Corrigendum: The landscape of somatic mutations in Down syndrome–related myeloid disorders

Kenichi Yoshida, Tsutomu Toki, Yusuke Okuno, Rika Kanezaki, Yuichi Shiraishi, Aiko Sato-Otsubo, Masashi Sanada, Myoung-ja Park, Kiminori Terui, Hiromichi Suzuki, Ayana Kon, Yasunobu Nagata, Yusuke Sato, RuNan Wang, Norio Shiba, Kenichi Chiba, Hiroko Tanaka, Asahito Hama, Hideki Muramatsu, Daisuke Hasegawa, Kazuhiro Nakamura, Hirokazu Kanegane, Keiko Tsukamoto, Souichi Adachi, Kiyoshi Kawakami, Koji Kato, Ryosei Nishimura, Shai Izraeli, Yasuhide Hayashi, Satoru Miyano, Seiji Kojima, Etsuro Ito & Seishi Ogawa *Nat. Genet.* 45, 1293–1299 (2013); published online 22 September 2013; corrected after print 30 October 2013

In the version of this article initially published, the discussion of cited reference 52 should also have noted that the work “reported accumulation of additional somatic mutations (including single cases of *SMC3* and *EZH2* mutation) during progression from TAM to DS-AMKL.” The error has been corrected in the HTML and PDF versions of the article.