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Clustered Genetic Algorithm to solve Multidimensional Knapsack Problem

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ABSTRACT

Genetic Algorithm (GA) has emerged as a powerful tool to discover optimal for multidimensional (MDKP). Multidimensional knapsack problem knapsack problem has recognized as NP-hard problem whose applications in many areas like project selection, capital budgeting, loading problems, cutting stock etc. Attempts has made to develop cluster genetic algorithm (CGA) by mean of modified selection and modified crossover operators of GA. Clustered genetic algorithm consist of (1) fuzzy roulette wheel selection for individual selection to form the mating pool (2) A different kind of crossover operator which employ hierarchical clustering method to form two clusters from individuals of mating pool. CGA performance has examined against GA with respect to 30 benchmark problems for multidimensional knapsack. Experimental results show that CGA has significant improvement over GA in relation to discover optimal and CPU running time. The data for **MDKP** is available set at http://people.brunel.ac.uk/ mastjjb/jeb/orlib/files/mknap2.txt

Keywords: Multidimensional knapsack problem, Genetic algo-rithm, Fitness function, Crossover, Mutation

1 INTRODUCTION

Multidimensional knapsack problem is one of the famous problem in applied mathematics belongs to NP- hard combi-natorial optimization [1]. This problem modeling has applied in several project selection and capital budgeting areas [2-4]. Meirier developed more mechanisms from real word practice to combine models of capital budgeting with novels, and practically applicable technique for evaluation of project [5]. They developed scenario dependent capital budgeting model that have MDKP as a subproblem associated with constraint of generalized

upper bound (GUB). Capital budgeting emerged as key challenge for not for profits multihospital healthcare units in United States. D.N. Kleinmuntz and C.E. Kleinmuntz developed a framework for capital budeting which used MDKP formulations [6]. MDKP have been employed to model prob-lems like investment policy in tourism area of developing country [7], database allocation and processor allocation in distributed computing environments [8], groceries delivery in vehicles with multiple compartments [9], cutting stock [10], loading problems [11] and approval voting [12]. Currently, MDKP has been applied in modeling of daily management of remote satellite like SPOT, which consisted in deciding each day what photographs will be attempted the next day [13].

Multidimensional knapsack problem is described by n objects and m knapsacks. Each knapsack has a capacity d_j (j = 1; 2; 3:::m). A number of binary variables x_i (i = 1; 2; 3::n) are used, that is set to 1 if ith object is chosen to inset in knapsacks otherwise it is set to 0. Each object has a profit value p_i (i = 1; 2; 3::n) and a weight w_{ij} corresponding to knapsacks. Thus mathematically this optimization problem can be modeled as :

$$\begin{array}{c} \underset{x_i}{\text{Maximize}} & \underset{=1}{\overset{n}{\text{pixi}}} & (1) \end{array}$$

Subject to :

$$\underset{=1}{\overset{M}{\underset{wijxi}{x}}} \underbrace{wijxi}_{j} \underbrace{j}_{j=1;2;...m} (2)$$

 $x_i 2 f0; 1g$ i = 1; 2; 3::n (3)

The goal of MDKP is to insert a subset of objects into knapsacks that obtain maximum profit without

capacity viola-tion of knapsacks. This characteristic reveal this problem has all nonnegative entries. More specifically, we can say $p_i > 0$,

$$d_j > 0, 0$$
 w_{jj} d_j and w_{jj} d_j for all $i 2 n$ and $j 2 m$

Rest of the paper is structured as follows. Genetic algorithm for MDKP is described in Section II. Section III give the overview of related work for MDKP in domain of genetic algorithm. Section IV explain CGA to solve multidimensional knapsack problem. Experimental environments and results are given in Section V. At last conclusion is made in section VI.

II. GENETIC ALGORITHM FOR MULTIDIMENSIONAL KNAPSACK PROBLEM

Genetic algorithm is computation optimization technique which mimic the evolutionary principle i.e. fittest individual will survive in each generation. GA is guided by three main operation: selection, crossover and mutation. Individuals re-production are governed by crossover and mutation operation. Selection operation choose fittest individuals from population to form mating pool. Algorithm run as follow : A set of random individuals are produced which serve as the initial population of GA. The individual can be expressed in real number, binary number, integer number, character based on nature of problem. MDKP uses the binary representation. Individual goodness measurement is estimated through fitness function. MDKP fitness function is maximize the objective function.

