CORRIGENDA

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

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Correction to: *European Journal of Human Genetics* (2016) **24,** 1396–1402; doi:10.1038/ejhg.2016.36; published online 11 May 2016

The formula on page 2 should read:

$$4.2 \times prevalence \times pop(1994)$$
$$+ \sum_{y=1995}^{2014} 4.2 \times prevalence / 18.8 \times pop(y).$$

The error is that the division by 18.8 is missing. The authors would like to apologise for their error.

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

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.