## Econometric Genetic Programming Outperforms Traditional Econometric Algorithms for Regression Tasks

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### ABSTRACT

Econometric Genetic Programming (EGP) evolves multiple linear regressions through Genetic Programming (GP), which is responsible for model selection, aiming to generate high accuracy regressions with potential interpretability of parameters. It uses statistical significance as a feature selection tool, directly and efficiently identifying introns and controlling bloat. In this paper, EGP is tested against traditional feature-selection econometric algorithms in regression tasks - namely Partial Least Squares Regression, Ridge Regression and Stepwise Forward Regression outperforming them in all three datasets. The way EGP explores search space of possible regressors and models is crucial for its results. EGP is carefully constructed considering econometric theory on cross-sectional datasets, giving rigorous treatment on topics like homoscedasticity and heteroscedasticity, statistical inference for estimated parameters and sampling criteria. It also benefits by the mathematical proof on accuracy and statistical significance: accuracy will only increase if the regressor presents a test's statistics module in a two-sided hypothesis testing higher than a predefined value.

#### **KEYWORDS**

Genetic Programming, Multiple Regression, Model Selection, Feature Selection

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#### **1 INTRODUCTION**

The very first form of regression was the method of Least Squares (LS), published by [1] and [2]. The term "regression" was coined by Francis Galton in the nineteenth century to describe a biological phenomenon. As stated in [3], the method for Symbolic Regression (SR) proposed by [4] is an alternative approach to curve fitting. The technique creates mathematical expressions to fit a set of data points using the evolutionary process of GP.

Past literature relates some pioneer works that hybridize linear regression with GP. For a generous list on it, see [5]. In the following, just papers that most influenced EGP will be described.

Works [3] and [6,7] create polynomial regression models for SR tasks. The Weierstrass approximation theorem (1885) states that every continuous function defined on a closed interval [a, b] can be uniformly approximated as closely as desired by a polynomial function. By itself, the theorem would be sufficient for a great effort on approximating the dependent variable in a regression task by polynomials. EGP follows this direction.

EGP, which was first and partially introduced in [8] for regression tasks and tested against exhaustive search algorithms, is carefully constructed considering econometric theory on crosssectional datasets. Rigorous treatment on topics like homoscedasticity [9], statistical inference for estimated parameters and sampling criteria are made. These considerations represent a significant difference with its predecessors, which relax some hypothesis or even do not test them on datasets and models, although each of them has its own contribution.

Kaizen Programming (KP) [13] is an interesting evolutionary tool based on concepts of continuous improvement from Kaizen methodology, which was successfully tested against traditional SR benchmark functions. EGP is similar to KP in the sense that both are concerned about bloat, introns and use  $\overline{R}^2$  (see section 2) as a comparison metric between models. But they also differ in several aspects, in particular the fact that EGP exclusively evolves polynomials (enhancing potential interpretability of parameters),

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while KP evolves linear-in-parameters individuals in a more general shape.

In this paper, EGP is tested against traditional featureselection econometric algorithms in regression tasks outperforming them in all three datasets. The main reason EGP outperforms traditional econometric algorithms is its capability to explore the regressors' and models' search space. While the number of regressors for each of the feature-selection algorithms are just a few for each dataset, EGP generates models with 50 statistically significant regressors or more. EGP explores nonlinearity of features, by multiplying different features, maintaining models with linear structure.

This paper is organized as follows: Section 2 describes the elements of econometrics used by EGP: there is no intention to fully exhaust the theme; justification on these elements is presented when necessary. Section 3 succinctly describes EGP. Sections 4 proposes experiments and discusses results. Conclusion is done in Section 5.

## **2** ECONOMETRICS

## 2.1 Linear Regression Model, Least Squares, QR Decomposition

As in [9], the multiple linear regression model with k parameters and n observations is:

$$y = X\beta + u \tag{1}$$

where  $y_{n \times 1}$  is the dependent variable vector,  $X_{n \times (k+1)}$  is the regressor's matrix,  $\beta_{(k+1) \times 1}$  is the vector representing the terms that adjust X to y and  $u_{n \times 1}$  is an error vector.

