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2000

Online at <https://mpra.ub.uni-muenchen.de/77790/>  
MPRA Paper No. 77790, posted 03 Apr 2017 10:13 UTC

**Evaluating Neural Spatial Interaction  
Modelling by Bootstrapping**

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Paper presented at the 6th World Congress  
of the Regional Science Association International,  
Lugano, Switzerland, May 16-20, 2000

## **Abstract**

This paper exposes problems of the commonly used technique of splitting the available data in neural spatial interaction modelling into training, validation, and test sets that are held fixed and warns about drawing too strong conclusions from such static splits. Using a bootstrapping procedure, we compare the uncertainty in the solution stemming from the data splitting with model specific uncertainties such as parameter initialization. Utilizing the Austrian interregional telecommunication traffic data and the differential evolution method for solving the parameter estimation task for a fixed topology of the network model [ i.e.  $J = 9$ ] this paper illustrates that the variation due to different resamplings is significantly larger than the variation due to different parameter initializations. This result implies that it is important to not over-interpret a model, estimated on one specific static split of the data.

**Keywords:** Neural spatial interaction modelling, model evaluation, bootstrapping, interregional telecommunications

## 1. Introduction

Spatial analysis is currently entering a period of rapid change leading to what is termed *intelligent spatial analysis*, sometimes referred to as *geocomputation*. The driving forces are a combination of huge amounts of digital spatial data from the GIS data revolution, the availability of attractive softcomputing tools, the rapid growth in computational power, and the new emphasis on exploratory data analysis and modelling. *Intelligent* spatial analysis has the following properties. It exhibits computational adaptivity, computational fault tolerance in dealing with incomplete, inaccurate, distorted, missing, noisy and confusing data; speed approaching human-like turnaround and error rates that approximate human performance. The use of the term 'intelligent' is, thus, closer to that in computational intelligence than in artificial intelligence. The distinction between artificial and computational intelligence is important because our semantic descriptions of models and techniques, their properties, and our expectations of their performance should be tempered by the kind of systems we want, and the ones we can build.

Much of the recent interest in intelligent spatial analysis stems from the growing realization of the limitations of conventional spatial analysis tools as vehicles for exploring patterns in data-rich GI (geographic information) and RS (remote sensing) environments and from the consequent hope that these limitations may be overcome by judicious use of computational intelligence technologies such as evolutionary computation and neural network modelling. Neural network models may be viewed as non-linear extensions of conventional statistical models that are applicable to two major domains: *first*, as universal approximators to areas such as spatial regression, spatial interaction, spatial choice and space-time series analysis; and *second*, as pattern recognizers and classifiers to intelligently allow the user to sift through the data, reduce dimensionality, and find patterns of interest in data-rich environments.

Neural spatial interaction models are termed neural in the sense that they are based on neural computational models, inspired by neuroscience. They are more closely related to spatial interaction models of the gravity type, and under commonly met conditions they can be understood as a special class of general feedforward neural network models with a single hidden layer and sigmoidal transfer functions (Fischer 1998). This class of

networks can provide approximations within an arbitrary precision (i.e. it has universal approximation property), as proven by Hornik et al. (1989).

Learning from examples, the problem for which neural networks were designed for to solve, is one of the most important research topics in computational intelligence. A possible way to formalize learning from examples is to assume the existence of a function representing the set of examples and, thus, enabling to generalize. This can be called a *function reconstruction from sparse data* (or in mathematical terms, depending on the required precision, approximation or interpolation problem, respectively). Within this general framework, the main issues of interest are the representational power of a given network model and the procedures for obtaining the optimal network parameters.

It is the objective of this article to evaluate the out-of-sample or forecast performance of neural spatial interaction approximators in a real world application environment. Training a neural spatial interaction model is not hard, but once we have a spatial interaction approximator, how much can we trust the forecast for truly new data? A standard procedure for evaluating the performance is to split the data into one training set (used for parameter estimation, e.g., through gradient descent or the differential evolution method), one validation set (used to determine the stopping point before overfitting occurs and/or used to set additional parameters or hyperparameters, such as the importance given to the amplification of the differential variation in the case of the differential evolution method), and one test set. This procedure has been used for many years in the connectionist community [see, e.g., Weigend et. al. (1990) in general and in neural spatial interaction modelling in particular [see Fischer and Gopal (1994)]. Recent experience has found this approach, along with conclusions drawn from it, to be very sensitive to the specific splitting of the data [see, Fischer and Gopal (1994)]. Thus, usual tests of forecast reliability appear over-optimistic in general.

This paper addresses the problem of evaluating a neural spatial interaction model with a bootstrapping procedure. The approach we present combines the purity of splitting the data into three disjoint sets - as suggested in Fischer and Gopal (1994) - with the power of a resampling procedure. This allows us to get a better statistical picture of forecast variability, including the ability to estimate the effect of the randomness of the splits of the data versus the randomness of initial conditions of the model parameters. We

compare the uncertainties in the solution stemming from the data splitting with neural network specific uncertainties such as parameter initializations. To demonstrate the procedure, we use the Austrian interregional telecommunication traffic data and the differential evolution method, a randomized parallel multipoint search procedure, for solving the parameter estimation task as suggested in Fischer et al. (1999).

The remainder of the paper is as follows. The next section provides a summarized description of single hidden layer neural spatial interaction models followed by a brief characteristic of the training procedure used for determining the model parameters. Section 4 describes the experimental design and the bootstrapping procedure while section 5 presents the empirical results of the study. The final section draws some conclusions.

