Characterization of transcriptional networks in blood stem and progenitor cells using high-throughput single-cell gene expression analysis

Victoria Moignard, Iain C. Macaulay, Gemma Swiers, Florian Buettner, Judith Schütte, Fernando J. Calero-Nieto, Sarah Kinston, Anagha Joshi, Rebecca Hannah, Fabian J. Theis, Sten Eirik Jacobsen, Marella F. de Bruijn and Berthold Göttgens

Nat. Cell Biol. **15**, 363–372 (2013); published online 24 March 2013; corrected online 2 April 2013

In the version of this Article originally published, Fig. 6d was incorrect; the red label should have read GFI1. This has now been corrected in the HTML and PDF versions.