit*



**Figure 3.** Examples of the evolution of the network error during training. The dotted (solid) line is the error evaluated on the training (test) data (except in case D where the lower error is for the training data). (A),(B) and (C) are for networks with a single continuous output. (D) is for a 50:5:5:3 network with probabilistic outputs. All cases used gradient descent with constant  $\mu$  and  $\nu$  terms. See Bailer-Jones (1996) for more details.

the data). Training could then be stopped just before the error starts to rise. However, experience shows that in some cases the training error never rises, even after a very large number of iterations have occurred and a small error has been achieved.

Another approach to regularization is *weight decay*. A high order fit to data is characterized by large curvature of the mapping function, which in turn corresponds to large weights. We can penalise large weights by adding the following term to the training error in equation 8

$$\alpha \frac{1}{2} \sum_i w_i^2, \quad (9)$$

where the sum is over all weights in the network and  $\alpha$  is some constant.

Many neural network applications in astronomy have used gradient descent for training, to the extent that some people think gradient descent and neural networks to be synonymous. This is not the case: the feedforward neural network is a general means of representing a nonlinear relationship between two domains, with the feature that the error gradient is easily calculable in terms of the free parameters (the weights) of the model. What optimization technique is then used to determine the best solution for the



weights is really a separate issue. Gradient descent is often used because it is easy to code, but it may not be the best choice.

#### 4. A Bayesian perspective

A rather different approach to interpolating data with a feedforward network is provided within a Bayesian probabilistic framework. If we go back to first principles, then we see that the problem we are trying to answer with the neural network is fundamentally probabilistic: Given the set of training data,  $D$ , and some background assumptions,  $H$ , what is the most probable output vector,  $\mathbf{y}^*$ , for some given input vector,  $\mathbf{x}$  (not in  $D$ )? When phrased this way, we see that we are not really interested in the network weights, and, moreover, an “optimal” set of network weights is not actually required. For example, if we have trained a network and found a good mapping, we may retrain it and find a different set of weights which give a mapping which is almost as good. Thus, while a given input will yield much the same output as before, the weights may be very different. In the optimization approach we ignore all but one solution (except when using committees), whereas in a Bayesian approach we combine all such solutions by the process of *marginalization*. Specifically, if the output of the network with weights  $\mathbf{w}$  and input  $\mathbf{x}$  is written  $\mathbf{y}(\mathbf{x}, \mathbf{w})$ , then

$$\mathbf{y}^* = \int \mathbf{y}(\mathbf{x}, \mathbf{w})P(\mathbf{w}|D, H)d\mathbf{w} . \quad (10)$$

In other words, we weight each possible output by the probability,  $P(\mathbf{w}|D, H)$ , that the training data would produce those weights. This *posterior* probability for the weights is given by Bayes’ Theorem

$$P(\mathbf{w}|D, H) \propto P(D|\mathbf{w}, H)P(\mathbf{w}|H) . \quad (11)$$

The first term on the right hand side of this equation is the *likelihood*. It is the probability that a network with a given set of weights produces the targets in the training data,  $D$ , when the inputs from these data are fed in. If we assume a Gaussian error model for the discrepancies between outputs and targets, this is just

$$P(D|\mathbf{w}, H) \propto \exp -E , \quad (12)$$

where  $E$  is the sum of squares error from equations 8 and 4. The other term in equation 11 is the *prior probability* over the weights, and represents our prior belief of what the weights should be. In this Bayesian framework, the weight decay regularizer mentioned earlier (equation 9) is naturally incorporated into this prior.

