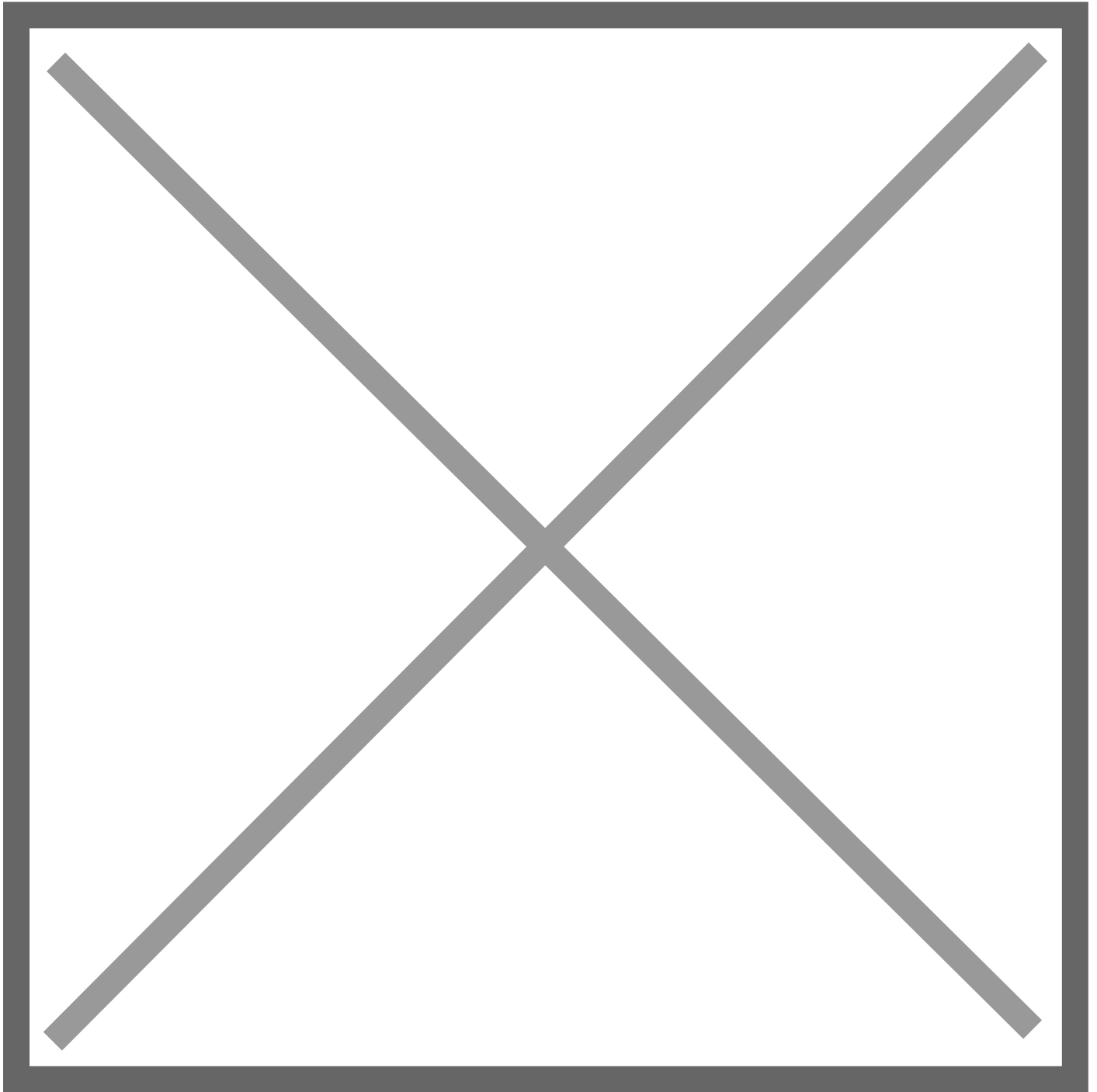


Pfizer Logo

How Human 'Gene Knockouts' Are Pointing the Way to New Medicines

Thursday, April 13, 2017



As more people sequence their genomes, scientists are studying genetic mutations in hopes of identifying new treatments.

Since the human genome was first sequenced 15 years ago, the field of genetic research has exploded, according to Morten Sogaard, vice president and head of Genome Sciences & Technologies at Pfizer, who also oversees Human Genetics, Computational Biomedicine, Computational Target Validation and Diagnostics research. “Drug discovery has always been painstaking, in large part due to the lack of correlation between efficacy in animal models and success in humans. Now, we’re at a point where human genetics has become so robust and

affordable that we can use it as a powerful new tool for drug discovery and development.”

As the cost of sequencing the entire genome of one person has dropped from about \$1 billion to \$1,000, and the speed of sequencing has grown many times faster, scientists across the medical field have been given an entirely new and powerful way of understanding how genetic variation may affect health and disease.

Knockouts Add Punch for Potential New Treatments

Since the 1980s, scientists have studied the role genetics plays in conditions like cancer, heart disease and obesity by inactivating or “knocking out” specific genes in laboratory mice. Now researchers are studying gene knockouts in a new model: humans.

Some people are naturally missing the working copy of a gene, the result of which can range from no effect at all to, occasionally, profound effects, such as very low cholesterol or congenital insensitivity to pain. Such significant, unusual medical characteristics are commonly referred to as “extreme phenotypes.” Advances in gene sequencing now enable scientists to discover the genetic underpinnings of these extreme phenotypes, often finding inactivation of a gene – or a “knockout” – in humans at a scale and speed never before possible.

These extreme phenotypes and their underlying genetic causes represent, in a sense, experiments of nature that point the way toward a new understanding of disease, and potentially reveal powerful new drug targets.

“It’s something we’re increasingly looking at as more individuals have their genomes sequenced,” said Sogaard. “Human gene knockouts could give an indication of what could be a good drug target. It could either tell us, before we start a program, that this looks like it could be a valid mechanism for us to pursue, or it may reveal possible side effects.”

Knockouts Prove Useful

A small segment of the population lacks a working copy of the gene that encodes for ketohexokinase (KHK), an enzyme that is involved in metabolizing fructose, a type of sugar. People with this genetic mutation have no noted health effects. And in a case of gene knockouts helping guide medicine, researchers have been studying the KHK enzyme as a potential target to treat metabolic diseases. Researchers have more confidence to move forward with a KHK-blocker knowing that those who naturally lack the enzyme can live without health complications.

Another example is Janus kinase inhibitors (JAK inhibitors), a class of drugs used to treat cancer and inflammatory diseases like rheumatoid arthritis. JAK inhibitors work by blocking a group of enzymes involved in the cell signaling process that leads to immune and inflammatory responses. Researchers first discovered the importance of these enzymes as a drug target after studying people with an immunodeficiency disorder that had a human knock-out of a JAK gene.

Harnessing the knowledge gained from studying human gene knockouts gives scientists a tool to identify potential new therapeutic targets and better understand safety concerns of therapies in development.

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