

MODREV - Model Revision tool for Boolean logical models of biological regulatory networks

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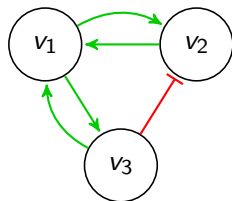
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Regulatory Networks

- Biological processes arise at the cellular level, governed by complex regulatory networks
- Regulatory network
 - ▶ Collection of molecular compounds (e.g. proteins, genes)
 - ▶ Compounds interact with each other
- Computational modelling allows
 - ▶ Functional understanding of the network
 - ▶ Test hypotheses
 - ▶ Identify predictions *in silico*
 - ▶ ...

Boolean Logical Model

- Different formalisms can be used [KS08]
 - ▶ We consider the Boolean logical formalism [Tho73].
- Compounds represented by a Boolean variable
 - ▶ active/inactive
- Interactions defined as positive (activation) or negative (inhibition)
- Regulations defined as Boolean functions



$$f_{v_1} = v_2 \wedge v_3$$

$$f_{v_2} = v_1 \vee \neg v_3$$

$$f_{v_3} = v_1$$

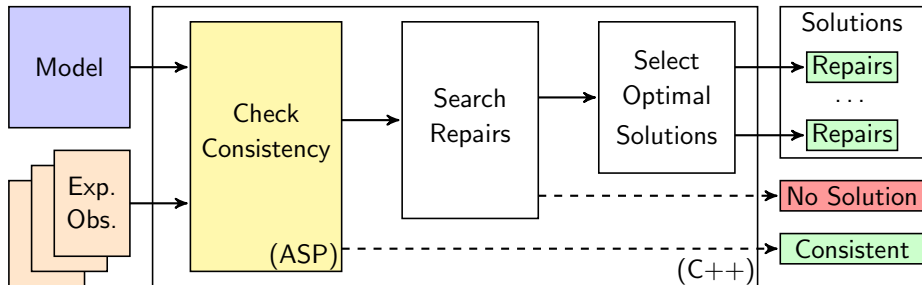
- As new experimental data becomes available, models may become **inconsistent**
 - ▶ Models may not be able to reproduce the new information
 - ▶ Models need to be **revised**
- Model Revision is mainly a manual task
 - ▶ Performed by a modeler
 - ▶ Prone to error
- How can we repair an inconsistent model?
 - ▶ Change a regulatory function?
 - ▶ Change the type of interaction?
 - ▶ Add or remove interactions?

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- MODREV is a freely available model revision tool
 - ▶ <https://filipegouveia.github.io/ModelRevisionASP/>
- Confronts a Boolean Logical Model with experimental observations
 - ▶ Stable state observations
 - ▶ Time-series observations
 - ★ Synchronous
 - ★ Asynchronous
- MODREV implements the following methods
 - ▶ Consistency check and reasons of inconsistency [GLM18]
 - ▶ Revision under stable state observations [GLM19]
 - ▶ Search for function repairs [GLM20a]
 - ▶ Revision under Time-series observations [GLM20b]

MODREV architecture



Repair Operations:

- Change regulatory Function
- Change interaction type
- Remove interaction
- Add interaction

Repair Operations:

- Change regulatory Function
- Change interaction type
- Remove interaction
- Add interaction

Optimization Criteria:

- 1 Minimize interaction addition/removal
- 2 Minimize interaction type changes
- 3 Minimize Boolean function changes

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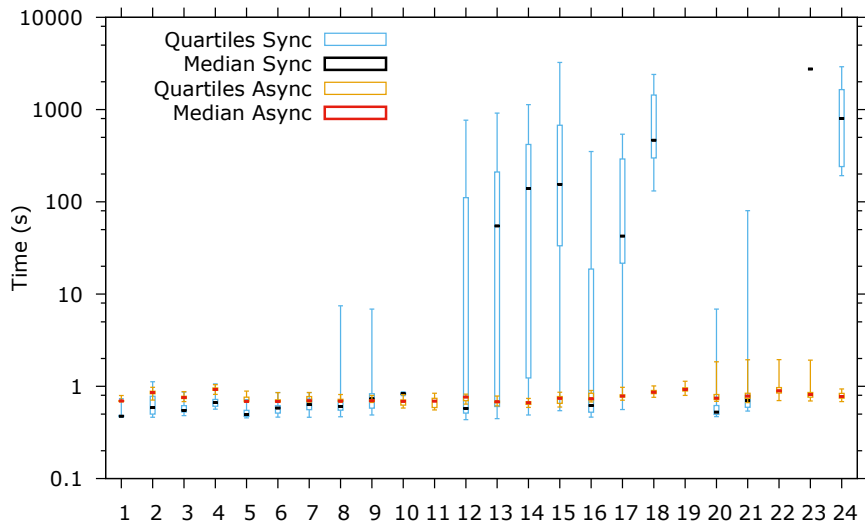
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- Segment Polarity (SP) network [SCT02]

