

LETTER TO THE EDITOR

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Leukemia cutis with *IDH1*, *DNMT3A* and *NRAS* mutations conferring resistance to venetoclax plus 5-azacytidine in refractory AML

JingHan Wang^{1,2†}, Xingnong Ye^{1,2,3†}, Cuihua Fan^{4†}, Jie Zhou³, Shuna Luo³, Jingxia Jin³, Dan Chen³, Yan Zheng³, Cai Wu³, Xiaoqiong Zhu³, Jie Jin^{1,2,3*}  and Jian Huang^{1,2,3*}

Abstract

Recently, novel drugs like venetoclax plus 5-azacytidine (VA) were reported to have promising efficacy in refractory acute myeloid leukemia (AML). However, there are still some cases presented with novel drugs resistance, and its genetics composition and clinical phenotype are urging to study. We described a 58-year-old patient who was resistant to intensive chemotherapy. This refractory AML was presented with the persistence of *RUNX1*, *IDH1* and *DNMT3A* mutations. *RUNX1* mutations disappeared and leukemia cutis ensued after multiple chemotherapies. Leukemia cutis exhibited *NRAS* mutations in addition to *IDH1* and *DNMT3A* mutations. With the VA salvage treatment, platelets were recovered to the normal level and blasts in bone marrow and peripheral blood were moderately controlled. However, leukemia cutis did not resolve. Unexpectedly, BM blasts obtained the new *NRAS* mutations after VA treatment, and consequently experienced leukostasis with two distinct leukemia clones. After survival of 230 days, this patient died because of spontaneous cerebral hemorrhage. This case highlights presentation of leukemia cutis with simultaneous mutations of *IDH1*, *DNMT3A* and *NRAS* in AML patients might act as a resistant niche to avoid the toxicity of multiple drugs including VA. There is unmet need to validate this result in the clinical trials or a large cohort of patients in the future.

Keywords: Acute myeloid leukemia, BCL-2 inhibitors, Leukemia cutis

To the Editor:

Approximately 30% of newly diagnosed patients with acute myeloid leukemia (AML) do not achieve complete remission with intensive induction therapy, and therefore are classified as refractory or resistant disease (RRD) [1]. RRD is among the most challenging scenarios in AML management. With the

growing clinical translation of genomics into daily routine [2–5], RRD has been becoming an important field for novel drug investigation. Recently, the well-tolerated regimens venetoclax plus 5-azacytidine (VA) were proved to be highly effective in these patients [6]. However, the features related to VA resistance are still under investigation. Here, we present with a RRD patient with a clinical and molecular picture of VA resistance.

The patient is a 58-year-old man with a morphological and immunological diagnosis of AML-M2 (Fig. 1a–b) and a past history of myocardial infarction (MI). His physical examination was

