

An Experimental Analysis of Explorative and Exploited Operators of Genetic Algorithm for Operating system Process Scheduling Problem.

Rajiv Kumar¹

^{*1}Computer Science & Engg. PhD. Scholar
Singhania University Jhunjhunu, Rajasthan, INDIA
rajiv_kumar_gill1@yahoo.co.in

Abstract— Many Research have be done for Process scheduling Problem. Most of the time parameters of the Scheduling problem have changed but the problem remain same as to find out the optimal schedule which can optimize the problem under consideration. Implementation of genetic algorithm for operating system process scheduling is a new idea . Genetic Algorithm is a robust technique for solve process scheduling and optimization problem. The performance of any genetic algorithm is depend on the operators used for the GA Simulation. In this paper we will analyse the performance of modified cross over genetic algorithm for operating system process scheduling problem. We take two operator of GA i.e. explorative operator and exploited operator. The performance of the genetic algorithm is greatly depends upon these two operator. As the probability of explorative and exploited operator is changed the convergence of Genetic algorithm is also changed.

Keywords: Genetic algorithm , NP-hard , Operating system, Exploited, Explorative, Scheduling.

1.Introduction

Process Scheduling is concern with allocation of hardware to the ready process. The scheduling problem is consider as NP-hard. There are many techniques which have been developed to handle the scheduling problem. But no is the best. The implementation of genetic algorithm for operating system is a new idea[1]. Over many years, most of the researchers thought that optimal scheduling is very difficult task to achieve. With the growing interest in Genetic Algorithms (GAs) and the GA is another promising global optimization technique[3] be used for operating process scheduling problem. The modified cross over genetic algorithm is first proposed by Davis[2] and apply this algorithm to operating system process scheduling problem by rajiv et al[3]. Genetic algorithms(Gas) are adaptive methods which may be used to solve search and optimization problems. Genetic algorithms (GAs) were first proposed by the John Holland[4] in the 1960s

This Genetic algorithm have been applied to many scientific and engineering problems[5][6][7]. The performance of the genetic algorithm is limited by some problem, typically premature convergence. This happens simply because of the accumulation of stochastic errors. If by

chance, a gene becomes predominant in the population, then it just as likely to become more predominant in the next generation as it is to become less he predominant. If an increase in predominance is sustained over several successive generations and population is finite, then a gene can be spread to all members of the population. Once gene has converged in this way, it is fixed then crossover cannot introduce new gene values. The crossover operator and mutation operator is consider the basic operator of GA . The cross over operator is consider as explorative operator and mutation operator is consider as exploited operator. The cross over operator diversify the population and mutation operator exploit the new result . when the population is premature converge then mutation exploit the population and reduce the premature convergence situation and hence the searching power of genetic algorithm.

1.2 The Paper Description

The major part of this paper, contained in section 2, will explain working of genetic algorithm and their application in process scheduling problem. The GA is robust techniques and it has no. of operators which have their own properties .The parameter setting in the genetic algorithm is concerned with the setting of applicable static values of the operators used. Ie crossover probability , inversion rate , population size etc. Accessible introduction can be found in the books by Davis [8] and Goldberg[9] .Section 3 describe the proposed structure of genetic algorithm. Section 4 explain the experimental setup for analysis and Section 5 shows the experimental results of the problem under consideration and section 6 is conclusion .

2. Introduction of Genetic Algorithm

2.1 overview

The evaluation function, or objective function, provides a measure of performance with respect to a particular set of parameters. The fitness function transforms that measure of performance into an allocation of reproductive opportunities. The evaluation of a string representing a set of parameters is independent of the evaluation of any other string. The fitness of that string, however, is always defined with respect to other members of the current population. In the genetic algorithm,

fitness is defined by: f_i / f_A where f_i is the evaluation associated with string i and f_A is the average evaluation of all the strings in the population. Fitness can also be assigned based on a string's rank in the population or by sampling methods, such as tournament selection. The execution of the genetic algorithm is a two-stage process. It starts with the current population. Selection is applied to the current population to create an intermediate population. Then recombination and mutation are applied to the intermediate population to create the next population. The process of going from the current population to the next population constitutes one generation in the execution of a genetic algorithm. In the first generation the current population is also the initial population. After calculating f_i / f_A for all the strings in the current population, selection is carried out. The probability that strings in the current population are copied (i.e. duplicated) and placed in the intermediate generation is in proportion to their fitness.

