

Exploiting DNA damage repair defects for effective targeting of acute myeloid leukaemia by PARP inhibitors

Maria Teresa Esposito^{1,5}, Lu Zhao¹, Tsz Kan Fung¹, Jayant K Rane¹, Amanda Wilson¹, Nadine Martin^{2,6}, Jesus Gil², Anskar Y Leung³, Alan Ashworth⁴, Chi Wai Eric So¹

¹Leukemia and Stem cell biology group King's College London, London

²Cell Proliferation group, Medical Research Council Clinical Sciences Centre, Imperial College London

³Department of Medicine, The University of Hong Kong,

⁴University of California, San Francisco (UCSF) Hellen Diller Family Comprehensive Cancer Center

⁵Present address School of Health Sport and Bioscience, University of East London

⁶Present address: Senescence Escape Mechanisms Lab, Centre de Recherche en cancerologie de Lyon

Introduction

Inhibitors of Poly-ADP-Ribose Polymerase (PARPi) have been successfully developed and approved by FDA for treatment of ovarian cancer carrying mutations in DNA damage response (DDR) genes *BRCA1/BRCA2*[1-3]. In Acute Myeloid Leukemia (AML) patients, these mutations are extremely rare. However, chromosomal rearrangements generate chimeric oncofusion proteins [4] that, by acting as transcriptional regulators, impair DDR gene expression[5, 6]. This prompted us to test the efficacy of using PARPi in AML.

Materials and Methods

PARPi were tested *in vitro* and *in vivo*. *In vitro* experiments were carried out in mouse and human leukemic cell lines. Mouse leukemic cells expressing the oncofusion gene of interest were generated by Retroviral Transduction Transformation Assay (RTTA)[7].

Results

Leukemic cells driven by the oncofusion genes, AML1-ETO and PML-RAR α , are sensitive to PARP inhibition whereas cells harbouring MLL-AF9 translocation are resistant[8]. Treatment of AML1-ETO and PML-RAR α leukemic cells with PARPi induces apoptosis, senescence, cell cycle arrest and differentiation *in vitro* and significantly prolongs survival *in vivo*. By using γ H2AX and RAD51 as markers of DNA damage and HR (Homologous Recombination)

we showed that AML1-ETO and PML-RAR α cells accumulate DNA damage and are defective in recruiting RAD51 to DNA damage foci upon PARPi treatment. Further analysis revealed that the expression of a number of genes that are involved in the HR pathway are reduced in AML1-ETO and PML-RAR α cells including *Rad51*, *Brca2* and *Rpa1*. This suggests that AML1-ETO and PML-RAR α are sensitive to PARPi as result of defective DDR. We showed that HOXA9, a key downstream target of MLL-fusions plays a critical role in promoting expression of HR genes and thus providing evidence by which MLL-AF9 are resistant to PARPi[9]. Depletion of *Hoxa9* reduces the expression of *Rad51* and *Brca2* in MLL-AF9 cells and confers PARPi sensitivity in MLL-AF9 leukemic cells, compromising its ability to form colonies, repair DNA damage and prolongs survival in mouse models. Conversely, HOXA9 overexpression rendered AML1-ETO and PML-RAR α cells resistant to PARPi. Likewise, pharmacological suppression of *Hoxa9*, using GSK3 inhibitor LiCl, can also sensitize MLL-AF9 cells to PARPi and prolongs survival in our mouse model.

Discussion

Our data indicate that PARPi might offer a new therapeutic strategy for patients with AML1-ETO or PML-RAR α translocations. More importantly, we showed for the first time that HoxA9 can activate a potential DNA repair back-up pathway. PARPi in combination with pharmacological inhibitors of HOXA9 may represent a novel avenue for tailored therapeutic targeting of the aggressive MLL leukaemia[8].

1. Bryant, H.E., et al., *Specific killing of BRCA2-deficient tumours with inhibitors of poly(ADP-ribose) polymerase*. Nature, 2005. **434**(7035): p. 913-7.
2. Drew, Y., *The development of PARP inhibitors in ovarian cancer: from bench to bedside*. Br J Cancer, 2015. **113 Suppl 1**: p. S3-9.
3. Farmer, H., et al., *Targeting the DNA repair defect in BRCA mutant cells as a therapeutic strategy*. Nature, 2005. **434**(7035): p. 917-21.
4. Zeisig, B.B., et al., *SnapShot: Acute myeloid leukemia*. Cancer Cell, 2012. **22**(5): p. 698-698 e1.
5. Alcalay, M., et al., *Acute myeloid leukemia fusion proteins deregulate genes involved in stem cell maintenance and DNA repair*. J Clin Invest, 2003. **112**(11): p. 1751-61.
6. Esposito, M.T. and C.W. So, *DNA damage accumulation and repair defects in acute myeloid leukemia: implications for pathogenesis, disease progression, and chemotherapy resistance*. Chromosoma, 2014.
7. Zeisig, B.B. and C.W. So, *Retroviral/Lentiviral transduction and transformation assay*. Methods Mol Biol, 2009. **538**: p. 207-29.
8. Esposito, M.T., et al., *Synthetic lethal targeting of oncogenic transcription factors in acute leukemia by PARP inhibitors*. Nat Med, 2015. **21**(12): p. 1481-90.
9. Armstrong, S.A., et al., *MLL translocations specify a distinct gene expression profile that distinguishes a unique leukemia*. Nat Genet, 2002. **30**(1): p. 41-7.

