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Author(s): Carrum S. Elias, Torres -T Luis, Pérez V. Pedro Title: Group technology: Technique with comparisons

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Once again, Congratulations on your CERM 2007 acceptance. We are very excited to be able to include your research and ideas in the Conference, and we are looking forward to seeing you in September in Cuernavaca.

Best Regards

Prof. Andrea Magadán Technical Committee Chair

Majadij

Group technology: Technique with comparisons

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Abstract

In this paper it is presented a new technique to form group technology (GT), this methodology is based in the Hamming distance, also a comparison of deferent approaches to form GT is used to validate the proposed technique like neuronal networks, genetic algorithms, and other classic techniques. Results indicate a better performance of Hamming approach.

1. Introduction

In the today's manufacturing systems they are some different problems like high production time, numerous work inventory process (WIP), between others, one of the most used methodology to solve some problems like the production time and the WIP is group technology (GT) [1].

Group technology consists in identify and group families of parts from the product based in they similitude's. Each family has some similitude's this can be in the machine that is process it or in the design of the product, also GT have some benefits like [1]:

- Reduction of the starting time.
- · Reduction of the WIP.
- Reduce the transportation cost between operations.
- Reductions in warehouse space.

Also this methodology is not only for solve the cell formation problem, GT is also used to improve the performance in the mix productions systems [2], also this methodology is been used to solve some logistic and distributions problems [3], and it is not only in the manufacturing systems where this methodology was be used also in image processing and recognition [3]. This paper will be presented in the followed

sections. In the first one will be present the new technique and a summary of the deferent's techniques, then in section two will be presented the different techniques of evaluation and finally a comparison will be present with the results.

2. Summary of techniques

In this part of the paper will be presented a new technique, also will be presented a summary of classical and soft computing techniques.

Hamming method

This methodology was inspired in Hamming distance by observing the formation of clusters obtained by the result of the application this distance on a binary matrix. For the application of this technique have two cases the first one can be define as:

For quadratics matrix (m = n)

Given an initial matrix of size m * n, in where m represent the machines and the n represent the process, were it is marked with 1 if the machine makes the process, distance of Hamming is calculated in the next form:

$$d_{ij} = \sum_{k=1}^{n} z(a_{ik}, a_{jk})$$
where
$$z(a_{ik}, a_{jk}) = \begin{cases} 1 & \text{if } a_{ik} \neq a_{jk} \\ 0 & \text{on the other way} \end{cases}$$

Where a new matrix d_{ij} of size m*n ones created the new matrix it needs to obtain the values located under the sum $\sum_{i=1}^{m} d_{ij}$ for each

process and select their position with respect to the machine, in this way, the process family (PF) for the machine family (MF) need to do the same proceeding but this time with respect to the machines of the matrix.

Example 1:

	Process						
7		0	1	0	1		
1ac		1	0	1	0		
fachines		0	1	0	1		
		1	0	1	0		
	Matrix	1 ex	amp!	le1			

Given this matrix of size 4 x 4 and applying the Hamming distance the resultant matrix will seems like this:

	Process						
7	0	4	0	4			
Machines	4	0	4	0			
	0	4	0	4			
S	4	0	4	0			
	Matrix	2					

Start to obtain the values under the highest value

of the sum $\sum_{i=1}^{m} d_{ij}$ for this example the values

under 2. For the first process it is obtain the values and take this position with respect to the machines this must do it until finish all the process and it will be look like this:

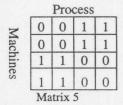
		Pro	cess	
Machines	1	2	1	2
ines	3	4	3	4
	Matrix	3		

It is taken the position that we already have it of the family witch belongs so the PF=1 & 3; 2 & 4. Now the same proceeding has to repeat with the machine families in the matrix 2 in this way we must follow the first line an take the values under 2 and take is position with respect to the machines there are left to finally obtain this:

	Machines			
_ [1	3		
Pro	2	4		
rocess	1	3		
"	2	4		

Now we take the families in with belong so the final families will be MF= 1 & 3; 2 & 4.

Finally in this way we can have a new matrix using the PF and MF like this:



For cases of matrix non-quadratics ($m \neq n$)

Given an initial matrix of size m * n, in witch the m represent the machines and the n process in were is marked with 1 if the machine makes the process. The matrix of the distance of Hamming is created of the next form:

$$d_{ij} = \sum_{k=1}^{n} z(a_{ik}, a_{jk})$$
 where
$$z(a_{ik}, a_{jk}) = \begin{cases} 1 & \text{if } a_{ik} \neq a_{jk} \\ 0 & \text{on the other way} \end{cases}$$

In this way it is create a matrix d_{ij} of size n*n to obtain the PF it need to be selected one part of the matrix d_{ij} this part must be of size n*n-1 and repeat the same steps using in the quadratic matrix. For the formation of MF needs to be selected a part of the matrix d_{ij} this part of the matrix must be of size m*m and make the same proceeding.

