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Constructing artificial features with Grammatical Evolution for earthquake prediction

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Abstract

Over the course of centuries, humanity has evolved, acquired knowledge, and developed an understanding of the geological phenomenon known as the earthquake. Earthquakes are not the result of the wrath of mythological beings, but rather of the dynamic processes occurring beneath the Earth's crust specifically, the movement and interaction of tectonic / lithospheric plates. When one plate shifts relative to another, stress accumulates and is eventually released as seismic energy. This process is continuous and unstoppable. This phenomenon is well recognized in the Mediterranean region, where significant seismic activity arises from the northward convergence (4–10 mm per year) of the African plate relative to the Eurasian plate along a complex plate boundary. Consequently, our research will focus on the Mediterranean region, specifically examining seismic activity from 1990 - 2015 within the latitude range of 33–44° and longitude range of 17–44°. These geographical coordinates encompass 28 seismic zones, with the most active areas being Turkey and Greece. In this paper we applied Grammatical Evolution for artificial feature construction in earthquake prediction, evaluated against machine learning approaches including MLP(GEN), MLP(PSO), SVM, and NNC. Experiments showed that Feature Construction (FC) achieved the best performance, with a mean error of 9.05% and overall accuracy of 91%, outperforming SVM. Further analysis revealed that a single constructed feature (Nf = 1) yielded the lowest average error (8.21%), while varying the number of generations indicated that Ng = 200 provided an effective balance between computational cost and predictive accuracy. These findings confirm the efficiency of FC in enhancing earthquake prediction models through artificial feature construction. Our results, as will be discussed in greater detail within the research, yield an average error of approximately 9%, corresponding to an overall accuracy of 91%.

Keywords: Earthquakes; Machine learning; Neural networks; Grammatical Evolution; Feature Construction

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1. Introduction

When entering the single keyword "earthquakes" into Google Scholar, more than 3,910,000 results are retrieved, demonstrating the intense interest that exists in the field of seismology. Thanks to these studies and investigations, which evolve from an initial idea or theory into practical applications, it can be stated with confidence that humanity is now capable of achieving timely early warning before seismic events occur. Consequently, the pursuit of sustainability strengthens our resilience against seismic phenomena. This has

been achieved through the implementation of early warning systems established across the globe, particularly in technologically advanced countries that are also seismically vulnerable regions. For this purpose, the UN Disaster Risk Reduction was established, which is "aimed at preventing new and reducing existing disaster risk and managing residual risk, all of which contribute to strengthening resilience and therefore to the achievement of sustainable development "[1]. An early achievement of Disaster Risk Reduction took place in Japan in 1960, when seismic sensors were installed along the railway infrastructure to ensure the automatic immobilization of trains [2]. The Japanese UrEDAS (Urgent Earthquake Detection and Alarm System) has been described as the "grandfather" of earthquake early warning systems in general, and of onsite warning systems in particular [3]. Since then, techniques and methods have advanced through technological progress. The next achievement in early earthquake warning was accomplished in Mexico in 1989 with the establishment of the Seismic Alert System (SAS) [4]. In 2006, Japan launched the Earthquake Early Warning system initially for a limited audience and subsequently for the general public, in order to ensure the effectiveness of EEW in disaster mitigation [5]. This allows an earthquake warning to be disseminated between several seconds and up to one minute prior to the occurrence of the event [6]. In Bucharest, Romania, an earthquake warning system, in 1999, was also developed, providing a preparation window of 25 seconds [7]. Also in Istanbul, in preparation for the anticipated earthquake, an early warning system was implemented in 2002 [8]. In Southern Europe, at the University of Naples in Italy, a software model called PRESTo (PRobabilistic and Evolutionary early warning SysTem) was developed, designed to estimates of earthquake location and size within 5–6 seconds [9]. Furthermore, the United States, through the U.S. Geological Survey, has established its own earthquake warning system. Since 2016, the ShakeAlert system has been operational along the West Coast [10,11]. Subsequently, a map is presented in Figure 1, illustrating Earthquake Early Warning Systems worldwide, with colors indicating the operational status of each system, provided in [12]. Purple denotes operative systems that provide warnings to the public, black represents systems currently undergoing real-time testing, and gray is used for countries where feasibility studies are still in progress.

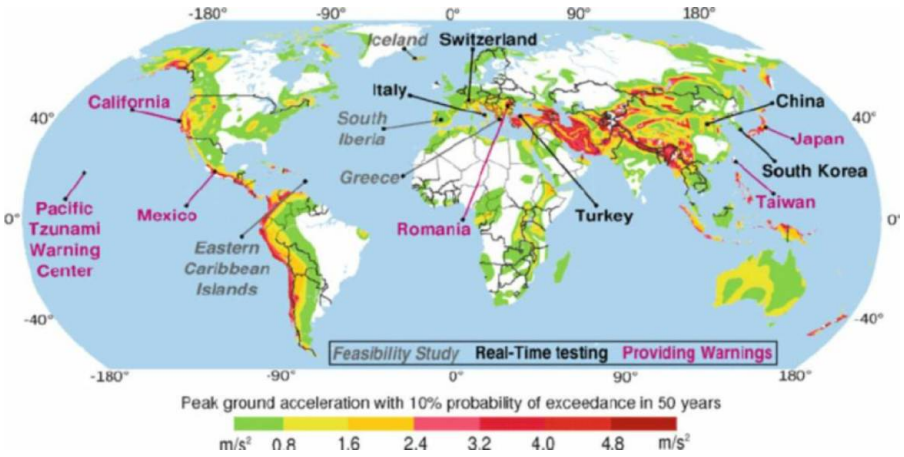


Figure 1. The map shows Earthquakes Early Warning Systems around the World

Within this framework, it is important to highlight that in recent years considerable emphasis has been placed on the advancement of diverse models for the early detection of seismic events, which have become available to the wider public through mobile applications, television broadcasts, and radio communication [13]. The primary function of Android applications is that, once an earthquake is detected, an alert is transmitted to all smartphones located within the affected area. Provided that the user is not in close proximity to the epicenter, the notification can be received in advance, allowing sufficient

time to take protective action before the destructive seismic waves arrive [14–17]. By harnessing technology as an ally against natural disasters, humanity can move beyond the devastating consequences of major earthquakes, such as the 2004 Indian Ocean event with more than 220,000 fatalities, the 2011 Tōhoku earthquake in Japan with over 19,000 losses, and the 2023 Turkey–Syria earthquake with more than 43,000 deaths. Accordingly, resilience and sustainability for populations affected by seismic events encompass both physical and social infrastructures capable of withstanding earthquakes, while simultaneously safeguarding long-term well-being through disaster risk reduction, community preparedness, and equitable recovery.