From equations 11 and 12 we can evaluate all of the terms under the integral in equation 10. However, this integral must usually be evaluated numerically with Monte Carlo methods. This involves replacing the integral with a summation over  $\mathbf{y}(\mathbf{x}, \mathbf{w})$  for

a large number of values of  $\mathbf{w}$  drawn from the distribution  $P(\mathbf{w}|D, H)$ . In this Bayesian approach, the optimization of the weights is replaced with the integration over all possible weight values, so no “optimum” weight vector is (or has to be) calculated. While this approach is theoretically more satisfying than an optimization, it is time consuming, as the marginalization must be carried out for each new input vector we wish to classify. Partly for this reason there exists an approximation to this approach which, although still within the Bayesian framework, returns the problem to one of optimization. There are several advantages to the Bayesian approach, such as automatic complexity control (avoidance of overfitting) and a natural way for determining the uncertainty in each network output. More on the Bayesian approach for feedforward neural networks can be found in (MacKay 1995). For a general introduction to Bayesian methods, Sivia (1996) is recommended.

## **5. Implementation issues**

Now that the basic principles of neural networks have been laid out, it is necessary to ask how they are used in practice to solve a problem. We illustrate this with the problem of stellar spectral classification.

### **5.1. Training and testing data**

The first thing which must be specified is the training data set, consisting of spectra preclassified on the classification system of interest. It must be emphasized that the network cannot be more accurate than these preclassifications, so considerable effort must often be spent in assembling the training data. We must also ensure that the examples in the training data set cover the full range of types to which we will want to apply the trained network: We cannot expect a network trained only on B stars to classify M stars well, as the characteristic features of M stars are not present in B star spectra. Conversely, we do expect a network trained only on early and late B stars to classify mid-B stars well, as B stars all have qualitatively similar spectra.

Once a network has been trained, we want to evaluate its performance, and this must be done using a separate data set. As mentioned earlier, it is possible for a network to fit the noise and thus memorize the specific training data set rather than generalize from it. This can be a particular problem if the data are noisy or if the network is very complex (i.e. has many hidden nodes so is flexible to model many nonlinearities). We can test for this by training the network on only a subset of the available data and then evaluating it on the rest, ensuring that the two data subsets have similar distributions of objects.

### **5.2. Form of the inputs**

The input to the network is some measured feature vector. In the case of stellar classification this will be a spectrum or a list of fluxes through various filters, or both. We could also input other measurements we believe to be relevant, such as the parallax or interstellar extinction. The inputs need not be contiguous or in any particular order. A stellar spectrum typically contains a large amount of redundant information (from the

point of view of general classification), so using the entire spectrum as the input may not be necessary. If we can reduce the number of input nodes, we reduce the number of weights and hence the dimensionality of the mapping function. This results in faster training, and may reduce the chance of getting stuck in local minima. It also increases the data/weights ratio (see section 5.5) and density of the data, thus providing a more reliable mapping. There are a number of approaches to *dimensionality reduction*. The simplest is to remove any inputs which we know, or suspect, to have little or no influence on the problem in hand. A more sophisticated approach is with Principal Component Analysis (PCA), which finds the most discriminatory linear combinations of the inputs. By removing the less significant principal components, we can compress the data with negligible loss of discriminatory information, and filter out some noise too. PCA is discussed in the article by Singh, Bailer-Jones & Gupta in these proceedings.

### 5.3. Output modes

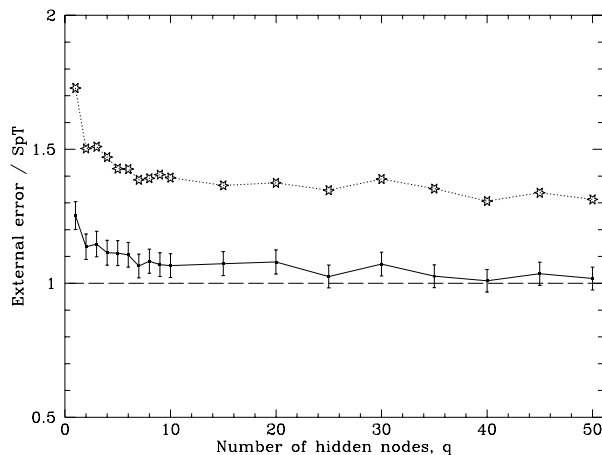
The output vector is the set of classifications. For example, we may have one output for MK spectral type (SpT) and another for luminosity class (LC). We could alternatively use separate networks for each parameter. As the network outputs are real numbers, these classes are represented on a continuous scale. Thus the network will be able to produce intermediate classes. This is useful for the spectral type parameter, as spectral types are simply points on what is really a continuous scale (closely related to effective temperature). We refer to this way of representing the outputs as *continuous mode*. Luminosity classes, on the other hand, have more claim to being discrete, for which we may want to use the network in *probabilistic mode*, whereby each class is represented by a separate output. By confining each output to the range 0–1 (e.g. by replacing equation 3 with the function  $0.5 \times [1 + \tanh]$ ), we can interpret the output as the probability that the input is of that class. For a five class problem, the target vector for an object in class 2 would be (0, 1, 0, 0, 0). When we apply the trained network to new data, we would assign the class with the largest probability, although if that probability is below some threshold, or not much larger than the probability of some other class, then we may prefer to label this object as uncertain.