## 2. Neural Spatial Interaction Predictors

Suppose we are interested in approximating an  $N$ -dimensional spatial interaction function, where  $\mathfrak{F} : \mathfrak{R}^N \rightarrow \mathfrak{R}$ , where  $\mathfrak{R}^N$  as  $N$ -dimensional Euclidean real space is the input space and  $\mathfrak{R}$  as 1-dimensional Euclidean real space is the output space. This function should estimate spatial interaction flows from regions of origin to regions of destination. In practice, only bounded subsets of the space are considered.

The function  $\mathfrak{F}$  is not explicitly known, but given by a finite set of samples  $S = \{z_k = (x_k, y_k), k = 1, \dots, K\}$  generated by a process that is governed by  $\mathfrak{F}$ , that is  $\mathfrak{F}(x_k) = y_k, k = 1, \dots, K$ . The set  $S$  is the set of pairs of input and output vectors. The role of network modelling is to provide a specific form of a continuous function which approximates (or interpolates) set  $S$ . The advantages of neural network modelling have to do with the virtues associated with such specific forms.

To approximate  $\mathfrak{F}$ , we consider the class of neural spatial interaction models  $\Omega$  with one hidden layer,  $N$  input units,  $J$  hidden units and one output unit.  $\Omega$  consists of a composition of transfer functions so that the single output  $y$  of  $\Omega$  is:

$$y(\mathbf{x}, \mathbf{w}) = \Omega(\mathbf{x}, \mathbf{w}) = \psi \left( \sum_{j=0}^J w_j \varphi_j \left( \sum_{n=0}^N w_{jn} x_n \right) \right) \quad (1)$$

Vector  $\mathbf{x} = (x_1, \dots, x_N)$  is the input vector augmented with a bias signal  $x_0$  which can be thought as being generated by a ‘dummy unit’ whose output is clamped at 1. The  $w_{jn}$ 's represent input to hidden connection weights and the  $w_j$ 's hidden to output weights (including the bias). The symbol  $\mathbf{w}$  is a convenient shorthand notation of the  $d = (J(N+1) + J + 1)$  - dimensional vector of all the  $w_{jn}$  and  $w_j$  network weights and biases (i.e. model parameters).  $\varphi_j(\cdot)$  and  $\psi(\cdot)$  are differentiable transfer functions of the hidden units  $j = 1, \dots, J$  and the output unit, respectively. Following Fischer and Gopal (1994), we will consider only the case  $N=3$ , i.e. the input space will be a closed interval of the three-dimensional Euclidean space  $\mathfrak{R}^3$ . The three input units correspond to the independent variables of the classical unconstrained spatial interaction model of the gravity type. They represent measures of origin propulsiveness, destination attractiveness and spatial separation. The output unit corresponds to the dependent variable of the classical model and represents the spatial interaction flows from origin to destination.

Without loss of generality we, moreover, assume the transfer functions  $\varphi_j(\cdot) = \varphi(\cdot)$  for all  $j = 1, \dots, J$ , and equal to the logistic function and,  $\psi(\cdot)$  to be the identity function, and, thus consider the special class  $\Omega_L(\mathbf{x}, \mathbf{w})$  of functions  $\Omega(\mathbf{x}, \mathbf{w})$  in this contribution:

$$\Omega_L(\mathbf{x}, \mathbf{w}) = \sum_{j=0}^J w_j \left( 1 + \exp \left( - \sum_{n=0}^3 w_{jn} x_n \right) \right)^{-1} \quad (2)$$

Network output can be generally be expressed in terms of an output function mapping inputs and network weights into network output. Formally,  $\Omega_L : \mathfrak{R}^3 \times \mathbf{W} \rightarrow \mathfrak{R}$  where  $\mathbf{W}$  is a weight space appropriate to the network architecture embodied in  $\Omega_L$ . We take  $\mathbf{W}$  to be a subset of  $\mathfrak{R}^d$  with  $d = 5J + 1$ . Given targets  $y$ , model performance is then  $E(y, \Omega_L(\mathbf{x}, \mathbf{w}))$ .

The process of determining optimal parameter values is called training or learning and may be formulated in terms of the minimization of  $E$  given a fixed model complexity (i.e.  $J$ ). The function that is minimized in this study is squared error

$$\min_{\mathbf{w}} E(y, \Omega_L(\mathbf{x}, \mathbf{w})) = |y - \Omega_L(\mathbf{x}, \mathbf{w})|^2 \quad (3)$$

For any combination of  $y$  and  $\mathbf{x}$ , and for any choice of parameters  $\mathbf{w}$ , we can now measure model performance. It can happen that there is no unique solution to the minimization problem. There are two main reasons for this possibility. The first refers to the case of redundant inputs and the second to the case of irrelevant hidden units. The case of redundant inputs occurs when one or more of the model inputs is an exact linear combination of the other inputs, including the bias input. The case of irrelevant hidden units occurs when identical optimal model performance can be achieved with fewer hidden units. Both redundant inputs and irrelevant hidden units generate entire manifolds in  $\mathbf{W}$  on which the performance statistic is flat and minimal. Because finding and eliminating redundant inputs and irrelevant hidden units both involve some computational efforts, network models are usually trained without regard to their possible presence.

### 3. Global Optimization over the Samples and Model Performance

Gradient descent procedures in combination with the backpropagation technique provide one approach to attempting to find a solution to the problem (3) what is typically a highly nonlinear optimization problem. In this study a global search approach based upon the differential evolution method (DEM) is used that appears to provide out-of-sample performance superior to gradient descent procedures such as the conjugate gradient procedure (see Fischer et al. 1999) and requires no computation of the gradient of the network output function.

