

Study advances fight against leading infectious cause of congenital birth defects

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A virus most people probably have never heard of, but that the majority of us carry, is the No. 1 infectious cause of congenital birth defects in the U.S. today. Because of cytomegalovirus (CMV) infection during in utero development, 1 in 750 children are born with or develop permanent disabilities such as hearing loss or brain damage. But efforts to develop a first-ever CMV vaccine are gaining ground.

Researchers from the La Jolla Institute for [Allergy](#) & Immunology and Cardiff University in Wales (UK) have discovered a previously unknown cellular mechanism that could prove critical in creating a CMV vaccine. "We have identified a novel trick that this virus uses to hide from immune detection," says La Jolla Institute scientist Chris Benedict, Ph.D., a CMV expert. "By uncovering this mechanism, we've provided an important piece of the CMV puzzle that could enable vaccine counter strategies that flush out and eliminate virus-infected cells." The finding was published online today in the journal *Cell, Host & Microbe*.

Ed Mocarski, Ph.D., a scientist at the Emory University School of Medicine's Vaccine Center, praised the discovery for identifying some of the key cells involved in the ongoing tug-of-war between CMV and the immune system. "This finding puts on the table the importance of TRAIL signaling in host (our body's) defense and how the virus works to block these efforts," says Dr. Mocarski, whose research focuses on new ways to combat CMV. "This knowledge could set the stage for developing ways to boost the adaptive immune response which could ultimately aid in developing an effective vaccine."

Now listed as a 'high priority' for vaccine development by the National Institute of Medicine, an advisory organization on health issues, cytomegalovirus is very familiar to scientists, but little known by the public. "More children have [disabilities](#) from this disease than other well-known congenital problems, such as Down's syndrome or fetal alcohol syndrome," says Dr. Benedict, who notes that the virus' stealthy nature keeps it under wraps. CMV is part of the herpes family of viruses that cause cold sores, the chicken pox, shingles, mononucleosis, and other maladies, and once contracted it never goes away. Transmission is believed to occur through saliva, urine, and other body fluids, with about 50 percent of Americans infected by age 20, increasing to more than 80 percent by age 80.

Most people are unaware they have contracted the virus, nor generally do they exhibit any visible symptoms although they carry it for life. However, research shows that the immune system has to fight extremely hard to keep CMV at bay. "In addition to causing terrible [birth defects](#) and severe disease in immune compromised patients, CMV induces changes in the immune system in anyone carrying the [virus](#)," says Professor Gavin Wilkinson at the Cardiff University School of Medicine and co-leader of the research team. This constant struggle to control CMV is believed to contribute to tiring the immune system over time.

In their study, the researchers found that a specific CMV protein, known as UL141, blocks the ability of two key immune pathways to kill CMV-infected cells. As their name implies, these two pathways – known as TRAIL "death receptor" 1 and 2 – are normally quite effective at destroying infected cells. In addition, the TRAIL death receptors have been actively targeted in the context of anti-tumor therapies for the last decade, with many phase I and II clinical trials underway or completed.

"While important in cancer, it has also become quite clear that TRAIL signaling by the [immune system](#) is very significant in viral infections,

such as CMV," says Dr. Benedict. "By discovering that this protein (UL141) inhibits TRAIL's ability to carry out its killing function, we believe we have revealed a pivotal piece of the cellular apparatus that needs to be considered when developing an effective vaccine."

As part of exploring CMV, Dr. Benedict also teamed up with La Jolla Institute scientist Dirk Zajonc, Ph.D., a structural immunologist who used X-ray crystallography to determine the three dimensional architecture of the viral protein UL141 bound to the cellular receptor TRAIL-R2 at the atomic level. The results of their study are to be published in another research journal near the end of March.

The TRAIL molecule and its receptors were discovered about 15 years ago and are members of the tumor necrosis factor family of molecules, which have proven extremely important in medical research efforts to control or prevent many diseases. The research team's finding was the first to identify a viral protein as interacting directly with the TRAIL death receptors.

Provided by La Jolla Institute for Allergy and Immunology

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