

# REX - A TOOL FOR DISCOVERING EVOLUTION TRENDS IN ONTOLOGY REGIONS

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# **ONTOLOGY EVOLUTION**

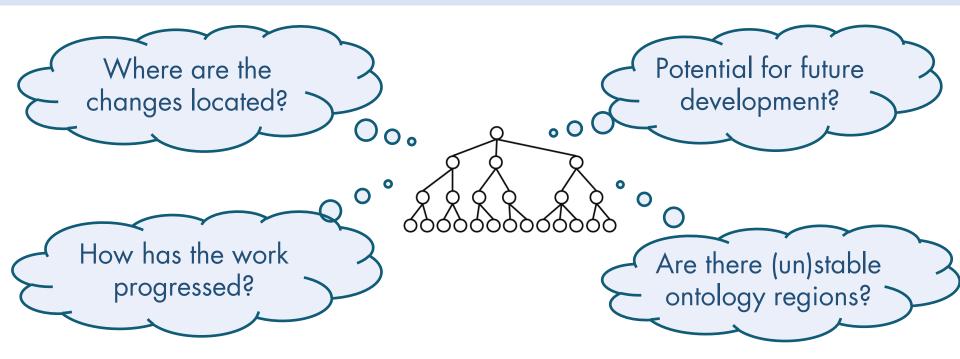
- Heavy usage of ontologies in the life sciences, e.g. annotations, semantic search, ...
- Ongoing research, new findings  $\rightarrow$  Ontology changes
- Continuous release of new versions
   + S
   \*
- Collaborative, distributed development of large ontologies

Ontology developers with different fields of expertise

Coordination by

project leader

# **ONTOLOGY EVOLUTION**



- Unstable ontology regions
  - $\rightarrow$  Many modifications  $\rightarrow$  focus of recent development
  - $\rightarrow$  Impact of changes on ontology-based algorithms or applications  $\rightarrow$  redo analyses?
- Stable ontology regions
  - $\rightarrow$  Already completed?
  - $\rightarrow$  Low interest so far  $\rightarrow$  further changes necessary?

# **CURRENT STATE & CONTRIBUTIONS**

- BioPortal, OBO Foundry , ...
  - Provide life science ontologies with versions
- Diff computation (PromptDiff, COntoDiff, ...)
- Web tools to analyze ontology evolution (GOChase, OnEX, ...)
- $\rightarrow$  No tool to analyze and compare evolution in different ontology parts

### Web application Region Evolution Explorer (REX)

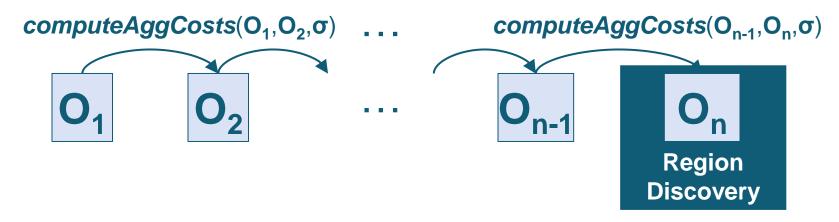
- Determination of differently changing ontology regions
- Interactive exploration of regions w.r.t. their change intensity
- Comparative trend analysis
  - $\rightarrow$  Monitor long-term evolution for regions of interest
  - $\rightarrow$  Track the work or coordinate future development

## AGENDA

- Region Discovery Method
- Trend Discovery
- REX
  - Infrastructure
  - Analysis workflows
- Conclusions and Outlook

# **REGION DISCOVERY METHOD**

• **Basic idea**: Compute change intensities for regions based on changes between succeeding ontology versions  $O_1 \dots O_n$  using change cost model  $\sigma$ 



- 1) Compute changes between succeeding versions
- 2) Propagate change costs within the is-a hierarchy
- 3) Transfer aggregated costs from the first to the last version
- 4) Compute change intensities to discover differently evolving regions

