

# Classification of earthquakes using Grammatical Evolution

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## Abstract

Earthquake predictability remains a central challenge in seismology. Are earthquakes inherently unpredictable phenomena, or can they be forecasted through advances in technology? Contemporary seismological research continues to pursue this scientific milestone, often referred to as the ‘Holy Grail’ of earthquake prediction. In the direction of earthquake prediction based on historical data, the Grammatical Evolution technique of GenClass demonstrated high predictive accuracy for earthquake magnitude. Similarly, our research team follows this line of reasoning, operating under the belief that nature provides a pattern that, with the appropriate tools, can be decoded. What is certain is that, over the past 30 years, scientists and researchers have made significant strides in the field of seismology, largely aided by the development and application of artificial intelligence techniques. Artificial Neural Networks (ANNs) were first applied in the domain of seismology in 1994. The introduction of Deep Neural Networks (DNNs), characterized by architectures incorporating two hidden layers, followed in 2002. Subsequently, Recurrent Neural Networks (RNNs) were implemented within seismological studies as early as 2007. Most recently, Grammatical Evolution (GE) has been introduced in seismological studies (2025). Despite continuous progress in the field, achieving the so-called “triple prediction” — the precise estimation of the time, location, and magnitude of an earthquake — remains elusive. Nevertheless, machine learning and soft computing approaches have long played a significant role in seismological research. Concerning these approaches, significant advancements have been achieved, both in mapping seismic patterns and in predicting seismic characteristics on a smaller geographical scale. In such a way, our research will analyze historical seismic events from 2004 to 2011, for the Latitude 21°-79° & Longitude 33°-176°. The data will be categorized and classified, with the aim of employing Grammatical Evolution techniques to achieve more accurate and timely predictions of earthquake magnitudes. This paper presents a systematic effort to enhance magnitude prediction accuracy using GE, contributing to the broader goal of reliable earthquake forecasting. Subsequently, in this paper we present the superiority of GenClass, a key element of the Grammatical Evolution techniques, with an average error of 19%, indicating an overall accuracy of 81%.

**Keywords:** Earthquakes; Machine learning; Neural networks; Grammatical Evolution;

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## 1. Introduction

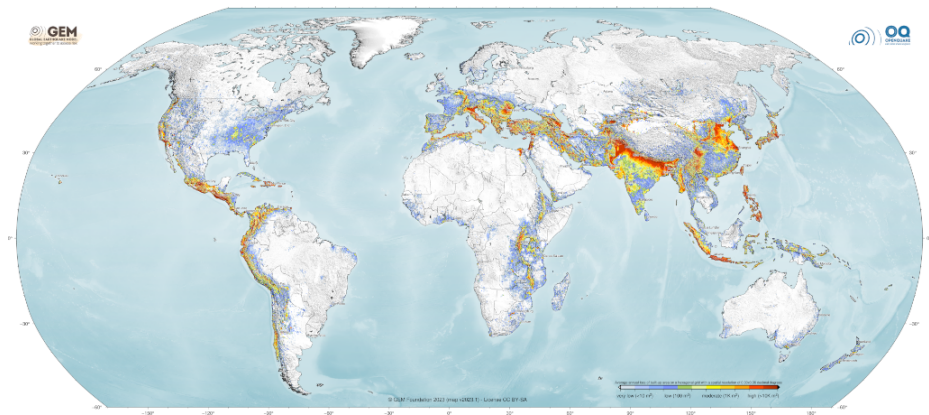
Seismology was first established as an academic discipline in 1876 at the Imperial College of Engineering in Tokyo, when John Milne (1850-1913) was invited to teach there, and in 1886, S. Sekiya (1855-1896) was appointed as the world’s first professor dedicated

specifically to the field of seismology [1]. Nevertheless, the seismologists who left a lasting legacy lending their names to the classification of earthquakes, were Charles Richter (1900-1985) and Giuseppe Mercalli (1850-1914). The former developed, in 1935, a scale measuring the magnitude of seismic events ranging from 1 to 9 [2] while the latter devised a scale that categorizes the destructive effects of an earthquake and it is shown in Table ranging from levels 1 to 12 [3]. The introduction of the Richter scale marked a significant advancement in seismology by establishing a standardized method for quantifying earthquake magnitude. This standardization enabled consistent comparisons between seismic events, thereby enhancing the analysis, preparedness, and management of earthquake-related disasters [4]. On the other hand, the Mercalli scale assessed the effects of an earthquake using qualitative criteria based on observed impacts on buildings, infrastructure, and human perception.