Subsequently, we will proceed with the presentation of related studies alongside our own, which progressively enhance both our sustainability and our capacity for prevention against seismic phenomena. Housner, in 1964, concluded that artificial earthquakes constitute adequate representations of strong-motion events for structural analysis and may serve as standard ground motions in the design of engineering structures [18]. Adeli, in 2009, proposed a novel feature extraction technique, asserting that when combined with a selected Probabilistic Neural Network (PNN), it can yield reliable prediction outcomes for earthquakes with magnitudes ranging from 4.5 to 6.0 on the Richter scale [19]. Zhou, in 2012, introduced a robust feature extraction approach, the Log-Sigmoid Frequency Cepstral Coefficients (LSFCC), derived from the Mel Frequency Cepstral Coefficients (MFCC), for the classification of ground targets using geophones. Employing LSFCCs, the average classification accuracy for tracked and wheeled vehicles exceeds 89% across three distinct geographical settings, achieved with a single classifier trained in only one of these environments [20]. Martinez-Alvarez, in 2013, investigated the utilization of various seismicity indicators as inputs for artificial neural networks. The study proposes combining multiple indicators—previously shown to be effective across different seismic regions—through the application of feature selection techniques [21]. Schmidt, in 2015, proposed an efficient and automated method for seismic feature extraction. The central concept of this approach is to interpret a two-dimensional seismic image as a function defined on the vertices of a carefully constructed underlying graph [22]. Narayanakumar, in 2016, extracted seismic features from a predetermined number of events preceding the main shock in order to perform earthquake prediction using the Backpropagation (BP) neural network technique [23]. Cortes, in 2016, sought to identify the parameters most effective for earthquake prediction. As various studies have employed different feature sets, the optimal selection of features appears to depend on the specific dataset used in constructing the model [24]. Asim, in 2018, developed a hybrid embedded feature selection approach designed to enhance the accuracy of earthquake prediction [25]. Chamberlain, in 2018, demonstrated that synthetic seismograms, when applied with matched-filter techniques, enable the detection of earthquakes even with limited prior knowledge of the source [26]. Okada, in 2018, employed observational data either by calibrating parameters within existing models or by deriving models and indicators directly from the data itself [27]. Lin, in 2018, employed the earthquake catalogue from 2000 to 2010, comprising events with a Richter magnitude (ML) of 5 and a depth of 300 km within the study area (21°–26° N, 119°–123° E). This dataset was utilized as training input to develop the initial earthquake magnitude prediction backpropagation neural network (IEMPBPNN) model, which was designed with two hidden layers [28]. Zhang, in 2019, proposed a precursory pattern-based feature extraction approach aimed at improving earthquake prediction performance. In this method, raw seismic data are initially segmented into fixed daily time intervals, with the magnitude of the largest earthquake within each interval designated as the main shock [29]. Rohas's, in 2019, paper reviewed the latest uses of artificial neural networks for automated seismic-data interpretation, focusing especially on earthquake detection and onset-time estimation [30]. Ali, in

2020, generated synthetic seismic data for a three-layer geological model and analyzed using Continuous Wavelet Transform (CWT) to identify seismic reflections in both the temporal and spatial domains [31]. Bamer, in 2021, demonstrated through comparison with several state-of-the-art studies, that the convolutional neural network autonomously learns to extract the pertinent input features and structural response behavior directly from complete time histories, rather than relying on a predefined set of manually selected intensity measures [32]. Wang, in 2023, reports that the accuracies of various AI models using the feature extraction dataset surpassed those obtained with the spectral amplitude dataset, demonstrating that the feature extraction approach more effectively emphasizes the distinctions among different types of seismic events [33]. Ozkaya, in 2024, developed a novel feature engineering framework that integrates the Butterfly Pattern (BFPat), statistical measures, and wavelet packet decomposition (WPD) functions. The proposed model achieved an accuracy of 99.58% in earthquake detection and 93.13% in three-class wave classification [34]. Sinha's review, in 2025, offers valuable insights into cutting-edge techniques and emerging directions in feature engineering for seismic prediction, highlighting the importance of interdisciplinary collaboration in advancing earthquake forecasting and reducing seismic risk [35]. Mahmoud, in 2025, investigates the application of machine learning approaches to earthquake classification and prediction using synthetic seismic datasets [36].

In contrast to the aforementioned studies, this paper introduces a novel approach constructing artificial features with Grammatical Evolution [37] for earthquake prediction, marking the first application of this evolutionary technique in the field of seismology. While the method does not directly address a specific gap in the literature, the experiments demonstrated that feature construction achieved an overall accuracy of 91%, thereby contributing to the advancement of earthquake prediction research. We consider this contribution significant in enhancing the understanding of seismic phenomena and in highlighting the potential of Grammatical Evolution as a promising tool for feature engineering in geophysical datasets. In particular, the method of Grammatical Evolution can be considered as a genetic algorithm [38] with integer chromosomes. Each chromosome contains a series of production rules from the provided Backus-Naur form (BNF) grammar [39] of the underlying language and hence this method can create programs that belong to this language. This procedure have been used with success in various cases, such as data fitting problems [40,41], problems that appear in economics [42], computer security problems [43], problems related to water quality [44], problems appeared in medicine [45], evolutionary computation [46], hardware issues in data centers [47], solution of trigonometric problems [48], automatic composition of music [49], dynamic construction of neural networks [50,51], automatic construction of constant numbers [52], playing video games [53,54], problems regarding energy [55], combinatorial optimization [56], security issues [57], automatic construction of decision trees [58], problems in in electronics [59], automatic construction of bounds for neural networks [60], construction of Radial Basis Function networks [61] etc. This research work focuses on the creation of artificial features from existing ones, aiming at two goals: on the one hand, it seeks to reduce the required information required for the correct classification of seismic data and on the other hand, it seeks to highlight the hidden nonlinear correlations that may exist between the existing features of the objective problem. In this way, a significant improvement in the classification of seismic data will be achieved.

Beyond the Grammatical Evolution approach proposed in this study, which achieved an accuracy of 91%, we will also provide a concise comparison with results from related research that has employed alternative machine learning techniques in the field of seismology. In the study by Adeli et al. [19] the PNN model demonstrated satisfactory predictive accuracy for earthquakes with magnitudes between 4.5 and 6.0, but its performance de-

clined considerably for events exceeding magnitude 6.0. Zhou et al. [20] reported that the application of the Log-Sigmoid Frequency Cepstral Coefficients method achieved an average classification accuracy exceeding 89%. Narayanakumar et al.[23] investigated the prediction of moderate earthquakes (magnitude 3.0–5.8). While the seismometer recorded an event of magnitude 4.0, the BP-ANN model predicted magnitudes in the range of 3.0–5.0, achieving a success rate between 75% and 125%. According to Asim et al., [25] the SVR-HNN prediction model achieved its highest performance in Southern California, with MCC, R score, and accuracy values of 0.722, 0.623, and 90.6%, respectively. The Chilean region ranked second, yielding an MCC of 0.613, an R score of 0.603, and an accuracy of 84.9%. In contrast, the Hindukush region exhibited the lowest performance, with MCC, R score, and accuracy values of 0.600, 0.580, and 82.7%, respectively. Chamberlain et al. [26] reported that template-based detections produced 7,340 events, of which 3,577 were identified as duplicates, whereas the STA/LTA method yielded 682 detections over the same time interval. Lin et al. [28] reported that the average magnitude error in Taiwan was $\Delta ML = \pm 0.3$, while the training errors of the EEMPBPNN model (<0.25) remained below this average error threshold. Zhang et al. [29] demonstrated that the proposed method achieved prediction accuracies of 93.26% and 92.07% across the two datasets examined. Rojas et al. [30] reported recognition performance of approximately 99.2% for P-wave signals and 98.4% for pure noise. Ali et al. [31] conducted a statistical analysis using a 95% confidence interval for the normalized CWT coefficients of P-wave velocity, seismic trace, acoustic impedance, and synthetic seismic trace. Wang et al. [33] found that, in the model generalization evaluation, the two-class models trained on the 36-dimensional network-averaged dataset achieved test accuracies and F1-scores exceeding 90%. Ozkaya et al. [34] reported that their model achieved an accuracy of 99.58% in earthquake detection and 93.13% in three-class wave classification. While comparable studies have demonstrated strong performance, many of them are limited to specific regions of interest or focus on particular signal types such as acoustic wave detection. In contrast, the Grammatical Evolution approach presented here not only achieves competitive accuracy (91%) but also demonstrates broader applicability across diverse seismic datasets, thereby highlighting its potential as a more generalizable solution in earthquake event discrimination.

The rest of this manuscript is organized as follows: the used dataset and the incorporated methods used in the conducted experiments are outlined in section 2, the experimental results are shown and discussed in section 3 and finally a detailed discussion is provided in section 4.