* Correspondence: jiej0503@zju.edu.cn; househuang@zju.edu.cn

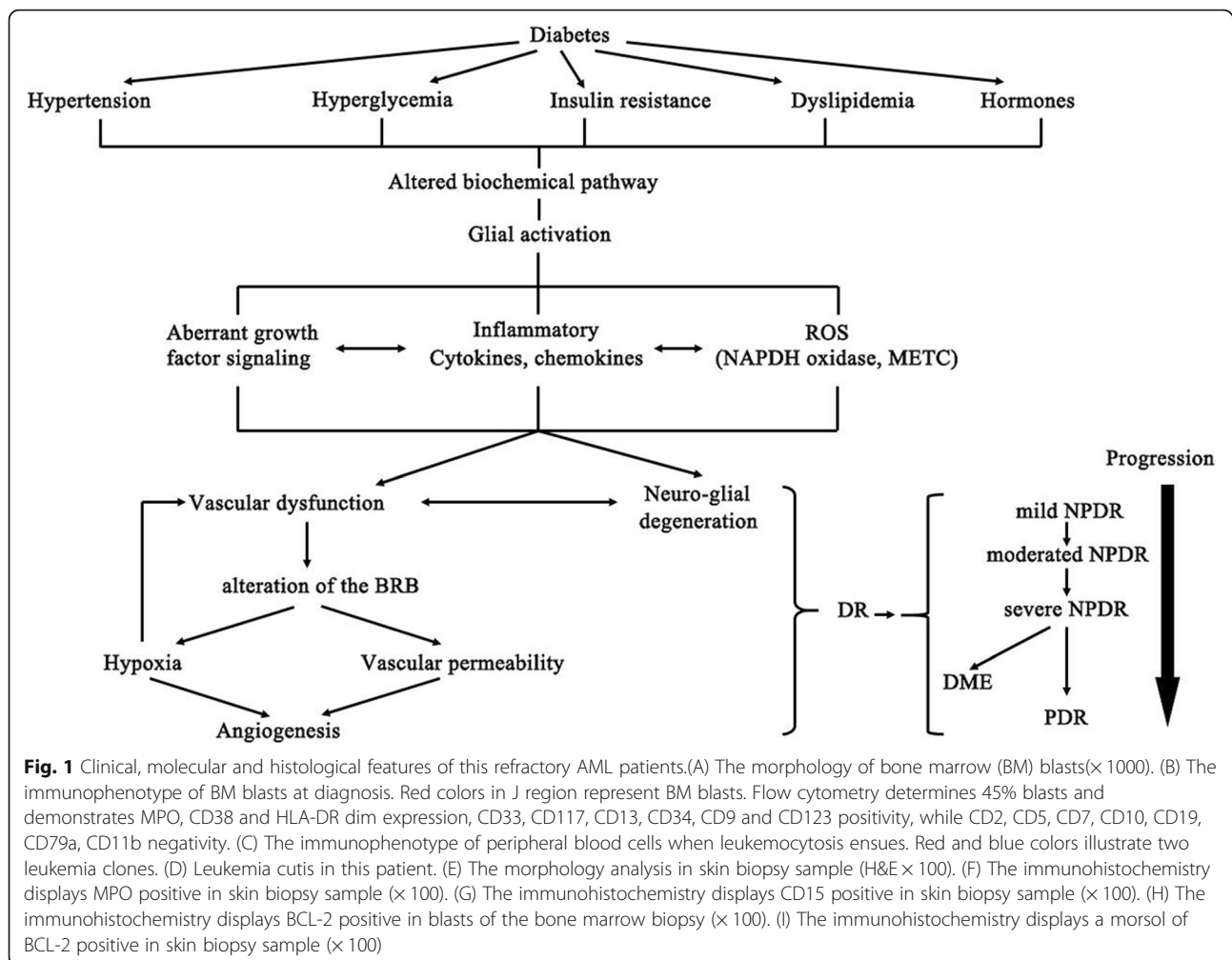
[†]Jing Han Wang, Xingnong Ye and Cuihua Fan contributed equally to this work.

¹Department of Hematology, The First Affiliated Hospital, Zhejiang University College of Medicine, Hangzhou, P.R. China

