# **Improving SLEUTH Calibration with a Genetic Algorithm**

Keith C. Clarke

Department of Geography, University of California, Santa Barbara, Santa Barbara CA 93106-4060, U.S.A.

Keywords: Land Use Change, Model, SLEUTH, Calibration, Cellular Automata, Genetic Algorithm.

Abstract: A review of calibration methods used for cellular automaton models of land use and land cover change was performed. Calibration advances have been achieved through machine learning algorithms to either extract land change rules, or optimize model performance. Many models have now automated the calibration process, reducing the need for subjective choices. Here, the brute force calibration procedure for the SLEUTH CA-based land use change model was replaced with a genetic algorithm (GA). The GA calibration process populates a "chromosome" with five parameter combinations (genes). These combinations are then used for model calibration runs, and the most successful selected for mutation, while the least successful are replaced with randomly selected values. Default values for the constants and rates of the genetic algorithm were selected from SLEUTH applications. Model calibrations were completed using both brute force calibration and the GA. The GA model performed as well as the brute force method, but used vastly less computation time with speed up of about 3 to 22. The optimal values for GA calibration are set as the defaults for SLEUTH-GA, a new version of the model. This paper is a contraction of Clarke (in press), which reports on the full set of results.

# **1 INTRODUCTION**

Land use change is driven by the conversion of natural lands to agriculture, and increasingly by the expansion of built-up land. Cities expand impervious surfaces outward and inward and create other land use changes at a distance. Land use and land cover change modeling attempts to simulate these changes, and asks how they can be modified, diverted or prevented so that future cities are more sustainable.

Modeling can seek to gain an understanding of a process, usually as revealed by spatial forms (Clarke 2014a). Modeling seeks to forecast a process, and so predict where and when changes will occur (NRC 2014). Models allow exploration of alternative futures by varying the forecasts to embody different anticipated circumstances (Xiang and Clarke 2003; Houet et al. 2016). A model can also help others understand the process, its outcomes and its consequences, and so educate. These purposes are dependent on the accuracy, reliability and effectiveness of the model.

Good models make their assumptions about a process explicit, use facts and data as inputs, then create accurate forecasts of future system states. To be accurate, models must use real data to fine tune the controls that create model behavior. The model design should make careful choices of constants and variables; and the model should use hindcasting, that is, be applied to historical data to effectively replicate the present. Accuracy can then be assessed as the level of agreement between the forecast and the actual (Pontius et al. 2007). The model's level of accuracy, reliability and effectiveness can then be measured and optimized. This stage is called model calibration, and calibration remains the most critical phase of model design and application.

### 2 CALIBRATION

Calibration uses a vast array of tools and techniques to optimize a model and seeks to determine the impacts of changes in a specific constant or variable upon the model outputs. Constants are the values that remain internal to the model, and may be choices of particular values or more structural elements of the model. The determination of constants is the first stage of calibration during model design. Methods include inspection of the correspondence of outputs, match statistics and the computation of many outputs across a range of

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DOI: 10.5220/0006381203190326

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In Proceedings of the 3rd International Conference on Geographical Information Systems Theory, Applications and Management (GISTAM 2017), pages 319-326 ISBN: 978-989-758-252-3

constant values. Critical in calibration are threshold values, where a small change in the constant produces large differences in the output--what Houet et al. (2016) call "non path-dependent" and contrasted/breaking trends. Simple models avoid these values, while complex systems models exploit them. Crossing these thresholds is called phase change in complexity theory, and leads to emergence (Holland 1998).

Calibration also involves repeated application of the model, the measurement of model performance, degree of fit, and the adjustment of input variables and data until the performance is maximized. This may involve accuracy of the model outputs as measured using historical data, or achievement of some other goal. A model is started at some point in the past, and executed without further input until the last period of known data (the present), periodically matching its numerical and spatially distributed outputs with real data.

Given the matches described above, measures can be compiled that represent multiple performance parameters. Changing parameters and repeating the model application allows retention of the best performing settings. One way to optimize is to repeat the parameter changes for all possible combinations and permutations of their values, socalled brute force. Models increasingly use machine learning algorithms to optimize. For example, weights assigned in agent based models can be selected using support vector machines, or cellular automata behavior rules selected using genetic algorithms (Clarke 2014b). Good calibrations derive the best set of input parameters that determine the model's performance, accuracy and behavior. Good models are always well calibrated.

