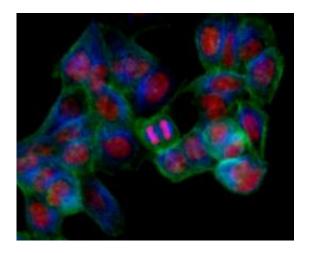


## **Suggesting genes' friends, Facebook-style**

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By silencing genes two at a time in cells like these, the scientists can analyze the genes' combined effects. In this microscopy image of human cells, nuclei are shown in red, cell membranes in green, and the cellular scaffolding in blue. Credit: © Christina Laufer/ DKFZ

Scientists at the European Molecular Biology Laboratory (EMBL) and the German Cancer Research Centre (DKFZ), both in Heidelberg, Germany, have developed a new method that uncovers the combined effects of genes. Published online today in *Nature Methods*, it helps understand how different genes can amplify, cancel out or mask each others' effects, and enables scientists to suggest genes that interfere with each other in much the same manner that facebook suggests friends.

To understand the connections between genetic make-up and traits like disease susceptibility, scientists have been turning to genome-wide



association studies, in which they compare genetic variants of people with a particular disease to those of healthy people. Such studies have linked many <u>genes</u> to diseases, but these links were often weak and not clear-cut, possibly because individual genes often do not act alone. The effects of a particular gene can depend on what other genes a person carries, and the new method developed by the teams of Wolfgang Huber at EMBL and Michael Boutros at DKFZ enables scientists to uncover and measure those combined effects.

The scientists took a set of genes that are important for cell signalling and, using a technique called <u>RNA interference</u>, silenced those genes two at a time, and compared the effect to what happens when you silence only one or the other member of each pair. In so doing, they were able to identify a new component in a cell-signalling process known as the Ras pathway, which is involved in <u>cellular proliferation</u>, and is known to go awry in <u>tumour cells</u>.

If two people have many friends in common on facebook, the odds are that those two people know each other – even if they themselves are not facebook friends. Similarly, genes that have similar genetic interaction profiles are likely to influence each other's effects, and Huber, Boutros and colleagues can now suggest such 'friends' – i.e. genes that are likely to affect the same cellular processes. In the long run, this could help predict patient outcomes and adapt treatments for diseases such as cancer.

## Provided by European Molecular Biology Laboratory

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