Clearly, the probabilistic mode offers more information than the continuous one. For example, if the input is a composite of two classes (e.g. two stars of different brightnesses), we can represent both in the target vector (e.g. 0,0.25,0,0.75,0). Thus if the network thinks an input is composite, it can (in principle at least) tell us this. The only way we could model binary star spectra in continuous mode is to have two spectral type outputs, although then we also need some means of knowing when the input spectrum is just a single star. See Weaver (2000) for an attempt to classify binary spectra with neural networks. A disadvantage of the probabilistic mode is that it may result in many more network weights than a network in continuous mode.

### 5.4. How many hidden nodes and layers are required?

Deciding upon the appropriate number of hidden nodes and layers is largely a matter of experience. With many problems, sufficient accuracy can be obtained with one or two

hidden layers and 5–10 hidden nodes in those layers. There is a theorem which states that a network with one hidden layer can approximate any continuous function to arbitrary accuracy, provided it has a sufficient number of hidden nodes (Hornick, Stinchcombe and White 1990). In practice, such a large number of nodes may be required that it is more efficient to go to a second hidden layer. This can be seen in the work of Bailer-Jones et al. (1998b). Fig. 4 shows how the error in determining the spectral type (in continuous mode) varies as a function of the number of hidden nodes in a network with just one hidden layer. The error is already reasonably small for a network with just one hidden node, which is equivalent to a network with no hidden layer. Adding a few hidden nodes improves the error, but once there are 5–10 nodes, the error does not decrease further. However, if we add a second hidden layer, then a network with 5 nodes in each of these layers produces a  $\sigma_{68}$  error of 0.85 SpT, a statistically significant improvement. A similar improvement using a second hidden layer was seen again in a related (but more complex) problem (Bailer-Jones 2000).



**Figure 4.** Variation of the network classification error as a function of the number of hidden nodes,  $q$ . The solid line is for  $\sigma_{68}$  and the dotted line for  $\sigma_{RMS}$ ; they represent the core and outliers of the error distribution respectively (see Bailer-Jones et al. 1998b). The network architecture is 50:q:1 with a continuous spectral type output. The error bars are  $3 \times \delta$ , where  $\delta$  is the standard deviation in  $\sigma_{68}$  due to a finite sample size. The mean RMS error in the target classifications themselves is 0.65 SpT.

### 5.5. Ensuring a well-determined network solution

An important consideration in setting up the network is the ratio of data to weights. If we train a network with one output using  $N$  training vectors, we have a total of  $N$  error measurements with which to determine the  $W$  weights. Thus we have a system of  $N$  equations with  $W$  unknowns, and, if the weights and training vectors are independent of each other, we require  $N > W$  to find a solution. With fewer data the solution for the

weights will be underdetermined. Thus a 10:5:5:1 network (indicating 10 inputs, 5 nodes in each of two hidden layers, and one output) has 91 weights (including the bias nodes), so will require at least as many training examples. The data/weight ratio estimation is more complex with multiple outputs (especially if they are correlated, as will be the case in probabilistic mode), but if there are two independent outputs, then the number of error measures is  $2N$ .