Models of land use and land cover change have a vast literature, with periodical reviews and surveys of the models and their applications (NRC 2014). All land change models require calibration, but these calibrations are a function of the model type and its intended purpose. A subset of land use change models is cellular automata (CA) models, discussed at length (Torrens and O'Sullivan 2001) and divided into types (Sante et al. 2010). This short paper focuses on CA models only, then a particular model and its improvement using a genetic algorithm (GA) to replace its current brute force calibration method. An advantage of this approach is that it removes human interaction and judgement entirely from the calibration process (Jafarnezhad et al. 2015).

#### 2.1 Cellular Automata Models

CA models are complex system models consisting of: (1) a set of mutually exclusive and nonoverlapping states; (2) a framework of points, cells or a grid in which each element is in one and only one state; (3) a defined neighborhood, consisting of a set of cells usually surrounding or adjacent to a cell; (4) a set of rules that govern state changes as a function of the other states within the neighborhood; (5) a relation to discrete time, such that all cells are evaluated in each time step; and (6) an initial arrangement of the states within each of the cells.

In CA land use change models, the states are standard land use classes, such as forest, agriculture, urban and wetlands; the framework is a map, a grid of raster cells within a GIS; the neighborhood is the adjacent cells of the Moore, Von Neumann or other neighborhood; the time steps are annual increments from a start time to a stop time; and the initial arrangements are mapped distributions at some point in past time. The rules are determined during the model design stage by following those of other models, using some a priori assumption about behavior, derived statistically system using probabilities or from exogenous quotas, or derived from data mining of past land use changes as functions of location, type and quantity.

The rule sets associated with land use and land cover change are often chosen by analysis of the driving factors of land use change. The factors that prove significant are then prioritized and assigned weights. Modeling then consists of taking an input model, combining the weighed input factors, deciding probabilistically whether a change from type A to type B could occur, then enacting the change at the most probable locations.

## 2.2 Calibrating CA Models

Using two land use maps as inputs to derive a rule set for CA by data mining has led to numerous attempts to calibrate CA models with data reduction methods. These include multi-criterion evaluation (MCE) (Wu and Webster 1998), multi-objective optimization (Cao et al 2014), logistic regression (Wu 2002) and decision trees (Li and Yeh 2004). Most successful among these methods have been neural networks (Yang and Li 2007). Some models use neural networks as the entire basis for land use change modeling (e.g. ANN-CA by Li and Gar-On Yeh 2002; and LTM by Pijanowski et al. 2002).

Other machine-learning algorithms have been used to help calibrate (and derive CA rules for) CA

models of land use and land cover change. Long et al. (2009), Hu and Lo (2007) and Liu and Phinn (2003) used logistic regression to select CA transition rules in the model design stage. Guan et al. (2005) used artificial neural networks for the same purpose. Another method is the support vector machine (Yang et al. 2008). Others have used neural networks to optimize CA control parameters (Li and Yeh 2004). More recently, such methods as particle swarm optimization (Feng et al. 2011) and ensemble learning strategies (multiple methods in parallel) have also been introduced (Gong et al. 2012).

Among the most successful machine learning methods for CA rule selection and parameterization are genetic algorithms (GA). A GA is a method for solving optimization problems based on a process of natural selection that mimics evolution in plants and animals. The algorithm starts with an approximate initial set of solutions, and then repeatedly modifies the population of genes while assessing fitness. Each iteration, changes are made to create better solutions (evolution and mutation) and to allow new random solutions that may outperform the current best "gene." Studies that have used GA to calibrate CA include Colonna et al. (1998), Goldstein (2004), Yang and Li (2007), Yang et al. (2008), Shan et al. (2008), Cao et al. (2011), Feng and Liu (2012), Clarke-Lauer and Clarke (2011), Garcia et al. (2013) and Jafarnezhad et al. (2015).

