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IMPACT OF MULTI-OBJECTIVE GENETIC PROGRAMMING TREE REPRESENTATIONS ON FEATURE EXTRACTION AND CLASSIFICATION

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ABSTRACT

In this paper three different genetic programming tree representations are used for classification. Two of them are employed as direct classifiers while the last one works as feature extractor before applying a simple threshold classifier. Each type of representation is discussed with the needed modification required for applying the evolutionary operators including tree generation, crossover and mutation. The three GP methods are applied to real world five datasets from the UCI machine learning database to verify approaches. The performance of the three approaches is compared to conclude the most suitable tree representation for feature extraction and classification

Keywords: Genetic Programming, Feature Extraction, Classification, Tree Representation, Multi-Objective.

1. INTRODUCTION

1.1. Feature Extraction

Many theoretical results have been obtained in the classification domain. Nonetheless, feature extraction retains a key position in the field since the performance of a pattern classifier is wellknown to be enhanced by proper preprocessing of the raw measurement data – this topic is the main focus of this work. Fig. 1 shows a prototypical pattern recognition system in which a vector of raw measurements is mapped into a decision space. In this paper focuses on comparing between using Multi-objective Genetic Program (MOGP) as a feature extractor and employing the same technique of MOGP to directly generate classifiers.



Figure 1. Prototypical Pattern Recognition System.

Designing feature extraction stage of a classifier usually requires deep domain-specific knowledge. Ideally, it is required to have some measure of class separability in the transformed decision space to be maximized. Most importantly, domain- independent feature extraction methodology exists to create or search for good feature extractors.

Often the focus here is on reducing the dimensionality of the problem by projecting the data down onto a sub-space which captures the greatest amount of variability. Even within the constraint of a fixed training set, optimality is hard to guarantee with such methods.

For feature selection, wrappers methods are used. Wrapper methods, as illustrated in Figure 2, use a search strategy to iteratively navigate in feature subset space until a stopping condition is met, while evaluation of the subset selection uses the feedback obtained from the classifier used in order to optimize the performance of the classifier. Wrapper methods use cross-validation techniques or set performance bounds to validate the performance of the classifier. Wrapper methods can find the most useful features but are prone to over-fitting.



Figure 2. Prototypical Wrapper Feature Selection Approach[14].

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In the sense that the feature extraction preprocessing stage is a transformation or mapping from input space to decision space, for a given classification problem the mapping which maximizes the separability of the classes in decision space is investigated. Thus the feature extraction can be regarded as finding an optimal sequence of operations subject to some criterion

1.2. Genetic programming (GP)

It's an evolutionary learning technique that offers a great potential for classification. GP is a very flexible heuristic technique that allows us to use complex pattern representations such as trees. For example, any kind of operation or function can be used inside that representation and domain knowledge can be used in the learning process.

The application of GP to classification offers some interesting advantages, the main one being its flexibility, which allows the technique to be adapted to the needs of each particular problem. GP can be employed to construct classifiers using different kinds of representations (decision trees, classification rules and discriminant functions). GP can be useful not only for inducing classifiers, but also for other preprocessing and post processing tasks aimed at the enhancement of classifiers.

Indeed, GP has been used before to optimize feature extraction and selection [2][3]. Sherrah et al. [4] proposed an Evolutionary Pre-Processor (EPrep) system which used GP to evolve a good feature mapping by minimizing misclassification error.

1.3. Bloat in Genetic Programming

The chromosomes in a genetic programming (GP) population will 'bloat' – that is, grow without limit without any accompanying improvement in fitness. Research on the causes of bloat has been recently summarized in [5]. Indeed, Langdon and Poli [6, 7] have shown that any variable-length representation suffers from bloating. They have summarised the three main approaches to control bloat in GP:

• Limiting the maximum permissible tree depth (or size) to a pre-defined value.

- Tailoring the genetic operators.
- Employing parsimony to exert selective pressure which favours smaller trees.