Algorithm 1 Genetic Algorithm (GA)

Initialize the control parameters ;

Initialize randomly N individuals as initial population P

(G); while stopping criterion(s) doesn't meet do

- i. Choose individuals to create mating pool via selection operator.
- ii.Produce new individuals via crossover operator.
- iii. Mutate obtained individuals using mutation operator.

iv. Evaluate fitness value of new individuals.

- v. Add new individuals with population.
- vi. Sort population with respect to fitness value.
- vii. Choose top N individuals as population of next generation.

end while

Return the fittest individual as the solution.

Every individual are transformed from one generation to next generation depending on fitness value i.e individual with high fitness value has high probability to participate in upcoming generation. GA employee crossover operator for exploration and mutation operator for exploitation. MDKP uses one point crossover and random mutation. Number of crossover operations are govern through crossover rate while number of mutation operations are govern through mutation rate. A cycle of GA comprise of applying GA operations i.e. evaluation, reproduction, crossover and mutation. Each generation produce a set of individuals. GA termination return the fittest individual as solution. The details of GA operators i.e. selection, crossover, and mutation are explained in [14].

III. RELATED WORK

J.P. Martin and C.B Neto and M.K. Crocomo made compar-ison among four linkage learning based genetic algorithm for multidimensional knapsack problem namely, extended com-pact genetic algorithm (eCGA), bayesian optimization algo-rithm (BOA) with detection graphs, BOA with community detection and linkage tree genetic algorithm (LTGA) [15]. eCGA and LTGA have better exploration and exploitation capabilities than BOA versions with small size population (N=100) and produce better quality solutions for MDKPs. All algorithms with large population size produce approxi-mately similar quality solutions for MDKPs but these have different running time and function evaluations. When consider solutions quality, functions evaluations and running time of algorithms, BOA versions have better running time than eCGA and LTGA. It is very difficult to make comparison for computational time between eCGA and LTGA because these have different reproduction operators.

Khuri, Back and Heitkotter proposed a GA for 0/1 MDKP [16]. Algorithm allowed breeding and participation of infeasi-ble solutions during search. In this GA, reproduction proceeds by combining the partial information from all element of population. A test suite with few problems were used to test this GA; only moderate quality solutions were produced. Hoff, Lokentangen and Mittet proposed a GA with proper tuned parameters and search mechanisms in which only feasible solutions were allowed during search [17]. This GA discovered optimal for 54 problems out a suite of 55 problems.

Rudolph and Sprave proposed a GA in which selection of parent is not unrestricted as in standard GA but is restricted be-tween neighboring solutions [18]. In their GA every infeasible solutions were penalized as GA of Khursi, Back and Heitkotter [16]. The premier component of this GA be solutions local interaction with in a spatial structured population and self ad-justed controlling mechanism of selection pressure. P. C. Chu and J.E. Beasley proposed heuristic dependent GA for MDKP [19]. In their GA heuristic operator was employed which work with problem specific knowledge. Heuristic operator repaired infeasible solutions and algorithm discover optimal solution among feasible solutions. C. Cota and J.M. Troya proposed another GA which have an improvement mechanism whose objective is to convert infeasible solutions in search space to feasible solutions [20]. Experimental results were better to basic GA, but not as good as Chu and Beasley's GA results.

IV. CLUSTERED GENETIC ALGORITHM FOR^{two clusters.}

MULTIDMENSIONAL KNAPSACK PROBLEM

In this work, clustered genetic algorithm is developed for multidimensional knapsack problem. CGA is explained in algorithm 2. CGA uses an advanced selection method which employee fuzzy and roulette wheel selection called fuzzy roulette wheel selection. Selection mechanism always choose first individual and assigned it to first partition of roulette wheel. Partition size is relative to fitness value. Next similarity value between first individual and remaining individuals are estimated. The individual with highest similarity value occupy the second partition of roulette wheel. This procedure repeat until all individuals are allocated to partition in roulette wheel. Figure 1 show the 4 individuals in binary representation and procedure of assigning individual to the partition of roulette wheel. Then mechanism determined overlapping area between adjacent individuals. Overlapping area size is correlated by similarity value between adjacent individuals. In mating pool formation, a random number is generated. After for each individual membership degree to which generated number belongs is calculated. Individual with highest membership degree is selected. This process repeat until mating pool size is equal to population size. Figure 2 show the fuzzy roulette wheel selection mechanism for 4 individuals in Figure 1.



Fig. 1: Procedure of assigning individual to partition of roulette wheel .