2.2 Coding

Before a GA can be run, a suitable coding (or representation) for the problem must be devised. We also require a fitness function, which assigns a figure of merit to each coded solution. During the run, parents must be selected for reproduction, and recombined to generate offspring. It is assumed that a potential solution to a problem may be represented as a set of parameters (for example, the parameters that optimize a neural network). These parameters (known as genes) are joined together to form a string of values (often referred to as a chromosome. For example, if our problem is to maximize a function of three variables, $F(x; y; z)$, we might represent each variable by a 10-bit binary number (suitably scaled). Our chromosome would therefore contain three genes, and consist of 30 binary digits. The set of parameters represented by a particular chromosome is referred to as a genotype. The genotype contains the information required to construct an organism which is referred to as the phenotype. For example, in a bridge design task, the set of parameters specifying a particular design is the genotype, while the finished construction is the phenotype.

The fitness of an individual depends on the performance of the phenotype. This can be inferred from the genotype, i.e. it can be computed from the chromosome, using the fitness function. Assuming the interaction between parameters is nonlinear the size of the search space is related to the number of bits used in the problem encoding. For a bit string encoding of length L ; the size of the search space is 2^L and forms a hypercube. The genetic algorithm samples the corners of this L -dimensional hypercube. Generally, most test functions are at least 30 bits in length; anything much smaller represents a space which can be enumerated. Obviously, the expression 2^L grows exponentially. As long as the number of "good solutions" to a problem are sparse with respect to the size of the search space, then random search or search by enumeration of a large search space is not a practical form of problem solving. On the other hand, any search other than random search imposes some bias in terms of how it looks for better solutions and where it looks in the search space. A genetic algorithm belongs to the class of methods known as

"weak methods" because it makes relatively few assumptions about the problem that is being solved. Genetic algorithms are often described as a global search method that does not use gradient information. Thus, no differentiable functions as well as functions with multiple local optima represent classes of problems to which genetic algorithms might be applied. Genetic algorithms, as a weak method, are robust but very general.

2.3 Fitness Evolution

A fitness evolution function must be devised for each problem to be solved. Given a particular chromosome, the fitness function returns a single numerical "fitness," or "figure of merit," which is supposed to be proportional to the "utility" or "ability" of the individual which that chromosome represents. For many problems, particularly function optimization, the fitness function should simply measure the value of the function.

2.4 Selection

Determine which strings are "copied" or "selected" for the mating pool and how many times a string will be "selected" for the mating pool. Higher performers will be copied more often than lower performers. Example: the probability of selecting a string with a fitness value of f is f/f_t , where f_t is the sum of all of the fitness values in the population.

Individuals are chosen using "stochastic sampling with replacement" to fill the intermediate population. A selection process that will more closely match the expected fitness values is "remainder stochastic sampling." For each string i where f_i/f_t is greater than 1.0, the integer portion of this number indicates how many copies of that string are directly placed in the intermediate population. All strings (including those with f_i/f_t less than 1.0) then place additional copies in the intermediate population with a probability corresponding to the fractional portion of f_i/f_t . For example, a string with $f_i/f_t = 1:36$ places 1 copy in the intermediate population, and then receives a 0:36 chance of placing a second copy. A string

with a fitness of $f_i/f_t = 0:54$ has a 0:54 chance of placing one string in the intermediate population. Remainder stochastic sampling is most efficiently implemented using a method known as stochastic universal sampling. Assume that the population is laid out in random order as in a pie graph, where each individual is assigned space on the pie graph in proportion to fitness. An outer roulette wheel is placed around the pie with N equally-spaced pointers. A single spin of the roulette wheel will now simultaneously pick all N members of the intermediate population.

