



Press Release

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AI to Predict the Protein Structure

Determination of the Protein Structure Has Been Difficult and Expensive so Far. KIT Researchers Have Now Developed a More Efficient Method.



Fibronectin plays an important role in wound healing. The figure shows an important part of the protein with contact pairs (spheres of the same color). (Figure: Ines Reinartz, *KIT*)

Proteins are biological high-performance machines. They can be found in every cell and play an important role in human blood coagulation or as main constituents of hairs or muscles. The function of these molecular tools is obvious from their structure. Researchers of Karlsruhe Institute of Technology (KIT) have now developed a new method to predict this protein structure with the help of artificial intelligence.

Depending on their structure, proteins can interact with other molecules by penetrating or enclosing them. This is very difficult to detect, the experiments needed for this purpose are expensive and complex. Researchers of the Steinbuch Centre for Computing (SCC), the computing center of KIT, have searched the databases for protein sequences and compared the same proteins of different species. "Hemoglobin that is responsible for transporting oxygen in our body can also be found in insects, voles, and chimpanzees," says Markus Götz, data analyst of SCC. The protein structure resembles a string of pearls, with the string consisting of the protein components, the amino acids. Its three-dimensional structure and the associated properties



KIT Information · Systems · Technologies Center

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Further materials:

Video showing the unfolding and folding of the fibronectin protein (Animation: Ines Reinartz/KIT): https://www.kit.edu/downloads/proteinfaltung.mp4

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result from some distant "pearls" forming pairs, thus folding the protein. These pairs may differ in different organisms. The properties of the protein, however, remain the same. "Harmful mutations are sorted out in the course of evolution," Götz says.

Now, Götz's research team has taught an artificial intelligence (Al) system which pairs proved to be successful in known protein sequences during evolution. "We expect the system to draw conclusions with respect to the structure of unknown protein sequences as well," Götz says. The benefit: "It is easy to determine the amino acids which form the protein chain. However, it is very complex and costs millions to directly determine protein structures experimentally," Alexander Schug, SCC, adds.

The use of AI to predict contacts in proteins is not new. "Currently, image processing methods are applied for this purpose," Götz says. Such neural networks can recognize patterns well. When determining the protein structure, however, contacts of protein components located far away from each other are of crucial importance, because they have a stronger impact on the structure during folding than those that are located closely to each other. "For this reason, we use an approach from automatic language translation. We consider the amino acid chains sentences that have to be translated into another language." So-called "self-attention neural networks" are applied in popular translation programs. They can identify which parts of the sentence are linked or, in the protein context, which amino acids form a pair.

More about the KIT Information · Systems · Technologies Center: <u>http://www.kcist.kit.edu</u>

Being "The Research University in the Helmholtz Association," KIT creates and imparts knowledge for the society and the environment. It is the objective to make significant contributions to the global challenges in the fields of energy, mobility and information. For this, about 9,300 employees cooperate in a broad range of disciplines in natural sciences, engineering sciences, economics, and the humanities and social sciences. KIT prepares its 25,100 students for responsible tasks in society, industry, and science by offering research-based study programs. Innovation efforts at KIT build a bridge between important scientific findings and their application for the benefit of society, economic prosperity, and the preservation of our natural basis of life.





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This year's **anniversary logo** recalls the milestones reached by KIT and its long tradition in research, teaching, and innovation. On October 1, 2009, KIT was established by the merger of its two predecessor institutions: the Polytechnic School and later University of Karlsruhe was founded in 1825, the Nuclear Reactor Construction and Operation Company and later Karlsruhe Research Center in 1956.