The differential evolution method, originally developed by Storn and Price (1996, 1997), is a global optimization algorithm that employs a structured, yet randomized parallel multipoint search strategy which is biased towards reinforcing search points at which the error function  $E(\mathbf{w})$  being minimized has relatively low values. The DEM is



similar to the method of simulated annealing (Kirkpatrick et al. 1983) in that it employs a random (probabilistic) strategy. But one of the apparent distinguishing features of DEM is its effective implementation of parallel multipoint search. DEM maintains a collection of samples from the search space rather than a single point. This collection of samples is called *population* of trial solutions.

To start the stochastic multipoint search, an initial population  $\mathbf{P}$  of, say  $M$ ,  $d$ -dimensional parameter vectors  $\mathbf{P}(0) = \{\mathbf{w}_0(0), \dots, \mathbf{w}_{M-1}(0)\}$  is created with  $M = 300$  and  $d = 5J + 1$  in the current study. The initial population is drawn at random from a uniform distribution between  $-0.3$  and  $0.3$ . From this initial population, subsequent populations  $\mathbf{P}(1)$ ,  $\mathbf{P}(2)$ , ...,  $\mathbf{P}(t)$ , ... will be computed by a scheme that generates new parameter vectors by adding the weighted difference of two vectors to a third. If the resulting vector yields a lower error function value than a predetermined population member, the newly generated vector will replace the vector which it was compared to, otherwise the old vector is retained. Similarly to evolution strategies, the greedy criterion is used in the iteration process and the probability distribution functions determining vector mutations are not a priori given. Different strategies arise depending upon whether some specific type of crossover is introduced to increase the diversity of the new parameter vectors or not. The scheme for generating  $\mathbf{P}(t+1)$  from  $\mathbf{P}(t)$  with  $t \geq 0$  as utilized in the current study may be summarized by three major stages:

*Stage 1:* For each population member  $\mathbf{w}_m(t)$ ,  $m = 0, 1, \dots, M-1$ , a perturbed vector  $\mathbf{v}_m(t+1)$  is generated according to:

$$\mathbf{v}_m(t+1) = \mathbf{w}_{best}(t) + \kappa(\mathbf{w}_{r1}(t) - \mathbf{w}_{r2}(t)) \quad (4)$$

with  $r1, r2$  integers chosen randomly from  $\{0, \dots, M-1\}$  and mutually different. The integers are also different from the running index  $m$ .  $\kappa \in (0, 2]$  is a real constant factor that controls the amplification of the differential variation  $(\mathbf{w}_{r1}(t) - \mathbf{w}_{r2}(t))$ . The parameter vector  $\mathbf{w}_{best}(t)$  which is perturbed to yield  $\mathbf{v}_m(t+1)$  is the best parameter vector of population  $\mathbf{P}(t)$ .

*Stage 2:* The decision whether or not the  $\mathbf{v}(t+1)$  should become a member of  $\mathbf{P}(t+1)$ , is based on the greedy criterion. If

$$E(\mathbf{v}_m(t+1)) < E(\mathbf{w}_m(t)) \quad (5)$$

then  $\mathbf{w}_m(t+1)$  is replaced by  $\mathbf{v}_m(t+1)$ , otherwise the old value  $\mathbf{w}_m(t)$  is retained as  $\mathbf{w}_m(t+1)$ .

*Stage 3:* The iteration process continues until the error measured using an independent validation set starts to increase.

Heuristically, one might expect an optimal  $\mathbf{w}_m$  to emerge from this process as the number  $[t]$  of generations becomes large. But to date, there does not appear to be a general theoretical result guaranteeing that the optimal solution is indeed produced in the limit. Such a result would be highly desirable. Nevertheless, the differential evolution method does seem to perform reasonably well in applications.

Model performance is measured as normalized mean squared error [i.e. mean squared error divided by the overall variance of the target] or in other words as *the average relative variance*  $ARV(S)$  of a set  $S$  of patterns given by (see Fischer and Gopal 1994):

$$ARV(S) = \frac{\sum_{(x_k, y_k) \in S} (y_k - \Omega_L(x_k, \mathbf{w}))^2}{\sum_{(x_k, y_k) \in S} (y_k - \bar{y})^2} \quad (6)$$

where  $\bar{y}$  denotes the average over the target values in  $S$ . The averaging makes  $ARV(S)$  independent of the size of the set  $S$ . Thus,  $ARV(S)$  provides a normalized mean squared error metric for assessing the in-sample and out-of-sample performance of trained neural spatial interaction models.  $ARV(S) = 1$  if the estimate is equivalent to the mean of the data (i.e.,  $\Omega_L(x_k, \mathbf{w}) = \bar{y}$ ).

#### 4. Experimental Design and Bootstrapping Methodology

The standard approach for finding a good neural spatial interaction model [see Fischer and Gopal 1994] is to split the available set of samples into three sets: training, validation, and test sets. The training set is used for parameter estimation. In order to

avoid overfitting, a common procedure is to use a network model with sufficiently large  $J$  for the task, to monitor – during training – the out-of-sample performance on a separate validation set, and finally to choose the network model that corresponds to the minimum on the validation set, and employ it for future purposes such as the evaluation on the test set. It has been common practice in the neural network community to fix these sets. But recent experience has found this approach to be very sensitive to the specific splitting of the data. Thus, usual tests of out-of-sample or generalization reliability may appear over optimistic.