2.5 Reproduction

After selection has been carried out the construction of the intermediate population is complete and recombination can occur. This can be viewed as creating the next population from the intermediate population. Crossover is applied to randomly paired strings with a probability denoted P_c . (The population

should already be sufficiently shuffled by the random selection process.) Pick a pair of strings. With probability p_c "recombine" these strings to form two new strings that are inserted into the next population. In the proposed algorithm we use the modified crossover operator.

Good individuals will probably be selected several times in a generation, poor ones may not be at all. Having selected two parents, their chromosomes are recombined, typically using the mechanisms of crossover and mutation. The previous crossover example is known as single point crossover. Crossover is not usually applied to all pairs of individuals selected for mating. A random choice is made, where the likelihood of crossover being applied is typically between 0.6 and 1.0. If crossover is not applied, offspring are produced simply by duplicating the parents. This gives each individual a chance of passing on its genes without the disruption of crossover.

Mutation is applied to each child individually after crossover. It randomly alters each gene with a small probability. The next diagram shows the fifth gene of a chromosome being mutated: The traditional view is that crossover is the more important of the two techniques for rapidly exploring a search space. Mutation provides a small amount of random search, and helps ensure that no point in the search has a zero probability of being examined.

2.6 Convergence

The convergence of the genetic algorithm is concern with the uniformity in the population of solution. when 95 % of the population has the same result then we can say that the gene is converge. As the population converges, the average fitness will approach that of the best individual. A GA will always be subject to stochastic errors. One such problem is that of genetic drift. Even in the absence of any selection pressure (i.e. a constant fitness function), members of the population will still converge to some point in the solution space.

This happens simply because of the accumulation of stochastic errors. If, by chance, a gene becomes predominant in the population, then it is just as likely to become more predominant in the next generation as it is to become less predominant. If an increase in predominance is sustained over several successive generations, and the population is finite, then a gene can spread to all members of the population. Once a gene has converged in this way, it is fixed; crossover cannot introduce new gene values. This produces a ratchet effect, so that as generations go by, each gene eventually becomes fixed. The rate of genetic drift therefore provides a lower bound on the rate at which a GA can converge towards the correct solution. That is, if the GA is to exploit gradient information in the fitness function, the fitness function must provide a slope sufficiently large to counteract any genetic drift. The rate of genetic drift can be reduced by increasing the mutation rate. However, if the mutation rate is too high, the search becomes effectively random, so once again gradient information in the fitness function is not exploited.

3. Structure of Proposed GA-Based Algorithm

Algorithm GA (Modified crossover GA)

- (1) Begin
- (2) Initialize Population (randomly generated);
- (3) Fitness Evaluation;
- (4) Repeat
- (5) Selection(Roullete wheel Selection) ;
- (6) Modified crossover;
- (7) Inversion();
- (8) Fitness Evaluation;
- (9) Elitism replacement with Filtration;
- (10) Until the end condition is satisfied;
- (11) Return the fittest solution found;
- (12) End

4. Experimental Setup

The Individual are randomly generated to form an initial population. Successive generations of reproduction and crossover produce increasing numbers of individuals . Modified crossover operator with crossover probability P_c is 0.6 is taken and then change the inversion probability with respect to crossover probability. The experieient is perform with two crossover probability (p_c) i.e.0.6 and 1.0 Crossover operator exploited the population or you can say that it can diversify the population. But due to the genetic drift some time the population is converge to the local optimal point, At that time crossover operation can not diversify the population. The inversion operator is explorative in nature ,it diversify the population ,but in general the probability of inversion is very low . so in our simulation We first have 0.1 inversion probability then we proceed with .01,.001.

5. Simulation Results

In this experiment we have compare modified crossover GA with two crossover probability P_c ie 0.6 and 1.0 also with variable inversion probability .we find that the performance and convergence state of the GA is greatly effected .

