

D5 Medical & Life Science Seminar (Elective 2 credits)

Academic Year 2018 "International Biomedical Research Seminars"

Insights into the Function of SETD2 and H3K36 Trimethylation in Leukemia, Chemo-resistance and Hematopoietic Stem Cells

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Date: **November 21, 2018 (Wednesday)**

Time: 17:30 –

Venue: IRCMS 1F Meeting Lounge



Abstract

The SET domain containing 2 (*SETD2*) gene encodes the histone 3 lysine 36 (H3K36) methyltransferase which is responsible for tri-methylation (H3K36me₃). Previously, we identified *SETD2* loss-of-function mutations in 22% of MLL-rearranged acute leukemia, implicating a mechanism for cooperativity between *SETD2* mutations and MLL fusions. Indeed, we found a global crosstalk between the oncogenic DOT1L-H3K79me₂ axis and the tumor suppressive SETD2-H3K36me₃ axis in gene regulation. Moreover, recurrence of cooperative mutations in *SETD2* is enriched in refractory/relapsed leukemia. We generated two novel loss-of-function *Setd2* mutation alleles, both of them cooperate with MLL-AF9 to accelerate leukemia development. Importantly AML with *Setd2* mutation allele results in resistance to standard chemo treatment by altered cell-cycle checkpoints. Checkpoint inhibition by either CHK1 or WEE1 inhibitor could re-sensitize resistant AML cells to chemotherapy. Thus, exploring this tumor vulnerability with synthetic lethal approach could be an effective therapeutic strategy to overcome chemo-resistant. Last, *Setd2* knockout in the adult hematopoiesis results in leukopenia, anemia, increased platelet, and erythroid dysplasia in bone marrow. *Setd2* knockout hematopoietic stem cells lose quiescence and self-renewal, but increase differentiation and apoptosis. Mechanistically, we found a critical role of *Setd2* in maintaining the fitness of HSCs and adult hematopoiesis through restricting transcriptional elongation of RNA polymerase II.

◆ Essay / To IRCMS: ircms@jimu.kumamoto-u.ac.jp

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