Randomness enters in two ways in neural spatial interaction modelling: in the splitting of the data samples on the one side and in choices about the parameter initialization and the control parameters of the estimation approach utilized [such as  $\kappa$  in the global search procedure] on the other. This leaves one question widely open. What is the variation in out-of-sample performance as one varies training, validation and test sets? This is an important question since real world problems do not come with a tag on each pattern telling how it should be used. Thus, we will vary both the data partitions and parameter initializations to find out more about the distributions of out-of-sample errors.

Monte Carlo experiments can provide certain limited information on the behaviour of the test statistics for fixed  $\kappa$ . The limitation of Monte Carlo experiments is that any results obtained pertain only to the environment in which the experiments are carried out. In particular, the data-generating mechanism has to be specified, and it is often difficult to know whether any given data-generating mechanism is to any degree representative for the empirical setting under study. Motivated by the desire to obtain distributional results for the test statistics that rely neither on large size approximations nor on artificial data generating assumptions, statisticians have developed resampling techniques such as bootstrapping that permit rather accurate estimation of finite sample distributions for test statistics of interest.

Bootstrapping is a computer intensive non-parametric approach to statistical inference that enables to estimate standard errors by re-sampling the data in a suitable way (see Efron and Tibshirani 1993). This idea can be applied to neural spatial interaction models in two different ways. One can consider each input-output pattern as a sampling

unit, and sample with replacement from the input-output pairs to create a bootstrap sample. This is sometimes called bootstrapping pairs (Efron and Tibshirani 1993) since the input-output pairs remain intact, and are resampled as full patterns. On the other hand, one can consider the predictors as fixed, treat the model residuals  $y_k - \Omega_L(x_k, \hat{\mathbf{w}})$  as the sampling units, and create a bootstrap sample by adding residuals to the model fit  $\Omega_L(x_k, \hat{\mathbf{w}})$ . This is termed the bootstrap residual approach. In this approach, the residuals obtained from one specific model are used in rebuilding patterns to obtain error bars reflecting all sources of error, including model misspecification. In the current contribution, we are primarily interested in variation due to data samples rather than error bars. Thus, the bootstrapping pairs approach is more appropriate here.

FIGURE 1 ABOUT HERE

The bootstrapping approach as utilized in this study is illustrated in Figure 1. The idea behind this approach is to generate many pseudo-replicates of the training, validation and test sets, then re-estimating the model parameters  $\mathbf{w}$  on each training bootstrap sample, and testing the out-of-sample performance on the test bootstrap samples. In this bootstrap world, the errors of forecast, and the errors in the parameter estimates, are directly observable. The Monte-Carlo distribution of such errors can be used to approximate the distribution of the unobservable errors in the real parameter estimates and the real forecasts. This approximation is the bootstrap: it gives a measure of the statistical uncertainty in the parameter estimates and the forecasts. Focus is laid in this study on the performance of the forecasts measured in terms of *ARV*.

In more detail the approach may be described by the following steps: The details may be a bit complicated, but the main idea is straightforward:

*Step 1:* Conduct three totally independent resampling operations in which

- (i)  $B$  independent *training bootstrap samples* are generated, by randomly sampling  $KI$  times, with replacement, from the observed input-output pairs  $S = \{(x_1, y_1), (x_2, y_2), \dots, (x_K, y_K)\}$ :

$$S_{Train}^{*b} = \{(x_{k_1}^{*b}, y_{k_1}^{*b}), (x_{k_2}^{*b}, y_{k_2}^{*b}), \dots, (x_{k_{K1}}^{*b}, y_{k_{K1}}^{*b})\} \quad (7)$$

for  $k_1, k_2, \dots, k_{K1}$  a random sample of integers 1 through  $K1 < K$  and  $b = 1, \dots, B$  [in this study:  $K=992, K1=480, B=376$ ],

- (ii)  $B$  independent *validation bootstrap samples* are formed, by randomly sampling  $K2$  times, with replacement, from the observed input-output pairs  $S$ :

$$S_{Valid}^{*b} = \{(x_{k_1}^{*b}, y_{k_1}^{*b}), (x_{k_2}^{*b}, y_{k_2}^{*b}), \dots, (x_{k_{K2}}^{*b}, y_{k_{K2}}^{*b})\} \quad (8)$$

for  $k_1, k_2, \dots, k_{K2}$  a random sample of integers 1 through  $K2 < K1 < K$  and  $b = 1, \dots, B$  [in this study:  $K2=256$ ],

- (iii)  $B$  independent *test bootstrap samples* are formed, by randomly sampling  $K3$  times, with replacement, from the observed input-output pairs  $S$ :

$$S_{Test}^{*b} = \{(x_{k_1}^{*b}, y_{k_1}^{*b}), (x_{k_2}^{*b}, y_{k_2}^{*b}), \dots, (x_{k_{K3}}^{*b}, y_{k_{K3}}^{*b})\} \quad (9)$$

for  $k_1, k_2, \dots, k_{K3}$  a random sample of integers 1 through  $K3 = K2 < K1 < K$  and  $b = 1, \dots, B$  [in this study:  $K3=256$ ],

*Step 2:* For each training bootstrap sample  $S_{Train}^{*b}$  ( $b=1, \dots, B$ ) minimize

$$\sum_{k=1}^{K1} |y_k^{*b} - \Omega_L(x_k^{*b}, \mathbf{w})|^2 \quad (10)$$

with the differential evolution procedure. The training process is stopped when  $ARV^*(S_{Valid}^{*b})$  defined by (6) starts to increase. This yields bootstrap parameter estimates  $\hat{\mathbf{w}}^{*b}$ .

