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The Prospector - Beyond Classification

Summary

The Prospector is an algorithm for data description standing

on the border between clustering and classification:

We are searching for clusters with respect to a target variable.

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Classification

The target variable is predicted from the explanatory variables. We use an algorithm based on averaging of classification trees called Random Forests proposed by Professor Emeritus Leo Breiman from University of California. This method benefits from averaging a lot of classification trees, each of them being trained on different subsample of the data. Moreover, the best splitting variable in each node of the tree is selected from a random subset of all possible variables that preserves from overfitting.

Categorization

Continuous variables are categorized into a few distinct categories and ordinal categorical variables are possibly recoded into fewer number of categories. This can be efficiently done by genetic algorithm: - pool of chromosomes 001100100010001...00, each gene stands for a distinct

value of variable, 0 = no split, 1 = split - fitness of chromosomes (given the target variable) is equal to either its

correlation or Cramer's V c^{2}

 $V = \sqrt{\frac{1}{n \cdot \min(M-1, N-1)}}$

where n is number of cases, M, N are number of categories - offspring is created using crossover operator, combination of parent genes step by step, or *mutation*

Clustering

Categories of each variable are sorted with respect to the target variable. This creates *n*-dimensional grid. Each grid point is evaluated by a criterion based on its radius, purity, weight and then we try to incorporate its nearest neighbours to improve the criterion. This creates so called nuggets and they are compared to each other and (a few) best ones are selected

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Case study

A brewery wants to define status of its brand of beer. Outlets are described according to their type, size, volume of all beer they sell, type of cuisine they provide and so on. Some of the outlets are scored as either "relevant to the brand" or "not

relevant to the brand". The task is to find the status (characterization, description) of the outlets that are "relevant to the brand".

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Fig. 4: XML output from the categorization



Fig. 3: Fitness of the best chromosome in population

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Fig. 5: F	Projecti	ion of th

arid into 2-dimension

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