# Inhibition of the mutated c-KIT kinase in AML1-ETO–positive leukemia cells restores sensitivity to PARP inhibitor

Margaret Nieborowska-Skorska,<sup>1</sup> Elisabeth M. Paietta,<sup>2</sup> Ross L. Levine,<sup>3</sup> Hugo F. Fernandez,<sup>4</sup> Martin S. Tallman,<sup>3</sup> Mark R. Litzow,<sup>5</sup> and Tomasz Skorski<sup>1</sup>

<sup>1</sup>Sol Sherry Thrombosis Research Center and Fels Institute for Cancer Research and Molecular Biology, Lewis Katz School of Medicine at Temple University, Philadelphia, PA; <sup>2</sup>Albert Einstein College of Medicine–Montefiore Medical Center, Bronx, NY; <sup>3</sup>Center for Hematologic Malignancies, Memorial Sloan Kettering Cancer Center, New York, NY; <sup>4</sup>Moffitt Malignant Hematology and Cellular Therapy at Memorial Healthcare System, Pembroke Pines, FL; and <sup>5</sup>Division of Hematology, Department of Internal Medicine, Mayo Clinic, Rochester, MN

#### **Key Points**

- c-KIT activating mutations cause resistance to PARP inhibitor in AML1-ETO-positive leukemias.
- c-KIT inhibitor avapritinib downregulates
  BRCA1/2 and DNA-PK
  catalytic subunit to
  restore the sensitivity
  to PARP inhibitor.

## Introduction

Numerous reports indicate that acute myeloid leukemia (AML) cells accumulate high levels of spontaneous and genotoxic agent-induced DNA lesions, but they are able to survive because of enhanced/ altered DNA repair activities.<sup>1-5</sup> Because DNA damage may constrain survival and proliferation of leukemia cells, transformed cells need to be protected from the lethal effects of DNA damage, such as DNA double-strand breaks (DSBs).<sup>6</sup> Thus, leukemia cells may be highly dependent on specific DSB repair mechanisms, and targeting these pathways could sensitize them to DNA-damaging agents.<sup>7</sup>

DSBs, the most lethal DNA lesions, are repaired by 2 major mechanisms: homologous recombination (HR; major proteins: BRCA1, BRCA2, PALB2, RAD51B, RAD51C, RAD51D, XRCC2, XRCC3, RAD54, and RAD51) and DNA-dependent protein kinase (DNA-PK)-mediated nonhomologous endjoining (D-NHEJ; major proteins: DNA-PK catalytic subunit, Ku70, Ku80, NHEJ1, Artemis, LIG4, and XRCC4).<sup>8</sup> PARP1-dependent alternative NHEJ (Alt-NHEJ; major proteins: PARP1 and LIG3) serves as a back-up pathway.<sup>9,10</sup>

Chromosomal translocations involving the core binding factor (CBF) family members, such as AML1-ETO (RUNX1-RUNX1T1) and CBFB-MYH11, are among the most frequent cytogenetic aberrations found in AML.<sup>11</sup> We and other investigators have shown that AML1-ETO-positive cells display BRCA1/2 deficiency, which diminishes HR activity and predisposes leukemia cells to synthetic lethality triggered by DNA repair inhibitors, such as the PARP inhibitor (PARPi) olaparib.<sup>12,13</sup> These data suggested that PARPi's, which are approved by the US Food and Drug Administration for the treatment of BRCA1/2-mutated breast and ovarian cancers,<sup>14</sup> can be used to treat AML1-ETO-positive AMLs.

Additional mutations (eg, in c-KIT and NRAS) often accompany AML1-ETO<sup>+</sup> and CBFB-MYH11<sup>+</sup> AMLs.<sup>15,16</sup> c-KIT mutations (c-KITmuts) in AMLs harboring AML1-ETO or CBFB-MYH11 are associated with poor disease outcome,<sup>17</sup> warranting novel therapeutic approaches. In this study, we tested whether c-KITmuts can modulate the response of AML1-ETO- or CBFB-MYH11-positive AML cells to PARPi.

## **Methods**

#### Primary AML cells and cell lines

Genetic aberrations in diagnostic primary AML samples collected for the Eastern Cooperative Oncology Group and the American College of Radiology Imaging Network (ECOG-ACRIN) E1900 clinical trial<sup>18</sup> are described in supplemental Table 1. Samples of normal hematopoietic cells were purchased from STEMCELL Technologies (Vancouver, BC, Canada). Lin<sup>-</sup>CD34<sup>+</sup> cells were obtained from mononuclear fractions by magnetic sorting using EasySep negative selection human progenitor

