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RCAExplore, a FCA based Tool to Explore Relational Data

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Abstract. Relational Concept Analysis (RCA) is one variant of Formal Concept Analysis for multi-relational dataset exploration. The tool RCAExplore is an implementation of the RCA process where several choices can be made before each iteration: the structure to be used (concept lattice, AOC-Poset, Iceberg lattice), the scaling quantifier (*exist*, *forall, contains, percentage-quantifiers*, etc.), and the considered formal and relational contexts. RCAExplore was developed during Fresqueau ANR 11 MONU 14 project, in order to explore relational hydroecological data, and has also been used on several datasets in different other domains. The source code and a standalone *jar* application are available online. An integration of the tool is ongoing within a platform for data exploration and knowledge representation.

Keywords: Data exploration \cdot Formal Concept Analysis \cdot Relational Concept Analysis

1 Relational Concept Anaysis (RCA) basics

Relational Concept Analysis (RCA) is an extension of Formal Concept Analysis [5] which considers relational data, formalized within a *relational context family* (RCF), i.e. a set \mathbf{K} of object-attribute contexts (object categories) and a set \mathbf{R} of object-object contexts (relations between objects of various categories).

For example, Figure 1 sketches a hydroecological dataset of Fresqueau project. It describes through different relations river stations and taxons living there. Taxons are characterized by life traits (e.g. their locomotion mode). Stations belong to a certain kind of watercourse. Physico-chemical parameters (temperature, oxygen, nitrite, etc.) have been measured on the stations, and their values segmented into two levels. Taxons have been counted from samples and their number segmented into three intervals.

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Fig. 1. A schema of data analysed by RCAExplore, from [4]

Exploring such a relational dataset allows to answer several questions, e.g. what are the links between life traits of the taxons and values of physico-chemical parameters on river stations? Answers can take several forms, such as extracting rules that involve the relations, or grouping objects from the different categories (like physico-chemical parameters, or taxons) depending on their attributes and on the objects of another category they are connected with.

RCA can be used for such tasks. First, the user has to build object-attribute contexts, here, life traits, taxons, stations, physico-chemical characters and watercourse characteristics. Relations between objects are then formalised into seven object-object contexts, e.g. presence of taxons1-19 goes from taxons to stations: a taxon is linked to a station, if the sample size for this taxon on this station is between 1 and 19 (see RCAExplore editor open on this relation, in Fig. 3).

The principle of RCA consists in integrating object-object relations as new attributes (called *relational attributes*) in formal contexts thanks to scaling quantifiers, such as the existential (*exist*) and universal strict (*exist+forall*) scaling quantifiers. It produces iteratively a set of concept lattices (one lattice per object category) interconnected through relational information. The concepts in a given lattice group objects according to the shared attributes and to the connections they have with objects of another category. The result is a family of concept lattice, as sketched out in Fig. 2: a concept C_s reveal a group of stations (1) where at least one physico-chemical value from a group of physico-chemical characteristics (concept C_p) has been measured with level 2, (2) being in a kind of watercourse from a watercourse characteristics group (concept C_w), (3) hosting more than 60% of numerous (100+) taxon types from a group (concept C_t).

The process is as follows: first, lattices are built on the object-attribute contexts K_i ; second, object-attribute contexts are extended with new attributes built using the object-object relations and the concept sets of the lattices; third, new lattices are built on the extended contexts. For instance let $\mathbf{K} = \{K_1, K_2\}$, $\mathbf{R} = \{R\}$, where $K_1 = (G_1, M_1, I_1), K_2 = (G_2, M_2, I_2)$ and $R = (G_1, G_2, r)$. Let C = (X, Y) be a concept of the K_2 lattice (i.e. $X \subseteq G_2$); \exists is the existential quantifier. A relational attribute $\exists r(C)$ is added to the K_1 context, and is owned by an object $g \in G_1$ if $r(g) \cap X \neq \emptyset$. A whole construction process consists in building a finite sequence of contexts and concept lattices based on (\mathbf{K}, \mathbf{R}) and

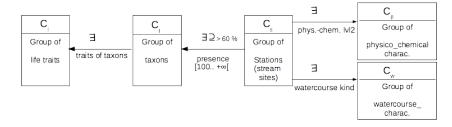


Fig. 2. Illustration of the results of RCA on the dataset described in Fig. 1: concepts are linked by relational attributes, \exists , and $\exists \supseteq_{\geq 60\%}$ are scaling quantifiers

chosen scaling quantifiers. The last sequence is obtained when the fixed point is reached. Details about RCA are given in [6].