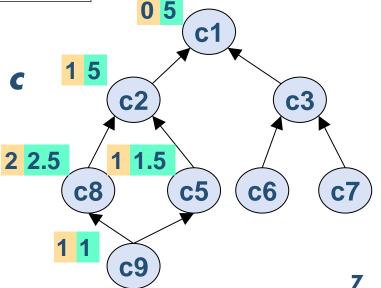
# **ONTOLOGY CHANGES AND COSTS**

	Change operation	Change costs
Concepta	addC	1
Concepts	delC	2
Relationships	addR	0.5/0.5
	delR	1.0/1.0
Attributes	addA	0.5
	delA	0.5
	chgAttValue	0.5

Cost model can be adapted!

### • Costs of an ontology concept c

- Local costs
   Ic(c)
  - Changes that directly affect *c*
- Aggregated costs *ac(c)*
  - Changes on *is-a* descendants of *c*



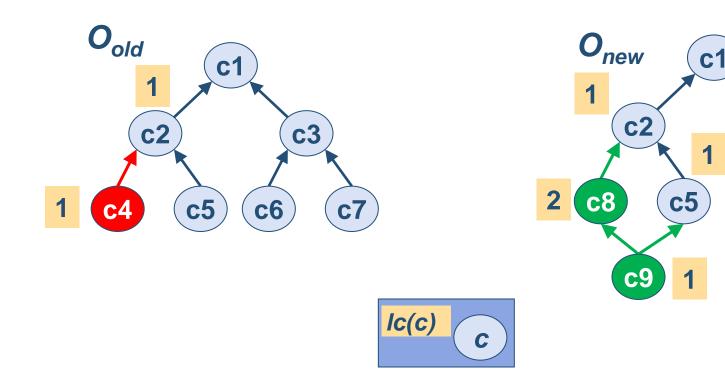
## **COMPUTE AGGREGATED COSTS FOR TWO VERSIONS**

## Algorithm: compute Aggregated Costs ( $O_{old}$ , $O_{new}$ , $\sigma$ )

- Input: Two ontology versions  $O_{old}$  and  $O_{new}$ , change costs  $\sigma$
- Output: Onew with computed aggregated costs (ac)
  - 1. Detect changes: diff (O<sub>old</sub>, O<sub>new</sub>)
  - 2. Assign local costs to  $O_{old}$  and  $O_{new}$
  - 3. Aggregate costs in  $O_{old}$  and in  $O_{new}$
  - 4. Transfer costs  $O_{old} \rightarrow O_{new}$

# **CHANGE DETECTION AND LOCAL COSTS**

- Simple diff computation: based on accession numbers used in ontology elements → addC, delC, addR, delR, ...
- Example
  - Uniform change costs of 1
  - Relationship costs to target



**c**3

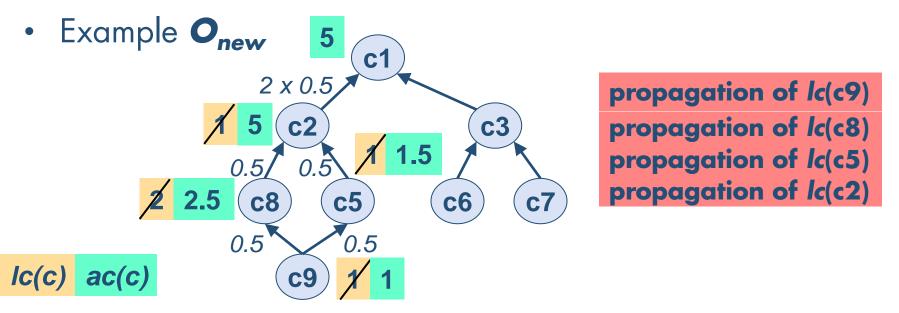
**c7** 

**c6** 

# **COST PROPAGATION AND AGGREGATION**

- Propagation of local costs Ic
- Aggregated costs *ac(c)* of concept *c*:
   Weighted sum of all children *ac*'s plus own local costs *lc(c)*

$$ac(c) = \sum_{direct \ children \ c'of \ c} \frac{ac(c')}{|parents(c')|} + lc(c)$$