**Table 1.** The Modified Mercalli Intensity Scale (near the epicenter of the earthquake).

Scale Level	Consequences / Effects
I	Felt only by a very small number of people and only under particularly favorable conditions.
II	Experienced by only a small number of people.
III	Felt quite noticeably by people indoors
IV	Felt by many people indoors, but only by a few outdoors.
V	Felt by nearly everyone
VI	Felt by all
VII	Damage is insignificant in well-designed and well-constructed buildings.
VIII	Damage is slight in specially designed structures
IX	Damage is considerable in specially designed structures
X	Some well-built structures are destroyed
XI	Few, structures remain standing
XII	Damage is total

These measurements proved particularly applicable in urban environments, especially within Western societies, where systematic documentation of earthquake damage was feasible [5]. The classification of earthquakes has made it possible not only to develop seismic hazard maps but also to organize data systematically for the development of algorithms aimed at predicting seismic events. In Figure 1a hazard map issued by the Global Earthquake Model(<https://www.globalquakemodel.org/>) is outlined, indicating a 10% probability of being exceeded within 50 years.



**Figure 1.** Hazard map from Global Earthquake Model.

In this framework, researchers are increasingly employing artificial intelligence and other advanced methodologies to improve the timely prediction of earthquakes and other natural hazards, aiming ultimately to support communities in implementing protective actions and reducing potential impacts[6]. A series of machine learning techniques were applied on data gathered from earthquakes. For example, Artificial neural networks (ANNs) [7] were first applied to the field of seismology in 1994, specifically through research focused on earthquake prediction using seismic electric signals [8]. A more advanced form of an Artificial Neural Network (ANN) was employed in the related study [9]. Furthermore, Mousavi et al. [10] suggested an attentive deep-learning model for simultaneous earthquake detection. Moreover, researchers employed convolutional neural networks (CNNs)[11] [11] trained on over 18 million manually annotated seismograms from Southern California to directly infer earthquake parameters from raw waveform data, thereby eliminating the need for manual feature extraction. The model demonstrated exceptional accuracy, achieving a standard deviation of just 0.023 seconds in arrival-time estimation and a 95% success rate in polarity classification [12].

The earliest application of Deep Neural Networks (DNNs) [13], incorporating two hidden layers, was introduced in 2002 [14]. Subsequently, a high-resolution earthquake catalog was generated through the application of deep neural network (DNN) techniques, providing valuable insights into the complexity and temporal characteristics of earthquake sequences, as well as their connections with recent nearby seismic events, as outlined in related studies [15]. Additionally, Recurrent Neural Networks (RNNs) [16] within the field of seismology was introduced in 2007 [17]. Furthermore, a subsequent study regarding earthquake magnitude prediction using machine learning techniques was introduced recently [18], concentrated on forecasting earthquake magnitudes in the Hindukush region. Four machine learning approaches neural network-based pattern recognition, Recurrent Neural Networks, Random Forests [19], and a Linear Programming Boost ensemble classifier, were individually applied to model the relationships between calculated seismic parameters and the likelihood of future earthquake occurrences. Additionally, a subsequent study showed that nearest-neighbor diagrams offer a straightforward yet effective method for distinguishing between different seismic patterns and evaluating the reliability of earthquake catalogs[20]. Another research team, used Nearest Neighbor method, determined that the Weibull model offered a more accurate fit for seismic data in California, showing well-structured tail behavior. They further confirmed the model's robustness by successfully applying it to independent datasets from Japan, Italy, and New Zealand, as reported in the related publication [21].

Building on recent advancements, Rouet-Leduc et al. applied machine learning techniques to datasets obtained from shear laboratory experiments, with the objective of

identifying previously undetected signals that could potentially precede seismic events [22]. In a subsequent study, the same research team applied a machine learning-based method, initially developed in the laboratory, to analyze large volumes of raw seismic data from Vancouver Island. This methodology enabled the differentiation of pertinent seismic signals from background noise and holds promise for evaluating whether, and under what conditions, a slow slip event might be associated with or evolve into a major earthquake [23]. Additionally, a recent study developed a predictive model capable of estimating both the location and magnitude of potential earthquakes for the following week. The model utilized seismic data from the current week and targeted seismogenic regions in southwestern China. It achieved a testing accuracy of 70%, with precision, recall, and F1 score values of 63.63%, 93.33%, and 75.66%, respectively. [24]. Moreover, in another publication, the research team demonstrated that machine learning techniques can effectively predict the timing and magnitude of laboratory-induced earthquakes by reconstructing and interpreting the system's complex spatiotemporal loading history [25].

Noteworthy, is the study by Zhang et al., which offers new perspectives in the seismology field. Specifically, the authors succeeded in documenting the directionality of coseismic acoustic waves generated by a major earthquake, and in elucidating the coupling between seismic activity, the ionosphere, and space weather. Their findings are presented in the study titled "Successively equatorward propagating ionospheric acoustic waves and possible mechanisms following the Mw7.5 earthquake in Noto, Japan, on 1 January 2024"[26]. In contrast, the current work focuses primarily on the classification of earthquakes using Grammatical Evolution techniques.