Similar Coefficients:

This technique is based in the similitude between two machines this similitude is defined as [3]:

$$s_{ij} = \frac{\sum_{k=1}^{n} d^{1}(a_{ik}, a_{jk})}{\sum_{k=1}^{n} d^{2}(a_{ik}, a_{jk})}$$

Where

$$d^{1}(a_{ik}, a_{jk}) = \begin{cases} 1 & \text{if } a_{ik} = a_{jk} = 1 \\ 0 & \text{on the other hand} \end{cases}$$

$$d^{2}(a_{ik}, a_{jk}) = \begin{cases} 0 & \text{if } a_{ik} = a_{jk} = 0\\ 1 & \text{on the other hand} \end{cases}$$

The GT will be formed when the families have the same coefficient of similarities.

Sorting-based algorithms or ROC

This technique is based on classify from more to less all the columns and the lines of the initial matrix using this expression [3]:

$$p_i = \sum_{k=1}^n a_{ik} 2^{n-k}$$

The P-median model

This technique is used to group parts into PF, this technique use the concept of distance between parts d_{ij} , this distance can be defined in two deferments ways one is the hamming distance and the other is the distance Mintowski for this work it is used the hamming distance. The GT problem can be solved used the follow binary program:

$$\min \sum_{i=1}^n \sum_{j=1}^n d_{ij} x_{ij}$$

Subject to:

$$\sum_{j=1}^{n} x_{ij} = 1; \ i = 1, \dots, n$$

$$\sum_{j=1}^{n} x_{jj} = p$$

$$x_{ij} \le x_{jj}$$
 for all $i = 1,...,n$; $j = 1,...,n$
 $x_{ij} = 0,1$ for all $\begin{cases} i = 1,...,n \\ j = 1,...,n \end{cases}$

Where p is the specify number of PF.

ART

The figure 1 shows the general structure of an artificial neuronal network ART-1 [4][5].

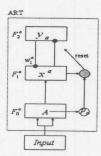


Figure 1. Basic structure of the ART-1

Genetics Algorithms

The type of genetic algorithm (GA) used to form GT was the ranking order genetic algorithm [6] the scheme is presented below:

- Initialized the population random.
- Evaluate each one of the solutions and assign the fitness.
- Choose the population witch have the best fitness.
- Mutate and make crossover to al solutions chosen.
- Repeat all this one considerate number of times

For see the behavior of the partially separable clusters in this part of the experimentation was used the ranking order genetic algorithm with the follows parameters:

- Number of individuals = 1000
- Crossover PMX
- Mutation interchange
- Number of generations = 100
- Crossover probability = .85
- Mutation probability = 0.1

The evaluation function is a classic method used to form group technology called Bond Energy algorithm given by:

MIN ME

$$ME = 1/2 \sum_{i=1}^{m} \sum_{j=1}^{n} a_{ij} \left[a_{i,j-1} + a_{i,j+1} + a_{i-1,j} + a_{i+1,j} \right]$$

3. Description of techniques of evaluation

Hamming Distance

This criterion of comparison consist in select the matrix with less distance and is given by:

$$d_{ij} = \sum_{k=1}^{n} z(a_{ik}, a_{jk})$$
where
$$z(a_{ik}, a_{jk}) = \begin{cases} 1 & \text{if } a_{ik} \neq a_{jk} \\ 0 & \text{on the other way} \end{cases}$$

Mitowski Distance

This criterion of comparison consist in select the matrix with less distance and is given by:

$$d_{ij} = \left[\sum_{k=1}^{n} \left| a_{ik} - a_{jk} \right|^{r} \right]^{1/r}$$

Where r is an entire positive and n is the number of parts.

Mitowski Euclidean Distance

This criterion of comparison consist in select the matrix with less distance and is given by:

$$d_{ij} = \left[\sum_{k=1}^{n} \left| a_{ik} - a_{jk} \right|^{2} \right]^{1/2}$$

Where n is the number of parts.

Bond Energy

This criterion of comparison consist in select the matrix with most ME and is given by:

$$ME = 1/2 \sum_{i=1}^{m} \sum_{j=1}^{n} a_{ij} \left[a_{i,j-1} + a_{i,j+1} + a_{i-1,j} + a_{i+1,j} \right]$$
 Check of efficiency

Using the criteria of efficiency employ by Chandrasaekharaq & Rajagopalan [7]. Where we use the global efficiency denoted by η this method is calculate in the follow way:

$$\eta = qn_1 + (1-q)n_2$$

Where:

$$n_1 = \frac{e_d}{\sum_{r=1}^k M_1 N_1}$$

$$n_2 = 1 - \frac{e_0}{mn - \sum_{r=1}^{k} M_1 N_1}$$

In this method it is selected the matrix with more global efficiency.

