DREAM_CANCER_REPORT

TEAM

April 2015

1 Introduction

2 Data preparation

We had to deal with very messy data because of the presence of NA's at high rate and also missing values.

Variable CREACLCA SMOKSTAT CREACL SMOKFREQ TSTAG_DX	Pourcentage_NA 96.75 96.125 94.0625 92.8125 72.3125
GLU REGION_C NA. TBILI NEU TRT3_ID	30.6875 30.125 30.0625 1.4375 1.3125 1.1875
ALT CREAT WEIGHTBL WGTBLCAT ECOG_C	0.3125 0.1875 0.125 0.125 0.0625

The first Work was to complete and clean the data in a way to keep the useful information. We choose to keep SMOKE since this variable is a often kept in studies on other types of cancers.

3 Benchmark Model

On the clean data we apply a first benchmark model wich is the Cox model:

4 Feature selection

Goal:

• Find a sparse representation of the model

- Find new usefull variables
- Express qualitative data in numeric ones in order to use the know model

All along this work we use a criteria to determine the effectiveness of the construction

5 Predictive model for goal 1a

- 5.1 Models considered
- 5.2 Score and comparison
- 6 Predictive model for goal 1b
- 6.1 Models considered
- 6.2 Score and comparison
- 7 Conclusion