

Artifacts of “End-to-end Scheduling of Real-time Task Pipelines on Multiprocessors”

Constraint Solvers

1. CoPi: Our heuristic constraint solver algorithm.
2. GEKKO: The APOPT Solver: <https://machinelearning.byu.edu/>

Most experiments compare the above two. We have also experimented with two other MINLP solvers but only for the first experiment:

1. `pyomo`: <http://www.pyomo.org/>
2. `scipy`: <https://docs.scipy.org/doc/scipy/reference/optimize.html>

Requirements

Please follow the installation procedure of `GEKKO`, `pyomo`, `scipy`, `pickle` (for reading and writing dataset). We believe that the code should work for python version 3.5 and 3.6. We have run all the experiments with python 3.6. If not sure, please use `virtualenv` to install python 3.6 .

Algorithms and Files

The CoPi algorithms are at the root location of the source directory. The MINLP solver implementations are in `ilp/`.

CoPi files are: 1. `copi_e2e.py`: For only E2E Delay Constraint implementation. 2. `copi_all.py`: For all the constraints. 3. `copi_lib.py`: Contains the CoPi algorithm that solves both constraints. 4. `multi_pipeline.py`: For the multiprocessor experiments with CoPi.

MINLP Solvers: 1. `ilp/ilp_gekko.py`: GEKKO constraint optimization solution. 2. `ilp/ilp_pyomo.py`: `pyomo` constraint optimization solution. 3. `ilp/ilp_scipy.py`: `scipy` constraint optimization solution.

Datasets Random pipelines are generated using UUnifast algorithm. All the dataset files are `pickle` files. 1. `dataset_*` files are for uniprocessor and runtime experiments. 2. `pipelines_*` files are for multiprocessor experiments. 3. `watersdata_*` files are for WATERS 2015 Workshop paper dataset.

Helper function files: 1. `utility.py`: Some helper functions for utilization, RMS bound and other calculations. 2. `pipeline.py`: Most functions related to a pipeline are here. For example, end-to-end delay, loss-rate, etc.

Figures The figures are provided in the **Figures** directory.

All the figures could be generated as PDF files by:

```
cd Figures
make
```

Experiments

The next subsections correspond to the subsections in the Evaluation section of the paper.

Uniprocessor Acceptance Ratio Experiments

Only for End-to-end Constraint

1. Run all the experiments for CoPi:

```
./run_exp1_copi.sh
```

2. Run all the experiments for GEKKO (APOPT):

```
./run_exp1_gekko.sh
```

3. Run all the experiments for pyomo (IPOPT):

```
./run_exp1_pyomo.sh
```

4. Run all the experiments for scipy (trust-constr):

```
./run_exp1_scipy.sh
```

Result All the data will be written to `accepted_sets_<solver>.txt` in space separated format for $LBG = \{11, 12, 14, 15, 16, 18\}$. For example, `accepted_sets_copi.txt` is for CoPi.

Paper results are summarized in `Figures/accept.csv`. (3rd, 5th, 6th, 7th lines are respectively CoPi, GEKKO, scipy and pyomo.)

*The results may vary a bit depending on the changes in the random dataset, but the basic overview of the result should remains same as the result in the paper and in **Figures** folder.*

Both End-to-end Delay and Loss-rate Constraints

1. Generate CoPi data points by the following script:

```
./run_loss_rate_copi.sh
```

The `accepted_lr_copi.txt` will have the number of accepted (schedulable) pipelines in space separated format for {0, 25, 50, 75}% loss-rate

2. Generate GEKKO data points by the following script (result will be appended to `accepted_sets_gekko.txt`, so if you can remove the file if you do not want to see previous results):

```
./run_gekko_lr.sh
```

All the data will be appended to `accepted_sets_gekko.txt`.

3. Generate GEKKO (with BAC) data points by the following script:

```
./run_gekko_lr_bac.sh
```

All the data will be appended to `accepted_sets_gekko.txt`.

