

# A New Multi-Objective Optimization in Solving Graph Coloring and Wireless Networks Channels Allocation Problems

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## ABSTRACT

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Graph coloring problem, a combinatorial optimization problem is being widely applied in solving the channels allocation in wireless networks. This paper exhibits a new evolutionary genetic multi-objective strategy that uses the combined single and multi-parent conflict-gene crossover and combined single and multi-parent conflict-gene mutation operators with clique partitioning to solve the graph coloring and channel allocation problems. The proposed operators minimize problem search space by reducing the expected number of genetic generations. A general fitness function is defined on finding the total conflicting edges in the graph for the initial and particularly the successive generations of individuals in the population. The outcomes of this proposed method are better than the well-known methods and are compared with some of the benchmark graph coloring and channel allocation problems. The devised method of clique partitioning with genetic operators also enhances the successful runs.

Keywords - approximation methods, channel allocation, chromatic number, genetic algorithm, graph coloring, wireless networks

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## I. INTRODUCTION

For a simple graph  $G$  consists of  $m$  edges &  $n$  vertices, the edge & vertex set are represented as  $E(G): \{e_1, e_2, \dots, e_m\}$  and  $V(G): \{v_1, v_2, \dots, v_n\}$  respectively. Also, every  $e_i \in E(G)$  defines a unique end vertex pair  $(v_j, v_k) \in E(G)$ ,  $1 \leq i \leq m$ ,  $1 \leq j, k \leq n$ . The adjacency matrix of  $G$  is represented in  $A(G)$  which is a symmetric matrix of order  $n \times n$ .  $A(j, k) = 1$  if there exists an edge joining  $v_k$  and  $v_j$ . Otherwise,  $A(j, k)$  is set to zero. Graph coloring problem (GCP) finds the least number of colors,  $\chi(G)$ , which are used to assign its  $V(G)$  without assigning the same colors to the adjacent vertices [1]. Recently wireless network are emerging as an important areas in which different soft computing models are required in solving the real-world problems [16]. There are many genetic methods available to find  $\chi(G)$  which requires the search space of  $n!$  [2-3]. The wireless networks capacity is increased when assigning the proper  $m$  channels or edges to  $n$  cells, called Channel Assignment Problem (CAP) which is defined with the constraints: co-channel, adjacent channel, and co-site constraints (CCC, ACC & CSC). If  $G=(V, E)$  is an undirected graph then its clique,  $C$ , is a subgraph that consists of the vertex set  $V'$  such that there is an edge between every vertex pair in  $V'$ .  $C$  is the complete subgraph of  $G$ . The split of  $G$  into a minimum number of cliques such that each vertex belongs to exactly a single clique is called clique partition. A maximum clique is a clique that consists of the maximum number of vertices. Few other methods to find  $\chi(G)$  are Particle Swarm Optimization (PSO), Ant Colony Optimization (ACO), and Cuckoo search [3]. Different methods are discussed to

solve graph coloring and channel allocation problems [9-15].

This paper presents a new genetic method in solving GCP and CAP with a clique partition strategy. Combined Single and Multi-Parent Conflict Gene crossover (CSMPGCGX), Combined Single and Multi-Parent Conflict Gene Mutation (CSMPGGM) operators are newly devised in this paper. These operators are targeted to minimize the expected genetic generations and also reduce the exploration of search space. Section II formulates the mathematical model of CAP. The evolutionary method with partitions of  $V(G)$  into different cliques in assigning proper channels is presented in Section III. Section IV analyses the theoretical results and global convergence of the devised operators. The well-known problems have experimented and their outcomes are compared with some of the existing methods and are presented in Section V. Section VI concludes the research.

## II. FORMULATING A MATHEMATICAL MODEL OF CAP

For the wireless network with  $m$  channels and  $n$  cells, channel assignment is an optimization problem that is defined with respect to some interference constraints. The channels assignment is defined in the vector  $F$  which represents a symmetric matrix of size  $n \times m$ . For each  $j$  and  $k$  ( $1 \leq k \leq m$  and  $1 \leq j \leq n$ ),  $F(j, k) = 1$  if cell  $j$  is assigned the  $k$ th channel; otherwise  $F(j, k) = 0$ .

