IkBα and IkBβ possess injury context–specific functions that uniquely influence hepatic NF-κB induction and inflammation

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Erratum

Original citation: J. Clin. Invest. 113:746–755 (2004). doi:10.1172/JCI17337 Citation for this erratum: J. Clin. Invest. 115:477 (2005). doi:10.1172/JCI17337E1 During preparation of this manuscript for publication, an error was introduced into Figure 5B. The correct figure appears below. We regret this error. Figure 5A: KBI mice have improved survival following I/R injury. KBI or WT littermates either (A) received a lethal dose of LPS (4 μg/g body weight, i.v.) or (B) underwent partial lobar liver I/R. Survival was assessed for 14 days. Only times of survival are given for which deaths occurred. No deaths occurred past the plotted times. Survival curves were significantly different between KBI and WT mice (P < 0.05) as assessed using the log-rank test.

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Erratum

**IkBα and IkBβ possess injury context–specific functions that uniquely influence hepatic NF-κB induction and inflammation**

Chenguang Fan, Qiang Li, Yulong Zhang, Xiaoming Liu, Meihui Luo, Duane Abbott, Weihong Zhou, and John F. Engelhardt


During preparation of this manuscript for publication, an error was introduced into Figure 5B. The correct figure appears below. We regret this error.

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**Erratum**

**Folate pathway gene expression differs in subtypes of acute lymphoblastic leukemia and influences methotrexate pharmacodynamics**

Leo Kager, Meyling Cheok, Wenjian Yang, Gianluigi Zaza, Qing Cheng, John C. Panetta, Ching-Hon Pui, James R. Downing, Mary V. Relling, and William E. Evans


During preparation of this manuscript for publication, errors were introduced into the abstract and introduction. The second sentence of the abstract incorrectly included the phrase “fewer than.” The sentence should read:

“We measured in vivo MTXPG accumulation in leukemia cells from 101 children with acute lymphoblastic leukemia (ALL) and established that B-lineage ALL with either TEL-AML1 or E2A-PBX1 gene fusion, or T-lineage ALL, accumulates significantly lower MTXPG compared with B-lineage ALL without these genetic abnormalities or compared with hyperdiploid (greater than 50 chromosomes) ALL.”

The third sentence of the introduction also incorrectly stated “fewer than.” This sentence should read:

“Subtypes with a relatively unfavorable prognosis on many treatment protocols include T-lineage ALL (T-ALL) and ALL with rearranged MLL genes or with BCR-ABL gene fusion, whereas ALL with either TEL-AML1 or E2A-PBX1 gene fusions, or hyperdiploid karyotypes (greater than 50 chromosomes), have a relatively good prognosis with most treatment protocols . . .”

We regret these errors.