

# Deriving an intelligent model for soil compression index utilizing multi-gene genetic programming

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**Abstract** Multi-gene genetic programming (MGGP) is a new nonlinear system modeling approach that integrates the capabilities of standard GP and classical regression. This paper deals with the prediction of compression index of fine-grained soils using this robust technique. The proposed model relates the soil compression index to its liquid limit, plastic limit and void ratio. Several laboratory test results for fine-grained soils were used to develop the models. Various criteria were considered to check the validity of the model. The parametric and sensitivity analyses were performed and discussed. The MGGP method was found to be very effective for predicting the soil compression index. The prediction coefficients of determination were 0.856 and 0.840 for the training and testing data, respectively. A comparative study was further performed to prove the superiority of the MGGP model to the existing soft computing and traditional empirical equations.

**Keywords** Multi-gene genetic programming · Soil compression index · Soil engineering properties · Prediction

## Introduction

Soil compressibility is the volume reduction under load due to the drainage of pore water. Precise determination of this property is important for the analysis of the soil layer settlements (Tiwari and Ajmera 2012; Singh and Noor 2012). This issue becomes more challenging for fine-grained soils due to their low permeability compared to coarse-grained soils. Compression index ( $C_c$ ) is one of the main soil compressibility indicators. This parameter is usually used for the measurement of the soil layer settlement (Carter and Bentley 1991; Gulhati and Datta 2005; Singh and Noor 2012; Mohammadzadeh et al. 2014). The settlement is typically obtained using the logarithm of the normal compressive stress ( $\sigma'$ ) against soil void ratio ( $e$ ) curve (Gregory et al. 2006).  $C_c$  is considered as the slope of the virgin region of the compression curve. This curve can be obtained from a standard consolidation (oedometer) test on an undisturbed sample (Gregory et al. 2006; Mohammadzadeh et al. 2014). To avoid cumbersome consolidation tests, various empirical equations have been developed to predict  $C_c$  (Skempton 1944; Nishida 1956; Cozzolino 1961; Terzaghi and Peck 1967; Sowers 1970; Azzouz et al. 1976; Wroth and Wood 1978; Mayne 1980; Park and Lee 2011). The existing models are mainly developed based on traditional regression analyses that have notable modeling drawbacks such as large uncertainties or oversimplification of the complicated mechanism of the consolidation process. Also, the regression method assumes the structure of the model in advance by merely a limited number of linear or nonlinear equations (Alavi and Gandomi 2011; Mohammadzadeh et al. 2014). Consequently, more sophisticated methods are needed for the analysis of the complex behavior of  $C_c$ .

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In this context, soft computing techniques may be considered as efficient alternatives to traditional methods. These techniques automatically specify the structure of a prediction model by learning from data (Salehi and Taghikhany 2012; Salehi et al. 2014; Alavi et al. 2016a, b). Different well-known branches of soft computing have been successfully applied to engineering problems (e.g., Das and Basudhar 2008; Samui 2008a, b; Kayadelen et al. 2009; Günaydın 2009; Kolay et al. 2010; Das et al. 2010, 2011a, b, c; Yilmaz 2010a, b; Akgun and Türk 2010; Kaunda et al. 2010; Azamathulla and Wu 2011; Azamathulla et al. 2012; Mert et al. 2011; Yilmaz et al. 2012; Sattari et al. 2012; Tasdemir et al. 2013; Ocak and Seker 2012, 2013; Isik and Ozden 2013; Wu et al. 2013; Salehi et al. 2015; Park et al. 2013; Ceryan et al. 2013). As a robust soft computing tool, artificial neural networks (ANNs) are widely utilized to predict the  $C_c$  of soil layers (Desai et al. 2009; Jianping et al. 2010; Farkhonde and Bolouri 2010; Daryaee et al. 2010; Kumar and Rani 2011; Park and Lee 2011; Kumar et al. 2012; Rani et al. 2013). A limitation of the ANNs and many of the other soft computing methods is that they do not usually generate prediction equations (Alavi and Gandomi 2011). To deal with this issue, genetic programming (GP) was established by Koza (1992). GP is a soft computing method that works based on the Darwinian evolution theory for creating computer programs. The computer programs evolved by GP can be converted to highly nonlinear prediction equations (Javadi et al. 2006; Guven 2009; Guven et al. 2009; Gandomi et al. 2010; Gandomi and Alavi 2011; Rezanian et al. 2011; Azamathulla et al. 2011; Alavi et al. 2011; Mousavi et al. 2012; Azamathulla 2012; Mahmood et al. 2013).

Multi-gene GP (MGGP; Searson et al. 2007, 2010) is a new subset of GP. While classical GP output is merely a single program as the optimal solution, the MGGP optimal solution includes several genes that are each a computer program (Searson et al. 2010; Gandomi and Alavi 2012a, b). As opposed to classical GP and other soft computing techniques, application of MGGP in the engineering domain is new (Gopalakrishnan et al. 2010; Desai and Shaikh 2012; Gandomi and Alavi 2012a, b; Muduli et al. 2013a, b; Muduli and Das 2013a, b; Mohammadi et al. 2014). This paper proposes the MGGP technique to develop a predictive equation for the compression index of soils. A comprehensive set consolidation test results was established for the development of the models. The proposed model was verified through different validation phases.