2. Materials and Methods

In this section, a detailed presentation of the datasets used as well as the machine learning techniques used in the experiments performed will be provided.

2.1. The Dataset Employed

In this study, we made use of open data from the NSF Seismological Facility for the Earth Consortium (SAGE) available from https://ds.iris.edu/(accessed on 25 November 2025), which is a platform offering an interactive global map that facilitates both data visualization and the real-time extraction of datasets from the displayed geographic regions. The area defined by latitudes 33°–44° and longitudes 17°–44° covers 28 seismic zones, with Turkey and Greece identified as the most seismically active regions.

Regarding the 28 seismic zones listed in Table 1, these correspond to specific geographic coordinates (latitudes and longitudes) provided through the interactive seismic platform previously referenced. The platform grants direct access to these spatial data,

ensuring that the delineation of the zones is both systematic and consistent with the established geospatial framework.

Table 1. Regions codes for the 28 seismic zones.

Region	Region Code
Turkey	1
Crete Greece	2
Greece	3
Dodecanese Islands, GREECE	4
Ionian Sea	5
Aegean Sea	6
Southern Greece	7
GREECEALBANIA border region	8
GREECEBULGARIA border region	9
Cyprus region	10
Albania	11
NORTHWESTERN Balkan Region	12
Central Mediterranean sea	13
Jordan Syria region	14
GEORGIA ARMENIA TURKEY Border Region	15
TURKEY IRAN border region	16
Iraq	17
Eastern Mediterranean sea	18
NORTHWESTERN CAUCASUS	19
Black Sea	20
Romania	21
Bulgaria	22
Crimea Region Ukraine	23
ARMENIA AZERBAIJAN IRAN border region	24
Adriatic sea	25
UKRAINE MOLDOVA RUSSIA border region	26
Southern Italy	27
IRAN IRAQ border region	28

Furthermore, the selection of NSF data was driven by its advanced functionality and broad accessibility. Notably, it supports the download of up to 25,000 records per file, thereby streamlining the workflow and improving the efficiency of information retrieval. The region under investigation is presented in Figure 2.

Figure 2. The study area.

2.2. Dataset Description

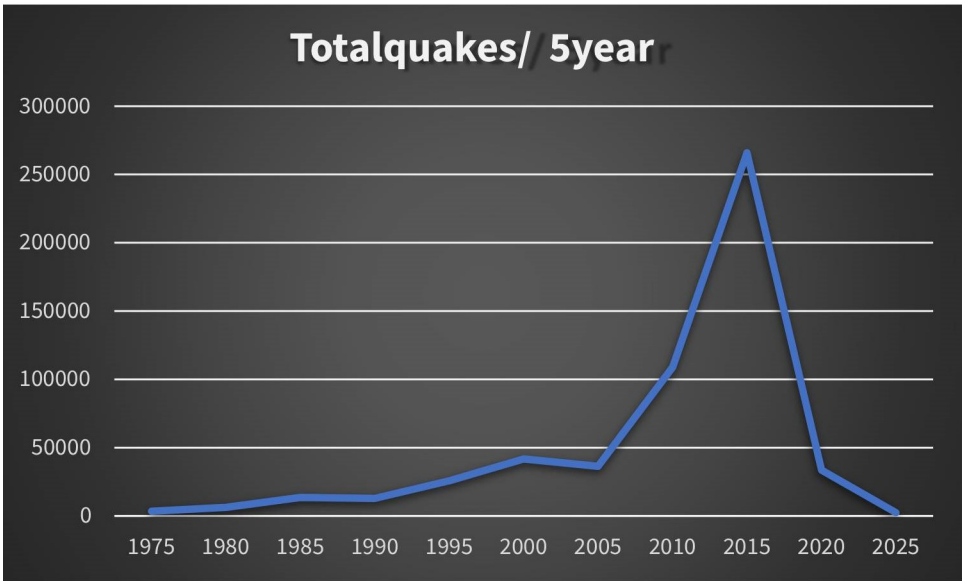
We obtained and systematically analyzed 511,064 earthquake events from 1990 to 2015, as this time period is of particular scientific interest due to the surge in seismic activity, as also illustrated in the graph of Figure 3. Specifically, we selected this time period because it encompasses a wide range of earthquake magnitudes, which provides a diverse dataset conducive to algorithm training and supports the construction of artificial features via Grammatical Evolution.

In the years under examination, earthquakes are distributed across all magnitude classes, exhibiting an almost ideal proportion consistent with the Gutenberg–Richter law [63]. For instance, within the first two five-year periods of our dataset, 2 events above magnitude 8.0, 3 events above magnitude 7.0, 19 events above magnitude 6.0, 165 events between 5.0–5.9, 3,048 events between 4.0–4.9, and 26,301 events between 3.0–3.9 were recorded, thereby confirming the expected logarithmic relationship between frequency and magnitude. Such a balanced and representative dataset further enhances the effectiveness of the Grammatical Evolution approach, as it provides a comprehensive distribution of seismic events across the full spectrum of magnitudes. Moreover, our analysis led us to the conclusion that the datasets from 1970–1989 and 2016–2025 lack the diversity observed in the dataset we selected. Specifically, classes 6 and 7 are entirely absent from the 1970–1989 records, while in the post-2016 data class 7 is missing and class 1 contains only a limited number of instances, namely 25.

Table 2. Raw Data from NSF Interactive Earthquake Browser (1990 - 2015)

Raw Data	
Features	**Range**
Year	1990 - 2015
Month	1 - 12
Day	1- 31
Time	00:00:00 - 23:59:59
Latitude	33 - 44
Longitude	17 - 44
Region	1 - 28
Depth	0.00 - 800.00
Magnitude	1 - 10
Timestamp	

Figure 3. Graphs seismic events from 1970 - 2025 (Area Study)



On the platform that provides us with the Interactive Earthquake Browser we employed coordinates spanning latitudes 33°–44° and longitudes 17°–44°, considered magnitude values from 1.0 to 10.0, and incorporated all available depths by default within the depth range. The raw dataset included the following variables: year, month, day, time, latitude (Lat), longitude (Lon), depth, magnitude, region, and timestamp. Accordingly, Table 2 provides a detailed overview of the raw dataset.

Subsequently, a preprocessing procedure was applied to the dataset. This included the identification of the lithospheric plate associated with each earthquake, to which a unique code was assigned. Furthermore, the months were categorized according to the four seasons, the days were grouped into ten-day intervals, and the time of occurrence was classified into four periods (morning, noon, afternoon, and night). The focal depth was divided into six categories. In addition, a new column was created to indicate, with a binary value (0 or 1), whether an earthquake had previously occurred in the same region during the same season. Finally, the dataset was merged with the Kp index, representing geomagnetic storm activity, which was further classified into six distinct categories. The final dataset was further processed, including the following: Year, Epoch Code, Day Code, Time Code, Latitude, Longitude, Depth Code, Previous Magnitude Code, Same Region Code, Lithospheric/Tectonic Plate, Kp Code. This information is outlined in Table 3.

