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Tony N. Frudakis, in Molecular Photofitting, 2008.
RECENT HISTORY OF ASSOCIATION MAPPING RESULTS. Candidate gene analysis is a hypothesis-driven approach, whereby genes thought to influence variability in trait value are selected based on prior relevant research.

[Genetic analysis - Wikipedia](#)

Genetic analysis is the overall process of studying and researching in fields of science that involve genetics and molecular biology. There are a number of applications that are developed from this research, and these are also considered parts of the process.

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[Gene chip analysis - Wikipedia](#)

Proper data processing and quality control are critical to the validity and interpretability of gene chip analysis. Data processing includes data normalization , flagging of the data, averaging the intensity ratio for replicates , clustering of similarly expressed genes, etc. Data also must be normalized before further analysis.

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Transcript abundance is in many ways an extraordinary phenotype, with special attributes that confer particular importance on an understanding of its genetics. The primary transformative potential of genome-wide gene expression genetics is the sheer number of traits thousands that can be assayed simultaneously.

