

Use of CART and CHAID Algorithms in Karayaka Sheep Breeding

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Abstract

The aim of this study was to determine the effect of some factors (sex, birth type, farm type, birth weight and weaning time) on weaning weight through CART and CHAID data mining algorithms. The classification and regression trees are modern analytic techniques that construct tree-based data-mining algorithms. Regression trees are used for the purpose of preliminary selection of the traits affecting the continuous dependent variable. The studied data were consisted of 366 records from Karayaka sheep breed. The CHAID algorithms results revealed that; predictors such as weaning time, sex and farm type statistically influenced weaning weight. Regression tree diagram constructed by CART algorithm depicted that birth type was effect the weaning weight, and in this tree weaning time of single born lambs was affected the birth type. The predicted values and original values were correlated ($P<0.05$). As a result, it could be suggested that CHAID algorithm was found more useful biologically than CART.

Keywords: CART, CHAID, Karayaka, Weaning weight

CART ve CHAID Algoritmalarının Karayaka Koyun Islahında Kullanımı

Öz

Bu çalışma, süttten kesim ağırlığı üzerine bazı faktörlerin (cinsiyet, doğum tipi, işletme tipi, doğu ağırlığı ve ölçüm zamanı) CART ve CHAID veri madenciliği algoritmaları ile belirlenmesini amaçlamaktadır. Sınıflandırma ve regresyon ağaçları veri madenciliği kapsamında olan modern analitik yöntemler sınıfında yer almaktadır. Regresyon ağaçları, sürekli bağımlı değişkeni etkileyen özelliklerin ön seçimi amacıyla kullanılmaktadır. Çalışmada Karayaka koyun ırkına ait 366 kayıt veri olarak kullanılmıştır. Sonuç olarak; CHAID algoritmasına göre ölçüm zamanı, cinsiyet ve işletme tipi süttten kesim ağırlığı üzerinde önemli derecede etkili bulunmuştur. CART algoritmasına ait sonuçlar ise süttten kesim ağırlığı üzerine doğum tipinin etkili olduğunu göstermiştir. Bu ağaçta tekiz kuzuların ölçüm zamanının doğum tipinden etkilendiği anlaşılmıştır. Tahmin edilen ve gözlenen değerler yüksek ilişkili bulunmuştur ($P<0.05$). Sonuç olarak, CHAID algoritmasının CART algoritmasına göre biyolojik olarak daha kullanışlı olduğu belirlenmiştir.

Anahtar sözcükler: CART, CHAID, Karayaka, Süttten kesim ağırlığı

INTRODUCTION

In general, the aim of animal breeding is to genetically improve populations of livestock so that they produce more efficiently under the expected future production circumstances. Genetic improvement for economic traits is achieved by selecting the best individuals of the current generation and by using them as parents of the next generation ^[1]. To achieve the aim of animal breeding, evaluation of the data is very important manner.

Necessary data for animal breeding is generally consisting of many factors which make the data multidimensional. When the number of factor increases, interpretation of

the results become difficult. In this case, overlooking the important details will be unavoidable. Fail to satisfy of conventional analysis tools on complex data, new approaches such as data mining have begun to use. Data mining is an entire process of applying a computer-based methodology, including new technologies, to discover knowledge from data ^[2].

The classification and regression trees are modern analytic techniques which are member of data-mining. They allow for building graphic easily-comprehensible models used to describe and to predict the phenomenon expressed in both the nominal and the ordinal scale. The classification and regression trees can also be used for the purpose



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of preliminary selection of the traits which have a statistical effect on the dependent variable [3]. Classification and regression trees are machine-learning methods for constructing prediction models from data. The models are obtained by recursively partitioning the data space and fit a simple prediction model within each partition. The partitioning can be represented graphically as a decision tree. Classification trees are designed for dependent variables that take a finite number of unordered values, with prediction error measured in terms of misclassification cost. Regression trees are for dependent variables that take continuous or ordered discrete values, with prediction error typically measured by the squared difference between the observed and predicted values [4-6].

