

Metamodel matching for automatic generation of model transformation Or other purposes

Sattose

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Context

- Increasing use of MDE approaches ;
- Many tools for MDE ;
- Many models and metamodels.

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- Many tools for MDE ;
- Many models and metamodels.

Explosion of the metamodel number

Several metamodels for a same domain

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Trace metamodels

- ETrace metamodel
- EML trace metamodel (Epsilon Merging Language)
- ATL trace metamodel
- ...

Class metamodels

- Ecore metamodel
- UML metamodel
- Kermeta metamodel
- Minjava metamodel
- ...

About metamodel compatibility

Ideally:

- Tools should allow to use all the models conform with common metamodels of a given domain
- For instance: a graphical editor for class models with UML, Kermet, or ECore

But:

- Compatibility is ensured by hand-written transformations
- tedious, error-prone, cost

Towards automatic generation of model transformations (for similar meta-model)

Metamodel alignment

- Generation of mappings between metamodel elements \longrightarrow metamodel alignment
- The alignment is used to produce the transformation code

Using schema *matching* techniques

Coming from:

- Database domain *matchings* of database schemas
- Semantic web : ontology alignment, XML schema *matching*

The proposed approach

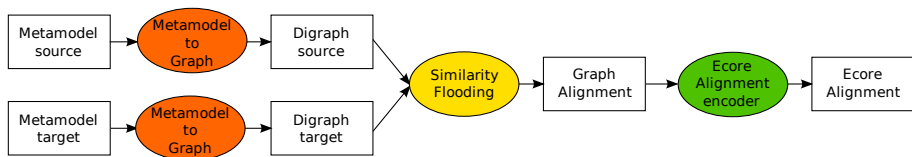
Similarity Flooding [Melnik and al., 2002] for *matching*

- Similarity flooding works on labeled directed graphs
- Similarity flooding is easily tunable

Using matching

- Testing several configurations for Similarity Flooding use
- Definition of a metamodel alignment
- Automatic construction of alignment models

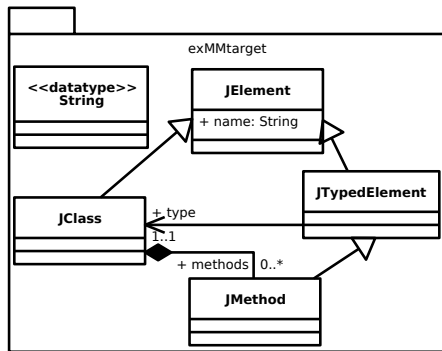
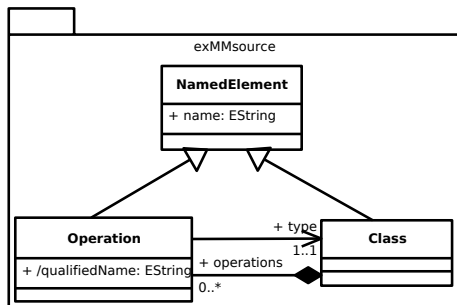
Three steps



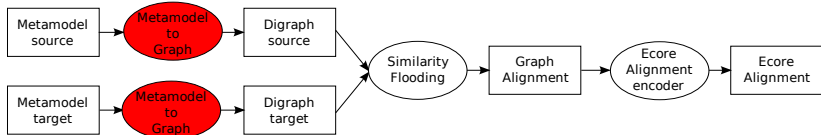
The three steps

- 1 From metamodels to graphs
- 2 Application of Similarity Flooding
- 3 Construction of an alignment metamodel using the result of Similarity Flooding

An example



1. From metamodels to graphs



Transform a metamodel into a labelled directed graph

Input

A metamodel

Output

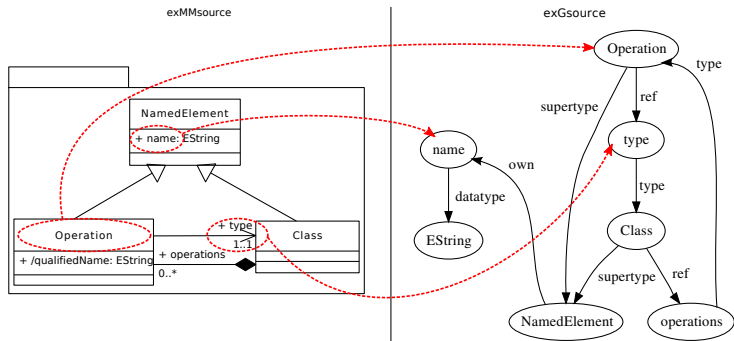
A directed labelled graph representing the model