There are many possible measures of goodness of fit between a real map and a modeled map (fitness of the gene or chromosome), including producers and users accuracy, various Kappa measures, matching of landscape metrics, correlation, the Receiver Operating Characteristics curve and others. Many calibrations simply use the percent correct as a measure. As an example, the SLEUTH model produces 13 regression-based fit measures, which in the past were combined by multiplication, although many studies have used the Lee-Sallee metric alone (Silva and Clarke 2002). Current practice uses the Optimal SLEUTH Metric (OSM) (Dietzel and Clarke 2007). This measure uses a subset of 7 of the 13 metrics, also combined by multiplication, selected to reduce interdependencies among the 13 metrics. The study reported here used the OSM as the fitness measure for calibrating SLEUTH.

Use of GA implies creation of the equivalent of a chromosome, with individual genes reflecting traits of an individual. SLEUTH has five control parameters, which vary from 0-100, termed diffusion, breed, spread, slope and road growth. A single run is controlled by the five values within the integer range  $\{0,0,0,0,0\}$  to  $\{100,100,100,100,100\}$ .

The single set of five values forms a gene, and a population of P such sets is the chromosome. Each gene is evaluated, i.e. the model is run and the fitness calculated. The genes are then sorted by fitness, so that those that performed best rise to the top. This is termed a generation. Between generations, new genes are created by combining the values of the best performing genes, after having pairs of chromosomes "compete" to reproduce, and so share their genes. Some of the genes in the chromosome are mutated, by altering their values. The mutation rate is the proportion of the chromosome subjected to change. Mutation can be by switching values or replacing values with random numbers. There are two levels of fitness associated with each generation: the total fitness of the chromosome and the specific fitness of a gene. In our case, we are interested in maximizing both total fitness to move the training process forward, and the fitness of the best performing gene, which is the best model fit at that generation. Evolution ends when a maximum number of generations is reached, or when successive generations have no better total fitness than their parents.

The chief variables in a GA include choosing the size of the population (number of genes in the chromosome), the maximum number of generations (or minimum improvement in fitness to continue evolution), the mutation rate, number of crossovers, the number of offspring, and the number of replacements. A second stopping criterion is the maximum number of evaluations of genes for possible inclusion as replacements. The GA populates the initial chromosome with genes using random numbers, standardizing values between zero and one hundred. In one generation, each of the genes is used as model input, and the fitness criterion calculated. In Blecic et al.'s (2010) study, the fitness values used were the Kappa coefficient and the Lee-Sallee metric (Silva and Clarke 2005), others have used the Optimal SLEUTH Metric (Dietzel and Clarke 2007). This is repeated for all genes in the chromosome, and the results ranked.

Each generation some proportion of the genes are crossed over. For example a set of SLEUTH input parameters may be {10, 20, 30, 40, 50}. After mutation, it may be {10, 20, 50, 40, 30} with 2 values switched and 3 remaining. Another form of mutation simply randomly or incrementally changes one or more gene values. Lastly, the lowest performing genes in terms of fitness are "killed off" and replaced with new random values. Such a choice increases the number of evaluations, when a maximum number is reached or a maximum number of generations pass, the winning genome is selected.

This final replacement stage is important because there is always a possibility that the chromosome with the highest total fitness is not a global but only a local maximum. Mutation and replacement ensure that a superior value either evolves or arrives by chance. The altered chromosome is then subjected to the next generation, and the process is repeated either until no further gain in fitness is achieved, or a maximum number of generations exceeded.

While research continues on using GA as a means to calibrate CA models, relatively few studies have examined how the specifics of the GA impacts the performance, accuracy and tractability of model calibrations. Obviously this can only be answered in the context of a single model. SLEUTH will be used for this purpose because it is one of the few instances where both brute force and GA calibration options are available in open source code.

### 2.3 Calibrating SLEUTH

SLEUTH is a land use and land cover change model based on two tightly coupled CA models: the Urban Growth Model, that simulates how urban areas expand and change; and the Deltatron model that propagates urban changes into other land use types. The model was originally developed and applied to the San Francisco Bay area (Kirtland et al. 1994; Clarke et al. 1997) and then to the Washington-Baltimore area (Clarke et al. 1998). SLEUTH's initial calibration was by monolooping (trying all possible settings for each parameter, holding the others constant), but this was replaced by brute force calibration (Clarke at al. 1996). The calibration methods were systematically improved over decades (Clarke et al. 2007; 2008a; 2008b; Chaudhuri and Clarke 2013). Recently, research has examined the goodness of fit between SLEUTH simulations and actual data, usually using hindcasting and spatial metrics of various kinds (Wu et al. 2009; Rienow and Goetzke 2014; Sakieh 2013).