Paper results are summarized in `Figures/accept_models_lr.csv`. 3rd, 4th, 5th lines are respectively: CoPi, GEKKO, GEKKO (with BAC).

The exact data of this experiment and the figure in the submitted version of the paper may differ depending on the changes in the dataset. However, the main outcome of both the results remains same.

Solver Runtime Overhead

1. Runtime vs pipeline length experiment (Figure 10c):

```
./time_measure_copi_pipelength.sh  
./time_measure_gekko_pipelength.sh
```

The results will be available in `scheduled_times_copi.txt`, `failed_times_copi.txt`, `accepted_time_gekko.txt` and `failed_time_gekko.txt` in space separated format. You may want to delete these files manually before running the test as results from the earlier experiments may already be there.

Paper results are summarized in `Figures/runtime_pipe_length.csv`. (1st, 2nd, 3rd, 4th, 5th line: pipeline length, CoPi times for schedulable, CoPi times for unschedulable, GEKKO times for schedulable and CoPi times unschedulable).

Exact results may vary depending on a machine's performance, but the main outcome of the results should remain the same.

2. Run CoPi experiments for Figure 9a and 9b:

```
./run_solver_exp2_copi.sh
```

3. Run GEKKO experiments for Figure 9a and 9b:

```
./run_solver_exp2_gekko.sh
```

The results will be available in `scheduled_times_copi.txt` and `failed_times_copi.txt` for CoPi, and `accepted_time_gekko.txt` and `failed_time_gekko.txt` for GEKKO, in space separated format. You may want to delete these files manually before running the test as results from the earlier experiments may already be there.

Paper results are summarized in `Figures/lbg_time.csv` and `Figures/lbg_time.csv`. In each file: (1st, 2nd line, 3rd line: LBG, times for CoPi and times for GEKKO)

CoPi Performance Insight

CoPi's AR vs. pipeline length and NLBG

```
./run_perform_insight.sh
```

The results will be written to `accepted_sets_copi.txt` file in space separated format. Paper results will be available in `Figures/increasing_task.csv`. 1st line is NLBG, 2nd line is the number of total tasks, 3rd, 4th, 5th, 6th, 7th lines are for 3, 5, 10, 15 and 20 tasks. In `accepted_sets_copi.txt`, every five entries should correspond to 1 line in `Figures/increasing_task.csv`. You may want to delete the `accepted_sets_copi.txt` file as it may contain results from previous experiments.

Exact result may vary depending on the change of datasets.

Multiprocessor Experiments

Multiprocessor experiments are run by the following command:

Run the processor based experiments with the following script:

```
./run_multiproc_2cores.sh
./run_multiproc_4cores.sh
./run_multiproc_8cores.sh
```

Results will be written to `accepted_multiproc_\<number of cores>_\<number of tasks in a pipeline>`.

Paper results are summarized in `Figures/accept_multiprocessor_2core.csv`, `Figures/accept_multiprocessor_4core.csv`, `Figures/accept_multiprocessor_8core.csv`

Lower level python script:

```
python multi_pipeline.py -p <number of pipelines> -t <number of tasks in each Pipeline> -c <number of cores>
```

Utilization Experiments and Results:

```
./run_util.sh
```

The result is written to `multi_util_result.txt`.

Migration Experiments:

```
./run_mig.sh
```

The result is written to `migrations_result.txt`.

WATERS 2015 Workshop Paper Experiments:

For the experimental results with the dataset provided in the WATERS 2015 workshop paper by Bosch (*Kramer, Simon, Dirk Ziegenbein, and Arne Hamann. "Real world automotive benchmarks for free." In 6th WATERS Workshop. 2015.*), run the following script:

```
./run_waters.sh
```

The results of the CoPi runtimes will be in `accepted_waters_copi.txt` file. The results of the derived E2E delay for schedulable pipelines will be populated in `accepted_waters_avge2e.txt` file. Match these numbers with the files provided in `Figures/waters_ar.csv` and `Figures/waters_avge2e.csv`. The exact result may vary a bit for the runtimes experiment depending on the machine where the code is being run. However, the overall trend should still match.