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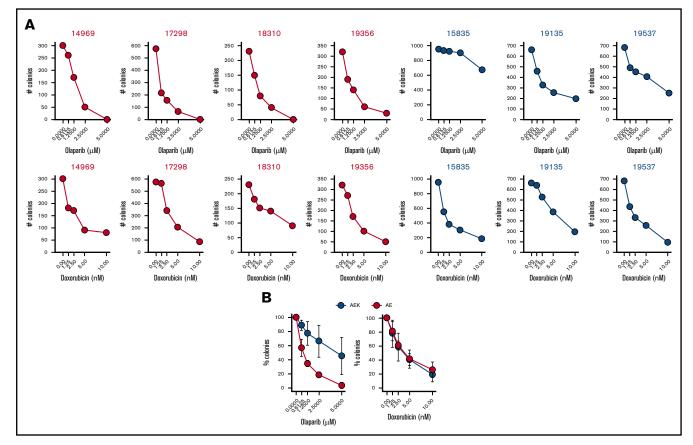


Figure 1. c-KITmut is associated with resistance to the PARPi olaparib, but not to doxorubicin, in AML1-ETO–positive AMLs. Clonogenic potential of Lin<sup>-</sup>CD34<sup>+</sup> cells from AML patients harboring AML1-ETO (AE) or AML1-ETO + c-KITmut (AEK) and treated with the indicated concentrations of olaparib or doxorubicin. (A) Mean number of colonies from individual samples tested in triplicates. (B) Mean percentage  $\pm$  standard deviation of colonies compared with untreated counterparts from the samples harboring the same mutations.

cell enrichment cocktail, followed by human CD34 positive selection cocktail (STEMCELL Technologies), as described previously.<sup>12</sup> The Kasumi-1 AML cell line harboring AML1-ETO + c-KIT(N822K) was purchased from the American Type Culture Collection.

## **Clonogenic assay**

Cells (10<sup>4</sup> per 0.1 mL) were treated with the PARPi olaparib, the c-KIT inhibitor (c-KITi) avapritinib, and/or doxorubicin (all from Selleckchem) for 72 hours, followed by plating in methylcellulose, as described previously.<sup>12</sup> Colonies were counted after 7 to 10 days.

## Western blot

Kasumi-1 cells were left untreated or were treated with the c-KITi avapritinib (5  $\mu$ M) for 48 hours. Total cell lysates and nuclear lysates were examined by western blot, as described previously.<sup>12</sup>

## **DSB** repair

HR, D-NHEJ, and Alt-NHEJ were measured in Kasumi-1 cells that were treated or not with the c-KITi avapritinib (5  $\mu$ M) for 72 hours using DR-GFP (HR), EJ2-GFP (D-NHEJ), and EJ5-GFP (Alt-NHEJ) reporter cassettes, as described previously.<sup>12</sup>

## **Results and discussion**

To test whether a c-KITmut (D816V or  $\Delta$ Y418+D419; supplemental Table 1) modulates the sensitivity of individual AMLs harboring AML1-ETO to PARPi, Lin<sup>-</sup>CD34<sup>+</sup> cells were incubated with increasing concentrations of the PARPi olaparib or the cytotoxic drug doxorubicin, followed by clonogenic testing. Results clearly show that the presence of c-KITmut is accompanied by reduced sensitivity to olaparib but not to doxorubicin (Figure 1). This observation is supported by another report that c-KIT(N822K) rescued BRCA2 expression and HR activity in AML1-ETO-positive cells.<sup>19</sup> In addition, although CBFB-MYH11-positive Lin<sup>-</sup>CD34<sup>+</sup> cells appeared less sensitive to olaparib compared with their AML1-ETO counterparts, c-KITmut further diminished their sensitivity to the drug, whereas mutated NRAS exerted the opposite effect, without affecting their response to doxorubicin (supplemental Figure 1). Altogether, these results suggest that c-KITmut was associated with resistance to the PARPi olaparib without affecting the sensitivity to doxorubicin.

To determine whether constitutive activation of c-KITmut tyrosine kinase was responsible for olaparib resistance, avapritinib was used to inhibit the mutated kinase in AML1-ETO + c-KIT(N822K)-positive

