



# BiVeS & BudHat

## Version Control for Computational Models

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University of Rostock

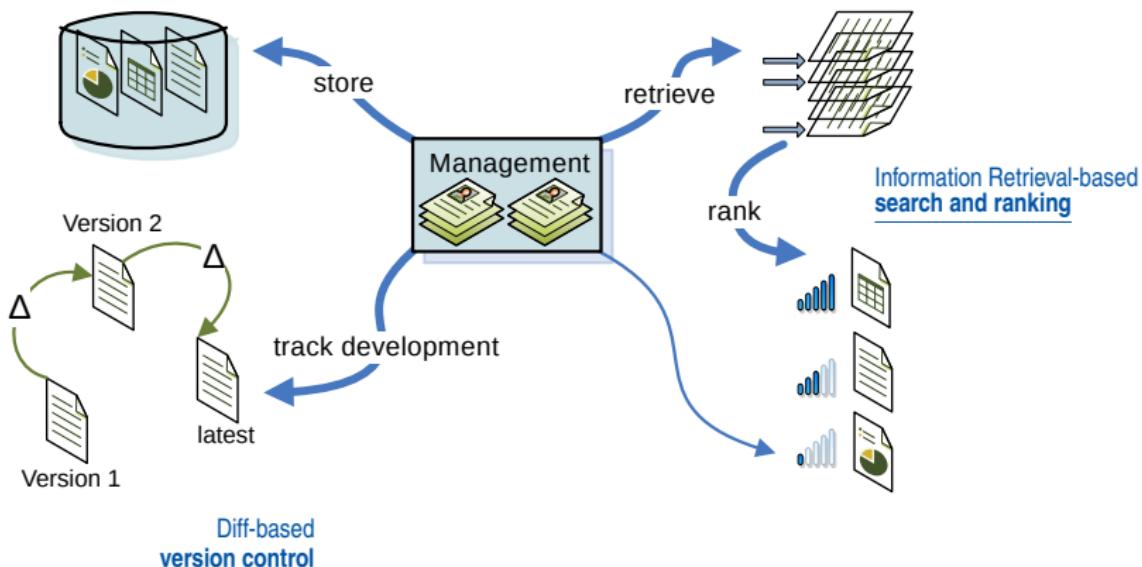
<http://sems.uni-rostock.de>

Gatersleben 2013





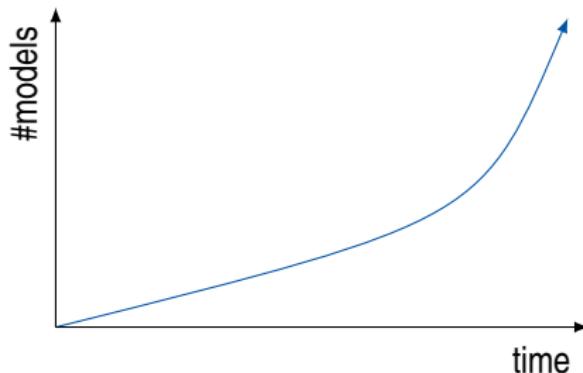
Format-independent,  
graph-based storage