# **COST PROPAGATION AND AGGREGATION**

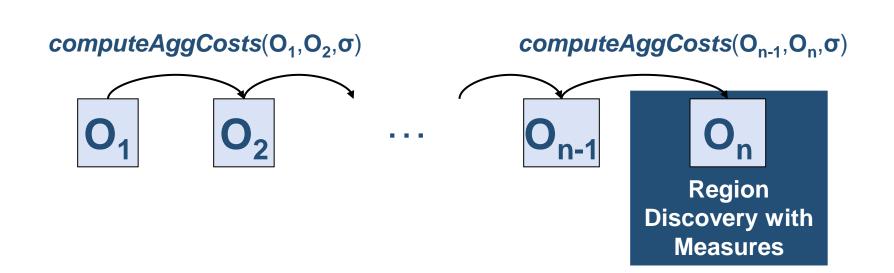
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$$ac(c) = \sum_{direct \ children \ c'of \ c} \frac{ac(c')}{|parents(c')|} + lc(c)$$

- Example Onew 5 cl
   2 × 0.5 cl
   5 c2 c3 propagation of *lc*(c9) propagation of *lc*(c8) propagation of *lc*(c5)
  - Transfer aggregated costs  $O_{old} \rightarrow O_{new}$
  - $\rightarrow$  Costs of del changes reflected in new(er) version(s)
  - Costs of equal concepts are summed up

## **DISCOVERY ALGORITHM FOR MULTIPLE VERSIONS**

- Reuse of computeAggregatedCosts algorithm
  - Successive computation and transfer of aggregated costs
  - Apply region discovery on latest version

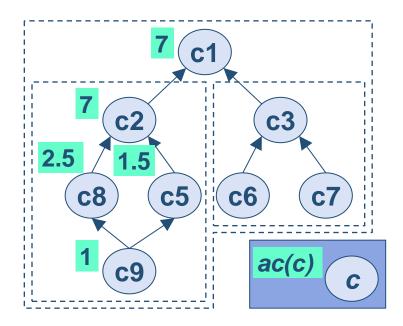


# **ONTOLOGY REGIONS AND MEASURES**

#### Ontology region OR

- Subgraph of an ontology consisting of a single root concept and all concepts in its *is-a* subgraph
- **Region measures** to quantify change intensity
  - Absolute change costs
  - Absolute / relative region size
  - Average change costs

region	abs_costs	rel_size	avg_costs
<b>c1</b>	7	8/8=1	7/8=0.875
<b>c2</b>	7	4/8=0.5	7/4=1.75
<b>c</b> 3	0	3/8=0.375	0/3=0



# **TREND DISCOVERY**

- Trend discovery based on sliding windows
- Monitor region changes over long periods of time
  - Ontology O, region of interest OR
  - Time interval (t<sub>start</sub>, t<sub>end</sub>)
  - Sliding window of size  $\boldsymbol{\omega}$
  - Step width  $\Delta$
- $\rightarrow$  Call region discovery algorithm within  $\omega$