*Step 3:* Calculate the bootstrap  $ARV$ -Statistic of generalization performance,  $\hat{ARV}^*(S_{Test}^{*b})$ , for each test bootstrap sample. The distribution of the pseudo-

errors  $A\hat{R}V^* - A\hat{R}V$  can be computed, and used to approximate the distribution of the real errors  $A\hat{R}V^* - ARV$ . This approximation is the bootstrap.

*Step 4:* The variability of  $A\hat{R}V^*(S_{Test}^{*b})$  for  $b=1, \dots, B$  gives an estimate of the expected accuracy of the model performance. Thus, estimate the standard error of the generalization performance statistic by the sample standard deviation of the  $B$  bootstrap replications:

$$\hat{s}e_B = \left\{ \sum_{b=1}^B [A\hat{R}V^*(S_{Test}^{*b}) - A\hat{R}V^*(.)]^2 / (B-1) \right\}^{\frac{1}{2}} \quad (11)$$

where

$$A\hat{R}V^*(.) = \sum_{b=1}^B A\hat{R}V^*(S_{Test}^{*b}). \quad (12)$$

Note that the name 'bootstrap' refers to the use of the original sample pairs to generate new data sets. The procedure requires to retrain the neural spatial interaction model  $B$  times (see *Step 2*). Typically, the  $B$  is in the range  $20 \leq B \leq 200$  (see Tibshirani 1996). Increasing  $B$  further does not bring a substantial reduction in variance (Efron and Tibshirani 1993).

## 5. Data and Experimental Results

To demonstrate the bootstrapping approach, we use the Austrian telecommunication flow data (see Fischer and Gopal 1994 for details). The data set was constructed from three data sources: a (32, 32)-interregional telecommunication flow matrix, a (32, 32)-distance matrix, and gross regional products for the 32 telecommunication regions. We have  $K = 992$  input-output patterns.  $K1=480$ ,  $K2=K3=256$ , and  $B=376$ . All inputs and the target data were scaled to have zero mean and unit variance over the entire original data set. In contrast to Fischer et al. (1999) we did not use a log-transformation of the input and output data.

We use that model specification with  $J = 8$  from the class  $\mathcal{Q}_L(\mathbf{x}, \mathbf{w})$  with one hidden layer of logistic units and a linear output unit. The initial parameters were drawn at random from an uniform distribution between -0.3 and 0.3. The differential evolution method with  $\kappa=0.9$  was used for the task of parameter estimation. All computations were done on a DEC Alpha Cluster 350 MHz.

## FIGURE 2 ABOUT HERE

Figure 2 shows the learning curves of a typical run, for  $S_{Train}^{*b}$ ,  $S_{Valid}^{*b}$  and  $S_{Test}^{*b}$  in terms of the  $ARV(S_{Train}^{*b})$ ,  $ARV(S_{Valid}^{*b})$  and  $ARV(S_{Test}^{*b})$  respectively. The term learning curve is used to characterize the as a function of iterations of the differential evolution method. Figure 1(a) plots the  $ARV$  performance on the training bootstrap set, Figure 1(b) the  $ARV$  performance on the validation bootstrap set and Figure 1(c) the  $ARV$  performance on the test bootstrap set. Note the clear increase of validation and test errors after passing through minima, usually called overfitting or overtraining. At some stage - in this specific run with a very small number of iterations around 70 - the model extracts a feature of the training set that helps the testset, but hurts the validation set. Note also that the minima of the validation set and the test set do not occur at the same learning time. From each of these sets of learning curves, only a single number is used for the subsequent analysis and comparisons in this paper: the  $ARV$  performance value on the test set at that time that has the minimum of the validation set.

Several experiments were conducted to get a better statistical picture of model performance measured in terms of out-of-sample performance, including the ability to estimate the effect of the randomness of the splits of the data versus the randomness of initial conditions of the spatial interaction model. The experiments involve a nested iteration. At the 'outer loop' training, validation and test bootstrap data sets are built up one after another  $B$  times. The training sets are presented to the 'inner loop' for random parameter initialization of the spatial interaction model. At each pass through the outer loop training, validation and test bootstrap samples are generated as independent draws from  $S$ ,  $B$  times. On each pass through the inner loop the  $d=41$  individual weights of the network model are then initialized randomly from a uniform distribution over

$[-0.3, 0.3]$ ,  $B$  times, the model being re-estimated, and the  $ARV$ -statistic calculated on the test sample.

TABLE 1 and FIGURE 3 ABOUT HERE

Table 1 along with Figure 3 summarize the results of a [degenerated bootstrap] experiment, with  $B=1$  and  $\beta=376$ . The experiment serves to illustrate the impact of parameter initialization on the out-of-sample performance of the neural spatial interaction models. Figure 3 (b)-(f) shows the empirical density from 376 simulations based upon  $B=1$  and  $\beta=376$ , while Figure 3(a) displays the empirical density from  $B=376$  bootstrap resamplings with  $\beta=1$ . The fact that the width of the histograms (b)-(f) tends to be smaller than the histogram (a) indicates that the randomness in network model initialization generates less variability than the randomness in the splitting of the data. Table 1 presents the out-of-sample model performance in terms of both the mean of 376 simulations (column 1) and the standard deviation (column 2). The Mann-Whitney U test statistic (column 3), a robust non-parametric test for the equality between two independent distributions, provides evidence that the empirical densities from 376 simulations based upon  $B=1$  and  $\beta=376$  are significantly different from that based upon  $B=376$  bootstrap replicates and  $\beta=1$ . The test statistic has a distribution that is asymptotically normal with zero mean and unit variance under the null hypothesis that the distributions are the same.

