

The Histone Demethylase KDM1A Sustains the Oncogenic Potential of MLL-AF9 Leukemia Stem Cells

William J. Harris,¹ Xu Huang,¹ James T. Lynch,¹ James R. Hitchin,² Yaoyong Li,³ Filippo Ciceri,¹ Julian G. Blaser,¹ Brigit F. Greystoke,¹ Allan M. Jordan,² Donald J. Ogilvie,² and Tim C.P. Somerville^{1,*}

¹Cancer Research UK Leukaemia Biology Laboratory

²Cancer Research UK Drug Discovery Unit

³Cancer Research UK Applied Computational Biology and Bioinformatics Group

Paterson Institute for Cancer Research, University of Manchester, Manchester, M20 4BX, United Kingdom

*Correspondence: tsomerville@picr.man.ac.uk

DOI [10.1016/j.ccr.2012.05.035](https://doi.org/10.1016/j.ccr.2012.05.035)

(Cancer Cell 21, 473–487; April 17, 2012)

Following the publication of this paper, the authors discovered that the list of genes shown in [Table S5](#) as “KIM MYC CORE MODULE HUMAN ORTHOLOGS” was in fact the set of murine genes prior to conversion to their human orthologs. This error does not affect the conclusions, but the authors apologize for the mistake. The online [Table S5](#) has been corrected.