R S D

A System for Relational Subgroup Discovery through First-Order Feature Construction

v1.0 – release candidate version

– User's Manual –

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1 Introduction

RSD is a system for relational subgroup discovery in individual-centered domains, based on principles described in

N. Lavrač, F. Zelezný, P. Flach: Relational subgroup discovery through first-order feature construction, in: Stan Matwin and Claude Sammut (eds.): Proceedings of the 12th international conference on inductive logic programming. 6/2002, Springer-Verlag.

An extended treatment of the theoretical principles is presently being submitted. RSD, however, can be **useful as well in tasks other than subgroup discovery**, as long as they require **to generate a propositionalized representation** of classified relational data achieved **through constructing firstorder features**.

RSD is implemented in YAP Prolog. YAP can be obtained at http://source forge.net/projects/yap in versions for various operating systems. The RSD package can be downloaded from http://labe.felk.cvut.cz/~zelezny/rsd. Questions, comments and suggestions should be sent to zelezny@fel.cvut.cz. Following are the main advantages of RSD.

- Syntactical feature construction controlled by mode declarations very similar to those used in the popular systems Progol and Aleph.
- The user can specify various constraints (syntactical or data-related) used by RSD's powerful pruning rules towards performance gains.
- Filtering of irrelevant (unneeded) features.
- Generation of propositionalized data representation compatible with the popular systems CN2 and Weka.
- Automated train-test splitting and stratified cross-validation process in RSD's rule induction component.

RSD consists of three Prolog programs. Their respective function is briefly as follows.

featurize.pl Using specified mode-language declarations, this program identifies all first-order literal conjunctions that by definition form a feature (1. all output variables¹ must also appear as input variables, 2. feature must not be decomposable into two separate features), and at the same time comply to user-defined syntactical constraints. Such features do not contain any constants and the task is completed independently of the input data.

 $^{^{1}}$ I will clarify in a while what *input* and *output* variables mean here.

- process.pl Here the features get confronted with the input data. This component extends the feature set by variable instantiations; certain features are copied several times with some variables (specified by the user) substituted by constants detected by inspecting the input data. During this process, some irrelevant features are identified and eliminated. The program also generates propositionalized representations of the input data using the generated feature set, i.e., it creates a relational table consisting of binary attributes corresponding to the truth values of features with respect to instances of data. The resulting table can be produced in a plain text format or in the format acceptable by CN2, Weka or the following component.
- rules.pl Finds interesting subgroups in the propositionalized data set. An interesting subgroup is a subset of instances which is sufficiently large and has a statistical distribution of target attribute values significantly different than that found in the entire data set. By a simple configuration of settings, this program can also be forged to behave as a standard inducer of rule-based predictive models, such as CN2.

Mode declarations, background knowledge (i.e. all data excluding the main table containing the target attribute) and all parameter/constraint settings are specified in the <filename>.b file by means of Prolog facts/clauses (settings can also be applied on-line when running the respective program components). The .b file is loaded by all three program components.

The relation, whose instances are classified, is represented by

- either the **<filename>.f** and **<filename>.n** files, containing positive and negative (respectively) instances in the form of unary Prolog facts of the target predicate. This variant is suitable only for two-class data.
- or the <filename>.pl file containing binary Prolog facts of the target predicate, where the first argument of each fact is assumed to define the class of the instance. This variant applies to general multi-class problems.

The files relevant to one of the two above itemized options are loaded only by the **process.pl** component. It detects automatically which file(s) is (are) available in the current (or specified) directory and consequently which option of those above applies.