In practice, this “overdetermination” requirement may not be so strict. For example, Gulati et al. (1994) used only 55 training examples to train a 161:64:64:55 probabilistic network. This network has about 18,000 free parameters (each output is parametrized by about 15,000 weights), yet only of order 55 error measures were available to determine them. In spite of this, a reasonably small error on a test data set was obtained. We suspect that many weights were essentially not trained, and, by staying at their small initial values, played no role in the network. Additionally, correlations between the inputs (flux bins of stellar spectra) could result in correlated weights in the input–hidden layer. Both of these reduce the effective number of free parameters. Other cases of apparently good results with formally underdetermined weights appear in the literature, implying that correlated or unused weights are not uncommon. Nonetheless, in many of these cases simpler networks could probably have been used without loss of accuracy.

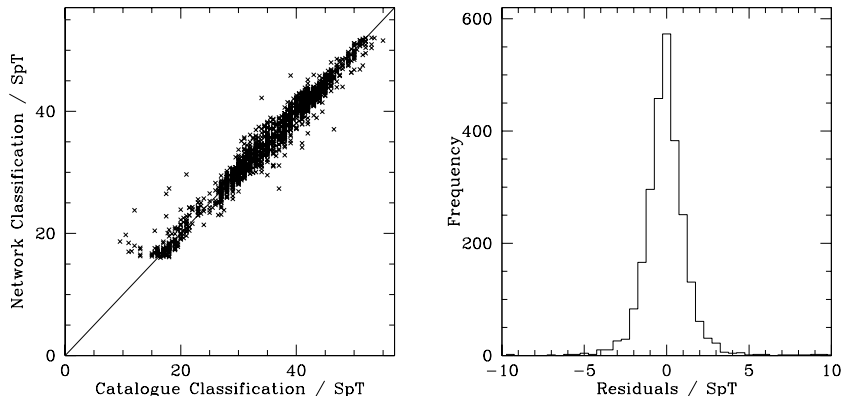
## 6. Example Applications

The paper by Gupta et al. in these proceedings gives an application of a 93:16:16:17 network to classify 2000 bright sources from IRAS LRS spectra.

Fig. 5 shows the performance of a committee of ten 50:5:1 networks applied to the spectral classification problem, trained using gradient descent. It shows the correlation between the network classifications and the “true” classifications (those from the training catalogue), and the histogram of the error residuals, for a test data set. There is an even scatter about the ideal classifications in the left hand plot, except for the hottest (early type) stars, where the network assigns cooler classes.

A similar plot to that in Fig. 5 is not very useful for a network used in probabilistic mode when there are only a handful of classes. In such cases it is more useful to look at the *confusion matrix*. Examples are shown in Fig. 6 for networks with three outputs, corresponding to the three luminosity classes III, IV and V. In probabilistic mode, we can also examine the confidence with which the network classifies objects. An example is shown in Fig. 7, where we plot the cumulative fraction of objects classified with a probability greater than some amount. This plot is for the 50:5:5:3 network with the line+continuum spectra from Fig. 6. Thus while 98.1% of the class V objects were correctly classified (i.e. a fraction 0.981 were classified with a probability,  $p > 1/3$ ), only 77.5% have a probability of over 0.90. Part of the reason for this is that a network with sigmoidal outputs can never produce an output of exactly 0.0 or 1.0, as that requires very large weights, something which regularization tries to avoid. (See Bailer-Jones 1996 and Bailer-Jones et al. 1998b for more details on these examples.)