Some studies reported significant information on usability of data mining algorithms in sheep and goat breeding. In recent times, many studies have been reported on applying CART (Classification and Regression Trees), CHAID (Chi-square Automatic Interaction Detection), exhaustive CHAID and MARS (Multivariate Adaptive Regression Splines) algorithms for various animal species in animal science for regression type problems [7-10]. However, the application of the mentioned algorithms is still scarce for Karayaka sheep. Therefore, in this study, we aimed to compare the CART and CHAID algorithms of classification and regression trees to predict weaning weight by means of the significant ones among sex, birth type, farm type, birth weight and weighting time in Karayaka sheep.

MATERIAL and METHODS

In this study, to predict weaning weight through CART and CHAID tree-based algorithms, 366 records of Karayaka sheep breed were taken from three different farms in the year 2017. Sex, birth type, farm type, birth weight and weighting time (age as days) variables were used as possible explanatory variables to predict weaning weight. Descriptive statistics for dependent and explanatory variables were given in Table 1.

This method was modified and extended by using some algorithms to minimize an estimate of misclassification error [11].

The classification and regression tree algorithm contains three important tasks. The first task is how to segment data at each step?, the second task is when to stop segmentation?, and the last one is how to predict the value Y for each X in a segment? [5,11,12]. The classification and regression trees begin with a single root node for response variable. The tree is constructed by splitting the whole data into nodes or sub-groups by using all the independent variables. This process goes on until the requirements of homogeneity are met on any child node [13,14]. It is aimed with obtaining terminal nodes in order to increase proportion of variance among nodes [15].

The most popular algorithms used in decision trees are CART and CHAID tree-based algorithms. These algorithms are nonparametric methods which allow using nominal, ordinal and continuous variables [12]. The term "Regression tree" is used for the tree that its response variable is continuous [16].

The response variable is a continuous and explanatory variables can be continuous or categorical in CART algorithm and it creates binary split. In the CHAID algorithm, the response variable can be continuous and categorical. But, explanatory variables are categorical variables only (can be more than 2 categories) and it can create multiple splits [17,18]. The explanatory variable has continuous structure will turn into categorical structure with use of CHAID algorithm [19]. CART algorithm has a characteristic of continuation of CHAID algorithm [19,20]. The aim is to produce homogeneous data sets as much as possible. CART algorithm produce more homogeneous groups than CHAID using pruning [21,22]. Gini, Twoing and Ordered Twoing impurity or diversity measures can be used for categorical response variables, but for continuous

Table 1. Descriptive statistics

Parameter		N	%	Birth Weight (Mean±Std. Deviation)	Weaning Weight (Mean±Std. Deviation)	Weighting Time (Mean±Std. Deviation)
Sex	Male	198	54.1	3.42±0.48	23.28±4.58	100.49±13.76
	Female	198	45.9	3.23±0.54	22.22±4.71	101.43±13.58
	Total	366	100	3.33±0.51	22.79±4.66	100.92±13.66
Birth type	Single	332	90.7	3.45±0.35	23.62±4.05	100.53±13.96
	Twin	34	9.3	2.10±0.09	14.63±0.45	104.70±9.70
	Total	366	100	3.33±0.51	22.79±4.66	100.92±13.66
Farm type	Farm 1	95	26.0	3.12±0.75	21.3±5.37	97.87±11.67
	Farm 2	110	30.0	3.4±0.45	23.89±4.33	107.09±11.32
	Farm 3	161	44.0	3.41±0.34	22.93±4.21	98.51±14.87
	Total	366	100	3.33±0.52	22.79±4.66	100.92±13.67

variables, Least-Squared Deviation (LSD) or Least Absolute Deviation (LAD) can be used [19,23].

The best algorithm produces minimum SD_{Ratio} (Standard Deviation Ratio), MAD (Mean Absolute Deviation), RMSE (Root Mean Square Error) and coefficient of determination (R^2) criteria for goodness of fit. Standard Deviation Ratio, MAD and RMSE can be written as [24,25];

$$SD_{Ratio} = \sqrt{\frac{\frac{1}{n-1} \sum_{i=1}^n (\varepsilon_i - \bar{\varepsilon})^2}{\frac{1}{n-1} \sum_{i=1}^n (Y_i - \bar{Y})^2}},$$

$$RMSE = \sqrt{\frac{\sum_{i=1}^n (Y_i - \hat{Y}_i)^2}{n}},$$

$$MAD = \frac{1}{n} \sum_{i=1}^n |Y_i - \hat{Y}_i|$$

$$R^2 = \left[1 - \frac{\sum_{i=1}^n (Y_i - \hat{Y}_i)^2}{\sum_{i=1}^n (Y_i - \bar{Y})^2} \right].$$

All statistical analysis was performed using IBM SPSS 20.0 via Ondokuz Mayıs University license. The best algorithm will produce the lowest goodness of fit value for mentioned statistics.

