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Collana	Disease markers, , 0278-0240 ; ; v. 17, no. 2
Altri autori (Persone)	StrausbergRobert L
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Nota di contenuto	Cover; Guest-Editorial: Talkin' Omics; Using Serial Analysis of Gene Expression to identify tumor markers and antigens; Cancer proteomics: The state of the art; Analysis of expression patterns: The scope of the problem, the problem of scope; Alternative spliced transcripts as cancer markers; Searching for pharmacogenomic markers: The synergy between omic and hypothesis-driven research; Candidate genes and single nucleotide polymorphisms (SNPs) in the study of human disease; A comparison of gene expression signatures from breast tumors and breast tissue derived cell lines C282Y and H63D mutation frequencies in a population from central Spain ANX7 as a bio-marker in prostate and breast cancer progression
Sommario/riassunto	This work states that we are no longer satisfied to study a gene or gene product in isolation, but rather we strive to view each gene within the complex circuitry of a cell. It states that as a family of diseases, all cancer results from changes in the genome.