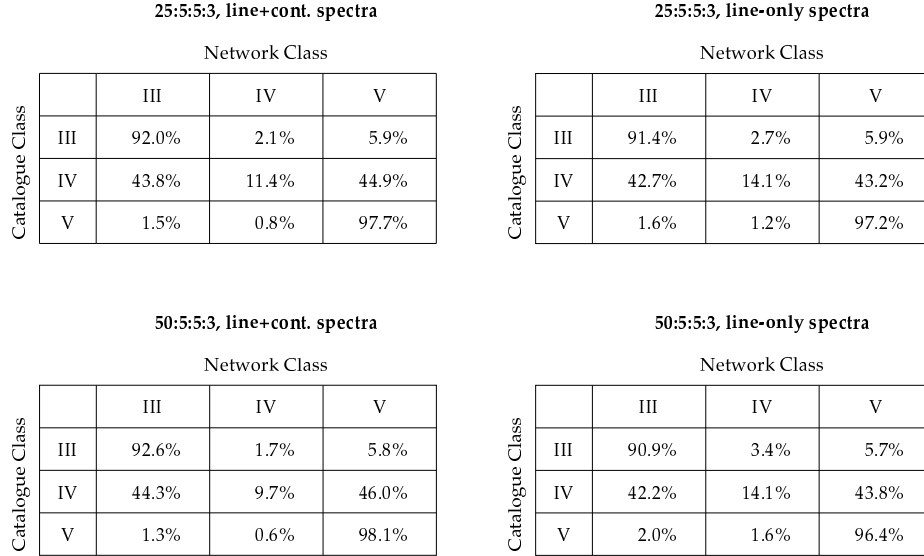
## 7. Understanding the network



**Figure 5.** Results from a committee of ten 50:5:1 networks with a single continuous spectral type output. The left plot shows the correlation between network and catalogue classifications (perfect classifications would fall on the diagonal line), and the residuals from this are plotted on the right. The spectral types O3 to M9 have been coded by the integers 1 to 57. The summary results are  $\sigma_{68} = 1.03$  SpT and  $\sigma_{RMS} = 1.35$  SpT. The internal error of the networks is 0.41 SpT. This is a measure of the average differences in classifications which members of the committee assign to a given spectrum.

The underlying principle of a neural network is that it learns its input–output mapping from the relationship intrinsically present in a preclassified data set. Unlike rule-based classification algorithms, the networks are not explicitly told anything about the relevance of certain inputs in determining the output. This relevance information is expressed in the optimized weight vectors, and in principle we can look at the values of the weights to find out which inputs are significant in determining the outputs. In practice, however, this may be rather difficult on account of there being more than one hidden node (which is necessary to get a nonlinear mapping). For example, weights leading from a relevant input to different nodes in the hidden layer could be both small and large in magnitude. There is no constraint on the network training to provide easily-interpretable weights, although such a one may be possible. Furthermore, there may be many minima in the weights space which have approximately equal errors, and the corresponding “optimal” weight vectors may give conflicting opinions about the relevance of the inputs. A degeneracy in the weights is made more likely if there are redundant, or correlated, inputs, which is often the case with stellar spectra. Ultimately, obtaining both an accurate model of a complex problem *and* a good understanding are conflicting demands, as understanding often requires simplification.

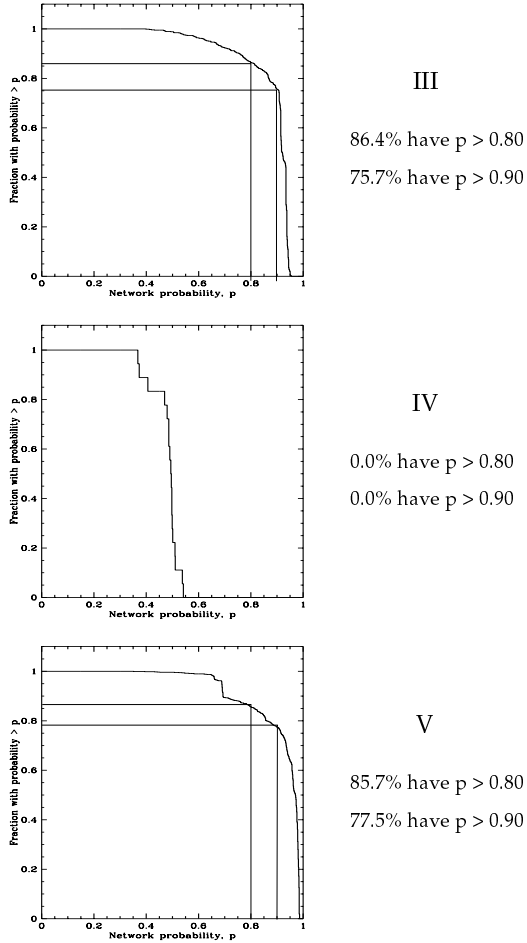
We can nonetheless get some idea of *how* a network produces its classifications. Fig. 8 shows the correlations between the values on the hidden nodes of a network and the outputs from that network. Looking at the first hidden layer, we see that no single node shows a low-scatter, monotonic relationship with spectral type. If one did, then that alone could be used for spectral type classification and we could dispense with the network altogether. Generally, the correlations are nonlinear, and with large scatter. However, node 1 does show linearity over a limited part of the spectral range (SpT =