CGA adopt a special crossover operator which uses the hierarchical algorithm to form clusters. Individuals in the mating pool are divided into two clusters using hierarchical clustering algorithm. In the hierarchical algorithm, single linkage procedure given in eq (2) is apply to estimate similarity between the two clusters.



Fig. 2: Fuzzy roulette wheel.

$$S(X; Y) = M AX fs(x; y)g$$
(4)

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Where X and Y are two clusters and s(x,y) stand for similarity between individuals x and y. A pair of offsprings are generated by mating of a random parent from first cluster and a random parent from second cluster. One point crossover operator is employed. This step is continued N/2-1 times. Random mutation is performed for each offspring to keep diversity and to defeat the problem of premature convergence.



Fig. 3: Selection of parent pair for crossover.





V. EXPERIMENTAL RESULTS AND DISCUSSION

To show CGA is better technique to GA, experiments has carried out over 30 standard problems for multidimensional knapsack.

Algorithm 2 Clustered Genetic Algorithm (CGA)

Initialize the control parameters ;

Initialize randomly N individuals as initial population P (G); while stopping criterion(s) doesn't meet do

- i.Choose individuals to create mating pool via fuzzy roulette wheel selection operator.
- ii. Create two clusters of individuals using hierarchical clustering algorithm .
- iii. (iii)Produce new individuals by using crossover operator with a random parent from cluster 1 and a random parent from cluster 2.
- iv. Mutate obtained individuals using mutation operator.

v. Evaluate fitness value of new individuals.

- vi. Add new individuals with population.
- vii. Sort population with respect to fitness value.
- viii. Choose top N individuals as population of next generation.

end while

Return the fittest individual as the solution.

A. Experimental Environment

GA and CGA depict in previous section were implementd for multidimensional Knapsack. Experiments were de-signed to estimate average mean error and average mean execution time for GA and CGA. Data set for multidimen-sional knapsack is available at http://people.brunel.ac.uk/ mastjjb/jeb/orlib/files/mknap2.txt. Crossover rate CR = 0.7 and mutation rate MR = 0.01. The experiments were conducted on 7th generation intel core i-7-7700k processor (4.2 GHz) and coded in c++ whose compilation is done using Dev c++ compiler.



Fig. 5: Error comparision between GA and CGA.

B. Results and Analysis

Experimental results of GA and CGA for MDKPs are given in Table I and Table II respectively. Column 1 denote instance no and column 2 represent MDKP, column 3 and 4 show number of knapsacks and number of objects corresponding to problem, columns 5 denotes the optimal value, columns 6-10 keeps the output value of five run, simulation statical information is given in columns 11-14 and column 15 show mean time taken in execution. GA and CGA has average mean error 4.49% and 1.764% respectively while GA and CGA has average mean execution time 2.34 seconds and 2.34 seconds respectively. CGA has considerable less average mean error in comparison of GA but CGA has slightly higher average mean execution time in comparison of GA. Figure 5 plot the average mean error against the problem instance and Figure 6 plot the average mean execution time against the problem instance.



Fig. 6: CPU running time comparison between GA and CGA.

VI. CONCLUSION AND FUTURE WORK

This paper addressed multidimensional knapsack problem via the cluster genetic algorithm to find optimal. CGA employ the fuzzy roulette selection mechanism for individual selection to form mating pool. CGA adopts a special kind of crossover operator in which uses hierarchical clustering algorithm to form two clusters from mating pool. A pair of offspring is generated by mating of a random parent from first cluster and a random parent from second cluster. CGA performance has examined against GA w.r.t 30 standard MDKPs over two criterion i.e average mean error and average mean execution time. As GA and CGA experimental results are shown, CGA has better performance than GA for MDKP. A comparative study of different mutation operators with CGA will be studied in the future work.

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Sr no	Test Proble ms	m	n	Opt ima l			Numb Ru				Statis Inforn			Time (s)
					1	2	3	4	5	Max	Min	Avg	% Err	
1	SENT01	30	60	7772	7712	7736	7725	7719	7723	7736	7712	7723	0.6	4.27
2	SENT02	30	60	8722	8690	8662	8702	8669	8682	8702	8662	8681	0.4 6	4.23
3	WEING 1	2	28	1412 78	1379 02	1388 04	1386 02	1399 03	13990 4	13990 4	13790 2	1390 23	1.6 0	1.58
4	WEING 2	2	28	1308 83	1244 53	1257 52	1249 73	1239 89	12498 3	12575 2	12398 9	1248 30	4.6 2	1.60
5	WEING 3	2	28	95667	92296	9199 8	9229 2	92398	92451	92451	9199 8	92287	3.5 3	1.68
6	WEING 4	2	28	1193 37	1166 25	1164 41	1159 98	1164 59	11679 2	11679 2	11599 8	1164 63	2.4 1	1.71
7	WEING 5	2	28	98796	93667	9366 2	9344 2	93778	93631	93778	9344 2	93636	5.2 2	1.53
8	WEING 6	2	28	1306 23	1244 50	1260 93	1260 92	1261 72	12615 8	12617 2	12445 0	1257 93	3.7 0	1.59

