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COMPARISON OF ARTIFICIAL NEURAL NETWORK WITH REGRESSION MODELS FOR PREDICTION OF SURVIVAL AFTER SURGERY IN CANCER PATIENTS

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Abstract

Cancer survival prediction in patients who had undergone surgical intervention is an important step in the decision process. The present study investigates the effects of prognostic variables on the breast cancer survival after surgery, over a period of 5-years using feed forward neural network. The neural network was trained and tested using 413 breast cancer patients for the survival prediction with 16 prognostic variables as inputs. The artificial neural network (ANN) proves to be better than regression based models in survival prediction.

1. Introduction

The use of artificial neural networks in biological and medical research has increased enormously in the recent years [11, 17, 20]. ANNs are non linear regressions which have been used in survival prediction and in many bio medical $\overline{2010 \text{ Mathematics Subject Classification: Kindly Provide,}}$

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research and breast cancer diagnosis [7, 21, 23, 25, 26, 30]. Neural network generalizes from the input data to patterns inherent in the data, and uses these patterns to make predictions. Breast cancer is one of the most common cancers in women [29]. The survival prediction after post surgery breast and lung carcinoma using artificial neural network was analyzed by many authors [10, 13, 19, 22]. For analysis of prognostic factors of patients with cancer, models such as Classification and Regression Trees (CART) and the Logistic Regression (LR) are widely used [9, 12, 16]. The challenge of training the neural network using the clinical data is to extract information hidden in the risk factors at the time of forecast. Recently, neural networks are compared with other statistical methods for survival prediction [2, 6, 14, 15, 24]. This study used artificial neural network for finding the accuracy of survival prediction of breast cancer patients who had undergone surgery and the results are compared with logistic regression and CART.

2. Breast Cancer Data

The data consists of patients registered for suspected breast cancer who had undergone surgery from 2000 to 2003. The descriptions of the database are already given elsewhere [27]. Four hundred and thirteen patients who followed 5 years treatment after breast cancer surgery were considered for this study. The covariates comprise of 16 disease and demographic variables which include age of the patient, age at menarche, age at marriage, number of children, breast feed, age at first child birth, family history, age at menopause, tumor size, nuclear grade, tumor stage, chemotherapy, radiotherapy, hormone therapy, the ratio of the number of positive lymph nodes affected to the total number of nodes examined and post tumor node. Staging index is with 3 grades according to the severity of the tumor, on admission. Some of the prognostic variables are binary. The baseline demographic and disease characteristics of the patients for the three stages are given in Tables 1 and 2.

Variable	Stage II n(143)	Stage III n(251)	Stage IV n(19)	All n(413)
AGE	49.3 (12.5)	48.1 (10.6)	48.8 (11.3)	48.5 (11.3)
AMA	19.7 (5.5)	19.6 (4.9)	19.2 (3.4)	19.6 (5.1)
FCB	21.1 (6.7)	20.5 (7.7)	19.6 (9.5)	20.7 (7.5)
NOC	2.4 (1.1)	2.4 (1.5)	1.6 (1.2)	2.36 (1.4)
TSZ	4.1 (1.6)	7.5 (2.7)	6.8 (1.6)	6.3 (2.8)
RAT	.10 (.17)	.23 (.27)	.26 (.24)	.18 (.25)

Table 1. Mean (SD) of covariates according to stage

Variable		Stage II n(143)	Stage III n(251)	Stage IV n(19)	All n(413)
		%	%	%	%
AMR (<13)		14.0	9.6	5.3	10.9
AMO (>54)		5.0	4.0	0	4.1
	Ι	24.5	16.3	26.3	19.6
NGR	II	39.2	31.5	47.4	34.9
	III	36.4	52.2	26.3	45.5
FAH (yes)		7.0	7.6	10.5	7.5
BFD (yes)		91.6	85.7	68.4	87.0
CT (yes)		92.3	53.4	89.5	68.5
RT (yes)		92.3	55.0	94.7	69.7
HT (yes)		81.8	32.3	68.4	51.1
PTN (yes)		68.5	76.1	89.5	74.1

Table 2. Distribution of categorical covariates according to stage

The average age and the age at marriage (AMA) were 48.5 and 19.6 years, respectively. The mean tumor size (TSZ) of the breast cancer patients was 6.3cm and the ratio of the positive number lymph nodes was .184. From Table 2, we see that 11% of the patients had the age at menarche (AMR) less than 13 years and 4% of patients have the menopause age (AMO) greater than 54. About 8% of patients have their family members affected by breast caner and 87% of patients had done the breast feeding (BFD). The tumor was present in 74% of patients even after the surgery (PTN).

3. Regression Models

Logistic regression. Regression methods are essential to describe the casual relationship between a response variable and predictor variables. Logistic regression analysis extends the technique of multiple regression analysis to situations in which the outcome variables are categorical. The binary logistic model was used to identify the significant prognostic variables. The goal of the logistic regression analysis is to find the best fitting and the most parsimonious, yet biologically reasonable model to describe the relationship between an outcome variable and a set of independent variables.

The logistic regression model indirectly models the response variable based on probabilities associated with the values of *Y*. The model is given by

$$\log\left(\frac{p}{1-p}\right) = x_0 + \sum_{i=1}^n a_i x_i,\tag{1}$$

where p = p(Y = 1/X). Logistic regression identifies five variables namely, age at marriage, number of children (NOC), tumor size, nuclear grade (NGR) and lymph node ratio (RAT) are the most significant variables for survival beyond 5 years. Hormone therapy (HT) was more effective in treating the breast cancer patients after surgery. The logistic regression resulted in about 71% and 74% correct prediction for survival after 4 and 5 years.