Note that the fact that the main relation predicate must be unary (or binary if the first argument is the class value) means that the predicate serves merely to address instances by their name (or by an identifying constant, in general), whereas their structural description is to be found in the background knowledge. If, however, the description is simple, the user may as well incorporate this description into a compound term, and use such terms as the main predicate argument instead of the name. This is the case of the East-West Trains example presented in the following Quick Start section. Let us digress a bit. If you find it inconvenient that RSD wants you to provide data in Prolog, you may want to try the freely available data convertor Sumatra TT, capable of connecting to data sources of an extensive variety of types (exceeding the command of a mortal person) and producing a Prolog form thereof. The system is described in the award-winning paper

P. Aubrecht, F. Železný, P. Mikšovský, O. Štěpánková: SumatraTT: Towards a Universal Data Preprocessor. In: Proceedings of the 16th European Meeting on Cybernetics and System Research, vol. 2, p. 818-823. Vienna, Austria, 4/2002, Austrian Society for Cybernetics Studies.

This software comes with a fancy graphical interface and can be down-loaded from http://krizik.felk.cvut.cz:8080/SumatraReg/index.html.

Some additional files are engaged in a typical process of using RSD. The featuri ze.pl program component produces a set of features without constants in the file <filename>_frs_noinst.pl and further a 'symbolic' representation of these features in the file <filename>_frs.smb (this file is not significant to the user). The component process.pl loads and uses both of these files to produce the 'instantiated' and filtered feature set <filename>_frs.pl and also the propositional representation of a chosen form; for example a <filename>.cov file is created to feed the rules.pl program, the rule induction component of RSD.

Figure 1 shows the interconnection of individual components and files. Note that the program CN2-SD is a variant of the CN2 program suited for subgroup discovery, described in N. Lavrač et al.: *Rule Induction for Subgroup Discovery with CN2-SD*, IDDM 2002.

2 Quick Start

I assume you decompressed the RSD download package into a chosen directory, where an rsd subdirectory is created. By <rsd> I denote the complete path to RSD, including the rsd directory. Change to <rsd>/samples/trains, containing an example data set representing the well-known East-West trains. I also assume the YAP Prolog is in your path of executables. Run YAP, obtaining roughly the following in the console:

```
$ yap <Enter>
[ Restoring file /usr/local/lib/Yap/startup ]
[ YAP version Yap-4.3.23 ]
?-
```

Proceed as follows to consult the f.pl file, which is actually a symbolic link to <rsd>/code/featurize.pl.

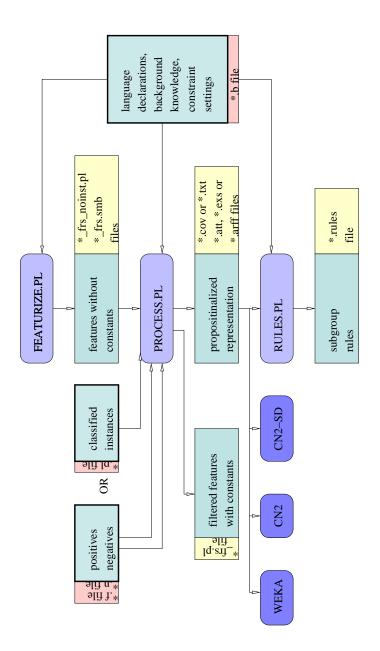


Figure 1: The cooperation of individual RSD (and external) components through files. Rounded boxes correspond to program codes, other boxes represent data files. Bold-framed boxes denote input files provided by the user.

```
?- [f]. <Enter>
  [ consulting /rsd/trains/f.pl... ]
  [ consulted /rsd/trains/f.pl in module user, 30 msec 51512
  bytes ]
yes
?-
```

Now to read the trains.b file present in the current directory, type²

r(trains). <Enter>.

When the file is read, type

s. <Enter>

to have the program show you the features constructible on the basis of the declarations in the trains.b file. The following should appear.

```
f(1,A):-hasCar(A,B),carshape(B,C),instantiate(C).
f(2,A):-hasCar(A,B),carshape(B,C),carlength(B,D),
instantiate(C),instantiate(D).
f(3,A):-hasCar(A,B),carshape(B,C),carlength(B,D),has_sides(B,E),
instantiate(C),instantiate(D),instantiate(E).
f(4,A):-hasCar(A,B),carshape(B,C),carlength(B,D),has_roof(B,E),
instantiate(C),instantiate(D),instantiate(E).
f(5,A):-hasCar(A,B),carshape(B,C),carlength(B,D),has_wheels(B,E),
instantiate(C),instantiate(D),instantiate(E).
f(6,A):-hasCar(A,B),carshape(B,C),carlength(B,D),has_load(B,E),
instantiate(C),instantiate(D),loadshape(E,F),instantiate(F).
```

(...etc.)