Objective

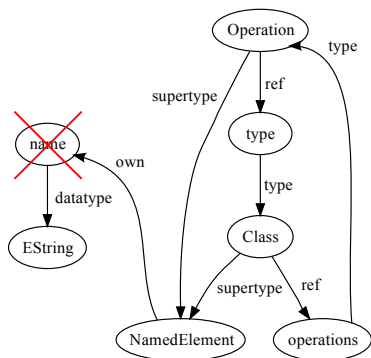
- Study the impact on Similarity Flooding of the configuration choice
- Six tested configurations
- Comparison of the results

Configuration *Minimal*

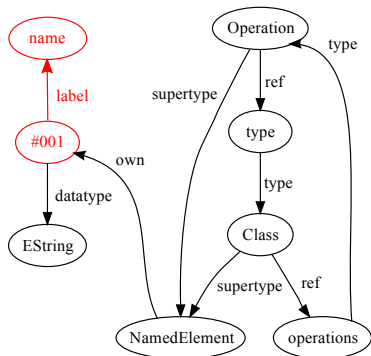
- Metamodel elements are converted into labelled nodes
- Relations are converted into labelled edges
- Derived attributes, references, operations and parameters are ignored



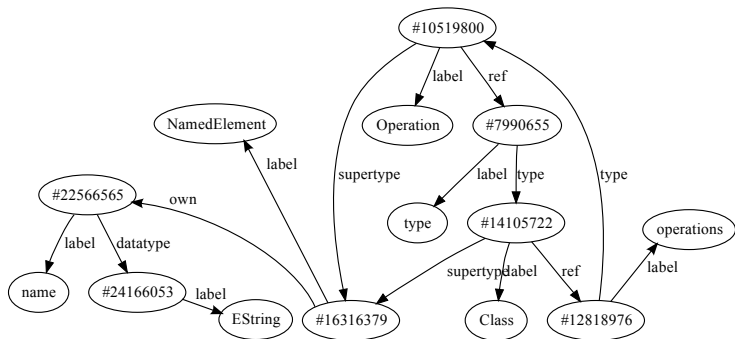
Like *Minimal*, but elements and names are separated



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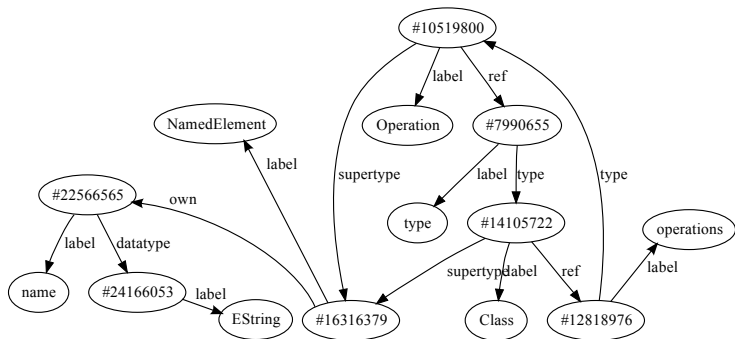


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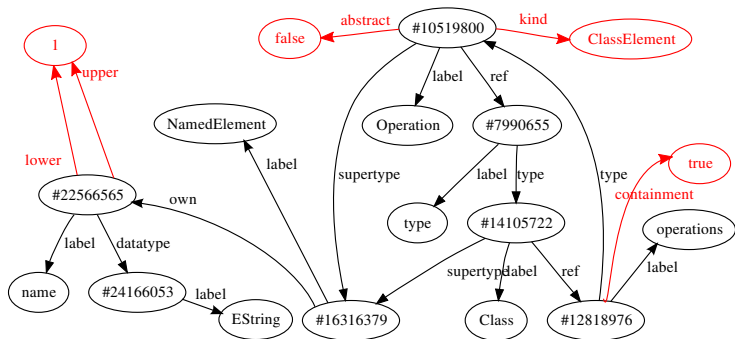
Configuration Standard