The demand constraint defines the maximum number of channels to be assigned to each cell, that is, the total

assigned channels for each cell  $i$  should not exceed its demand value  $d_i$ . That is,  $F_{i1} + F_{i2} + F_{i3} + \dots + F_{im} - d_i = 0$ .

For any two channels  $q$  and  $r$ , if  $c_{ii} > |q - r|$ , then channel  $q$  should not be assigned to  $i^{\text{th}}$  cell. Also For any two channels  $q$  and  $r$ , if channel  $q$  is assigned to  $j^{\text{th}}$  cell and if  $c_{ij} > 0$  and  $i \neq j$  with  $c_{ii} > |q - r|$  then  $i^{\text{th}}$  cell violates the assignment of the  $q^{\text{th}}$  channel.

The objective function of CAP is finding  $F$  while minimizing the overall cost function,  $C(F)$  while fulfilling the constraints of the problem.

### III. THE PROPOSED GENETICAL ALGORITHM

This section presents an evolutionary method with clique partitioning in solving graph coloring and channel allocation problems and its flowchart is presented in Figure 1. Cliques are obtained and are assigned the proper channels in minimizing the network interference. The strategy splits  $G$  into several maximum cliques. For the interfering cliques, different channels are assigned to them. The overall skeleton to find the solution to CAP is given below:

- i. Apply partition of  $V(G)$  to obtain the different cliques, say  $c_1, c_2, c_3, \dots, c_q$ .
- ii. Apply the following for each of the cliques  $i=1$  to  $q$ :
  - a. Find the feasible solution for vertex partitions and cliques.
  - b. Arbitrarily check for the interference of any two cliques and assign the valid channel to the cells based on interference constraints.

Clique partition constructs a super graph  $G': (S, E')$  which is obtained from  $G: (V, E)$ . Every vertex  $s_i$  in  $S$  is a super vertex that consists of a set of finite vertices  $v_i$  in  $V$ . A vertex  $s_i$  in  $S$  is a common neighbor of  $s_j$  and  $s_k$  in  $S$  if there exist edges  $(i, j)$  and  $(i, k)$  in  $E'$ . The following procedure finds the maximum clique of  $G$ :

- a. Put each  $v_i$  in  $V(G)$  in a separate vertex  $s_i$  in  $S$  of  $G'$ .
- b. Identify  $s_{i1}$  and  $s_{i2}$  in  $S$  such that  $(s_{i1}, s_{i2})$  in  $E'(G)$  with the greatest number of common adjacent vertices.
- c. Join the identified vertices into a single vertex  $s_{i1i2}$  which have  $s_{i1}$  and  $s_{i2}$  in all.
- d. Determine the general set which consists of the entire general neighbors of  $s_{i1}$  and  $s_{i2}$ .
- e. Delete all edges which are emerging from  $s_{i1}$  or  $s_{i2}$  in  $G'$ .
- f. Include the new  $e_i$ 's from  $s_{i1i2}$  to every super vertex in the general set.
- g. Iteratively execute the procedure defined in (a) to (f) until  $G$  has no edges.
- h. Vertex set  $s_i$  in  $S$  defines the clique.

Flowchart to solve GCP & CAP using the proposed method is shown in Figure 1. The new genetic operators are represented below.

#### CSMPCGXoperation:

Apply steps (a) to (e) of SPCGX crossover [11].

Apply crossover at single, multiple points and find the better offspring.

Repeat the combined single and multipoint crossover for  $t$  times.

Obtain the offspring:  $i' = (i_1', i_2', i_3', \dots, i_n')$  &  $j' = (j_1', j_2', j_3', \dots, j_n')$ .

