



An overview on the use of neural networks for data mining tasks

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An Overview of the Use of Neural Networks for Data Mining Tasks

First author: Full name and affiliation; plus email address if corresponding author [*Frederic Stahl; University of Portsmouth; Frederic.Stahl@port.ac.uk]
Second author: Full name and affiliation; plus email address if corresponding author [Ivan Jordanov; University of Portsmouth]

Abstract

In the recent years the area of data mining has experienced a considerable demand for technologies to extract knowledge from large and complex data sources. There is a substantial commercial interest as well as research activities in the area that aim to develop new and improved approaches for extracting information, relationships, and patterns from datasets. Artificial Neural Networks (NN) are popular biologically inspired intelligent methodologies, whose classification, prediction and pattern recognition capabilities have been utilised successfully in many areas, including science, engineering, medicine, business, banking, telecommunication, and many other areas. This paper highlights the implementation of NN, trained by supervised and unsupervised learning, for pattern recognition, classification, prediction and cluster analysis, from a data mining perspective, and discusses some examples of their usage in bioinformatics and financial data analysis tasks.

Introduction

Currently, ubiquitous computer systems record our habits, decisions, behaviours and lives daily. We use debit and credit cards to pay and club cards to save on our shopping. We are recorded on CCTV cameras and leave our information in the World Wide Web using applications such as email, blogs, twitter or social networks. But it is not only us, the individuals that constantly generate data records in various large data bases every day, it is also scientific applications that record large amounts of data constantly. For example, the NASA's system of earth orbiting satellites which generates a terrabyte of data every day⁶¹; the human genome project¹ that stores the genetic blueprint of our bodies; molecular simulations that study the dynamic behaviour of molecules and generate large amounts of data stored in distributed data warehouses⁵³. Considerable advances in storage technology made it possible to store all this information at relatively low cost. However, just having the data available in databases does not make us more knowledgeable. We may be 'information rich', but still 'knowledge poor'¹⁵ and the gap between recording information and understanding it is growing⁷.

Data mining can be used to close the gap between collecting and understanding the information. It can be regarded as the extraction of meaningful, and potentially useful patterns from large data bases⁶⁵, or referring to the earlier used expressions, as the 'understanding' of recorded 'information'. For the extraction and discovery of knowledge, data mining employs a variety of intelligent methods, among which, NN are frequently used for different data mining tasks, including classification and cluster analysis. The authors of²⁸ highlight some of the advantages of using NN for data analysis: they are tolerant to noise; can deliver high accuracy on complex non linear mappings and at the same time can be implemented on parallel hardware; they are highly maintainable (can easily be updated with fresh data) and can easily be automated.

This paper highlights the use of NN in the field of data mining. First, the basic Knowledge Discovery process with data mining is outlined and the basics of NN are introduced. Next, the NN are put into the context of data mining and the Knowledge Discovery process and their usage for data mining tasks is described. This is followed by a summary of actual applications of NN, more specifically in bioinformatics and finance areas and finally, concluding remarks are made.

KNOWLEDGE DISCOVERY AND DATA MINING

Knowledge Discovery from Data

Data Mining is often described with the metaphor of 'gold mining'¹⁵. In it, the 'gold' is the actual valuable 'knowledge' to be extracted. Thus, according to this metaphor, 'data mining' should be better named 'knowledge mining' as pointed out in¹⁵. In fact, another popular term used, instead of 'data mining', is 'Knowledge Discovery from Data' (KDD). In the KDD process, data mining is often regarded as just one, but very important step.

The following steps (Fig.1) describe the main activities involved in the KDD process:

1. *Integration and Cleaning*: Integration of different data sources into one data store and removal of inconsistent data and noise;

2. *Selection and Transformation*: Selection of data that is relevant to the analysis task and its transformation into a standard format. Also, data reduction techniques may be applied in this step;
3. *Data Mining*: Applying intelligent methods to the preprocessed data in order to extract meaningful patterns, such as decision rules;
4. *Evaluation and Interpretation*: Identification of 'interesting' aspects of the extracted patterns and representing them in a meaningful way.

Data Mining

The data mining step in the KDD process aims to extract meaningful patterns from large amounts of data, using different intelligent techniques¹⁵. There are two typical fields of application that can be automated by the use of data mining of large databases: (a) the prediction of trends and behaviours; and (b) the extraction of unknown patterns and relationships.

With respect to (a), classifiers can be generated from labelled data samples. These classifiers can then be applied to previously unseen data instances in order to classify them. This is one of the most important data mining techniques. Typical applications of classifiers are in the area of marketing⁵⁸. Data mining can be applied on past promotional mailings in order to classify receivers into groups that are likely or unlikely to take advantage of the promotions. This information can be used in the future to save money by only targeting groups that are likely to respond to the promotion. Further applications of classification are risk assessment for credit card loan⁵⁰ and mortgage applications¹⁰, as well as bankruptcy predictions. However, prediction comprises not only classification of unlabelled data, but also prediction of future trends. For example, prediction of future trends could be useful for forecasting the future values of stock market papers, stock market indices, etc⁶⁹.

With respect to (b), there are applications that are concerned with the discovery of new, hidden knowledge/patterns in large data bases, rather than predictive modelling. For example, drug designers are interested in discovering knowledge from gene expression datasets about influences of the designed drugs on gene expression levels. Drugs may promote or suppress certain genes which in turn may influence further genes and thus create a complex pattern⁷⁰. Also in the business area patterns that indicate which products are likely to be bought together are useful for product placement⁷¹. Or in banking, discovered patterns may indicate fraudulent credit card transactions²².

Commonly used methods in data mining comprise:

- *Artificial Neural Networks (NN)*: NN are biologically inspired predictive models (mimicking the functioning of human brain), that can learn and map linear and nonlinear functions⁵²;
- *Genetic Algorithms (GAs)*: GAs are nature inspired evolutionary approaches, that simulate natural selection and survival of the fittest; they are often used for global optimization tasks, and also to optimise other pattern recognition algorithms⁵⁶;
- *Generalised Rule Induction (GRI)*: generates rules about significant relationships, rather than just predicting a class value^{51, 2};

- *Top Down Induction of Decision Trees (TDIDT)*: Induces classification rules in the intermediate form of a tree structure³⁸;
- *K-Nearest Neighbour (KNN)*: It classifies unlabelled data instances with a majority class of the k most similar data instances in the training set⁷.