CART. The classification and regression tree methodology is one of the most commonly used techniques in medical applications. Binary trees give an interesting and often illuminating way of looking data in classification and regression problems. A CART is a flexible nonparametric tool for classification and regression problem. The binary tree structured classifiers are constructed by repeated splits of subsets of prognostic variables into descendant subsets beginning with 5th year status [5]. The terminal subsets form a partition of tumor size. Each terminal subset is designed by a class label. There may be two or more terminal subsets with the same class label. The entire construction of tree is revolved around three elements: (i) selection of splits (ii) declare a node as terminal or not and (iii) assignment of terminal node to a class. Figure 1 gives the CART diagram for the breast cancer data. Figure 2 represents the relative importance of covariates.



Figure 1. Structure generated by CART at the end of 5th year.

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The tree had an initial split on the survival status at the end of the 5th year and 4 terminal subgroups were formed. The variables determining the structure of the tree included were all the breast cancer patient's undergone surgery at the end of the 5th year, the ratio of the number of positive lymph nodes affected to the total number of nodes examined and the tumor size.



Figure 2. Normalized relative importance of covariates.

The tumor size and the ratio of lymph nodes had 100% and 79.7% as the normalized importance at the 4th year. Number of children and family history (FAH) were the least important factors with 3.1% and .2%, respectively. In the 5th year, the ratio of lymph nodes and tumor size had 100% and 89.3% as the normalized importance. Hormone therapy was significant in treating the patients.

4. Artificial Neural Network

A multilayer perceptron (MLP) is a feed forward neural network (FFNN) with at least one hidden layer. This class of neural networks is used in the most of the applications in medicine and other applications [1, 4, 8, 18]. Multilayer perceptron consists of multiple layers of computational units interconnected in a feed forward manner. In MLP the weighted sum of the inputs and bias term are passed to activation level through a transfer function to produce the output. Multilayer perceptron network uses supervised learning technique and back propagation as error correction method. By using the back propagation algorithm the weights are adjusted until the error becomes negligibly very small. The multilayer perceptron neural network used in this application is shown in Figure 3. The output layer had one output node and one hidden with 9 nodes. The choice of the number of hidden layers, hidden nodes and type of activation function plays an important role in model construction [3, 28].



Figure 3. Survival prediction using FFNN.

A multiplayer perceptron using sigmoid function was constructed as shown in Figure 3. To minimize the risk of over fitting and to test the networks generalizing ability, the dataset was divided into training set (70%) and testing set (30%). The training data were used to train the model and the test data were used to test the measure the performance of the network. The output was calculated using

$$y_k = \phi \left(\sum_{h=1}^H w_{hk} \phi \left(\sum_{i=1}^l w_{ik} x_i \right) \right), \tag{2}$$

where ϕ is the sigmoid function given by

$$\phi(x) = \frac{1}{1 + e^{-\lambda x}} \tag{3}$$

whose output lies between 0 and 1, and λ is known as the slope parameter. In (2), x_i 's are input values, w_{ih} are the weights from input to the hidden layer. The error is calculated using

$$E(w) = \frac{1}{2} [t_k - y_k]^2, \qquad (4)$$

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where t_k and y_k are target and the output values. The neural network software NeuNet Pro was used to fit the ANN model.

The survival prediction accuracy is given in Tables 3 and 4. The comparative survival prediction for 4th year and 5th year status for the regression models and the neural network is given in Table 5.

Actual/Predicted	Dead	Alive	Total	Prediction Error
				(%)
Dead	66	10	76	13.2
Alive	03	59	62	4.8
Total	69	69	138	9.4
Actual Error (%)	4.4	14.5	9.4	

Table 3. ANN accuracy measures (4th year)

Actual/Predicted	Dead	Alive	Total	Prediction Error
				(%)
Dead	74	06	80	7.5
Alive	05	52	57	8.8
Total	79	58	137	8.0
Actual Error (%)	6.3	10.3	8.0	

 Table 4. ANN accuracy measures (5th year)

Table 5. Comparative survival prediction of Regression Models and NN

Model/status	LR	CART	NNW
4th year	70.9	67.6	90.6
5th year	73.6	71.2	92.0

5. Discussion

This work attempts to compare the predictive accuracy of the neural network

with the regression based approaches. The results show that the ANN model predicts survival after the surgery better than the regression models. The regression model identified age at marriage, number of children, tumor stage, tumor size, nuclear grade and the ratio of lymph nodes as the most significant factors for survival. Hormone therapy was the significant factor in treating the patients. Because of the clinical complexity and pathologic heterogeneity of cancer, correct identification of patients with active disease is unlikely to depend on the presence or absence of a single defining feature. In our study the sensitivity and the specificity for the 4th year survival status are 93% and 81% and for the 5th year are 97% and 78%, respectively. Burke et al. concluded that neural networks are more accurate in predicting the breast cancer than LR and CART models for 5th year survival. Even our findings are similar to [7, 9].

Neural networks have the ability to approximate predictive output to any desirable degree of accuracy when provided with enough running time. This could result in over fitting, particularly when there is an attempt to increase the processing power of the network by adding a large number of hidden neurons. In this case, the network will end up learning not only the training set but also the noise in the data, which leads to poor generalization. Until the model is tested on a different population set, the study can be viewed only as the first attempt in the use of connectionist models in the predictions of breast cancer survival. Our study has several implications regarding the clinical application of artificial neural networks as a diagnostic tool for breast cancer. The use of the neural network could provide physicians and health-care workers with a simple and fast tool with which to assess the survival after surgery for breast cancer. Clinically prognosis is an important indicator to determine the therapies given to the breast cancer patients after surgery. The genetic algorithm can be used along with ANN to improve classification and prediction. Fuzzy neural networks are the other choice when the cancer data are imprecise in nature.

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