	Internatio	nai J	ourm				/ Itesea	on and	Develop			JUIN 27	50 01	10
9	WEING	2	10	1095	1054	1057	1059	1057	1056	10598	1054	1057	3.4	5.72
	7		5	445	402	806	803	804	405	03	402	244	9	
10	WEING	2	10	6243	6214	6217	6206	6218	62176	62188	62064	6215	0.4	5.64
10	8		5	19	22	98	42	82	1	2	2	01	5	••••
11	WEISH	5	30	4554	4214	4298	4288	4178	4292	4298	4178	4254	6.5	1.62
11	01		50	1551	1211	1290	1200	11/0		1290	1170	1231	9	1.02
12	WEISH	5	30	4536	4197	4182	4178	4181	4187	4197	4178	4187	7.7	1.58
14	02	5	50	7550	1)/	7102	170	4101	4107	7177	4170	4107	4	1.50
13	WEISH	5	30	4115	3990	3987	3976	3973	3984	3990	3973	3982	3.2	1.64
15	03	5	50	4115	3990	3907	3970	3973	3904	3990	3973	3962	3.2	1.04
14		5	20	45(1	4214	4217	4222	4202	4204	4222	4202	4212		1.((
14	WEISH	5	30	4561	4314	4317	4323	4302	4304	4323	4302	4312	5.4	1.66
	04	-	•			10.00	1000	10.00	40.00	1000	12.0.0	12.0.1	5	1.(0
15	WEISH	5	30	4514	4311	4309	4388	4303	4309	4388	4303	4324	4.2	1.63
	05												1	
16	WEISH	5	40	5557	5331	5333	5307	5228	5291	5333	5228	5298	4.6	2.08
	06												6	
17	WEISH	5	40	5567	5186	5193	5177	5244	5180	5244	5177	5196	6.6	2 <mark>.1</mark> 2
	07												6	
18	WEISH	5	40	5605	5302	5368	5352	5214	5309	5368	5214	5329	4.9	2.06
	08												2	-
19	WEISH	5	40	5246	4901	4848	4898	4886	4812	4901	4812	4869	7.1	2.31
	09												9	
20	WEISH	5	50	6339	5916	5923	5978	5981	5972	5981	5916	5954	5.9	2.28
	10	_											9	
21	WEISH	5	50	5643	5387	5362	5324	5336	5361	5387	5324	5354	5.1	2.70
	11			0010	0007	0002	0021	0000	0001	2201	0021	0001	2	2.70
22	WEISH	5	50	6339	5984	5972	5988	5977	5974	5988	5972	5979	5.6	2.66
	12	5	50	0557	5704	5712	5700	5711	5774	5700	5712	5715	8	2.00
23	HP1	4	28	3418	3220	3198	3224	3278	3215	3278	3198	3227	5.5	1.54
23	111 1	4	20	3410	5220	3190	3224	5270	5215	3278	5190	3221	9	1.54
24	1100	4	35	2106	2016	2010	3078	2999	2012	2079	2912	2002	5.7	1.00
24	HP2	4	33	3186	3016	3010	30/8	2999	2912	3078	2912	3003		1.80
25	DD 1	4	07	2000	2016	2024	20.42	2010	2020	20.42	2016	2026	4	1.((
25	PB1	4	27	3090	2916	2924	2942	2918	2930	2942	2916	2 <mark>92</mark> 6	5.3	1.66
• (224					• • • • •		A A F 1		0 00 C				1.5
26	PB2	4	34	3186	2978	2986	2962	2971	2963	2986	2962	2972	<mark>6</mark> .7	1.76
													2	
27	PB4	2	29	95168	92372	9270	9250	92666	92780	92780	9237	92606	2.6	1.64
						8	4				2		9	
28	PB5	10	20	2139	2037	2032	2043	2012	2041	2043	2012	2033	4.9	1.50
													5	
29	PB6	30	40	776	706	740	765	758	766	766	706	747	3.7	3.40
													4	
30	PB7	30	37	1035	977	992	966	973	932	992	932	968	6.4	3.02
													7	
													/	

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 TABLE I: Result of Genetic Algorithm For Multidimensional Knapsack Problems.

