LETTER TO THE EDITOR



TP53 mutated AML subclones exhibit engraftment in a humanized bone marrow ossicle mouse model

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Dear Editor,

Despite extensive efforts to develop novel therapies, the prognosis of patients with acute myeloid leukemia (AML) is still poor. One of the reasons is the genetic heterogeneity of AML with the majority of patients exhibiting distinct mutational subclones and diverse biological characteristics [1, 2]. *TP53* mutated AMLs are frequently resistant to intensive treatments and we recently showed that subclonal *TP53* mutations confer an equally devastating prognosis [3]. This observation lead us to hypothesize that *TP53* mutated subclones display characteristics of leukemic stem cells (LSCs) thus contributing to relapse or resistant disease. We, therefore, tested LSC properties of these subclones in a recently developed, highly sensitive humanized bone marrow (BM) ossicle xenotransplantation mouse model [4, 5].

Patient specimens, experimental methods, and ethical approvals are described in detail in the Supplementary Data. Briefly, BM-derived mesenchymal stromal cells (MSCs) from healthy donors were expanded in vitro and subcutaneously injected into four sites of immunodeficient

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NOD/SCID/ γ^{null} (NSG) mice. These MSCs underwent endochondral ossification leading to the formation of a humanized BM ossicle microenvironment. We transplanted three diagnostic, T cell depleted AML specimens into ossicle-bearing NSG mice either by tail vein or direct intraossicle injection. Two specimens showed subclonal TP53 mutations with a variant allele frequency (VAF) of <20%, one a clonal TP53 mutation serving as control (Supplementary Table 1). Mice were sacrificed 16-18 weeks post transplantation and humanized ossicles as well as mouse BM were analyzed for the engraftment of human leukemia. Leukemia engraftment was analyzed by multicolor flow cytometry, and different engrafted cell populations were sorted based on the expression of hCD45, hCD33 and hCD19 (Supplementary Fig. 1). gDNA of sorted cells was analyzed for patient-specific TP53 and cooperating mutations using high-resolution mutation profiling allowing for the detection of mutations with a VAF of <0.1% [6].

All three AML specimens showed engraftment in humanized ossicles as well as mouse BM. However, human

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Fig. 1 Engraftment of TP53 mutated AML specimens in a humanized bone marrow ossicle mouse model. a. Median human engraftment of AML specimens following tail vein injection in humanized bone marrow ossicles and mouse bone marrow of three mice 16 weeks post transplantation. b,c, Engraftment of two AML specimens with subclonal TP53 mutations. The left panels depict human hematopoietic cells being hCD45+ and HLA-ABC+ and sorted into CD33+ myeloid and CD19+ B-lymphoid cells. The right panels depict variant allele frequencies (VAFs) of the particular TP53 and cooperating mutations in the respective compartments. Error bars denote 95% confidence intervals. Abbreviation: BM, bone marrow





cells preferentially engrafted in the humanized ossicle microenvironment (Fig. 1a, Supplementary Fig. 2). For both AML samples with subclonal TP53 mutations, CD33+ leukemic cells constituted the predominant cell population within the graft. Interestingly, differentiation into CD19+ B-lymphoid cells was also observed and the subclonal TP53 mutations could be detected in both, the engrafted myeloid and lymphoid compartments (Figs. 1b, c; Supplementary Figs. 3,4; Supplementary Tables 2,3). These data indicate that subclones with TP53 mutations reveal characteristics of LSCs and pre-leukemic hematopoietic stem cells (pHSCs) being in line with our previous report on clonal TP53 mutations in AML [7]. Stem cell features may contribute to the fact that AML patients with subclonal TP53 mutations do also face an adverse prognosis since LSCs and pHSCs are considered less vulnerable to cytotoxic therapy ultimately giving rise to relapsed or resistant disease [8]. Similarly, expansion of clones with TP53 aberrations was also shown in murine models exposed to genotoxic stress [9, 10]. Finally, our data re-emphasize the usefulness of the humanized BM ossicle mouse model, particularly for the engraftment of small myeloid clones.

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Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

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