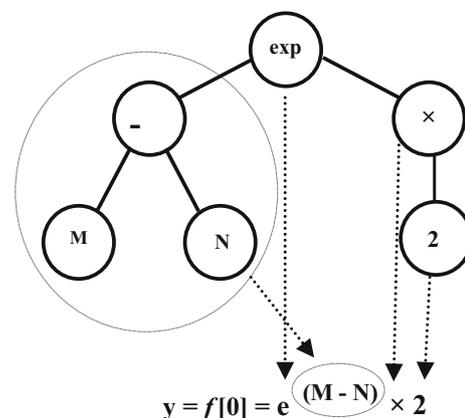
### Multi-gene genetic programming

GP generates computer models by simulating the biological evolution of living organisms (Koza 1992). The genetic operators of genetic algorithm (GA) and GP are almost the

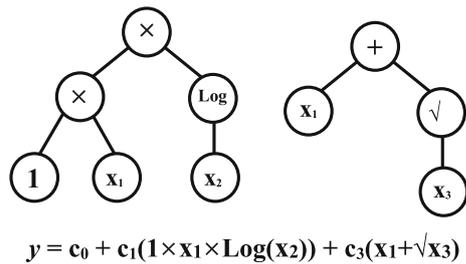
same. GA gives the optimal solution as a string of numbers while the solutions generated by GP are computer programs provided in a tree-like form (Koza 1992; Gandomi and Alavi 2012a, b). In GP, the main goal is to obtain a program connecting inputs to the corresponding output. This process is started by creating a population of programs. The tree-like programs created by GP include functions and terminals chosen randomly. Figure 1 illustrates a typical tree-like program evolved by GP. Further details about GP are given in Koza (1992), Alavi and Gandomi (2011) and Gandomi and Alavi (2012a, b).

MGGP (Searson et al. 2007, 2010; Searson 2009) is a new variant of GP. MGGP evolves a number of genes (sub-programs) that are each individual tree-like programs (Searson et al. 2007, 2010; Searson 2009). Then, it assigns weights to each of these genes to generate the final model as a linear combination of the sub-programs. Figure 2 shows a typical program evolved by MGGP. The inputs of the model are  $x_1$ ,  $x_2$  and  $x_3$  and the functions used for the evolution process are  $\times$ ,  $-$ ,  $+$ ,  $\log$ , and  $\sqrt{\quad}$ . The model is linear in the parameters regarding the coefficients  $c_0$ ,  $c_1$  and  $c_2$  despite using nonlinear terms (Searson et al. 2007, 2010; Gandomi and Alavi 2012a, b). To obtain the linear coefficients, an ordinary least squares analysis is performed on the training data. It is possible to use a partial least squares method to this aim (Searson et al. 2007, 2010; Gandomi and Alavi 2012a, b). The important MGGP parameters that need notable control are the number of genes and tree depth. Restricting the tree depth mostly results in generating more compact models (Searson et al. 2007, 2010; Gandomi and Alavi 2012a, b).

The initial population generated by MGGP contains GP trees with different randomly generated genes. MGGP employs a tree crossover operator as well as the classical recombination operators (Searson et al. 2007, 2010; Gandomi and Alavi 2012a, b). As an example, assume that two parent programs evolved by MGGP contain two



**Fig. 1** Tree representation of a GP model



**Fig. 2** A typical program evolved by MGGP

genes (G1 G2) and three genes (G3 G4 G5). The genes subjected to crossover are shown by [] as given herein: (G1 [G2]) and (G3 [G4 G5]). Thus, during the crossover operation, the genes are exchanged to create two new programs: (G1 G4 G5) and (G3 G2). Besides, there are different types of mutation in MGGP such as sub-tree mutation and mutation of constants using an additive Gaussian perturbation. Further details about MGGP and its operators can be found in Searson et al. (2007, 2010) and Gandomi and Alavi (2012a, b).

**Modeling of  $C_c$  of fine-grained soils**

Plasticity is one of the major physical properties of fine-grained soils. It is measured by Atterberg limits, i.e., liquid limit (LL) and plastic limit (PL). These parameters represent a threshold of water content at which soil is expected to have certain behavior types (e.g., viscous flow, plasticity). LL denotes the lower limit of viscous flow, or the water content, at which the soil behavior changes from plastic to liquid. On the other hand, PL represents the lower limit of the plastic state (Kim et al. 2013). The structure of the existing models (Skempton 1944; Nishida 1956; Cozzolino 1961; Terzaghi and Peck 1967; Sowers 1970; Azzouz et al. 1976; Mayne 1980; Park and Lee 2011; Mohammadzadeh et al. 2014) clearly indicates that  $C_c$  of fine-grained soils can be considered as a function of percentage of LL and PL, as well as initial void ratio ( $e_o$ ). The same strategy was considered in this study to develop a simplified prediction model for  $C_c$  using merely its basic physical properties. Determination of LL, PL and  $e_o$  is much more easier than performing the consolidation tests to find  $C_c$ . Hence, the adapted strategy would result in a notable cost savings for geotechnical engineering design. As it is, all these parameters have a rational relationship with the natural water content for saturated soils (Bartlett and Lee 2004; Kim et al. 2013; Mohammadzadeh et al. 2014).

To develop the MGGP-based model, a comprehensive database containing soil properties (LL, PL and  $e_o$ ) and corresponding  $C_c$  was developed. The database was

divided into three sub-divisions of learning, validation and testing subsets as will be described in “MGGP-based formulation of  $C_c$ ” section. The MGGP algorithm was run on the learning data and checked on the other two subsets. The optimal model was selected considering its simplicity as well as its performance on the learning and validation data.

