

A mutation in the α tropomyosin gene *TPM3* associated with autosomal dominant nemaline myopathy NEM1

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Nature Genetics 9, 75–79 (1995)

The primer (5'-CTTGTCTAACTTCAGCATAGGC-3') stated to introduce an *Nsi*I site into the normal *TPM3* sequence is in fact a primer which introduces a *Stu*I site into the mutant *TPM3* sequence. This works very well for detecting the mutation since incomplete digestion does not give a control pattern. The correct primer for introducing a *Nsi*I site into the normal *TPM3* sequence is:

5'-CTTGTCTAACTTCAGCATATG-3'.

Stability of an expanded trinucleotide repeat in the androgen receptor gene in transgenic mice

Peter M. Bingham, Marion O. Scott, Suping Wang, Michael J. McPhaul, Elizabeth M. Wilson, James Y. Garbern, Diane E. Merry & Kenneth H. Fischbeck
Nature Genetics 9, 191–196 (1995)

On page 192 in the second paragraph under the cross heading "AR transgene expression", Figure 3*a* should read Figure 3*b*, and Figure 3*b* should read Figure 3*a*. (The figure legend correctly refers to *a* and *b*.)

Also, the authors unintentionally omitted the following credit in the Acknowledgements section: Injection of transgene fragments and implantation of ova was carried out by the Transgenic Core Facility of the University of Pennsylvania's School of Medicine.

Inherited microdeletions in the Angelman and Prader-Willi syndromes define an imprinting centre on human chromosome 15

Karen Buiting, Shinji Saitoh, Stephanie Gross, Bärbel Bittrich, Stuart Schwartz, Robert D. Nicholls & Bernhard Horsthemke

Nature Genetics 9, 395–400 (1995)

In the legend to Fig. 3, the last sentence should read: The patients lack the paternal band, whereas the father of family U and the paternal grandmother of family S lack the maternal band.