Like *Basic*, but includes metaclasses, cardinality, abstract et containment



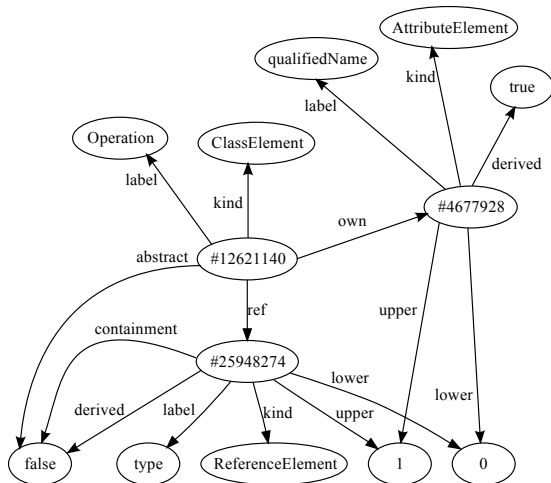
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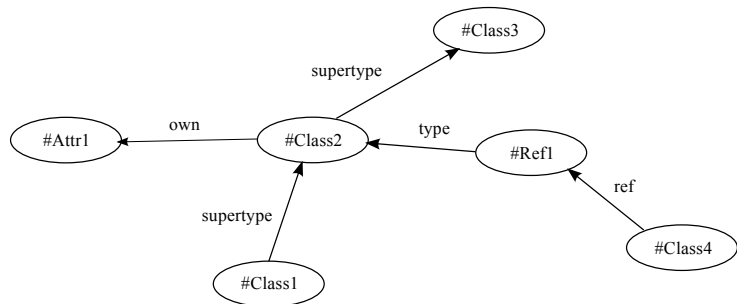
Configuration *Full*

- Based on *Standard* configuration,
- with nodes for derived attributes and references



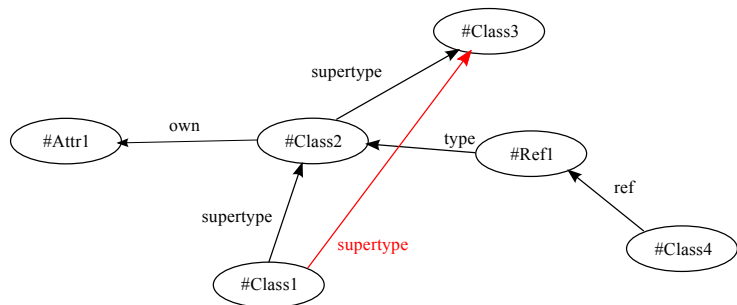
Configuration *Saturated*

- Based on *Standard*.
- *supertype* is closed by transitivity
- For a class node, we add *own* and *ref* towards inherited attributes and references
- For a reference node, we add *type* towards all superclasses



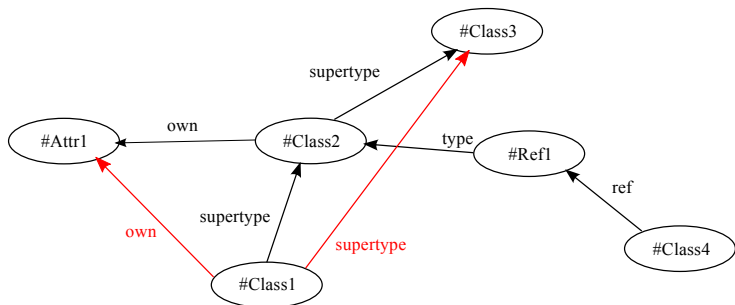
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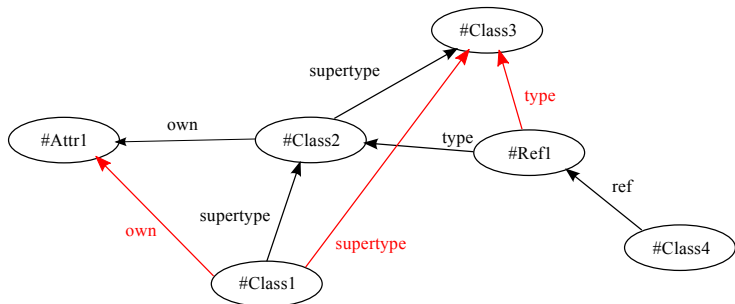
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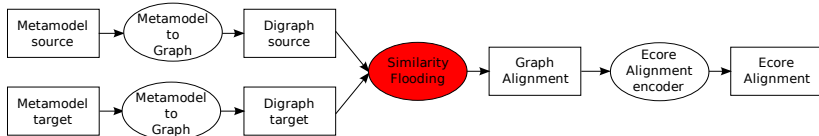
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- Based *Standard*.
- Abstract class nodes and *supertype* edges are removed,
- references typed by an abstract class are replaced by a set of references typed by concrete classes
- Links *own* and *ref* are added towards attributes and references of superclasses when a *supertype* edge is removed