Determination coefficient ( $R^2$ ), root mean squared error (RMSE) and mean percent error (MAE) were used to assess the performance of the models (Alavi and Gandomi 2011):

$$R^2 = \frac{(\sum_{i=1}^n (h_i - \bar{h}_i)(t_i - \bar{t}_i))^2}{\sum_{i=1}^n (h_i - \bar{h}_i)^2 \sum_{i=1}^n (t_i - \bar{t}_i)^2}, \tag{1}$$

$$RMSE = \sqrt{\frac{\sum_{i=1}^n (h_i - t_i)^2}{n}}, \tag{2}$$

$$MAE = \frac{1}{n} \sum_{i=1}^n |h_i - t_i|, \tag{3}$$

where  $h_i$  is the measured  $C_c$ ,  $t_i$  is the predicted  $C_c$ ,  $\bar{h}_i$  is the average of the measured  $C_c$ , and  $\bar{t}_i$  is the average of the predicted  $C_c$ .

**Laboratory test**

Several laboratory test results were performed on samples obtained from different locations in Khorasan Razavi Province, Iran to develop the database. The soil samples were picked up from test pits at 0.5–1.0 m depth. The physical and plastic characteristics of soil were determined through geotechnical laboratory tests. Conventional oedometer tests were carried out to determine  $C_c$ . The testing procedure was as described in Abbasi et al. (2012) and Mohammadzadeh et al. (2014). A conventional oedometer apparatus having brass ring, 75 mm in diameter and 20 mm in height was used to perform the tests. The vertical dead load was applied using a loading device until there was no change in dial gauge reading for two consecutive hours. Other details of the test were performed in general accordance with ASTM D2435 procedure (2000). The variation of void ratio versus pressure was plotted for each specimen on a semilogarithmic scale to obtain the  $C_c$  values (Abbasi et al. 2012). One-dimensional consolidation and basic geotechnical characterization tests are very well-known tests. Thus, for brevity, some basic information was provided herein. Table 1 presents the descriptive statistics of the parameters (LL, PL,  $e_o$ ,  $C_s$ ) used in this study. There were a total of 101 laboratory tests on fine-grained soil samples. To enrich the database, seven consolidation test results done by Malih (2006) were added to the available data. Different soil types used in this study were silty clay with sand (CL–ML), gravelly lean clay with sand (CL), and silty, clayey sand (SC–SM).

**Table 1** Descriptive statistics of the variables used for the model development

Parameters	$e_0$	LL (%)	PL (%)	$C_c$
Mean	0.745	36.156	22.605	0.171
Standard deviation	0.123	12.786	5.639	0.047
Range	0.516	52.600	29.200	0.176
Minimum	0.514	19.400	14.800	0.077
Maximum	1.030	72.000	44.000	0.253

**Table 2** Parameter considered for the implementation of the MGGP algorithm

Parameters	Settings
Population size	200, 400, 800, 1200, 1500, 2000, 3000
Generations	200, 400, 800, 1000
Maximum number of genes	4, 6, 10, 12
Maximum tree depth	4, 6, 10, 12
Tournament size	25
Crossover events	0.85
Mutation events	0.1
Function set	+, -, ×, /, ^2, ^3, tanh, sin, cos, exp

To avoid overfitting, the data were classified into three subsets: (1) learning, (2) validation, and (3) test subsets. The models were trained using the learning data. Then, their performance was checked on the validation data. The model with the highest  $R^2$  and lowest RMSE and MAE values on the learning and validation data was chosen as the optimal model. Eventually, the optimal model was tested on un-seen testing data to guarantee that it has generalization capacity. Since both the learning and validation sets were included in the model selection, they were grouped together as “training data” (Alavi et al. 2011; Mohammadzadeh et al. 2014). The training, validation and testing data are commonly taken as 50–70, 15–25 and 15–25 of all data, respectively (Shahin and Jaksa 2005; Alavi et al. 2011). Herein, about 65 % (71 sets), 15 % (16 sets) and 20 % (21 sets) of the data were considered for the training, validation and testing processes, respectively.

**MGGP-based formulation of  $C_c$**

To find the optimal model, the MGGP algorithm was run several times with different combinations of the parameters. This process was based on both an extensive trail study and referring to the suggested values in previous studies (Searson et al. 2010; Gopalakrishnan et al. 2010; Desai and Shaikh 2012; Gandomi and Alavi 2012a, b). Table 2 presents the parameters considered for different

**Table 3** Individual genes/model terms for the prediction of  $C_c$

Terms	Values
Bias	-0.176
Gene 1	$-0.0227x_2 - 0.0113x_3$
Gene 2	$-(0.162 (x_3 + x_2 \tanh(x_3)))/x_2$
Gene 3	$0.016x_2 - 0.194x_3 + (0.016x_3)/x_1$
Gene 4	$0.00468x_1x_2$
Gene 5	$0.87x_1$
Gene 6	$-(0.0191x_2)/(x_1 + x_3 + \tanh(\tanh(\exp(x_2 - \exp(x_1))))))$
Gene 7	$0.187x_3$
Gene 8	$(0.014x_3)/x_2$
Gene 9	$7.33 \times 10^{(-4)}x_2 + 7.33 \times 10^{(-4)} \sin(x_2) + 7.33 \times 10^{(-4)} \tanh(x_2)^3 - (4.23 \times 10^{14}x_2 + 4.23 \times 10^{14}x_3 + 4.23 \times 10^{14}\tanh(x_1))/ (1.15 \times 10^{18}x_1 + 1.15 \times 10^{18}x_2 + 5.76 \times 10^{17}\exp(\tanh(x_1x_3)))$