2 RCAExplore

RCAExplore¹ is a reference implementation of RCA offering new ways of dealing with relations. RCAExplore is developed in Java. It proposes a relational context family editor (see Fig. 3, highlighting an object-object context linking taxons and stations), an interactive concept lattice family generator (see Fig. 4), and a concept lattice family browser. The tool is developed under the LGPL license.

Context name:				Column actions:			Row actions:	
presence-of-taxons1-19				Add	Del Na	ame Ad	ld Del	Name
🔗 stations 🕜 watercourseKind 🔗 taxons 🔗 presence-of-taxons1-19 🔗 prese 📧								
scaling operator: exist 💌 description:								
	BREI0001	BRUMB001	BRUN001	BRUN002	DOLL001	FECH001		
Asellus			V	V		V		
Athericidae	V		V			V		
Baetis	V		×.	v	V	V		
Bithynia			×.	¥	~			
Boreobd	V					V	1	
Bithynia Boreobd	v					×		



Fig. 3. Editing a relational context family; files are saved in a readable format .rcft

The novelty behind RCAExplore, with respect to other RCA tools such as Galicia², is to provide, besides an automatic mode, a manual mode where it is possible to modify the data considered before each new iteration step of the RCA process. At each step of the process, the user can choose (Fig. 4):

 $^{^{1}\} http://dataqual.engees.unistra.fr/logiciels/rcaExplore$

² http://www.iro.umontreal.ca/~galicia/

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- the conceptual structure (concept lattice, AOC-poset, Iceberg lattice),
- the scaling quantifier (in $\{\exists, \exists\forall, \exists\supseteq, \exists\forall_{\geq n\%}, \exists\supseteq_{\geq n\%}\}$, n = 30 and n = 60),
- the considered contexts (among the available object-attribute and objectobject contexts),
- if they want the simplified or the full intents and extents, and other options.

The tool checks whether the choices made by the user are consistent. For example, a user cannot choose at step n a relational context with the target formal context K, if K has not been chosen at step n - 1. Files are generated at each step, that contain all the conceptual structures of this step in a format (.svg and .dot, from Graphviz³) which is readable and exploitable by other programs, and it is possible to also see the extended contexts in LATEX and HTML. A trace file records the choices made during the process, and can be used to replay the same configuration in automatic mode, namely without again manually selecting structures, quantifiers or contexts. A visualization option also allows to see graphically, at the end of the process, the conceptual structures that have been generated at all steps.

The concept lattice family browser allows, from a given step number, a given context and a given concept name, to look at the parents, children, intent, simplified intent and extent of a concept. If two attributes are selected in the simplified intent and the intent respectively, the corresponding implication rule is displayed.

3 Application examples

As previously introduced, RCAExplore has been primarily used to explore hydroecological data in the framework of Fresqueau ANR 11 MONU 14 project. The question was to link the characteristics (traits) of taxons to the physicochemical state of a river station where they live. Data have been explored according to different paths: focusing on river stations and trying to characterize them both with taxon traits and physico-chemical parameters, focusing on specific relations (e.g. presence of rare taxons in river stations) by varying and combining

fca V IIfeTraits	for30percent containsExist
fca 💌 🕛 🚈 taxons	exist contains60percent presence-of-taxons100
iceberg 💌 50 🗧 physicoChemicalCharacter	contains30percent
fca vatercourseCharacteristics	forall for60percent
acposet 👻 🛛 🗁 stations	
back	back

Fig. 4. Choosing the conceptual structure and scaling operators to be used

³ https://www.graphviz.org/

the scaling operators, etc. Rules have been computed [3], e.g. a straightforward rule⁴, could be (meaning that if a station hosts many of numerous taxon types that appreciate slow current, then the river has calm water):

```
 \exists \supseteq_{\geq 60\%} \text{ presence}_{[100,..+\infty[}(\exists \text{traits_of_taxons}(\texttt{appreciate slow current})) \\ \rightarrow \qquad \exists \text{watercourse_kind}(\texttt{calm water})
```

The features of RCAExplore have been useful to support FCA and several research experiences, to mention just a few: effect of varying the scaling quantifiers on a subset of the Fresqueau dataset is shown in [1], assessing the relevance of using AOC-posets during gradual class model refactoring has been shown in [7] on industrial UML and Java class models. In [2], the authors use RCAExplore to assess the scalability of RCA on connected product comparison matrices. RCA-Explore has also been used to analyse spatio-temporal data characterizing river networks [8] and is currently used to teach FCA and RCA at Montpellier and Strasbourg Universities. It is under integration in the COGUI platform⁵ under the guidance of Alain Gutierrez.

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⁴ Such straightforward rules are useful to understand and to build confidence on other, more informative, rules.

⁵ https://www.lirmm.fr/cogui/3/