

Figure 3. Demonstration of the inconsistency in sample labeling: The labeling described in Table S3a) (Ooi *et al.*) differs from that in the input file (NCI_full_train.txt) online at the author's website

http://www.omniarray.com/bioinformatics/GA/full_data/

Graphical illustration in the next page

Figure 3

Table S3a) Cross validation results of the best predictor set obtained ($E_C = 14.63\%$) for the NCI60 dataset. Each of the values in gray box is the percentage of training samples predicted correctly in cross validation (m_{iq} = number of training samples actually belonging to class q , m'_{iq} = number of training samples classified as class q)

