Estimation of Pan Evaporation Coefficient using Neuro-Genetic Approach

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Abstract: Problem statement: The pan evaporation coefficient (Kp) is used to convert pan Evaporation (Ep) to reference Evapotranspiration (ETo) due to its simplicity and suitability for locations with limited availability of meteorological data. Approach: This study presents the use of neuro-genetic approach for estimating Kp for Class A pan and Colorado Sunken pan under green and dry fetch conditions. Results: Representative values were used to represent the category data, i.e., wind run and relative humidity. It was found that the genetic algorithm helped automatically search for the optimal structure of the back-propagation network, replacing the very tedious trial and error approach. Conclusion: A comparative analysis showed that the neural-genetic approach fairly outperformed previous proposed Kp equations for both green and dry fetch conditions.

Key words: Informatics technology, statistical analysis, pan evaporation, colorado sunken, comparative analysis, indicator regression

INTRODUCTION

Hydrological extreme events are typically defined as floods and droughts. Floods are associated with high rainfall and may cause dam break events (Tingsanchali and Chinnarasri, 2001) while drought is associated with lack of precipitation and high evaporation. For modeling purpose and decision making, it is therefore a need for a study to estimate hydrological data for areas with lack of data and different climatic conditions (Hosseini et al., 2011).

Pan Evaporation (Ep) has become a widespread method for estimating reference Evapotranspiration (ETo) due to its simplicity, low cost, ease of data interpretation and application and suitability for locations with limited availability of meteorological data (Phene and Campbell, 1975; Trajkovic, 2009). To convert Ep to ETo, a pan coefficient (Kp) is necessary. Although the FAO-24 Kp Table 1 provides Kp values, several Kp equations have been developed for estimating Kp values (Cuenca, 1989; Snyder, 1992; Allen, 1998; Raghuwanshi and Wallender, 1998; Abdel-Wahed and Snyder, 2008). Those equations were developed for two types of pan, i.e., Class A and Colorado sunken pans and for two conditions, i.e., a pan placed in a short green cropped area and a pan placed in a dry fallow area.

Several Kp equations have been suggested based on the FAO-24 Kp tables using linear, nonlinear and indicator regression techniques or combinations thereof. There are a few regression equations for predicting the Kp values for a FAO Class A pan placed in a short green cropped area based on the FAO-24 Kp Table (Frevert et al., 1983; Cuenca, 1989; Snyder, 1992; Raghuwanshi and Wallender, 1998).

Cuenca (1989) modified Kp equation as proposed by Frevert et al. (1983) by rounding off the coefficients of the equation. Snyder (1992) used the representative values to represent the category data of wind run and relative humidity and applied a least-squares regression approach for predicting Kp values.

To develop a Kp equation for Class A pan placed in short green cropped areas, Raghuwanshi and Wallender (1998) applied the indicator regression technique, which is a widely accepted approach for developing a relationship between categorical and quantitative data.

Allen (1998) and Abdel-Wahed and Snyder (2008) proposed Kp equations for an FAO Class A pan placed in a dry fallow area.
The main function of this model is based on ANNs while a GA helps to automatically find a suitable structure for the network, i.e., the decision variables are the number of hidden layers, the number of nodes in each hidden layer and the transfer functions used. A neuro-genetic process includes two sub-processes: A Neural Network (NN) process (an inner iterative loop) and a genetic algorithm process (an outer iterative loop). Detailed information for neuro-genetic processes can be seen in Fig. 1.

A typical Back Propagation (BP) neural network structure, which includes one input layer, one or more hidden layers and one output layer, is shown in Fig. 2. Each layer is made up of several neurons and the layers are interconnected by sets of weights. The neurons in the input layer receive input directly from the input variables. The neurons in the hidden and output layers receive input from the interconnections. Neurons operate on the input and transform it to produce an analogue output. The transformation is performed in two stages. First, the input from each neuron is multiplied by weights and a weighted sum is taken. Next, an activation function such as the sigmoid function converts such a weighted sum to be the output of each neuron, which becomes the input to neurons of the succeeding layer.

The training of a BP neural network involves two stages (Kumar et al., 2002). In the first stage (forward pass), the input signals propagate from the network input to the output. The calculation of the output is carried out, layer by layer, in the forward direction. The output of one layer is the input to the following layer.

### MATERIALS AND METHODS

The neuro-genetic approach is a hybrid model between Artificial Neural Networks (ANNs) and Genetic Algorithms (GAs).
In the second stage (backward pass), the calculated error signals propagate backward through the network, where they are used to adjust the weights of the connections in order to improve a predefined performance measure. Initially, since the weights to the interconnections are randomly generated, the difference between the predicted and desired output values can be large. Learning therefore involves iteratively adjusting the connection weights to minimize these differences.

