Lab 2 Association Rule Mining

The Titanic Dataset

 downloaded "titanic.raw.rdata" from http://www.rdatamining.com/data.

```
> str(Titanic)

table [1:4, 1:2, 1:2, 1:2] 0 0 35 0 0 0 17 0 118 154 ...
- attr(*, "dimnames")=List of 4
    ..$ Class : chr [1:4] "1st" "2nd" "3rd" "Crew"
    ..$ Sex : chr [1:2] "Male" "Female"
    ..$ Age : chr [1:2] "Child" "Adult"
    ..$ Survived: chr [1:2] "No" "Yes"
```

The Titanic Dataset

```
> str(titanic.raw)
data.frame:
            2201 obs. of 4 variables:
 $ Class : Factor w/ 4 levels "1st", "2nd", "3rd", ...: 3 3 3 3 3 3 3 3 3 ....
 $ Sex : Factor w/ 2 levels "Female", "Male": 2 2 2 2 2 2 2 2 2 ...
 $ Age : Factor w/ 2 levels "Adult", "Child": 2 2 2 2 2 2 2 2 2 ...
 $ Survived: Factor w/ 2 levels "No", "Yes": 1 1 1 1 1 1 1 1 1 1 ...
> head(titanic.raw)
  Class Sex Age Survived
   3rd Male Child
                          No
   3rd Male Child
                          No
```

Basic arules () function

```
> install.packages("arules")
> library(arules)
> rules.all <- apriori(titanic.raw)
> rules.all
> inspect(rules.all)
```

arules () parameters

```
> rules<-apriori(titanic.raw, control=list(verbose=F),
                parameter=list(minlen=2, supp=0.005, conf=0.8),
                appearance = list(rhs=c("Survived=No",
                                          "Survived=Yes"),
                default="lhs"))
> quality(rules)<-round(quality(rules),digits=3)</pre>
> rules.sorted <- sort(rules,by="lift")</pre>
> inspect(rules.sorted)
```

arules () parameters

Some key parameters:

- rhs containing survival information only, the rest not interested: rhs=c("Survived=No", "Survived=Yes")
- Default ="lhs" => All other items can appear in the lhs.
- To suppress the details of progress use verbose=F.
- minlen=2 means lhs consists of at least two items.

Removing Redundancy

• Rule 2 is a redundant rule for Rule 1. Adding sex=Female will not affect the rule.

Removing Redundancy

To find redundant rules:

```
> # find redundant rules
> subset.matrix <- is.subset(rules.sorted, rules.sorted)
> subset.matrix[lower.tri(subset.matrix, diag=T)] <- NA
> redundant <- colSums(subset.matrix, na.rm=T) >= 1
> which(redundant)

[1] 2 4 7 8

> # remove redundant rules
> rules.pruned <- rules.sorted[!redundant]
> inspect(rules.pruned)
```

Matrix: lower.tri()

```
> m2 <- matrix(1:16, 4, 4)
> lower.tri(m2)
> m2[lower.tri(m2)] <- NA
> m2
```

```
> m2
     [,1] [,2] [,3] [,4]
[1,]
                     13
[2,]
            6 10
       NA
[3,]
                      15
       NA
            NA
[4,]
                      16
       NA
                 NA
            NA
```

Interpreting Rules

Interpreting Rules

```
lhs
                rhs
                                    support confidence
                                                            lift
1 {Class=2nd,
   Age=Child} => {Survived=Yes} 0.010904134 1.0000000 3.0956399
2 {Class=1st,
  Age=Child} => {Survived=Yes} 0.002726034 1.0000000 3.0956399
3 {Class=1st,
  Age=Adult} => {Survived=Yes} 0.089504771 0.6175549 1.9117275
4 {Class=2nd,
  Age=Adult} => {Survived=Yes} 0.042707860 0.3601533 1.1149048
5 {Class=3rd,
   Age=Child} => {Survived=Yes} 0.012267151 0.3417722 1.0580035
6 {Class=3rd,
  Age=Adult} => {Survived=Yes} 0.068605179 0.2408293 0.7455209
```

Visualization Rules

```
> install.packages("arulesViz")
> library(arulesViz)
> plot(rules.all)
> plot(rules.all, method="grouped")
> plot(rules.all, method="graph")
```

Visualization Rules

Grouped matrix for 27 rules