Table 3. Utilized Data from NSF Interactive Earthquake Browser (1990 - 2015)

Utilized Data		
Features	**Range**	**Class**
Year	1990 - 2015	
Epoch Code	1 - 12	0 -3
Day Code	1 - 31	0 - 2
Time Code	00:00:00 - 23:59:59	0 - 3
Latitude	33 - 44	
Longitude	17 - 44	
Depth Code	0.00 - 800.00	0 - 5
Previous Magnitude Code	1 - 10	1 - 7
Same Region Code	1 - 28	1 - 28
Lithospheric Code	1 - 7	1 - 7
Kp Code	0.000 - 9.000	0 - 5

In the processed dataset, categorical classes were introduced to facilitate the analysis. Specifically, the months were initially ordered numerically and subsequently grouped according to the corresponding season. Thus, class 0 represents the winter months, class 1 corresponds to the spring months, class 2 to the summer months, and class 3 to the autumn months. The days comprising each month were divided into three ten-day intervals, with the first interval assigned to class 0, the second interval to class 1, and the final interval to class 2. Accordingly, the hours of the 24 hour day were divided into distinct periods, with the morning zone assigned to class 0, midday to class 1, afternoon to class 2, and night to class 3. For the depth code, earthquakes occurring near the surface (0–32.9 km) were assigned to class 0, those recorded at depths between 33–69.9 km to class 1, events within 70–149.9 km to class 2, those between 150–299.9 km to class 3, earthquakes at depths of 300–499.9 km to class 4, those between 500–799.9 km to class 5, and finally, events occurring at depths greater than 800 km were assigned to class 6. Continuing, earthquakes with magnitudes below 2.9 were assigned to class 1, those with magnitudes between 3.0–3.9 to class 2, magnitudes between 4.0–4.9 to class 3, magnitudes between 5.0–5.9 to class 4, magnitudes between 6.0–6.9 to class 5, magnitudes between 7.0–7.9 to class 6, while events with magnitudes of 8.0 and above were assigned to class 7. With regard to the categorization of the geographical region and the lithospheric plate, each region was assigned a code ranging from 1 to 28. Similarly, a code from 1 to 7 was allocated to the lithospheric plates corresponding to each seismic event. Finally, the Kp code was classified as follows: values ranging from 0.000–4.000 were assigned to class 0, those from 4.100–5.000 to class 1, values between 5.100–6.000 to class 2, those from 6.100–7.000 to class 3, values between 7.100–8.000 to class 4, and finally, values from 8.100–9.000 were assigned to class 5.

At the following stage of data processing, we elected to focus on earthquakes with a magnitude code of 2 and above, since the inclusion of lower-magnitude events would bias the model toward predicting minor seismic occurrences. Notably, the small-magnitude category (1–2.9 mag) prevailed as the majority class with 407,144 records, creating a significant imbalance in the dataset. By excluding this dominant class, we aimed to achieve a more balanced distribution across magnitudes and to enhance the representational capacity of the model. This approach is consistent with previous studies that have similarly excluded small earthquakes to mitigate bias and improve predictive performance. Wang et al., in 2023, reports in Dataset and Feature Engineering " The seismic catalog used in this study was obtained from the China Earthquake Data Center (CEDC, http://data.earthquake.cn/, last accessed on 12 December 2025) and includes earthquake events with a magnitude greater than 3.0 in the Sichuan–Yunnan region from 1970 to 2021" [62]. It is noteworthy that the division into two classes within the field of seismology has previously been employed,

as in the study of Zhang et al. (2023), which investigated seismic data by categorizing events into high magnitude ($M \geq 5.5$) and low magnitude ($M < 5.5$) [64]. Following these steps, we proceeded with our experiments, utilizing approximately 10,000 seismic events in order to generate artificial features through Grammatical Evolution.

In summary, the seismic data employed in this study were recorded on the magnitude scale, covering events from 1 to 10. Initially, we excluded small earthquakes, which predominated in number and affected the balance of the dataset. The data source was the EarthScope Consortium, which operates the NSF Geodetic Facility for the Advancement of Geoscience (GAGE) and the NSF Seismological Facility for the Advancement of Geoscience (SAGE). These records were obtained from thousands of stations worldwide, constituting the primary NSF SAGE archive.

2.3. Global optimization methods for neural network training

In the current work two well - known global optimization methods were incorporated for neural network training [65,66], the Genetic Algorithm and the Particle Swarm Optimization (PSO) method. The Genetic algorithms is an evolutionary optimization method designed to minimize an objective function defined over a continuous search space. It operates on a population of candidate solutions, each represented as a vector of parameters and this population is evolved through a series of steps, where in each step some process that resemble natural processes are applied to the population. Genetic algorithms have applied on a wide range applications, that include training of neural networks [68,69]. Beyond neural network training, Genetic Algorithms have demonstrated strong performance in physics, biotechnology, and medical physics. In bioinformatics and biotechnology, they are commonly applied to the analysis of biological data, gene selection, and the optimization of complex biological systems [70]. In medical physics and biomedical engineering, GAs have been successfully used for medical image segmentation [71], diagnostic modeling, and the optimization of treatment parameters [72]. They have also been applied to multi-objective problems related to energy efficiency, economic viability, environmental life-cycle assessment [73] and parameter identification in complex multi-physics energy systems [74]. Also, Figure 4 presents the basic steps of a genetic algorithm.

Algorithm 1 The main steps of a Genetic Algorithm..

INPUT
- f: objective function
- N_c: number of chromosomes
- N_g: maximum number of allowed generations
- p_s: the selection rate of the algorithm, with $p_s \leq 1$
- p_m: the mutation rate of the algorithm, with $p_m \leq 1$
- k: the generation counter
- a: uniformly distributed random numbers, in $\in [-0.5, 1.5]$

OUTPUT
- x_{best}, f_{best}

INITIALIZATION
- $k \leftarrow 0$

main pseudocode

```
01 while k < N_g do // termination check
02     for each g_i, i ∈ {1..N_c} do
03         f_i ← f(g_i)
04     endfor
05     Sort chromosomes by increasing fitness: N_b ← (1 − p_s) x N_c
06     Select parents w, z randomly among the best N_b chromosomes
07     Draw a_i ∈ U[−0.5, 1.5]
08         z_i ← a_i z_i + (1 − a_i) w_i
09         w_i ← a_i w_i + (1 − a_i) z_i
10     foreach g_i, i ∈ {N_b + 1..N_c} do
11         Replace g_i with z_i or w_i
12     endfor
13     foreach g_i, i ∈ {1..N_c} do
14         foreach gene, j ∈ {1....n} do
15             Draw rU[0, 1]
16                 if r ≤ p_m then
17                     Mutate gene j of g_i
18                 endif
19         endfor
20     end for
21     for each g_i, i ∈ {1..N_c} do
22         f_i ← f(g_i)
23         Draw ri ∈ U[0, 1]
24         if (f_i < f_best) then
25             xbest ← gi, fbest ← f i
26         endif
27     endfor
28     k ← k + 1
29 endwhile
30 return x_best, f_best
```

Figure 4. The main steps of the Genetic algorithm.

Particle Swarm Optimization (PSO) is a population-based search method inspired by how animals move and cooperate in groups like flocks of birds or schools of fish [76,77]. The PSO was widely used in a variety of practical problems as well as in neural network training [78,79]. Beyond neural network training, Particle Swarm Optimization

has been applied across a broad spectrum of scientific and technological domains. In particular, it has found significant use in economic and resource optimization[80], as well as in healthcare management and medical physics, where it supports efficient medical service allocation and the management of stochastic patient queue systems[81]. At the same time, PSO has been increasingly adopted in physics and chemical engineering for the optimization of catalytic processes and energy systems, often in combination with deep learning approaches to extract meaningful physical and chemical insights[82]. In more dynamic and decentralized settings, PSO has proven especially effective in robotics, where it is used to coordinate swarms of autonomous robots during search-and-rescue missions operating under uncertain and rapidly changing conditions[83]. Finally, its flexibility has also led to successful applications in bioinformatics and computational biology, where advanced multi-objective PSO variants are employed for community detection and the identification of disease-related functional modules within complex biological networks[84]. The main steps of the PSO method are outlined in Figure 5.