TABLE 2 and FIGURE 4 ABOUT HERE

Table 2 along with Figure 4 reports the results of a bootstrap experiment along with  $B=376$  and  $\beta=1$ . This experiment serves to illustrate the impact of variations in training, validation and test sets. In the simulation world of this experiment the standard deviations of the 376 bootstrap replications are substantially less variable, but on the whole larger than those appearing in the first experiment. This clearly indicates that the randomness in the splitting of the data generates more variability than the randomness in the network model initialization does. Figure 4 illustrates that the empirical densities from the 376 bootstrap replications are relatively similar and not significantly different in location as indicated by the Mann-Whitney U test statistic presented in Table 2.



## 6. Conclusions

This paper tried to combine bootstrap based statistical tests with neural spatial interaction modelling to get a better statistical picture of forecast variability of the model. Forecast variability can come from many sources. We focused on the noise due to the data set resampling which may be termed sample noise and parameter noise stemming from the choices in parameter initialization. Other noise sources, very different in nature, such as errors-in-variables or model misspecification had been outside of the scope of this paper. Utilizing the Austrian interregional telecommunication traffic data and the differential evolution method [ $M=300$ ,  $\kappa=0.9$ ] for solving the parameter estimation task for a fixed topology of the network model [ $J=9$ ] the study gave us important insights into the variability of forecast performance over changes in training, validation and test samples, and parameter initialization. For our example, most of the variability in forecast performance was clearly coming from sample variation and not from variation in parameter initializations. This implies that it is important not to over-interpret a model, estimated on one specific static split of the data.

The bootstrap approach proved to be extremely useful in getting a clearer picture of what might be real and what is noise. But it is important to keep in mind that each bootstrap iteration requires a run of the differential evolution method on the training bootstrap set. In very large real world problem contexts this computational burden may become prohibitively large.

Finally, it is worth mentioning that there are several other computationally intensive methods such as the jackknife and balanced repeated replications that are similar in spirit to the bootstrap procedure but quite different in detail (see Efron and Tibshirani 1993 for more details). Each of these procedures generates 'pseudo-data' sets from the original data and assesses the actual variability of a statistic from its variability over all the sets of pseudo-data. The procedures differ from the bootstrap and from one another essentially in the way the pseudo-data sets are constructed.

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**Table 1.** Variability of out-of-sample performance of the neural spatial interaction model (2) with  $J=8$ ,  $[M=300, \kappa=0.9]$  due to the randomness in parameter initialization

	Out-of-Sample Performance		Mann-Whitney U
	Average	Std.Dev.	Test Statistic
Data Samples A [ $B=1, \beta=376$ ]	0.2819	0.0888	11.72 [0.0000]
Data Samples B [ $B=1, \beta=376$ ]	0.3016	0.1150	9.24 [0.0000]
Data Samples C [ $B=1, \beta=376$ ]	0.4149	0.1015	-2.91 [0.0036]
Data Samples D [ $B=1, \beta=376$ ]	0.4440	0.0574	-7.14 [0.0000]
Data Samples E [ $B=1, \beta=376$ ]	0.5895	0.0514	-16.90 [0.0000]
Bootstrap A as Benchmark [ $B=376, \beta=1$ ]	0.4009	0.1541	- -

Average: Performance values [measured in terms of out-of-sample  $ARV$ ] represent the mean of  $\beta=376$  simulations differing in the initial parameter values randomly chosen from  $(-0.3, 0.3)$  in the case of data samples A-E.

Std.Dev.: Performance values [measured in terms of out-of-sample  $ARV$ ] represent the standard deviation of  $\beta=376$  simulations differing in the initial parameter values randomly chosen from  $(-0.3, 0.3)$  in the case of data samples A-E.

Mann-Whitney U Test Statistic: This statistic tests for distribution equality. The test statistic is  $N(0, 1)$  under the null hypothesis of distribution equality with the empirical distribution from  $B=376$  bootstrap replications. 10 and 5 percent critical values are 1.64 and 1.96 respectively for a two tailed test.

**Table 2.** Variability of out-of-sample performance of the neural spatial interaction model (2) with  $J=8$ ,  $[M=300, \kappa=0.9]$  due to sample variation

