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## LATEST **NEWS**

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MOLECULAR BIOLOGY

## 'Evolved' Enzyme Has Unique Action

Technique produced a protein that catalyzes RNA ligation

## Stu Borman

A WAY TO MAKE new types of enzymes has produced one that accelerates a ligation reaction that no other enzyme is known to catalyze.

The approach, which could create enzymes that cleave and modify substrates in addition to linking them together, may aid basic research and speed the synthesis of drugs and other chemicals, among other applications.

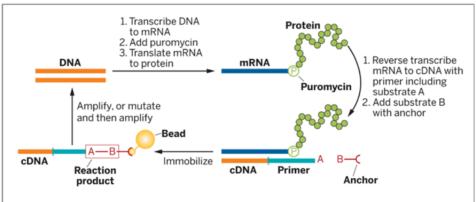
Research fellow Burckhard Seelig and molecular biology professor and Howard Hughes Medical Institute investigator <a href="Jack W. Szostak">Jack W. Szostak</a> of Massachusetts General Hospital, Boston, devised the approach, which is based on messenger RNA (mRNA) display and in vitro selection (*Nature* 2007, 448, 828). mRNA display generates very large libraries of proteins covalently linked to their encoding mRNAs, and in vitro selection is an iterative, cyclic way to screen molecules for desired properties.

Using mRNA display, Seelig and Szostak generated a library of more than 10<sup>12</sup> randomized "zinc finger" proteins, which are known to bind nucleic acids. To each mRNA-protein conjugate, they attached complementary DNA and an RNA substrate. Then they exposed the resulting structures to another RNA substrate bearing an anchor group. When an mRNA-protein conjugate catalyzed ligation of the RNA substrates, it could be isolated because it bound to a bead through the anchor group. The DNA for each "hit" was amplified (or mutated and amplified) and added to the library to enrich it for further selection cycles until an enzyme with unique RNA ligase activity was identified. The new enzyme accelerates the ligation more than 2 million-fold, which is not a tremendously high level of activity. "Biological ligases and polymerases can be 1,000- to 1 million-fold faster" than the new enzyme, Szostak says.

New enzymes generally have been created by modifying existing ones or by designing or "evolving" enzymes based on a detailed understanding of known catalytic mechanisms. The new technique, in contrast, relies only on in vitro evolution and not on known mechanisms.

Chemistry and immunology professor <u>Kim Janda</u> of Scripps Research Institute, La Jolla, Calif., comments that some understanding of the mechanism of ligation reactions is inherent in use of a zinc-containing protein library, since ligation reactions are known to require metal and/or acid-base catalysis. Nevertheless, he says that the new approach "is appealing in that it appears to be a powerful technology for catalyst procurement, yet inexpensive and simple enough for almost any lab to carry out."

In future work, Szostak and coworkers hope to improve the new enzyme's activity, obtain its 3-D structure, and determine its mechanism.



**ENZYME EVOLUTION** Large numbers of mRNA-protein conjugates are produced, and cDNA and an RNA substrate (A) are added. Complexes that catalyze the ligation of two substrates (A and B) are selected by immobilization. DNA for the selected enzymes is then fed into additional cycles.