Vector  $\boldsymbol{\beta}$  is an unknown statistical population parameter usually estimated by LS, which generates  $\hat{\boldsymbol{\beta}}$ , the ideal multiplier for  $\boldsymbol{X}$  on  $\delta(\boldsymbol{X})$  (the column space of  $\boldsymbol{X}$ ) that makes  $\boldsymbol{X}\hat{\boldsymbol{\beta}}$  the most closely projection to  $\boldsymbol{y}$  on  $\delta(\boldsymbol{X})$ . As  $\delta(\boldsymbol{X})$  is orthogonal to  $\boldsymbol{u}$ , some matrix manipulation leads to:

$$\widehat{\boldsymbol{\beta}} = (\boldsymbol{X}^{\mathrm{T}}\boldsymbol{X})^{-1}\boldsymbol{X}^{\mathrm{T}}\boldsymbol{y}$$
(2)

In [10], it is stated: "The most commonly used, and in many ways the most important, estimation technique in econometrics is LS". In general, calculations on (2) via matrix inversion are numerically unstable and QR Decomposition is recommended by [11,12] in such cases.

#### 2.3 Hypothesis Test

Under conditions stated in [14],  $\hat{\boldsymbol{\beta}}$  is a BLUE estimator for  $\boldsymbol{\beta}$  [9], which does not guarantee statistical significance for  $\boldsymbol{\beta} = [\beta_1 \cdots \beta_i \cdots \beta_k]^T$ . I.e., it is possible that some  $\beta_i$  in (1), or even all  $\boldsymbol{\beta}$ , is a pure random effect on  $\boldsymbol{y}$  and does not present any causal relationship with it. To check statistical significance, it is natural to perform HT on  $\beta_i$ , individually, or on  $\boldsymbol{\beta}$ . Just fully satisfiability

of conditions stated in [14] allows to perform HT as described in the following and that is the case of models generated by EGP and datasets used in this article.

HT is constructed with a null and alternative hypothesis ( $H_0$  and  $H_1$ , respectively), a test statistics and a decision criteria. For a two-sided HT for  $\beta_i$ , hypothesis  $H_0$  and  $H_1$  are frequently:

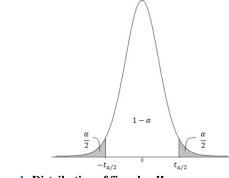
$$H_0: \beta_i = 0$$

$$H_1: \beta_i \neq 0$$
(3)

and  $\hat{\boldsymbol{\beta}} | \boldsymbol{X} \sim N(\boldsymbol{\beta}, \sigma^2(\boldsymbol{X}^T \boldsymbol{X})^{-1})$ , while *T* is a test statistic with a known probability distribution:

$$T = \frac{\hat{\beta}_i - \beta_i}{SE(\hat{\beta}_i)/\sqrt{n}} \sim t \ (n - k - 1) \tag{4}$$

with  $SE(\hat{\beta}_i)$  the standard error of  $\hat{\beta}_i$ . As *n* increases,  $T_n \xrightarrow{a} N(0,1)$ . Decision criteria is defined following Fig. 1.





The quantity  $T_{obs}$  is the observed value of the random variable T when all variables in (4) are substituted by their respective values. If  $|T_{obs}| > t_{\alpha/2}$ ,  $T_{obs}$  is far away from the average of the curve in Fig. 1 and thus is less likely that  $T_{obs}$  is indeed generated by the distribution of T under  $H_0$ . In this case,  $H_0$  is rejected and  $\beta_i$  remains in (1). Otherwise, if  $|T_{obs}| < t_{\alpha/2}$ ,  $T_{obs}$  is probably generated by the distribution of T under  $H_0$ . In this case,  $H_0$  is not rejected and  $\beta_i$  quits (1), because it is not statistical significant. Typically,  $\alpha = 5\%$ ,  $t_{\alpha/2} = 1.96$  and  $-t_{\alpha/2} = -1.96$ .