In this paper, a technique based on Grammatical Evolution [27] is proposed for the efficient creation of classification rules on seismic data that are widely available on the internet. Grammatical Evolution is a genetic algorithm [28], where the chromosomes are series of production rules from a provided Backus-Naur form (BNF) grammar [29]. Among the numerous cases of application of Grammatical Evolution, one can find problems such as function approximation[30,31], economic problems[32], network security issues [33], water quality problems [34], medical problems [35], evolutionary computation [36], prediction of temperature in data centers [37], solving trigonometric problems [38], composing music [39], construction of neural networks [40,41], numerical problems [42], video games [43, 44], energy issues [45], combinatorial optimization [46], security issues [47], evolution of decision trees [48], problems that appear in electronics [49] etc.

The main contributions of the current work are:

1. The proposed method investigates a wider geographical area than other related studies, incorporating 255 seismic regions out of the 708 regions identified worldwide.
2. The current work utilized a classification method based on Grammatical Evolution to properly classify the seismic events in some predefined classes. The use of this technique has two clear advantages: on the one hand, it isolates those characteristics of a seismic event that are deemed necessary for its effective classification and on the other hand, it can discover, through the generation of complex rules, hidden linear and nonlinear correlations between the characteristics of the problem.

The remainder of this paper is organized as follows: in section 2 the used dataset is described as well as the steps of the proposed method, in section 3 the experimental results are presented and finally in section 4 some conclusions are discussed.

## 2. Materials and Methods

In this section, an extensive presentation of the seismic data used, their post-processing, as well as the computational rule construction method applied for the effective classification of earthquakes will be given.

## 2.1. The Dataset Employed

In this study, we utilized open data provided by the NSF Seismological Facility for the Earth Consortium (SAGE), specifically accessed through the Interactive Earthquake Browser (<https://ds.iris.edu/>). The choice of NSF data was motivated by its enhanced functionality. The selection of NSF data was motivated by its high accessibility. In particular, it allows for the download of up to 25,000 records per file, substantially accelerating the workflow and enhancing information retrieval. Additionally, the platform provides an interactive global map, enabling both data visualization and the extraction of datasets directly from the displayed geographical regions in real time. While the GEOFON program offers comparable information, it limits the maximum number of earthquakes retrievable per query to 1,000 events, thereby constraining the temporal coverage of data extraction. This limitation is especially relevant, as nearly 1,000 seismic events may occur globally within a single day. The NSF SAGE Facility is acknowledged as a reliable data repository, having been certified by the CoreTrustSeal Standards and Certification Board.

## 2.2. Dataset Description

We downloaded and analyzed 1,035,971 earthquake events, from IEB, between 2004-2011 (starting date 2004/04/01 - ending date 2011/03/16), count days 2487. The dataset includes the following variables: Year, Month, Day, Time, Latitude (Lat), Longitude (Lon), Depth, Mag, Region, and Timestamp. We utilize the coordinates for (latitude 21°-79° & Longitude 33°-176°), select the magnitude measurements (range 1.0-10.0), and select by default for Depth Range, in order to use all available depths. From our dataset, the daily average is 41,655 earthquakes; the minimum earthquakes in a day was 33, on 2005/03/02, and the maximum in a day was 2326, on 2011/03/11 (Tohoku earthquake 9.1 mag). The regions encompassed within these geographical coordinates are 255, while globally approximately 708 seismogenic zones have been recorded since 1970, based on data from the Interactive Earthquake Browser. The features of the original dataset are shown in Table 2.

**Table 2.** The description of the original dataset.

FEATURE	RANGE
YEAR	2004-2011
MONTH	1-12
DAY	1-31
TIME	00:00:00 - 23:59:59
LATITUDE	21.00°-79.00°
LONGITUDE	33.00°-176°
DEPTH	0.00-800.00
MAGNITUDE	1.0-10.0
TIMESTAMP	

We selected the specific time period for two main reasons. First, it allowed us to manage the exceptionally large volume of seismic data, amounting 1,035,971 records. Second, we observed a distinct pattern in the seismic data from 2015 to 2025 that did not align with established seismic trends. More specifically, the number of recorded events rose to 46,400, a finding that prompted further consideration. Regarding the geographical region, we deliberately selected seismic zones with high activity, such as those in the Mediterranean including Greece and Turkey, which share the same longitudinal range as Japan.



### 2.3. Pre-processing steps

The initial dataset was enhanced after processing the original data. An important role in this process was played by the  $d_c$  distance, which will be used to determine when two earthquakes are close in distance. This process attempts to include critical information about seismic events that can potentially be extracted from seismic events that are relatively close in both kilometer distance and time. Hence, additional features were derived based on the critical distance  $d_c$  to capture spatial-temporal proximity of events. The proposed value in the present implementation was  $d_c = 5$  km. Based on the above distance, the following features were added to the datasets:

1. Number of seismic events at a distance less than the  $d_c$  that have occurred in a previous time.
2. Average magnitude of seismic events at a distance less than  $d_c$ , which have preceded the current seismic event in time.
3. The greatest magnitude of a seismic event recorded in the past, at a distance of less than  $d_c$  from the current seismic event.
4. The time in seconds since the largest seismic event that has occurred in the past at a distance less than  $d_c$  from the current seismic event.
5. The distance in kilometers from the largest seismic event that has occurred in the past at a distance less than  $d_c$  from the current seismic event.

