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Simulated Annealing Based Clustering

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Abstract - Literature review revealed that data Mining , clustering techniques has become an important topic in effective analysis of gene expression data due to its wide application in the biomedical industry . In this paper we propose an efficient and automatic simulated Annealing based clustering approach that is suitable for gene expression analysis . The purpose approach primarily employs similarity- matrix based clustering techniques. In particular a novel validation technique is incorporated for evaluating the quality of the discovered gene expression patterns . We have implemented proposed algorithm and tested their performance using real biological data of cancer of [118 x 60] samples and breast cancer data of [700x9] samples. The results of proposed algorithm were compared with the well - known k-means algorithm with very comparable results. This approach to gene expression analysis that integrates simulated-annealing, clustering and validation techniques simultaneously results in automated, high quality and highly efficient algorithm .