$$x_1 = e_0; x_2 = LL; x_3 = PL$$

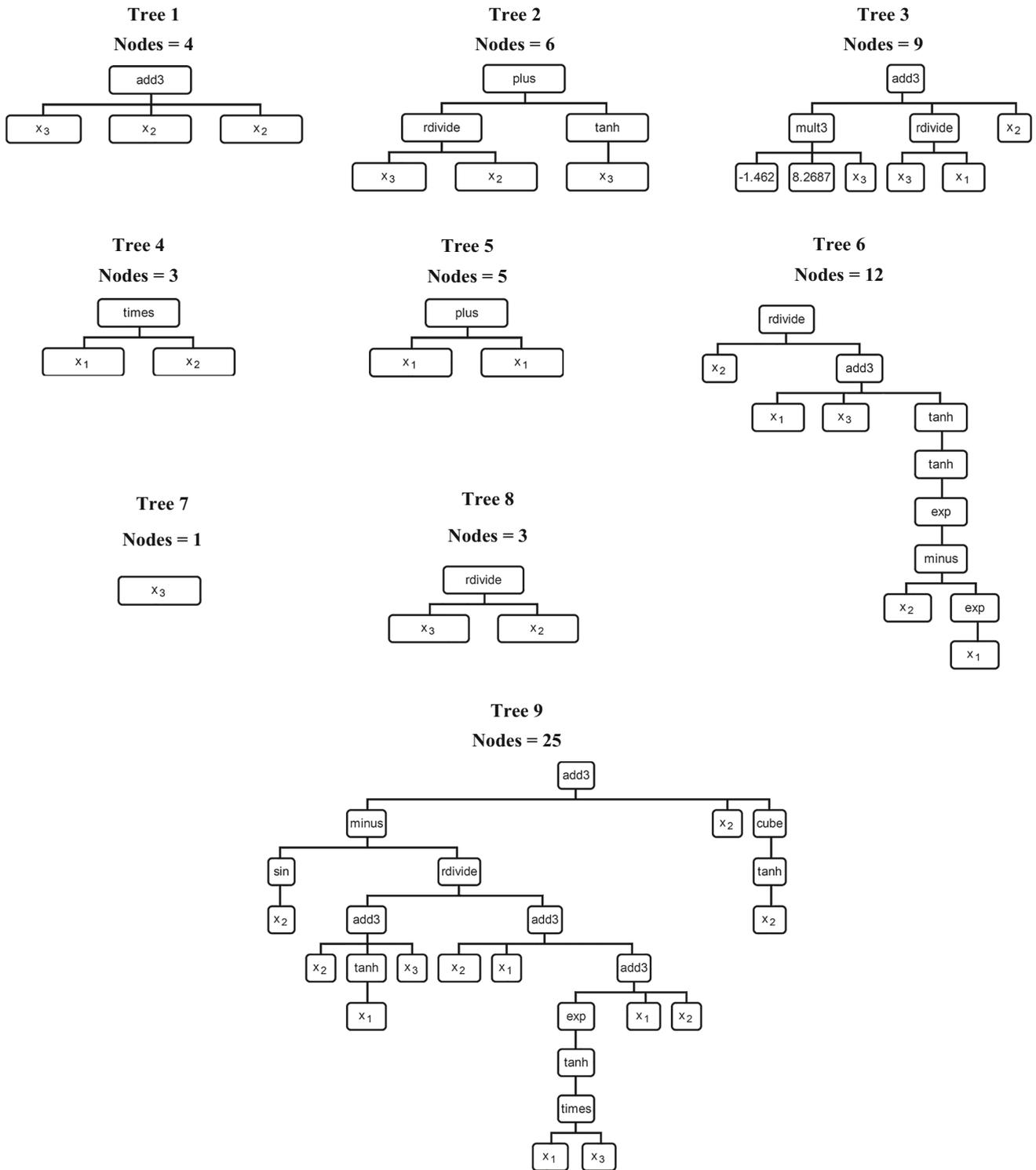
runs. Large values were taken for the population and generations to ensure that the best models were derived. As it is seen in Table 2, there were  $7 \times 4 \times 4 \times 4 = 448$  parameter arrangement. All these arrangements were tested with two replications for each. Thus, about  $448 \times 2 = 896$  optimal individual runs were performed. GPTIPS toolbox (Searson 2009) coded in MATLAB was utilized to run the algorithm. The RMSE function was adapted for error minimization during runs.

Table 3 shows the individual genes/model terms for the best models that were obtained during the conducted runs. Each gene includes its weighting coefficient. As can be observed from this table, the derived model is composed of complicated array of operators, variables, and constants to estimate  $C_c$ . Besides, Fig. 3 shows the expression trees (ETs) of the MGGP model. In these figures,  $x_1$ ,  $x_2$ , and  $x_3$  represent  $e_0$ , LL and PL, respectively.

