Zhang F, Michaelson JE, Moshiach S, et al. Tetraspanin CD151 maintains vascular stability by balancing the forces of cell adhesion and cytoskeletal tension. *Blood*. 2011;118(15): 4274-4284.

On page 4278 in the 13 October 2011 issue, there is an error in Figure 2B. An error in the traction-force coding resulted in reported traction forces being low by a factor of $2 \times \pi$. This correction is applied to all traction forces, so all comparisons made in this figure or in the article remain correct and statistically significant. The error does not affect any of the conclusions.

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Gay F, Magarotto V, Crippa C, et al. Bortezomib induction, reduced-intensity transplantation, and lenalidomide consolidation-maintenance for myeloma: updated results. *Blood*. 2013;122(8):1376-1383.

On page 1376 in the 22 August 2013 issue, the affiliation of the 14th and 20th authors (Vittorio Montefusco and Paolo Corradini) is incorrect. The affiliation reads, "Ematologia, Fondazione Istituto di Ricovero e Cura a Carattere Scientifico Istituto Tumori, Milano, Italy." The affiliation should have read, "Division of Hematology, IRCCS Istituto Nazionale dei Tumori Milano, University of Milano, Milano, Italy." The byline and affiliations should have read as shown below:

Francesca Gay,¹ Valeria Magarotto,¹ Claudia Crippa,² Norbert Pescosta,³ Tommasina Guglielmelli,⁴ Federica Cavallo,¹ Sara Pezzatti,⁵ Samantha Ferrari,² Anna Marina Liberati,⁶ Stefania Oliva,¹ Francesca Patriarca,⁷ Massimo Offidani,⁸ Paola Omedé,¹ Vittorio Montefusco,⁹ Maria Teresa Petrucci,¹⁰ Nicola Giuliani,¹¹ Roberto Passera,¹² Giuseppe Pietrantuono,¹³ Mario Boccadoro,¹ Paolo Corradini,⁹ and Antonio Palumbo¹

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Rowley JW, Oler AJ, Tolley ND, et al. Genome-wide RNA-seq analysis of human and mouse platelet transcriptomes. *Blood*. 2011;118(14):e101-e111.

On pages e101, e102, and e103 in the 6 October 2011 issue, there is a typographic error in a URL, which reads, "www.bioserver.hci.utah.edu: 8080/DAS2DB/genopub." The correct URL is "http://bioserver.hci.utah.edu:8080/DAS2DB/genopub." In "Introduction," the fifth sentence of the third paragraph reads, "The sequencing datasets from 2 independent isolations of mouse platelets (C57bl/6, 1 male pool and 1 female pool, 4-8 mice per pool), and platelets from 1 male and 1 female healthy human donor can be accessed at www.bioserver.hci.utah.edu:8080/DAS2DB/genopub (Login: guest. Password: guest. For human data: Homo sapiens > H_sapiens_Feb_2009 > Weyrichlab_RNA-Seq." The sentence should have read, "The sequencing datasets from 2 independent isolations of mouse platelets (C57bl/6, 1 male pool and 1 female pool, 4-8 mice per pool), and platelets from 1 male and 1 female healthy human donor can be accessed at http://bioserver.hci.utah.edu:8080/DAS2DB/genopub (Login: guest. Password: guest. For human data: Homo sapiens > H_sapiens_Feb_2009 > Weyrichlab_RNA-Seq." The sentence should have read, "The sequencing datasets from 2 independent isolations of mouse platelets (C57bl/6, 1 male pool and 1 female pool, 4-8 mice per pool), and platelets from 1 male and 1 female healthy human donor can be accessed at http://bioserver.hci.utah.edu:8080/DAS2DB/genopub (Login: guest. Password: guest. For human data: Homo sapiens > H_sapiens_Feb_2009 > Weyrichlab_RNA-Seq."

Furthermore, in "Methods," the second sentence of the third paragraph under the heading "Sequencing and analysis" reads, "Paired endreads and read coverage files were uploaded onto Genopub (www.bioserver.hci.utah.edu:8080/DAS2DB/genopub) for access and visualization in the Integrated Genome Browser.²⁸" The sentence should have read, "Paired end-reads and read coverage files were uploaded onto Genopub (http://bioserver.hci.utah.edu:8080/DAS2DB/genopub) for access and visualization in the Integrated Genome Browser.²⁸"

Moreover, in "Results," the last sentence of the first paragraph reads, "The datasets can be manually and programmatically accessed through a DAS/2 server (www.bioserver.hci.utah.edu:8080/DAS2DB/genopub; for access instructions, see supplemental data)." The sentence should have read, "The datasets can be manually and programmatically accessed through a DAS/2 server (http://bioserver.hci.utah.edu:8080/DAS2DB/genopub; for access instructions, see supplemental data)."

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