ARTIFICIAL NEURAL NETWORKS

This paper focuses on the use of NN in the data mining step of the KDD process (Fig. 1). This section highlights NN in general. The use of NN in data mining is discussed in the 'Neural Networks in Data Mining' Section. NN are computer paradigms inspired from the biological NN that can be used to generate predictive models from historical data.

They have been applied in data mining for prediction, classification and cluster analysis. Initially, the application of neural networks has faced strong opposition due to the fact that they are complex structures, difficult to interpret and need a long training time³³. Yet NN produce compelling accuracy and are highly adaptable across a large range of applications. There have been also developed technologies that extract explicit rules using NN, hence NN are currently experiencing a renaissance in data mining³³.

Analogy to Biological Neural Networks

The term artificial neuron can be traced back to the 1940s, in an early work of McCulloch and Pitts³², where it was described as a mathematical simulation of a biological neuron³². This paper highlights the most common form of an NN, the Multi Layer Perceptron (MLP) and briefly introduces alternative topologies.

A biological neuron, as illustrated in Fig. 2(a), sends a signal if the collective influence of all its inputs reaches a threshold level (*axon hillock's* threshold) – the neuron is said to **fire**. The impulse is sent from the *cell body (soma)*, through the *axon*, which influences the *dendrites* of the next neuron over narrow gaps called *synapses*. They translate the pulse in some degree into **excitatory** or **inhibitory** impulse of the next neuron. It is this receiving of signals, summation and transferring procedures that are emulated by the artificial neuron (a perceptron, Fig. 2(b)).

As in elaborate 'Lego' constructions and integrated circuits, which are built from a multitude of simple constituent units that interact with one another, the NN consists of organised in layers, interconnected neurons, as shown in Fig. **Error! Reference source not found..**

Although the units from Fig. 3 can perform very simple tasks (as summation and transferring signal depending on the sum), when organised in NN, they can solve complex mapping, classification and recognition tasks. In the NN literature these units are also often referred to as neurons, nodes, or perceptrons. Neurons influence each other using 'weighted' connections between them; positive weights have stimulating influence, while the negative ones have inhibiting influence. The number of neurons in the input and output layers depend on the problem's inputs and outputs, and the number of units in the hidden layer depends on the complexity of the problem or concept to be represented and solved. The higher the absolute value of the weight the stronger the influence of one neuron to another. The input layer receives data from the outside world in the form of a vector which is weighted and passed through the connections to the (first) hidden layer. In each hidden unit

the incoming products of the weights and the input values are summed and 'squashed' through an activation/transfer function that generates an output value, which in turn is input for the next layer and so on. Two commonly used transfer functions are the sigmoid and the hyperbolic tangent functions.

The calculation of weighted input for the k^{th} unit of the hidden layer is then:

$$v_k = \sum_{i=1}^n w_{ki} x_i, \quad k=1, \dots, m;$$

where $x_i, i=1, \dots, n$ are the inputs, w_{ki} are the weights and m is the number of units in the layer. The output y_k of neuron k is then calculated by a transfer function: $y_k = \phi(v_k)$ and y_k is then input value for the next layer of units.

Training and Testing of Neural Networks

An NN is typically learned using training, validation and testing phases. During the training phase the weights of the NN are adjusted using training data and the validation data subset is used to avoid so called 'overfitting' (when the NN starts learning more to memorise, rather than to generalise), and in the testing phase the NN is checked whether and how well it has learned the task. For predictive data mining, the training and testing in its basic form divides the dataset into two subsets, a train subset and a test subset. The train subset is used to train/learn the NN and the test subset is used to check whether the NN learned the knowledge and can generalise. Generalisation is the ability of a trained NN to correctly classify, fit, map, recognise input instances/samples/examples that have not been used during the training.

In the *training phase* there are several learning rules that can be used and they determine how the weights are being adjusted. However, most learning approaches can be categorised into supervised and unsupervised learning.

- In *supervised learning* the correct output for each input training data instance is a-priori known. The training input instances are fed into the network, propagated, the actual output of the network is compared with the target one, and if there is a difference, then the weights are adjusted in order to minimise this difference.
- In *unsupervised learning* the output is not known, and the network itself adjusts the weights in order to map inputs to outputs without any additional knowledge.

Backpropagation^{63, 45} is the most popular training technique for supervised learning of MLPs. It uses gradient based (deterministic) methods for minimising an error function, which represents the difference between the actual and the desired output, through adjusting the weights of the network. During this process there is a danger of overfitting, when the error function becomes zero, but the NN is learned to memorise, rather than to generalise. In this case, an additional validation subset is used in parallel to the training one and when the validation error starts to increase, the training should stop (regardless of further decrease of the training error).

In the *testing phase* there are no changes performed on the networks weights, once trained, the NN weights are fixed. During this phase, a testing (unseen during the training) data subset, is presented to the NN, in order to validate how well the NN can generalise (in other words, its ability to produce the target output when presented with unseen input instances).

Further possible learning rules that can be used for adjusting the weights of a NN are, for example, Hebb's rule¹², the Delta Rule⁶⁴, Competitive Learning²⁶, and global optimisation techniques, such as Genetic Algorithms⁷², Simulated Annealing²⁵, etc.

Artificial Neural Network Types

There are a number of variations of NN topologies and architectures. Most of them are feedforward NN (FFNN). In FFNN the information moves only in one direction from the input nodes through the hidden and to the output nodes. Contrary to the FFNN, recurrent NN have a bi-directional data flow, as they propagate data from later processing stages into earlier ones. MLPs can differ in the number of hidden, input and output units. In Radial Basis Function (RBF) networks⁴⁴ each neuron in the hidden layer represents a point in the input space. The activation of a hidden neuron depends on the distance of the input values to the point that the hidden neuron represents in the⁶⁵. However, Radial Basis Function networks will not be discussed further in this article, it is referred to⁴⁴ for more information. A special NN architecture for cluster analysis will be highlighted in the 'Cluster Analysis' Section.

NEURAL NETWORKS IN DATA MINING

This section describes the basic data mining process in the context of NN. Fig. **Error! Reference source not found.** takes a closer look into the data mining step of the KDD process illustrated in Fig. **Error! Reference source not found.**. This is done in the context of NN based on Ni's description in³³. Input to the data mining process is the preprocessed data and the patterns generated by the data mining step are called 'useful rules'.

