

New metabolic mechanisms discovered that regulate the macrophage's role in immune response

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Integrated metabolic-transcriptional profiling and analysis pipeline. Credit: Cell Press



A group of researchers from Washington University in St. Louis, Agios Pharmaceuticals and ITMO University has discovered new metabolic mechanisms that regulate macrophage polarization - the unique ability of these immune cells to change their specialization depending on the required task. The research opens new possibilities for the development of a new class of drugs based on controlling the metabolism of immune cells. The results were published today in the *Immunity* journal.

One of the key properties of macrophages is their ability to assume different states in accordance with the environmental factors. For example, resident macrophages (M0) can undergo classical activation and transform into proinflammatory macrophages (M1) that by all means seek to destroy pathogens. There exists, however, an alternative activation procedure, wherein macrophages acquire an anti-inflammatory healing function (M2). Understanding these processes is extremely important for understanding the immune system as a whole as well as for fighting autoimmune diseases.

Macrophage <u>polarization</u> is accompanied by changes on both transcriptional and metabolic levels. Activation significantly alters the expression of some genes and the concentration of used and produced compounds. While some aspects of activation are known, until recently, the big picture remained elusive.

To study macrophage polarization, the scientists developed the method of non-targeted analysis of metabolic changes based on data obtained from the simultaneous use of RNA sequencing and high-throughput mass spectrometry of metabolites. The researchers combined the data on the basis of the network of biochemical reactions, which allowed them to identify the most significant of the ongoing reactions.

The final set of reactions revealed unknown metabolic mechanisms associated with macrophage polarization. Based on this information, the



scientists determined the most probable metabolic pathways that manifest themselves in classical and alternative activations. The pathways were confirmed using 13C - 15N isotope labeling and pharmacological inhibition experiments. One of the new mechanisms described in the work is the dependence of alternative macrophage activation on glutamine concentration in the environment, which regulates the activity of TCA cycle, and the synthesis of UDP-Glc-Nac the compound necessary for post-translational protein modification.



Map of macrophage metabolism. Credit: Alexey Sergushichev



"We developed a method of simultaneous analysis of metabolic and transcriptional data. The method proved to be effective and, most importantly, yielded a concrete biological result. Now we continue to improve the method and apply it to other biological tasks such as studying cancer metabolism," says Alexey Sergushichev, PhD student at Computer Technologies Department of ITMO University.

In the long run, the ability to regulate pro- and anti-inflammatory activity can lead to the creation of a new class of drugs based on controlling the metabolism of <u>immune cells</u>. For the first time, macrophage polarization was described in such detail and on such a scale. The work made it possible not only to reveal new regulatory mechanisms of polarization, but also identify and describe the global restructuration of metabolic pathways during polarization. The researchers hope their work will give new impetus to the investigation of pharmacological potential of polarization regulators, which also represents the next step in the collaborative work of the group.

Provided by ITMO University

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