2. Similarity Flooding

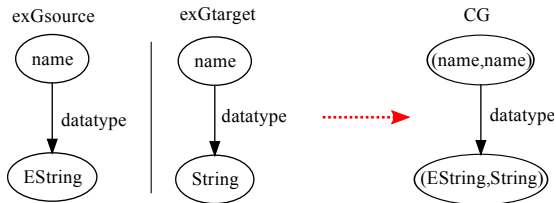


- Computes similarities between the nodes of two labelled directed graphs
- Input: two graphs G_{source} and G_{target} .
- Output: an association set between one node of G_{source} and one node of G_{target} .
- A 5-steps algorithm.
- Two nodes are similar when their neighbors are

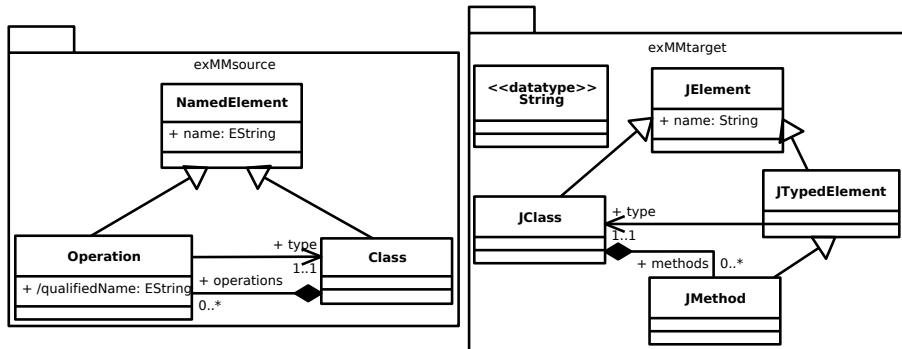
First step: The compatibility graph

Principe

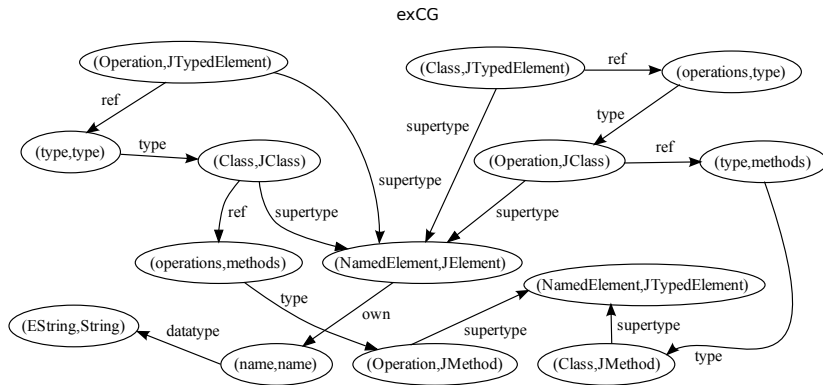
- G_{source} and G_{target} are composed of triplets (x, r, y) ,
- CG is the compatibility graph,
- $((x_1, x_2), r, (y_1, y_2)) \in CG$ iff $(x_1, r, y_1) \in G_{source}$ et $(x_2, r, y_2) \in G_{target}$



Example, with the configuration *Minimal*



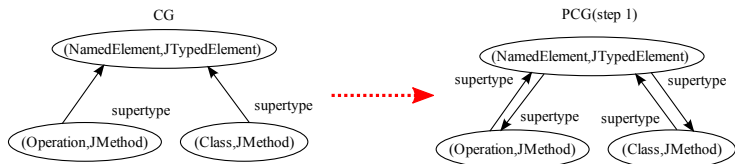
Example, with the configuration *Minimal*