INPUT
- f: objective function
- m: number of particles with $x_i \in S$
- u_i: number of velocities with $u_i \in S$
- x_i: number of positions in Ω
- w: inertia
- $c1$,$c2$: constant numbers
- $r1$,$r2$: random numbers
- $iter$: iteration counter
- $iter_{max}$: max iterations
- p_i: vectors are best located values for every particle i

OUTPUT
-p_{best}

INITIALIZATION
- **Set** $iter \leftarrow 0$
- **Set** positions $x_i \in S, i = 1.....m$
- **Set** velocities $u_i, i = 1.....m$
-**For** each particle $i \in \{1..m\}$ **do**
-$p_i \leftarrow x_i$
-**End for**
-$p_{best} \leftarrow argmin_i f(x_i)$// global best
01 **while** $generation < Gmax$ **do**// termination check
02 **For** each $i \in \{1..m\}$ **do**
03 draw $r1, r2 \backsim U(0, 1)$
04 $u_i \leftarrow wu + c1r1(p_i - x_i) + c2r2(p_{best} - x_i)$// update velocity
05 $x_i \leftarrow x_i + u_i$// update position
06 **if** $f(x_i) < f(p_i)$**then**
07 $p_i \leftarrow x_i$
08 **End if**
09 **End for**
10 $p_{best} \leftarrow argmin_i f(x_i)$// Update global best
11 $iter \leftarrow iter + 1$
12 **End while**

Figure 5. The main steps of the PSO algorithm.

2.4. The SVM method

The Support Vector Machine (SVM) is a supervised learning algorithm applied to both classification and regression problems [85]. The concept of Support Vector methods was introduced by V. Vapnik in 1965, in the context of addressing challenges in pattern recognition. During the 1990s, V. Vapnik formally introduced Support Vector Machine (SVM) methods within the framework of Statistical Learning. Since their introduction, SVMs have been extensively employed across diverse domains, including pattern recognition, natural language processing, and related applications [86]. For instance, the SVM algorithm has been employed both in studies on earthquake prediction and in research on the early detection of seismic phenomena, as demonstrated in the following works: Hybrid Technique Using Singular Value Decomposition (SVD) and Support Vector Machine (SVM) Approach for Earthquake Prediction [87], Using Support Vector Machine (SVM) with GPS Ionospheric TEC Estimations to Potentially Predict Earthquake Events [88], The efficacy of support vector machines (SVM) in robust determination of earthquake early warning magnitudes in central Japan [89]. As with other comparative analyses of models, SVMs

require greater computational resources during training and exhibit reduced susceptibility to overfitting, whereas neural networks are generally regarded as more adaptable and capable of scaling effectively. Beyond these applications, SVMs have also been widely adopted in several applied scientific and engineering fields. In bioinformatics, SVMs are commonly employed to analyze and interpret complex biological data, where their ability to handle high-dimensional feature spaces is particularly valuable[90]. In medical imaging and clinical applications, they have been used effectively for disease classification and staging, such as the prediction of lung cancer stages from medical imaging data[91]. In addition, SVM-based models are increasingly integrated into healthcare monitoring systems, including Internet of Medical Things (IoMT) frameworks, where reliability and robustness are critical requirements[92]. Moreover, SVMs have been applied in applied physics and engineering, including the prediction of electronic properties in advanced material structures and nanocomposites[93]. Their robustness has also made them suitable for safety-critical applications, such as fault detection and diagnosis in nuclear power plant systems[94].

2.5. The neural network construction method

Another method used in the experiments to predict the category of seismic vibrations is the method of constructing artificial neural networks using Grammatical Evolution [95]. This method can detect the optimal architecture for neural networks as well as the optimal set of values for the corresponding parameters. This method was used in various cases, such as chemistry problems [96], solution of differential equations [97], medical problems [98], educational problems [99], detection of autism [100] etc. This method can produce neural networks in the following form:

$$N(\vec{x}, \vec{w}) = \sum_{i=1}^{H} w_{(d+2)i-(d+1)} \sigma\left(\sum_{j=1}^{d} x_j w_{(d+2)i-(d+1)+j} + w_{(d+2)i} \right) \tag{1}$$

In this equation the value H represents the number of used computation units (weights) for the neural network. The vector w denotes the vector containing the parameters of the neural network and vector x represents the input vector (pattern). The function $\sigma(x)$ stands for the sigmoid function, which is defined as:

$$\sigma(x) = \frac{1}{1+\exp(-x)} \tag{2}$$

2.6. The proposed method

The method proposed here to tackle the classification of seismic events is a procedure that constructs artificial features from the original ones using Grammatical Evolution. The method was initially presented in the paper of Gavrilis et al [101] and used in various cases in the past [102–104]. The main steps of this method have as follows:

1. **Initialization** step.
   (a) **Obtain** the train data and denote them using the M pairs (x_i, t_i), $i = 1..M$. The values t_i represent the actual output for the input pattern x_i.
   (b) **Set** the parameters of the used genetic algorithm:N_g for as the number of allowed generations, N_c for the number of chromosomes, p_s for the selection rate and p_m for the mutation rate.
   (c) **Set** as N_f the number of artificial features that will construct the current method.
   (d) **Initialize** every chromosome $g_i, i = 1, \ldots, N_g$ as a set of randomly selected integers.

(e) **Set** $k = 1$, the generation counter.

2. **Genetic step**

(a) **For** $i = 1, \ldots, N_g$ **do**

i. **Construct** with Grammatical Evolution a set of N_f artificial features for the each chromosome g_i. The BNF grammar used for this procedure is shown in Figure 6.

ii. **Transform** the original set of features using the constructed features and denote the new training set as $(x_{g_i,j}, t_j)$, $j = 1, ..M$

iii. **Train** a machine learning C on the new training set. The training error for this model will represent the fitness f_i for the current chromosome and it is computed as:

$$f_i = \sum_{j=1}^{M} \left(C\left(x_{g_i,j}\right) - t_j\right)^2 \tag{3}$$

In the current work the Radial Basis Function networks (RBF) [105,106] were use d as the machine learning models. A decisive factor for this choice was the short training time required for these machine learning models.

iv. **Perform** the selection procedure: Initially the the chromosomes are sorted according to the associated fitness values and the best $(1 - p_s) \times N_C$ of them are copied intact to the next generation. The remaining chromosomes will be replaced by offsprings that will be produced during the crossover and the mutation procedure.

v. **Perform** the crossover procedure: The outcome of this process is a set of $p_s \times N_c$ new chromosomes. For every pair $(\tilde{z}, \tilde{w})$ of new chromosomes, two distinct chromosomes z and w are chosen from the current population using the process of tournament selection. Afterwards, the nee chromosomes are produced using the procedure of one - point crossover, that is illustrated in in Figure 7.

vi. **Execute** the mutation procedure: during this procedure a random number $r \in [0, 1]$ is selected for each element of every chromosome. The corresponding element is altered randomly if $r \leq p_m$.

(b) **EndFor**

3. **Set** $k = k + 1$
4. **If** $k \leq N_g$ goto **Genetic** Step
5. **Obtain the best chromosome** g^* from the current population.
6. **Create** the corresponding N_f features for g^* and apply this features to the set set and report the corresponding test error.

Also, a flowchart that presents the overall process of feature construction is presented in Figure 8.

