Leukemia (2008) 22, 889–890
© 2008 Nature Publishing Group All rights reserved 0887-6924/08 $30.00
www.nature.com/leu

CORRIGENDUM

Expression and mutation status of candidate kinases in multiple myeloma

JO Claudio, F Zhan, L Zhuang, R Khaja, YX Zhu, K Sivananthan, S Trudel, E Masih-Khan, R Fonseca, PL Bergsagel, SW Scherer, J Shaughnessy and AK Stewart

Leukemia (2008) **22,** 889–890; doi:10.1038/sj.leu.2405095

Correction to: *Leukemia* (2007) **21,** 1124–1127. doi:10.1038/sj.leu.2404612

The authors of the above paper inadvertently submitted the incorrect supplementary information. They have corrected this information in Supplementary Table 1, which shows the complete list of 280 probe IDs representing 185 unique kinases as presented in the heat map of Figure 2 below.

The authors apologize for any inconvenience caused.

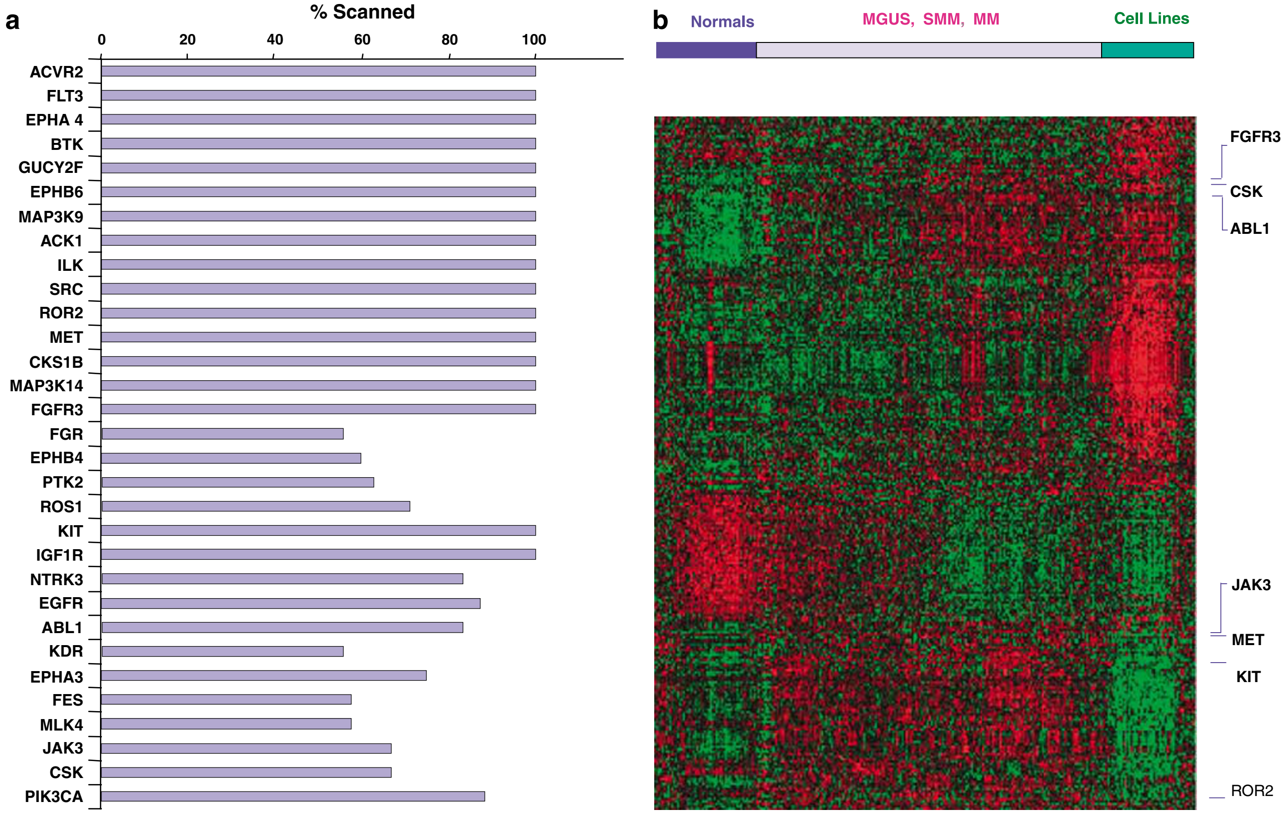

Figure 2 (**a**) Sequence screening of 31 genes in a panel of 32 myeloma cell lines. The graph shows the percentage of kinase domains screened in the 31 kinases. (**b**) Heat map showing the groups of normals, primary myeloma samples and cell lines when expression of kinase genes was used for cluster analysis. (**c**) Nucleotide sequence changes in FGFR3, EGFR, EPHB4 and GUCY2F showing changes in the boxed sequence.

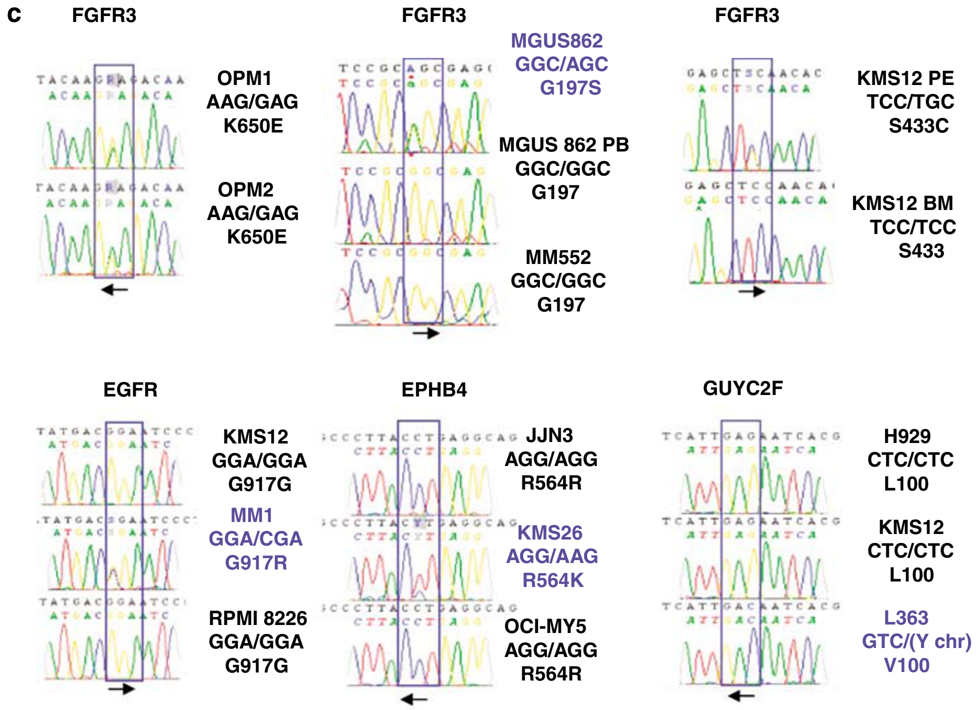

Figure 2 Continued.