**Figure 6.** Confusion matrices for four different network models differing in the number of PCA inputs and whether or not the continuum has been subtracted from the spectra. Each matrix lists the percentage of spectra which have been classified correctly and incorrectly: in the top-left matrix the network classifies 1.5% of spectra which are class V in the catalogue as class III, whereas it correctly classifies 97.7% of class Vs. In all four cases the networks' do not believe in the existence of luminosity class IV (random assignments would lead to about 33% in the middle row).

28–42, or F0–K0). This is interesting, as it means that for classes F and G this node could serve as the output node and we would get reasonable classifications. Looking back at equation 1, we see that if we were to re-parametrize the spectral type as  $\tanh^{-1}$  SpT, we could classify F and G stars using a simple linear regression on the spectral indices (as  $\tanh^{-1}$  is monotonic over the range of interest). Nodes 3–5 also show linear correlations over small portions of the spectral type range. However, all nodes show either some insensitivity (due to saturation of the node values) or large scatter in the correlation, for at least some part of the spectral type range.

When these values are passed to the second hidden layer, the situation is somewhat different. The correlations are still nonlinear, but there is much less scatter. Furthermore, node 5 has a low scatter, *linear* correlation for spectral types earlier than about F3 (SpT = 30), and node 4 shows a linear correlation for types later than about K0 (SpT = 42). Node 2 (or 3) alone could reasonably classify all intermediate types. Thus we see that there is some localization in the network, with certain nodes taking on the job of identifying certain ranges of the target variable. There is, of course, no significance in the numbering of the nodes within a layer, as the nodes could be shuffled with no change in the final output.