Sr	Test	m	n	Opt			Number of				Statis			Time
no	Proble ms			ima 1			Ru	ns			Inform	ation		(s)
				-	1	2	3	4	5	Ma x	Min	Avg	%Er r	
1	SENT01	30	60	7772	7762	7746	7758	7752	7747	776 2	7746	7754	0.23	4.95
2	SENT02	30	60	8722	8712	8707	8705	8716	8710	871 6	8705	8710	0.14	4.89
3	WEING1	2	28	1412 78	1401 77	1401 83	1401 96	1401 78	14077 1	140 771	14017	1403 01	0.69	1.83
4	WEING2	2	28	1308 83	1278 62	1278 04	1281 08	1279 03	12799 3	127 993	12780 4	1279 934	2.25	1.82
5	WEING3	2	28	95667	94570	9458 2	9444 8	94684	94556	945 82	9444 8	94568	1.15	1.85
6	WEING4	2	28	1193 37	1179 82	1179 47	1178 92	1179 32	11794 2	117 982	11789 2	1179 39	1.17	1.92
7	WEING5	2	28	98796	96694	9670 2	9688 4	96879	96861	968 79	9669 4	<mark>9</mark> 6804	2.02	1.77
8	WEING6	2	28	1306 23	1285 03	1286 64	1287 04	1286 30	12863 9	128 704	12850 3	1286 28	1.53	1.91
9	WEING7	2	10 5	1095 445	1074 998	1076 681	1075 662	1074 552	1073 902	107 668 1	1073 902	1075 159	1.85	6.10
10	WEING8	2	10 5	6243 19	6231 10	6229 90	6231 84	6232 08	62311 3	623 208	62299 0	6231 21	0.19	6.08
11	WEISH01	5	30	4554	4428	4462	4448	4436	4461	446 2	4428	4447	2.35	1.93
12	WEISH02	5	30	4536	4402	4398	4372	4410	4403	440 3	4372	4397	3.06	1.87
13	WEISH03	5	30	4115	4083	4076	4083	4071	4082	408 3	4071	4079	0.87	1.90
14	WEISH04	5	30	4561	4461	4448	4457	4434	4445	446 1	4434	4449	2.46	1.92
15	WEISH05	5	30	4514	4476	4487	4462	4462	4468	448 7	4462	4452	1.00	1.87
16	WEISH06	5	40	5557	5441	5436	5392	5446	5463	546 3	5392	5436	2.18	2.44
17	WEISH07	5	40	5567	5395	5407	5437	5437	5414	543 7	5395	5392	2.83	2.52
18	WEISH08	5	40	5605	5518	5507	5532	5522	5536	553 6	5507	5523	1.46	2.38
19	WEISH09	5	40	5246	5076	5084	5102	5078	5080	510 2	5076	5084	3.09	2.68
20	WEISH10	5	50	6339	6201	6198	6182	6180	6184	620 1	6180	6189	2.37	2.76
21	WEISH11	5	50	5643	5558	5537	5529	5568	5568	556 8	5529	5552	1.61	3.07
22	WEISH12	5	50	6339	6236	6199	6214	6208	6233	623 6	6199	6218	1.91	3.02

									-					
23	HP1	4	28	3418	3368	3332	3357	3362	3341	336	3332	3352	1.93	1.82
										8				
24	HP2	4	35	3186	3109	3112	3106	3104	3114	311	3104	3109	2.42	2.12
										4				
25	PB1	4	27	3090	3012	3001	3017	3005	3015	301	3001	3010	2.58	1.93
										7				
26	PB2	4	34	3186	3094	3088	3116	3098	3089	309	3088	3097	2.79	2.06
										8				
27	PB4	2	29	95168	94207	9399	9410	94103	94111	942	9399	<u>94105</u>	1.12	1.89
						8	6			07	8			
28	PB5	10	20	2139	2099	2111	2104	2106	2115	211	2099	2107	1.50	1.76
										5	-			
29	PB6	30	40	776	763	758	759	766	769	769	758	763	1.68	3.87
30	PB7	30	37	1035	1008	1017	1012	1006	1002	101	1002	1009	2.51	3.49
										7				

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TABLE II: Result of Clustered Genetic Algorithm for Multidimensional Knapsack Problems.