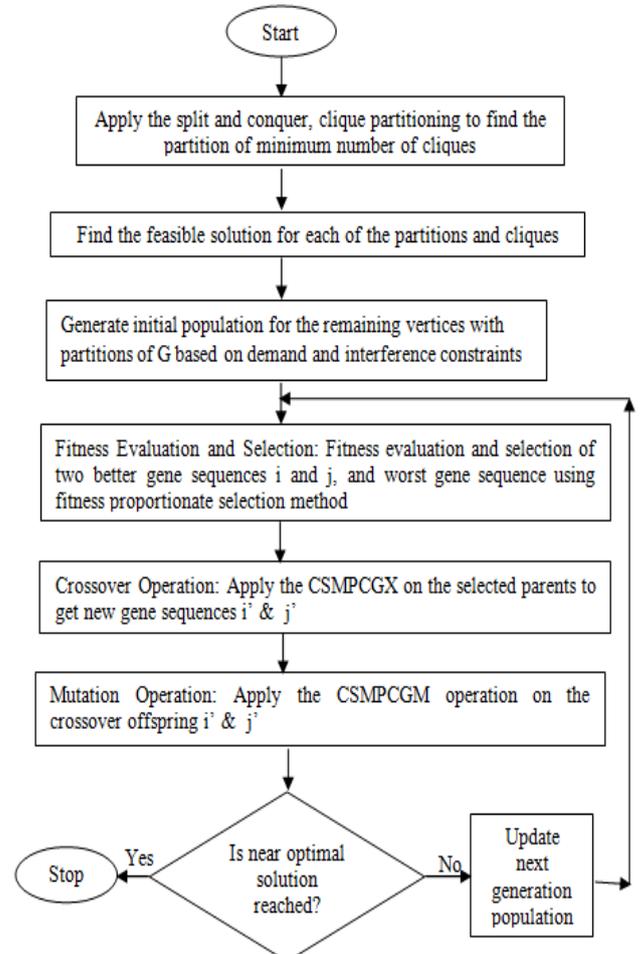


Fig. 1: Solving GCP & CAP using the proposed method - Flowchart

#### CSMPCGMoperation:

Apply the steps (a) to (d) of the mutation operator [11]. Apply mutation at single, multiple points and find the better offspring.

Repeat the combined single and multipoint crossover for  $t$  times.

Generate better offspring:  $i''$  and  $j''$ .

### IV. ANALYSIS AND GLOBAL CONVERGENCE

This section focus on the analysis and global convergence of the proposed genetic operators. The devised operators achieve quick stochastic convergence in finding the better near optimal solution. The stochastic convergence of the proposed operators are analyzed as follows:

The new genetic operators with clique partition strategy monotonically reduce the fitness values of gene sequences

$i$  and  $j$ ,  $f_g(i)$  &  $f_g(j)$  and the following analysis is observed during the generations:

Case (a): Either  $f_g(i)$  or  $f_g(j)$  decreases monotonically and converges. In this case,

$$f_0(i) \geq f_1(i) \geq f_2(i) \geq \dots \geq f_q(i) \geq f_{q+1}(i) = 0 \text{ or } f_0(j) \geq f_1(j) \geq f_2(j) \geq \dots \geq f_q(j) \geq f_{q+1}(j) = 0.$$

Case (b): At the initial stage, if  $f_g(i)$  or  $f_g(j)$  satisfies  $f_0(i) \geq f_1(i) \geq f_2(i) \geq \dots \geq f_q(i)$  and  $f_0(j) \geq f_1(j) \geq f_2(j) \geq \dots \geq f_q(j)$ , then after finite and smaller iterations, the following holds to converge.

$$f_q(i) < f_{q+1}(i) < \dots < f_t(i) \text{ and } f_q(j) < f_{q+1}(j) < \dots < f_t(j)$$

The global convergence of the devised method to achieve better near optimal solution while reducing the search space is proved in the following theorem.

**Theorem 1: The proposed genetic algorithm with CSMPCGX and CSMPCGM always converges stochastically.**

Proof: Let  $S$  represents the finite search space. The following conditions should be fulfilled for stochastic convergence [2]. For arbitrarily chosen individuals  $i, j \in S$ , individuals should be reachable and the population  $P_g$  satisfy the monotone property.

$p_c$  and  $p_m$  define the CSMPCGX & CSMPCFM probabilities for applying crossover and mutation respectively.

