

# **Bayesian Artificial Neural Networks in Water Resources Engineering**

by Greer B. Kingston

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# Abstract

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A new Bayesian framework for training and selecting the complexity of artificial neural networks (ANNs) is developed in this thesis, based on Markov chain Monte Carlo (MCMC) techniques. The primary motivation of the research presented is the incorporation of uncertainty into ANNs used for water resources modelling, with emphasis placed on obtaining accurate results, while maintaining simplicity of implementation, which is considered to be of utmost importance for adoption of the framework by practitioners in this field. By applying the Bayesian framework to a number of synthetic and real-world case studies and by comparison with a state-of-the-art ANN development approach, it is shown throughout this thesis how the Bayesian approach can be used to address the three most significant issues facing the wider acceptance of ANNs in this field; namely generalisability, interpretability and uncertainty. The state-of-the-art approach is devised through reviewing and, where necessary, improving current best practice deterministic ANN development methods, leading to the recommended use of the global SCE-UA optimisation algorithm, which has not been used before for ANN training, and the development of a modified connection weight approach for extracting knowledge from trained ANNs. The real-world case studies used in this research, which involve salinity forecasting in the River Murray at Murray Bridge, South Australia, and the forecasting of cyanobacteria (*Anabaena* spp.) in the River Murray at Morgan, South Australia, are used to demonstrate the practical value of the Bayesian framework, particularly when extrapolation is required and when the available data are of poor quality. These issues lead to poor model performance when deterministic ANN development methods are applied, yet as the generated predictions are deterministic, there is no direct way of assessing their quality. Application of the proposed Bayesian framework leads to better average performance of the ANN models developed, since a minimal ANN structure is selected and a more generalised input-output mapping is obtained. More importantly, prediction limits are provided which quantify the uncertainty in the predictions and enable management and design decisions to be made based on a known level of confidence.

*Abstract*

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# **Statement of Originality**

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*I Greer B. Kingston hereby declare that this work contains no material that has been accepted for the award of any other degree or diploma in any university or other tertiary institution. To the best of my knowledge and belief, it contains no material previously published or written by any other person, except where due reference is made in the text.*

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# List of Publications

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Kingston, G. B., H. R. Maier, and M. F. Lambert (2005), A probabilistic method to assist knowledge extraction from artificial neural networks used for hydrological prediction, *Mathematical and Computer Modelling*, *In press*, doi:10.1016/j.mcm.2006.01.008.

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# Nomenclature & Abbreviations

---

| Symbol                                   | Description  |
|--|--|
| <i>General</i>                           |  |
| $\Re^d$                                  | $d$ -dimensional set of all real numbers   |
| $\Theta$                                 | search space; $\Theta \in \Re^d$   |
| $\mathbf{I}$                             | identity matrix  |
| $\mathbf{H}$                             | Hessian matrix   |
| $\mu$                                    | scalar mean  |
| $\sigma^2$                               | variance ( $\sigma$ is the standard deviation)   |
| $\Sigma$                                 | covariance matrix  |
| $\lambda$                                | signal-to-noise ratio  |
| $r^2$                                    | coefficient of determination   |
| <i>Probability Distribution Notation</i> |  |
| $N(\mu, \sigma^2)$                       | Normal distribution with mean $\mu$ and variance $\sigma^2$                            |
| $U(a, b)$                                | Uniform distribution with boundaries $a, b$ , where $b > a$                            |
| $\text{Inv-}\chi^2(\nu, S)$              | Inverse chi-square distribution with $\nu$ degrees of freedom<br>and scale $S$         |
| <i>ANN and Modelling Notation</i>        |  |
| $f(\cdot)$                               | function modelled by an ANN  |
| $K$                                      | number of inputs   |
| $J$                                      | number of hidden layer nodes   |
| $M$                                      | number of outputs  |
| $I_k$                                    | $k$ th input node  |
| $H_j$                                    | $j$ th hidden node   |
| $O_m$                                    | $m$ th output node   |
| $w_i$                                    | $i$ th “true” weight   |
| $\hat{w}_i$                              | $i$ th estimated weight  |
| $\mathbf{w}$                             | “true” vector of connection and bias weights $\equiv (w_1, \dots, w_d)$                |
| $\hat{\mathbf{w}}$                       | estimated vector of connection and bias weights $\equiv (\hat{w}_1, \dots, \hat{w}_d)$ |

*continued on next page*

| Symbol             | Description                                       |
|--------------------|---|
| $d$                | dimension of weight vector                        |
| $g(\cdot)$         | activation function                               |
| $zin_j$            | summed input into hidden node $j$                 |
| $z_j$              | output from hidden node $j \equiv g(zin_j)$       |
| $\hat{y}in$        | summed input into output node                     |
| $\hat{y}_m$        | output from output node $m \equiv g(\hat{y}in_m)$ |
| $E_y$              | error, or objective, function                     |
| $E_w$              | penalty term used to regularise weights           |
| $\nabla E$         | gradient of error function                        |
| $\epsilon$         | model residuals                                   |
| $\sigma_y^2$       | scale (variance) of model residuals               |
| $\sigma_w^2$       | scale (variance) of weights                       |
| $\hat{\sigma}_y^2$ | scale of model residuals at optimum of $E_y$      |
| $\hat{\sigma}_w^2$ | scale of weights at optimum of $E_y$              |