To facilitate the use of the developed model, it was transformed into a functional form and thereafter simplified as follows:

$$C_c = 0.87e_0 - 0.00591LL - 0.0186PL + 7.33 \times 10^{-4} \sin(LL) - 0.162 \tanh(PL) + 7.33 \times 10^{-4} \tanh(LL)^3 + 0.00468e_0LL - (1.38 \times 10^{15}LL)/(7.21 \times 10^{16}e_0 + 7.21 \times 10^{16}PL + 7.21 \times 10^{16} \tanh(\tanh(\exp(LL - \exp(e_0)))))) - (4.22 \times 10^{14}LL + 4.22 \times 10^{14}PL + 4.22 \times 10^{14} \tanh(e_0))/(1.15 \times 10^{18}e_0 + 1.15 \times 10^{18}LL + 5.76 \times 10^{17} \exp(\tanh(e_0PL))) + (0.016PL)/e_0 - (0.148PL)/LL - 0.176. \tag{4}$$

Figure 4 shows the experimental against predicted  $C_c$  values. The numbers of population, generations, genes and tree depth for the model were equal to 1500, 800, 10 and 10, respectively. The performance of the model is good on

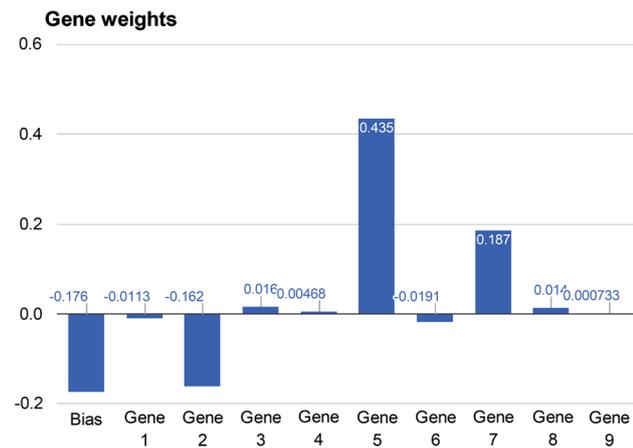
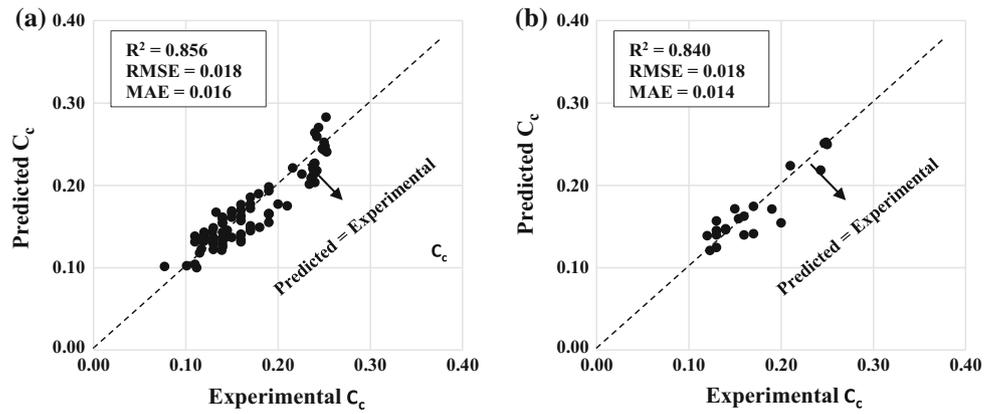


**Fig. 3** Expression trees of the best models for the prediction of  $C_c$  ( $ET = \sum \text{sub-ET}_i$ )

both the training and testing data. The importance of each of the nine genes of the derived model is shown in Fig. 5. It is seen that the weight of the genes (sub-programs) 5, 7, 2 and the bias terms are higher than the other genes. This

means that they have higher contribution to the prediction of  $C_c$ . Besides, Fig. 6 presents the accuracy against the complexity of the evolved models. As it is seen, the models with good performance seem to be much less complex. The

**Fig. 4** Performance of the MGGP model on: **a** training data, and **b** testing data

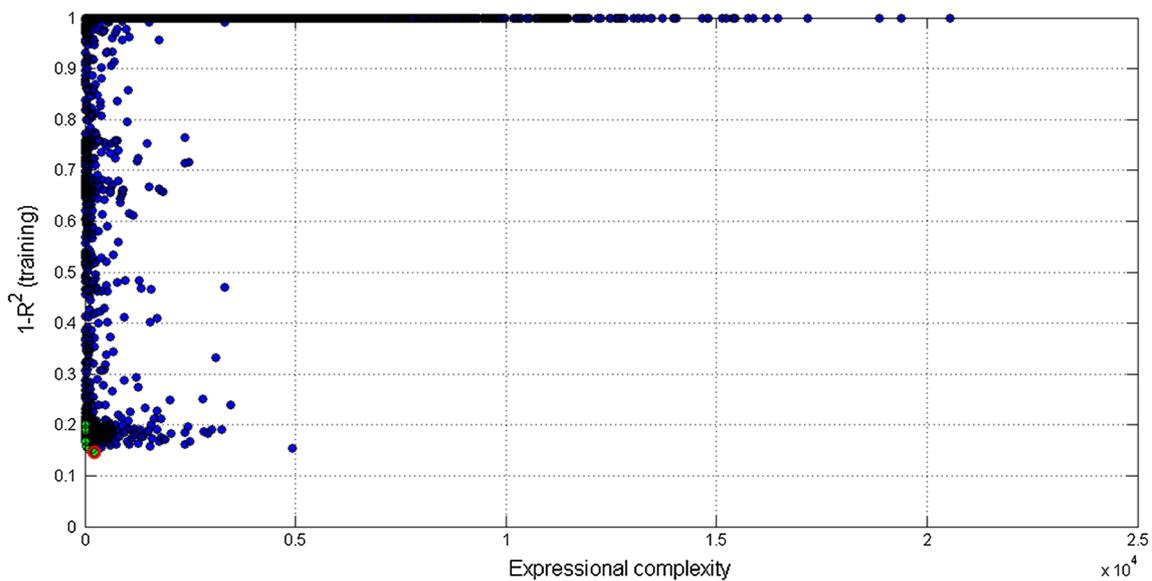


**Fig. 5** Weights of the genes (sub-programs) of the MGGP model

red circle in Fig. 6 designates the best model presented herein.