Figure 6. The grammar used for the proposed method.

```
S::=<expr>   (0)
<expr> ::=  (<expr> <op> <expr>)  (0)
           | <func> ( <expr> )    (1)
           |<terminal>            (2)
<op> ::=     +      (0)
           | -      (1)
           | *      (2)
           | /      (3)
<func> ::=   sin  (0)
           | cos  (1)
           |exp   (2)
           |log   (3)
<terminal>::=<xlist>                (0)
           |<digitlist>.<digitlist> (1)
<xlist>::=x1    (0)
           | x2 (1)
           .........
           | xN (N)
<digitlist>::=<digit>                  (0)
           | <digit><digit>            (1)
           | <digit><digit><digit>     (2)
<digit>  ::= 0 (0)
           | 1 (1)
           | 2 (2)
           | 3 (3)
           | 4 (4)
           | 5 (5)
           | 6 (6)
           | 7 (7)
           | 8 (8)
           | 9 (9)
```

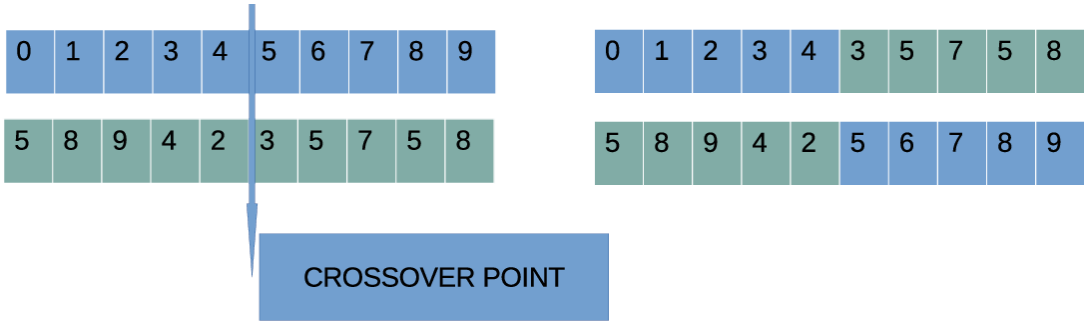


Figure 7. An example of the one - point crossover method.

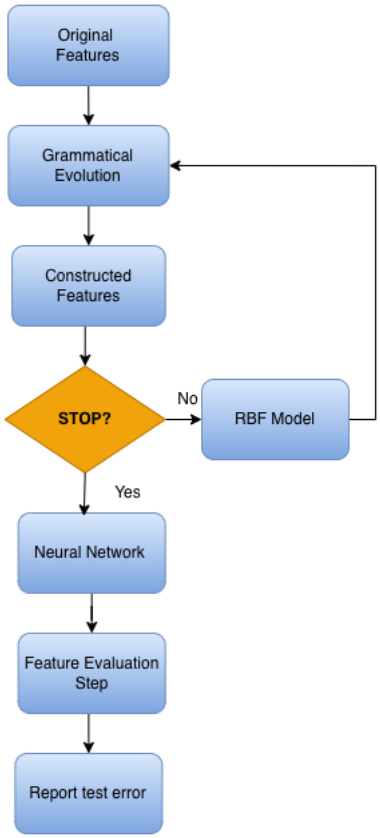


Figure 8. The steps of the feature construction procedure.

3. Results

The methods used in the conducted experiments were coded in ANSI C++, with the assistance of the freely available optimization package of OPTIMUS [107]. Moreover, for the feature construction procedure the freely available programming tool QFc, that can be downloaded from https://github.com/itsoulos/QFc(accessed on 12 December 2025)[108] has been used. Each experiment was conducted 30 times, using different seed for the random generator in each execution. Also, the procedure of ten - fold cross validation was incorporated to validate the conducted experiments. The values for the experimental parameters are listed in Table 4. The parameters in this table have been chosen to provide a compromise between speed and efficiency of the proposed procedure and were applied to all methods used in the experiments, so that there is fairness in the comparison of experimental results. The proposed method of feature construction was applied to each individual fold and averages of the classification errors in the control sets were obtained. In addition, the RBF machine learning model was used to construct the new features, due

to the significantly shorter learning time required for it compared to other models, such as artificial neural networks.

Table 4. The values of the experimental parameters.

PARAMETER	MEANING	VALUE
N_g	Number of generations	500
N_c	Number of chromosomes	500
p_s	Selection rate	0.9
p_m	Mutation rate	0.05
N_f	Number of features	2

Table 5. Experimental results using a series of machine learning methods.

DATASET	MLP(GEN)	MLP(PSO)	SVM	NNC	FC
GCT1990	39.98%	38.21%	14.30%	22.72%	13.50%
GCT1995	38.08%	33.22%	13.48%	21.50%	10.82%
GCT2000	26.70%	27.48%	10.19%	24.59%	6.52%
GCT2005	38.44%	33.19%	6.85%	27.85%	4.45%
GCT2010	52.89%	44.77%	14.46%	29.81%	9.97%
AVERAGE	**39.22%**	**35.37%**	**11.86%**	**25.29%**	**9.05%**

Table 5 reports the classification error rates for five temporal subsets of the seismic dataset (GCT1990, GCT1995, GCT2000, GCT2005, GCT2010), where the first column encodes the year of the data and the remaining columns correspond to the machine learning models MLP(GEN), MLP(PSO), SVM, NNC, and the proposed FC (Future Constructions) model. The values are expressed as percentage classification error, that is, the proportion of misclassified events between the two seismic classes defined during preprocessing (events with magnitude code 2–3 versus events with magnitude greater than 3). The following notation is used in this table:

1. The column MLP(GEN) denotes the error from the application of the genetic algorithm described in subsection 2.3 for the training of a neural network with 10 processing nodes.
2. The column MLP(PSO) represents the incorporation of the PSO method given in subsection 2.3 for the training of a neural network with 10 processing nodes.
3. The column SVM represent the application of the SVM method, described in subsection 2.4. In the current implementation the freely available library LibSvm [109] was used.
4. The column NNC represents the neural network construction method, described in subsection 2.5.
5. The column FC denotes the proposed feature construction technique, outlined in subsection 2.6.
6. The row AVERAGE denotes the average classification error for all datasets.

Also, Table 6 indicates the precision and recall values for SVM, NNC and the proposed method.

Table 6. Precision and recall values for the methods SVM,NNC and FC.

	SVM		NNC		FC	
DATASET	PRECISION	RECALL	PRECISION	RECALL	PRECISION	RECALL
GCT1990	0.798	0.837	0.684	0.728	0.800	0.874
GCT1995	0.811	0.830	0.684	0.729	0.801	0.886
GCT2000	0.835	0.844	0.662	0.700	0.835	0.901
GCT2005	0.856	0.869	0.628	0.627	0.856	0.904
GCT2010	0.802	0.824	0.633	0.628	0.804	0.805

The most striking observation is that FC achieves the lowest error for all five datasets, with no exceptions. For every GCT year, the FC column attains the minimum error among all models, demonstrating a consistent and temporally robust superiority. This pattern is clearly reflected in the AVERAGE row: the mean classification error of FC is 9.05%, whereas the corresponding mean error of the best conventional baseline, the SVM, is 11.86%. Moving from 11.86% to 9.05% corresponds to an error reduction of approximately 24-25%, meaning that roughly one quarter of the misclassifications made by the SVM are eliminated when using FC. Compared with the neural approaches, the gain is even more pronounced: FC reduces the mean error from 39.22% to 9.05% for MLP(GEN), from 35.37% to 9.05% for MLP(PSO), and from 25.29% to 9.05% for NNC, effectively absorbing about 64-77% of their errors.

Examining the behaviour per year reveals how the proposed model interacts with the specific characteristics of each temporal subset. Dataset GCT2005 appears to be the "easiest" case for all models, with the SVM dropping to 6.85% error and FC reaching 4.45%, which corresponds to an accuracy of about 95.5%. At the opposite end of the spectrum, GCT2010 is clearly the most challenging dataset, as indicated by substantially increased errors for the conventional models (MLP(GEN) 52.89%, MLP(PSO) 44.77%, NNC 29.81%) and by a higher error for the SVM (14.46%). Even in this difficult scenario, FC keeps the error below 10% (9.97%), preserving a clear qualitative advantage in the most demanding temporal setting. A similar pattern emerges for the intermediate years 1995 and 2000, where FC-SVM absolute differences are in the range of 2-4 percentage points, corresponding to roughly 20-35% relative error reduction. This suggests that as the data become more complex, the benefit of the automatically constructed features generated by Grammatical Evolution becomes increasingly pronounced.