RESULTS

Decision tree diagrams to estimate weaning weight by CART and CHAID algorithms are depicted in Fig. 1 and Fig. 2, respectively. CART was seen to produce more branches compared with CHAID tree-based algorithm. Both diagrams reflected environmental factors that can be effective on weight trait, more understandably.

For CART algorithm, Node 0 was split into two smaller subgroups as single birth (23.629±4.052; n=332) and twin birth (14.634±0.456; n=34) according to birth type. These results showed that the effect of single birth was higher than twins on weaning weight as expected. Nodes derived for single lambs had heavier average than those derived for twin lambs.

For single birth, the branch was divided into two smaller subgroups according to age predictor as Node 3 (the subgroup of the single lambs with the age≤91.5 [18.863±2.545 kg; n=94]) and Node 4 (the subgroup of the single lambs with the age>91.5 (25.512±2.796 kg; n=238)). For twin birth lambs, binary branching was occurred with Nodes 5 and 6. Node 5 represented the subgroup of the twin born lambs with the age≤108.0 (14.372±0.362 kg; n=20), whereas

the subgroup of the twin born lambs with the age>108.0 (15.008±0.283; n=14) was named Node 6. From Fig. 1, it was understood that the heaviest lamb average was obtained with the subgroup of the single lambs with birth weight>3.195 kg and age>112.500 days (28.344 kg).

Pearson correlation coefficient between weaning weight and predicted weaning weight for CART Algorithm was estimated as 0.938 (P<0.01).

For CHAID algorithm weaning weight 22.794±4.663 with n=366 began with Node 0 and the tree had 30 child nodes. The age variable divided into six child nodes at first depth of the tree structure. Predictive performances of the CART and CHAID algorithms are presented in Table 2. The influential predictors of the CHAID algorithm were found as age, birth weight and birth type, and farm. Nodes 1, 2, 7, 11, 12, 13, 15, 16, 17, 19, 20-30 were terminal nodes. Node 0 was divided by age predictor into six smaller subgroups i.e. (Nodes 1-6) at the first tree depth. Node 3 was split into three smaller subgroups i.e. (Nodes 7-9) at the second tree depth according to birth weight. Node 4 was branched into two smaller subgroups. Each of Nodes 5 and 6 was divided into four smaller subgroups (Nodes 12-15) and (Nodes 16-19) at the second tree depth according to birth weight. At the third tree depth, Nodes 8-10 were divided into two or three smaller subgroups according to farm. Nodes 14 and 18 were exposed to a binary partition at the third tree depth. At the first tree depth, Node 19 (the subgroup of those with age>114 days birth weight>3.5 kg) produced the heaviest average body weight with 29.327 kg, as expected.

Pearson correlation coefficient between weaning weight and predicted weaning weight for CHAID Algorithm was estimated as 0.937 (P<0.01).

Pearson correlation coefficient between weaning weight and predicted weaning weight for CHAID Algorithm.

DISCUSSION

Results displayed that the partition at the first depth in the tree structures was based on birth type in CART algorithm and weighting age in CHAID algorithm. So, the most predictor variables were different according to the specified algorithms. CART algorithm produced five levels of branching where CHAID algorithm produced three levels of branching. It could be said that CHAID algorithm could be interpreted more easily [12].

For these algorithms, the estimation of SD_{ratio} value smaller than 0.40 was an indicator of the good fit means [12,26,27]. Both MAD and RMSE produced nearly equal values, coefficient of determination were found equal for both algorithms.