**Validation verification**

Smith (1986) argues that for correlation coefficient ( $|R| > 0.8$ ), a powerful correlation exists between the predicted and actual values and, therefore, the model is suitable. In addition, the error values should be minimum in all cases (Alavi et al. 2011). Figure 4 indicates that the MGGP model has high  $R$  and low RMSE and MAE values. Besides, Golbraikh and Tropsha (2002) defined new statistical parameters of  $k$  and  $k'$  to validate the models on the testing data. Also, Roy and Roy (2008)



**Fig. 6** Complexity against accuracy of the models evolved by MGGP

**Table 4** Verifying the validity of the MGGP model on the testing data

Items	Formulas	Conditions	MGGP
1	$\frac{\sum_{i=1}^n (h_i - \bar{h}_i)(t_i - \bar{t}_i)}{\sqrt{\sum_{i=1}^n (h_i - \bar{h}_i)^2 \sum_{i=1}^n (t_i - \bar{t}_i)^2}}$	$0.8 < R$	0.916
2	$k = \frac{\sum_{i=1}^n (h_i \times t_i)}{h_i^2}$	$0.85 < k < 1.15$	1.001
3	$k' = \frac{\sum_{i=1}^n (h_i \times t_i)}{t_i^2}$	$0.85 < k' < 1.15$	0.989
4	$R_m = R^2 \times (1 - \sqrt{ R^2 - R_O^2 })$	$0.5 < R_m$	0.503
Where	$R_O^2 = 1 - \frac{\sum_{i=1}^n (h_i - h_i^o)^2}{\sum_{i=1}^n (t_i - t_i^o)^2}, h_i^o = k \times t_i$	Should be close to 1	1.000
	$R_O'^2 = 1 - \frac{\sum_{i=1}^n (h_i - t_i^o)^2}{\sum_{i=1}^n (h_i - h_i)^2}, t_i^o = k' \times h_i$	Should be close to 1	0.998

$h_i$  Actual output,  $t_i$  predicted output

**Table 5** Comparison of the predictions provided by different models

References	$R^2$	RMSE	MAE
Skempton (1944)	0.367	0.072	0.056
Nishida (1956)	0.752	0.301	0.285
Cozzolino (1961)	0.752	0.105	0.103
Terzaghi and Peck (1967)	0.367	0.110	0.077
Sowers (1970)	0.752	0.055	0.042
Azzouz et al. (1976)	0.752	0.036	0.032
Mayne (1980)	0.367	0.102	0.073
Park and Lee (2011)	0.752	0.089	0.085
MEP (Mohammadzadeh et al. 2014)	0.811	0.019	0.016
ANN (Mohammadzadeh et al. 2014)	0.859	0.017	0.014
MGGP (this study)	0.840	0.018	0.014

proposed a predictability indicator parameter called  $R_m$ , as well as two other related parameters ( $R_O^2$  and  $R_O'^2$ ).  $k, k', R_m, R_O^2$ , and  $R_O'^2$  are defined based on actual ( $h_i$ ) and predicted ( $t_i$ ) outputs. It is suggested that at least one slope of regression lines ( $k$  or  $k'$ ) through the origin should be close to 1.  $k$  is the slope of the regression line in plot of  $h_i$  against  $t_i$  values.  $k'$  is the slope of the regression line in plot of  $t_i$  against  $h_i$  values (Golbraikh and Tropsha 2002). Either the squared correlation coefficient (through the origin) between predicted and experimental values ( $R_O^2$ ), or the coefficient between experimental and predicted values ( $R_O'^2$ ) should be close to 1.  $R^2$  and  $R_O^2$  are used for calculating  $R_m$ . For  $R_m > 0.5$ , a prediction model can be considered valid. The above criteria were checked for the developed model and given in Table 4. This table indicates that the derived model meets the mentioned criteria and, therefore, is credible.

Furthermore, the MGGP model was compared with two other powerful soft computing methods, called multi-

expression programming (MEP) and ANN (Mohammadzadeh et al. 2014) and several other well-known empirical models by other researchers (Skempton 1944; Nishida 1956; Cozzolino 1961; Terzaghi and Peck 1967; Sowers 1970; Azzouz et al. 1976; Mayne 1980; Park and Lee 2011). The MEP and ANN models were established upon the same data used for the development of the MGGP-based model. The comparison study was conducted on the testing data and presented in Table 5. According to this table, the proposed model has a comparable performance with ANN. The MGGP formula outperforms the MEP model and has a notable better performance than the existing regression-based models. In particular, the prediction error values (RMSE and MAE) for the existing empirical models are much higher than those for the MGGP model. It is notable that the regression analysis used for developing the  $C_c$  empirical models assumes the structure of the model by a limited number of linear or nonlinear equations. Hence, such models cannot consider the highly nonlinear interactions between the soil parameters and  $C_c$  (Alavi et al. 2011). Conversely from the empirical and analytical methods, MGGP models the mechanical behavior directly from the experimental data with no prior assumptions. The best MGGP-based equation is chosen among numerous linear and nonlinear preliminary models (Alavi et al. 2011; Alavi and Gandomi 2011).