#### 2.4 Statistical Significance and Accuracy

The Root Mean Square Error (RMSE) is a typical accuracy measure used in SR experiments. Its relation with  $\overline{R}^2$ , a typical metric of fitness in cross-sectional econometrics that linearly penalizes by addition of non-statistically significant regressors with  $|T_{obs}| < 1.00$ , is given by:

$$\overline{\mathbf{R}}^{2} = 1 - \left(\frac{n(\mathrm{RMSE})^{2}}{(\boldsymbol{y} - \overline{\boldsymbol{y}})^{\mathrm{T}}(\boldsymbol{y} - \overline{\boldsymbol{y}})}\right) \left(1 + \frac{k}{n - k - 1}\right)$$
(5)

with  $\overline{y}$  the average of y. By (5), it is concluded that the RMSE minimization implies  $\overline{R}^2$  maximization, maintaining others factors constant.

It is stated in [9] that  $\overline{R}^2$  will increase if, and only if,  $|T_{obs}| > 1.00$  in a two-sided HT for  $\beta_i$ , with null and alternative hypothesis, test statistics and decision criteria stated as before. EGP uses this mathematical proof to increase accuracy of its individuals by statistical significance, while simultaneously controlling bloat. To achieve fitness improvement with statistical significance, EGP uses 1.96 as threshold in HT instead of 1.00.

## 3 ECONOMETRIC GENETIC PROGRAMMING

EGP evolves models in format of (1) through GP, which is responsible for model selection. GP is mainly based in configuration presented in [15], as well as EGP parameters.

## 3.1 Representation

Individuals / programs / regressions / models are multigenic. Any constant in any program comes from LS in (2), i.e. there are no ephemeral constants. The terminal set, namely  $\Omega$ , is purely composed by variables. The primitive set, namely  $\vartheta$ , is composed just by variables and operations of sum and multiplication, due (1) format.

## 3.2 Initial Population

EGP uses a probabilistic version of ramped half-and-half method. Fig. 2 shows a possible individual generated by EGP.

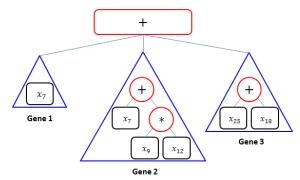


Figure 2: A possible individual generated by EGP.

Set  $\Omega$  is composed by *K* features (independent variables). Every individual has its own set of regressors, forming its own *X*, composed by simple or combined elements of  $\Omega$ . As an example, it is possible that  $x_1$ ,  $x_3x_{11}^2$  and  $x_3x_4x_6$  are regressors of a particular individual, formed by features  $x_1$ ,  $x_3$ ,  $x_4$ ,  $x_6$  and  $x_{11}$ .

#### 3.3 Accuracy

RMSE is the objective function. The  $\overline{\mathbb{R}}^2$  is just used to compare models. Ideally,  $\overline{\mathbb{R}}^2$  would be the objective function, but some issues, principally when k > n, makes it little suitable in practice.

To calculate accuracy in an EGP individual, the one showed in Fig. 2 needs to be transformed into a model like  $y = X\beta + u$ . Then, EGP will solve  $\hat{\beta}$  for  $X\hat{\beta} = \beta_1 x_7 + \beta_2 x_7$  $\beta_2 x_9 x_{12} + \beta_3 x_{18} x_{25}$ . If any of the regressors are not statistically significant, they will be removed from  $y = X\beta + u$ . In sequence,  $\hat{\beta}$  is recalculated just with statistically significant regressors. RMSE is finally calculated using  $\hat{\beta}$  after these steps. This routine is traditional in econometric studies, ensuring statistical significance over a determined significance level  $\alpha$ , and that is the way EGP performs feature selection. Modifications described are necessary just for accuracy calculation, therefore individuals will keep their multigene structure to mutation, crossover and elitism.

EGP is not a kind of stepwise regression, because it does not build a model sequentially, variable by variable, as described in [20]: "(Forward) Stepwise regression builds a model sequentially, adding one variable at a time. At each step, it identifies the best variable to include in the active set, and then updates the least squares fit to include all the active variables."

EGP does not estimate on genes, just on regressors, by two main reasons: possible multicollinearity problem, interfering on HT for  $\beta_i$ , and lack of interpretation for  $\hat{\beta}_i$  when it is related to a gene.