Conf.	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24
F	5	25	50	100	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	25	50	100	5	10
E	0	0	0	0	5	10	15	20	25	50	75	0	0	0	0	0	0	0	0	5	25	50	25	10
R	0	0	0	0	0	0	0	0	0	0	0	1	5	10	15	0	0	0	0	0	0	0	5	5
A	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	1	5	10	15	0	0	0	5	5

- Random changes were made according to probabilistic parameters
 - ▶ $F\%$: Change a **F**unction
 - ▶ $E\%$: Flip the sign of an **E**dge
 - ▶ $R\%$: **R**emove an existing edge
 - ▶ $A\%$: **A**dd a missing edge
- 100 corrupted models for each of the 24 configurations
- 5 time-series observations with 20 time-steps

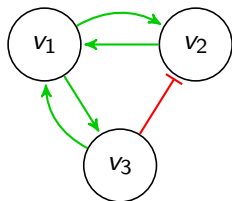
Evaluation



- MODREV repaired the model smaller # operations
- Models repaired mostly under 60 seconds
- Changing the topology of the network has the greatest impact
- Better performance under the asynchronous update scheme
 - ▶ Only on regulatory function is updated at each time step

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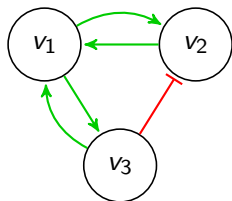


`vertex(v1).`
`vertex(v2).`
`vertex(v3).`

$$f_{v_1} = v_2 \wedge v_3$$

$$f_{v_2} = v_1 \vee \neg v_3$$

$$f_{v_3} = v_1$$



`vertex(v1).`

`vertex(v2).`

`vertex(v3).`

`edge(v1,v2,1).`

`edge(v1,v3,1).`

`edge(v2,v1,1).`

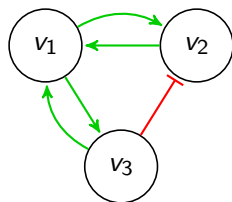
`edge(v3,v1,1).`

`edge(v3,v2,0).`

$$f_{v_1} = v_2 \wedge v_3$$

$$f_{v_2} = v_1 \vee \neg v_3$$

$$f_{v_3} = v_1$$



$$f_{v_1} = v_2 \wedge v_3$$

$$f_{v_2} = v_1 \vee \neg v_3$$

$$f_{v_3} = v_1$$

```
vertex(v1).
```

```
vertex(v2).
```

```
vertex(v3).
```

```
edge(v1,v2,1).
```

```
edge(v1,v3,1).
```

```
edge(v2,v1,1).
```

```
edge(v3,v1,1).
```

```
edge(v3,v2,0).
```

```
functionOr(v1,1..1).
```

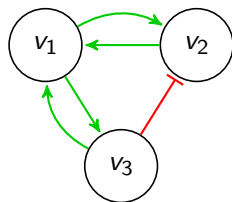
```
functionAnd(v1,1,v2). functionAnd(v1,1,v3).
```

```
functionOr(v2,1..2).
```

```
functionAnd(v2,1,v1). functionAnd(v2,2,v3).
```

```
functionOr(v3,1..1).
```

```
functionAnd(v3,1,v1).
```