 $\rightarrow$  Collect change intensities for region of interest over time

$$O_{1} O_{2} O_{3} O_{4} O_{5} O_{6} O_{7} O_{8} \dots O_{n-2} O_{n-1} O_{n}$$
  
$$\omega = 4, \Delta = 1$$

**t**start

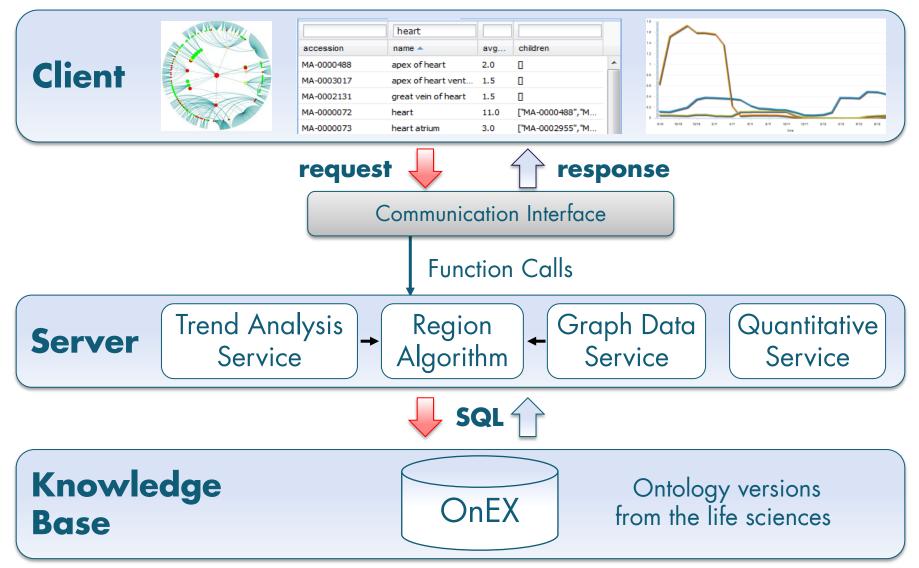
Tend

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## INFRASTRUCTURE





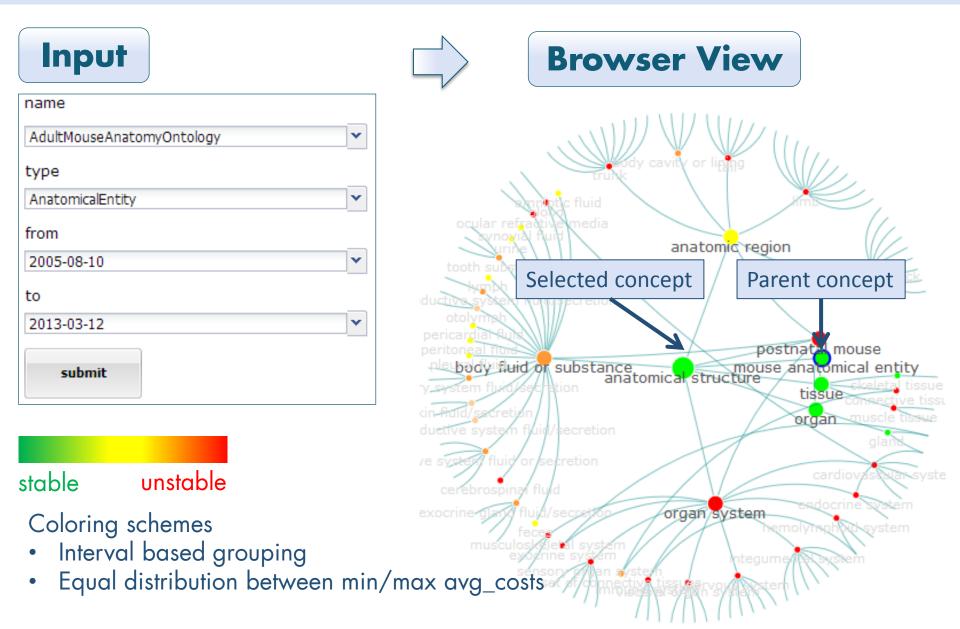
# **QUANTITATIVE CHANGE ANALYSIS**

al stati	stics							
on type	e: ®	simple	C	2 ontologi	es	2 interva	ls (	show
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					fr	om dat	е	
tology		~	r		2	007-01-01		
					u	ntil date	Э	
		*			2	012-12-01		
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Input	<b>Browser View</b>
name	
AdultMouseAnatomyOntology 🗸	
type AnatomicalEntity	()))))///(()
from	body fluid g stance
2005-08-10	Click node to navigate ontology root
to	
2013-03-12	
submit	anatomical structurese anatomical entity
	tissue
stable unstable	
sidble unsidble	organ
Coloring schemes <ul> <li>Interval based grouping</li> </ul>	
Equal distribution between min	/max avg_costs







