## CORRIGENDA

# 22 Years of predictive testing for Huntington's disease: the experience of the UK Huntington's Prediction Consortium 

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European Journal of Human Genetics (2017) 25, 1290; doi:10.1038/ejhg.2017.139
Correction to: European Journal of Human Genetics (2016) 24, The error is that the division by 18.8 is missing. 1396-1402; doi:10.1038/ejhg.2016.36; published online 11 May 2016 The authors would like to apologise for their error.

The formula on page 2 should read: $4.2 \times$ prevalence $\times p o p(1994)$

$$
+\sum_{y=1995}^{2014} 4.2 \times \text { prevalence } / 18.8 \times \operatorname{pop}(y)
$$

## Glaucoma spectrum and age-related prevalence of individuals with FOXC1 and PITX2 variants

Emmanuelle Souzeau, Owen M Siggs, Tiger Zhou, Anna Galanopoulos, Trevor Hodson, Deepa Taranath, Richard A Mills, John Landers, John Pater, James E Smith, James E Elder, Julian L Rait, Paul Giles, Vivek Phakey, Sandra E Staffieri, Lisa S Kearns, Andrew Dubowsky, David A Mackey, Alex W Hewitt, Jonathan B Ruddle, Kathryn P Burdon and Jamie E Craig

European Journal of Human Genetics (2017) 25, 1290; doi:10.1038/ejhg.2017.147

Correction to: European Journal of Human Genetics (2017) 25, 839-847; doi:10.1038/ejhg.2017.59; published online 3 May 2017

Post publication, the authors realised that the genomic coordinates for two cases were incorrect in their paper.

Table 2, case 19: the genomic coordinates should be g. 110618453 _ 110618454del and not g.110618315_110618316del. Table 2, case 23: the genomic coordinates should be g.(110505201_ 110607760)_(110967245_111069815)del and not g.(111426357_ 111528916)_(111888401_111990971)del.

