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## Corrigendum: A genome-wide association study identifies susceptibility loci for Wilms tumor

Clare Turnbull, Elizabeth R Perdeaux, David Pernet, Arlene Naranjo, Anthony Renwick, Sheila Seal, Rosa Maria Munoz-Xicola, Sandra Hanks, Ingrid Slade, Anna Zachariou, Margaret Warren-Perry, Elise Ruark, Mary Gerrard, Juliet Hale, Martin Hewitt, Janice Kohler, Sheila Lane, Gill Levitt, Mabrook Madi, Bruce Morland, Veronica Neeffjes, James Nicholson, Susan Picton, Barry Pizer, Milind Ronghe, Michael Stevens, Heidi Traunecker, Charles A Stiller, Kathy Pritchard-Jones, Jeffrey Dome, Paul Grundy & Nazneen Rahman

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In the version of this article initially published, the following statement was omitted from the Acknowledgments: “We acknowledge use of data from the database of Genotypes and Phenotypes (dbGaP) from the NCBI, US National Library of Medicine. Research support to collect data and develop an application to support High Density SNP Association Analysis of Melanoma (phs000187) was provided by grants 3P50CA093459, 5P50CA097007, 5R01ES011740 and 5R01CA133996 from the US National Institutes of Health. This study also used, in part, data from the National Institute of Neurological Disorders and Stroke (NINDS) dbGaP database from the CIDR:NGRC Parkinson’s Disease Study (phs000196).” The error has been corrected in the HTML and PDF versions of the article.

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## Corrigendum: Mutations in *TUBG1*, *DYNC1H1*, *KIF5C* and *KIF2A* cause malformations of cortical development and microcephaly

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In the version of this article initially published, the label “– Tub” in Figure 3d was placed incorrectly. The label should appear adjacent to the lower bracketed panels rather than adjacent to the bottom portion of the upper bracketed panels. The error has been corrected in the HTML and PDF versions of the article.