

Scheduling on restricted uniformly related machines.

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1 Abstract

In this project we try to solve scheduling problem using three heuristics: Local search, Branch and bound, and genetic heuristic. We write three unattached projects in each one we use one heuristic; we can get better and faster algorithms by combining more than one heuristic or by using more complex algorithms like evaluating pairs of chromosomes instead of chromosomes, and using more complex target functions in genetic. But the main idea of the project was to learn the basic concept of the following heuristics.

The problem is Scheduling on restricted uniformly related machines. where the Input is An integer number of machines $m \geq 2$. A set of n jobs $J = \{1, 2, \dots, n\}$ where job j has an integer processing time $P_j > 0$. Machines speeds $s_i \in \{1, 2, 4\}$ for $i = 1, 2, \dots, m$, we solve the same problem in the three heuristics.

2 Local Search

The local search algorithm It consists of several steps we will Explain the algorithm step by step

2.1 pre preparing(initial solution)

to start running the local search algorithm first we need initial solution. we have as input two files the first file is for tasks where we have on each line task time (integer value), the second file is for machines where each line contains machine speed (integer 1, 2, 4). from the input we build two vectors the first one holds the tasks and the second holds the machines. after having data structure hold the tasks and the machines we should Hand out the tasks into the machines and make initial solution, so we need greedy algorithm its not important how many the greedy algorithm is good. also if we hand out the task randomly at the end we will have the same result, so for convenience we insert the numbers to minimum heap to sort the tasks and then we Turned on the tasks into machines In a fair way (In a circular) and on each time we give the same amount tasks as the current machine speed value to the machine, for example if we have machine with speed 2 we will give it two task each time, and if the machine speed 4 we will give it 4 tasks each time. we do that while we still have tasks, and when we hand out all the tasks we have an initial solution.

2.2 Local Search for two machines

on this subsection and on the next subsections we will solve small problems and then we will combine all of them together in the last subsection creating the full problem solution.

the current problem is given two machines m_1 and m_2 we want to find the best tasks from m_1 and m_2 that if we swap them between m_1 and m_2 we get the best Improvement to achieve the Goal . so now we should define what is the goal that we want to achieve.

giving two machines m_1 and m_2 where the time of m_1 is X and the time of m_2 is Y . let assume that we chose k tasks from m_1 and r tasks from m_2 now we should evaluate the time after swapping the tasks (simulation) between the two machines by passing k tasks from m_1 to m_2 and removing the k tasks from m_1 ,and the same by passing r tasks from m_2 to m_1 and removing the tasks from m_2 . lets define the time of m_1 after the change is A and the time of m_2 after the change (swapping the tasks) is B . the our goal that we want to achieve:

$(\max\{A,B\} < \max\{X,Y\} \ || \ (\max\{A,B\} = \max\{X,Y\} \text{ and } (A+B) < (X+Y))$

that means if we save the best solution and we find new solution by swapping k, r tasks between m_1 and m_2 and that solution better than the best solution for now, we save it as the best solution and we save r, k tasks and then we check the other k, r combination with the new best solution after we check all possible to choose r, k we have the best r, k that by swapping them we achieve our goal NOTE(k and r is constant).

the idea of adding $(\max\{A,B\} = \max\{X,Y\} \text{ and } (A+B) < (X+Y))$ is to give priority to machines with high speed we want to fill machines with high speed also if the final solution still the same $(\max\{A,B\})$, because that help us to find bigger task to swap in the future.

2.3 All possible combination of r,k

as we saw in previous section we need to go through all possible combinations of choosing k tasks from m_1 (the same about r) as we know from mathematics we have $\binom{n}{k}$ different combination. where n is the number of tasks on m_1 and k is the number of task that we want to choose from m_1 to swap them.

we can solve the combination problem easily by using recursion and array with size k (int data[k]) , we go through all the n tasks we start from first task (first function call) and we call the function recursively twice the first call with array data that we add to it task1 and the second one with array data without adding to it task1, in general in the p function call we call the function recursively twice ,the first time with adding $task_k$ to data array and the second with out adding it. and when data array full we fill all the k cells we have new combination.

in other words the idea that each task has two possibilities the first one is to be part of the combination and the second its not part of the combination.

