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In the present study, Byrd et al report the long-term results of acalabrutinib monotherapy in 99 patients with previously untreated CLL (median age, 64 years; unmutated IGHV gene, 62%; and TP53 aberration, 18%.). The starting dose of acalabrutinib was 200 mg once daily (n = 37) or 100 mg twice daily (n = 62), and therapy continued until progression or unacceptable toxicity; twice daily dosing resulted in better trough BTK occupancy and was therefore selected for long-term therapy in all patients. Overall response rate was 97% (complete remissions, 7%); at the median of 53 months, 86% of patients continued to receive therapy, and the 4-year PFS rate was 96%. Acalabrutinib was generally well tolerated. The occurrence of grade \geq 3 side effects of interest appears to be similar to the acalabrutinib arm in the ELEVATE-TN randomized trial: atrial fibrillation developed in 2%, bleeding in 3%, and infections in 15%.

efficacy but lower incidence of typical

side effects associated with ibrutinib.

This study is important for several reasons: first, 100 mg twice daily has been identified as the optimal acalabrutinib dosage for subsequent trials. Second, the results show excellent long-term efficacy of acalabrutinib in untreated CLL, with median follow-up almost twice that reported in the randomized ELEVATE-TN study. Third, it provides crucial data on the safety profile of acalabrutinib, which seems to be favorable compared with that of ibrutinib with regard to the typical off-target side effects (eg, atrial fibrillation or bleeding). Of note, the occurrence of second primary malignancies (SPMs) in this study was relatively high at 26%, compared with 9% in the ELEVATE-TN trial. However, 58% of the malignancies were nonmelanoma skin cancers, which are usually easily cured by surgery; grade \geq 3 SPMs were reported in 5% of patients. Nevertheless, this finding certainly merits further exploration in larger trials, with longer follow-up and vigilance in routine practice. The only additional caveat of the study by Byrd et al appears to be a relatively vague indication of ineligibility for chemoimmunotherapy (CIT). This was defined as a patient's refusal to undergo CIT or the patient's comorbidity profile assessed by an investigator; however, comorbidities and creatinine clearance were not recorded, which makes the fitness and organ function of the study population less clearly defined for the purpose of comparison with other trials. Basic demographic data, efficacy, and safety of acalabrutinib vs ibrutinib in the first-line treatment of CLL are summarized in the table. It needs to be emphasized that, although the overall safety profile of acalabrutinib looks better than that of ibrutinib, it is the result of indirect cross-trial comparison with all of the wellknown limitations, such as different trial populations, length of follow-up, and other factors. Therefore, it is essential to wait for the results of the randomized phase 3 ELEVATE-RR trial directly comparing acalabrutinib vs ibrutinib in previously treated CLL.¹⁰ In summary, Byrd et al have provided important data on the long-term efficacy and safety of acalabrutinib in the first-line therapy of CLL.

Conflict-of-interest disclosure: L.S. has received honoraria and travel grants from and has served on advisory boards of Roche, Janssen, Gilead, AbbVie, and AstraZeneca.

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HEMATOPOIESIS AND STEM CELLS

Comment on Hu et al, page 3339

Associated guilt: radiation/bystanders

Hal E. Broxmeyer | Indiana University School of Medicine

Radiation-induced bystander effects (RIBEs) is a neglected, but crucial, area of radiation response. In this issued of *Blood*, Hu et al¹ have provided important new information and mechanistic insights into RIBE-impairment of hematopoietic stem (HSC) and progenitor (HPC) cells in hematopoietic cell transplantation (HCT), with implications for the mitigation of RIBEs.

Although known, RIBE is not a wellstudied area. However, it has significant consequences after exposure of humans, animals, and cells to radiation.²⁻⁶ Following lymphoid leukemia. N Engl J Med. 2014; 371(3):213-223.

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up on their studies of negative bystander

effects on mouse HSC/HPC,⁷ Hu et al

report the RIBEs on human cells trans-

planted into irradiated and nonirradiated



RIBEs on transplanted human HSCs and HPCs in the context of HCT: knowledge and questions remaining.

NOD/Shi-scid/IL-2Ry^{null} (NOG) mice (see figure). They observed the human transplanted cells in both primary and secondary recipients and demonstrated a significant impairment of HSCs/HPCs in irradiated NOG mouse recipients, using human CD45 chimerism and limiting dilution analysis to calculate the number of SCID-repopulating cells (SRCs; a quantitative measure for human HSCs). They also evaluated how RIBEs influenced the cell cycle, apoptosis, and senescence of human HSCs and HPCs by in vivo and in vitro assessments. They demonstrated that excessive production of reactive oxygen species (ROS) led to HSC/HPC DNA damage associated with upregulation of DNA damage response markers, including ATM, CHK1, CHK2, P53, P16^{INK4a}, P21^{cip1/waf1} and apoptosis-related caspases. They identified increases from bystander cells of interleukin-1 (IL-1), -6, and -8 and tumor necrosis factor- α , and discuss their roles in context of RIBE impairment of functional populations of HSCs and HPCs. Importantly, they were able to apply this information to mitigate RIBEs in vitro and in vivo by using the following antioxidants: N-acetyl-L-cysteine (NAC), sulforaphane (SF), and resveratrol (Res). NAC, SF, and Res each mitigated RIBEs in vitro and in vivo. In vitro, the antioxidants improved HSC-enrichment, but only SF and Res improved engraftment of HPCs. In vivo, engraftment of HSC-enriched cells was improved by the 3 antioxidants, but only Res improved HPC engraftment (see figure). Although more work is clearly needed, the use of select antioxidants, alone or in combination with other reagents, such as epigenetic modifiers, may be effective in dampening RIBEs and improving clinical HCT. Whether this therapeutic approach is efficacious will have to be determined in human HCT trials. The authors also suggest that mitigation of the adverse effects of RIBEs may have implications for patients who undergo radiotherapy.