*Data Notation*

|  |   |
|--|---|
| $N$  | number of samples in the data set   |
| $y$  | general term for observed target data   |
| $y_i$  | $i$ th observed target data scalar  |
| $\mathbf{y}$                                   | general term for set of scalar target data $\equiv (y_1, \dots, y_N)$                   |
| $\mathbf{y}^M$                                 | general term for observed target data vector $\equiv (y_1, \dots, y_M)$                 |
| $\mathbf{y}_i^M$ or $\mathbf{y}_i$             | $i$ th target vector $\equiv (y_{1,i}, \dots, y_{M,i})$                                 |
| $\mathbf{y}_m$                                 | $m$ th target variable $\equiv (y_{m,1}, \dots, y_{m,N})$                               |
| $\mathbf{Y}$                                   | general term for set of target vectors $\equiv (\mathbf{y}_1^M, \dots, \mathbf{y}_N^M)$ |
| $\hat{y}_i$                                    | $i$ th predicted data scalar  |
| $\hat{\mathbf{y}}$                             | general term for set of scalar predicted data $\equiv (\hat{y}_1, \dots, \hat{y}_N)$    |
| $\hat{\mathbf{y}}^M$                           | general term for predicted data vector $\equiv (\hat{y}_1, \dots, \hat{y}_M)$           |
| $\hat{\mathbf{y}}_i^M$ or $\hat{\mathbf{y}}_i$ | $i$ th predicted output vector $\equiv (\hat{y}_{1,i}, \dots, \hat{y}_{M,i})$           |
| $\hat{\mathbf{y}}_m$                           | $m$ th predicted output variable $\equiv (\hat{y}_{m,1}, \dots, \hat{y}_{m,N})$         |
| $\mathbf{x}$                                   | general term for input variable $\equiv (x_1, \dots, x_N)$                              |
| $\mathbf{x}^K$                                 | general term for input vector $\equiv (x_1, \dots, x_K)$                                |
| $\mathbf{x}_i^K$ or $\mathbf{x}_i$             | $i$ th input vector $\equiv (x_{1,i}, \dots, x_{K,i})$                                  |
| $\mathbf{x}_k$                                 | $k$ th input variable $\equiv (x_{k,1}, \dots, x_{k,N})$                                |
| $\mathbf{X}$                                   | general term for set of input vectors $\equiv (\mathbf{x}_1^K, \dots, \mathbf{x}_N^K)$  |
| $\mathcal{D}$                                  | set of data pairs $\equiv [(\mathbf{x}_1, y_1), \dots, (\mathbf{x}_N, y_N)]$            |

*continued on next page*

| Symbol                                     | Description  |
|--|--|
| <i>Probability Notation</i>                |  |
| $\theta$                                   | general term for model parameters  |
| $\mathcal{H}$                              | general term for model structure, including implicit assumptions   |
| $p(\theta)$                                | probability density of model parameters $\theta$ , also known as the prior probability density   |
| $p(\theta, \mathbf{y})$                    | joint probability of the model parameters $\theta$ and the data $\mathbf{y}$   |
| $p(\theta \mathbf{y})$                     | conditional probability of $\theta$ , given the data $\mathbf{y}$ , also known as the posterior probability density of $\theta$                      |
| $p^*(\theta \mathbf{y})$                   | unnormalised posterior density   |
| $p(\mathbf{y} \theta)$                     | conditional probability of the data $\mathbf{y}$ , given the model parameters $\theta$ , also known as the likelihood function                       |
| $L(\theta)$                                | the likelihood function, as above  |
| $p(\mathbf{y} \mathcal{H})$                | conditional probability of the data $\mathbf{y}$ , given the model $\mathcal{H}$ , also known as the marginal likelihood or evidence                 |
| $\hat{p}(\mathbf{y} \mathcal{H})$          | approximate evidence   |
| $p(\mathcal{H} \mathbf{y})$                | conditional probability of the model $\mathcal{H}$ , given the model $\mathbf{y}$ , also known as the posterior probability density of $\mathcal{H}$ |
| <i>Training Notation</i>                   |  |
| <b>Backpropagation (BP)</b>                |  |
| $\gamma$                                   | stepsize of gradient descent   |
| $\mathbf{d}$                               | direction of gradient descent  |
| $\eta$                                     | learning rate  |
| $\phi$                                     | momentum rate  |
| $\Delta \mathbf{w}$                        | weight increment   |
| $\delta$                                   | delta function   |
| $\kappa$                                   | epoch size   |
| <b>Genetic Algorithm (GA)</b>              |  |
| $G$  | population of chromosomes  |
| $s$  | population size  |
| $\rho_{cross}$                             | crossover rate   |
| $\rho_{mut}$                               | mutation rate  |
| $\tau$                                     | mutation stepsize  |
| $gene'$                                    | mutated gene value   |
| <b>Shuffled Complex Evolution (SCE-UA)</b> |  |
| $p$  | number of complexes  |
| $m$  | number of points in a complex  |