In addition, the size of the seismic recordings was divided into two large classes:

- The first class contains all the events with magnitude  $\leq 3$
- The second class contains all the remaining events, with magnitude  $> 3$ .

### 2.4. The used method

The method that constructs classification rules was initially presented in [50] and an implementation in C++ was suggested later by Anastasopoulos et al. [51]. The method can produce a series of classification rules in a human readable form in order to effectively classify patterns in predefined classes. This method does not require prior knowledge of the specifics of a dataset and can furthermore discover hidden correlations between the features of the dataset or even isolate only those features that play the most significant role in the successful classification of patterns. The method has been successfully applied in many cases, such as pollution detection [52], network problems [53] etc. The main steps of this method have as follows:

1. **Initialization step.**
  - (a) **Set** the parameters of the genetic algorithm:  $N_c$  as the number of chromosomes,  $N_g$  as the number of allowed generations,  $p_s$  as the selection rate and  $p_m$  as the mutation rate.
  - (b) **Perform** the initialization of chromosomes  $g_i$ ,  $i = 1, \dots, N_c$ . Every chromosome is considered as a set of randomly selected integers.
  - (c) **Set**  $k = 0$ , the generation counter.
2. **Fitness calculation step.**
  - (a) **For**  $i = 1, \dots, N_c$  **do**
    - i. **Create** a classification program  $C_i$  for the associated chromosome  $g_i$ . The construction of this program is performed using the BNF grammar of Figure 2.
    - ii. **The** produced classification program is applied to the train set of the objective problem. Denote with  $f_i$  (fitness value) the classification error for this program.

- (b) **End For** 220
3. **Application of genetic operations.** 221
- (a) **Selection:** The chromosomes are ranked according to their fitness values. 222  
The  $(1 - p_s) \times N_c$  chromosomes with the lowest fitness are carried over to the 223  
next generation without modification, while the remaining chromosomes are 224  
replaced by offspring generated through crossover and mutation. 225
- (b) **Crossover:** In this procedure a series of offspring are produced, through a 226  
process similar to biological crossover in nature. For every couple  $(\tilde{z}, \tilde{w})$  of pro- 227  
duced offsprings two chromosomes are selected from the current population, 228  
using tournament selection. Subsequently, these chromosomes will produce 229  
the set  $(\tilde{z}, \tilde{w})$  with the application of one-point crossover. A graphical example 230  
of the one-point crossover method is outlined in Figure 3. 231
- (c) **Mutation:** During this procedure, a random number  $r \in [0, 1]$  is chosen for 232  
every element  $g_{i,j}$  of each chromosome  $g_i$ . Subsequently, this element is altered 233  
randomly when  $r \leq p_m$ . 234
4. **Termination check step.** 235
- (a) **Set  $k = k + 1$**  236
- (b) **If  $k \leq N_g$  then go to Fitness Calculation step else terminate.** 237

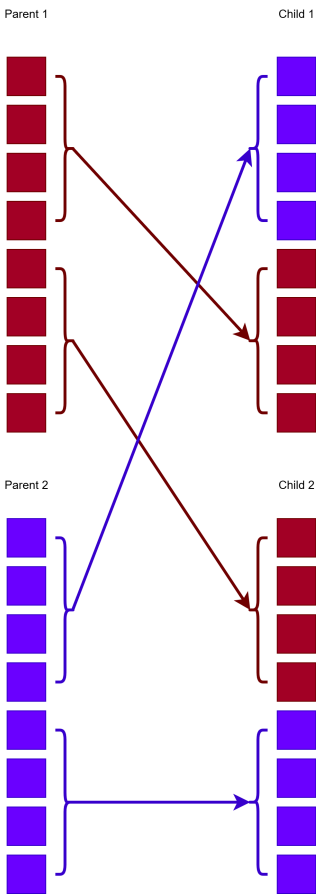
```

<S> ::= if(<BEXPR>) CLASS=0 else CLASS=1 (0)
<BEXPR> ::= <XLIST><BOOLOP><EXPR> (1)
           | !(<BEXPR>) (1)
           | <XLIST><BOOLOP><EXPR>&<BEXPR> (2)
           | <XLIST><BOOLOP><EXPR>|<BEXPR> (3)
<BOOLOP> ::= > (0)
           | >= (1)
           | < (2)
           | <= (3)

<EXPR> ::= (<EXPR><BINARYOP><EXPR>) (0)
           | <FUNCTION>(<EXPR>) (1)
           | <TERMINAL> (2)
<BINARYOP> ::= + (0)
           | - (1)
           | * (2)
           | / (3)
<FUNCTION> ::= sin | cos | exp | log (0-3)
<TERMINAL> ::= <XLIST> (0)
           | <DIGITLIST>.<DIGITLIST> (1)
           | (-<DIGITLIST>.<DIGITLIST>) (2)
<XLIST> ::= x1 | x2 | ... | xD (0-D-1)
<DIGITLIST> ::= <DIGIT> (0)
           | <DIGIT><DIGIT> (1)
           | <DIGIT><DIGIT><DIGIT> (2)
<DIGIT> ::= 0 | 1 | 2 | 3 | 4 | 5 | 6 | 7 | 8 | 9 (0-9)