In this paper the parsimony pressure approach is addressed. It use a multi-objective method in which the (strictly) non-commensurable objectives of problem-specific error and tree complexity are handled in a Pareto optimisation framework [8]. The results from the Pareto framework is not a single unique solution but a set of equivalent solutions which lie on a Pareto front (or surface) in objective space and which delineate the fundamental trade- offs in the problem. No point on the Pareto front can be modified to improve one objective without simultaneously another. degrading Multi-objective GP (MOGP) has a number of advantages: As well as controlling bloat very effectively, it does not require a pre-determined depth-limit parameter and the tree depth is free to adjust to suit the problem at hand.

This paper is organized as follows: a generic framework to evolve optimal feature extractors with multiple objectives is presented in Section 2. Section 3 and section 4 represent GP implementation and experimental results. Comparison with three Genetic programming classifiers is also introduced by other researchers is presented in Section 5. Conclusion is given on Section 6.

2. METHODOLOGY

2.1. Multiple Objectives

Within the multiobjective framework, a twodimensional fitness vector of objectives is used comprising: Tree complexity and misclassification error as follows:

2.1.1. Tree complexity measurement

As pointed-out above, there is a danger that trees evolved by GP will become very large due to tree bloat. The huge trees could produce an extremely small error over the training set but a very poor error estimated over an independent validation set.Broadly, for a given training error, the simpler individual is preferred. Thus node count in the tree as a straightforward measure of tree complexity is used as one of the fitness vector elements driving the evolution. Thus a selective pressure that favors small trees is imposed, all other things being equal.

2.1.2. Misclassification error

The second element in the fitness vector is the conventional one of the fraction of misclassified patterns counted over the training set.



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2.2. Multiple Objectives Genetic Programming (MOGP)

In this work MOGP is employed in two ways. The first one is to evolve the whole classifier. While in the second it is employed to find optimal feature extraction for pattern recognition system. Since the input pattern is projected to a one- dimensional decision space, a simple threshold classifier is used. The threshold is adapted as part of the fitness value to give the minimum error. This means its tried to evolve a feature extractor which maps the original pattern space into a new feature space where thresholding is able to yield the smallest possible misclassification.

Kumar and Rockett [9] proposed the Pareto Convergence Genetic Algorithm (PCGA) used an elitist, steady-state strategy. In each generation, only two individuals are generated and added to the population to replace the worst two individuals. The ranking mechanism of Fonseca and Fleming [10] was used. Roulette wheel selection was employed to select two individuals for crossover. Mutation was always applied to the results of the crossover operation. Their method improved the sampling on the Pareto front to ensure convergence of the evolution without the need for sharing/niching techniques with significantly less computational effort.

In this work non-destructive, depth-dependent crossover [11] is used in order to avoid the breaking of building blocks.. The sub-tree is chosen based on its complexity (i.e. the number of nodes) using the depth-fair operator . Thus one of the sub-trees at the chosen depth will be picked by roulette wheel selection, biased in its complexity. Having chosen depth, d in a tree, there are N sub- trees, each comprising M1, M2,..., MN nodes, respectively. The probability of selecting the i-th sub-tree is given

$$\Pr[i] = \sum_{j=1}^{i} M_j / M \qquad i \in [0...N]$$
1
(1)

The crossover operation is self-explanatory; in mutation, the selected sub-tree is then replaced by a new, randomly created sub-tree attached at the original mutation point. Only those offspring which dominate either of their parents are retained. In this way, the algorithm was able to maintain diversity in the population and avoid being trapped in local minima in the early stages.

3. GENETIC PROGRAMMING IMPLEMENTATION

As a basis for comparison with MOGP, three model for extractions are used [15] (Decision trees, Rule base and Discrimint Functions).

3.1. Decision tree

A decision tree contains zero or more internal nodes and one or more leaf nodes. All internal nodes have three children nodes. All internal nodes contain splits, which test the value of an expression of the attributes(X) as show in figure.4. Arcs from an internal node t to its children are labeled with distinct outcomes of the test at t.



3.2. Rule-Based

Rules are a simple and easily interpretable way to representation. The rule has two parts, the antecedent and the consequent. The rule antecedent contains a combination of conditions for the predicting attributes(X). Typically, conditions form a conjunction by means of the AND logical operators, but in general any logical operator can be used to connect elemental conditions. The rule consequent contains the value predicted for the class. This way, a rule assigns a data instance to the class pointed out by the consequent if the values of the predicting attributes satisfy the conditions expressed in the antecedent; hence a classifier is represented as a rule set.