A second important finding is that the ranking of the conventional models remains stable across all years. The SVM is consistently the best among the standard baselines, followed by NNC and the two MLP variants, with MLP(PSO) systematically outperforming MLP(GEN). This stability reinforces the credibility of the table as a benchmark, indicating that the results are not driven by noise or random fluctuations but instead reflect a coherent hierarchy of model performance. On top of this stable baseline, FC does not merely swap the winner for a single dataset, it establishes a new performance level by consistently pushing the error rates to substantially lower values in every year.

From a practical standpoint, given that the final processed dataset contains on the order of 10,000 seismic events, an average error of 11.86% compared with 9.05% implies hundreds fewer misclassified instances when FC is used instead of a plain SVM trained on the original features. This has direct implications for early warning and risk assessment applications, where each reduction in misclassification translates into more reliable decision-making. The evidence from Table 5, combined with the methodological description, strongly supports the view that the feature construction phase based on Grammatical Evolution is not a minor refinement over existing classifiers, but rather a key component that reshapes the feature space so that the seismic classes become much more separable. This explains why the proposed FC model achieves an average error of approximately 9%, in full agreement with the reported overall accuracy of 91% in the abstract.

Overall, Table 6 shows that FC is not only the best-performing model in each individual year but also the most stable across different temporal subsets, confirming that the future-construction approach with Grammatical Evolution yields a substantial and consistent improvement over standard machine learning models for the discrimination of seismic events.

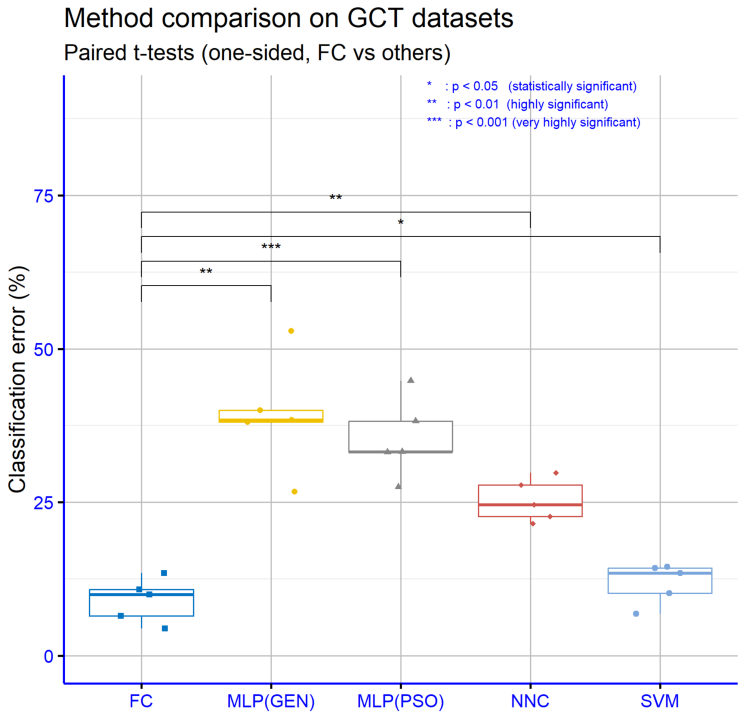


Figure 9. Comparison of FC and baseline classifiers on GCT datasets (paired one-sided t-tests on classification error)

The repeated-measures ANOVA with Method as the within-subject factor and DATASET (year) as the blocking factor confirms that the choice of classifier has a statistically significant effect on the classification error, indicating that the models are not equivalent in terms of predictive performance. Building on this global result, we conducted pairwise paired t-tests between the proposed FC model and each baseline, using one-sided alternatives that explicitly test the hypothesis that FC achieves lower mean error than its competitors. The resulting p-values, visualised as significance stars on the boxplot, show that the superiority of FC is statistically supported across all comparisons (Figure 9).

In particular, the comparisons FC vs MLP(GEN) and FC vs NNC yield ** ($p < 0.01$), indicating that the probability of observing differences of this magnitude in favour of FC purely by chance is below 1%. An even stronger effect emerges for FC vs MLP(PSO), which is marked with *** ($p < 0.001$), reflecting the very large performance gap between FC and the PSO-trained MLP. Finally, the comparison FC vs SVM is annotated with * ($p < 0.05$), showing that, although the numerical difference in error is smaller than in the neural baselines, it remains statistically significant in favour of FC. Overall, the pattern of significance codes (*, **, ***) corroborates the message of Table 6: FC is not only the best-performing model in terms of average classification error, but its advantage over all other methods is statistically significant, with particularly strong evidence against the MLP-based models and clear, though more moderate, evidence against the already very competitive SVM baseline.

Moreover, the proposed method can identify the hidden relationships between the features of the objective dataset. As an example of constructed features consider the following two features created for a distinct run on the GCT2010 dataset:

$$\begin{aligned} f_1(x) &= \sin((28.841/9.28) * x_{10} + (-78.60) * x_3 + 16.12 * x_7 + 6.3 * x_{11}) \\ f_2(x) &= \cos(458.6 * x_2 + (-18.809) * x_4) \end{aligned}$$

3.1. Experiments with the number of features

An additional experiment was conducted, where the number of constructed features was changed from 1 to 4 for the proposed feature construction method. The corresponding experimental results are outlined in Table 7.

Table 7. Experimental results using the proposed method and a series of values for the number of constructed features N_f.

DATASET	$N_f = 1$	$N_f = 2$	$N_f = 3$	$N_f = 4$
GCT1990	11.03%	13.50%	14.62%	15.40%
GCT1995	9.41%	10.82%	9.98%	10.05%
GCT2000	6.51%	6.52%	6.75%	6.54%
GCT2005	4.40%	4.45%	4.42%	4.38%
GCT2010	9.72%	9.97%	9.82%	10.32%
AVERAGE	**8.21%**	**9.05%**	**9.12%**	**9.34%**

Table 7 investigates the behavior of the proposed FC model as the number of constructed features N_f varies from 1 to 4. The results show that performance is generally very stable, with the mean classification error ranging within a narrow band between 8.21% (for N_f = 1) and 9.34% (for N_f = 4). The lowest average error is obtained with a single constructed feature (N_f =1), whereas the configuration N_f =2, which is adopted as the default setting in Table 5, yields a mean error of 9.05%, very close to the optimum and with highly consistent behaviour across all years. In some datasets, such as GCT2005, slightly better values appear for larger N_f (e.g., 4.38% for N_f = 4 versus 4.40% for N_f = 1), but these gains are marginal and do not change the overall picture. Taken together, the results indicate that FC does not require a large number of future constructions to perform well: one or two constructed features are sufficient to achieve very low error rates, while further increasing N_f does not lead to systematic improvements and likely introduces redundant information that does not translate into better generalization. This supports the view that the quality of the features generated by Grammatical Evolution is more important than their quantity, and that the proposed choice N_f = 2 offers a well-balanced compromise between performance and model simplicity.