```

**Figure 2.** The Backus-Naur form grammar used to generate classification rules for seismic events.



**Figure 3.** An example of the one - point crossover method.

**3. Experiments**

The code employed in the experiments was implemented using the C++ programming language, along with the freely available optimization tool [54], which can be downloaded from <https://github.com/itsoulos/GlobalOptimus.git> (accessed on 12 October 2025). Additionally, the WEKA programming tool [55] was utilized. C++ was chosen due to its high computational efficiency. Each experiment was repeated 30 times, using a different random seed for each run, and the average classification error was reported. The experimental results were validated using the ten-fold cross-validation method. The parameter values for the employed methods are presented in 3.

**Table 3.** The values for the experimental settings.

PARAMETER	MEANING	VALUE
$N_c$	Chromosomes	500
$N_g$	Generations	2000
$p_s$	Selection rate	0.1
$p_m$	Mutation rate	0.05
$d_c$	Critical Distance	5km
$H$	Number of weights	10

*3.1. Experimental results*

The following machine learning methods were used in the conducted experiments as denoted in Table 4.

1. RBF, a Radial Basis Function (RBF) network [56,57] was incorporated with 10 weights.

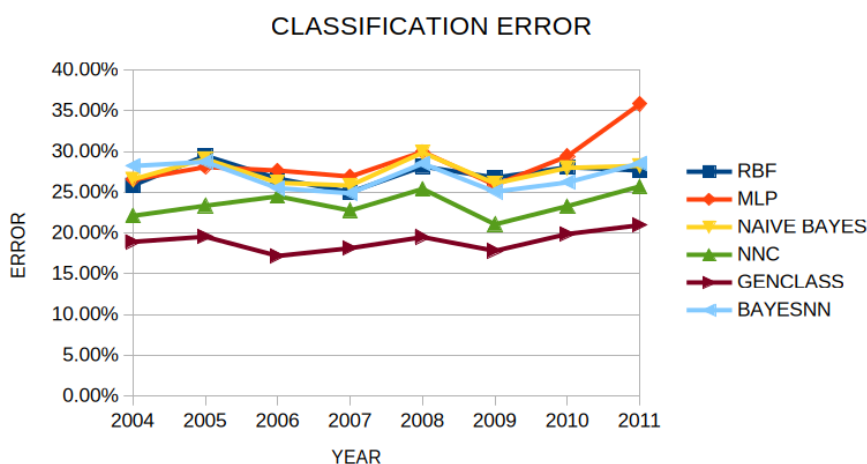


2. MLP, an multilayer perceptron neural network [58,59] with 10 processing nodes. The neural network was trained using the BFGS optimization method [60].
3. BAYES, where the Naive Bayes method [61] was utilized on the dataset.
4. The column BAYESNN denotes the results from the application of the Bayesian optimizer as implemented in BayesOpt [62] library to train a neural network with  $H = 10$  processing nodes. The code can be downloaded from <https://github.com/rmcantin/bayesopt> (accessed on 12 October 2025).
5. NNC, used to represent the application of Neural Network Construction method [63], which creates the architecture of neural networks using Grammatical Evolution.
6. GENCLASS, represents the application of the proposed method.

**Table 4.** Experimental results on the obtained datasets with the incorporation of the mentioned machine learning methods. The acronyms are defined as follows: RBF(Radial Basis Function networks), MLP (Multilayer Perceptron Network), NNC (Neural Network Construction).

YEAR	RBF	MLP	BAYES	BAYESNN	NNC	GENCLASS
2004	25.85%	26.55%	26.59%	28.23%	22.12%	18.92%
2005	29.49%	28.10%	29.09%	28.73%	23.35%	19.56%
2006	26.69%	27.67%	26.17%	25.50%	24.51%	17.19%
2007	25.02%	26.93%	25.83%	24.89%	22.77%	18.14%
2008	28.17%	29.97%	29.88%	28.53%	25.42%	19.51%
2009	26.80%	25.90%	26.09%	25.08%	21.07%	17.81%
2010	28.08%	29.43%	28.00%	26.22%	23.31%	19.89%
2011	27.67%	35.83%	28.25%	28.56%	25.73%	20.97%
AVERAGE	<b>27.22%</b>	<b>28.80%</b>	<b>27.49%</b>	<b>26.97%</b>	<b>23.54%</b>	<b>19.00%</b>

Also, the classification error for all methods per year is presented in Figure 4.

