Machine Learning for Data Analysis Assignment 1

This assignment has been completed with SAS-studio and edited with MS word for explanations and analysis. The SAS Code is added as an APPENDIX at the end of this document.

Response Variable: Life Expectancy,

1: High – Life expectancy greater or equal to 60 years

2: Low – Life expectancy less than 60 years

Explanatory Variables:

- country
- incomeperperson
- alcconsumption
- breastcancerper100th
- femaleemployrate
- hivrate
- suicideper100th
- employrate
- urbanrate

The decision tree analysis tests the non-linear relations among the specified variables.

Entropy and cost complexity criteria are included to obtain final subtree.

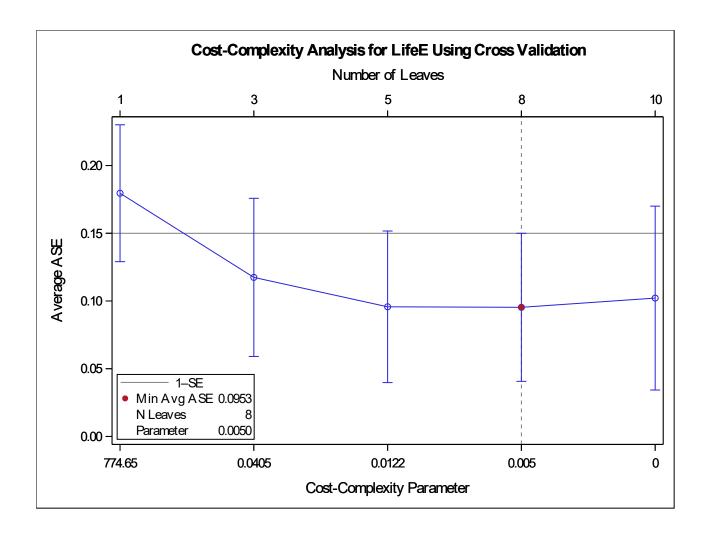
[&]quot;LifeExpectancy" is converted to a categorized variable yielding values of 'high' and 'low' with a threshold at 60

Performance Information			
Execution Mode	Single-Machine		
Number of Threads	2		

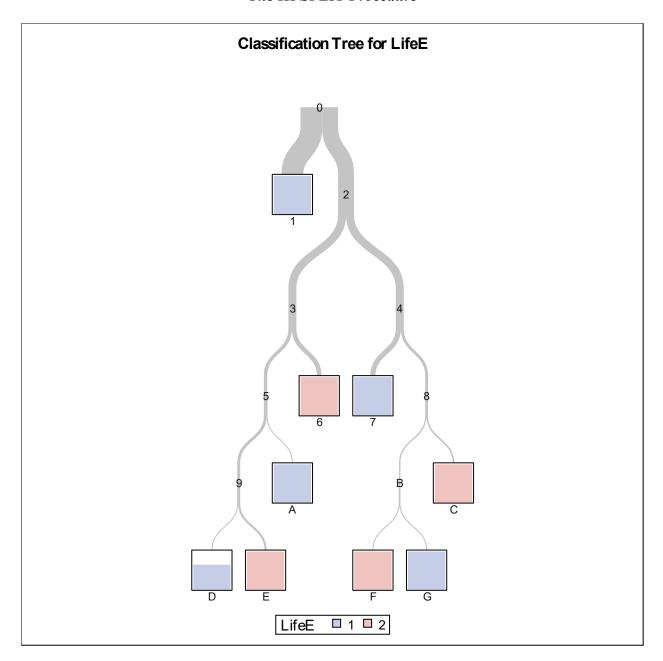
Data Access Information					
Data	Engine	Role	Path		
WORK.NEW	V9	Input	On Client		

Model Information				
Split Criterion Used	Entropy			
Pruning Method	Cost-Complexity			
Subtree Evaluation Criterion	Cost-Complexity			
Number of Branches	2			
Maximum Tree Depth Requested	10			
Maximum Tree Depth Achieved	7			
Tree Depth	5			
Number of Leaves Before Pruning	11			
Number of Leaves After Pruning	9			
Model Event Level	1			

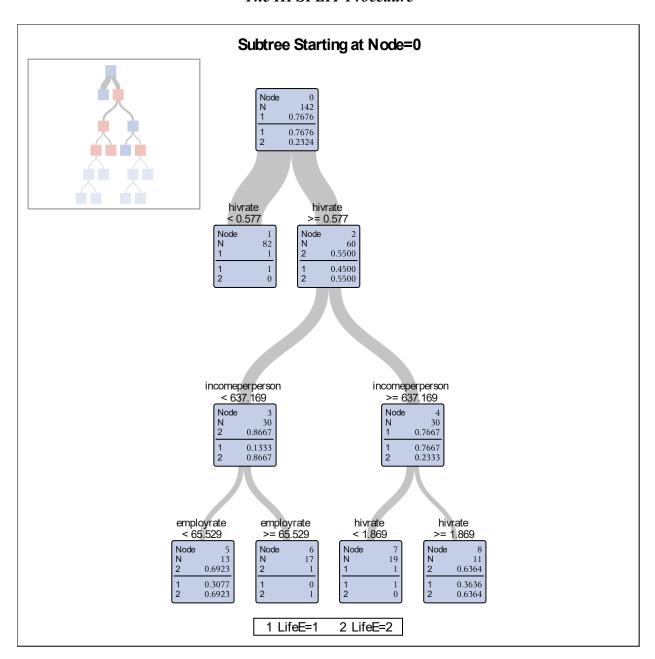
Number of Observations Read	213
Number of Observations Used	142