Below is a description of the steps in the data mining process based on NN (Fig. 4):

- *Data Preparation*: converts the data into a suitable format for the particular neural network being used. This involves replacing missing values or deleting data instances with missing values; filtering out instances that introduce noise; and converting categorical data values into numerical ones as NN process numerical data;
- *Learn NN*: The NN is trained, validated and tested for generalization on the prepared data and rules are generated in the form of network weights. In data mining, these learned NN can be treated as 'black boxes' and already be applied;
- *Rule Generation*: Often rules are extracted from NN in order to be able to assess and refine the generated rule set;
- *Rule Assessment*: Loosely speaking, rule assessment validates the rule sets (the NN quality) and possibly refines certain rules. Possible assessment criteria for a rule set or single rules may be the rules, rule sets or the NN's accuracy; finding the optimal order in which rules are being applied; detecting contradicting rules in order to manipulate them, etc.

The following sections highlight some data mining tasks, in which NN are used.

Classification and Numeric Prediction

Two of the most important tasks in data mining are classification and numeric prediction of variables of data instances. An example for *classification* is the analysis of credit card applications. Credit card institutions wish to assess whether issuing a credit card is ‘safe’ or ‘risky’. Usually, a classifier is generated that can be used to predict a categorical class label such as ‘risky’ or ‘save’ for previously unseen data instances. An example of *numeric prediction* is a sale at a mail-order house. A marketing analyst is interested in predicting how much a certain customer or a group of customers is likely to spend. A model or predictor is generated that predicts a numeric value.

Classification and numeric prediction usually happen in a two step process as shown in Fig. **Error! Reference source not found.** First, the classifier is generated on a training data set, of which the class label or the numeric value of interest is known. This is also known as the *training phase*. In the second step, the model is used for classification or numeric prediction. However, these classifiers tend to overfit on the training data, meaning that they memorise anomalies and noise rather than learning the general trend. Thus, this second step is often used to validate the classifier/predictor using test data with known class labels, in order to verify the classifier’s accuracy and ability to generalise. There is a variety of techniques used for training and testing (depending on the size and quality of the available data sets), e.g., *split sample*, *k-fold crossvalidation*, *bootstrapping*, etc.^{7, 15, 65}, which discussion is however outside the scope of this article.

Three popular approaches for classification/predictions are *Bayesian Classification*, *Rule Based Classification* and *Classification by Backpropagation*. Bayesian classifiers use statistical methods based on the ‘Bayes theorem’²⁷, which do not predict a particular class value, but the likelihood of a data instance belonging to a particular class. Furthermore, they are fast and thus can be applied to large data volumes. Rule based classifiers are in the form of decision rules, these can be decision trees or IF-THEN rules as illustrated in Fig. **Error! Reference source not found.** A notable classifier for Decision trees is ‘C4.5’³⁸, and for IF-THEN rules ‘Prism’⁸. Decision trees are often converted into IF-THEN rules. Both approaches have the problem that they suffer from noisy training datasets, tend to learn the noise, hence the generated rule sets need to be manipulated in order to gain a high classification accuracy. However, they are often used if insights are needed about the predictions that have been made.

NN, as outlined earlier, can be used for classification purposes. In particular, Backpropagation MLPs are used as classifiers and predictors¹⁵. They have been initially criticised as less suitable for data mining due to their poor interpretability. This is because their symbolic meaning is hidden in the neurons’ connections and thus is difficult to be assimilated by humans, hence NN are often referred to as a ‘black box’¹⁵. Their high tolerance to noisy data and the fact that they can classify patterns they have not been trained on, makes them an excellent technology for data mining. Furthermore, NN are well suited to learn continuous data without prior discretisation and predict numeric values, which is not possible with most rule based and Bayesian classifiers. Their ability to produce non linear models enables them to approximate any function with a reasonable accuracy^{20, 11, 19}. One of their disadvantages is that they are computationally not very efficient, for example, Backpropagation can be entrapped in local minima, yet they are inherently parallel and thus can easily be executed on

parallel computer architectures. In order to do so for decision trees and rule based algorithms, parallel versions need to be developed first, such as the discussed in⁴⁹.

NN are gaining popularity in data mining due to advances in rule extraction techniques from trained NN, which enhance the NN's suitability for classification and numeric prediction. A fully connected NN is usually too complex to be described in the form of rule sets, hence weighted connections between nodes that have least effect on the trained NN are removed prior to the rule extraction. A weighted link may be removed if it, for example, does not decrease the networks classification accuracy during testing. A popular approach to extract rules from NN is the NeuroRule algorithm that expresses rules as binary input^{48, 29}. The basic steps of NeuroRule are outlined below:

1. Train the NN;
2. Prune the network by removing redundant connections whilst maintaining its accuracy;
3. Discretise activation values of the neurons in the hidden layer and the variable values of the input layer;
4. Generate rules that explain the relationship between the output and the hidden layer;
5. Generate rules that explain the discretised activation values of the neurons in the hidden layer in relation of the network input values;
6. Generate a set of rules that relate the inputs and outputs of the NN by merging the two rule sets created in step 4 and 5.

Further rule extraction methods comprise the SUBSET algorithm^{46, 13} and the MofN algorithm⁵⁹.

Cluster Analysis

Cluster analysis aims to organise a dataset into groups of 'similar' data objects according to some distance function. Cluster analysis is an unsupervised technique, which is exploratory in its nature, thus usually performed when there exist no or very little prior knowledge about the hidden concepts in the dataset. There are many application areas, such as finance, (e.g., an analyst may want to find clusters of companies that have similar financial performances); health and medical applications, (an analyst may want to cluster patients with similar symptoms or reactions of patients to a certain drug); marketing, (building clusters of customers with similar shopping preferences), etc.⁷.

The authors of⁶⁷ describe the clustering process in four steps as shown in Fig. **Error! Reference source not found.**

1. Usually, the first step is to select and extract features – a very important part of the process, especially if there is a large amount of features in the dataset. It improves the likelihood of finding useful clusters and also increases the algorithms' speed. Feature selection aims to only select the most relevant characteristics for distinguishing a cluster, whereas feature extraction aims to create novel features from two or more original characteristics.
2. Clustering algorithms usually use proximity metric in order to measure the resemblance of individual data instances to each other. The clustering techniques use this proximity metric in

order to build groups of data instances that are 'similar' (in other words, being close in the feature space).