<http://sems.uni-rostock.de/>

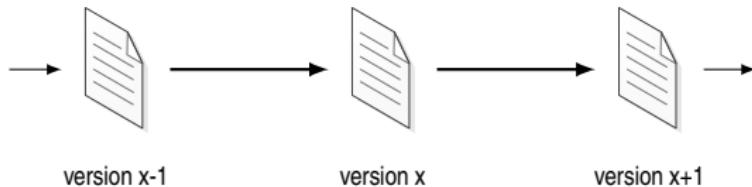


- many models
- multiple versions of each model
- several repositories



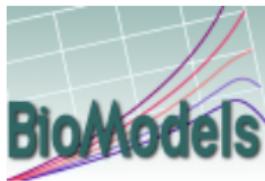


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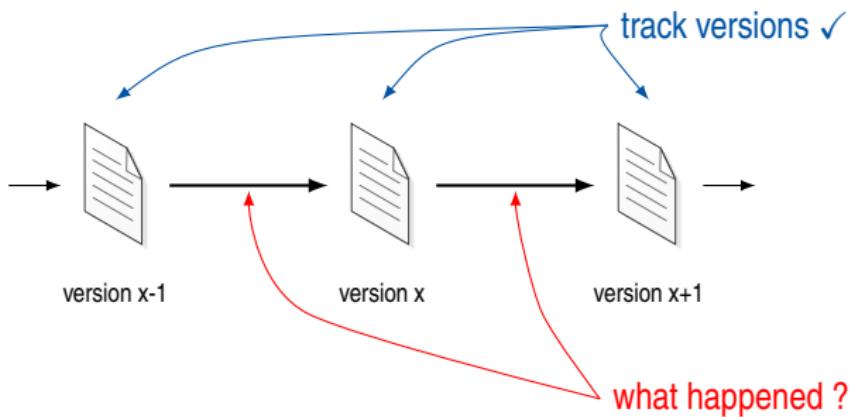


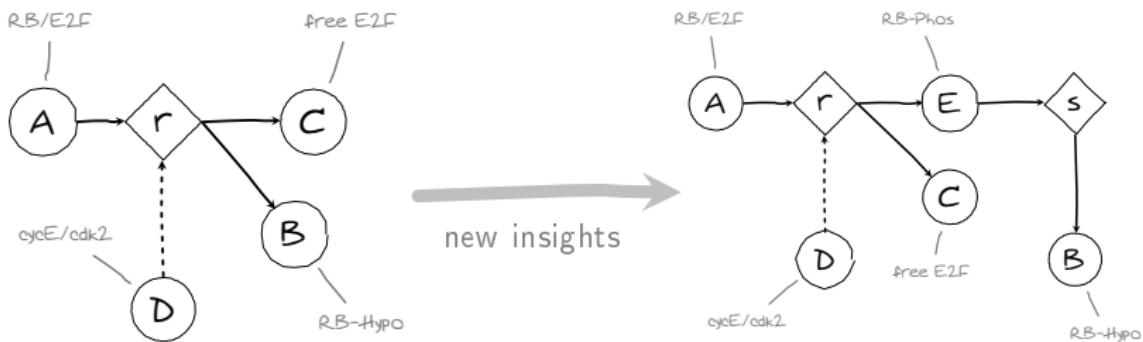
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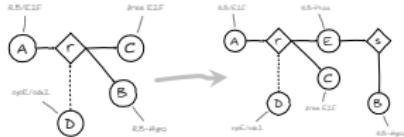


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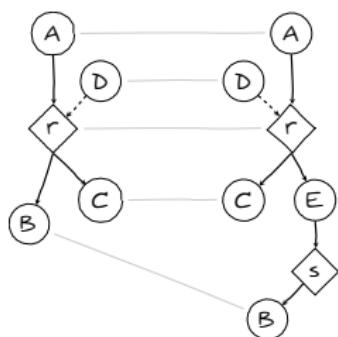


Waltemath *et al.*: Improving the reuse of computational models through version control. *Bioinformatics* (2013) 29(6): 742-728;



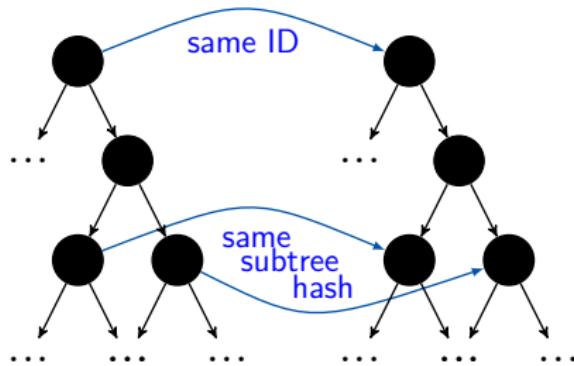
## Biochemical Model Version Control System

- compares models encoded in standadized formats (currently: SBML and CellML)
- maps hierarchically structured content



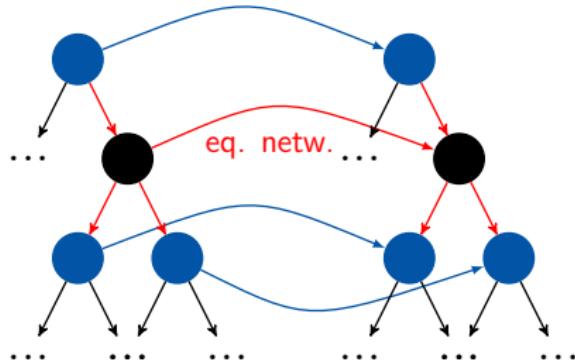


- input: 2 XML documents
- map unambiguous nodes/subtrees
- propagate mapping up/down
- exploit further knowledge of biological systems



