



Running COPASI biochemical simulations with HTCondor

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http://www.comp-sys-bio.org http://copasi.org



- Biochemical system simulator, directed at biologists/chemists (ie little computing or maths expertise)
- GUI-based standalone software (QT)
- Has now a well-established worlwide user-base
- Around 100 papers published yearly that use COPASI
- Over 10,000 downloads per year



publications per year



http://copasi.org

@COPASI_software



- ODE, SDE, or SSA
- Discrete events
- Steady states
- Stability analysis
- Parameter scans
- Sensitivity analysis
- Metabolic Control Analysis
- Time scale separation analysis

- Optimisation,
- Parameter estimation
- Cross-sections
- Lyapunov exponents
- Network diagrams
- 2D and surface plots
- GUI and command line versions
- Reads/writes SBML



http://copasi.org

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University of Manchester FEPS HTCondor pool

- Setup by Ian Cottam around 2009
- 3000 CPU cores mostly from student terminal room workstations
- During daytime about 100 machines available
- Overnight, weekends and holidays, full power of 3000 CPUs

Using COPASI with HTCondor

- 1) Edit model in COPASI
- 2) Set appropriate report format
- 3) Mark job for execution
- 4) Prepare HTCondor job specification file

- 5) Prepare output directories
- 6) Submit to HTCondor
- 7) Edit output to match required order
- 8) Create summary plots

File: stochastic_test_1.job

```
executable = /home/ed/software/copasi/bin/CopasiSE
universe = vanilla
arguments = --nologo --home . stochastic_test_batch_mode.cps
transfer_input_files = stochastic_test_batch_mode.cps
log = stochastic_test.log
error = stochastic_test.err
output = stochastic_test.out
Requirements = (OpSys == "LINUX" && Arch == "X86_64" && Memory > 0 )
should_transfer_files = YES
when_to_transfer_output = ON_EXIT
queue
```

User-friendly use of COPASI with HTCondor ?

- Yes!
- Condor-COPASI is a package that will do that
- Needs to be installed in a Condor pool
- Registered users can then use it easily
- Condor-COPASI written in python, uses a web server
- Manages all Condor issues and emails user when job is complete
- Makes summary reports and plots automatically

github.com/copasi/condor-copasi

Kent E, Hoops S, Mendes P (2012) Condor-COPASI: High-Throughput Computing for Biochemical Networks. *BMC Systems Biology* **6**:91



Ed Kent

Kent *et al. BMC Systems Biology* 2012, **6**:91 http://www.biomedcentral.com/1752-0509/6/91



SOFTWARE

Open Access

Condor-COPASI: high-throughput computing for biochemical networks

Edward Kent¹, Stefan Hoops² and Pedro Mendes^{2,3*}



Condor-COPASI Server



Condor-COPASI

- Registered users (email needed)
- COPASI files with required tasks are uploaded to web server
- (Optionally) Condor-COPASI measures speed of each subtask
- Condor-COPASI breaks tasks in several subtasks, minimizing overhead costs
- Jobs are submitted to HTCondor
- Return results are collated in correct order
- Further processing of some results
- All results zipped
- User alerted by email that results are ready

Condor-COPASI Tasks

- **Stochastic simulation repeats**: same simulation repeated to draw time-dependent distributions
- **Parameter scans**: parameters varies and simulated repeated
- Sensitivity optimization: finding maximum and minimum values of local sensitivities in wide parameter space
- **Parameter estimation repeat**: repeats same (stochastic) optimization algorithm
- Optimization repeat: repeats same (stochastic) optimization algorithm
- **Parameter estimation with all algorithms**: runs the same problem with all algorithms available

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🗍 Condor-COPASI Web Frontend - ...

