Measuring the effects of drugs on cancer cells
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A new approach established at the University of Zurich sheds light on the effects of anti-cancer drugs and the defense mechanisms of cancer cells. The method makes it possible to quickly test various drugs and treatment combinations at the cellular level.

Cancer cells are cells over which the human body has lost control. The fact that they are transformed body cells makes it all the more difficult to combat them effectively—whatever harms them usually also harms the healthy cells in the body. This is why it is important to find out about the particular weaknesses.

In certain types of breast and ovarian cancer, for example, such weakness is caused by mutations in genes that play a role in DNA repair. Treating cancer cells of this kind with a group of newly approved PARP inhibitors makes it difficult for these cells to replicate their DNA, and they ultimately perish. Normal cells, however, can solve such problems using their intact DNA repair machinery.

The new approach enables researchers to analyze the initial reaction of cancer cells to PARP inhibitors with great precision. What's special about the very sensitive procedure is the high number of individual cells that can be analyzed concurrently with high resolution using the automated microscopes at the Center for Microscopy and Image Analysis of UZH. Cancer cells vary and thus react differently to drugs depending on their mutations and the cell cycle phase they are in. The UZH researchers have now found a way to make these differences visible and quantify them precisely.

Rapid and precise testing of cancer cells

Outside of the laboratory, the success of PARP inhibitors and other cancer medication is complicated by the fact that in some patients the cancer returns—after a certain point, the cancer cells become resistant and no longer respond to the drugs. The high-throughput method employed by UZH researchers is particularly useful for this kind of problem: Cells can be tested in multiple conditions with short turnover times, and specific genes can be eliminated one by one in a targeted manner. Doing so can reveal which cell functions are needed for a certain drug to take effect.
In addition, mechanisms of drug combinations can be analyzed in great detail. In her study, Jone Michelena has already identified such a combination, which inhibits cancer cell proliferation to a significantly higher extent than the combination's individual components by themselves. "We hope that our approach will make the search for strategies to combat cancer even more efficient," says Matthias Altmeyer, head of the research group at the Department of Molecular Mechanisms of Disease at UZH.


Provided by University of Zurich


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