

# Associative Semantic Ranking Models of Satellite Images

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**ABSTRACT** -Associative procedures for content-based image position by semantics are gorgeous due to the correspondence of produced models to human models of considerate. Though they tend to return results that are better unstated by image analysts, the induction of these models is problematic to build due to factors that affect training density, such as synchronicity of visual patterns in same images, over-fitting or under-fitting and semantic representation differences among image analysts. This article proposes a methodology to reduce the complexity of ranking satellite images for associative methods. Our method employs genetic operations to afford faster and more correct models for ranking by semantic using low level features. The added exactness is provided by a decrease in the likelihood to reach local minima or to over fit. The investigates show that, using genetic optimization, associative methods perform better or at similar levels as state-of-the-art ensemble methods for ranking. The mean average precision (MAP) of ranking by semantic was improved by 14% over similar associative methods that use other optimization procedures while maintaining smaller size for each semantic model.

**Keywords:** content-based image ranking, data mining, genetic, satellite images.

## I. INTRODUCTION

Estimation of geospatial imagery is interesting due to high dimensionality of spatial data and to the synchronicity of visual patterns interrelated to several semantics in images. As the rate of image assemblage grows exponentially, it is fetching exceptionally challenging for image analysts to manually cutting knowledge from geospatial descriptions in order to distribute focused information for decision making. This needs the need for systematizing remote sensing data analysis and evaluation. Outdated data approaches, such as statistical methods, have restrictions in terms of distributional assumptions and boundaries on data input which may prevent them from investigating unknown and unexpected relationships in geospatial images. Other customary methods of data mining such as Artificial Neural Networks and Genetic Algorithms (GA) have a black-box distinguishing which makes it difficult for users to apply removed rules to other cases. Besides, data values gain connotation only in the context of the geospatial domain and the reality of multiple semantic interpretations for the same image, which makes it difficult to apply traditional data analysis methods to images. Consequently, new approaches that consider unique appearances of image data have emerged for mining patterns from images.

In content-based image reposition, images are indexed by their visual contents such as color and shapes. Though, these low-level features cannot properly capture the high-level image semantics in a user's mind. Accordingly, recent studies on content-based image retrieval focus on reducing the semantic gap between low-level features and high-level human

semantics by accumulating semantic models that can be used for prediction. A complete review of various semantic models are providing in, where methods for plummeting the semantic gap include using object ontology to define concepts, using machine education methods to associate low-level features to users' semantics, disseminating relevance feedback to learn users' intentions, producing a semantic template to map low-level features to high-level concepts, and merging visual and text content for web image retrieval.

Recent research in the geospatial area providing a variety of in-depth solutions, to epitomize the complex, often covering geospatial knowledge and to assist image predictors in generating necessary domain specific metadata. The research in designates a framework for modeling and image retrieval using directional spatial associations among objects. Content-based image retrieval (CBIR) methods were pragmatic to position satellite images using opportunity associations among low-level features and semantics of interest. The researchers use Latent Dirichlet Allocation (LDA) semi-supervised methods to interpret images with semantic classes. Both supervised and unsupervised methods are shared in the I3KR framework to increase image penetrating capabilities using semantic and content-based information. The professionally retrieve images using indexing structures on the feature space. The application of self-organizing maps to the investigation of man-made structures in multispectral imagery is investigated. The research in proposes the amalgamation of a multi-modal content-based system with complex methods of querying on shape, multi-

object associations, and semantics, while the research in repeatedly detects variations in geospatial images and applies clustering techniques to organize visual pattern variations. The approach in uses ontological knowledge and artificial neural networks to figure semantic models of visual patterns using both low-level and descriptive image features.

Evolutionary algorithms are self-adaptive optimization methods that accomplish the global search in a solution space. They tend to execute the better with attribute interactions when associated to greedy decision algorithms. Genetic Algorithms (GAs) model the space of candidate solutions in chromosome structure where the success of each chromosome is considered with a fitness function. The best solution or most satisfactory solution is based on ordinary selection methods that combine successful features existent in a set of formerly generated models by selection, crossover and mutation. Since knowledge about the search space is gathered during the search process, GAs can eliminate local-maxima traps by adaptively moving the solution space to methodology a global optimal. GAs is applied in various spatial data mining domains.

