

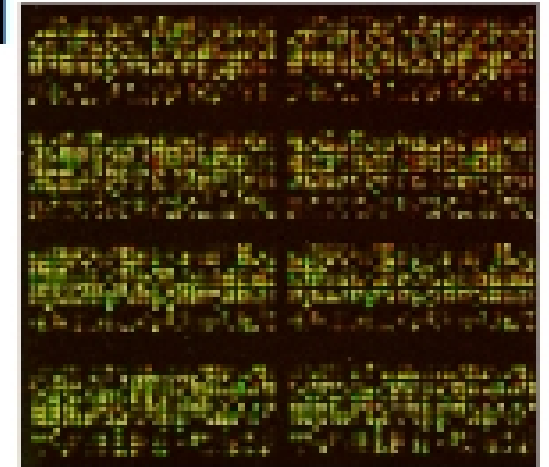
Margin Trees for High-dimensional Classification

Tibshirani and Hastie

Errata (confirmed by Tibshirani)

- Section 2 (a) about the property of 'single linkage'. **M should be M_0**
- Section 2.1 close to the last line of second paragraph. **“at least” should be “at most”**
- The statements about complete/single linkage are misleading. In fact, they use **standard definition** of complete/single linkage except the distance metric is replaced with **margin between pairwise classes**.
(I traced their code to confirm this).

Targeted Problem



- Multi-class
 - #class $\gg 2$
- High-dimensional, few samples
 - #features \gg #data \rightarrow linear separable
 - already good accuracy, need interpretable model
- Ex. micro-array data
 - feature : gene expression measurement
 - class: type of cancer
 - Instances: patients