2.4 combination of two machines

i go throw all possible combinations of two machines twice on different order the code is so easy so writing the code more sample than to describe it :

```
1 for (int offset = 1; offset < M.size(); offset++) {
2     for (int i = 0; i < M.size() / 2; i++) {
3
4         // the index of first machine i * 2
```

```

5 // the index of second machine (i * 2 + offset) % M.size()
6 //M.size() is number of machines
7 GetBestOfNxM(i * 2, n, (i * 2 + offset) % M.size(), m, 0, d, 0, true);
8 if(!GetBestOfNxMbool)
9 SwapmTasks(NxMcom1Best, i * 2, NxMcom2Best, (i * 2 + offset) % M.size());
10 flag = flag && GetBestOfNxMbool;
11
12
13 }
14 }
15
16 for (int offset = 0; offset < M.size(); offset++) {
17     for (int i = 0; i < M.size() ; i++) {
18
19         // the index of first machine offset
20         // the index of second machine i
21         //M.size() is number of machines
22         GetBestOfNxM(offset, n, i, m, 0, d, 0, true);
23         if (!GetBestOfNxMbool)
24             SwapmTasks(NxMcom1Best, offset, NxMcom2Best, i );
25         flag = flag && GetBestOfNxMbool;
26     }
27 }

```

2.5 all together

we show above how to throw all posible combination of pair of two machine. and on each combination of two machines how to go throw all possible combination of k tasks from m_1 and r tasks from m_2 (where k and r is constant) and how to choose the best combination of choosing k, r (how we apply LocalSearch on this two machines) (we check all the combination of choose k from $machine_1$ with all the combination of choose r from $machine_2$ "each combination from $machine_1$ with all the combination of $machine_2$ ")

so we can write one function $LocalSearchNxM(k, r)$ and applied LocalSearch that swap k tasks from first machine with r from second machine and we go throw all compination of two machine like we said above

this is the main LocalSearch Function:

Where:

LevelZero(): is function implement local search that only pass one task each time from $machine_1$ to $machine_2$ with the same rolles above.

LocalSearchNxM(k,r): function implement localsearch where we want swap each time k tasks from m_1 with r tasks from m_2

if the there is no improvement in the current level we did not run the next level , and if there is no level on this itertion has improvement we stop the while .and after we stop we print the result

```

1 void LocalSearch() {
2     bool flag = true;
3     bool temp;
4     while (flag) {
5
6         flag = flag && LevelZero();
7         for (int i = 1; i <= maxLevelSearch; i++) {
8             for (int j = i; j <= maxLevelSearch; j++) {

```

```

9      if (j == 1 && i == 1) {
10         temp = LocalSearchNxM(i, j);
11         if (temp) {
12             i = maxLevelSearch+1; j = maxLevelSearch+1;
13         }
14         flag = flag && temp;
15     }
16     else {
17
18         if (j == 1) {
19             if(TasksTable[i-1][maxLevelSearch]>0)
20                 temp = LocalSearchNxM(i, j);
21             if (temp) {
22                 i = maxLevelSearch+1; j = maxLevelSearch+1;
23             }
24             flag = flag && temp;
25         }
26         else {
27             if (TasksTable[i][j-1] > 0)
28                 temp = LocalSearchNxM(i, j);
29             if (temp) {
30                 i = maxLevelSearch+1; j = maxLevelSearch+1;
31             }
32             flag = flag && temp;
33         }
34     }
35
36
37
38
39
40     }
41 }
42
43
44
45     flag = !flag;
46
47 }
48 }

```

3 Branch and bound

on this heuristic we should build tree, each node on the tree have lower bound and higher bound , each edge bettween father and son represent handing out task to the son, where each node represent machine , so each father has at most children's as the number of machines, so the path from the root to each node in the level- k (node with deapth k) represent handing out the first k task where $edge_j$ represent handing out $task_j$, but we will not build all the tree because if there node that have lower bound equal or greater than any upper bound on any node on the tree there is no need to continue to his son's because we will not get better soultion than the node with the upper bound, for that reason we have also a global upper bound that his value is the minimum value on all the upper bounds that we have on all the nodes that we build to that point.(on building each node

we update his value if the node upper bound is less than the global). upper bound can be greedy algorithm result but unlike Local search the greedy algorithm is important because bad greedy algorithm mean that we need to build more nodes on the tree, and as result we will have to wait more time to get the solution, so here for example its good idea to set the result of local search as upper bound but we did not do that because most of the time we get optimal solution using local search and our goal to learn the heuristic and to find the best way to calculate the upper and lower bounds.

so our greedy algorithm is LPT (Longest Processing Time Algorithm) . also the sequence of handing out the tasks on branch and bound algorithm is important by handing out the big tasks first can help us to build less nodes because the lower bound grow and get bigger faster, also this issue in line with LPT ,so $task_1$ is the biggest task and $task_n$ is the smallest task , and on each path $edge_p$ represent handing out the $task_p$.