Efficacy Criteria

Using criteria of efficiency employ by Chandrasaekharaq & Rajagopalan [8]. As the criteria efficacy we select the matrix with more criteria of efficacy and is given by:

$$\mu = \frac{N_1 - N_1^{Out}}{N_1 + N_0^{In}}$$

Distance to diagonal

This method consists in found the less distance between the group formed and the diagonal matrix.

Machine Utilization

This method consist in select the most machine utilization [9]:

$$Mu = \frac{e_1}{\sum_{k=1}^{g} j_k i_k}$$

Group efficiency

This method consist in select the most Group Efficiency [9]:

$$Ge = 0.5Mu + .5[1 - \frac{e_e}{(MN - \sum_{k=1}^{g} j_k i_k)}]$$

4. Experimentation

To make the comparisons between techniques and techniques of evaluation were selected nine different matrixes they are listed below:

Name	Size Referen		
Kusiak	4x5	Kusiak 1998 [3]	
Problem B	6x7	Created	
Problem C	6x7	created	
Rajagopalan	6x11	Rajagopalan 1989 [7]	
Pump Machine	15x8	Internet	
Factory	25x27	Carrum 2005 [10][11]	
Carrie	20x35	Carrie 1973 [12]	
Rajagopalan 1989	24x40	Rajagopalan 1989 [8]	
Rajagopalan 1987	40x100	Rajagopalan 1987 [7]	

Table 1 matrix of experimentation

From the nine different matrixes in this work where obtain several tables of results showed in the annexes of this paper. To select the techniques that made a better grouping selecting those ones who have been more scored in the techniques of evaluation, this was made selecting between all the techniques those how have better evaluation in each matrix (table 2)

Name	Winner Genetic Algorithm ART, ROC, P-median Genetic Algorithm				
Kusiak					
Problem B					
Problem C					
Rajagopalan	Genetic Algorithm, ROC				
Pump Machine	Genetic Algorithm, Hamming Method ART, ROC, Hamming Method				
Factory					
Carrie	ART, ROC, Similar Coefficients, Hamming Method				
Rajagopalan 1989	ART, ROC, Genetic Algorithm, Similar Coefficients, Hamming Method				
Rajagopalan 1987	Genetic Algorithm, Hamming Method				

Table 2 experimentations results

5. Conclusion and Future Work

Where can find that the better techniques to form group technology based in the size of the matrix and in the better group evaluation are:

- Genetic Algorithm
- ART
- Hamming Method

As future work it is necessary to make an autonomous technique, make a new ways of using the genetics algorithms combined with other techniques or make hybrids techniques to use more parameters like cost, time of movement of the pieces, and make some simulations to se how it is the behavior of the new technique.

