



## Guest editorial: chronic myeloid leukemia

Yosuke Minami<sup>1</sup>

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The emergence of resistance to molecular target therapies such as ABL-kinase inhibitors in chronic myeloid leukemia (CML) has become a significant problem despite remarkable and successful clinical aspects with treatment-free remission (TFR) [1]. In this issue, Dr. D. Rea and Dr. J.-M. Cayuela summarize clinical results and current issues regarding TFR in patients with CML [2]. The most common cause of resistance in CML is supposed to be the selection of leukemic clones with point mutations in the ABL-kinase domain, and overexpression of target molecules is another reason of resistance [3].

Comprehensive genetic analyses have shaped a deeper understanding of hematologic malignancies. It is also becoming clear that cancer cells display features of normal tissue organization in the microenvironment, where cancer stem cells (CSCs) can drive tumor growth in the tumor environment. It has been proposed that the genetic and CSC models of cancer can be harmonized by considering the role of genetic diversity and tumor heterogeneity [4, 5]. Each clone contains a mixture of cells that vary with respect to their stemness and/or proliferative ability. Standard chemotherapy can reduce tumor burden by eliminating the highly proliferative cells within each subclone, while sparing the relatively dormant cells; following therapy, these cells can seed a renewed cancer. Subclonal diversity can be altered with chemotherapy and can allow for the selection of cells with additional genetic mutations that confer a survival advantage [6]. The concept of leukemia stem cells (LSCs) also becomes critical in understanding the pathogenesis of leukemia. In 1990s, Dick et al. showed that the CD34<sup>+</sup>CD38<sup>-</sup> population in bone marrow cells from acute myelogenous leukemia (AML) patients can reconstitute

human AML in immunodeficient mice, demonstrating clear evidence for the presence of LSCs [7, 8]. Xenogeneic transplantation of leukemia cells into immunodeficient mice has made it possible to identify cell fractions enriched for LSCs, and to characterize an LSC-specific gene expression signature that is strongly correlated with patient survival following standard chemotherapy [9]. There have been steady improvements to the xenograft assay, including development of more immune-deficient recipient mice, better methods for transplantation, and humanizing recipients with human tumor environment and/or growth factors [10].

Persistent disease, especially retention of LSCs, is another therapeutic challenge also in CML. Novel anti-metabolic signal agents and allosteric inhibitors such as Asciminib are under research & development for overcoming resistance and disease persistence [11–13]. In this issue, Dr A Inoue et al. describe the cutting-edge research progress regarding CML stem cells and the further clinical possibilities [14].

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✉ Yosuke Minami  
yominami@east.ncc.go.jp

<sup>1</sup> Department of Hematology, National Cancer Center Hospital East, 6-5-1 Kashiwanoha, Kashiwa 277-8577, Japan

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