## 3.4 Selection

Tournament selection with  $n_{tourn} = 7$  and repetitions allowed, with a variation on lexicographic parsimony pressure of [16], is used. Individuals with a large number of statistically significant regressors will be preferred over others with a few number, in the same range of fitness. Therefore, EGP is parsimonious in its nature, because it avoids the individuals with a large amount of *introns* (in this case, non statistically significant regressors).

#### 3.5 Mutation, Crossover and Elitism

Types of mutation used: traditional mutation proposed by [4] and mutation by regressors' substitution. Types of crossover used: intergenic and intragenic crossovers. Mutation and crossover rates vary through evolution following automatic adaptation of operators as described in [17]. Elitism rate is settled to 5% of individuals by generation.

#### 3.6 Tools and Parameters

EGP is implemented through a modification on GPTIPS, a *Matlab* toolbox, presented in [18]. Information on EGP parameters are shown in Table 1.

Table 1	l: EGP	Parameters
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Parameters	
- Population size	150.
- Generations	50.
- Maximum gene Depth	5.
- Maximum number of genes by individual	5.
- Probability of traditional mutation [4]	95%.
- Probability of intragenic crossover	50%.

## 4 EXPERIMENTS AND RESULTS

Table 2 presents the results. The training set contains 70% of the data samples. Results for EGP ("EGP-Regression" in the table) are the average of 50 runs for best individuals. EGP is carefully constructed considering econometric theory on cross-sectional datasets and thus needs cross-sectional datasets to be tested. Time series datasets are largely available, while the same is not true for cross-sectional datasets. Considering availability and the quality of datasets, the following ones have been chosen to compare EGP with traditional feature-selection econometric algorithms: "Concrete Compressive Strength"; "Housing" and "Airfoil Self-Noise (Nasa)". All information on datasets can be found in UCI Machine Learning Repository [19].

# Table 2: EGP and traditional feature-selection econometric algorithms

Position	Algorithm	Adjusted R*2 Training Set	Adjusted R*2 Test Set
1	EGP-Regression	0,799	0,822
2	Partial Least Squares Regression	0,597	0,658
3	Ridge Regression	0,604	0,643
4	Stepwise Forward Regression	0,604	0,640

Dataset : Housing

Position	Algorithm	Adjusted R*2	Adjusted R*2
		Training Set	Test Set
1	EGP-Regression	0,769	0,746
2	Stepwise Forward Regression	0,739	0,710
3	Ridge Regression	0,699	0,673
4	Partial Least Squares Regression	0,624	0,572
(to details on the dataset: https://archive.ics.uci.edu/ml/datasets/Housing			

Dataset : Airfoil Self-Noise (Nasa)

Position	Alith	Adjusted R*2	Adjusted R <sup>*</sup> 2
	Algorithm	Training Set	Test Set
1	EGP-Regression	0,629	0,673
2	Stepwise Forward Regression	0,516	0,451
3	Partial Least Squares Regression	0,480	0,418
4	Ridge Regression	*1	"1

(to details on the dataset: https://archive.ics.uci.edu/ml/datasets/Airfoil+Self-Noise) (\*1: Adjusted R\*2 for Training and Test Sets were inconclusive in sinal and were not reported)

The main reason EGP outperforms traditional econometric feature-selection methods is its capability to explore the regressors' and models' search space. The number of regressors for each of the algorithms shows it. Partial Least Squares Regression, Ridge Regression and Stepwise Forward Regression have at most 8, 13, and 5 regressors, respectively, in its generated regressions for each dataset. For Housing Dataset, as an example, EGP generates models with 50 statistically significant regressors

or more. EGP explores non-linearity of features, by multiplying different features, maintaining models with linear structure.

## 5 CONCLUSION

EGP was successful in achieving its objective of generating high accuracy regressions with potential interpretability of parameters. Feature and model selection performed well, when comparing with traditional methods, as previously shown in the results. Statistical significance proved to be a powerful feature selection tool, directly and efficiently identifying introns and controlling bloat.

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