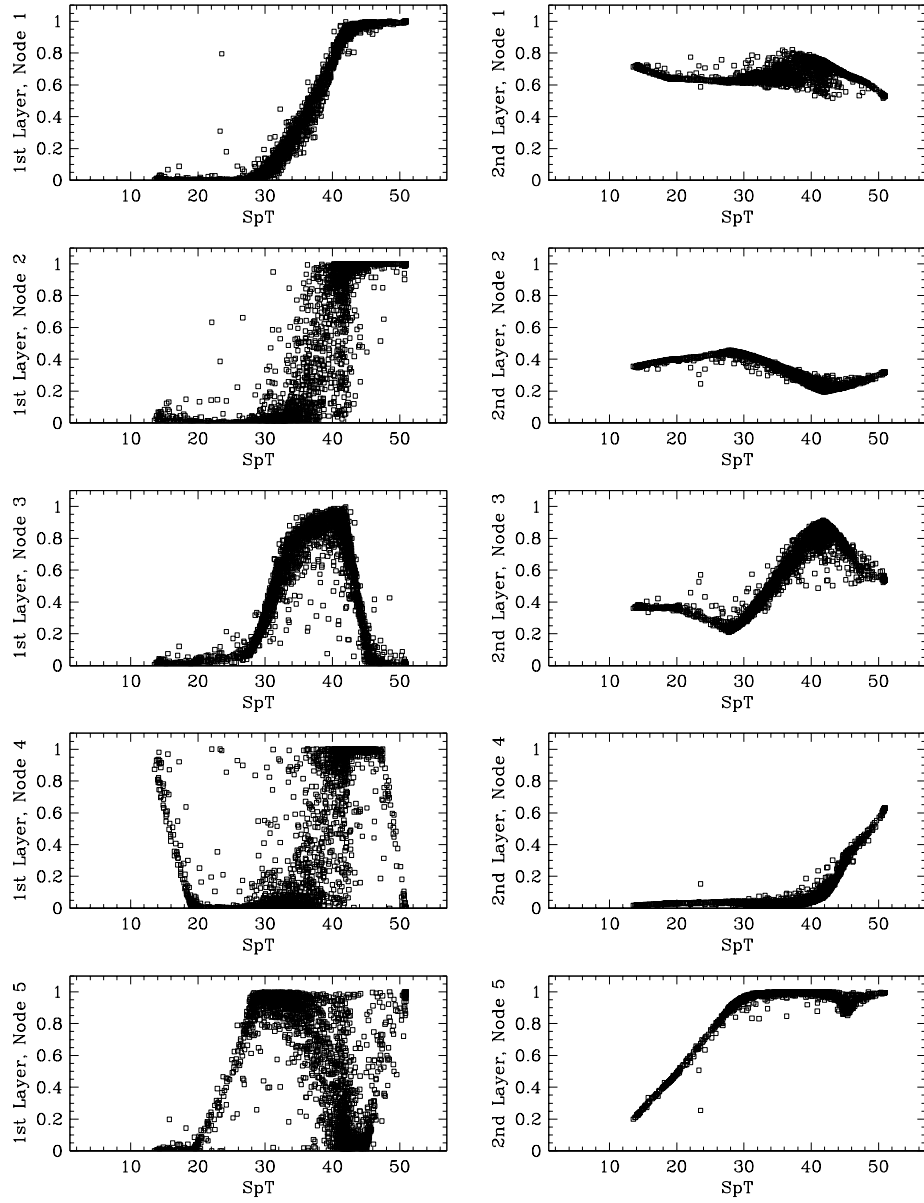


**Figure 7.** Confidence of correct network classifications based on network output probability. Each figure is a cumulative plot of the fraction of spectra which have been classified with a probability  $> p$ , for the luminosity classes III, IV and V.

## 8. Summary

We have introduced the basic concepts of feedforward neural networks. Although the type introduced, the multilayer perceptron, is only one of many types, it is very important as judged by its wide and numerous applications. We have described the backpropagation algorithm for training the network, as well as some of the ways in which it can be implemented to optimize the weights. We also briefly looked at a Bayesian approach to modelling with a network, which marginalizes over the weights rather than optimizing





**Figure 8.** Correlation between the node values and outputs for a network trained to give spectral types in continuous mode. The column on the left (right) represents the first (second) hidden layer of nodes in a 50:5:5:1 network. Each point is the node value produced by one input vector in a test data set. The transfer function in each layer (including the subsequent output layer) is a sigmoid function, which restricts the node values to the range 0–1.

them. A number of implementation issues were discussed and example applications were presented.

Although it is interesting to see how these models were originally inspired by attempts to model brain behaviour, these artificial neural networks really have very little in common with biological neural networks, and, in our opinion, are better understood in purely mathematical terms. Whilst they may be powerful models, there is nothing “mysterious” about them. They are simply parametrized nonlinear models: the input–output function of a two-hidden layer model is described entirely by equations 1–3. They are no more a “black box” than is a straight line fit to two-dimensional data. The nonlinearity and multidimensionality of neural networks are present because real-world problems are often complex. If we have difficulty understanding how they work, then it is because an *accurate* description of real-world problems often requires that we avoid the simplifying assumptions which typically make easily-understood descriptions only approximate.

We finish by stressing that the purpose of these models is to generalize the relationship between a data domain and a parameter domain which is intrinsically present in a preclassified data set. They are useful because they can model complex relationships in a large number of parameters, and generalize this to areas of the input space where training data are sparse. This has potential applications to large, multiparameter astronomical surveys (see Bailer-Jones, these proceedings). But the models can only be as good as the data on which they are trained (although they can be a lot worse if trained badly), so construction of an accurate and relevant training data set is of fundamental importance.

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