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**Heuristic Optimisation**  
**Implementation Exercise 1**  
*Iterative improvement algorithms for the PFSP*

# 1 Introduction

The goal of this assignment was to...

## 2 Code Use

### 3 Exercise 1.1

#### 3.1 The Results

We list here the average percentage deviation from the best solutions for each algorithm tested, along with the average computation time:

Algorithm	APD	ACT(ms)
-random-exchange-best	26	1362
-random-exchange-first	10	5744
-random-insert-best	16	2909
-random-insert-first	6	12869
-random-transpose-best	354	28
-random-transpose-first	340	37
-slack-exchange-best	32	1244
-slack-exchange-first	11	3944
-slack-insert-best	18	2523
-slack-insert-first	6	8661
-slack-transpose-best	243	43
-slack-transpose-first	240	47

Figure 1: The APD and ACT for each algorithm.

We see that the transpose mode for the neighbourhood generation is the worst, and by far, but is the less costly in time. The insert mode is the best when it comes to the quality of the solution, but is the worst in time cost. The remaining exchange mode is between the other ones, half less time than the insert mode, and a quality of solution a bit lower than the insert mode, but very far from the quality of the transpose mode.

The reason behind the difference between these 3 modes becomes clear when we consider the size of the respective neighbourhood. If  $N$  is the number of jobs and  $p$  is the current permutation, we get these sizes:

$$\begin{aligned} |N_{transpose}(p)| &= N - 1 \\ |N_{exchange}(p)| &= \sum_{i=0}^{N-2} N - i - 1 = (N - 1)^2 - \frac{(N - 1)(N - 2)}{2} \\ |N_{insert}(p)| &= N(N - 1) \end{aligned}$$

We observe that the SLACK heuristic makes the time cost decrease everywhere, except for the transpose mode. As an attempt to explain this, we could say that the size of the neighbourhood, being a lot smaller than the other mode (linear in the number of jobs), has a much lower impact on the time cost. The real impact comes from the heuristic at the beginning.

The APD is always lower for the "first" mode than the "best" mode. Would this be explained by the fact that more diverse solutions are found with the first mode ? Indeed, we see that the time consumed for the first mode is much superior than the time used for the best mode. And if we look at the number of intermediate solutions, the number is much higher. We could say that with the best mode, we explore more locally than with the first mode, and get stuck quickly in a local maximum.

### 3.2 Difference between the solutions

We used the Student t-test to determine whether there is a statistically significant difference between the solutions generated by the different perturbative local search algorithms. The obtained p-values for the Student t-test are shown in figure 2. The algorithms corresponding to the numbers are the ones from figure 1, in the same order (alphabetically sorted).

The Student t-test can be used to test statistically the mean equality null hypothesis. The p-value corresponds to the probability that the null hypothesis is incorrectly rejected. If this value is below the significance level, the null hypothesis is rejected and thus the means are very different. If the value is above the significance level, the hypothesis is incorrectly rejected, thus accepted, and the means are approximately equal. The higher the value, the closer the means.

Here is the R code used to write the p-values to a file. The results matrix was created by taking the average percentage deviation of each algorithm and each instance (matrix of dimension 60x12).

```
1 a <- read.table("R-avRelPer—random—exchange—best.dat")$V1
2 ...
3 l <- read.table("R-avRelPer—slack—transpose—first.dat")$V1
4
5 results <- c(a,b,c,d,e,f,g,h,i,j,k,l)
6 results <- array(results, dim=c(60,12))
7
8 test = numeric(length = 60)
9 test2 = numeric(length = 60)
10
11 x = 1
12 for (i in 1:11) {
13   for (j in (i+1):12) {
14     test[x] <- t.test(results[,i], results[,j], paired=T)$p.value
15     test2[x] <- wilcox.test(results[,i], results[,j], paired=T)$p.value
16     x = x + 1
17   }
18 }
19
20 write(test, file = "p values st test", ncolums = 1)
21 write(test2, file = "wilcox p-values", ncolums = 1)
```

We take a significance level of 0.05 ( $\alpha = 0.05$ ).

	1	2	3	4	5	6	7	8	9	10	11	12
1		2.325677e-10	0.0001764137	1.492921e-10	2.324708e-07	1.397426e-08	0.009320388	1.814296e-10	0.0009812694	1.651045e-12	1.002514e-09	3.956704e-09
2			1.380619e-07	5.676628e-05	1.667657e-07	1.022813e-08	1.119052e-10	0.2236329	1.656816e-12	1.904083e-08	6.981541e-10	2.61601e-09
3				2.616446e-12	2.784541e-07	1.86176e-08	4.364759e-06	6.281086e-05	0.2084699	9.391734e-12	2.116133e-09	7.112663e-09
4					1.560699e-07	1.017382e-08	5.164549e-10	4.937766e-05	6.505518e-17	0.3771135	7.162175e-10	2.629253e-09
5						0.3131815	3.198631e-07	1.624651e-07	2.896557e-07	1.378438e-07	0.0002024202	5.743684e-05
6							1.81788e-08	1.005804e-08	2.054526e-08	8.50651e-09	1.793673e-05	7.765756e-07
7								4.646304e-11	3.193926e-05	1.330835e-11	1.541866e-09	5.902744e-09
8									6.202973e-11	1.221381e-07	6.366285e-10	2.46713e-09
9										8.756858e-19	1.987018e-09	7.336251e-09
10											5.066557e-10	1.945128e-09
11												0.4696791
12												

Figure 2: P-values for each combination of algorithms (Student t-test). Paired test, with each possible pair of algorithms.

4

	1	2	3	4	5	6	7	8	9	10	11	12
1		1.6479e-11	6.566262e-07	1.650747e-11	1.66871e-11	1.668972e-11	0.001887301	3.558256e-11	3.880961e-05	1.649194e-11	1.66871e-11	1.66688e-11
2			1.401733e-07	5.791151e-06	1.670281e-11	1.670805e-11	1.646866e-11	0.7562894	7.25813e-09	1.681026e-08	1.668972e-11	1.667926e-11
3				1.218116e-10	1.669234e-11	1.670281e-11	2.685764e-09	9.410355e-06	0.07339366	1.376776e-09	1.669757e-11	1.667664e-11
4					1.669757e-11	1.667664e-11	1.648417e-11	4.229889e-07	2.965186e-11	0.861429	1.670281e-11	1.668449e-11
5						0.7742204	1.669757e-11	1.669495e-11	1.670019e-11	1.670543e-11	4.520244e-11	3.454208e-11
6							1.66871e-11	1.669495e-11	1.669495e-11	1.669757e-11	1.66688e-11	2.449249e-11
7								1.642735e-11	5.420053e-09	1.657235e-11	1.670281e-11	1.669757e-11
8									2.515346e-08	3.401544e-08	1.670019e-11	1.670019e-11
9										5.411351e-11	1.66871e-11	1.667403e-11
10											1.670019e-11	1.669495e-11
11												0.0002718787
12												

Figure 3: P-values for each combination of algorithms (Wilcoxon test). Paired test, with each possible pair of algorithms.

The Wilcoxon test does not assume that the population is normally distributed. We see on figure 3 that the p-values give the same idea of the difference between the solutions, except for the combination 11x12. We can see that the hypothesis is rejected in the Wilcoxon test but not in the Student t-test.

We can see on the figures that a lot of values are below the threshold  $\alpha$ . We can thus conclude that most of the algorithms generate solutions of significantly different quality. We can check it in figure 1.

## **4 Exercise 1.2**