

and another set in which a lack of objective, or even subjective, knowledge prevents any hypothesis from being put forth. Most of the papers dealing with gene expression based on "gene arrays" present a list of activated or repressed genes only because the majority of links and nodes in the net(work) remain unknown, and discussion would at this stage be premature. This situation, of course, holds true only at present, and progress in knowledge and tools (bioinformatics) will soon make such a comment of historical value only. The authors were also limited by the space available for the discussion, in which only a few genes were referred to.

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