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		App	pendix					
					Kusiak			
		Inicial	ART	ROC	GA	P m e d ia	S C 4 8	Hamming 48
Distancia Hamming		4 8	4 8	4 8	48	4 8	4 8	4 8
Distancia Minkowski Distancia Minkowski Euclidiana	_	4 8	8 4	8 4	8 4	8 4	8.4	8 4
E fficacy	_	14	1 4	1.4	1 0	10	10	10
Rajagopalan		-1.2	-1.2	-1.2	-0.64	-0.64	-0.64	-0.64
Dr. Luis		18	22.66	22.66	6.66	8	8	8
Bond Energy		0.5	0.5	0.5	4 . 5	4.5	4.5	4 . 5
					Problem			
		Inicial 80	A R T 8 O	ROC	6 A	P m e d ia	8 C	Hamming 80
Distancia Hamming Distancia Minkowski		80	80	8 0	8 0	8 0	8 0	8 0
Distancia Minkowski Euclidiana		80	8 0	8 0	8 0	8 0	8 0	8 0
E ffica cy		- 11	1 0	1 0	1 2	1 0	1 1	1 2
Rajagopalan		-1.5	-1.4	-1.4	- 1 . 6	-1.4	-1.5	-1.6
Dr. Luis		10.2	5 . 4	5 . 4	1 0 . 2	5 .4	1 0 . 2	1 0 . 2
Bond Energy		0	0.5	0 . 5	0.5 Proble	0.5 m a C 6 x 7	0	0.5
		Inicia	I ART	ROC		Pmedia	sc	Ham ming
Distancia Hamming		1 2 8	1 2 8	1 2 8	128	1 2 8	1 2 B	1 2 8
Distancia Minkowski		1 2 8	1 2 8	1 2 8	1 2 8	1 2 8	1 2 8	1 2 8
Distancia M.inkowski Euclidiana		2 4 0	2 4 0	2 4 0	2 4 0	2 4 0	2 4 0	2 4 0
E Ifica cy		1 7	1 6	2 1	2 1	1 5	2 1	1 7
Rajagopalan		-1.4	-1.3	-1.8	-1.8	-1.2	-1.8	-1.4
Dr. Luis Bond Energy		29.2	1 5 .6	5.5	4 9 . 2	1 6 5 . 5	5.5	2 0
Bond Energy			3.5	3.3	Brown		3.3	3.0
		Inicial	ART	ROC	G A	Pmedia	s c	Hamming
Distancia Hamming		3 2 8	3 2 8	3 2 8	3 2 8	3 2 8	3 2 8	3 2 8
Distancia Minkowski		3 2 8	3 2 8	3 2 8	3 2 8	3 2 8	3 2 8	3 2 8
Distancia Minkowski Euclidiana		5 5 6	5 5 6	5 5 6	5 5 6	5 5 6	5 5 6	5 5 6
Efficacy Rajagopalan		-2.68	-2.68	-2.8	-2.74	-2.68	-2.68	-2.74
Dr. Luis		9 3	9 3	9 3	7.4	9 3	9 3	9 1
Bond Energy		5 . 5	5 . 5	7 . 5	7 . 5	5 . 5	5 . 5	6
						hine 15x8		
		Inicial	ART	ROC	G A	P m e d ia	s c	Ham ming
Distancia Hamming Distancia Minkowski		302	3 0 2	3 0 2	3 0 2	302	3 0 2	3 0 2
Distancia Minkowski Distancia Minkowski Euclidiana		999	999	9 9	999	999	999	999
E Ificacy		3 3	3 3	3 3	3 3	3 3	3 3	3 3
Rajagopalan		-1.9	-1.9	- 1 . 9	-1.9	- 1 . 9	- 1 . 9	-1.9
Dr. Luis		5.8	3 9	6 7	1 0 5	5 8	5 8	3 6 . 5
Bond Energy		3	1 3	1 0	1 5	3	3	1 2 . 5
		nicia I	ART	ROC	Planta 2	P m e d ia	s c	Ham ming
Distancia Hamming		1532	1532	1532	1532	0	1532	1532
Distancia Minkowski		1 5 3 2	1 5 3 2	1 5 3 2	1532	0	1532	1 5 3 2
Distancia Minkowski Euclidiana		1922	1922	1922	1922	0	1922	1 9 2 2
E flica cy		7.74	5 5	5 5	56.96	0	5 4 .9 6	5 5
Rajagopalan Dr. Luis	-	66.25	87.6	87.6	302.8	0	149.33	87.6
Bond Energy		2.5	4 . 5	4 . 5	4 . 5	0	4	4 . 5
					Carrie 20			
Distancia Hamming	In ic		3 4 4	8 O C	G A 4 3 4 4	P m e d ia	8 C	Hamming 4344
Distancia Minkowski	4 3		3 4 4	4344	4344	0	4344	4344
Distancia Minkowski Euclidians	293		9 3 2 4	29324	2 9 3 2 4	0	29324	2 9 3 2 4
E fficacy	15		152	1 5 2	154	0	152	1 5 2
Rajagopalan Dr. Luis	883			4 3 3 . 1 1	883.52	0	4 3 3 .1 1	4 3 3 .1 1
Bond Energy	2		6.5	96.5	2 0	0	96.5	96.5
		AR			palan (198			
Distancia Hamming	Inicial 5330	5 3 3		5 3 3 0	5 3 3 0	Pmedia	5 C	Hamming 5330
Distancia Minkowski	5 3 3 0	5 3 3	0	5 3 3 0	5 3 3 0	0	5330	5 3 3 0
Distancia Minkowski Euclidiana	30031	300		30031	3 0 0 3 1	0	30031	3 0 0 3 1
E fficacy Rajagopalan	151	.6.		151	1 5 2	0	152	1 5 1
Dr. Luis	1064	898	07179	8 9 8 0 7 1 7 9			1092.1	
Bond Energy	2 3	9 9		9 9	77.5	0	7 7 -5	9 9
	-	Inicial	ART	Rajag	opalan (1	987)100x4	0	1
Distancia Hamming	-	29736	2957			A Pme 36 0		2 9 5 7 0
Distancia Minkowski		29736	2957					29570
Distancia Minkowski Euclidiana		3 2 5 9 8 8	3 2 2 4					3 2 2 4 9 9
E Ifica cy		0	0	0				0
Rajago palan Dr. Luis	_	5 2 6 5	0 496	1.3 0	0			0
Bond Energy		78.5	280.		191	96.8 . 0		4961.3
					. 9 1			200.5