Noah Goldstein was the first to experiment with GAs to calibrate SLEUTH (Goldstein 2004). Others tried the same approach with more sophisticated tools (Clarke-Lauer and Clarke 2011; Jafarnezhad et al. 2015). Clarke-Lauer and Clarke used the OSM as the fitness criterion and replaced the brute force module in SLEUTH with a new code routine that employed a GA that was posted to SourceForge. Values that could be varied included choices on encoding, fitness evaluation, crossover, mutation and survival selection. Coding involved a

random number between 0 and 4 to index the five SLEUTH control parameters (diffusion, breed, spread, slope and road growth) and to decide how many elements from the parent were to be reproduced in the offspring. Remaining elements were selected from the second parent, with the second offspring using the opposite genes used for the first. Parents were selected by tournament selection, with a random set selected and the parents chosen with the highest fitness. Each generation replaces the weakest genes in the old population with the strongest in the new. The SLEUTH-GA was tested using the demo city sample data set available on the SLEUTH website. Mutation rates of 0.10 to 0.16 were found satisfactory, with a population size of 25. The paper concluded that the GA produced a speed up by a factor of 5 over brute force.

Jafarnezhad et al. (2015) used the SLEUTH-GA code to apply SLEUTH to 3 cities in Golestan Province, Iran. They calibrated SLEUTH first using the standard brute force procedure, then used GA with the fitness metric as the OSM. They coded their own GA procedures based on Goldstein's method (Goldstein 2004). Model outputs were then compared using the Receiving Operator Statistic (ROC), landscape metrics and two Kappa coefficients. Speed up over brute force was 4-5 times, and the authors noted that the results could be improved by "testing different values for mutation rate and decreasing model tendency to elitism."

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## **3 RESULTS**

Existing SLEUTH data for San Diego, California and Andijan, Uzbekistan were used (Syphard et al, 2011). The Andijan data set produced the lowest OSM fits achieved by SLEUTH. In both cases these were the best model calibrations, but they varied substantially in predictive power. This is believed to be because of Andijan's extraordinary urban growth history. The full set of results and data details are published in Clarke (in press).

Both cities were then used with identical inputs in the SLEUTH-GA version of the model code. The SourceForge version was adjusted slightly to take six parameters from the shell to be passed to the code. These were the population size (genes in the chromosome), the maximum number of generations, the mutation rate, the maximum number of evaluations per gene, the number of offspring, and the maximum replacement number. Population size, mutation rate, number of offspring, the replacement number and the maximum number of evaluations were varied, while the other values were held constant. The maximum number of generations was set to 100, but in fact the GA rarely used more than 20 generations in the calibration, contrary to the higher numbers determined by Jafarnezhad et al. (2015). The maximum number of evaluations for substitution per chromosome was found to give peak fitness at about 900, and this did not affect the calibration process, other than increasing the number of generations and CPU time.

Table 1: Brute Force Calibration Results. Values for constants are after calibration, with high and low coefficients in the top 8 solutions given, then after averaging to the last time period.

	San Diego	Andijan
Calibration	1960-1999	1934-2013
period		
Best OSM	0.7414836	0.0773797
Diffusion/derive	(100:98-100)	(63:60-63)
d	100	100
Breed/ derived	(97:97-99)	(100:85-100)
	100	100
Spread/ derived	(25:24-25)	(1:1-2)
	25	3
Slope/derived	(15:15-18)	(80:75-79)
	1	1
Road gravity/	(53:45-53)	(25:15-25)
derived	53	38
Calibration time	175589	440715
(s)	E AND	TECH

For Andijan the fitness was very low, with a slight peak at a population size of 70. For San Diego, the peak fitness occurred at a population size of 55, so this value was then used for the next monoloops. Similarly for mutation rates, the peak fitness for both San Diego and Andijan was at a rate of 0.13, so this value was used for all further calibrations.

The information on calibration fine tuning for the GA was rather limited from the Andijan case, so testing of the ranges of the number of offspring and the replacement number were restricted to the San Diego data. Their best fitness values were 55 and 50 respectively. The final set of input parameters is shown in Table 2. In particular, the maximum number of evaluations sets the computation cost for the run, and there appears to be a fine balance between too many generations versus achieving a good fit. A best value of 900 was selected, which creates about 10-12 generations of evolution.