Full list of author information is available at the end of the article

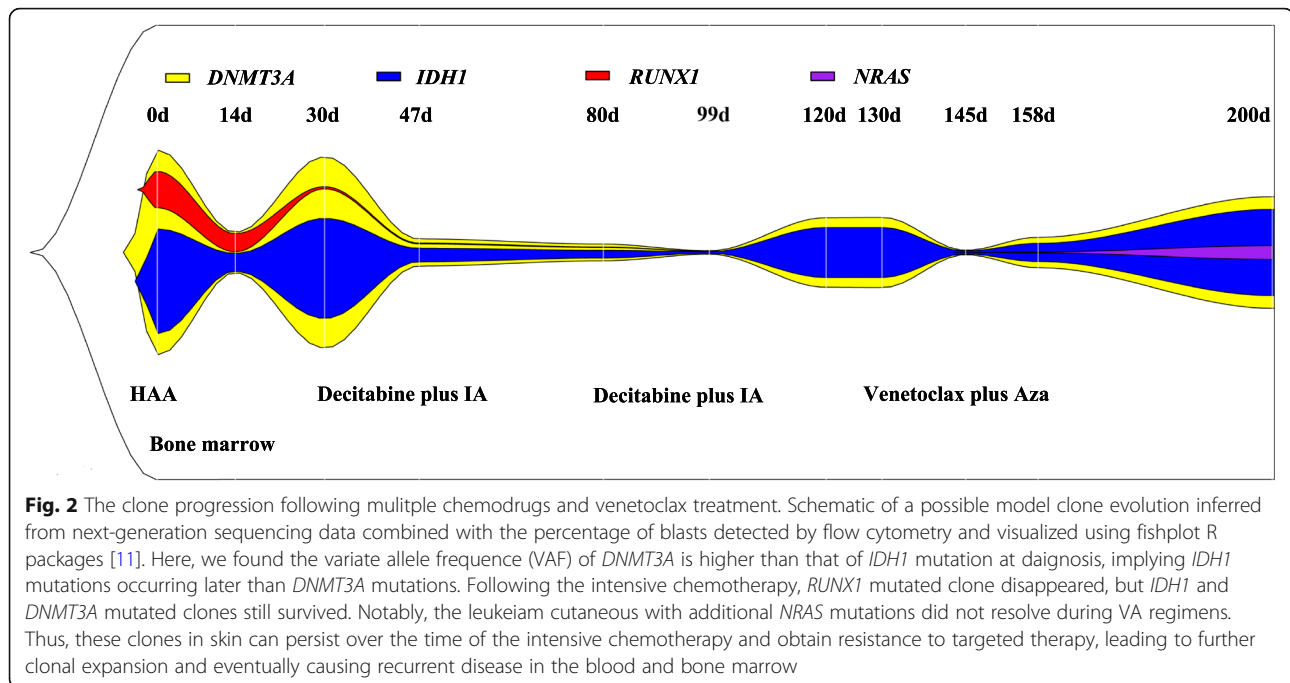


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unremarkable. At the time of diagnosis, the percentage of blasts was 66%, and the karyotype was normal (Figure S1). A peripheral blood test was notable for a substantially leukocytosis with WBC $104 \times 10^9/L$, hemoglobin 104 g/L, and platelets $60 \times 10^9/L$. As shown in treatment flowchart (Figure S2), induction chemotherapy with HAA based regimens (homoharringtonin 2 mg/m² daily for 7 days, cytarabine 100 mg/m² daily for 7 days, aclarubicin 20 mg daily for 7 days) was started [7], but bone marrow (BM) blasts reached 9% on day 15 and surged up to 36% on day 30 indicating poor response (Figure S3). NGS analyses had revealed *IDH1* (exon4:c.394C > G:p.Arg132Gly), *DNMT3A* (exon19:c.2078G > A:p.Arg693His) and *RUNX1* (exon1:c.86 T > C:p.Leu29Ser) mutations (Fig. 2 and Table S1). Based on genetic results, decitabine plus standard IA regimen (decitabine 20 mg/m², days 1–5; idarubicin 10 mg/m² daily for 3 days and cytarabine 100 mg/m² daily for 7 days) were used as the re-induction therapy. About 1 month later,

bone marrow smear revealed a morphological complete remission (CR) with 3% blasts, while platelet was not recovered (Figure S3B). Thus, CR with incomplete platelet recovery was rendered. Therefore, treatment with decitabine plus IA was immediately initiated as a bridge to allogeneic hematopoietic stem cell transplantation. Unfortunately, he began to note skin lesion, although BM blasts were stable for approximately 2 months. After treatment of 130 days, leukemia cells increased up to 20% in the peripheral blood and 6% in bone marrow with a normal karyotype (Figure S1). Physical examination showed numerous dermal gray-blue papules (Fig. 1d). No evidence of leukemia blasts involvement was observed in the Computed Tomography lung screening, hepatic ultrasound, and cerebral Magnetic Resonance Imaging, respectively (Figure S4). Biopsy of the skin lesion demonstrated a dermal infiltration of myeloblast population, which was illustrated by diffuse reactivity for CD15 and MPO immunostains (Fig.