Note that the instantiate/1 literals, which appear in the features, identify variables that will be substituted by constants in a following step described further on. To write the features into the output files (named trains_frs_noins .pl and trains_frs.smb), do as follows

?- w. <Enter>

0.781 seconds taken to construct features.

yes ?-

²Note that if a filename was used containing special symbols (unlike here) that disqualify the name from being a Prolog atom, it would have to be enclosed in apostrophes, e.g. r('trains-version1'). etc.

Now exit YAP by pressing CTRL+D, and run it again. This time you invoke p.pl linked to <rsd>/code/process.pl

```
$ yap <Enter>
[ Restoring file /usr/local/lib/Yap/startup ]
[ YAP version Yap-4.3.23 ]
?- [p]. <Enter>
[ consulting /rsd/trains/p.pl... ]
[ consulted /rsd/trains/p.pl in module user, 40 msec
80232 bytes ]
yes
?-
```

Then again you read the declaration file along with the outputs of the previous step and the classified data, as follows.

```
?- r(trains). <Enter>
20 examples read.
[ reconsulting /rsd/trains/trains.b... ]
[ reconsulted /rsd/trains/trains.b in module user, 10 msec
13220 bytes ]
[ reconsulting /rsd/trains/trains_frs_noinst.pl... ]
[ reconsulted /rsd/trains/trains_frs_noinst.pl in module user,
20 msec 43880 bytes ]
Congratulations! You succeeded loading all neccessary files.
yes
?-
```

Now again, if you type

s. <Enter>

the program will show you a set of features. This time this is the set of nonredundant features which contain constants extracted from the data set and comply to constraints related to feature's coverage on the data. You should see the following:

```
f(1,A):-hasCar(A,B),carshape(B,u_shaped).
f(2,A):-hasCar(A,B),carshape(B,bucket).
f(3,A):-hasCar(A,B),carshape(B,hexagon).
f(4,A):-hasCar(A,B),carshape(B,ellipse).
f(5,A):-hasCar(A,B),carshape(B,rectangle),carlength(B,long).
f(6,A):-hasCar(A,B),carshape(B,rectangle),carlength(B,short).
```

(...etc.)

Now write the expanded feature file as illustrated below. The program comes up with a progress meter accompanied with a very funny message, which fortunately appears only once.

?- w. <Enter>

Warning: progress meter is approximate and takes no legal responsibility for accurate estimates.

Finally let us create a propositional representation (named trains.cov) acceptable by the RSD rule inducer.

Note that changing the first argument of w/2 to weka or cn2 would produce files acceptable by the respective systems, text would produce a plain text file, and that appropriate file name extension is added automatically. To proceed to the rule induction component, exit YAP (CTRL+D), relaunch it, and in a manner analogous to the above procedures, type

[r]. <Enter>

to consult the linked file <rsd>/code/rules.pl. To read the train.b file along with the result of the preceding step (i.e. the propositionalized representation trains.cov), you again - surprise - use the command

r(trains). <Enter>

To induce rules, type

i. <Enter>

obtaining the following in response.

Inducing rules.

```
Class: east (prior = 0.5)
IF 93 AND 69 AND 16 THEN Class = east
Evaluation: 0.225 Significance: 9.0 Class distribution: [10,1]
Remaining % of initial total weight = 75.0
IF 93 AND 112 AND 16 THEN Class = east
Evaluation: 0.15 Significance: 12.0 Class distribution: [9,0]
Remaining % of initial total weight = 68.0
IF 106 THEN Class = east
Evaluation: 0.062 Significance: 7.0 Class distribution: [5,0]
Remaining % of initial total weight = 65.0
IF 9 AND 106 THEN Class = east
Evaluation: 0.048 Significance: 7.0 Class distribution: [5,0]
Remaining % of initial total weight = 64.0
```

(...etc.)

