TSC and PTEN genes important in controlling blood precursor cells in Drosophila

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UCLA stem cell scientists have shown that two common tumor suppressor genes, TSC and PTEN, are vital to regulating the stem cell-like precursor cells that create the blood supply in Drosophila, the common fruit fly.

The researchers examined a signaling pathway called TOR that the cells use to gauge nutrition levels and stress, said study senior author Dr. Julian A. Martinez-Agosto, an assistant professor of human genetics and pediatrics and a researcher with the Eli and Edythe Broad Center of Regenerative Medicine and Stem Cell Research at UCLA.

"We wondered how an organism knows how many blood cells to make and when to make them in the context of injury and repair to tissue. In particular, we wondered how the blood progenitor cells sense that change and know when it's time to make more blood cells," Martinez-Agosto said. "We found that the TOR pathway uses these two genes to regulate its function and, when activated, it expands or increases the number of blood progenitor cells in the fly's blood."

The study appears Sept. 5, 2012 in the advance online issue of the peer-reviewed Journal Development.

Michelle Dragolovic-Munther, a graduate student in the Martinez-Agosto laboratory and first author of the study, found that cells with increased activity of TOR have a competitive advantage, allowing them to divide and make more of themselves so they can make blood. These progenitors, Dragolovic-Munther found, also have high levels of reactive oxygen species (ROS) - ions or very small molecules that include free radicals - which are known to damage cells and can predispose humans to aging and heart disease. But in this case, the ROS proved valuable.

The precursors, Martinez-Agosto said, were producing ROS all the time and when TOR was activated, the levels increased dramatically. Too much ROS caused them to divide more than normal. If they treated the flies with antioxidants, which reduce ROS levels, the cells would develop normally.

The finding could be important because the TOR pathway is abnormally activated in many cancers, and it may be possible to target the levels of ROS, which may help regulate the pathway.

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"What this study may be telling us is that too much ROS is causing more cells to divide and we may be able to target therapies that reduce ROS to significantly improve the condition," Martinez-Agosto said, adding that specifically targeted antioxidants might be a potential treatment in certain subsets of blood disorders. "Sometimes that pathway is working more than it should, and we need the right amount of ROS for balance. It's like Goldilocks, there can be too little or too much. We need it just right."

Going forward, Martinez-Agosto and his team will try to determine where the ROS is coming from and perhaps discover an enzyme that may be a good target for therapeutics. They know that the higher ROS levels in blood progenitors are not coming from mitochondrial, the cells power source, but have not identified how they are being produced.

"This study highlights mechanistic differences between TSC and PTEN on TOR function and demonstrates the multifaceted roles of a nutrient-sensing pathway in orchestrating proliferation and differentiation of myeloid-specific blood progenitors through regulation of ROS levels and the resulting myeloproliferative disorder when deregulated," the study states.

Source: University of California - Los Angeles Health Sciences

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