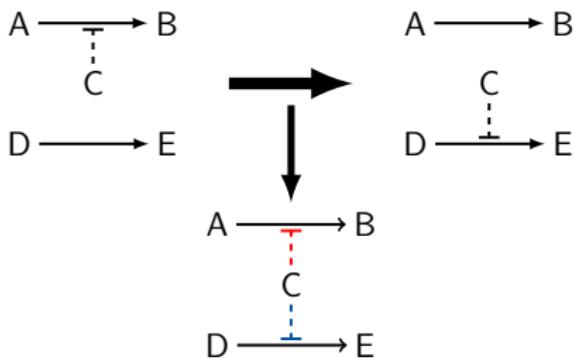


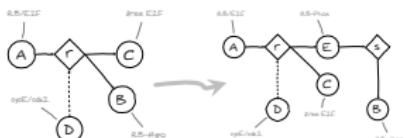
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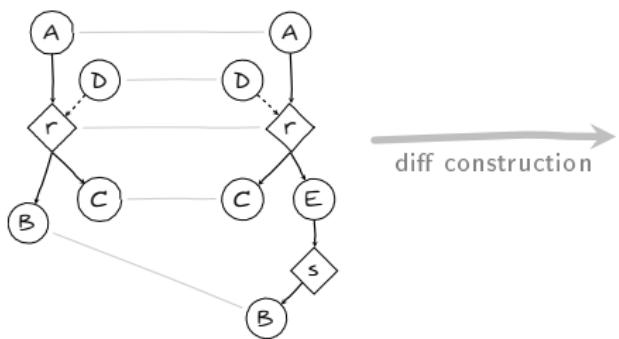
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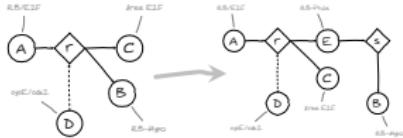


## Biochemical Model Version Control System

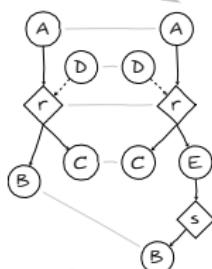
- compares models encoded in standadized formats (currently: SBML and CellML)
- maps hierarchically structured content
- constructs a diff (in XML format)
- is able to interpret this diff



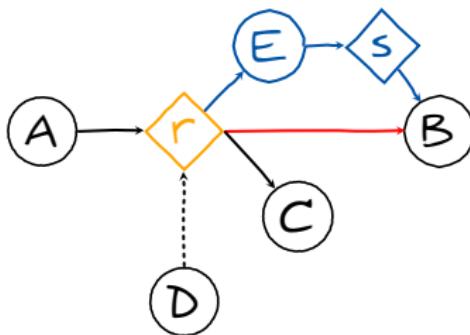
```
<XML>
  Diff
    moves
      product of r: C
    deletes
      product of r: B
    inserts
      species: E
      product of r: E
    reaction s
  </XML>
```