When rules have been generated, you may save them by typing

w. <Enter>

(this will create the trains.rules file). When you run rules.pl newly, you may load these rules back by typing

rr. <Enter>

(after the initial command r(trains).) instead of re-inducing them. This may be used for later evaluation purposes (explained further in this text). Finally, the command

s. <Enter>

serves to show the rules currently in the memory.

3 Using RSD

3.1 Language declarations

Every effort has been taken to make mode-language declarations as similar as possible to the popular systems Progol and Aleph, so that existing declarations and background knowledge can be recycled.

All predicates which can appear in the features are declared in the ***.b** file. There is exactly one *head* declaration describing the predicate of the classified (main) relation. It has the form :-modeh(1,<S>(+<T>)).

where $\langle S \rangle$ is the predicate symbol of the main relation (found in the .f and .n, or .pl file(s)) and $\langle T \rangle$ specifies the *type* of the variable that addresses learning instances. It may be an arbitrary atom (a self-explaining name is recommended though). An appearance of this variable type in body predicates (see below) will denote that an instance-addressing variable should occur in the place of that appearance. The number 1 and the + sign in the head declaration are included only for compatibility sakes. Note that a unary predicate is declared by the head declaration even if one works with a .pl data file containing binary facts; the declaration merely serves to identify the main predicate symbol and the instance key.

The body declarations specify predicates which can appear as literals in the feature body. They have the form

```
:-modeb(<R>,<S>(<M1><T1>,<M2><T2>, ... ,<Mn><Tn>)).
```

where

- <R>, the *recall*, is a number which specifies how many times the predicate can appear in one feature body with the same input variables (those whose *mode* is +, see below). This can be used for predicates giving multiple answers (outputs) to a given input.
- <S> is the symbol of the declared predicate.
- <Mi> is the mode of the i^{th} argument variable, which may either be + (input variable) or (output variable). I shall explain the meaning of the mode in a moment.
- <Ti> is the *type* of the *i*th argument variable, which is an arbitrary atom.

The modes and types are used to constrain the feature-language bias, similarly again to the systems Progol and Aleph. They are employed as the following rule dictates.

A predicate will only be considered as a literal at a given place in the feature definition, if each of its input arguments is of a type equal to the type of an output argument of some preceding literal, or equal to the type T found in the modeh declaration.

Predicates, which are declared by modeb/2 with no output variables, are called *property predicates*. Other predicates declared by modeb/2 are called *structural predicates*.

Unlike Progol, the user does not specify value-domains of variable types. Types are here used solely for purposes of variable matching. Unlike both Aleph and Progol, there is no variable mode labelled by the sign **#**, which in these systems means that a constant value should be put in the given argument place. RSD does not extract constants from a single example, as in the mentioned systems, but rather selects them 'carefully' from the whole data set in the following manner.

There is one special reserved unary property predicate called instantiate/1, which may not occur in the background knowledge. It specifies a variable that should be substituted with a constant during the subsequent feature processing by the process.pl component. Let us illustrate this on an example. Out of the following declarations³ in the East-West trains domain

```
:-modeh(1, train(+train)).
:-modeb(1, hasCar(+train, -car)).
:-modeb(1, hasLoad(+car, -load)).
:-modeb(1, hasShape(+load, -shape)).
:-modeb(*, instantiate(+shape)).
```

exactly one feature would be generated:

f(1,A) :- hasCar(A,B),hasLoad(B,C),hasShape(C,D),instantiate(D).

However, the component process.pl would take f1 and substitute it by a set of features, in each of which the instantiate/1 literal is removed and the D variable is substituted by a constant, making the body of f1 provable in the data. Provided they contain a sufficient number of trains with a rectangle load, the following feature will appear among those created out of f1:

f(1,A) :- hasCar(A,B),hasLoad(B,C),hasShape(C,rectangle).