Then  $p(i'') \geq p_c p\{i' = \text{crossover}(i)\} p_m p\{i'' = \text{mutation}(i')\}$

CSMPCGX operation implies  $p\{i' = \text{crossover}(i)\} = 1 / c^k > 0$  and

CSMPCGM results in  $p\{i'' = \text{mutation}(i')\} = 1 / f(G)^l > 0$ .

Thus  $p(i'') \geq (p_c p_m) / (c^k f(G)^l)$  is in  $[0, 1]$ .

The elitism operation replaces the worst individual with a better one and  $P_g$  gets updated for the monotone property.

## V. RESULTS OF BENCH MARK INSTANCES

The maximum number of cells considered is 25 and the maximum number of channels considered for the simulation of the proposed method is 91. The devised operators are tested in solving some standard benchmark graphs and CAPs using Intel Core i5-2450M 2.5GHz system using Java. The graph parameters  $n$  and  $m$  are evaluating the output performance measurements. The outcomes are analyzed with the following inferences.

1. Stochastic convergence is reached even to the high-density graphs, such as miles1500.col.
2. Expected generations are reduced compared to other genetic methods and the obtained values are shown in figures, Fig. 2 to Fig. 4.
3. The solution is obtained with minimal complexity for the queen, register allocation, Miles graphs.
4. When the number of cells increase, the frequency of convergence will be affected.
5. The frequency of convergence depends on the number of channels. When the channels count decreases, near

optimal solution with less frequency of convergence is attained.

When comparing the results of this method with some of the existing methods [9-15], this new method offsets the problems such as improving the near-optimal solution, minimize the problem search space, and achieve fast stochastic convergence with smaller population size  $N$ . The major inferences to target these justifications are:

1. The solution is obtained for different benchmark random graphs.
2. Stochastic convergence is achieved for small  $N$  compared to other methods [9-21].
3. The solution is obtained for queen, Mycielski graphs compared to the near-optimal performance of existing methods.
4. Provides better solution than DSATUR, FINOCCHI, and FROGSIM methods.

Table 1 represents the different CAP instances and the expected generations obtained are given in Figure 5. The convergence of near-optimal solutions for these problems is shown in Table II and these outcomes are matched with some existing methods [4-7] and are indicated in Table III. This new method significantly reduces the complexity in getting the solution with a minimum fitness threshold and higher frequency of convergence (FOC).