## II. METHODOLOGY

In this section we present our approach for classification of satellite image regions using genetic operations. For each image in the database we generate a feature space  $F$ . The key feature of the algorithm is that we use sets of suggestion rules among feature subspaces and semantics in a semantic space  $S$  to rank images by semantic. Each set of connotations is engendered and evolved using genetic operations at two levels: the feature and subspace levels. At the feature level, we vary the set of features used to recognize association rules, while at the subspace level we vary the region for the same

feature set that will be used in position. For example, for a 38-dimensional space there  $2^{38}$  are unique arrangements of features. Using genetic operations we aimlessly choose and evolve combinations of features using methods such as crossover, shrink, constant, or grow mutations. Once a grouping of features is selected, we indiscriminately generate and evolve features' subspaces exhibited by sigmoid possibility functions. Additional, sets of feature spaces are used additively to model correlation to a semantic of interest. To appraise which subspace is the most applicable we also apply genetic operations at this level.

### 2.1. Fitness Function

The fitness function for each semantic model is used by the optimization algorithm to regulate which groupings of association rules will better model the suggestion among feature subspaces and semantics of interest. In our study, we use the MAP to regulate the relevance of each feature subspace, a set of families that will form a semantic model.

### 2.2. Encoding

Each produced membership function is painstaking an exon  $\epsilon$  and it is encoded as a decimal string for the sequence  $(\phi, \lambda_1^L, \lambda_2^L, \lambda_3^R, \lambda_4^R)$  using a total 20 decimal digits. The feature  $\phi$  is recorded as the index of the feature in the feature space using four decimal digits, while for each of the sigmoid parameters we store the most significant four digits after the decimal point that resulted after the process of normalization.

### 2.3. Genetic Operations

We perform genetic operations at three levels: exon, gene, and chromosome. Underneath we enumerate the genetic operations that are accomplished on each population which are exemplified in Figure 1 on a shortened two-dimensional feature space composed of object convex area kurtosis (F1) and orientation skewness (F2). In this figure, the perpendicular axis is the relevance feature points to a semantic of interest.