3. Clustering methods tend to build clusters regardless of whether there is an apparent structure in the data that determines different clusters or not. Also, different clustering algorithms tend to converge to different solutions, regardless of using the same datasets. Hence, clusters need to be validated in order to strengthen the users or analysts confidence in the found solution.
4. Finally, the built clusters need to be interpreted in order to provide insights into the meaning of the original data.

Two popular clustering methods comprise *k-means*¹⁶ and *agglomerative hierarchical clustering*^{15, 74}. K-means is one of the simplest clustering techniques, an *exclusive clustering algorithm*, meaning that each data object can only be assigned to one cluster. The analyst has to decide the number of clusters *k* to be built, and usually they are initialised randomly. Next, each data point is assigned to a cluster whose centroid (the centre of the cluster according to the used proximity metric) is nearest. After all data objects are assigned to clusters, the centroids are recomputed, and each data object is reassigned. This is repeated until there are no changes in the clusters. The disadvantage of the *k-means* algorithm is that the analyst needs to have some prior knowledge of the number of clusters in order to generate meaningful ones.

In *agglomerative hierarchical clustering*, each data object is initially considered as its own cluster. Next, the distance between each pair of clusters is calculated and the closest pair of clusters is merged into a single cluster. This is repeated until all data objects are assigned to a single cluster. Agglomerative hierarchical clustering produces a dendrogram, a binary tree of all clusters build in each iteration.

Kohonen networks (also known as *feature maps*, or *topology-preserving* maps, or simply SOMs)²⁶, are very popular methods used for cluster analysis¹⁵. Each output node will represent one cluster (class) of inputs and is associated with a *weight vector*, which size is the same as the *input vector*. There are also lateral connections between the neighbouring output neurons (Fig. 8).

Unlike the FFNN, there is no hidden layer and no transfer function used. Also, during the unsupervised learning neither the clusters, nor which node corresponds to which cluster is specified. The SOM network should work out the clusters for itself and organize the output nodes accordingly. For any given sample input, the output units compete to represent it, and the node with the closest weight vector becomes the winner.

The essential process in the formation of the SOM comprises the following steps:

1. *Competition*: For each input pattern the neurons in the output layer will compete to represent this pattern. The neuron with the closest weight vector to the input vector will update its weights and will become a winner. The closeness is defined by a distance metric (e.g., *Euclidian*, *Hamming*, or others);
2. *Cooperation*: The winning neuron determines the topological neighbourhood, by exciting its surrounding neurons. This is called cooperation, because it strengthens the neuron that is the winner and to some extend the neurons that are close to it;

3. *Synaptic Adaptation*: It adjusts the weights of the connections in order to enable the excited neurons to update their weight vectors to become 'closer' to the input vector.

Cluster analysis using SOMs does not require prior knowledge about the number of clusters, which represents a clear advantage compared with the *k-means* technique. SOMs can also be used to detect outliers in the data and clusters are built in order to isolate instances that are distinctly different from the remaining instances. Sometimes outliers are of more interest than the actual clustered data instances. For example, let the data instances represent branches of a retailer chain. If there are several clusters that comprise branches with different increases of revenue, then it is likely that a branch whose revenue did not increase is not inside any cluster. A closer look into this branch and comparison with branches in clusters that have had an increase in revenue may reveal the reason for its bad performance. For example, this could be a result of distinctly different advertising strategy (the outlier branch may have advertised in the local newspaper, whereas the remaining (non outlier) branches used radio advertisements, or, there may be a smaller customer parking space, comparing with the others, etc.).

APPLICATIONS OF NEURAL NETWORKS

Artificial neural networks have been used in many areas of bioinformatics⁷⁵ and financial forecasting⁶⁹. This section highlights general problems in these two chosen areas of NN application and discusses some particular examples.

Bioinformatics

Bioinformatics is an interdisciplinary science (which encompasses computer science, mathematics and statistics), that aims to refine biological information into biological knowledge using computer based methodologies. It has been coined in 1978 by Paulien Hogeweg^{17, 18} and gained popularity with the mapping of the human genome in 2001¹. Dynamic modelling of complex biosystems, sequence analysis, protein structure analysis are just few problem areas that fall within the scope of bioinformatics, but it could be broadly categorised into genomics and proteomics. Genomics investigates the structure and function of a very large number of genes, including mapping sequencing of genes for the entire genome of organisms, and proteomics studies the protein-protein and protein-DNA interactions, enzymatic and biochemical pathways of an organism⁵⁴. Proteins are the building blocks of an organism; the blueprint for a particular protein is encoded in a particular gene of the organism's genome. Proteins are composed of a chain of amino acids and genes are essentially composed of base pairs, consisting thymine, adenine, cytosine and guanine⁷³. A gene is a substring of the DNA that codes for a particular protein. Within a gene, a sequence of three consecutive base pairs codes for a particular amino acid in the protein. Now, loosely speaking, in genomics the sequence of base pairs and expression levels of genes are analysed, whereas in proteomics the protein's chemical structure and thus its function is analysed. The authors of^{39, 47, 55, 66} outline approaches of using NN for the analysis of protein structure prediction and gene sequence analysis.

One of the biggest problems associated with data mining algorithms in bioinformatics is the curse of dimensionality⁵. While the number of genes in the *Escherichia coli* bacterium is just 4288, the relevant number of genes in a human genome is estimated about 35000, meaning that there are 35000 genes that code for about 35000 proteins. This way microarray datasets can easily reach up to

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35000 features (depending on the microarray chip). Yet patterns in this data are often described by a small subset of the feature space. Also, measurements of gene expression levels are generally very noisy. Genomics data is described by a very large number of features compared to the limited number of samples. This makes achieving high accuracy very difficult for task for the predictive algorithms. Hence, feature selection strategies³¹ are used to overcome the curse of dimensionality.

Sequence Alignment

In bioinformatics, a sequence alignment is a way of arranging the sequences of DNA base pairs, or protein amino acids, to identify regions of similarity that may be a consequence of functional, structural, or evolutionary relationships between the sequences. Following are two references to particular applications where NN have been successfully used for such sequence alignments^{4, 23}. The authors of⁴ developed a molecular alignment method using a Hopfield NN. They successfully reproduced the three-dimensional structure of the real molecular alignments, which have been obtained experimentally using X-ray crystallography. The authors of²³ developed a neural network architecture named GenTHREADER that was used to predict the similarity between gene sequences. In two cases GenTHREADER successfully predicted the similarity between two structures.

