

Unveiling how lymph nodes regulate immune response

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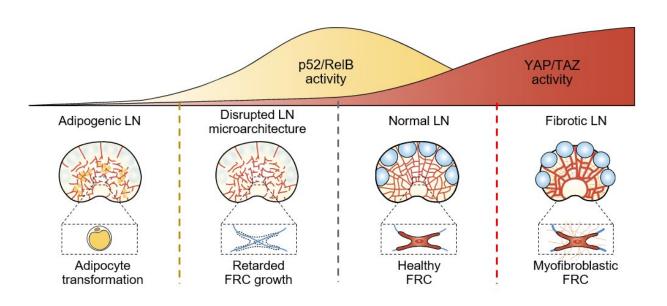


Figure 1: Schematic images proposing the importance of coordination of YAP/TAZ activity and p52/RelB activity during lymph nodes' growth and maintenance. Deletion of YAP/TAZ in fibroblastic reticular cells (FRCs) during development impairs their growth and differentiation, compromising the structural organization of lymph nodes (LNs) and transforming FRCs precursors into fat cells (adipocytes). On the contrary, hyperactivation of YAP/TAZ in FRCs during the developmental period severely impairs FRCs' differentiation and maturation, leaving non-functional and fibrotic LNs. Credit: Institute for Basic Science

Pathogens such as severe acute respiratory syndrome (SARS), Middle



East respiratory syndrome (MERS), and recently the novel coronavirus in Wuhan, China (2019-nCoV) have been a global threat. Lymph nodes (LNs) fight against infectious diseases by providing a shelter for immune cells to grow and launch an attack against pathogens. However, LNs' particular inner workings are poorly understood.

Scientists led by KOH Gou Young at the Center for Vascular Research, within the Institute for Basic Science (IBS), with collaborators of the Korea Advanced Institute of Science and Technology (KAIST), South Korea, have found that a chain of chemical reactions, known as the Hippo-YAP/TAZ signaling pathway, that plays a dominant role in the formation and maintenance of LNs. Their findings have been reported in the journal *Nature Communications*.

One of the key components of LNs are fibroblastic reticular <u>cells</u> (FRCs), which form LN's basic infrastructure and trigger immune responses by releasing cytokines, which are proteins important for immunity. Functional FRCs form during LN's development: a poorly defined population of mesenchymal cells differentiate into FRC precursors, which further develop into mature FRCs with immune functions. Whereas the molecular details involved in the latter process, such as lymphotoxin- β receptor (LT β R) signaling, have been thoroughly described, the details of the commitment steps of FRC development are still unclear.

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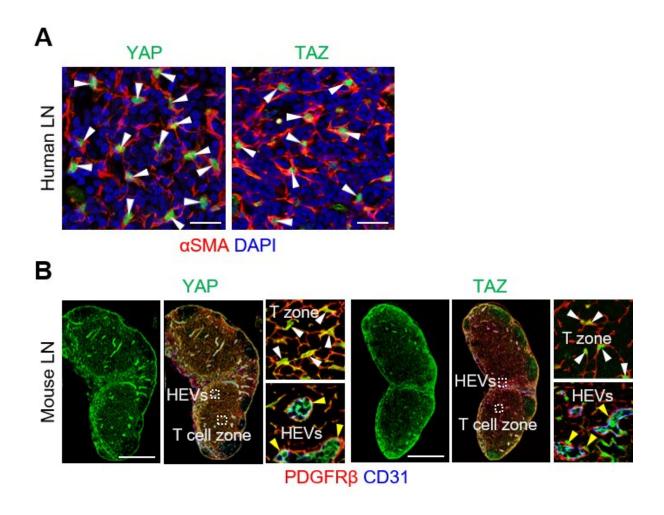


Figure 2: Expression of Yap/Taz in human/mouse FRCs. A shows representative images of YAP and TAZ (green) expression (arrowheads) in FRCs (red) of human cervical lymph nodes. B shows representative images of YAP and TAZ (green) in FRCs (red) in T cell zone (white arrowheads) and around blood venules (high endothelial venules, yellow arrowheads) within the mouse inguinal lymph nodes. Credit: Institute for Basic Science

The research team confirmed the importance of the Hippo pathway—a key regulator of cellular proliferation and organ size control—in FRCs' maturation. The researchers used more than 20 different genetically modified mouse models to characterize the Hippo pathway at specific



time points, depleting the proteins YAP/TAZ at various stages of FRC development.

