

DOI: 10.1038/ncomms10203

Corrigendum: Genome-wide association study identifies variants at 16p13 associated with survival in multiple myeloma patients

Elad Ziv, Eric Dean, Donglei Hu, Alessandro Martino, Daniel Serie, Karen Curtin, Daniele Campa, Blake Aftab, Paige Bracci, Gabriele Buda, Yi Zhao, Jennifer Caswell-Jin, Robert Diasio, Charles Dumontet, Marek Dudziński, Laura Fejerman, Alexandra Greenberg, Scott Huntsman, Krzysztof Jamroziak, Artur Jurczyszyn, Shaji Kumar, Djordje Atanackovic, Martha Glenn, Lisa A. Cannon-Albright, Brandt Jones, Adam Lee, Herlander Marques, Thomas Martin, Joaquin Martinez-Lopez, Vincent Rajkumar, Juan Sainz, Annette Juul Vangsted, Marzena Watek, Jeffrey Wolf, Susan Slager, Nicola J. Camp, Federico Canzian & Celine Vachon

Nature Communications 6:7539 doi: 10.1038/ncomms8539 (2015); Published 22 Jul 2015; Updated 9 Dec 2015

In this Article, members of the UCSF cohort who had been alive for longer than two years were inadvertently included in the data presented in Table 3. USCF/old treatments should have 109 patients with a hazard ratio of 3.35 and a *P* value of 0.00028 instead of the 124 patients with a hazard ratio of 3.37 and a *P* value of 0.00026. The USCF/new patients should have 187 patients with a hazard ratio of 3.57 and a *P* value of 0.0007 instead of the 208 patients with a hazard ratio of 3.62 and a *P* value of 0.0006. Finally, in the table legend, the first line should read 'All models are adjusted for age, gender and principal components 1–3'. The exclusion of these individuals does not change the conclusions of the study. The correct version of Table 3 appears below.

	HR*	95% CI	P value [†]
Mayo Clinic			
Old treatments [‡]		N = 136, 102 deaths	
RS72773978	1.90	0.98-3.83	0.057
New treatments§		N = 93, 64 deaths	
RS72773978	2.71	1.56-4.70	0.00045
Entire sample adjusted for treatment		N = 229, 166 deaths	
RS72773978	2.18	1.43-3.32	0.00028
UCSF			
Old treatments [‡]		N = 109 60 deaths	
RS72773978	3.35	1.74-6.44	0.00028
New treatments§		N=187, 30 deaths	
RS72773978	3.57	1.71-7.43	0.0007
Entire sample adjusted for treatment		N = 296, 90 deaths	
RS72773978	3.35	2.07-5.41	$8.2 \times 10^{-}$

§Treatments containing at least one of the following agents: thalidomide, botezomib or lenalidomide. ‡Regimens including vincristine/adriamycin/dexamethasone or melphalan/prednisone.