Second step: propagation graph

Principle

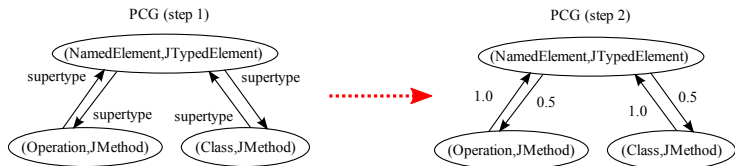
- For each (s, r, t) in the compatibility graph, we add the reverse triplet (t, r, s) .
- $O_l^n = \# \text{edges with label } l \text{ and leaving } n$.
- Triplets (s, r, t) are replaced by $(s, \frac{1}{O_s^r}, t)$



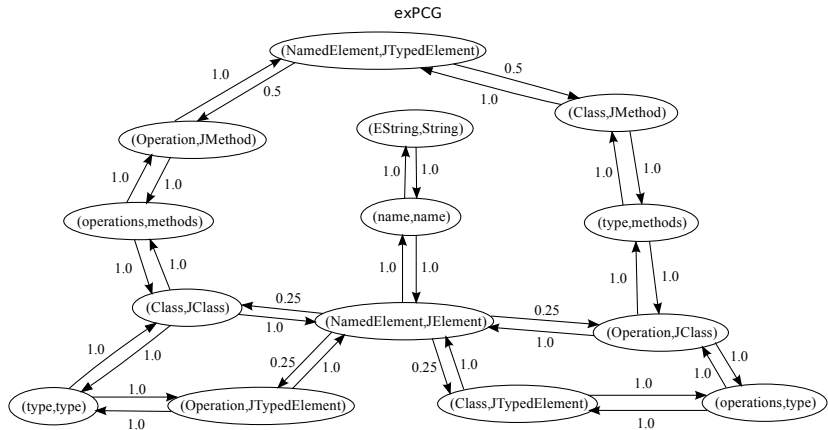
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Example



Third step: assigning initial similarity values

Principe

For a compatibility node $n = (x, y)$, initial value s_n^0 is given by:

- 0 if x or y is an identifier (label begins with #)
- $1 - lev(x, y) / \max(len(x), len(y))$ otherwise

With:

- $lev(x, y)$ Levenshtein distance [Levenshtein, 1966] between label of x and y (editing distance between two strings)
- $len(x)$ the length of x

Example

- 0 if x or y is an identifier (label begins with #)
- $1 - lev(x, y) / \max(len(x), len(y))$ otherwise

Compatibility node	Initial similarity value
(NamedElement, JElement)	0.5833334
(name, name)	1.0
(EString, String)	0.85714287
(NamedElement, JTypedElement)	0.6923077
(Operation, JTypedElement)	0.23076922

Principe

- Propagation of similarity values in the propagation graph, until finding a fix point

- Propagation formulae : at step i ,

$$s_n^{i+1} = s_n^i + s_n^0 + \sum_{m \in I^n} w(m, n) \times (s_m^0 + s_m^i)$$

- Fixpoint: when similarity values differences is less than ϵ during two successive steps.

Example

Compatibility node	Final similarity value
(NamedElement, JElement)	1.0
(name, name)	0.7771561
(EString, String)	0.5186626
(NamedElement, JTypedElement)	0.2985012
(Operation, JTypedElement)	0.44552472

Principe

- To keep best matches.
- A node of G_{source} can match with several nodes of G_{target} .
- A relative similarity value is computed for each node
- Pairs with a similarity under a threshold are eliminated



