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GENETIC DISTANCE MEASURE FOR K-MODES ALGORITHM

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ABSTRACT. K-means algorithm has been shown to be an effective and efficient algorithm for clustering. However, the k-means algorithm is developed for numerical data only. It is not suitable for the clustering of non-numerical data. K-modes algorithm has been developed for clustering categorical objects by extending from the k-means algorithm. However, no one applies this technique for classification of categorical data. In this paper, the k-modes algorithm is introduced for the classification of categorical objects based on Soybean and Nursery databases. Especially, a genetic algorithm is proposed for designing the dissimilarity measure termed Genetic Distance Measure (GDM) such that the performance of the K-modes algorithm may be improved by 10% and 76% for Soybean and Nursery databases compared with the conventional k-modes algorithm. **Keywords:** K-means, K-modes, Genetic algorithm, Categorical data, Numerical data

1. Introduction. In the clustering area, k-means algorithm [6,8] has been adopted for wide applications for the data sets with numerical values. The k-means algorithm is a centroid-based clustering technique. The representatives of the clusters are the centers of the clusters. The applications include the clustering of customers for supermarket, clustering of patients for hospital, codebook design for vector quantization [3,7], and the texture segmentation [10]. Huang proposed the k-modes algorithm for clustering the data sets with categorical values [4]. In fact, the original k-modes algorithm is extended from the k-means algorithm by applying the Hamming distance between objects for the categorical data. In k-modes algorithm, the modes are analogue to the means in the