To predict weaning weight from sex, birth type, farm type,

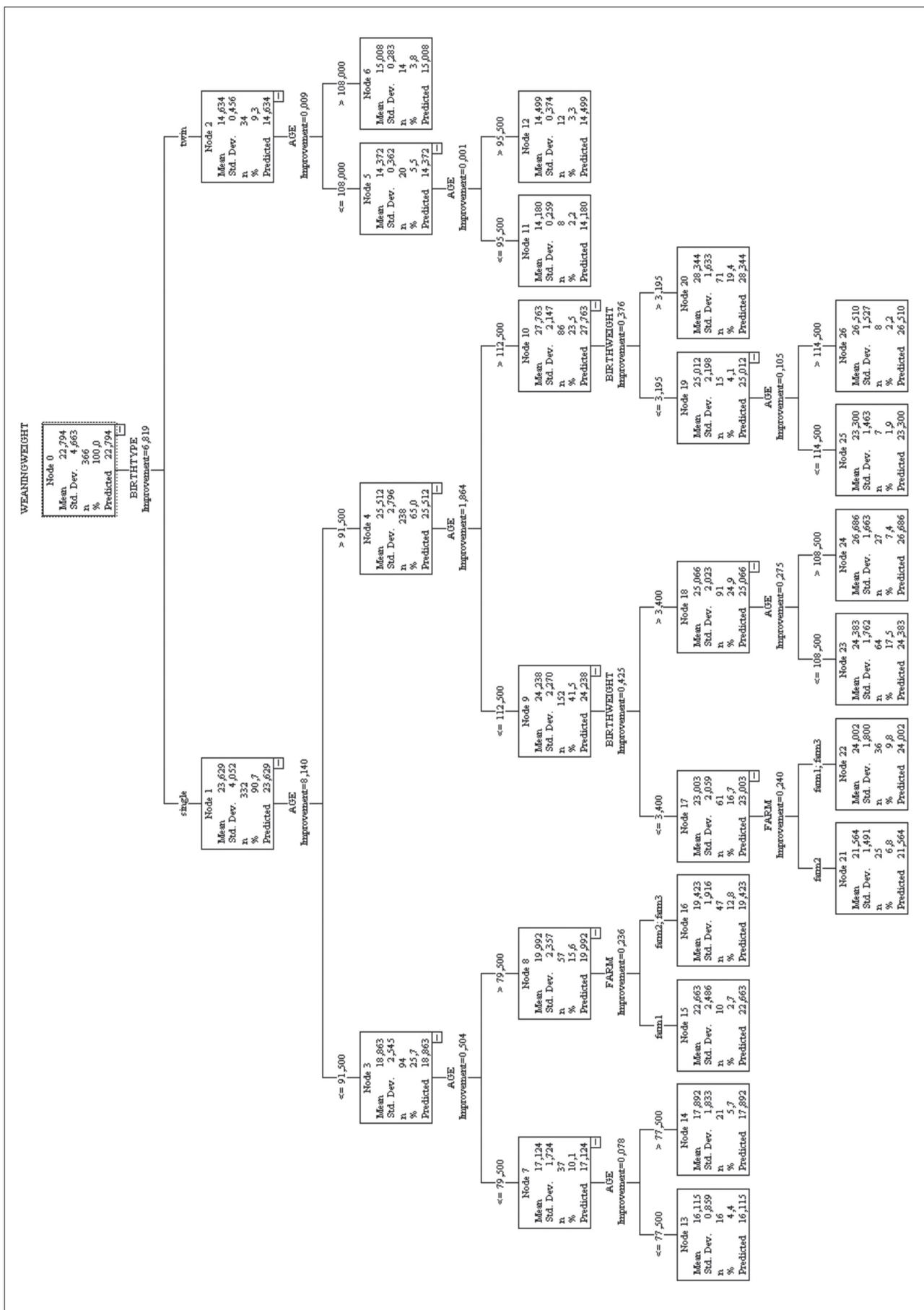


Fig 1. The decision tree diagram obtained by CART

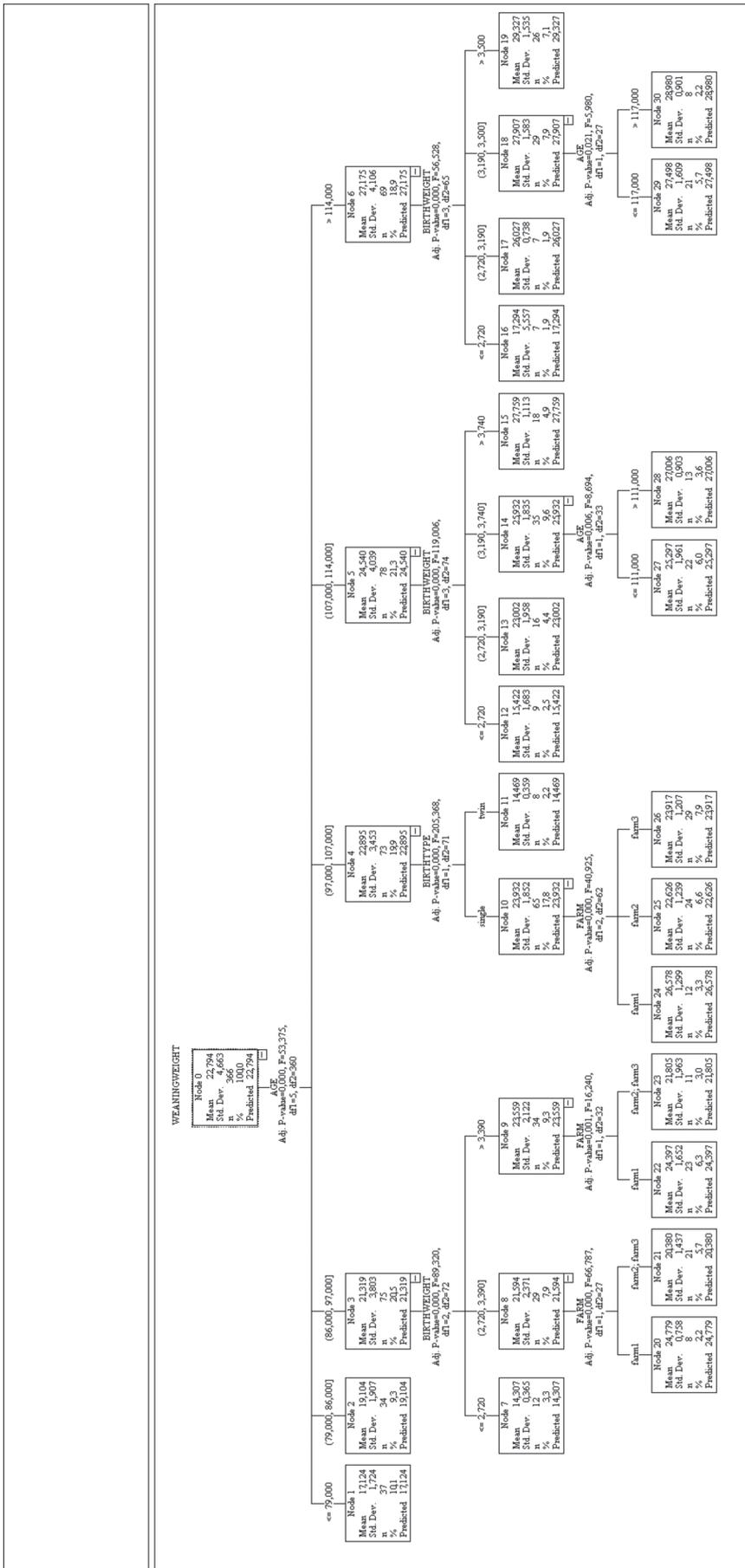


Fig 2. The decision tree diagram obtained by CHAID algorithm

Table 2. Comparison criteria for CART and CHAID algorithms

Algorithm	SD ratio	MAD	RMSE	R ²
CART	0.347	1.195	1.612	0.88
CHAID	0.348	1.192	1.623	0.88

birth weight and weighting time (age) variables, CART and CHAID algorithms can be used interchangeable. The sex variable was excluded from the model in both algorithms. This result was biologically interesting for sheep breeding. Both two methods had similar fitting criteria; however, interpretation of CHAID algorithm because of less branching was more user friendly than CART algorithm. CART algorithm had sub branching over the same variable, which obstruct the interpretation. Similar findings also supported the statements of some earlier authors^[10,12,15,25].

Also, the-tree based algorithms could be applied as a remarkable alternative for the data of response surface designs^[10].

Regression tree aims to repeatedly partitioning the population into different child nodes where the variation of response variable is minimum within and maximum between the child nodes. Also, it aims to balance predictive accuracy and complexity with interpretation of model. Advantage of easy interpretation for both response and predictor variables with visual diagrams are superiority of regression tree. Due to nonparametric properties, it does not require any parametric assumptions. Results of this study showed that CART and CHAID algorithms can be used interchangeable.

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