- calls BiVeS to construct the diff
- displays the result in various formats
  - the XML diff
  - a reaction network highlighting the changes using Cytoscape Web
  - a human readable report



```
<XML>
  Diff
    moves
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```





## The Power of BudHat!

<http://budhat.sems.uni-rostock.de>



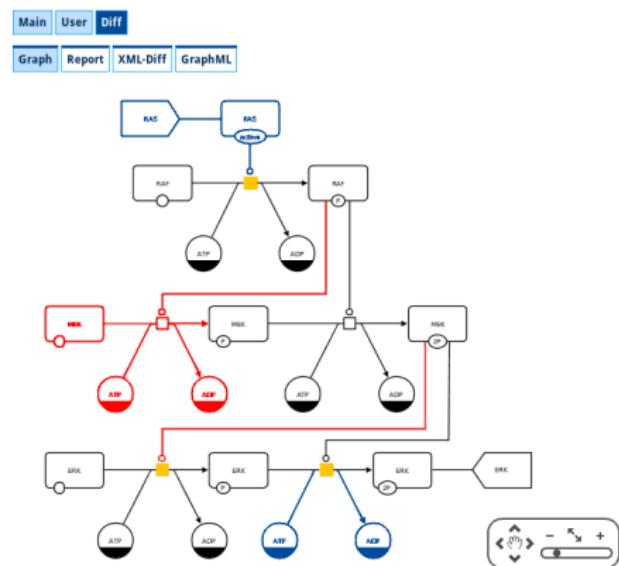
VANTED → BiVeS

**Project:BudHat**

available models

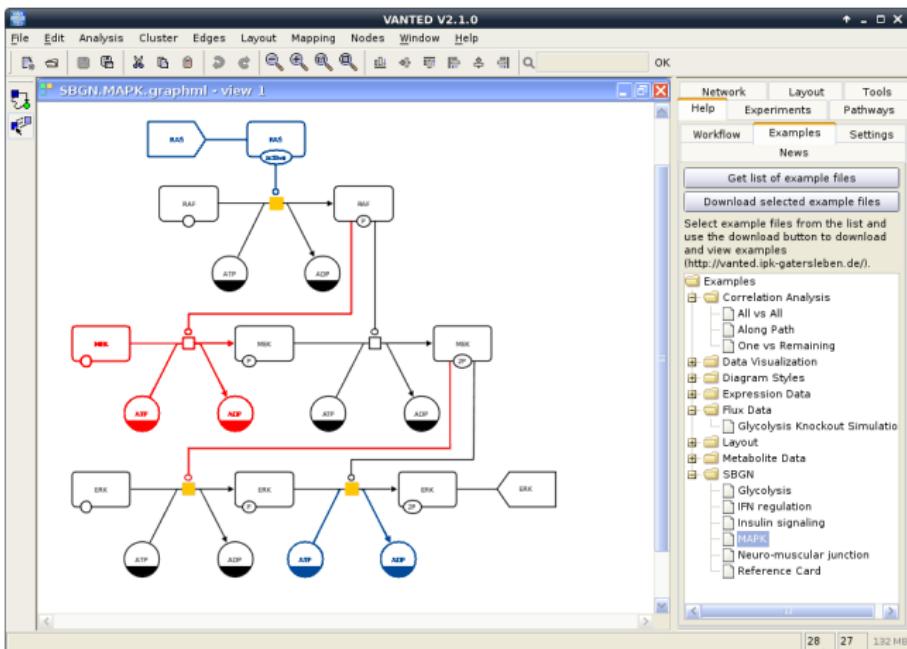
selected models

compare





VANTED ← BiVeS





VANTED ↔ BiVeS

let's collaborate!

We VANT BiVeS!



jvm

network

cmd

```
import de.unirostock.sems.bives.api.SBMLDiff;  
[...]  
SBMLDiff differ = new SBMLDiff (sbmlFileA, sbmlFileB);  
differ.mapTrees ();  
String graph = differ.getGraphML ();  
[...]
```



jvm

network

cmd

```
curl -d "file1=http://some.server/model/version1"
-d "file2=http://other.server/model/version2"
-d "get=graphml" bives.server/api
```

```
Transmission Control Protocol, Src Port: 42889 (42889), Dst Port: http (80), Seq: 1, ACK: 1, Len: 142
└ Hypertext Transfer Protocol
  └ POST /api HTTP/1.1\r\n
    └ [Expert Info (Chat/Sequence): POST /api HTTP/1.1\r\n]
      Request Method: POST
      Request URI: /api
      Request Version: HTTP/1.1
      User-Agent: curl/7.30.0\r\n
      Host: bives.server\r\n
      Accept: */*\r\n
    └ Content-Length: 92\r\n
      Content-Type: application/x-www-form-urlencoded\r\n
      \r\n
      [Full request URL: http://bives.server/api]
  └ Line-based text data: application/x-www-form-urlencoded
    file1=http://some.server/model/version1&file2=http://other.server/model/version2&get=graphml
```



jvm

network

cmd

```
java -jar BiVeS.jar path/to/version1 path/to/version2
```



```
git diff 88fealcddf b64477d742 model.file
```



- BiVeS = Difference detection for hierarchically structured content
- BudHat = Prototype to demonstrate the capabilities
- Both tools are open source
- Our goal is to extend existing models repositories with a valuable version control mechanism



That's it! Stay tuned ;-)

 @SemsProject

<http://sems.uni-rostock.de>

<http://budhat.sems.uni-rostock.de>

Questions? Suggestions? Recommendations? Drop me an email:  
[<martin.scharm@uni-rostock.de>](mailto:martin.scharm@uni-rostock.de)