A similar principle applies for features with multiple occurrences of the literal instantiate/1. Arguments of this literal within the feature form a set of variables ϑ ; only those (complete) instantiations of ϑ making the feature's body provable on the input database will be considered.

3.2 Settings

Each of the three RSD components has a number of configurable parameters, which can be specified in the .b file by using the set/2 command in the following form

```
:-set(<parameter>,<value>)
```

The set of parameter settings in the .b file is common to all three RSD components that load the .b file; each one will obviously apply the relevant subset. Tables 1 to 4 provide explanations to these settings. For the components featurize.pl and process.pl there is a special group of formatting-related settings. These define, for example, whether constructed features should contain variables such as A, B, C, ... or Car1, Car2, Car3 ..., where the feature numbers should be put, etc.

³The recall parameter has no effect on the instantiate/1 predicate.

featurize.pl

Essential Settings Parameter Value and Function clauselength An integer specifying the maximum length of a feature body. **Default** is 8. An integer specifying the maximum *depth* of variables depth found in a feature body. See e.g. S. Muggleton: Inverse Entailment and Progol, New Gen. Computing 13, 1995 about variable depth. Default is 4. max_occ A compound term (Symbol/Arity, Occurrences) specifying the maximum number of occurrences of the predicate with given symbol and arity in a feature body. Not limited by default. pruning Value is on or off. If off, admissible search-space pruning functions are not used. This is not recommended. Default is on.

Formatting Specifications

Parameter	Value and Function		
head_name	An atom specifying the head symbol of features.		
	Default is f.		
feature_num	One of argument, end_of_name. In the former case,		
	feature heads are numbered as f(1,A), f(2,A), etc.,		
	otherwise f1(A), f2(A), etc.		
	Default is argument.		
format_vars	One of alphabet, all_cap_type and first_cap_type		
	alphabet: Names of used variables are subsequently		
	A, B, C,		
	all_cap_type: Variables have the name of their type,		
	followed by an integer, for the Trains domain e.g. CAR1,		
	CAR2,		
	first_cap_type: Same as above, but Car1, Car2,		
	Default is alphabet.		

Table 1: Parameter settings relevant to the featurize.pl component of RSD. Note that the default values of the formatting specifications should be used to create an output file correctly processable by process.pl.

process.pl

Essential Settings

Parameter	Value and Function				
negation	One of now, later, none.				
	now: To features generated by featurize.pl, process.pl				
	will also add their versions where the complete body is				
	negated. (Negations of individual literals can be done by				
	suitably defining background knowledge predicates.)				
	later: tells the program that an inducer capable of				
	negating features will be applied on the propositionalized				
	representation. This influences functions described below.				
	Default is none.				
min_coverage	An integer mc . All features (including negated versions)				
	covering fewer than mc instances will be discarded.				
	However, if negation is later (see above), a feature				
	is discarded only if both (a) coverage thereof and (b) the				
	coverage of its negated version are smaller than mc .				
	Default is 1.				
filtering	One of true, false. If true, each feature will be discarded				
	if (a) it covers the same set of instances as some previously				
	constructed feature, or (b) it covers all instances.				
	Default is true.				
splitting	Either false, or a number in $(0, 1)$, or an integer. If false: the resulting propositional representation (call it				
	RPR) will describe all instances of the input data.				
	If in $(0, 1)$: RPR will be split into train and test sets. The				
	number then indicates the proportion of instances to be put				
	in the test split. Supported for RSD- and CN2-formatted				
	RPR's. (For Weka, train/test splitting is a built-in).				
	If integer N : RPR is a set of N stratified cross-validation				
	folds. Supported only when creating RPR in an RSD rule-				
	inducer format. Cross-validation splits for the CN2 format				
	is not yet supported by RSD. For Weka, cross-validation				
	splitting is a built-in.				
	No kind of splitting is at the moment available for plain-				
	text RPR (it has to be done manually).				
	Default is false.				