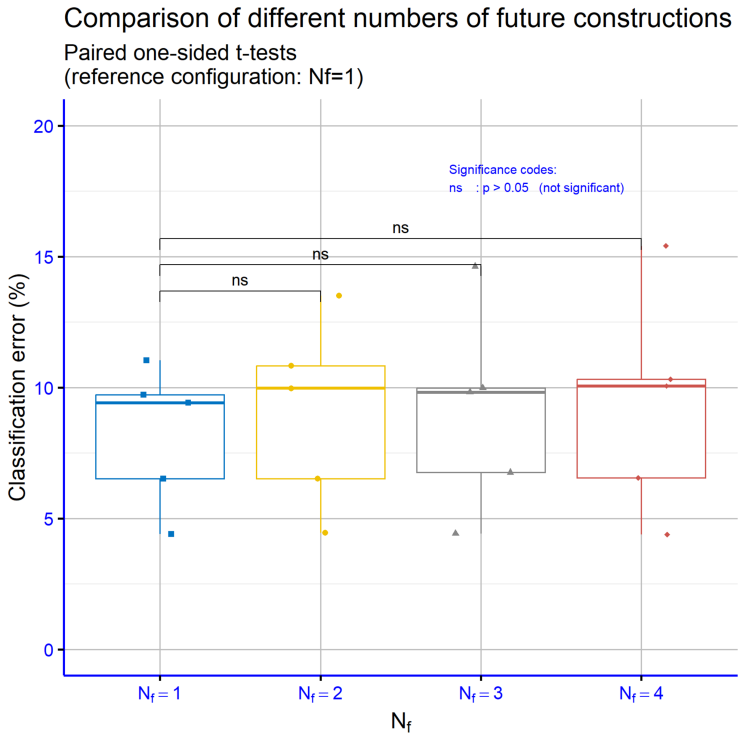


Figure 10. Comparison of FC and baseline classifiers on GCT datasets (paired one-sided t-tests on classification error)

The boxplot for Table 7, together with the paired t-tests, shows that none of the comparisons among the $N_f = 1$, $N_f = 2$, $N_f = 3$ and $N_f = 4$ configurations reaches statistical significance (all labelled as ns). This indicates that, given the available datasets, the performance of the FC model is essentially insensitive to the number of future constructions, and that using smaller values such as $N_f = 1$ or $N_f = 2$ achieves comparable accuracy without any statistically supported gains from increasing N_f (Figure 10).

3.2. Experiments with the number of generations

Moreover, in order to test the efficiency of the proposed method, an additional experiment was executed, where the number of generations was altered from 50 to 400.

Table 8. Experimental results using the proposed method and a series of values for the number generations N_g.

DATASET	$N_g = 50$	$N_g = 100$	$N_g = 200$	$N_g = 400$
GCT1990	13.61%	13.19%	13.50%	12.30%
GCT1995	11.06%	11.70%	10.82%	10.74%
GCT2000	6.53%	6.58%	6.52%	6.52%
GCT2005	4.58%	4.45%	4.45%	4.41%
GCT2010	9.88%	10.00%	9.97%	9.94%
AVERAGE	**9.13%**	**9.18%**	**9.05%**	**8.78%**

Table 8 focuses on the number of generations N_g of the evolutionary algorithm during feature construction and evaluates four values (50, 100, 200, 400). The results demonstrate that FC is remarkably robust with respect to this hyperparameter: the mean classification error remains very close across settings, ranging from 9.18% for $N_g = 100$ down to 8.78% for $N_g = 400$. The configuration $N_g = 200$, which is used as the default in the previous analysis, attains an average error of 9.05%, essentially indistinguishable from the more expensive setting $N_g = 400$, which improves the mean error only by about a quarter of a

percentage point. At the level of individual datasets there are cases where a larger number of generations yields more noticeable gains (for instance, GCT1990 improves to 12.30% at N_g = 400), but the overall pattern is that most of the benefit is already captured within 100-200 generations, with further iterations providing diminishing returns. Thus, Table 8 suggests that the Grammatical Evolution process converges to useful future constructions relatively quickly, and that the choice N_g = 200 offers a very good trade-off between computational cost and predictive performance, avoiding unnecessary increases in training time without delivering substantial accuracy gains.

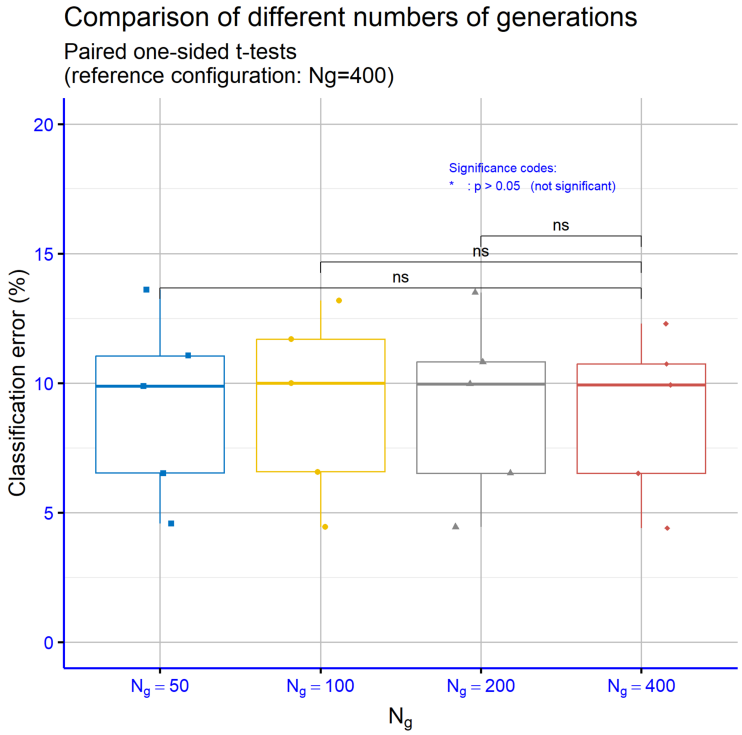


Figure 11. Comparison of FC and baseline classifiers on GCT datasets (paired one-sided t-tests on classification error)

In Figure 11, the paired t-tests for the different numbers of generations show that all pairwise comparisons are non-significant (ns), indicating that variations in N_g do not materially affect the classification error of the FC model.

4. Conclusions

This study investigates the use of Grammatical Evolution for constructing artificial features in earthquake prediction, applying several machine learning approaches, including MLP(GEN), MLP(PSO), SVM, and NNC, alongside Feature Construction (FC). The analysis is based on seismic data recorded between 1990 and 2015 within the geographical area defined by latitudes 33°–44° and longitudes 17°–44°. While all the aforementioned methods belong to the domain of machine learning, FC is distinguished as a feature engineering technique. Specifically, Feature Construction (FC) refers to the process of generating new, informative attributes from the existing dataset, thereby enhancing the representational capacity of the data and improving the performance of machine learning models. Following these steps, our experiments demonstrated that the FC technique yielded the best results, achieving the lowest mean error 9.05% corresponding to an overall accuracy of 91%. The SVM method achieved the second-best performance, with an average error of 11.86%. Consequently, we proceeded with the FC technique, which yielded the best results, and

implemented artificially constructed features N_f with ranging from 1 to 4. Furthermore, to evaluate the efficiency of the proposed method, an additional experiment was conducted in which the number of generations N_g was varied from 50 to 400. Consequently, we applied 1, 2, 3, and 4 constructed features within this technique, with the single constructed featureN_f =1 exhibiting superior performance compared to the others producing the minimal average error 8.21%. Our experiment was further extended by incorporating a series of values for the number of generationsN_g (50, 100, 200, and 400), and the results indicated that N_g 400 generations yielded the best performance 8.78%. In contrast, selecting N_g 200 provides also an effective balance between computational cost and predictive performance. This study was conducted as a direct response to the challenge identified in our previous research, thereby extending and refining the scope of the earlier findings.

Specifically, the study demonstrates the potential of Grammatical Evolution as a novel feature engineering tool in seismology, offering a new perspective on how artificial features can enhance earthquake prediction models. At the same time, certain limitations must be acknowledged, such as the imbalance in the dataset and the restriction to a specific geographical region, which may affect generalizability. Future research should therefore explore the application of this approach to diverse seismic catalogs, investigate its integration with machines learning architectures. By articulating these implications, limitations, and directions, the contribution of the study becomes clearer and more impactful.

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Conflicts of Interest: The authors declare no conflicts of interest.

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