### Parametric and sensitivity analyses

To ensure the validity of MGGP-based model, a comparative parametric analysis was performed. Figure 7 shows the results of the parametric analysis. As it is seen,  $C_c$  increases with increasing LL up to about 32 % and then it decreases for higher LL values. Figure 7b indicates that  $C_c$

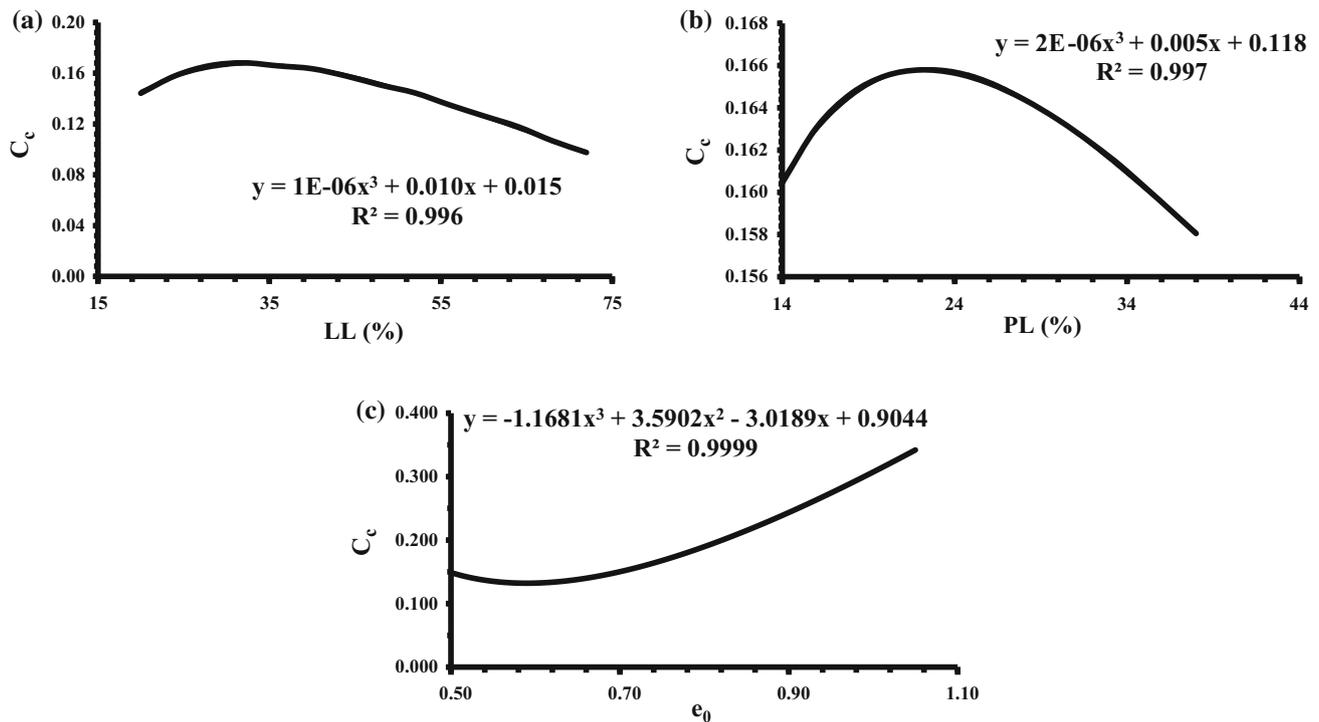


Fig. 7 Parametric analysis of  $C_c$  in the MGGP model

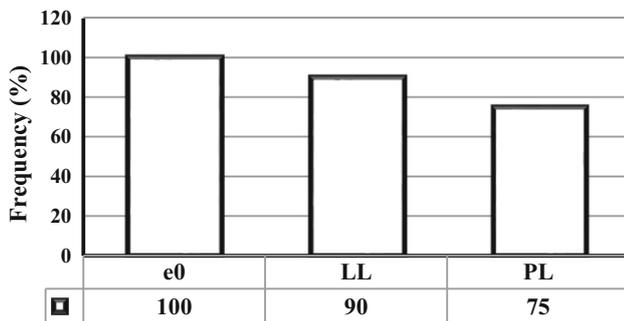


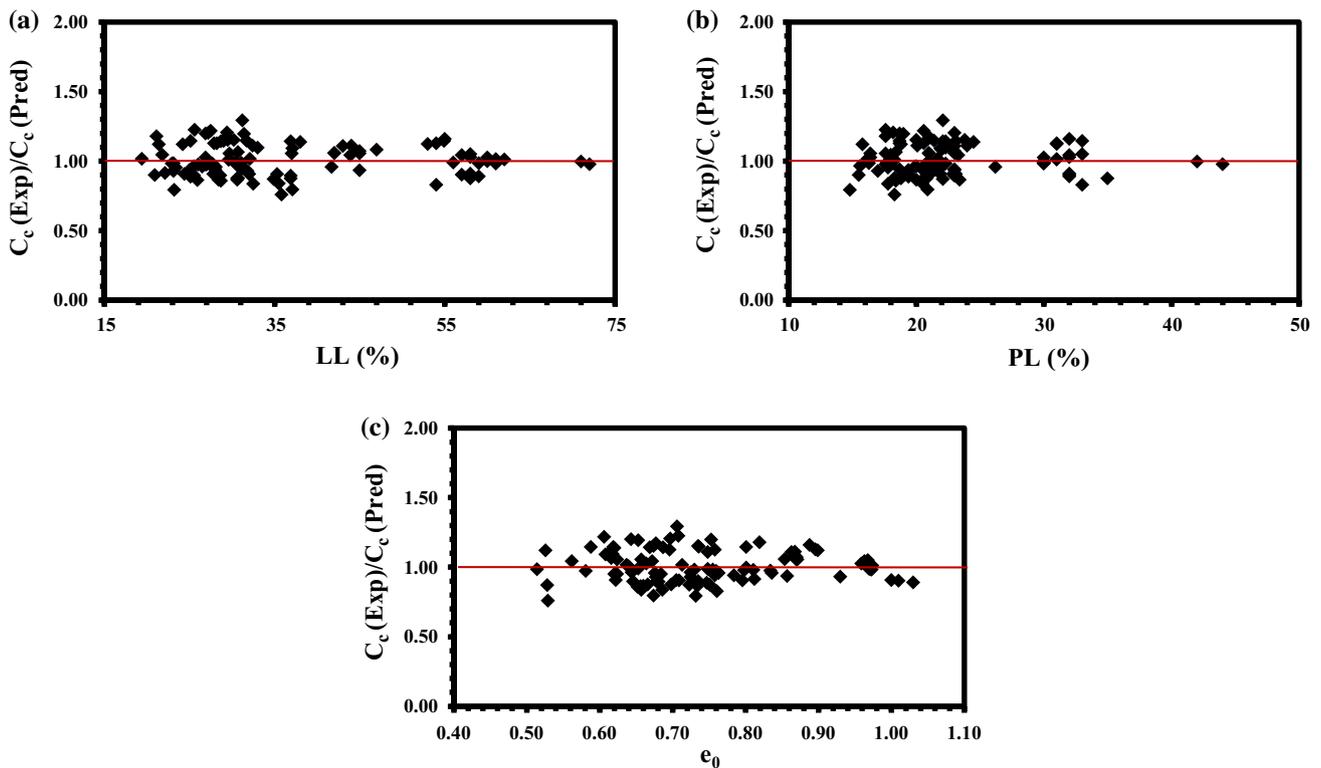
Fig. 8 Contributions of the predictor variables

is positively correlated with PL up to 22 % and for higher PL values  $C_c$  decreases. Referring to Fig. 7c,  $C_c$  decreases with increasing  $e_0$  up to about 0.6 and thereafter it starts increasing.

As discussed before, the effect of all the considered parameters (i.e., LL, PL and  $e_0$ ) on  $C_c$  is well understood. Excluding any of these three parameters from the modeling process decreased the accuracy of the derived models. Herein, a sensitivity analysis was conducted to provide a

better understanding of the contribution of these important parameters to the prediction of  $C_c$ . A common approach for the sensitivity analysis in the GP-based modeling is to obtain the frequency values of the input parameters (Francone 1998–2004; Alavi et al. 2011; Gandomi et al. 2011a, b; Mohammadzadeh et al. 2014). A parameter with a frequency equal to 100 % has been included in 100 % of the best 30 programs generated by MGGP. The sensitivity analysis results are shown in Fig. 8. As it is seen, the frequency of  $e_0$ , LL and PL is equal to 100, 100, and 85 %, respectively. Thus, the MGGP-based model is more sensitive to  $e_0$  and LL compared to PL. This is in agreement with the results reported by other researchers (Mohammadzadeh et al. 2014; Daryaei et al. 2010).