Figure 4. Rule Base Representation

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3.3. Discriminant functions

Discriminant functions are mathematical expressions in which different kinds of operators are applied to the attributes(X) of a data instance that must be classified. A single output value is computed from the operations performed on the values of the attributes. The value computed by the function indicates the class predicted. Usually, this is accomplished by means of a threshold or set of thresholds. For binary classification problems, a single function is enough; if the output value is greater than a given threshold, the example is assigned to a certain class, otherwise it is assigned to the other one.



Figure 5. Discriminant Functions representation.

For Each of these representations there are a set of constraints should be checked while generating trees or sub-trees during the evolution process. Also, a set of constraints should be taken in consideration while applying cross-over and mutations in order to keep the correct representation of the tree.

• *Generation Decision Tree representation*: has done with some rule where the tree divided to three leaf

Left leaf : should be constant node *Middle leaf* : Ternary Node or Class Node *Right leaf:* Ternary Node(feature) or Class Node

- *Generation Rule Based representation*: where tree divided in to binary subtree (leafs) where each leaf has logical function, for last sub tree has comparative function with binary terminal node which represented as feature or constant node.
- *Generation Discriminant Functions representation*: where tree divided in to binary subtree (leafs) where each leaf has mathematical function ,for last sub tree has terminal node which represented as feature or constant node.
- Crossover Decision Tree representation:

depends on selected leaf if

Left leaf : the crossover is performed only within corresponding constant node *Middle leaf* : the crossover is performed only within TernaryNode or ClassNode *Right leaf*: the crossover is performed within TernaryNode or ClassNode

- *Crossover Rule Based representation*: The crossover that performed within constant node should be to corresponding constant node. The crossover that performed within feature node should be to corresponding feature node author wise performed crossover sub tree to sub tree (comparative or logical function node).
- Crossover Discriminant Functions representation: The crossover that performed within constant node or feature node should be to corresponding constant or feature node. The crossover performed sub tree should be to sub tree (mathematical function node).
- *Mutation Decision Tree representation*: depends on selected leaf if :

Left leaf : the mutation is performed only within corresponding constant node Middle leaf : the mutation is performed only within Ternary Node or Class Node Right leaf: the mutation is performed within Ternary Node or Class Node

 Mutation Rule Based representation: depends on selected leaf if: Right leaf : if constant node the mutation is performed only within constant node. Left leaf: if feature node the mutation is

performed within feature Node For any leaf : if comparative or logical operation node the mutation is performed only within comparative or logical node.

• *Mutation Discriminant Functions representation:* The mutation that performed within constant node or feature node should be to constant or feature node. The mutation performed sub tree to sub tree (mathematical function node).

4. EXPERIMENTAL RESULTS

In this section the three techniques performance is examined across a representative range of two-class classification problems from the UCI Machine Learning database [1].

The work presented here considers an

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extensive set of comparisons across five datasets with two-class learning problems. For each dataset a statistical comparison of the classification performance between the three MOGP tree representations and a range of established classifiers is presented.

4.1. The UCI Datasets

The five datasets used in the current work are from the UCI Machine Learning database [1]:

Table 1. UCI Dataset

Name	Size and Distributions	Features
Glass	153 = 87 (float) + 76 (non-float)	9
BUPA	345 = 200 (Benign) + 145 (Malignant)	6
AUS	569 = 357 (Benign) + 212 (Malignant)	30
PID	532 = 355 + 177 (Diabetic)	7
WBC	699 = 458 (Benign) + 241 (Malignant)	10

4.2. Ten-Fold Cross Validation

This estimation will depend on the dataset partitioning method applied. TEN-fold crossvalidation is commonly used to obtain a more accurate estimate of the empirical risk. In Ten-fold cross-validation the dataset is divided into ten disjoint subsets. One of the subsets is used as the test set and the training dataset is formed using the other N-1 subsets. The cross-validation process is then repeated ten times and the average of the errors estimated over these ten folds is computed. This method reduces the sensitivity of the estimation to how the data gets divided as eventually every example in the dataset is used both for training and testing, but this increase the computational effort needed to make an evaluation. Another variation of this method can be done by splitting the dataset into two folds and repeating this for n different splitting. For each splitting, one of the datasets is used as a training set and the other as the test data. Then the experiment is repeated interchanging the roles of the datasets.

