A Genetic Algorithm for Physical Mapping Problems

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1 INTRODUCTION

In this paper, we study a genetic algorithm for solving physical mapping problem. First, the physical mapping problem is transformed to an optimization problem by incorporating biological knowledge and limitations into the objective function. Based on the idea of genetic algorithms, the proposed approach integrates Edge Assembly Crossover (EAX) and Inver-over genetic operators to get the optimal solution. We analyze essential components of the proposed approach as well as implementation details. Our approach is then applied to some widely used test sets and simulated data, real data of this problem. Experimental results indicate that the new approach performs efficiently and precisely to solve physical mapping problem.

2 PROBLEM AND APPROACH

In the physical mapping problem, we use the relation of probes and clones to get the order of probes and then give us the relative positions of the clones in the DNA sequences. There are chimeric, false negative and false positive errors that make it hard to solve. Based on the previous researches (Alzadeh et. al, 1993; Jain and Myers, 1997), we adapt the following function as our objective function for optimization:

\[ F(A) = X \cdot C(A) + Y \cdot P(A) + Z \cdot N(A) + T \cdot M(A) + P \cdot L(A) \]

where

- \( X = - \log \left( \frac{P}{N} \right) \)
- \( Y = - \log \left( \frac{F}{P} \right) \)
- \( Z = - \log \left( \frac{T}{M} \right) \)
- \( T = - \log \left( \frac{M}{A} \right) \)

The structure of our genetic algorithm is shown in Figure 1. We first use the efficient EAX operator (Nagata, 1997) to generate children. If two selected individuals are equal, we select one to perform Inver-over operator (Tao, 1999) to make a chance for generating different connection of genes to prevent from converging at local optimum. In other words, this operator used here to keep the diversity. We think this is a way to avoid the prematurity but also keeps the spirit of local searches.

3 EXPERIMENTAL RESULTS

We have tested our GA on TSP and physical mapping problems. The result is shown as follows. The graphs is the results of our GA tested on real data from chromosome 1. And the table is the results of our GA tested on some standard TSP problem.

<table>
<thead>
<tr>
<th>Problem</th>
<th>Gen</th>
<th>Opt/num</th>
<th>Error</th>
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</thead>
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<tr>
<td>Eil75</td>
<td>10</td>
<td>20/20</td>
<td>0.0000</td>
</tr>
<tr>
<td>Eil101</td>
<td>16</td>
<td>20/20</td>
<td>0.0000</td>
</tr>
<tr>
<td>Lin318</td>
<td>34</td>
<td>18/20</td>
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<tr>
<td>Pcb442</td>
<td>37</td>
<td>20/20</td>
<td>0.0000</td>
</tr>
</tbody>
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Figure 1: Overview of our GA

Figure 2: Graph of results

Figure 3: Table 1

References