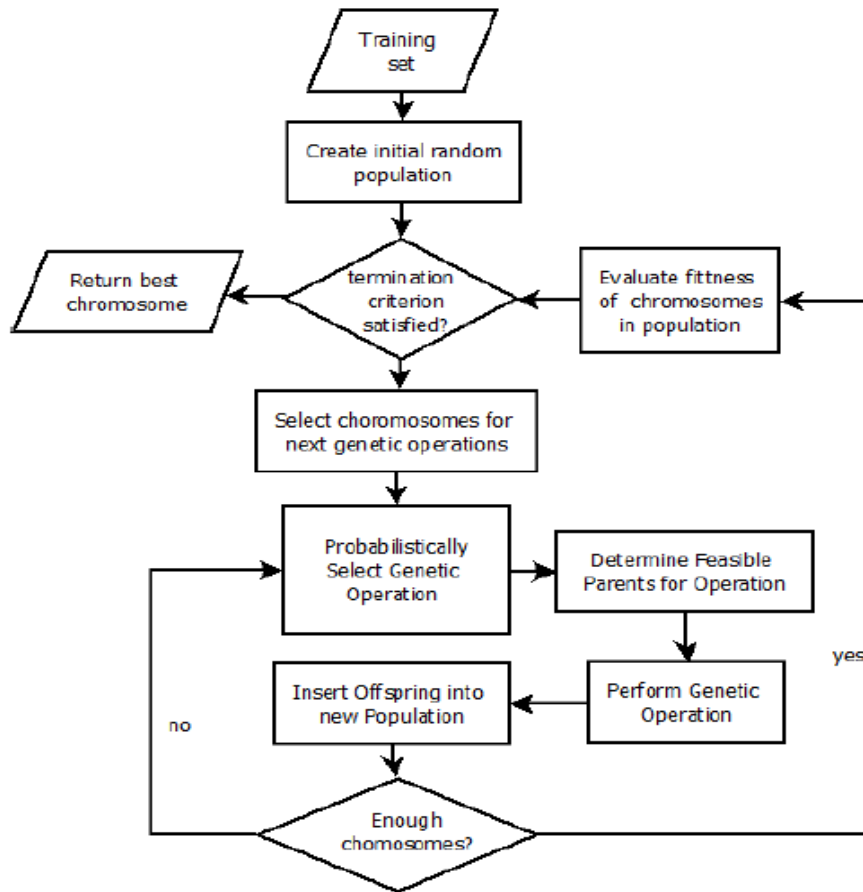


Fig 1. Flowchart for generating a semantic model using genetic operations

Figure 1 shows the flowchart for engendering a semantic model using genetic operations. The input parameters for this process are an exerciseset containing image features that were categorized by image analysts with one or multiple semantics. This algorithm also takes, as input, the following parameters: the number of chromosomes in each generation of population, the maximum number of generations (iterations) the algorithm will execute, and a beginning on the quality of ranking for which the algorithm would dismiss. The algorithm starts with a populace in which each chromosome, gene, and exon was randomly produced. The quality of status is then evaluated using the MAP quantity and it is shown in Equation. The top chromosomes are then nominated as parents for the chromosomes in the next generation, which is engendered using the genetic operation explained in Section 2.3. Lastly, when the termination criterion was met—either the quality of ranking of the top chromosome surpassed the preset threshold or the maximum number of iterations was accomplished—the algorithm returns the most fitted chromosome. This chromosome is

transformed to a semantic model that is used for ranking of new, unlabeled images.

### III. EVALUATION

We considered three investigates to evaluate the relevance of smearing genetic optimization methods to ranking images by semantics: (1) we evaluate the routine of the proposed approach over a large number of genetic operations; (2) we perform an in-depth proportional evaluation of Associative & SFFS and the proposed approach (Associative & Genetic); and (3) we compare the presentation of the proposed method with that of six other methodologies. For each experiment we followed the procedure shown in Figure 2: First, the original data was detached into ten subsets using a stratified strategy to ensure that each semantic class in correspondingly represented in each fold. Next, using a ten-fold iteration approach, data was detached into testing containing a different subset for each fold and training comprising the

remaining folds. Then, ranking models were built on the training data and appraised on testing data.

For the proposed method, we have documented each genetic operation that was performed on the genetic population. This resulted in a number of 90,000 genetic operations for the experimentations over the UCI Stat log Landsat data set and 120,000 genetic operations for the experiments over the WROC data set. The percentage for each specific operation performed is shown in Figure 2. For example, the crossover operations accounted for 57% of all the operations equally disseminated over chromosome and gene mutations. Due to the randomness of the genetic operations, we observed minimal percentile variations for the experiments on the two data sets.

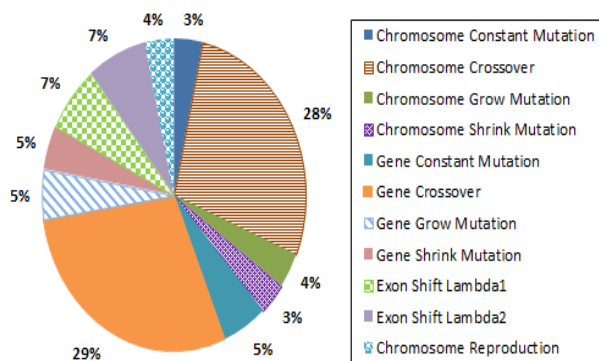


Fig 2 Genetic operations performed as percentage when ranking images by semantics.

#### IV. CONCLUSIONS

I have established an approach for engendering associative models for ranking satellite image regions by land cover. The results of our proportional studies show that the proposed method performs better or has similar presentation to that of other ensemble methods. Our method relates genetic methods to return better precision on new unapproved data while avoiding overfitting by reducing the local minima issues existent in additive models. Overall our results show that the genetic method revealed better association rules faster than the existent additive method. This shows that associative methods offer auspicious alternatives to visual patterns found in images, though they are prone to overfitting. The key to their success is an acceptable learning procedure that is able to evade local minima. Previous associative approaches use suggestion rule mining algorithms to identify relevant feature spaces but suffer from inadequate measure of suggestion rule relevance, such as support and confidence, which are

not optimal for ranking problems. Genetic models have also the advantage of aimlessly selecting and testing new feature subspaces which result in better models in shorter time. Though not precisely measured, training time is an imperative component in any ranking algorithm. As with any other collaborative method, training the proposed method is comparative to the size of the training set, number of rules in a semantic model and number of iterations. This is an enhancement over SFFS methods for which reducing the number of rules in a model requires quadratic complexity of number of rules.

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