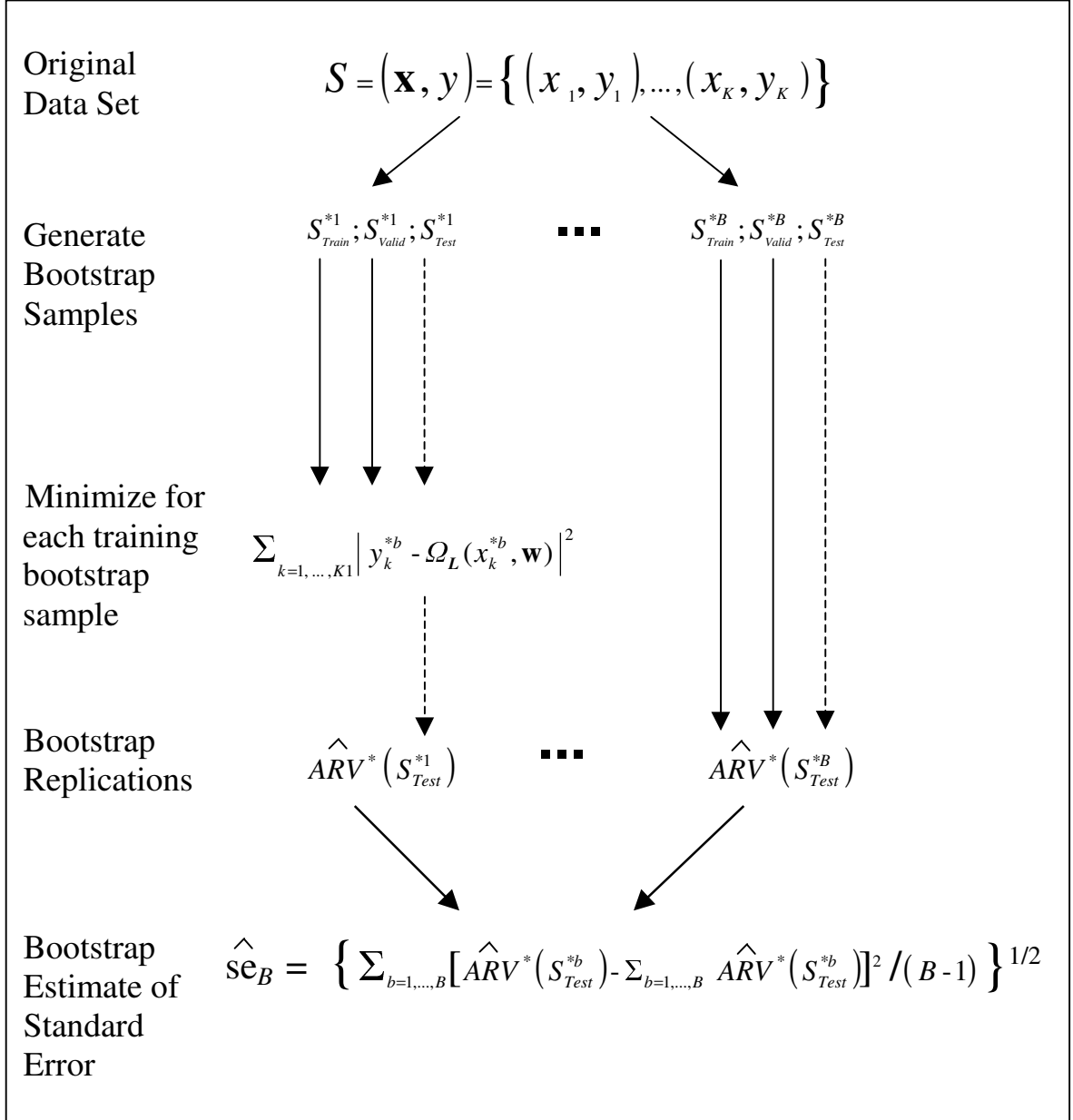
	Out-of-Sample Performance		Mann-Whitney U	
	Average	Std.Dev.	Test Statistic	
Bootstrap A [ $B=376, \beta=1$ ]	0.4009	0.1541	-	-
Bootstrap B [ $B=376, \beta=1$ ]	0.3974	0.1479	0.27	[0.7896]
Bootstrap C [ $B=376, \beta=1$ ]	0.4100	0.1595	-0.93	[0.3541]
Bootstrap D [ $B=376, \beta=1$ ]	0.3920	0.1409	0.21	[0.8359]
Bootstrap E [ $B=376, \beta=1$ ]	0.4064	0.1419	-1.17	[0.2410]
Bootstrap F [ $B=376, \beta=1$ ]	0.4057	0.1539	-0.76	[0.4458]

Average: Performance values [measured in terms of out-of-sample  $ARV$ ] represent the mean of  $B=376$  bootstrap replications with identical ( $\beta=1$ ) parameter initializations randomly chosen from  $(-0.3, 0.3)$

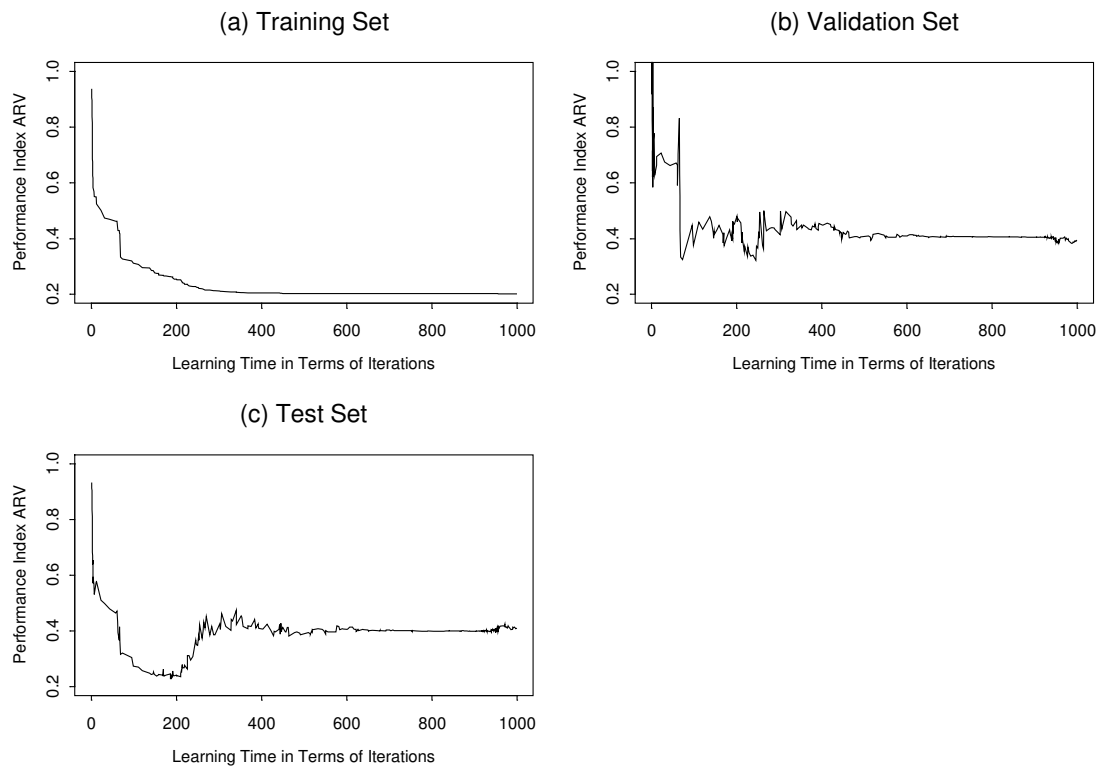
Std.Dev.: Performance values [measured in terms of out-of-sample  $ARV$ ] represent the standard deviation of  $B=376$  bootstrap replications with identical ( $\beta=1$ ) parameter initializations randomly chosen from  $(-0.3, 0.3)$