1e-f-g). Notably, NGS demonstrated *NRAS* (exon2: c.38G > A:p.Gly13Asp), *DNMT3A* and *IDH1* (Table S1) mutations coexisting in leukemia cutis, but *RUNX1* negativity and *DNMT3A* and *IDH1* positivity exhibiting in refractory BM samples, which was distinguished with the initially mutated pattern of BM blasts. Putting the leukemia cutis and the chemodrug resistant blasts together, AML refractory disease was definitely diagnosed. As DiNardo et al. reported using ivosidenib (an inhibitor of mutant *IDH1*) to treat *IDH1*-mutated relapsed or refractory AMLs, the median durations of responses were more than 8 months with 30.4% CR rate [8]. The major side effects were differentiation syndrome and prolongation of QT interval. Based on these studies, this patient might not fit for ivosidenib treatment due to the MI history. By contrast, another novel drug venetoclax was also sensitive in *IDH1* mutant primary AML cells with less drug toxicity [9, 10]. Thus, we treated this patient with venetoclax combining with 5-azacytidine (venetoclax 400 mg and intravenous azacitidine 75 mg/m² [days 1–7 of each 28-day cycle]) [6]. At venetoclax initiation, despite WBC up to $20 \times 10^9/L$, we did not observe tumor lysis syndrome. With this regimen, platelets were recovered to the normal level and blasts (4.5 and 6.7% respectively in bone marrow and peripheral blood) were moderately controlled. However, the patient's skin lesion did not resolve during the course of VA treatment. At the survival time of 200 days, WBC increased rapidly

up to more than $50 \times 10^9/L$ and the immunophenotypic data revealed two clonal architecture of neoplasia in the peripheral blood (Fig. 1c). In addition, NGS diagnosis showed the same as its initial mutated genes of *IDH1* and *DNMT3A* were still positive and unexpectedly *NRAS* mutation in blood was incurred after 1 months of VA treatment (Fig. 2 and Table S1). We hypothesized the action of myeloid leukemia clones transferring to skin tissue as a resistant niche to avoid the toxicity of venetoclax. Therefore, we measured the expression of BCL-2 of both BM biopsy (Fig. 1h) and cutaneous blasts (Fig. 1i). As a result, the more and higher expression of BCL-2 was observed in BM biopsy than in leukemia cutis. After survival of 230 days, this patient unfortunately died because of spontaneous cerebral hemorrhage.

Supplementary Information

The online version contains supplementary material available at <https://doi.org/10.1186/s40364-020-00246-9>.

Additional file 1 Figure S1. Normal karyotype was illustrated at the time of AML diagnosis (A), morphological complete remission (B), and leukemia refractory and resistance (C), respectively. **Figure S2.** Treatment flowchart illustrates in this study. **Figure S3.** The morphology of bone marrow (BM) blasts (x 1000) indicated no complete remission on day 30 after HAA treatment (A), a morphological complete remission after treatment with decitabine plus IA (B), leukemia relapse after the second course of decitabine plus IA (C), and moderately control under VA treatment. **Figure S4.** No evidence of leukemia blasts involvement was observed in the Computed Tomography lung screening (A), hepatic

ultrasound(B), and cerebral Magnetic Resonance Imaging(C), respectively.
Table S1. The detailed information of gene mutations.

Abbreviations

VA: Venetoclax plus 5-azacytidine; AML: Acute myeloid leukemia; BM: Bone marrow

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Availability of data and materials

The data analyzed was seen in this text and supplementary data.

Authors' contributions

J.H.W, J.J. and J. H planned the treatment, J.H.W wrote the manuscript, C.H.F, S. L, J.X.J, D. C, Y. Z, C. W, X.Q.Z followed the patient. Y. Z, C. W and J. Z were responsible for histopathology and flow cytometry, J.H.W, J. J and J. H revealed the mutations; and all authors critically read and approved the draft manuscript.

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Ethics approval and consent to participate

The study was approved by the Institutional Review Board of the Fourth Affiliated Hospital of Zhejiang University. This patient was informed about the study and gave an informed consent.

Consent for publication

Consent for publication was provided by this patient.

Competing interests

The authors declare that they have no competing interests.

Author details

¹Department of Hematology, The First Affiliated Hospital, Zhejiang University College of Medicine, Hangzhou, P.R. China. ²Key Laboratory of Hematologic Malignancies, Diagnosis and Treatment, Zhejiang, Hangzhou, PR China. ³Department of Hematology of the Fourth Affiliated Hospital Zhejiang University School of Medicine, Yiwu, P.R. China. ⁴Department of Hematology, Shulan Hospital, Hangzhou, P.R. China.

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