4.3. Statistic

The mean of the test errors for three classification algorithms across the five datasets are summarized in Table 2. The counted number of the node in the trees and the calculated average of the errors over each test fold of the dataset for every classifier is shown. Reassuringly, the MOGP algorithms return the lowest mean error for each dataset with number of nodes. Overall results show that the Discriminant Functions algorithm are better than of the other algorithms. Although this result needs to be treated -with some caution to be sure

the statistical significance of these differences is clear.

Table 2. Mean Error Comparisons Of Classifiers On
Each Dataset

	Classifiers					
Dataset	RB	node	DF	node	DT	node
Glass	0.385	7	0.134	55	0.172	34
BUPA	0.276	59	0.213	65	0.234	40
PID	0.265	55	0.190	65	0.211	43
WBC	0.086	13	0.015	15	0.019	25
AUS	0.129	33	0.132	22	0.115	25

In order to quantitatively compare statistical significance of the cross-validation experiments the Alpaydin F-statistic is computed for the three MOGP algorithms compared with each other. The comparisons are summarized in Table3 where MOGP_DF represents superiority over the other algorithms.

Table 3. F-Statistic Comparisons Of Classifiers On Each Dataset

	Classifiers		
Dataset	DF_DT	RB_DT	RB_DF
Glass	1.7	216.1	87.4
BUPA	2.57	9.73	32.66
PID	4.361	62.71	30.4
WBC	2.539935	135.93	597.4
AUS	3.16	1.557	16.9

5. COMPARISONS WITH INTERPRETATION

The mean error rates from these earlier studies are summarized, in Table 4. Typically, error rates were estimated using ten-fold cross-validation. The results of Bot & Langdon [6] are the mean validation error of the best individual of 30 runs. Table shows that MOGP has best result over all datasets.

Table 4. Reported Error Rates For Other Evolutionary
Feature Detection / Classification Algorithm [14]

Training	Dataset			
Algorithem	WBC	PID	BUPA	Glass
Muni	0.0281	N/K	0.3007	N/K
Bot	N/K	0.305	0.416	0.48
Bot &	N/K	0.25	N/K	0.368
Langdon				
Krawiec	N/K	0.2359	N/K	0.3361
Loveard	0.032	0.242	0.308	N/K
Lim MIN	0.03	0.22	0.29	N/K
MOGP	0.02634	0.2057	0.2644	0.2271

Although tree size is one of the multiple objectives that has been used to suppress tree bloat, a number of the trees are not of the

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absolute minimum size and contain a few redundant sub-trees. Figs. 6 - 10 show the trees which have been generated by the evolutionary algorithms.

It is clear, however, that these are not completely optimal in that the identical classification performance could be obtained in some cases with slightly smaller trees.

Nonetheless, the work presented here produces near- optimal trees which are reasonable for a stochastic search method such as genetic programming and an advance on previous work on feature extraction. In practice, any redundant sub-trees could be easily removed by hand from the final solution.

6. CONCLUSION AND FUTURE WORK

This paper a demonstrate using of multiobjective genetic programming (MOGP) to evolve an "optimal" feature extractor which transforms input patterns into a decision space such that class separability is maximized.

In comparison with a number of other representative classifier paradigms, the performance of MOGP method turns out to be better. The use of multiple objectives, particularly classification error objective has been shown to be effective in guiding and speeding the optimization.

The node number objective employed penalizes an individual according to its complexity. This appears to be essential in order to prevent tree bloat as well suppressing over-fitting of the training set leading to poor generalization.

The three MOGP algorithms were applied to machine learning datasets from the UCI database. The MOGP_DT has less error classification than MOGP_RB and it is shown that MOGP_DF has better result than others in multi objectives technique.

Finally, developmental GP has the potential to contribute to the discovery and exploitation of knowledge in databases in significant ways [9]. Although only binary classification problems are treated in this paper, extension to multiple classes is a logical development and is currently underway.

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