### Input

name	
AdultMouseAnatomyOntology	~
type	
AnatomicalEntity	×
from	
2005-08-10	~
to	
2013-03-12	~
submit	

- Filter / sort by accession number, name, avg\_costs ...
- View individual change history of concepts

### **Table View**

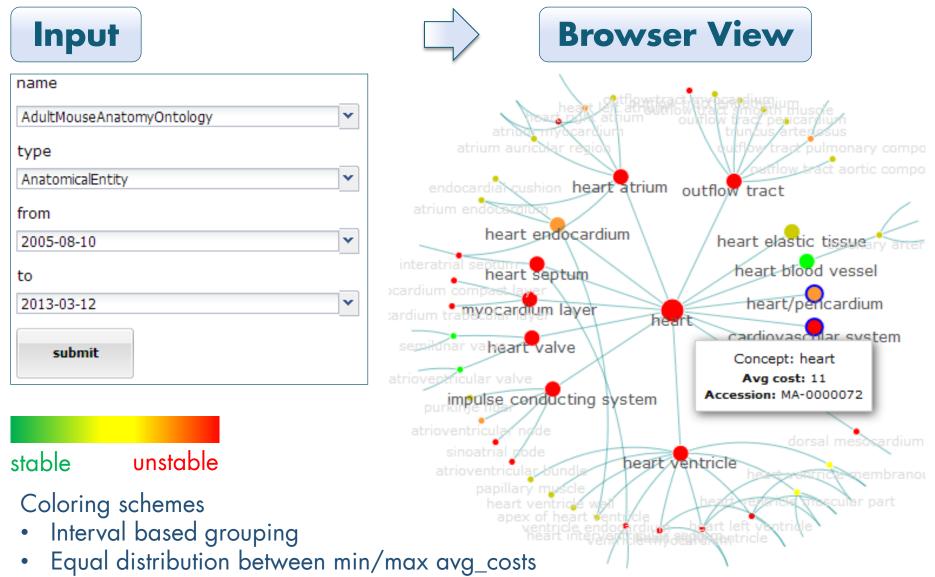
	heart		
accession	name	avgCo	children
MA-0000072	heart	11	["MA-0000488", "MA-00
MA-0000098	heart mesentery	7.5	["MA-0000482"]
MA-0000085	heart interventri	5.5	["MA-0002939", "MA-00

**Change History** 

~		/
	$\checkmark$	

Change history of concept MA-0000072							
date of cha	type of ch	type of i	changed i	new value(s)	old v		
2005-08	addition	attribute	name	heart			
2005-08	addition	relation	is_a	MA-0000557			
2005-08	addition	relation	part_of	MA-0000010			
2005-12	addition	relation	part_of	MA-0002449			
2005-12	deletion	relation	is_a		MA-0		







## **TREND ANALYSIS**

specificat	ion			immune system		
name:	GeneOntology	*	accession	name	accession	name
type:	Process	*	GO-0002376	immune system process	GO-0032502	developmental process
from:	2008-03-09	*	GO-0002520	immune system development	GO-0050896	response to stimulus
to:	2013-03-12	*	GO-0002682	regulation of immune system process	GO-0002376	immune system process
step size:	1	*	GO-0002683	negative regulation of immune system process		
window size	ze: 6	*	GO-0002684	positive regulation of immune system process		
1. <sub>submit</sub> 2	calculate trends	3. 📑 export				



# **CONCLUSIONS AND FUTURE WORK**

- Explore evolution of life science ontologies to discover (un)stable regions and trends
- Monitor development of large ontologies (developers, project coordinators)
- Support ontology-based algorithms and analysis
- Support versioning of individual ontologies (upload of own versions)
- Include user-based adaptation of cost model
- Web service interface