Promoter Sequence Detection

Promoters are regulatory sequences in the genome that are necessary for the initiation of the processes that read the DNA sequence of a particular gene in order to synthesize proteins. Promoters are regions on the gene that mostly control the biological activation of the gene⁷⁶. An early investigation³⁰ describes the problem of detecting promoter genes in the context of NN. Because only a small portion of promoter sequences are known, NN can only be trained on a limited subset of promoter sequences versus random ones; yet NN achieve a high accuracy in detecting real promoter sequences. The authors of²⁴ used a multilayered NN successfully for the prediction of a particular promoter sequence (the mycobacterial sequence). The authors of⁷⁸ developed a technique to differentiate promoter regions from the rest of the genome based on an unsupervised SOM, which leads to significantly fewer false predictions than existing promoter prediction techniques⁷⁸.

Gene Expression Analysis and Gene Regulatory Networks

The process that converts the gene's base pair sequence into proteins is called gene expression. The more and the faster proteins from a certain gene are synthesised, the higher the expression level of this gene. Gene expression levels can be measured using microarray technology³⁶. Gene expression levels are often used to compare regular and cancerous cells, e.g., the authors of^{34, 3, 62} have been using NN successfully on classification problems in the context of cancerous cells. Gene expression data is also used to create gene regulatory networks; these are maps that describe how genes influence each other's expression levels. In^{41, 43} NN have been utilised successfully to identify gene-gene interactions in order to create gene regulatory networks. The authors of⁴¹ developed clustering technique called adaptive double self-organizing map (ADSOM), which does not require prior knowledge about the number of clusters. The authors of⁴³ show that machine learning strategies for optimising a NN architecture, for the identification of gene-gene interactions in complex human diseases, may be preferable to traditional trial-and-error approaches.

Protein Structure Prediction

Proteins are composed of a chain of up to twenty different amino acids. A protein chain folds in a particular way due to chemical forces between amino acids. Important is the 3D structure of the protein chain as it determines the protein's function. Hence the prediction of the 3D structure of a protein from only knowing the amino acid sequence is important. However, predicting the 3D structure of a protein solely based on the chemical properties of the amino acid sequence is computationally too expensive; hence research follows alternative approaches^{53, 6}. One approach is the alignment of the amino acid sequence of the protein with the unknown structure with the amino acid sequences of homologous proteins with known 3D structures. Yet this method only works if there are homologues proteins in the database, hence alternative methods such as NN are needed. Following are two references to particular applications where NN have been successfully used for protein structure prediction^{42, 14}. The authors of⁴² use separate NN tailored for predicting three standard structures (alpha-helix, beta-strand, and coil). The authors of¹⁴ created a method based on single hidden layer feed-forward NN for predicting different types of beta-turn structures from the amino acid sequence of a protein. NN based technologies have also been used to fight diseases such as HIV. For example the authors of⁷⁷ propose a framework using NN based technologies in order to predict the specificity of HIV-1 protease cleave sites (enzymes that postprocess proteins).

Finance

NN are of practical interests in financial data mining as it has been shown in^{79, 40} that they often perform better than statistical approaches for financial forecasting, fraud detection, bankruptcy prediction and marketing.

Financial Market Forecasting

Data mining for financial data has been particularly interesting for financial time series prediction for stock markets in order to maximise the trading revenue. A recent study used a MLP architecture in order to predict the Shanghai composite index⁶⁹ based on historical data of the closing prices of the stock market. In order to overcome the lack of explanatory capabilities of the NN, trading strategies (for example buy & sell & hold) have been incorporated. Furthermore the system predicted possible returns. Another recent application for NN is the forecasting of the next day's electricity price in the Spanish electricity market³⁵. Their approach is based on MLPs and historical time series data about the electricity demand and price. For a comprehensive review about neural network technologies their application on forecasting stock markets, it is referred to⁸⁰.

Credit Card Fraud Detection and Risk Assessment

Credit card fraud is a crime that is defined as the use of credit cards or credit card information without the knowledge of the credit card's owner. Card issuers and merchants employ techniques that screen credit card transactions and extract customer behaviour patterns in order to detect fraud⁵⁷. Data mining techniques can be used to extract such patterns²², for example NN can be trained on historical transactions and are able to adjust to new forms of fraud. The authors of⁶⁸ use SOMs in order to create a profile of a typical cardholder's behaviour. If there are new transactions, then the deviation of the new transactions to the learned profile is used in order to detect suspicious transactions. Also³⁷ employs SOMs in order to detect fraud, but in real time without prior knowledge about spending patterns.

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The decisions on whether to or not to issue credit cards to certain applicants can be vital for financial institutions. This is because issuing to many credit cards to defaulters will result in considerable financial losses. Recently credit analysts are looking for alternative algorithms to statistical models⁸¹ that can provide a higher accuracy and NN are considered to be an adequate alternative⁸¹. The authors of⁸⁶ use logistic regression and MLPs in order to create a credit scoring model for lending decisions for agricultural loans in Thailand.

Bankruptcy Prediction

A company is bankrupt if it has serious and continuing losses and its debts are beyond its assets²¹. A reliable prediction of bankruptcy is obviously important to many stakeholders around this company such as banks, customers, creditors, governments etc.

The authors of⁹ adopted data mining techniques and NN for creating a predictor of financial distress using historic financial data (such as financial statements, balance sheets and income statements) of 68 Taiwanese companies. Most approaches of bankruptcy prediction using NN are based on a single NN, however, the authors of⁶⁰ considers the usage of an ensemble of several NN in order to predict financial distress. The authors of⁸² derived a method for assessing the corporate financial health of various companies. For this they use a hybrid of Multivariate Discriminate Analysis (MDA) and a SOM. Financial ratios drawn from balance sheets and the the MDA's Z-scores on these ratios are the input to the SOM. The system correctly estimated the corporate financial health in Serbia for 95% of companies in 2008 and for 94% of companies in 2009.

Marketing

The application of NN in marketing is popular due to their ability to model nonlinear relationships between variables⁸³. Applications of NN in marketing range from Market Segmentation, Market Response Prediction, New Product Launch, Sales Forecasting, Consumer Choice Prediction, etc.⁸³. The authors of⁸⁴ used a NN approach to examine consumer behaviour in choosing a hospital in southern Taiwan. The consumer behaviour was categorised in four different consumer styles. Then a MLP with two hidden layers using backpropagation was trained in order to derive a classification model for recognising different consumer styles. Their model achieved a classification accuracy of 85.1%. The authors of⁸⁵ propose the hierarchical self-organizing segmentation model (HSOS) approach as a system for visualising hierarchical clusters step by step. The system is intended to aid a manager as a visualisation tool for market segmentation. Their experimental evaluation shows that HSOS's visualisation capabilities allow a better understanding of market segmentatoin than traditional hierarchical clustering approaches. NN have also been used for analysing customer preferences towards certain kinds of shops. For example the authors of⁸⁷ have used an MLP in order to explain customers preferences towards traditional and web stores for certain products.