Table 2: Essential parameter settings relevant to the process.pl component of RSD. 'RPR' abbreviates 'resulting propositional representation'.

process.pl

Formatting Specifications						
Parameter	Value and Function					
true_sign	Determines the sign used to indicate that a feature holds					
	for an instance in the RPR attribute table. Default is '+'.					
false_sign	See above and guess :-). Default is '-'.					
separator	The symbol for separating attributes in a row of the RPR					
	attribute table. Default is ' ' (space).					
terminator	The symbol that should appear in the end of rows in the					
	RPR attribute table. Default is '' (empty char).					
meter_width	Integer specifying the width (in characters) of the progress					
	meter displayed when conducting an exhaustive job.					
	Default is 50.					

Formatting Specifications

Table 3: Formatting specifications relevant to the process.pl component of RSD. 'RPR' abbreviates 'resulting propositional representation'. Note that the first four settings apply only when generating a plain-text RPR, and are overridden when creating files for specified learning systems (CN2, Weka, RSD rule-inducer).

3.3 Basic Commands

Although each of the three components of RSD provides a different set of services, the basic user's control of all the components is quite unified. Therefore the description below is common to all the three programs. The commands are always issued in the YAP console after having consulted the respective program code (e.g. for the featurize.pl component this was done in the Quick Start section by [f]. <Enter> as the the file f.pl linked to the <rsd>/code/featurize.pl program was present in the current directory).

In the following, expressions of the form S/A correspond to a predicate of the symbol S and arity A.

- r(filename). Reads input file(s). All three program components read the filename.b file, but rules.pl will not complain if it does not find the file (it simply won't apply any settings then). The process.pl will also read the filename_frs_noinst.pl and filename_frs.smb files, which have been created by featurize.pl, and the classified data file(s) (see Section 1). The rules.pl will also read the filename.cov file containing the propositional representation created by process.pl.
- s. Shows the product on the screen: On this command, featurize.pl will create non-instantiated features and list them, process.pl will process features and list the resulting set. The component rules.pl upon this

rul	es.	pl
-----	-----	----

Essential Settings Parameter Value and Function beam_width An integer specifying the width of the beam in the beam-search performed to find rules. **Default** is 3. max_length An integer specifying the maximum length of an induced rule in literals. **Default** is 3. evalfn One of acc, wracc. Specifies the heuristic ruleevaluation function (accuracy, or weighted relative accuracy). **Default** is wracc. One of freq, laplace. Specifies how rule accuracy acc_est should be estimated (plain frequency count, or the Laplace estimate). **Default** is **freq**. One of cover, weighted. If cover, then the search standard instance-covering algorithm is used in the rule-induction cycle. If weighted, then the instanceweighting approach is taken. Default is weighted. weight_threshold Number $c \in (0, 1)$. In an instance-weighting induction, generation of rules for the current class is stopped once the total weight of the class instances falls below c.wwhere w is the initial total weight of the class instances. **Default** is 0.1. gamma In an *instance-weighting* induction, number specifying the γ parameter in the weight-decay formula. Default is 1. sig_threshold Number specifying the minimum significance value of an acceptable rule. **Default** is 0. eval_threshold Number specifying the minimum heuristic evaluation value of an acceptable rule. **Default** is 0. Integer specifying the maximum number of rules that max_rules_for_class should be generated for one class. **Default** is 10. stop_on_sig One of yes, no. If yes, then rule-generation for the current class terminates once the best rule in the search space *deceeds* the minimum significance value specified by sig_threshold. Default is no. *Remark:* look at http://130.88.203.73/asktheexperts/faq/ aboutwords/exceed if you doubt the verb to deceed (my spell-checker does). stop_on_eval Same as stop_on_sig, but related to the minimum evaluation determined by eval_threshold.

Table 4: Parameter settings relevant to the rules.pl component of RSD.

command will show the induced rules, but the rules must have been previously induced, or read from a .rules file (see the i/0 command below).