Mann-Whitney U Test Statistic: This statistic tests for distribution equality. The test statistic is  $N(0, 1)$  under the null hypothesis of distribution equality with the empirical distribution from bootstrap A. 10 and 5 percent critical values are 1.64 and 1.96 respectively for a two tailed test.

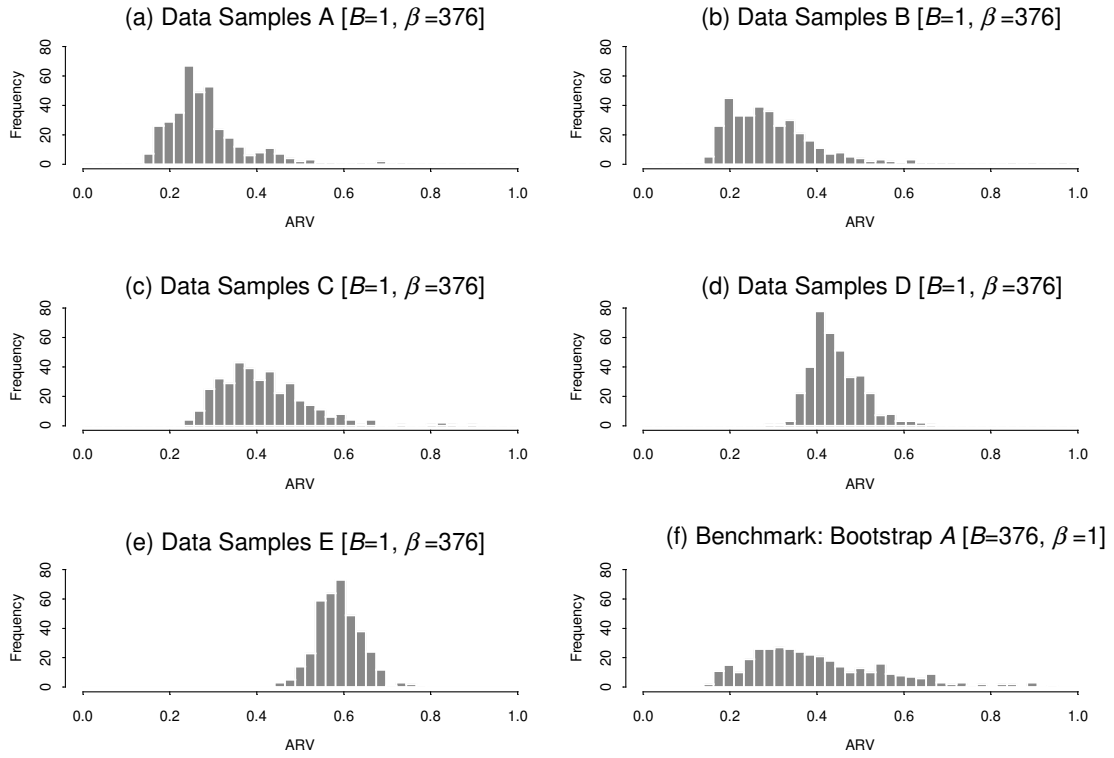
**Figure 1.** A diagram of the bootstrap procedure for estimating the standard error of the generalization performance of neural spatial interaction models



**Figure 2.** Learning curves of one specific neural spatial interaction model [ $J=8$ ] and parameter conditions [ $M=300$ ,  $\kappa=0.9$ ]. They show the performance versus the learning time in terms of iterations: (a) the performance on a training bootstrap set, (b) the performance on a validation bootstrap set, and (c) the performance on a test bootstrap set.



**Figure 3.** Histograms of out-of-sample performance [measured in terms of  $ARV$ ] of the neural spatial interaction model (2) with  $J=8$  [ $M=300$ ,  $\kappa=0.9$ ]: Variability due to the randomness in parameter initialization





**Figure 4.** Histograms of out-of-sample performance [measured in terms of  $ARV$ ] of the neural spatial interaction model (2) with  $J=8$  [ $M=300$ ,  $\kappa=0.9$ ]: Variability due to the randomness in data resampling