There are two parameters of concern for a BP neural network: the learning rate ($\alpha$) and the momentum rate ($\beta$). The learning rate controls the incremental change in the interconnection weights during iterative training as a percentage of the difference between the desired or target output and the NN computed output. Thus, a high learning rate generally results in a larger weight change and faster learning. The momentum rate is a means to increase the rate of learning and avoid the possibility of getting trapped in local optima. The momentum rate is a multiplication factor to the change in the previous interconnection weights (Sivapragasam and Muttil, 2005).

A Genetic Algorithm (GA) is a stochastic optimization model based on Darwin’s concept of natural selection. GA is a robust method for searching for the optimal solution of complex problem, although it may not always necessarily obtain the best possible solution (Goldberg, 1989). In a GA, the population of strings is used to represent possible problem solutions. Each string consists of a number of blocks, which represent the individual variables of the problem. The variables represented in the string can be processed in an evaluation function, or fitness function, which is in effect the objective function.

GA consists of three basic operations (selection, crossover and mutation) involved in manipulating strings and moving to a new generation. The selection operator is that through which strings are selected for inclusion in the reproduction process and for participation in the next generation. The fittest strings have the highest probability of being used in reproduction. The crossover operator permits the exchange of genes between pairs of chromosomes in a population. This operator offers the possibility of good genetic material from different individual strings being combined to create an even fitter individual. The mutation operator permits new genetic material to be introduced to a population. As shown in Fig. 1, replacement is used to refill the population that is dropped due to poor chromosomes during the selection process.

RESULTS

To estimate $K_p$ values based on the neuro-genetic approach, the data sets from Table 1 were used. This table shows $K_p$ values based on the FAO-24 $K_p$ table for Class A pans under green fetch conditions (case I), Class A pans under dry fetch conditions (case II), Colorado sunken pans under green fetch conditions (case III) and Colorado sunken under dry fetch conditions (case IV).

The $K_p$ values depends on the fetch distance ($F$), which is quantitative data and two qualitative (or categorical) data, i.e., the daily mean Relative Humidity (RH) and daily mean wind speed (U). Hence, the structure of the neural networks comprised three nodes of input layer, i.e., $F$, RH and U and one node of output layer, i.e., $K_p$ data.

Back-Propagation (BP) was selected as the supervised learning method and a Genetic Algorithm (GA) helped in finding a suitable network, i.e., the number of hidden layers, the number of nodes in each hidden layer and the transfer functions to be used for each node.
Table 2: Comparison of statistical indices in estimating \(K_p\) values using different methods

| Methods                                      | \(R^2\) | RMSQ  | MAE (%) | MARE (%) | MXARE (%) | DEV (%) | NE > 2% |
|----------------------------------------------|---------|-------|---------|----------|-----------|---------|---------|
| Class A pan under green fetch condition (case I) |         |       |         |          |           |         |         |
| Cuenca (1989)                                | 0.9601  | 0.0327| 2.69    | 4.41     | 13.25     | 3.31    | 27      |
| Snyder (1992)                                | 0.9745  | 0.0262| 2.12    | 3.29     | 9.84      | 2.48    | 21      |
| Allen (1998)                                 | 0.9822  | 0.0235| 2.72    | 4.07     | 11.46     | 2.85    | 26      |
| Raghuvanshi and Wallender (1998)             | 0.9796  | 0.0235| 1.88    | 2.97     | 8.46      | 2.22    | 21      |
| Neuro-genetic (present study)                | 0.9901  | 0.0167| 1.40    | 2.29     | 6.18      | 1.63    | 15      |
| Class A pan under dry fetch condition (case II) |         |       |         |          |           |         |         |
| Allen (1998)                                 | 0.9849  | 0.0383| 3.11    | 5.02     | 12.90     | 3.20    | 32      |
| Abdel-Wahed and Snyder (2008)                | 0.9868  | 0.0194| 1.59    | 2.87     | 10.00     | 2.18    | 16      |
| Neuro-Genetic (Present study)                | 0.9877  | 0.0188| 1.45    | 2.42     | 8.10      | 2.05    | 17      |
| Colorado sunken pan under green fetch condition (case III) |         |       |         |          |           |         |         |
| Allen (1998)                                 | 0.9840  | 0.0545| 4.40    | 5.20     | 11.59     | 3.01    | 27      |
| Neuro-Genetic (Present study)                | 0.9921  | 0.0201| 1.72    | 2.27     | 7.64      | 1.62    | 13      |
| Colorado Sunken pan under dry fetch condition (case IV) |         |       |         |          |           |         |         |
| Allen (1998)                                 | 0.9851  | 0.0425| 3.32    | 4.48     | 11.74     | 3.12    | 29      |
| Neuro-Genetic (present study)                | 0.9890  | 0.0246| 1.84    | 2.62     | 7.77      | 2.31    | 20      |