- w. The same as the s. command above, but instead of displaying on the screen, the results are written into appropriate files. File name is created by taking *filename* from the initial r/1 command (see above) and adding appropriate extensions.
- w(filename). Just like w/0 above, but allows to specify a different file name (extensions again added automatically).
- w(system, filename). Only for process.pl, after having used the w/0 or s/0 command: creates a file containing the propositional representation of the data using the generated features. The format is determined by the system argument, which can be one of rsd (for the RSD rule-inducer component), cn2, weka or text (see the formatting parameters in Table 3 for the last option). File name extension is added automatically.
- i. Only for rules.pl: induces rules. Each rule will have a conjunction of feature numbers in its antecedent (the meaning of individual features has to be looked up in the _frs.pl file), the class name in the consequent, and a few characteristic quantities attached, calculated on the training data.
- rr. Only for rules.pl: reads rules from the file filename.rules (erasing any rules currently in memory), where filename is taken from the initial r/1 command.
- rr(filename). Just like rr/0 above but allows to specify a different file name
 (extension added automatically).

Two special commands, common to all three components are the following.

- set (parameter, value). This command changes a parameter as described in Section 3.2. Another usual way to specify parameters is to put this command into the .b file in the spirit of Section 3.2. Obviously, issuing this command will overwrite the corresponding setting read previously from the .b file. For example, you may want to repeatedly create features, induce rules, etc., each time with altering constrain settings, in a single session with an RSD component.
- st. Shows current parameter settings on the screen (excluding those that currently acquire the default value). Note that this listing may also contain settings not relevant to the currently running RSD component, as the settings may have been specified commonly for all components in the .b file.

Finally, the two following commands (relevant to featurize.pl) may be in principle issued interactively, but it is not recommended to do so.

modeh/2 The head-literal declaration command.

modeb/2 The body-literal declaration command.

Rather, declarations should be a part of the .b file, as explained in Section 3.1.

3.4 Assessment Functions

The components process.pl and rules.pl have special built-in functions enabling for a statistical assessment of discovered rules, from the point of view of subgroup interestingness. Also, detecting and discarding rules not lying on the convex-hull of the ROC curve is supported.⁴

Although RSD has some tentative functions allowing to test the *predictive* classification capacity of induced rule models, they are not yet supported 'officially', as the primary scope of RSD is (surprise:) subgroup discovery.

We will now illustrate the use of the assessment functions again on the East-West trains example domain for clarity. Note, however, that data splitting (and especially cross-validation) on this toy domain counting 20 instances may yield funny results.

3.4.1 Quality Calculations and Train-Test Validation

The assessment procedure starts already when producing the propositional representation by the component process.pl, where the user specifies the ratio of the number of testing instances to the number of all instances. After features have been created by the command s. or w., specify the ratio in the YAP console (if you haven't set it in the .b file already), e.g. 0.2, as follows

```
?- set(splitting,0.2). <Enter>
```

yes

Then, after issuing the command

w(rsd,trains). <Enter>

two files are generated: trains.cov and trains_test.cov, of which the latter contains the description of 20% of instances.⁵ The two splits are stratified, which means that they have the same (or very close) distribution of the target-class values.

Subsequently, after launching the rules.pl component, the training file is read by

r(trains).

 $^{^{4}}$ An explanation of the method of creating subgroup-discovery suited ROC curves as well as of the significance and other characteristics can be found in the paper referred to in the Introduction, but is beyond the scope of this manual.

⁵When generating the split files for CN2 by w(cn2,trains)., the files will be: (a) trains.att - the attribute description file, (b) trains_train.exs - the training examples, and (c) trains_test.exs - the testing examples.

and rules are induced by the i. command. Say you first want to check the average quality of the induced rules, as it appears to be on the training data. Proceed as follows.

```
?- sq. <Enter>
Data file = trains
Number of rules = 20
Average Length of Rule = 1.95
Average Significance = 4.44
Average Coverage = 22.5% of instances
Average (over classes) Area Under Roc = 0.8336
```

yes

The sq. command calculates and reports self-explaining rule characteristics, averaged over all induced rules (subgroups). Since the ROC diagram is constructed for each class separately, the last reported number is further averaged over all classes.

