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CORRIGENDA

miR-194-5p/BCLAF1 deregulation in AML tumorigenesis

C Dell'Aversana, C Giorgio, L D'Amato, G Lania, F Matarese, S Saeed, A Di Costanzo, V Belsito Petrizzi, C Ingenito, JHA Martens, I Pallavicini, S Minucci, A Carissimo, HG Stunnenberg and L Altucci

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Correction to: *Leukemia* (2017) **31**, 2315–2325; doi:10.1038/leu.2017.64; published online 10 March 2017

Following the publication of this article, the authors noted a mistake in the author affiliations.

4. Centre for Research in Molecular Medicine, The University of Lahore, Lahore, Pakistan

Should be replaced with:

4. University Institute of Biochemistry and Biotechnology, PMAS Arid Agriculture University, Rawalpindi, Pakistan

The authors apologise for any inconvenience this might have caused.

OPEN

Whole-genome sequencing of chronic lymphocytic leukaemia reveals distinct differences in the mutational landscape between IgHV^{mut} and IgHV^{unmut} subgroups

A Burns, R Alsolami, J Becq, B Stamatopoulos, A Timbs, D Bruce, P Robbe, D Vavoulis, R Clifford, M Cabes, H Dreau, J Taylor, SJL Knight, R Mansson, D Bentley, R Beekman, JI Martín-Subero, E Campo, RS Houlston, KE Ridout and A Schuh

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Since the publication of this article, the authors have identified an omission in their paper, namely that two contributing authors were not included in the author list:

B Stamatopoulos¹, R Clifford²

¹Laboratory of Clinical Cell Therapy, Jules Bordet Institute, Université Libre de Bruxelles (ULB), Brussels, Belgium; ²Department of Haematology, University Hospital Limerick, Limerick, Ireland.

In addition, these authors have equally contributed to the study.

The full and correct author listing is reproduced above. This error has now been rectified and the corrected article appears in this issue together with this corrigendum.

The authors would like to apologise for any inconvenience this may have caused.

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