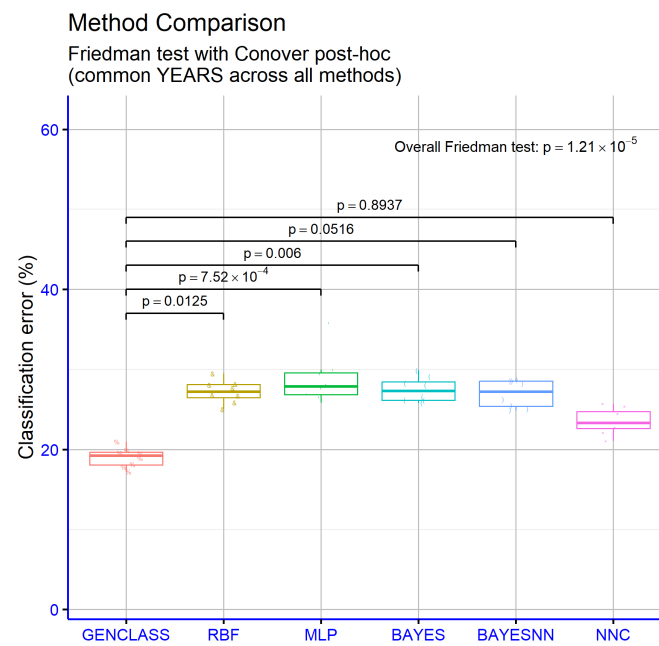


**Figure 4.** The classification error of each machine learning method per year.

Table 4 reports yearly classification error rates (2004-2011) for six machine-learning models. The proposed GENCLASS method consistently ranks first every year, achieving the lowest average error of 19.00%, which corresponds to an overall accuracy of about 81%. The runner-up is NNC with an average error of 23.54%, so GENCLASS reduces error by roughly 4.54 percentage points on average, about a 19% relative reduction versus the closest competitor. The annual margin of GENCLASS over NNC ranges from 3.20 to 7.32 points, peaking in 2006. Beyond accuracy, GENCLASS also shows the smallest across-year variability (standard deviation  $\approx 1.16$ ), indicating stable performance throughout the evaluation period. In contrast, MLP exhibits the highest mean error (28.80%) and the

largest variability, with a pronounced deterioration in 2011. Overall, these results support the article's claim that Grammatical Evolution is particularly effective for this seismic classification problem, as GENCLASS delivers both the lowest average error and the most consistent year-to-year behavior.

The R-based significance analysis reveals a clear overall difference among models (Friedman  $p = 1.21 \times 10^{-5}$ , very strong evidence), with pairwise tests indicating where these differences lie (Figure 5). GENCLASS outperforms MLP with  $p = 7.52 \times 10^{-4}$  (extremely significant) and BAYES with  $p = 0.006$  (highly significant), and the comparison with RBF is also statistically significant at  $p = 0.0125$ . In contrast, GENCLASS vs BAYESNN ( $p = 0.0516$ ) and GENCLASS vs NNC ( $p = 0.8937$ ) are not statistically significant at the 0.05 threshold. Overall, while the omnibus test confirms marked differences across models, the superiority of the proposed method is clearly supported against MLP, BAYES, and RBF, whereas the contrasts with BAYESNN and NNC do not reach conventional significance.



**Figure 5.** Statistical tests performed on the experimental results using the variety of machine learning methods.

### 3.2. Experiments with the critical distance $d_c$

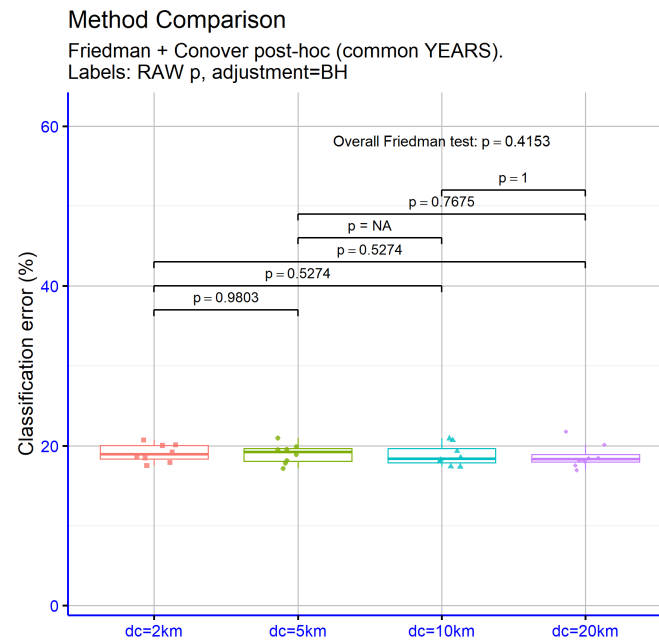
The critical distance  $d_c$  was used to generate the final datasets as the distance between seismic events. In order to determine the correlation of this parameter with the experimental results produced, another experiment was conducted where this distance ranged from 2 to 20km. In this experiment, the proposed classification rule generation technique was used. The experimental results from this experiment are presented in Table 5. The results indicate that the critical distance  $d_c$  influences performance, but the effect size is small. The average classification error decreases slightly as the distance increases: 19.08% for  $d_c = 2\text{km}$ , 19.00% for 5km, 18.82% for 10km, and 18.71% for 20km. The gap between the worst and best setting is 0.37 percentage points, i.e., about a 1.9% relative error reduction, so the overall advantage of the 20km setting is real but modest. Year-by-year,  $d_c = 20\text{km}$  achieves the lowest error in 4 out of 8 years (2004, 2005, 2009, 2010),  $d_c = 5\text{km}$  is best in 2008,  $d_c = 10\text{km}$  in 2007, and  $d_c = 2\text{km}$ , indicating no monotonic or universally optimal choice across years but a mild preference toward larger distances. Regarding stability,  $d_c = 2\text{km}$  shows the narrowest range over time (17.51%–20.70%), whereas  $d_c = 20\text{km}$  exhibits the largest spread mainly due to 2011 (16.97%–21.78%). Overall, the method benefits marginally from increasing  $d_c$ .

