The origins of blood: Researchers identify a gene critical to blood production

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Key findings of this study:
- Rasip1 is involved in the formation of hematopoietic stem cell-containing hematopoietic cell clusters in midgestation mouse embryo.
- Hematopoietic stem cells first arise in Sox17-expressing cells in hematopoietic cell clusters formed in the dorsal aorta at midgestation. Sox17 directly interacts with the Rasip1 gene promoter and induces the Rasip1 gene expression.
- Overexpression of Rasip1 in hematopoietic cluster forming cells led to
maintenance of the clusters with high hematopoietic activity, while knockdown of Rasip1 in Sox17-transduced cells impeded the cluster formation and diminished the hematopoietic ability. Credit: Department of Stem Cell Regulation, TMDU

Blood has long been a symbol of life and health, so it may be surprising that some aspects of blood production (hematopoiesis) remain incompletely understood. One such mystery is the role of a protein called SOX17. Blood cells are generated by hematopoietic stem cells (HSCs), and SOX17 seems to be important to the development of HSCs because SOX17 is expressed where HSCs first develop. What exactly SOX17 does, however, has remained unclear.

Now, a research team at Tokyo Medical and Dental University (TMDU) has discovered just that, revealing that SOX17 targets a gene called Rasip1. To appreciate their research achievement, we first need to know a little more about SOX17 and the origins of blood.

SOX17 is a "transcription factor," a special type of protein that regulates the activities of genes and whether genes are active at any given time. SOX17 is expressed in clusters of cells in a blood vessel called the dorsal aorta; these clusters, called "intra-aortic hematopoietic cell clusters," are where HSCs first develop in mice, around embryonic day 10.5. The researchers aimed to determine SOX17's role in these clusters.

To investigate potential targets of SOX17, the team first carried out RNA-sequencing analysis to see which genes were activated in two very similar populations of cells, one expressing SOX17 and one not.

"One gene that stood out during RNA-sequencing analysis was Rasip1," explains first author Gerel Melig. "This gene is known to be a regulator
in vascular cells, which line the blood vessels."

Significance of this study: Hematopoietic stem cells do not maintained and expanded in vitro. This study will contribute to the development of technology for in vitro proliferation of hematopoietic stem cells in bone marrow. Based on the findings of this study, a new technology to increase the number of hematopoietic stem cells from bone marrow in vitro is expected to be developed. Credit: Department of Stem Cell Regulation, TMDU

The research team therefore investigated Rasip1 further. They showed that SOX17 binds to the enhancer element of the Rasip1 gene to activate it, and then analyzed the effects of both knocking down and
overexpressing the gene. Rasip1 knockdown and subsequent loss of activity resulted in fewer clusters of cells with hematopoietic activity, whereas Rasip1 overexpression increased hematopoietic activity.

"We therefore have proposed a model of early hematopoiesis, in which SOX17 induces the expression of Rasip1, leading to the development of HSCs and associated hematopoietic activity in intra-aortic hematopoietic cell clusters," says senior author Tetsuya Taga.

This study increases our understanding of the early stages of hematopoiesis. This process, forming all cellular components of blood, occurs not only during embryonic development but also throughout adulthood, producing and replenishing the blood cells in the body. A greater understanding of the mechanisms and molecules involved will enhance our knowledge of the processes underlying disorders and cancers of the blood.

The study is published in the journal Inflammation and Regeneration.


Provided by Tokyo Medical and Dental University

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