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

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

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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:

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

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.

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