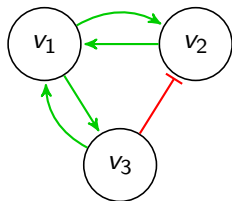
Stable State observation

```
exp(p1).  
obs_vlabel(p1,v1,0).  
obs_vlabel(p1,v2,0).  
obs_vlabel(p1,v3,1).
```

$$f_{v_1} = v_2 \wedge v_3$$

$$f_{v_2} = v_1 \vee \neg v_3$$

$$f_{v_3} = v_1$$



$$f_{v_1} = v_2 \wedge v_3$$

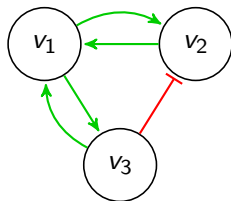
$$f_{v_2} = v_1 \vee \neg v_3$$

$$f_{v_3} = v_1$$

Stable State observation

```
exp(p1).  
obs_vlabel(p1,v1,0).  
obs_vlabel(p1,v2,0).  
obs_vlabel(p1,v3,1).
```

```
$ ./modrev -m model.lp -obs obsSS.lp -ss
```



$$f_{v_1} = v_2 \wedge v_3$$

$$f_{v_2} = v_1 \vee \neg v_3$$

$$f_{v_3} = v_1$$

Stable State observation

```
exp(p1).  
obs_vlabel(p1,v1,0).  
obs_vlabel(p1,v2,0).  
obs_vlabel(p1,v3,1).
```

```
$ ./modrev -m model.lp -obs obsSS.lp -ss
```

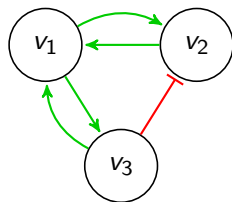
```
### Found solution with 1 repair operation.
```

```
Inconsistent node v3.
```

```
Repair #1:
```

```
Flip sign of edge (v1,v3).
```

Time-series observation



		Time		
		0	1	2
Node	v1	0	1	0
	v2	0	0	0
	v3	1	0	0

$$f_{v_1} = v_2 \wedge v_3$$

$$f_{v_2} = v_1 \vee \neg v_3$$

$$f_{v_3} = v_1$$

```
#const t = 2.
```

```
exp(p2).
```

```
obs_vlabel(p2,0,v1,0). obs_vlabel(p2,0,v2,0).
```

```
obs_vlabel(p2,0,v3,1).
```

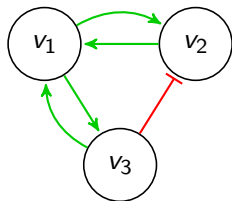
```
obs_vlabel(p2,1,v1,1). obs_vlabel(p2,1,v2,0).
```

```
obs_vlabel(p2,1,v3,0).
```

```
obs_vlabel(p2,2,v1,0). obs_vlabel(p2,2,v2,0).
```

```
obs_vlabel(p2,2,v3,0).
```

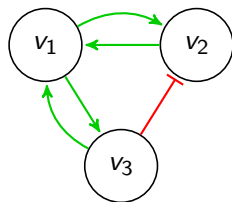
```
$ ./modrev -m model.lp -obs obsTS01.lp -up s
```



$$f_{v_1} = v_2 \wedge v_3$$

$$f_{v_2} = v_1 \vee \neg v_3$$

$$f_{v_3} = v_1$$



$$f_{v_1} = v_2 \wedge v_3$$

$$f_{v_2} = v_1 \vee \neg v_3$$

$$f_{v_3} = v_1$$

```
$ ./modrev -m model.lp -obs obsTS01.lp -up s
```

Found solution with 5 repair operations.

Inconsistent node v1.

Repair #1:

Change function of v1 to $(v_2) \vee (v_3)$

Inconsistent node v2.

Repair #1:

Change function of v2 to $(v_1 \wedge v_3)$

Flip sign of edge (v_1, v_2) .

Repair #2:

Change function of v2 to $(v_1 \wedge v_3)$

Flip sign of edge (v_3, v_2) .

Inconsistent node v3.

Repair #1:

Change function of v3 to $(v_1 \wedge v_2)$

Add edge (v_2, v_3) with sign 1.

Repair #2:

Change function of v3 to $(v_1 \wedge v_3)$

Add edge (v_3, v_3) with sign 1.

Thank you!