with 20km yielding the lowest average error, but the choice should also consider temporal stability and yearly idiosyncrasies, since the per-year optimum shifts across settings.

**Table 5.** Experiments with the GENCLASS method and different values of critical distance  $d_c$

YEAR	$d_c = 2km$	$d_c = 5km$	$d_c = 10km$	$d_c = 20km$
2004	18.69%	18.92%	18.54%	18.44%
2005	18.47%	19.56%	18.26%	17.56%
2006	17.51%	17.19%	17.38%	18.48%
2007	20.03%	18.14%	18.06%	18.20%
2008	20.70%	19.51%	20.69%	20.14%
2009	17.90%	17.81%	17.39%	16.97%
2010	19.21%	19.89%	19.32%	18.08%
2011	20.09%	20.97%	20.91%	21.78%
<b>AVERAGE</b>	<b>19.08%</b>	<b>19.00%</b>	<b>18.82%</b>	<b>18.71%</b>

The significance analysis across critical distance settings  $d_c$  provides no statistical evidence of an effect on classification error (Figure 6). The omnibus Friedman test is non-significant ( $p = 0.4153$ ), indicating no systematic differences across  $d_c$  levels. Pairwise contrasts corroborate this:  $d_c = 2km$  vs 5km ( $p = 0.9803$ ), 10km ( $p = 0.5274$ ), and 20km ( $p = 0.5274$ ) are not significant, nor are  $d_c = 5km$  vs 20km ( $p = 0.7675$ ) and  $d_c = 10km$  vs 20km ( $p = 1$ ). The  $d_c = 5km$  vs 10km comparison returned  $p = NA$ , which typically reflects a degenerate post-hoc case (e.g., complete rank ties or identical values per block) and does not alter the overall conclusion. In sum, within the 2–20km range,  $d_c$  does not yield statistically significant performance differences for the proposed method, so this hyperparameter may be selected based on practical or stability considerations rather than expected accuracy gains.



**Figure 6.** Statistical comparison for the results obtained by the application of the GENCLASS method, using a variety of values for the critical distance  $d_c$ .

### 3.3. Experiments with the number of generations $N_g$

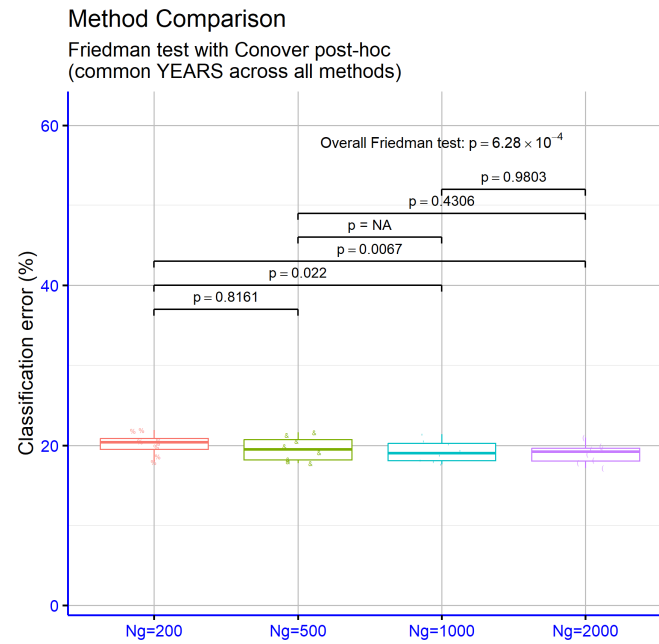
An additional experiment was conducted to verify the stability of the proposed classification rule generation technique. In this experiment, the maximum number of generations