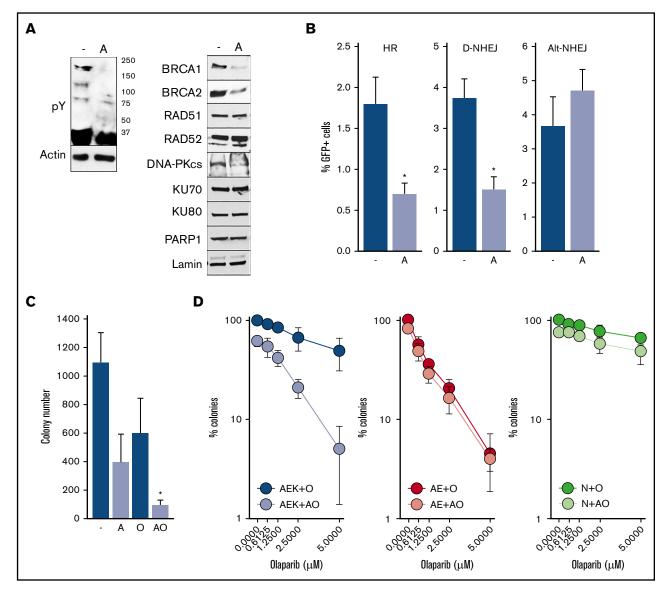


Figure 2. Inhibition of c-KITmut kinase causes DSB repair defects and restores sensitivity of AML1-ETO-positive AML cells to the PARPi olaparib. (A) Tyrosine-phosphorylated proteins (pY) and indicated DSB repair proteins were detected by western blot in total cell lysates (left panel) and nuclear cell lysates (right panel) from Kasumi-1 cells treated (designated by "A") or not (-) with avapritinib. Actin and lamin served as loading controls. (B) HR, D-NHEJ, and Alt-NHEJ activities in Kasumi-1 cells treated (designated by "A") or not (-) with avapritinib. Results represent mean percentage  $\pm$  standard deviation (SD) of GFP<sup>+</sup> cells in DsRed<sup>+</sup> population from 3 experiments. \*P < .001, Student *t* test. (C) Kasumi-1 cells were left untreated (-) or treated with 5-µM avapritinib (designated by "A"), 5-µM olaparib (O), or avapritinib + olaparib (AO). Mean  $\pm$  SD number of colonies from 3 experiments. \*P < .05, Mann-Whitney rank sum test. (D) Clonogenic potential of Lin<sup>-</sup>CD34<sup>+</sup> cells from 3 AML patients harboring AML1-ETO + c-KITmut (AEK), 3 AML1-ETO (AE)-positive AMLs, and from 3 healthy donors (N) treated with the indicated concentrations of olaparib (O) or 5-µM avapritinib + olaparib (AO). Mean percentage  $\pm$  SD of colonies compared with untreated counterparts.

Kasumi-1 cells (Figure 2A, left panel).<sup>20</sup> Inhibition of c-KIT(N822K) kinase by avapritinib (Figure 2A, left panel) was associated with downregulation of BRCA1 and BRCA2 (HR pathway) and the DNA-PK catalytic subunit (D-NHEJ pathway), but not PARP1 (Alt-NHEJ pathway) (Figure 2A, right panel). In concordance, avapritinib inhibited HR and D-NHEJ activity, but not Alt-NHEJ activity, in Kasumi-1 cells (Figure 2B) and restored their sensitivity to olaparib (Figure 2C). Moreover, avapritinib enhanced the sensitivity of primary AML1-ETO + c-KITmut-positive Lin<sup>-</sup>CD34<sup>+</sup> cells to olaparib (Figure 2D, left panel) but did not affect the sensitivity of

AML-ETO-positive cells bearing wild-type cKIT (Figure 2D, middle panel). In addition, avapritinib + olaparib exerted only a modest effect against Lin<sup>-</sup>CD34<sup>+</sup> cells from healthy donors (Figure 2D, right panel). Although it has been reported that avapritinib inhibits mutated cKIT and PDGFRA oncogenic kinases,<sup>20</sup> our results strongly support the role of cKITmut in modulation of the sensitivity of AML cells to PARPi.

In agreement with our observation presented here, oncogenic tyrosine kinase inhibitor-induced dual deficiency in HR and

D-NHEJ was also associated with highly effective elimination of FLT3(ITD)-positive AML cells and JAK2(V617F)-positive myeloproliferative neoplasm cells by PARPi.<sup>21,22</sup> In conclusion, AML1-ETO-positive AML cells harboring c-KITmut (N822K, D816V, and  $\Delta$ Y418+D419) were resistant to the PARPi olaparib but not to doxorubicin. Inhibition of c-KITmut kinase activity by avapritinib inhibited HR and D-NHEJ and restored the sensitivity of AML cells to olaparib. We postulate that PARPi combined with c-KITi can be effective against c-KITmut-positive AMLs, especially those harboring AML1-ETO. Moreover, PARPi may be useful for eliminating CBFB-MYH11 + mutated NRAS-positive AMLs.

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# Authorship

Contribution: M.N.-S. performed experiments and contributed to the writing of the manuscript; E.M.P. provided genetically characterized AML samples from E1900; R.L.L. performed the somatic sequencing studies of E1900 samples; H.F.F. led trial E1900; M.S.T. was ECOG-ACRIN Leukemia Committee chair when E1900 was activated; M.R.L. is the current ECOG-ACRIN Leukemia Committee chair; and T.S. designed the studies, supervised the experiments, and wrote the manuscript.

Conflict-of-interest disclosure: The authors declare no competing financial interests.

ORCID profile: M.R.L., 0000-0002-9816-6302.

Correspondence: Tomasz Skorski, Temple University School of Medicine, Sol Sherry Thrombosis Research Center and Fels Institute for Cancer Research and Molecular Biology, 3400 N Broad St, MRB 548, Philadelphia, PA 19140; e-mail: tskorski@ temple.edu.

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