The HPSPLIT Procedure



The HPSPLIT Procedure



The HPSPLIT Procedure

 Model-Based Confusion Matrix

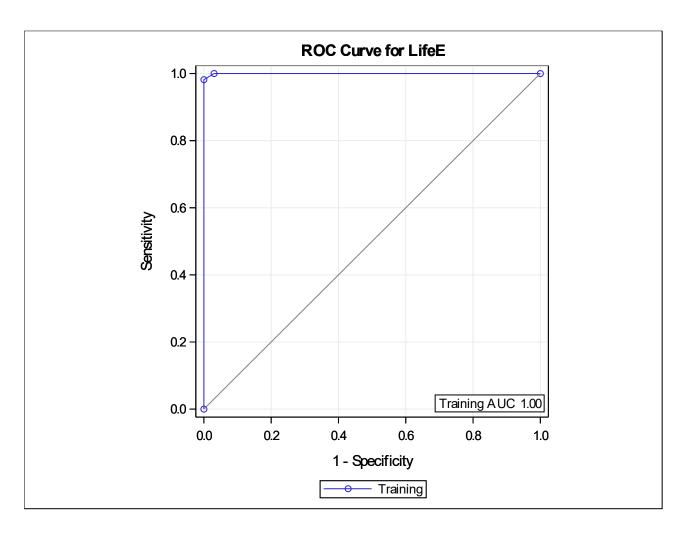
 Predicted
 Error Rate

 1
 109
 0
 0.0000

 2
 1
 32
 0.0303

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Model-Based Fit Statistics for Selected Tree								
N Leaves	ASE	Mis- class	Sensitivity	Specificity	Entropy	Gini	RSS	AUC
9	0.00469	0.0070	1.0000	0.9697	0.0194	0.00939	1.3333	0.9997



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Variable Importance						
	Variable	Tr				
Variable	Label	Relative	Importance	Count		
hivrate	HIVRATE	1.0000	5.1579	2		
incomeperperson	INCOMEPERPERSON	0.7159	3.6923	2		
employrate	EMPLOYRATE	0.3709	1.9133	2		
suicideper100th	SUICIDEPER100TH	0.3622	1.8684	1		
femaleemployrate	FEMALEEMPLOYRATE	0.2700	1.3926	1		

Analysis

142 were considered for the analysis, out of 213 observations

The initial tree yields 11 nodes. When pruning is applied, subtree yields 9 leaves.

I - The first classification is obtained when the HIV rate branches into two groups:

- a) 82 countries with HIV rate < 0.577 and 100% has a higher life expectancy
- b) 60 countries with HIV rate (>= 0.577) where 45% have high life expectancy and 55% have low.

II - This subgroup of 60 countries is then divided by income per person with a threshold of 637.169.

III - The subgroup tree reveals that:

- a) Countries with higher HIV rate, and
- b) Low income per person (30 countries) and
- c) Low employ rate (<65.529) (13 countries) results in:

Only 30.77% have high life expectancy and 69.23% has low life expectancy

IV - The tree also reveals that:

- a) Countries with higher HIV rate, and
- b) Higher income per person (30 countries), and
- c) lower HIV rate (19 countries) results in 100% higher life expectancy.
- d) This is as opposed to countries with higher HIV rate where 35.36% have a higher life expectancy and the rest have a low higher life expectancy.

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V - Confusion Matrix:

The total model correctly classifies those countries with high life expectancy 100%

The confusion matrix reveals the classification tree classifies It correctly classifies countries with low life expectancy 98% of the time (1 - 0.0303 = 0.0.9697 or 96.97%).

Finally, the model variable importance table. Due to the fact that decision trees attempt to maximize correct classification with the simplest tree structure, it's possible for variables that do not necessarily represent primary splits in the model to be of notable importance in the prediction of the target variable.

Potential explanatory variables are highly correlated, or provide similar information, for example: HIVRATE, INCOMEPERPERSON. EMPLOYRATE, SUICIDEPER100TH, FEMALEEMPLOYRATE are likely to be selected for the model. The absence of the alternate variable from the model does not necessarily suggest that it's unimportant, but rather that it's masked by the others.

Appendix 1

```
3/25/2017
                                         Code: W1.sas
  1 /* Machine Learning for Data Analysis */
    /* Running a Classification Tree*/
    /* AB Lopez*/
    LIBNAME mydata "/courses/d1406ae5ba27fe300 " access=readonly;
  4
  5
  6
    DATA new;
  7
        set mydata.gapminder;
  8
  9
         /*life expectancy can be classified in Low, average and High */
         if lifeexpectancy GE 60 THEN
 10
                 LifeE=1;
 11
 12
         /*above 60 high*/
 13
 14
         else
                 LifeE=2;
 15
 16
         /*low*/
 17
 18
    PROC SORT ;
 19
         BY country;
 20
         ods graphics on;
 21
 22 proc hpsplit seed=155311;
 23
         class LifeE country;
 24
         model LifeE=country incomeperperson alcconsumption breastcancerper100th
 25
                 femaleemployrate hivrate suicideper100th employrate urbanrate;
 26
         grow entropy;
 27
         prune costcomplexity;
 28 RUN;
```