At the end we want to run the program on personal computer and our challenge to build algorithm with good Space complexity (use less amount of memory space) . if we use BFS tree traversals algorithm we need a huge memory because on the first levels of the tree we can't determine if the path is good and we need to build and save more nodes that we can't remove node before we make sure that this path is bad path and we will not continue to his son's, also if we be satisfied to save only one level (depth) each time (save the last level of depth nodes and one each nood save the path from the root (each task to each machine)) the amount of nodes could be huge , despite that the BFS does not guarantee better solution, on other hand using DFS we can build solution with linear space complexity and also that help the global upper bound to decrease faster because we cant to determine tight upper bound if we still have many tasks to handing out because the upper bound will be for all possible different solution of handing out the other tasks (maximum possible upper bound), so going deeper help us make tighted upper bound, and DFS go deeper fast. we can build solution using DFS that his maximum space complexty $m * n$ where m is the number of machines and n is the number of tasks.

we have customized DFS in our solution the default is to going to the worst son first (son's with the highest time first) we have the option going to the best son's first (have lowest time first (where the time is the time of the worst machine)) by deleting (`#define BadFirst`) on file scheduling.h . the idea of choosing best son's first is to get tight upper bound faster and that help us to build less nodes, on the other hand the idea of choosing worst son's first is to make the lower bound grow faster and as result we build less nodes, we try the three options without sorting(original DFS) ,bad first and best first we get the result of bad son's first is the best. and we also have to option to sort the tasks at the beginning like we said above big tasks first or run without sorting by removing the define (`#define _WithSort`) but as we explain with sort is more better . to save time also i added some rules: if we have optimal solution we stop, and if there two son's or more with the same speed and the same time we call recursively only one of them, and if i have solution better than the global upper bound we update the upper bound to it.

4 Genetic algorithm

our default Genetic algorithm have on each generation 100 Chromosome ,and we build new generation by four mutation and 48 pairing ,where by mutation we can made one new Chromosome ,and by pairing we can made two new Chromosomes the default number of generation one thousand, all the parameters above you can change them on scheduling.h they they defined as `#define` . we

have also about twenty fitness function where the default is function number 17 you can also choose more than one function and the program print different files for each function as result, you can choose interval of function to run by changing the defines(`#define Bf` , `#define Ef`)

4.1 first generation

we have function that build vector with size m where m is machines number and the sum of the values on the vector is n where n is the number of tasks, assume the name of the vector is *vec* we give each *machin_j* , *vec[j]* tasks and on this way we have one solution and we build on Chromosome we repeat that 100 times making 100 Chromosomes (each time we build from scratch *vec* and handing out the tasks to the machines randomly) we make the survival Chromosome the best chromosome time

4.2 mutation

we peak random numbers k, m where $k, m \leq n$ and $m - k < n$ where n is the tasks number ,and in Chromosome class we have hashmap that have pairs <task index, machine index> we use that hash map to change the machine for each $k \leq \text{task index} \leq m$ for each task index we peak randomly a machine the default is that new machine should be different than the old ,but there is an option to enable peak the same machine (it will be like to randomly choose who from this interval we change).

its ok that we change interval of sequence of indexs and we didn't peak randomly tasks because the tasks it self random and doesn't sorted ,we change the machine index in the hash map and also pass the task from the old machine to the new one the we peak randomly.

4.3 pairing

as we saw in mutation we have hash map the same thing we peak randomly k, m but here if we have j where $k < j < m$ and in *Chromosome₁* *task_j* in machine m_A and in *Chromosome₂* *task_j* in machine m_C we swap bettween the machines in the two Chromosomes that mean after the change we will find *task_j* in *Chromosome₁* at m_C , and *task_j* in *Chromosome₂* at m_A there is two option of pairing the default that *Chromosome₁* \neq *Chromosome₁* (we can't pair Chromosome with it self) the second option allow that as kind of mutation

4.4 Fitness function

for each Chromosome we call the fitness function and build probability vector see 4.5 ,we have 20 Fitness function that they combination of:

Y: Chromosome time (the time of maximum machine)

X:Optimal time (the sum of tasks dividing the sum of machines speed)

a : median time (the machine that have the median time on this Chromosome) worsGM : the worst machine in all the Chromosomes

min : the machin with the smallest time in the current Chromosomes (local)

max (the same Y) : the machin with the biggest time in the current Chromosomes (local) == Y

4.5 randomly peaking Chromosome for mutation/pairing

for each Chromosome we call the fitness function and enter the result into vector then we normalize the vector to probability that the sum of all the cells on the vector is one $\sum_{i=0}^{vector.size} vector.at(i) == 1$ now we have on $vector_i$ the probability of $Chromosome_i$ after that we apply to the vector that code

```
1 for (int i = 1; i < generation.size(); i++)  
2     vector.at(i) = vector.at(i) + vector.at(i - 1);
```

now we have monotonically increasing vector between zero to one we peak random number between zero and one and find out where it falls in the vector and return the appropriate Chromosome here the default repeating random but there is an option to enable non-repeating random

4.6 build new generation

as we said above we build it by 4 mutation and 48 pairing.