

Dominant Mutations in KBTBD13, a Member of the BTB/Kelch Family, Cause Nemaline Myopathy with Cores

Nyamkhisig Sambuughin,^{1,*} Kyle S. Yau,² Montse Olivé,³ Rachael M. Duff,² Munkhuu Bayarsaikhan,¹ Shajia Lu,⁴ Laura Gonzalez-Mera,³ Padma Sivadorai,⁵ Kristen J. Nowak,² Gianina Ravenscroft,² Frank L. Mastaglia,⁶ Kathryn N. North,⁷ Biljana Ilkovski,⁷ Hannie Kremer,⁸ Martin Lammens,⁹ Baziel G.M. van Engelen,⁹ Vicki Fabian,⁵ Phillipa Lamont,¹⁰ Mark R. Davis,⁵ Nigel G. Laing,² and Lev G. Goldfarb¹¹

(The American Journal of Human Genetics 87, 842–847; November 24, 2010)

Dr. Kremer's affiliation information was incorrect in the original publication. The corrected affiliation list appears here. The authors apologize for this error.

¹Department of Anesthesiology, Uniformed Services University, Bethesda, MD 20814, USA; ²Centre for Medical Research, University of Western Australia, Western Australian Institute for Medical Research, QEII Medical Centre, Western Australia 6009, Australia; ³Institut de Neuropatologia, Department of Pathology and Neuromuscular Unit, Department of Neurology, IDIBELL-Hospital de Bellvitge and CIBERNED, Feixa Llargà s/n, Hospitalet de Llobregat, Barcelona 08907, Spain; ⁴National Institute of Arthritis, Musculoskeletal and Skin Diseases, National Institutes of Health, Bethesda, MD 20892, USA; ⁵Department of Anatomical Pathology, Royal Perth Hospital, Perth, Western Australia 6000, Australia; ⁶Centre for Neuromuscular and Neurological Disorders, University of Western Australia 6009, Australia; ⁷Institute for Neuroscience and Muscle Research, The Children's Hospital at Westmead, Westmead 2145, Australia; ⁸Department of Human Genetics and Department of Otorhinolaryngology, Radboud University Nijmegen Medical Centre, Nijmegen 6500, The Netherlands; ⁹Department of Neurology, Radboud University Nijmegen Medical Centre, Nijmegen 6500, The Netherlands; ¹⁰Division of Neurosciences, Royal Perth Hospital, Perth, Western Australia 6000, Australia; ¹¹National Institute of Neurological Disorders and Stroke, National Institutes of Health, Bethesda, MD 20892, USA

*Correspondence: nsambuughin@usuhs.mil

DOI 10.1016/j.ajhg.2010.12.013. ©2011 by The American Society of Human Genetics. All rights reserved.

To Identify Associations with Rare Variants, Just WHaIT: Weighted Haplotype and Imputation-Based Tests

Yun Li,^{*} Andrea E. Byrnes, and Mingyao Li

(The American Journal of Human Genetics 87, 728–735; November 12, 2010)

On page 729, the third formula on the left panel should be:

$$S_h = 1/\sqrt{N_{ct} \cdot f_{ct,h} \cdot (1 - f_{ct,h})}$$

The formula on the right panel should be:

$$WDS_i = \sum_{j=1}^M I(j \in \mathbf{M}_C) \cdot (-1)^{I(j \in \mathbf{M}_P)} \cdot W_j \cdot D_{i,j}$$

The authors apologize for the errors.

*Correspondence: yunli@med.unc.edu

DOI 10.1016/j.ajhg.2010.12.014. ©2011 by The American Society of Human Genetics. All rights reserved.