Condor-COPASI

Home >> Tasks >> Parameter Estimation Repeat >>

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Parameter Estimation Repeat

Tasks

Home

Sensitivity
 Optimization

My Account

- Stochastic Simulation
- Scan in Parallel
- Optimization Repeat
- Parameter
 Estimation
 Repeat
- Raw Mode
- Optimization Repeat (Different Algorithms)
- Usage Statistics
- Help

Select the COPASI model to submit. Before submitting, ensure the model has been correctly configured:

- Parameter Estimation task
 - At least one variable parameter must have been added to the Parameter Estimation task
 - · Any experimental data files must have been selected and configured
 - Note, all experimental data files must be placed in the same directory as the model . CPS file prior to setting up the Parameter Estimation task

Condor-COPASI will automatically generate an appropriate report; no report needs to be set for the Parameter Estimation task.

Please note - it is very important that the COPASI file is saved using a supported version of COPASI. At present, only Build 33 (version 4.6.33), Build 34 (version 4.7.34), and Build 35 (version 4.8.35) are supported.

Model file:	date2/TranslationModel2.cps Browse	
Job Name:	For your reference, enter a name for this job	
Rank:	0 Use rank from last job. If you are unsure how to use rank, then do not change from the default value	
Repeats:	600 The number of repeats to perform	
Parameter estimation data:	lation/Candidate2/data-t.zip Browse Select either a single data file, or if more than one data file is required, upload a .zip file containing multiple data files	
Use a custom report:	Select this to use a custom report instead of the automatically generated one. If you select this, Condor- COPASI may not be able to process the output data, and the job will fail. However, you will still be able download the upprocessed results for manual processing. For output processing to work, you must create a	



) c	ondor-COPASI Web Frontend - Stochastic Simulation - Mozilla Firefox 💿 💿 🖉	9	– c (Particle N	
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Condor-COPASI Web Front	end - Stochas 💥 🕂			
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me >> Tasks >> Stoc	welcome mendes Administration Logout	4000		
Home		3000	1	
My Account	Stochastic Simulation			
Tasks		2000		
 Sensitivity Optimization 	Select the COPASI model to submit. Before submitting, ensure the model has been correctly configured:			
 Stochastic Simulation 	Time Course task: A The Time Course should be set up as if a single rup were to take	N The	MUSA	
 Scan in Parallel 	place on the local machine	of		
 Optimization Repeat 	 An appropriate stochastic method must be selected 			
 Parameter Estimation Repeat 	Condor-COPASI will automatically generate an appropriate report; no report needs to be set for the Time Course task.	-1000	Ч.	
 Optimization Repeat (Different Algorithms) 	Please note - it is very important that the COPASI file is saved using a supported version of COPASI. At present, only Build 33 (version 4.6.33) and Build 34 (version 4.6.34) are supported.	-2000 0 5 10 15 7 Time	1 1 1 25 30	
Help	Compare Global Sensitivity Job Output			
	Model file: Browse		*• h	
	Job Name: For your reference, enter a name for this job	► 0.000 ► 7.000		
	Select this to skip the automatic load balancing step, and Skip load make the run time of each parallel inb as short as possible	6,000-		
	balancing step: Use with caution! This has the potential to overload the Condor system with huge numbers of parallel jobs. Not applicable for some job types - see documentation	(pyruvate decarbo	ixylase (PDC1)).Kpyr_max	
	Repeats: The number of repeats to perform	2,000		
	Submit	1,000-		

Load-balancing

- How many subtasks should be submitted to each node?
- If too few, lots of jobs with high overhead
- If too many, very few jobs and little benefit from parallelization
- Ideally jobs should execute for ~20min



Figure 4 Tuning the load balancing algorithm.

Work in progress: Cloud-COPASI



- Updated web front-end (updated Django and Python 3)
- Uses Bosco to support different task managers
- Allow execution on AWS
- Add new use cases



Hasan Baig

1D likelihood profiles for parameter identifiability analysis

$$LP(p_i) = min_{p_{j\neq i}}(SSR(p_j))$$

- Re-optimizing all parameters $p_{j\neq i}$ for fixed values of p_i around the solution
- Confidence intervals for each parameter can be estimated based on likelihood contours (C_{LC}) or on likelihood ratios (C_{LR}) , e.g.:

$$C_{LR} = \{ p: SSR(p) \leq SSR(\hat{p}) e^{\chi_{\alpha}^2/n} \}$$

Example: 1D likelihood profiles to assess parameter identifiability



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COPASI Team



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