

External links Category:Computer programming toolsNIH Uncovers Roles for Small RNA in Regulating Gene Expression John M. Bishop and colleagues at the National Institute of Diabetes and Digestive and Kidney Diseases (NIDDK) have identified four distinct modes of small-RNA-based regulation, which the researchers say may be a new mechanism for controlling gene expression in the body. These modes can be viewed as guidelines that the cell follows to find and interpret small RNAs that target gene-expression control elements and silence those genes. The research, "RNA-Guided Gene Expression Regulation," is published online in the July 16 issue of the journal Cell. The study, the first to elucidate three previously unrecognized small-RNA-guided gene-expression regulatory modes, was led by Peter F. Vogel at the National Cancer Institute, and co-led by Melissa A. Tomes, a graduate student at the Rockefeller University, and Jonathan A. Weissman, a research scientist at the NIDDK. "Our study helps explain the remarkable ability of RNAi to regulate and change the expression of genes at the molecular level," says Bishop, the lead author. RNAi is one form of gene silencing, and is frequently used in the laboratory to study gene function. Gene silencing involves the process in which small RNA molecules, called short interfering RNAs, or siRNAs, bind to specific regions of genes that control their expression, thereby preventing the expression of certain proteins that are encoded by that gene. Before the work by Bishop and colleagues, the study of siRNA-based gene regulation was limited to two modes: forward gene-regulation silencing, by which siRNA targets the gene-expression control element itself; and antisense gene-regulation silencing, in which the gene-expression control element is silenced by the binding of an antisense siRNA to a different, complementary region of the gene. "These two modes are but a few degrees removed from a perfect textbook example of the type of regulation that is seen in nature," says Tomes. "When two complementary complementary binding partners of the same strand of RNA bind, they are locked into an inactive state or, as the case may be, they can potentially unwind the molecule and allow for the production of new RNA, which will be complementary to the other binding partner." Using a method that Bishop and colleagues developed earlier, they



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