MODREV <https://filipegouveia.github.io/ModelRevisionASP/>

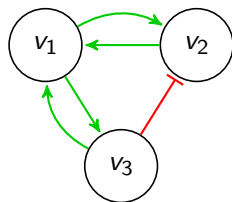
Acknowledgements:

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para a Ciência
e a Tecnologia



References

- [GLM18] Filipe Gouveia, Inês Lynce, and Pedro T Monteiro. "Model Revision of Logical Regulatory Networks Using Logic-Based Tools". In: *ICLP 2018 (Technical Communications)*. Schloss Dagstuhl-Leibniz-Zentrum fuer Informatik. 2018.
- [GLM19] Filipe Gouveia, Inês Lynce, and Pedro T Monteiro. "Model Revision of Boolean Regulatory Networks at Stable State". In: *International Symposium on Bioinformatics Research and Applications*. Ed. by Zhipeng Cai, Pavel Skums, and Min Li. Springer International Publishing. 2019, pp. 100–112.
- [GLM20a] Filipe Gouveia, Inês Lynce, and Pedro T. Monteiro. "Revision of Boolean Models of Regulatory Networks Using Stable State Observations". In: *Journal of Computational Biology* 27.2 (2020), pp. 144–155.
- [GLM20b] Filipe Gouveia, Ines Lynce, and Pedro Tiago Monteiro. "Semi-automatic model revision of Boolean regulatory networks: confronting time-series observations with (a)synchronous dynamics". In: *bioRxiv preprint doi:10.1101/2020.05.10.086900* (2020).
- [KS08] Guy Karlebach and Ron Shamir. "Modelling and analysis of gene regulatory networks". In: *Nature Reviews Molecular Cell Biology* 9.10 (2008), p. 770.
- [SCT02] Lucas Sánchez, Claudine Chaouiya, and Denis Thieffry. "Segmenting the fly embryo: logical analysis of the role of the Segment Polarity cross-regulatory module". In: *Int. J. Dev. Biol.* 52.8 (2002), pp. 1059–1075.
- [Tho73] René Thomas. "Boolean formalization of genetic control circuits". In: *J. Theor. Biol.* 42.3 (1973), pp. 563–585.



$$f_{v_1} = v_2 \wedge v_3$$

$$f_{v_2} = v_1 \vee \neg v_3$$

$$f_{v_3} = v_1$$

Incomplete time-series observation

		Time		
		0	1	2
Node	v ₁	0		1
	v ₂	1	0	0
	v ₃			

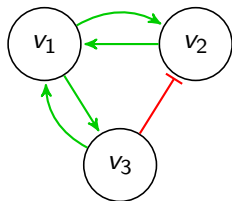
```
#const t = 2.
```

```
exp(p3).
```

```
obs_vlabel(p3,0,v1,0). obs_vlabel(p3,0,v2,1).
```

```
obs_vlabel(p3,1,v2,0).
```

```
obs_vlabel(p3,2,v1,1). obs_vlabel(p3,2,v2,0).
```

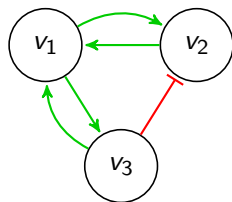


```
$ ./modrev -m model.lp -obs obsTS02.lp -up s
```

$$f_{v_1} = v_2 \wedge v_3$$

$$f_{v_2} = v_1 \vee \neg v_3$$

$$f_{v_3} = v_1$$



$$f_{v_1} = v_2 \wedge v_3$$

$$f_{v_2} = v_1 \vee \neg v_3$$

$$f_{v_3} = v_1$$

```
$ ./modrev -m model.lp -obs obsTS02.lp -up s
```

```
### Found solution with 3 repair operations.
```

```
  Inconsistent node v1.
```

```
    Repair #1:
```

```
      Change function of v1 to (v2 || (v3)
```

```
      Flip sign of edge (v2,v1).
```

```
  Inconsistent node v2.
```

```
    Repair #1:
```

```
      Change function of v2 to (v1 && v3)
```

- `vertex(V)` . : V is a node of the network
- `edge(V1,V2,S)` . : edge from $V1$ to $V2$ with sign $S \in \{0, 1\}$
- `functionOr(V,1..N)` . : regulatory function of V in DNF is represented by a disjunction of $N \in \mathbb{N}$ terms
- `functionAnd(V,T,R)` . : node R is present in the T -th term of the regulatory function of V

- `exp(E)` . : E is an experimental observation
- `obs_vlabel(E,V,S)` . : node V has an observed value of $S \in \{0, 1\}$ in experiment E (Stable State)
- `obs_vlabel(E,T,V,S)` . : in time-step T node V has an observed value of $S \in \{0, 1\}$ in experiment E (Time-series)