

Supplementary Materials: Comparison of small n statistical tests of differential expression applied to microarrays Additional File 2

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1 Source Code Availability

- R statistical language - version 2.5.1
(<http://www.r-project.org>)
- MAS 5.0:
bioconductor affy library - version 1.13
(<http://www.bioconductor.org>)
- RMA:
bioconductor affy library - version 1.13
(<http://www.bioconductor.org>)
- gcRMA:
bioconductor affy library - version 1.13
(<http://www.bioconductor.org>)
- dChip:
bioconductor affy library - version 1.13
(<http://www.bioconductor.org>)
- LPE z statistic:
bioconductor LPE library - version 1.10
(<http://www.bioconductor.org>)

- LMGene:
bioconductor LMGene library - version 1.6
(<http://www.bioconductor.org>)
- limma:
bioconductor limma library - 2.10.5
(<http://www.bioconductor.org>)
- BRB t-statistic:
Wright and Simon -BRB Array Tools
(<http://linus.nci.nih.gov/BRB-ArrayTools.html>)
- CyberT t-statistic:
(<http://cybert.microarray.ics.uci.edu/>)