ranged from 200 to 2000 and the experimental results per year are presented in Table 6. In this table, the results show a clear downward trend in classification error as the maximum number of generations increases. The average error decreases from 20.20% ( $N_g = 200$ ) to 19.58% ( $N_g = 500$ ), 19.25% ( $N_g = 1000$ ) and 19.00% ( $N_g = 2000$ ), i.e., a total gain of 1.20 percentage points or roughly a 6% relative reduction compared to  $N_g = 200$ . Improvements exhibit diminishing returns: the largest drop is from 200 to 500 ( $-0.62$ ), followed by 500 to 1000 ( $-0.33$ ) and 1000 to 2000 ( $-0.25$ ). On a per-year basis,  $N_g = 2000$  is best in six out of eight years (2006-2011 except 2004-2005), while in 2004-2005 the minimum error occurs at  $N_g = 1000$ . The 2005 value for  $N_g = 2000$  is noticeably higher than the other settings, which may reflect stochastic variability or model over-specialization for that year's data. The spread across years is broadly similar across settings, so the main benefit of increasing generations is a lower mean error rather than a dramatic change in variability. In practice, the 1000-2000 range offers the strongest performance,  $N_g = 1000$  comes very close to  $N_g = 2000$  (0.25 points apart) and is attractive under tighter computational budgets, whereas  $N_g = 2000$  yields the lowest average error when runtime is not a constraint.

**Table 6.** Experiments with the GENCLASS method and different values for the number of generations  $N_g$

YEAR	$N_g = 200$	$N_g = 500$	$N_g = 1000$	$N_g = 2000$
2004	19.85%	19.14%	18.76%	18.92%
2005	17.94%	17.80%	17.63%	19.56%
2006	20.33%	18.01%	17.95%	17.19%
2007	20.51%	19.91%	19.37%	18.14%
2008	21.83%	21.31%	20.16%	19.51%
2009	18.61%	18.25%	18.15%	17.81%
2010	20.59%	20.57%	20.55%	19.89%
2011	21.90%	21.63%	21.39%	20.97%
<b>AVERAGE</b>	<b>20.20%</b>	<b>19.58%</b>	<b>19.25%</b>	<b>19.00%</b>

The significance levels in Figure 7 indicate that the maximum number of generations has an overall effect on performance, as evidenced by a strongly significant Friedman test ( $p = 6.28 \times 10^{-4}$ ). In pairwise terms, moving from  $N_g = 200$  to  $N_g = 1000$  yields a statistically significant error reduction ( $p = 0.022$ ), and the contrast between  $N_g = 200$  and  $N_g = 2000$  is even more significant ( $p = 0.0067$ ), confirming that a low generation budget underperforms relative to higher budgets. By contrast,  $N_g = 200$  vs  $N_g = 500$  is not significant ( $p = 0.8161$ ), nor are  $N_g = 500$  vs  $N_g = 2000$  ( $p = 0.4306$ ) and  $N_g = 1000$  vs  $N_g = 2000$  ( $p = 0.9803$ ), suggesting diminishing returns beyond roughly 1000 generations. The  $N_g = 500$  vs  $N_g = 1000$  comparison returned  $p = \text{NA}$ , typically due to a degenerate post-hoc scenario (e.g., complete rank ties or identical per-block values) and does not alter the main conclusion. Overall, increasing generations above 200 significantly improves performance, with gains saturating around 1000 generations and no clear statistical advantage of 2000 over 1000.



**Figure 7.** Statistical comparison for the results obtained by the usage of the GENCLASS method, using different values for the maximum number of generations  $N_g$ .

#### 4. Conclusions

This study introduces an innovative methodology for earthquake classification employing various machine learning methods, including RBF, MLP, BAYES, NNC, along with the proposed GENCLASS approach. The analysis covers seismic data recorded between 2004-2011, within the geographical bounds of latitude  $21^\circ$ - $79^\circ$  and longitude  $33^\circ$ - $176^\circ$ . Our first process involved classifying the size of the seismic events into two large classes: The first class contains all the events with magnitude  $\leq 3$ . The second class contains all the remaining events, with magnitude  $> 3$ , followed by magnitude prediction based on the classified data.

To construct the final datasets, the critical distance  $d_c$  defined as the distance between seismic events, was applied. The results indicate that the proposed method benefits slightly from increasing  $d_c$ , with a value of 20km yielding the optimal average error. Also we Set as  $N_c$  the number of chromosomes in the genetic population. Furthermore, an additional experiment was conducted to evaluate the stability of the proposed generations mechanism  $N_g$ . In this test, the maximum number of generations was varied from 200-2000, indicating that the maximum number of generations has an overall effect on performance. Among all evaluated methods, GENCLASS consistently achieved the best performance each year, attaining the lowest average error rate of 19%, corresponding to an overall classification accuracy of 81%.

The main challenges encountered in this research were related to the large volume of data, exceeding one million records, and the adaptation of such data into a reliable classification system. In general, seismology still keeps many of its secrets well guarded, but not indefinitely. With the rapid advancement of machine learning and artificial intelligence techniques, these mysteries are gradually being unraveled. Future work in this area of research could include recent seismic events as well as the use of other advanced computational intelligence techniques such as feature construction from existing ones [64]. Furthermore, since Grammatical Evolution is essentially a genetic algorithm, parallel computing techniques can be used to speed up this process, such as the OpenMP method [65] or the OpenMPI programming library [66].

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