"As I witnessed the enriched expression of YAP/TAZ in fibroblastic reticular cells of <u>lymph nodes</u>, I knew there must be a role of the Hippo pathway in FRCs," says CHOI Sung Yong, first co-author of this study. By performing a careful examination of the mice's LNs, the team found that FRCs transform into fat cells when YAP/TAZ are reduced in FRC precursors.

BAE Hosung, first co-author of this study, explains, "It was like a mathematic equation, when we drew out the findings on the blackboard, we were sure that depleting YAP/TAZ in fibroblastic reticular cell precursors would show an effect on the lymph nodes."

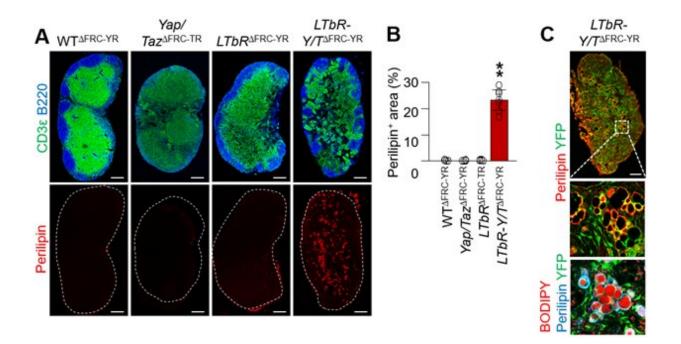


Figure 3: Depletion of Yap/Taz transforms FRC precursors into fat cells. These experiments show YAP/TAZ's role in the differentiation of mesenchymal cell to precursor FRC (first stage) and then in the steps that convert precursor FRC to



mature FRC (second stage). A and B show representative images and comparison of perilipin (protein found in fat cells, red) within the inguinal lymph nodes (LN, dashed-line) at different development phases, from the left: 8-weeks old mouse, YAP/TAZ depleted at the second stage, LTbR depleted at the second stage, and YAP/TAZ depleted at the first stage mice. C shows representative images of inguinal LN filled with fat cells (red) with YAP/TAZ depleted at the second stage. The entire LN is visible on top and a magnified view of the region within the white dashed box below. Credit: Institute for Basic Science

The researchers found that YAP/TAZ binding to p52 is required for maintaining FRC identity. JEONG Sun-Hye, first co-author of this study, notes, "I had this basic instinct that YAP/TAZ should bind with key components that regulate fibroblastic reticular cell identity, such as p52."

Future research will focus on determining whether diseases or conditions that affect systemic immune responses can be linked to alterations in the Hippo signaling pathway in FRCs, and whether modulating Hippo signaling within FRCs could serve as a viable therapeutic option. Beyond their importance in the <u>immune response</u> against flu, FRCs have recently gained considerable recognition for their role in cancer progression and patient outcome. The degree of stromal fibrosis within metastatic LNs is an important prognostic factor that significantly affects disease-free survival of cancer patients. "It definitely warrants more extensive investigation of fibroblastic reticular cells in patients with tumor lymph node metastases prior to clinical investigation," adds Koh.

More information: Sung Yong Choi et al. YAP/TAZ direct commitment and maturation of lymph node fibroblastic reticular cells, *Nature Communications* (2020). DOI: 10.1038/s41467-020-14293-1



Provided by Institute for Basic Science

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