



BiVeS & BudHat

Difference Detection for Computational Models

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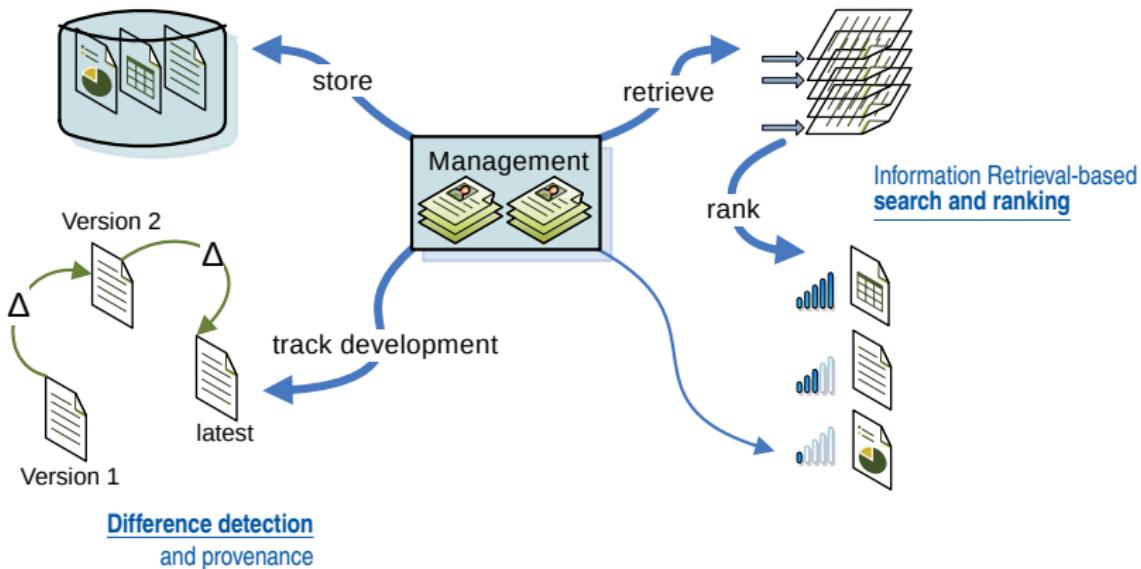
<http://sems.uni-rostock.de>



COMBINE 2013



Format-independent,
graph-based storage



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



Provenance represents the seven W's (Who, What, Where, Why, When, Which, (W)how).

– Goble 2002



Music
and Arts



Literature

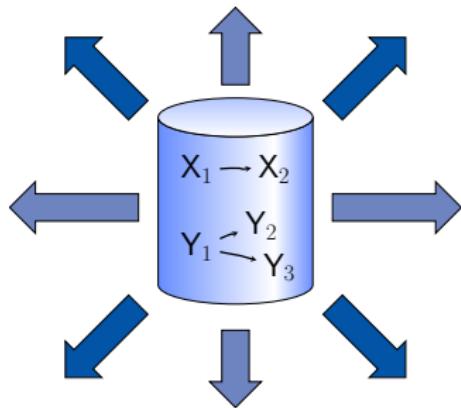
```

14 my <--> userAgent;
15 my $SEERER = " ";
16 my $GROPE = ("0", "0", "0", "0", "0", "0", "0");
17 my $GROPAIT = ("0", "0");
18 my $GROPLIT = ("0", "0", "0", "0", "0", "0", "0", "0", "0", "0", "0", "0", "0", "0", "0", "0");
19 my $GROPLIT = ("0", "0", "0", "0", "0", "0");
20
21 my $script = $a;
22 my $step = "$&#x2019;";
23
24 usage (Script) if ($step eq "?>/<");
```

