European Journal of Human Genetics (2013) 21, 1031–1034

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CORRIGENDA

Next-generation sequencing (NGS) as a diagnostic tool for retinal degeneration reveals a much higher detection rate in early-onset disease

Morag E Shanks, Susan M Downes, Richard R Copley, Stefano Lise, John Broxholme, Karl AZ Hudspith, Alexandra Kwasniewska, Wayne IL Davies, Mark W Hankins, Emily R Packham, Penny Clouston, Anneke Seller, Andrew OM Wilkie, Jenny C Taylor, Jiannis Ragoussis and Andrea H Németh

European Journal of Human Genetics (2013) **21,** 1031; doi:10.1038/ejhg.2013.91

Correction to: *European Journal of Human Genetics* (2013) **21,** 274-280; doi:10.1038/ejhg.2012.172

The authors would like to apologise for the following errors in Table 2.

Three references cited in Table 2 were absent from the final reference list:

ABCA4 N965S

Sun H, Smallwood PM, Nathans J: Biochemical defects in ABCR protein variants associated with human retinopathies. *Nat Genet* 2000; **26:** 242–246.

CRX Y258X

Chau KY, Chen S, Zack DJ, Ono SJ: Functional domains of the cone-rod homeobox (CRX) transcription factor. *J Biol Chem* 2000; **275:** 37264–37270.

GUCY2D R838H

Wilkie SE, Newbold RJ, Deery E *et al*: Functional characterization of missense mutations at codon 838 in retinal guanylate cyclase correlates with disease severity in patients with autosomal dominant cone-rod dystrophy. *Hum Mol Genet* 2000; **9:** 3065–3073.

The dbSNP entry rs61749449 refers to N965D rather than N965S.

CRB1T, 745K should read CRB1, T745K.

Also, the following errors were found in Supplementary Table 2:

ABCA4 c.322C>T should read c.3322C>T

ABCA4 c.5882G>C should read c.5882G>A

ABCA4 c.6379C>T should read c.6391G>A

PRPF8 c.6926T>G should read c.6926A>C

Mortality in neurofibromatosis 1: in North West England: an assessment of actuarial survival in a region of the UK since 1989

D Gareth R Evans, Catherine O'Hara, Anna Wilding, Sarah L Ingham, Elizabeth Howard, John Dawson, Anthony Moran, Vilka Scott-Kitching, Felicity Holt and Susan M Huson

European Journal of Human Genetics (2013) **21,** 1031; doi:10.1038/ejhg.2013.121

Correction to: *European Journal of Human Genetics* (2011) **19,** 1187–1191; doi:10.1038/ejhg.2011.113

The authors would like to apologize for the following error that has been brought to their attention.

In the first row of Table 4 the PMR (95% CI) should be 4.1 (2.4, 6.4) rather than 4.1 (2.4, 2.6).

Also, the first sentence on page 1189 should read:

'...cardiovascular disease was reported four times more frequently in NF1 males (PMR = 4.1; 95% CI, 2.4–6.4) than the general population.'