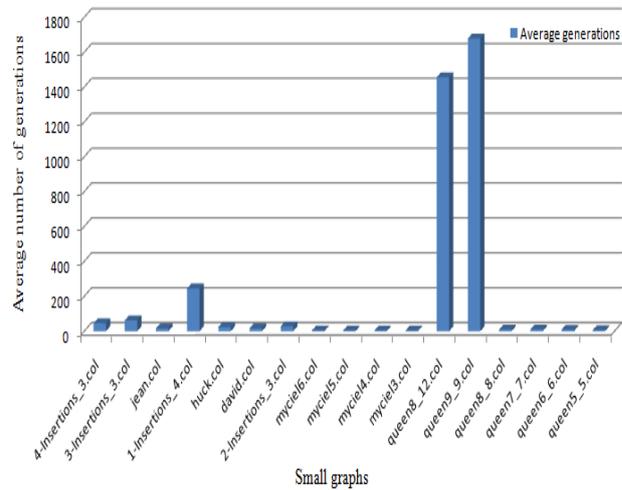


Fig. 2 Expected generations obtained for small graphs

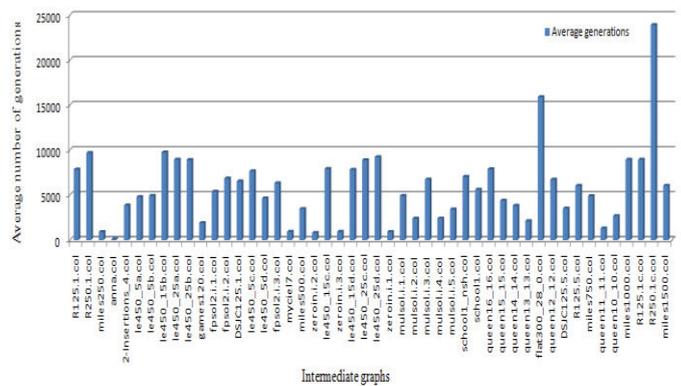


Fig. 3. Expected generations obtained for intermediate graphs

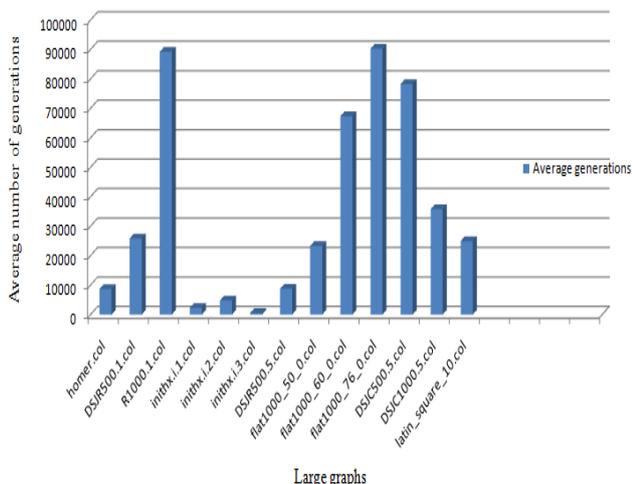


Fig. 4 Expected generations obtained for large graphs

Table I C(F) obtained for various benchmark instances

Problem	Number of Cells	Number of Channels	C(F)
EX1	4	11	0
Ex2	5	17	0
HEX1	21	37	0
HEX2	21	91	0
HEX3	21	21	0
HEX4	21	56	0
KUNZ1	10	30	0
KUNZ2	15	44	0
KUNZ3	20	60	0
KUNZ4	25	73	0

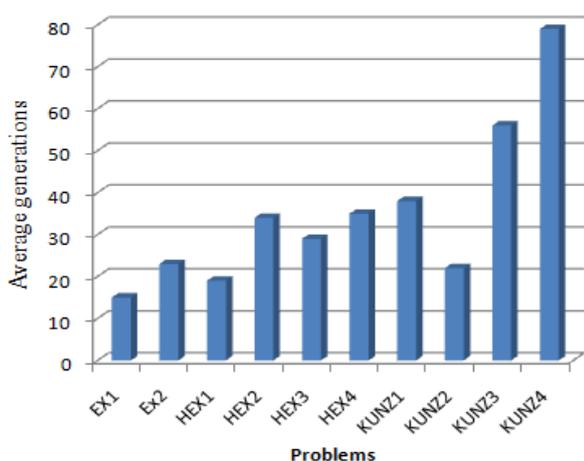


Fig. 5 Expected generations obtained for CAP instances

Table II Convergence of F for benchmark instances

Problem	Cells n	Channels m	Simulation Run Length	Fitness Threshold	FOC
1	4	11	100	0	100%
2	25	73	20000	0	100%
3	21	385	20000	0	100%
4	500	1250	40000	0.83	100%
5	1000	1500	45000	0.89	100%
6	2000	1750	50000	3.45	99.5%
7	5000	2000	50000	2.48	98.6%
8	10000	2500	50000	6.46	97.9%
9	20000	5000	50000	7.92	97.5%
10	50000	7500	50000	9.95	98.5%

Table III Comparison with Existing Methods

Problem	Existing Methods	Proposed Method		
	FOC	FOC	No. of Trials	Time(sec)
1	100%	100%	1	0
2	92%	100%	6	36
3	80%	100%	1	0
4	-	100%	23	762
5	-	100%	545	864
6	-	99.5%	453	342
7	-	98.6%	564	654
8	-	97.9%	233	544
9	-	97.5%	348	889
10	-	98.5%	655	903

VI. CONCLUSION

A new genetic method is designed for solving GCP & CAP using CSMPCGX & CSMPCGM operators. The experiments are conducted on difficult benchmark instances and showed that this method outperforms the various existing methods with the reduction in the exploration of search space and minimizing the expected generations. This evolutionary strategy reduces the interference of CAP with minimal N and hence better performance is obtained compared to the well-known methods. Stochastic convergence of higher frequency of convergence is attained for the standard and random CAPs of large sizes.

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