Conclusion

This paper presents the NN as useful technology for data mining tasks. It first highlights how data mining fits into the knowledge discovery process and then, in turn, how NN fit into the data mining process. Subsequently, the use of NN for two very important data mining tasks, namely classification and clustering is described. This is followed by a summary of actual applications of NN for classification and clustering problems in bioinformatics and financial data analysis.

Although formerly criticised for producing incomprehensible models and requiring too extensive learning approaches, NN have been experiencing a renaissance in the recent decades. Their successful use in the data mining field is a result of their ability to produce models and to extract explicit and meaningful rules from these models. Also, because the NN are in their nature highly parallel, the advances in shared memory parallel processor architectures allow to speed up the learning process considerably. However, the most important factor for NN' recent popularity in data mining and data analysis is due to their compelling classification accuracy, especially in domains with noisy datasets, such as in the discussed finance and bioinformatics areas. Speeding up NN using super computers may be unattractive for modest size organisations as they may not have the financial power to afford parallel machines, yet they may still want to process large datasets using NN (using their local area network to execute distributed NN algorithms concurrently). Hence a possible future direction of research for NN in data mining is the development of distributed NN algorithms that can be executed on loosely coupled systems using 'off the shelf' computer workstations.

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Figure captions

Figure 1: The Knowledge Discovery Process, comprising four important steps: data integration and cleaning; data selection and transformation; data mining; and evaluation and interpretation.

Figure 2: (a) A sketch of a biological neuron. The neurons are cells that can be electrically stimulated. Each neuron in the brain can take electrochemical signals as input via its *dendrites* and can process them before sending new signals along the *axon* and via the *dendrites* of the other connected neurons; (b) An artificial neuron (a perceptron) which emulates a biological neuron.

Figure 3: An NN – organised in interconnected neurons. The weighted input is squashed through a transfer function and the output from one layer is the input for the next one, and so on, until the NN output is produced (NN with only one hidden layer is shown).

Figure 4: The Data Mining Step of the KDD process using Neural Networks, comprises four important sub-steps: Prepare data for the NN; Learn the NN; Extract rules from the trained NN and Assess the extracted rules.

Figure 5: Training and testing of a Classifier/Predictor. First the classifier is generated from training data; then the classifier is either applied to new unlabelled data or refined by testing it on labelled test data.

Figure 6: The top half of this figure shows a rule based classifier in the form of a decision tree. The bottom half shows some of the rules, encoded in the tree, in the form of IF-THEN rules.

Figure 7: Cluster Analysis portrayed as a four step process comprising: the extraction and selection of features; application of the clustering technique; validation of the derived clusters; and the interpretation of the clusters.

Figure 8: Self Organising Map with two inputs and 49 outputs. Each output unit will represent one cluster (class) of inputs and is associated with a *weight vector*, which size is the same as the *input vector* size.

Further Reading/Resources

[Please insert any further reading/resources here]

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Article ID	Article title
214	Knowledge discovery
216	Self-organizing maps and data mining
217	Neural based learning classifiers


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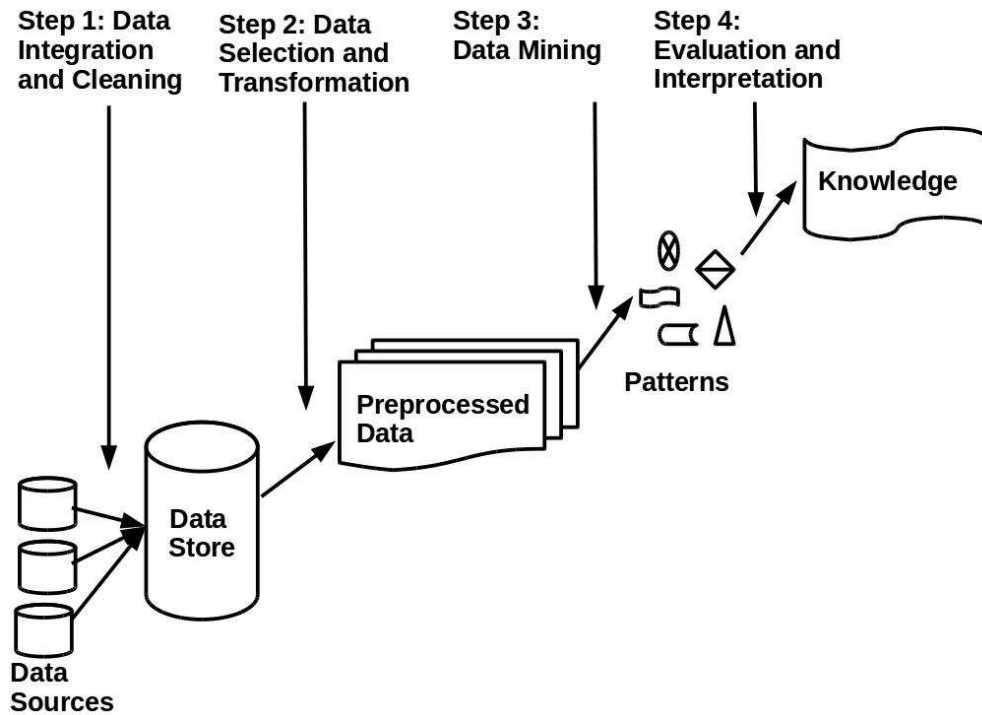
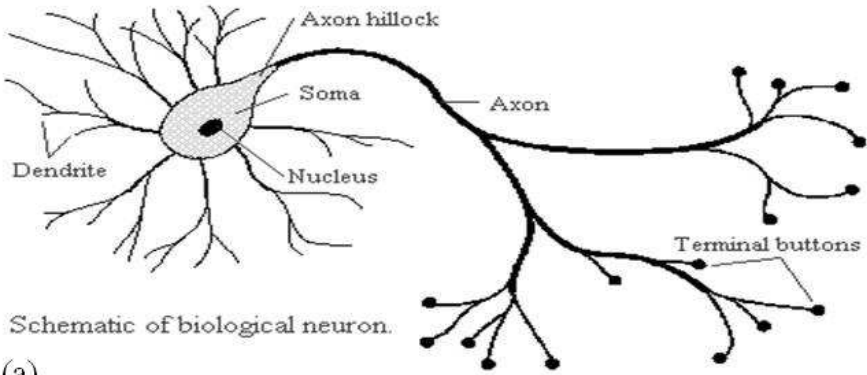
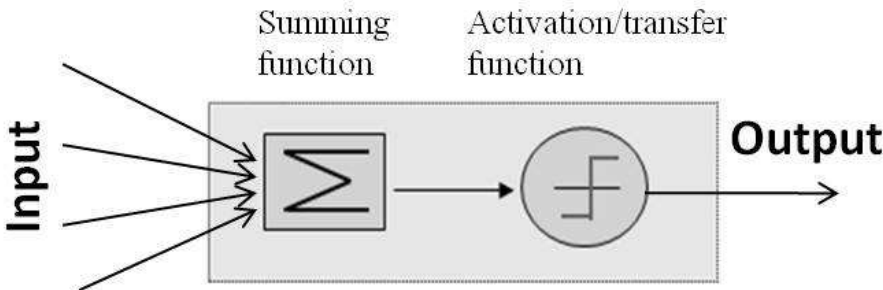


