

## Erratum: “Diversity Outbred Mice Identify Population-Based Exposure Thresholds and Genetic Factors that Influence Benzene-Induced Genotoxicity”

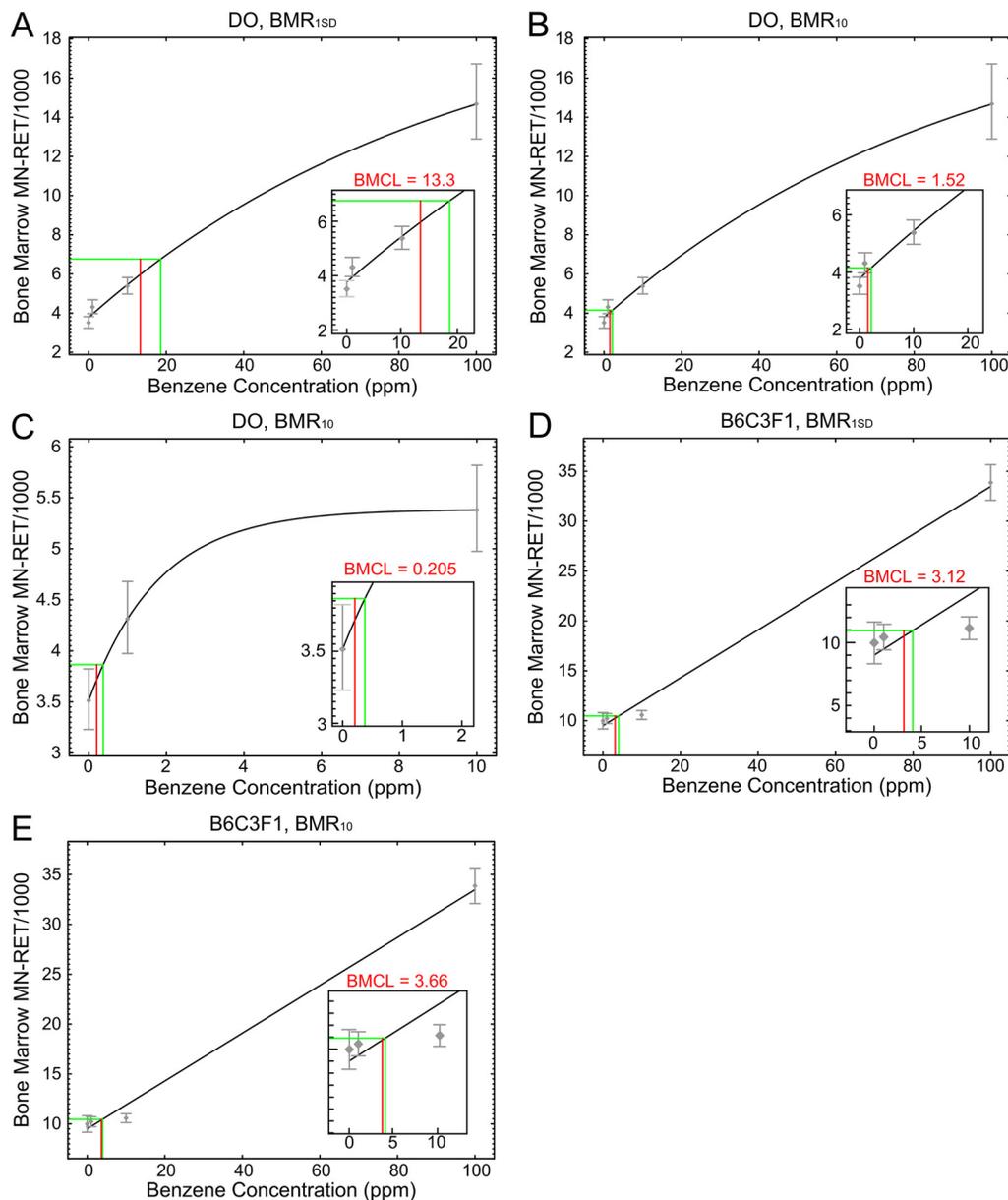
John E. French, Daniel M. Gatti, Daniel L. Morgan, Grace E. Kissling, Keith R. Shockley, Gabriel A. Knudsen, Kim G. Shepard, Herman C. Price, Deborah King, Kristine L. Witt, Lars C. Pedersen, Steven C. Munger, Karen L. Svenson, and Gary A. Churchill

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In the published paper, the *x*-axis of Figure 2C was incorrect; the concentration range should be 0–10 ppm, not 0–100. The corrected figure and legend are presented here.

*EHP* regrets the error.

Also, contact information for corresponding author J.E. French has changed to University of North Carolina Nutrition Research Institute, 500 Laureate Way, Room 3205, Kannapolis, NC 28081. Telephone: 703-250-5029. Email: [jefrench43@email.unc.edu](mailto:jefrench43@email.unc.edu).



**Figure 2.** BMC modeling of bone marrow MN-RET in DO mice using the 0-, 1-, 10-, and 100-ppm exposure groups to estimate a BMR<sub>1SD</sub> (A) and a BMR<sub>10</sub> (B). Values shown are the mean of each exposure group  $\pm$  SE. Curved black lines represent the model fit, green lines represent the BMC, and red lines indicate the BMCL. Insets show details of low concentration range. (C) BMC modeling of bone marrow MN-RET in DO mice using the 0-, 1-, and 10-ppm exposure groups using a BMR<sub>10</sub>. BMC modeling of bone marrow PCE in B6C3F1 mice using a BMR<sub>1SD</sub> (D) and a BMR<sub>10</sub> (E).