Parameter / Strategy	Setting
Population Size	30
Population Type	Generational
Initialization	Random
Selection	Roulette wheel
Crossover	Two Parents, Modified crossover
Crossover Probability	0.6 and 1.0
Variable Inversion Probability	0.1,0.01,0.001
Replacement strategy	Keep 80 % Best
Stopping Strategy	85 % Population converge
No. of process to be Schedule	5
Fitness criterion	Minimum Weighted Turn Around Time

Table 1. Parameters of GA

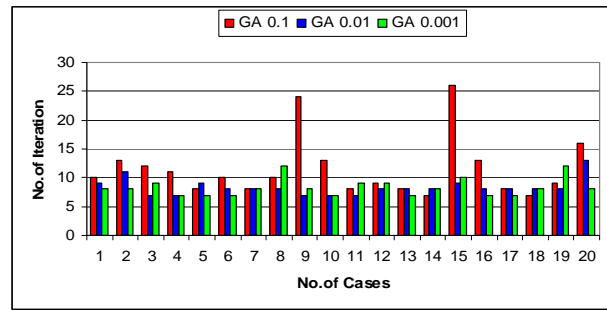


Figure 1. Comparison of Pi (0.1,0.01,0.001) with Cp=0.6, between No. of case and No. of Iteration

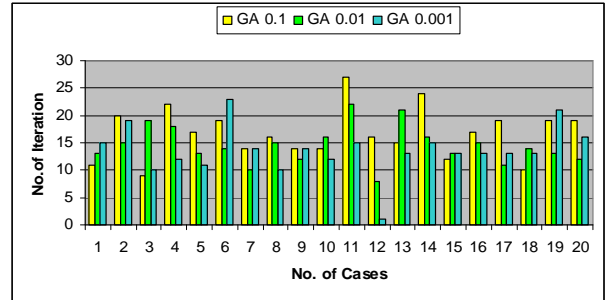


Figure 2. Comparison of Pi (0.1,0.01,0.001) with Cp=1.0, between No. of case and No. of Iteration

Sr.No.	Job Service Time in Second					C _{P=0.6}			C _{P=1.0}		
	J1	J2	J3	J4	J5	I _{Pi=0.1}	I _{Pi=0.01}	I _{Pi=0.001}	I _{Pi=0.1}	I _{Pi=0.01}	I _{Pi=0.001}
1	17	21	5	10	24	11	13	15	10	9	8
2	12	24	4	16	19	20	15	19	13	11	8
3	11	27	3	23	18	9	19	10	12	7	9
4	14	19	21	2	7	22	18	12	11	7	7
5	22	11	10	19	1	17	13	11	8	9	7
6	12	14	17	27	13	19	14	23	10	8	7
7	15	7	24	28	30	14	10	14	8	8	8
8	17	11	17	26	10	16	15	10	10	8	12
9	24	15	16	23	18	14	12	14	24	7	8
10	10	9	27	21	19	14	16	12	13	7	7
11	25	16	15	23	12	27	22	15	8	7	9
12	20	17	11	14	21	16	8	1	9	8	9
13	9	29	22	10	14	15	21	13	8	8	7
14	21	15	24	16	20	24	16	15	7	8	8
15	14	24	13	28	17	12	13	13	26	9	10
16	26	30	29	19	20	17	15	13	13	8	7
17	23	17	20	16	27	19	11	13	8	8	7
18	23	22	17	28	21	10	14	13	7	8	8
19	14	20	9	10	20	19	13	21	9	8	12
20	18	27	5	29	23	19	12	16	16	13	8
Total No. of Iteration = $\sum_{s=1}^{s=20} I_s$						334	290	273	230	166	166
Mean Iteration = $\frac{\sum_{s=1}^{s=20} I_s}{20}$						16.7	14.5	13.65	11.5	8.3	8.3
Cp = Crossover Probability=0.6,1.0 Pi = Probability of Inversion =0.1,0.01,0.001											

Table 2. Comparison Results

In this paper, we examine the effect of varying inversion probability with respect to variable crossover probability. The combine variation of the probability of the two operator will have enhance the performance of GA. We have found that when the probability of inversion is decrease with constant crossover probability ie $cp=0.6$ then the no of iteration is also decrease and we get good result by decreasing the inversion probability. The second thing that examine is that as the probability of cross over increase i.e. from 0.6 to 1.0 and inversion probability decrease from 0.1 to 0.001. the performance of the genetic algorithm is enhanced . so it is clear that decreasing inversion probability and increasing crossover probability increase the performance of genetic algorithm under operating system process scheduling problem.

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