


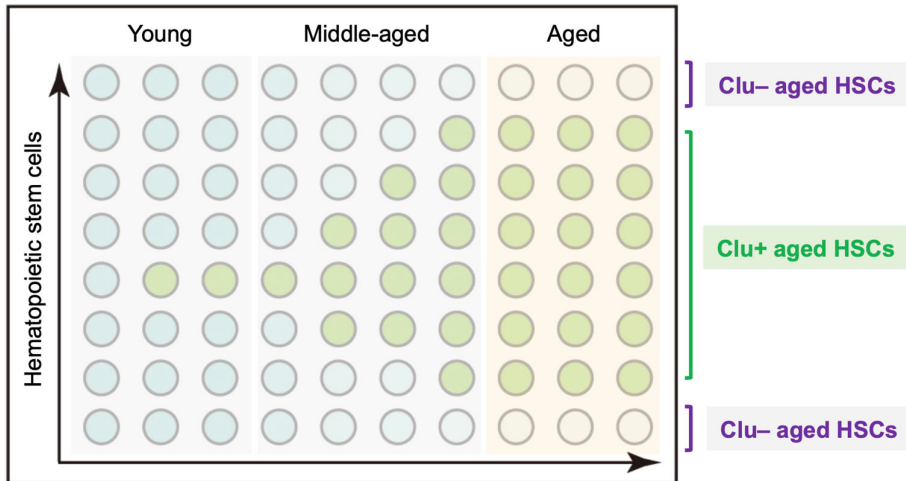
A Study of hematopoietic stem cell aging and its implication in the pathogenesis of hematological malignancies

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Purpose and Background of the Research

● Outline of the Research

- ✓ In the Scientific Research (S) (2019~2023), we identified an hematopoietic stem cell (HSC) aging marker gene *Clusterin* and identified that the reporter mice of *Clusterin*, an aging HSC signature gene, and demonstrated the reporter mice of *Clusterin* identify and prospectively track the two functionally distinct mouse HSC subsets throughout (Figure 1).
- ✓ In this study, using this mouse system, we understand the HSC aging and the heterogeneity of aged HSCs. Furthermore, we identify the stem cell subpopulations, from which clonal hematopoiesis and hematological malignancies originate.
- ✓ We integrate the data from mouse models with the data obtained from human aged HSCs and myeloid malignancies and decipher the role of HSC aging in the pathogenesis of age-associated hematological malignancies.



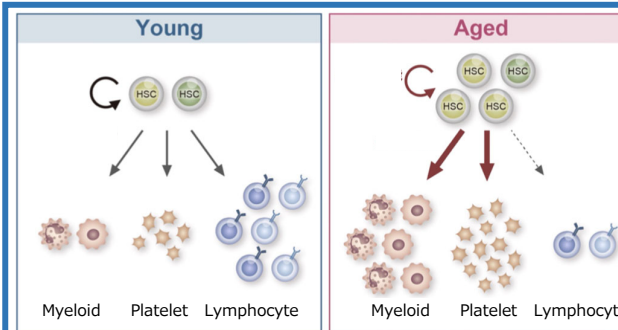
Clu+ aged HSCs
 Myeloid-biased differentiation
 Self-renewal > Differentiation
 Low output of mature cells
 Accumulation of DNA damage
 Mitochondrial dysfunction
 Active inflammatory signals

Clu- aged HSCs
 Balanced differentiation
 High output of lymphocytes
 High output of mature cells
 mild DNA damage
 Retained mitochondrial function
 Suppressed inflammatory signals

→ largely define aged HSC properties

→ Young HSC-like properties

Figure 1. Heterogeneity of aged hematopoietic stem cells



1. Impaired repopulating capacity of hematopoiesis, increase in proportion
2. Myeloid/platelet-biased differentiation, low output of lymphocytes, anemia
3. Dysregulated immune function (Immunosenescence)
4. Clonal hematopoiesis and increased risk of myeloid malignancies

Figure 2. The characteristics of aged HSCs

● Aged HSCs

The following issues remain to be clarified (Figure 2);

1. Their differentiation properties
2. Their heterogeneity
3. The relationship between HSC aging and clonal hematopoiesis by HSCs that acquired somatic driver gene mutations

Expected Research Achievements

● Monitoring of aging process of HSCs

We determine whether Clusterin-negative HSCs become Clusterin-positive HSCs or a small population of Clusterin-positive HSCs expands with aging.

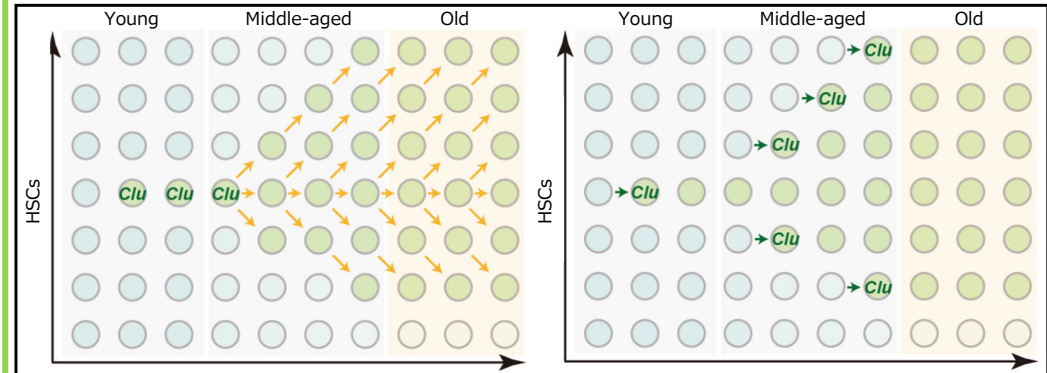


Figure 3. Model of HSC aging process

● Identification of the HSC subpopulations responsible for clonal hematopoiesis

Using *Tet2*-deficient mice, we monitor the process of clonal expansion and transformation of Clusterin-positive and -negative HSCs and determine which HSCs are more prone to transformation.

● Development of intervention approaches to HSC aging

We develop a model mice in which Clu-positive HSCs can be conditionally eliminated (Figure 4). Ablation of Clu-positive HSCs may allow expansion of Clu-negative HSCs, thereby rejuvenating hematopoiesis and preventing the development of clonal hematopoiesis and hematological malignancies.

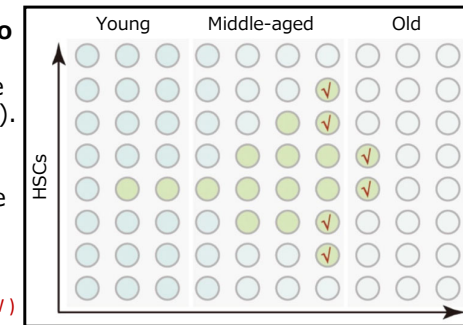


Figure 4. Ablation of Clu-positive HSCs (v)