Figure 1: The Knowledge Discovery Process, comprising four important steps: data integration and cleaning; data selection and transformation; data mining; and evaluation and interpretation.



(a)



(b)

Figure 2: (a) A sketch of a biological neuron. The neurons are cells that can be electrically stimulated. Each neuron in the brain can take electrochemical signals as input via its dendrites and can process them before sending new signals along the axon and via the dendrites of the other connected neurons; (b) An artificial neuron (a perceptron) which emulates a biological neuron.

123x96mm (150 x 150 DPI)

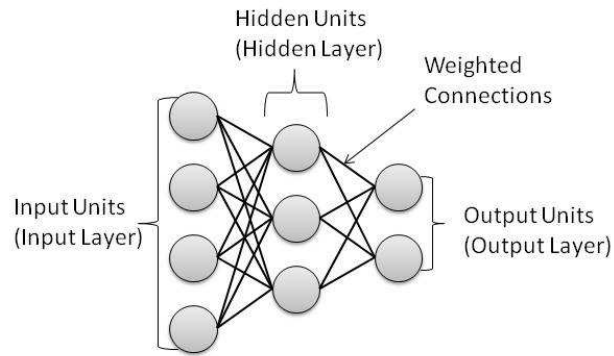


Figure 3: An NN – organised in interconnected neurons. The weighted input is squashed through a transfer function and the output from one layer is the input for the next one, and so on, until the NN output is produced (NN with only one hidden layer is shown).

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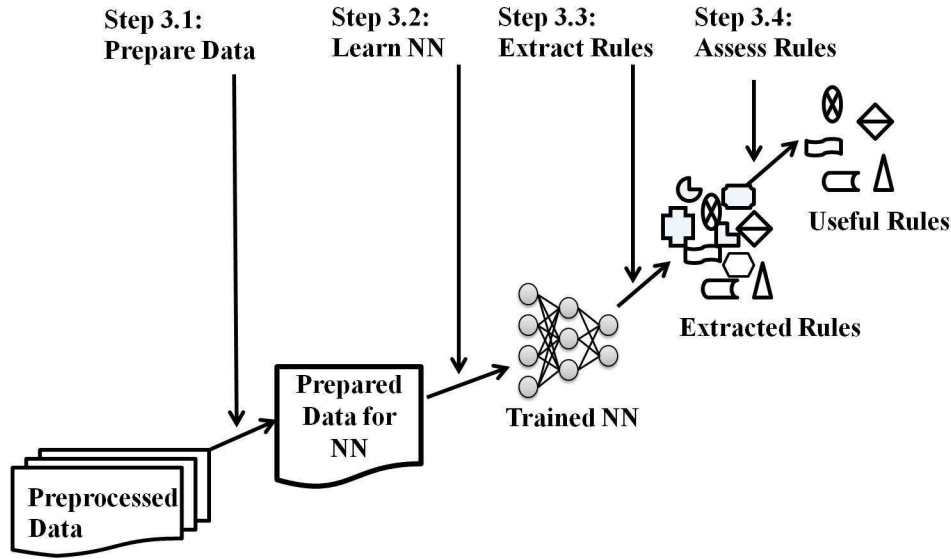


Figure 4: The Data Mining Step of the KDD process using Neural Networks, comprises four important sub-steps: Prepare data for the NN; Learn the NN; Extract rules from the trained NN and Asses the extracted rules.
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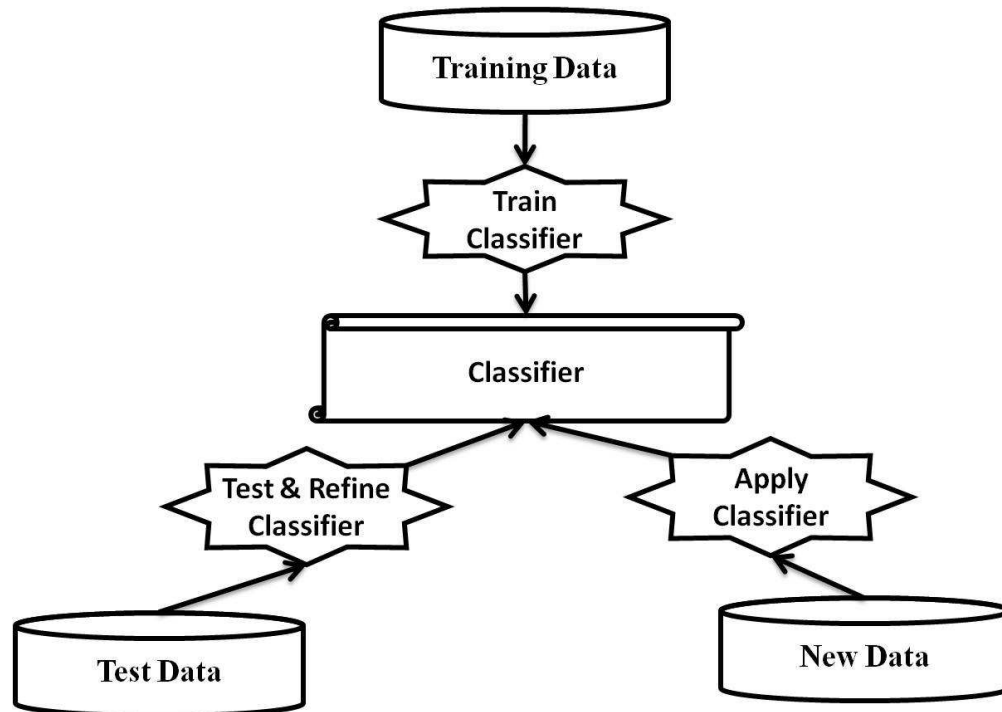


Figure 5: Training and testing of a Classifier/Predictor. First the classifier is generated from training data; then the classifier is either applied to new unlabelled data or refined by testing it on labelled test data.