 Table 2: Genetic Algorithm Parameter Monolooping Calibration Results.

City	San Diego	Andijan
Calibration period	1960-1999	1934-2013
Best OSM	0.72972	0.07292
Maximum # of	900	900
evaluations		
Population	55	55
(Chromosome size)		
Mutation Rate	0.13	0.13
Number of offspring	55	55
Replacement per	50	50
generation		
Calibration time (s)	55588	19866

This set of GA control parameters possibly provides universal application for SLEUTH-GA modeling. The values have been integrated into the SLEUTH-GA code as defaults. This goes a long way toward the fully automated and objective calibration of SLEUTH, without user intervention (Straatman et al 2004).

What range of parameters is there within the chromosome that might still be improved by brute force calibration over a smaller range, and what is the impact of this difference on the actual forecasts spatially? Table 3 shows the ranges of parameters in the first gene subpopulation (highest performing individuals of the 8 most fit parents) for the best GA derived parameters. The maximum, average and total fitness of a chromosome tend to peak simultaneously, indicating that the best performing chromosome is led by the most fit gene.

Table 3: Genetic Algorithm Calibration Results.

	San Diego	Andijan
C I'I C	Ŭ	2
Calibration	1960-1999	1934-2013
period		
Best OSM	0.729724	0.072920
Diffusion/	(90: 79-90)	(54:53-94)
derived	100	82
Breed/ derived	(23: 22-25)	(2:0-2)
	26	3
Spread/ derived	(89:74-98)	(88:62-94)
	100	82
Slope/derived	(13:2-32)	(70:9-70)
	1	3
Road gravity/	(19:19-98)	(47:14-47)
derived	30	75
Calibration time	55588	19866
(s)		
Speed Up	3.16	22.18

To investigate the spatial impact of the differences in calibration mode, maps of forecast

urbanization with a likelihood of over 50% were created for the two cities and shown for both methods of calibration (Figure 1). It is evident that as in the calibrations, both cities are forecast with higher uncertainty and greater spread using brute force calibration, while the forecasts for both cities are more constrained but with greater certainty using GA. This appears to be the case both for high and low model fit, and may be a robust way of providing better forecasts.

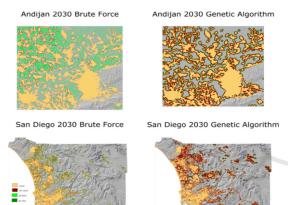


Figure 1: Spatial extent of SLEUTH forecasts and Actual Urban Growth During the Calibration Period.

## 4 CONCLUSION

Santé et al. (2010) pointed out the "need of making urban CA more flexible while keeping their simplicity bv developing better calibration methods." This study has been in response to this challenge. An important move, suggested by Jafarnezhad et al. (2015) is to eliminate human choices and judgements during the calibration process, replacing the subjective with the objective (Goldstein 2004). On the surface, replacing the brute force calibration method for SLEUTH calibration just substitutes a new set of calibration problems, i.e. dealing with the characteristics of the gene and determining how the evolutionary process yields the best results. Prior work cited above, and now this study, show that GA leads to at least equal, and often superior calibration results while considerably speeding the process. The results here also indicate lower modeling uncertainty. The differences in the calibration parameter sets are small, and the differences among model forecasts are also small. The advantages are the objectivity, and the benefits of speed-up. At the least, GA can provide a convergent set of genes that can be further optimized by brute force over a much more limited parameter set, such as the range over the top 8 genes listed in table 3.

This study reviewed the importance of calibration for CA land use change models. Calibration performs important functions for models because it ensures the model's accuracy, integrity, reliability and trustworthiness. Well calibrated models are defensible and objective, and use real world data instead of assumptions in their properties, constants, variables and behavior types. There is an obligation to perform sensitivity analyses and to run controls. Moving SLEUTH calibration from brute force to GA, the level of objectivity is further improved. As a bonus, the amount of CPU time devoted to calibration was reduced by about a factor of 3 for San Diego and 22 for Andijan. Hopefully this latter fact will enable new applications and new cities to be simulated. The final version of the posted SLEUTHGA software is at. www.ncgia.ucsb.edu/projects/gig/Dnload/download. htm and is available as open source code for modelers.

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