That radiation has effects on accessory cells for HPCs was noted by us about 45 years ago.⁸ We noted that amounts of stimulation (denoted by colony-stimulating activity, years before isolation, identification, purification, and cloning of many different cytokines/chemokines, and other growth factors), when applied to cell cultures, influenced the apparent death of human colony forming cells (CFCs), induced by increasing doses of ¹³⁷Cs irradiation. These effects were not confined to CFC death, as medium conditioned by cells during in vitro irradiation elicited respective stimulating and inhibitory properties at 600 and 1000 rads. With the current knowledge that there are hundreds of cytokines, chemokines, and other growth-regulating factors, it is time for rigorous identification of the factors produced by accessory cells in the context of RIBEs. Likewise, identification of which accessory cells are involved (eg, monocytes, macrophages, lymphoid cells, other myeloid cell subsets, stromal cells, and other nonhematopoietic cells within the bone marrow microenvironment) is needed (see bottom of figure).

Radiation can be a double-edged sword, having both helpful and detrimental effects.^{9,10} Every day, we are exposed to background levels of radiation, and this exposure is greatly increased once one leaves earth's atmosphere. This is problematic for astronauts and is something I became acutely aware of during the 10 years I served as Chairman of the Board of Scientific Counselors and on the Executive Committee of the National Space Biomedical Research Institute (NSBRI, National Aeronautic and Space Administration). Being able to mitigate not only the direct effects of radiation on sensitive tissues and cells such as HSCs and HPCs, but also RIBEs, is crucial for future health and treatment efforts and is not limited to HCT. There are ongoing efforts to prevent RIBEs by using different mediators,¹¹ in addition to those discussed herein.

Several other factors come to mind when evaluating RIBEs on HSCs, HPCs, and other cells affected by radiation. Most studies entail analysis of cells collected and processed ex vivo in ambient air oxygen (\sim 21% O₂) or cultured in ambient air. Ambient air O_2 levels are not the same as physoxia (lowered O₂ levels present in vivo that are relevant to in vivo physiology). Collecting and processing cells in ambient air results in a phenomenon termed extra physiological shock/stress (EPHOSS).¹² EPHOSS is associated with enhanced mitochondrial ROS, and is linked with P53, opening of the mitochondrial permeability transition pore, ROS, hypoxia inducing factor-1 α , and the hypoximir miR210. Mitigating EPHOSS by collecting and processing cells at a lowered O_2 allows for detection of increased HSCs and decreased slow- cycling HPCs.12,13 Hence, it is appropriate to reevaluate the effects of radiation and RIBEs by collecting/ processing and culturing cells at lowered O₂/physoxia to remove the confounding influences of EPHOSS on collected cells. This approach will provide a more accurate reading of the results and may be especially important for evaluation of the effects of radiation and RIBEs on cells of aged mice¹³ and humans.

Another area that could use reanalysis in this context is that of RIBE-induced cytokines, chemokines, and growth factors and whether these molecules, produced or released in response to radiation, are in a full-length or truncated form. Dipeptidylpeptidase4 (DPP-4) is an enzyme that truncates selected proteins with an alanine, proline, or other amino acid at the penultimate N terminus.¹⁴ Full-length and DPP-4-truncated proteins do not have similar functional activities. In certain situations, DPP4-truncated proteins (such as granulocyte colony-stimulating factor, granulocyte-macrophage colony-stimulating factor, IL-3, TPO, and EPO) have no or less activity than their full-length forms, but can block activities of their full-length forms.14

In their article, Hu et al add to and bring us closer to a more in-depth analysis of the impact of RIBEs on human HSCs and HPCs in the context of HCT. It is clear that there is much more to be learned about health benefits involving radiation (see bottom of figure). This is a virgin field, ready for continued rigorous evaluation in terms of cell responses and in-depth mechanistic insight.

Conflict-of-interest disclosure: H.E.B. is on the scientific advisory board of Elixell Therapeutics.

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LYMPHOID NEOPLASIA

Comment on Domostegui et al, page 3351

Blocking RiBi to suppress MYC lymphomagenesis

Jean-Francois Peyron | INSERM

In this issue of *Blood*, Domostegui and colleagues demonstrate that in MYCdriven B-cell lymphoma, activation of the ribosome biogenesis (RiBi) checkpoint triggers an apoptotic response, through the p53-induced, proteasome-dependent degradation of MCL-1.¹

The MYC bHLH (basic Helix-Loop-Helix) transcription factor may be the most deregulated oncogene in all of human cancers, where it promotes increased cell metabolism and growth, enhanced survival, and abnormal proliferation.² In particular, MYC is a strong stimulator of RiBi and is unique by its simultaneous enhancing effect on the activity of the RNA polymerases pol-I, pol-II, and pol-III. They respectively enhance transcription of the 47S precursor ribosomal RNA (rRNA), of messenger RNAs (mRNAs) for the 80 ribosomal proteins (RPs), and of the 55 rRNA, in order to boost the production of ribosomes.³

MYC is the hallmark oncogene amplified in aggressive diffuse large B-cell lymphoma (DLBCL) and Burkitt lymphoma (BL). Myc-driven lymphomas can be modeled in $E\mu$ -MYC mice, where MYC is overexpressed in B lymphocytes under the control of the strong immunoglobulin heavy chain enhancer. Not surprisingly, it was observed that $E\mu$ -MYC lymphomas are addicted to RiBi and protein synthesis. Indeed, the survival of $E\mu$ -MYC mice was greatly increased by inducing a haploinsufficiency of either of the genes coding for the L24/RPL24 or L38/RPL38 RPs, correcting the enhanced protein