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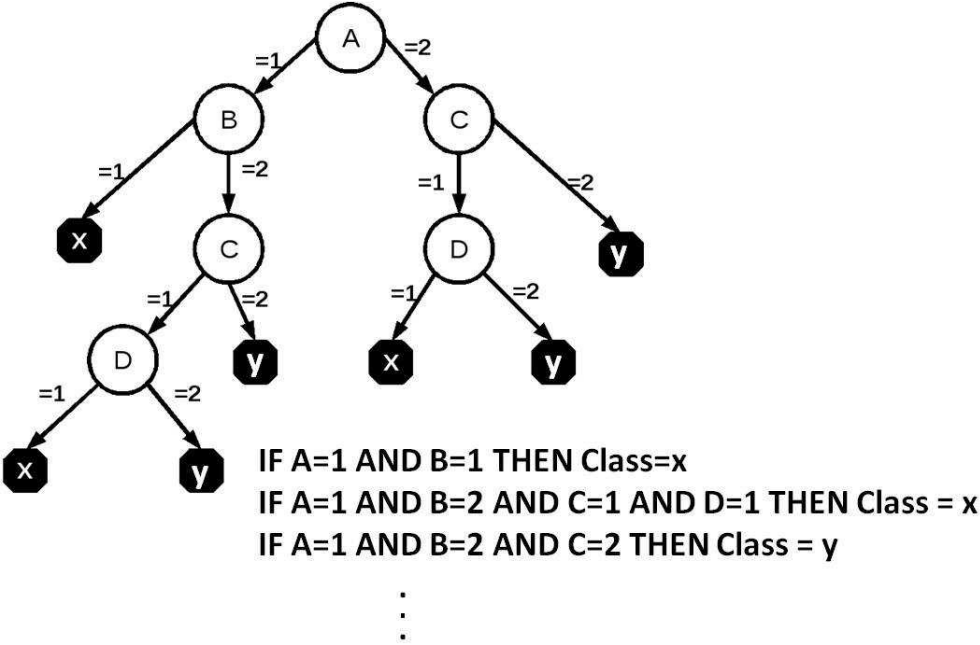


Figure 6: The top half of this figure shows a rule based classifier in the form of a decision tree. The bottom half shows some of the rules, encoded in the tree, in the form of IF-THEN rules.
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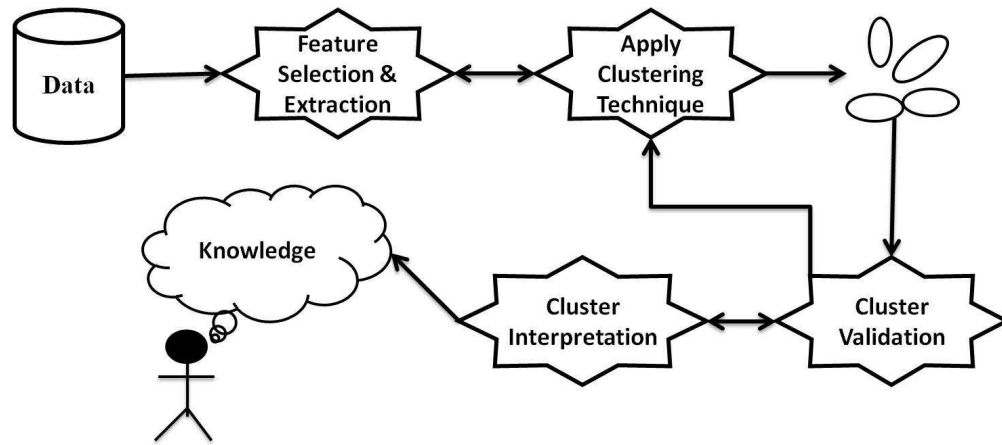


Figure 7: Cluster Analysis portrayed as a four step process comprising: the extraction and selection of features; application of the clustering technique; validation of the derived clusters; and the interpretation of the clusters.
249x110mm (150 x 150 DPI)

Input nodes Weights Output nodes

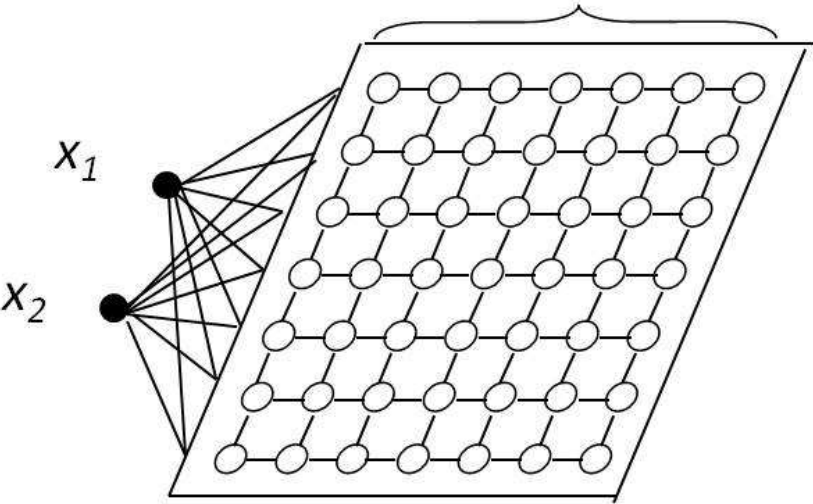


Figure 8: Self Organising Map with two inputs and 49 outputs. Each output unit will represent one cluster (class) of inputs and is associated with a weight vector, which size is the same as the input vector size.
122x77mm (150 x 150 DPI)