NeuroGenetic Optimizer (demo version) was selected as the learning tool. The default parameters of neural networks and genetic algorithm as provided by the NeuroGenetic Optimizer software package were selected in this study as follows. The input and output data scaling were between -1 and 0.1 and 0.9, respectively. The multiple hidden layers with maximum of 2 hidden layers and the maximum number of nodes in each hidden layer of 128 were selected. Three transfer functions (Tan-Hyperbolic, Logistic and Linear transfer functions) were used. The initial weights were randomly assigned between -0.3-0.3. Learning rates of between 0.1-0.4 for the first and second hidden layers and between 0.1-0.2 for the output layer were set. A momentum between 0.1-0.2 was set.

A population size (the number of chromosomes in the population) of 200 was used. The percent selection method with 50% selection was chosen. That is, the networks with fitness greater than average will be selected to survive while those less than average will be dumped into the bit bucket. Since this system uses a selection technique that drops poor chromosomes, the population needs to be refilled every generation. In this study, a cloning technique, whereby the survivors of the selection process are cloned to refill the population, was used. The tail swap mating method (also known as crossover) was selected. Hence, the system picks a cut point and exchanges genetic material between the cut point and the end of the string of the parents, essentially swapping tails. A random exchange mutation method with a probability of 0.25 was selected. With this method, two points in a given chromosome string are randomly selected and exchanged their values with a probability of 0.25.

Seven statistical indices, including the determination coefficient \((R^2)\), root Mean Square Error (RMSE), Mean Absolute Error (MAE), mean absolute relative error (MARE), maximum absolute relative error (MXARE), standard deviation of absolute relative error (DEV) and the number of samples with an error greater than 2% (NE> 2%) were used to evaluate the performance in estimating \(K_p\) values. The \(R^2\) measures the degree to which two variables are linearly related and should optimally be one. The RMSE is a measure of the residual standard deviation and should be as small as possible (optimally 0). The MAE, MARE and MXARE measure the difference between actual and estimated \(K_p\) values and should be as small as possible (optimally 0).

**DISCUSSION**

The statistical indices in estimating \(K_p\) values using neuro-genetic and previous equations for all study cases are presented in Table 2. For case I, the neuro-genetic approach gave a higher performance \((R^2 = 0.9901, \text{RMSQ} = 0.0167, \text{MAE} (%) = 1.40, \text{MARE} (%) = 2.29, \text{MXARE} (%) = 6.18, \text{DEV} (%) = 1.63)\) and \(K_p\) values than other methods. For case II, although the values of \(R^2 (0.9877)\) and \(\text{NE}> 2\% (17)\) of neuro-genetic approach were lower than those of the equation proposed by Abdel-Wahed and Snyder (2008) other statistical indices \((\text{RMSQ} = 0.0188, \text{MAE} (%) = 1.45, \text{MARE} (%) = 2.42, \text{MXARE} (%) = 8.10)\) were better.

For cases III and IV, the neuro-genetic approach obviously outperformed the equations proposed by Allen (1998). It gave \(R^2 = 0.9921, \text{RMSQ} = 0.0201, \text{MAE} (%) = 1.72, \text{MARE} (%) = 2.27, \text{MXARE} (%) = 8.10\).
7.64, DEV (%) = 1.62 and NE> 2% = 13 for case III and R² = 0.9890, RMSQ = 0.0246, MAE (%) = 1.84, MARE (%) = 2.62, MXARE (%) = 7.77, DEV (%) = 2.31 and NE> 2% = 20 for case IV.

The structures of BP neural networks (# input layer nodes-1st hidden layer nodes-2nd hidden layer nodes- # output layer nodes-the hidden layers being grouped by type of transfer function) obtained by the GA for cases I-IV, respectively, were (3-17 nodes with Lo, 5 with T, 5 with Li-50 with Lo, 28 with T, 46 with Li-1), (3-46 with Lo, 4 with T, 24 with Li- 95 with Lo, 27 with T, 6 with Li-1), (3-5 with Lo, 6 with T, 5 with Li-15 with Lo, 2 with T, 2 with Li-1) and (3-15 with Lo, 44 with Li-47 with Lo, 24 with T, 3 with Li-1), respectively. Lo, T and Li stand for Logistic, Tan-Hyperbolic and Linear transfer functions, respectively and the GA always selected Tan-hyperbolic for the output node.

**CONCLUSION**

In this study, a Neuro-genetic approach was successfully applied to estimating Kp values for the cases of Class A pans under green fetch conditions, Class A pans under dry fetch conditions, Colorado sunken pans under green fetch conditions and Colorado sunken pans under dry fetch conditions. A comparative analysis showed that this approach fairly outperformed the previous proposed Kp equations in estimating Kp values.

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