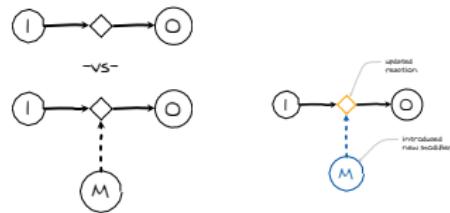
Code

REFERENCES

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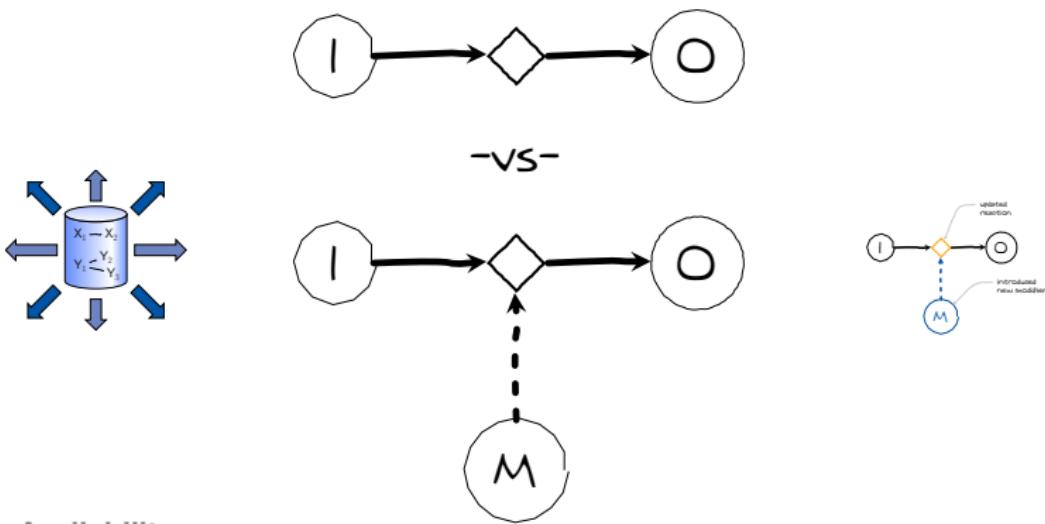


