

## Gene Expression Studies Using Affymetrix Microarrays Chapman Hallerc Mathematical And Computational Biology By Hinrich Gohlmann 2009 07 15

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COMMAND Core: Differential Gene Expression Analysis using R

Gene Expression analysis using **R****Gene expression + summary statistics = identify genes for 30 complex traits** *Gene Omnibus Expression (GEO) - NCBI - microarray - affymetrix - expression Differential expression analysis* DNA Microarray

RT-PCR for Gene Expression*Heatmaps course - Introduction DNA Microarray Methodology Microarrays How to Make an R Heatmap with Annotations and Legend Gene Expression Data Microarray Lab Part 1 Gene Expression*

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Results. The present study implements seven methods commonly used in the literature to calculate expression indices from Affymetrix microarray gene expression data, which was collected from a well-designed genome-wide microarray hybridization experiment with eight genetically divergent barley cultivars.

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We made a thorough performance comparison of the three most recent human gene expression chips available from Affymetrix: Human Genome U133 Plus 2.0, Human Exon 1.0 ST and Human Gene 1.0 ST. First, we have compared reproducibility at the probe-, exon- (where possible) and gene-level using publicly available data on technical and biological replicates.

[A comparison of Affymetrix gene expression arrays](#)

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