The following commands will read⁶ the testing file (training instances are automatically erased from the memory), and calculate the subgroup characteristics on the testing instances.

The message "trains_test.b file not found, no settings changed." is not an error, it merely means that only data, not settings, have been replaced.

 $^{^6\}mathrm{Remind}$ that that quoting marks must now be used in the argument of r/1 due to the underline character in the name.

Now you may discard all rules, which do not lie on the convex hull of the ROC curve belonging to the class present in the consequent of the respective rule, which is done by the command

dr. <Enter>

(it can only be issued *after* the sq. command which constructs the ROC curves). In the case of this example, only one rule is discarded and the new average subgroup characteristics can again be checked by the sq. command. However, this approach is not totally correct because rules would be filtered according to the ROC curves generated on the basis of the *testing*, not *training* data. The correct and exhaustive sequence of commands (which is also followed by the automated validation procedure described in the next Section) would thus be as follows.

- 1. r(trains). <Enter> (Reads the training data.)
- 2. i. <Enter> (Induces rules.)
- 3. w. <Enter> (Writes the induced rules into the file trains.rules.)
- 4. sq. <Enter> (Calculates and shows characteristics of unfiltered rules on training data.)
- 5. dr. <Enter> (Discards rules below ROC convex hull.)
- 6. w('trains-ch'). <Enter>
 (Writes the filtered rules into the file trains-ch.rules.)
- 7. sq. <Enter> (Calculates and shows characteristics of filtered rules on training data.)
- 8. rr. <Enter> (Reads back the unfiltered rules, erasing the filtered rules from memory.)
- 9. r('trains_test'). <Enter> (Reads the testing data.)
- 10. sq. <Enter> (Calculates and shows characteristics of unfiltered rules on testing data.)
- 11. rr('trains-ch'). <Enter> (Reads back the filtered rules.)
- 12. sq. <Enter> (Calculates and shows characteristics of filtered rules on testing data.)

3.4.2 Automating the Cross-Validation Procedure

Performing the validation procedure for a number of cross-validation folds would be tiresome, therefore RSD provides means of automation.

First of all, the stratified cross-validation folds are again produced by the **process.pl** component as in the example above, however, the **splitting** parameter is set this time to an integer N defining the number of folds. For example, using

set(splitting,5). <Enter>

and then

w(rsd,trains). <Enter>

5 cross-validation folds will be produced, represented by 5 pairs of files

```
trains1.cov, trains1_test.cov ... trains5.cov, trains5_test.cov.
```

When the rule inducer rules.pl is launched, one may read in the trains.b file (if there are any settings to be applied) by

['trains.b']. <Enter>

instead of using the r(trains). command (this is because one does not assume the file trains.cov to exist in the cross-validation case). The rest of the job is very simple. Just type

validate(trains). <Enter>

The program will automatically detect the number of cross-validation folds present in the current directory (and even if there are none, it will check if files of a single train-test split exist) and perform an adequate number of induction - validation procedures, with progress commented on the screen. After they are completed, the following message will appear

```
Cross-validation completed with 5 folds. Results stored in files: 'report.txt', 'report-test.txt', 'report-ch.txt' and 'report-ch-test.txt'. yes
```

The mentioned files located in the current directory then contain tabulated results measured for (respectively): all rules on the train set, all rules on the test set, convex-hull rules on the train set and convex-hull rules on the test set. Here the convex-hull rules are always obtained by inspecting the ROC curves created on the basis of the train sets.

Furthermore, induced rules are stored for each training fold. All rules generated on fold number 1 are stored in the file trains1.rules, convex-hull rules for this fold are found in trains1-ch.rules, and similarly for the rest of folds.

4 Frequently Asked Questions (none yet)

Got any? Mail me at zelezny@fel.cvut.cz. All questions, besides being answered, will enter a lottery with a valuable first (and only) prize (yet to be determined). Drawing on April 30, 2003.