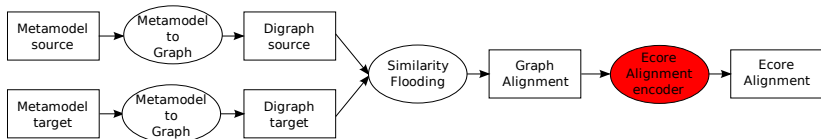
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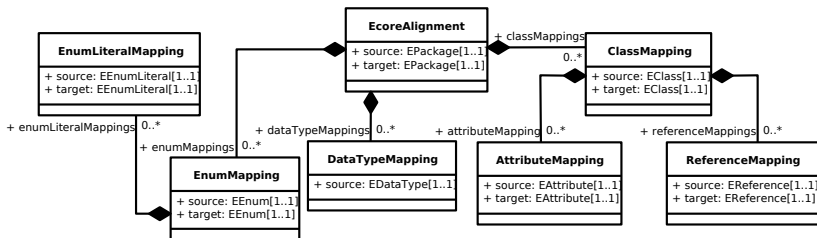
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3. Encoding into an Ecore alignment: reorganizing the alignment graph



The alignment metamodel in ECore



Constraints

- Two classes that match are not abstract.
- Two attributes or references that match are not derived.

Étude de cas

Objectif

Test the six configurations

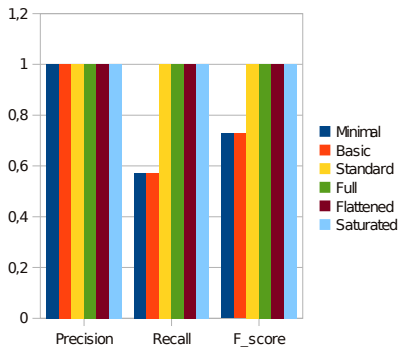
Data

- exMMSource → exMMTarget
- Ecore → Minjava
- Ecore → Kermeta
- Ecore → UML

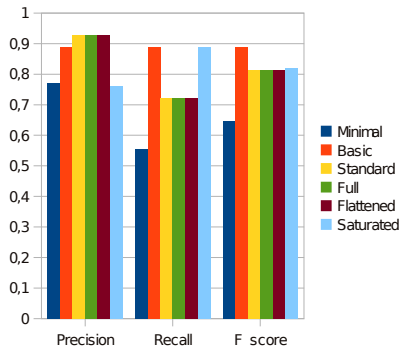
precision, recall et f_score:

- $precision = \frac{\text{Number_of_Correct_Found_Mappings}}{\text{Number_of_Total_Found_Mappings}}$
- $recall = \frac{\text{Number_of_Correct_Found_Mappings}}{\text{Number_of_Total_Existing_Mappings}}$
- $f_score = \frac{2 \times recall \times precision}{recall + precision}$

exMMSource → exMMTarget

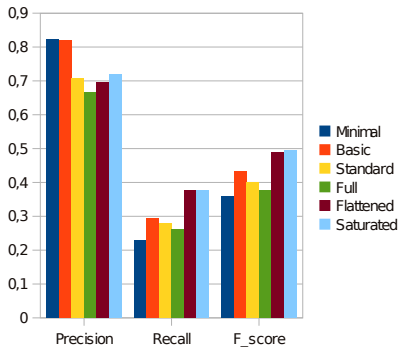


Ecore → Minjava

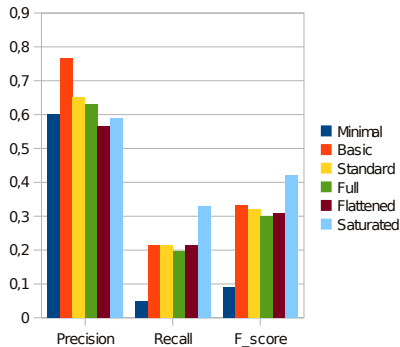


- Good results
- Better results for similar metamodel size
- Configurations Saturated et Basic give good results

Ecore → Kermeta



Ecore → UML



- Good results
- Better results for similar metamodel size
- Configurations Saturated et Basic give good results

- A tool that automatically aligns two metamodels
- Assessment of different configurations
- alignments can be used for code generation

- Code generation
- more complex alignement
- Testing other tunings
- Generic tool



Sergey Melnik, Erhard Rahm and Philip A. Bernstein

Similarity Flooding: A Versatile Graph Matching Algorithm and Its Application to Schema Matching

International Conference on Data Engineering, 2002



Vladimir I. Levenshtein

Binary codes with correction of deletions, insertions and substitution of symbols

Soviet Physics Doklady, 1966

GUM

Gumm (Generic and Useful Model Matcher)

<http://code.google.com/p/gumm-project>

J.R. Falleri

<http://www.lirmm.fr/falleri>