continued on next page

| <b>Symbol</b>                          | <b>Description</b>   |
|--|--|
| $s$                                    | population size = $m \times p$                                   |
| $q$                                    | number of points in a subcomplex                                 |
| $\alpha$                               | number of offspring generated by a subcomplex                    |
| $\beta$                                | number of evolution steps taken by each complex                  |
| <b>Markov chain Monte Carlo (MCMC)</b> |  |
| $T(\cdot)$                             | transition distribution  |
| $Q(\cdot)$                             | proposal density   |
| $\alpha(\cdot)$                        | acceptance probability distribution                              |
| $\theta^*$                             | candidate parameter state  |
| $c$                                    | adaptive scaling parameter                                       |
| $T$                                    | temperature used for simulated annealing                         |
| $\varphi$                              | simulated annealing schedule parameter                           |
| $t_0$                                  | number of iterations for which $\Sigma$ is held constant         |
| $t_{\sigma_0^2}$                       | number of iterations for which hyperparameters are held constant |
| $t_b$                                  | number of burn-in iterations                                     |

| <b>Abbreviation</b> | <b>Description</b>                                  |
|---------------------|---|
| AM                  | Adaptive Metropolis                                 |
| AIC                 | Akaike's information criterion                      |
| ANN                 | Artificial neural network                           |
| ARD                 | Automatic relevance determination                   |
| AWQC                | Australian Water Quality Centre                     |
| BF                  | Bayes factor  |
| BIC                 | Bayesian information criterion                      |
| BMS                 | Bayesian model selection                            |
| BP                  | Backpropagation                                     |
| CCE                 | Competitive complex evolution                       |
| CDF                 | Cumulative distribution function                    |
| CE                  | Coefficient of efficiency                           |
| C-J                 | Chib-Jeliazkov                                      |
| DWR                 | South Australian Department for Water Resources     |
| EBMLP               | Evolutionary backpropagation multi-layer perceptron |
| EP                  | Evolutionary programming                            |
| GA                  | Genetic algorithm                                   |
| G-D                 | Gelfand-Dey   |

*continued on next page*

| <b>Abbreviation</b> | <b>Description</b>                                 |
|---------------------|--|
| GLUE                | Generalized likelihood uncertainty estimation      |
| GRNN                | General regression neural network                  |
| HMC                 | Hybrid Monte Carlo                                 |
| L1LF                | Lock 1 lower flow                                  |
| L1LL                | Lock 1 lower river level                           |
| L1UL                | Lock 1 upper river level                           |
| L7F                 | Lock 7 flow  |
| LOL                 | Loxton river level                                 |
| LOS                 | Loxton salinity                                    |
| MAE                 | Mean absolute error                                |
| MAL                 | Mannum river level                                 |
| MAS                 | Mannum salinity                                    |
| MBL                 | Murray Bridge river level                          |
| MBS                 | Murray Bridge salinity                             |
| MCMC                | Markov chain Monte Carlo                           |
| MDB                 | Murray-Darling Basin                               |
| MDBC                | Murray-Darling Basin Commission                    |
| MDBMC               | Murray-Darling Basin Ministerial Council           |
| MI                  | Mutual information                                 |
| MLP                 | Multi-layer perceptron                             |
| MOL                 | Morgan river level                                 |
| MOS                 | Morgan salinity                                    |
| MSE                 | Mean squared error                                 |
| MSRE                | Mean squared relative error                        |
| MT                  | Model tree   |
| NGO                 | NeuroGenetic Optimizer                             |
| OCF                 | Overland Corner flow                               |
| OCL                 | Overland Corner river level                        |
| OCW                 | Overall connection weight                          |
| PC                  | Principle component                                |
| PCA                 | Principle component analysis                       |
| PDF                 | Probability density function                       |
| PMI                 | Partial mutual information                         |
| RI                  | Relative importance                                |
| RMSE                | Root mean squared error                            |
| SCE-UA              | Shuffled complex evolution - University of Arizona |
| SOM                 | Self-organising map                                |

*continued on next page*

| <b>Abbreviation</b> | <b>Description</b>     |
|---------------------|------------------------|
| SSE                 | Sum squared error      |
| SVM                 | Support vector machine |
| WAL                 | Waikerie river level   |
| WAS                 | Waikerie salinity      |