Besides, Fig. 9 shows the ratios of the experimental  $C_c$  values to those predicted by the MGGP model with respect to LL, PL and  $e_0$ . As the scattering increases in these figures, the model accuracy consequently decreases. It can be observed from these figures that the predictions obtained by the proposed model have a very good accuracy with no significant trend with respect to the design parameters. In the case of LL and  $e_0$ , the scattering slightly decreases with increasing these parameters.



**Fig. 9** The ratio of the experimental to the predicted  $C_c$  values with respect to the input parameters

**Conclusion**

This paper presented the MGGP method for the prediction of  $C_c$  of fine-grained soils.  $C_c$  was formulated in terms of basic properties of soil, i.e., LL, PL and  $e_0$ . The model was established upon a comprehensive experimental database. Extensive simulations were carried out to find the optimal model. To verify the validity of the model, various criteria were adapted. The proposed model has a good performance and outperforms the existing soft computing and empirical models. Contrary to the classical models for  $C_c$ , the MGGP model is highly nonlinear and includes complex combination of the predictor variables. For further evaluation of the behavior of the model, parametric and sensitivity analyses were performed. It was found that  $C_c$  increases with increasing LL and PL up to 32 and 22 %, respectively. In general,  $C_c$  was positively correlated with  $e_0$ . More, LL and  $e_0$  seem to be more influencing on  $C_c$  than PL. Within its calibrated range, the proposed model can be considered as an efficient alternative to cumbersome testing procedures for the determination of  $C_c$ .

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