Availability & Identity



Difference
Detection

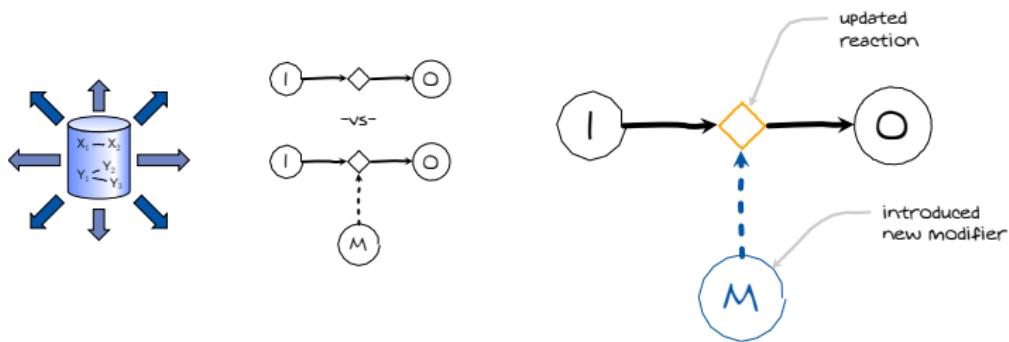
Interpretation



Availability
& Identity

Difference Detection

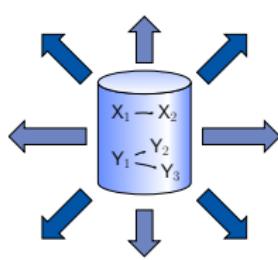
Interpretation



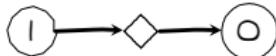
Availability
& Identity

Difference
Detection

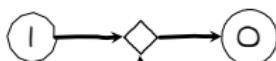
Interpretation



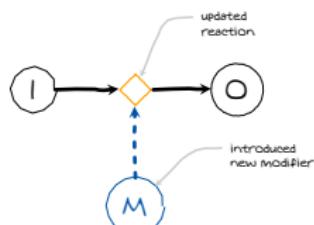
**Availability
& Identity**



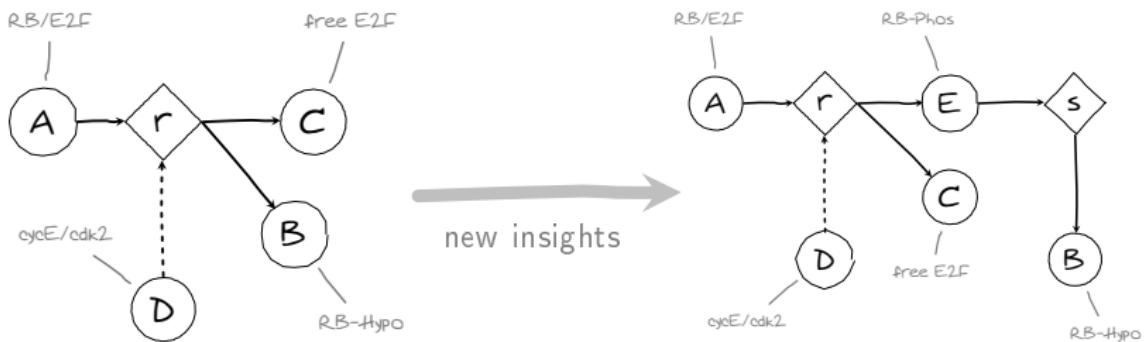
-vs-



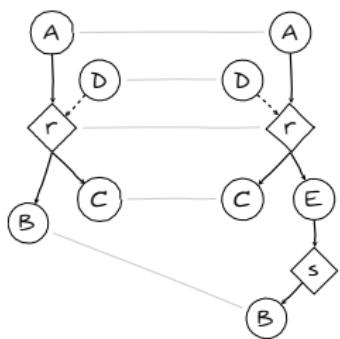
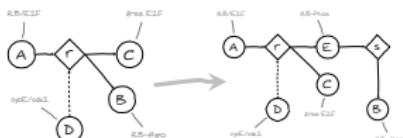
**Difference
Detection**



Interpretation



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

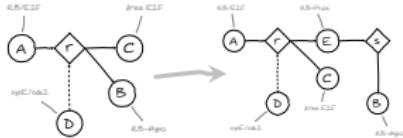


diff construction

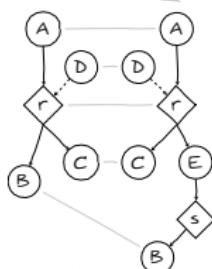
Biochemical Model Version Control System

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

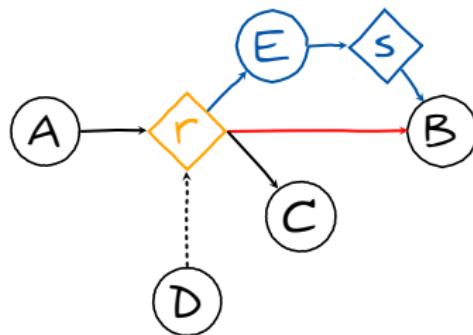
```
<XML>
  Diff
    moves
      product of r: C
    deletes
      product of r: B
    inserts
      species: E
      product of r: E
    reaction s
  </XML>
```



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



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





BudHat in action!

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



jvm

network

cmd

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



jvm

network

cmd

```
curl -d '{
  "get": [
    "documentType",
    "xmlDiff"
  ],
  "files": {
    "versionA": "http://your.db/path/to/versionA.sbml",
    "versionB": "http://your.db/path/to/versionA.sbml"
  }
}' http://bives.server.tld
```



jvm

network

cmd

```
java -jar BiVeS.jar path/to/versionA path/to/versionB
```



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



- BiVeS = Difference detection for hierarchically structured content
- BudHat = Prototypic web interface to demonstrate how BiVeS can be used to communicate the differences



That's it! Stay tuned ;-)

 @SemsProject

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

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

Questions? Suggestions? Recommendations? Drop me an email:

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