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## ERRATUM

# Erratum: Long-range massively parallel mate pair sequencing detects distinct mutations and similar patterns of structural mutability in two breast cancer cell lines

Oliver A. Hampton, Maxim Koriabine, Christopher A. Miller, Cristian Coarfa, Jian Li, Petra Den Hollander, Caroline Schoenherr, Lucia Carbone, Mikhail Nefedov, Boudewijn F.H. Ten Hallers, Adrian V. Lee, Pieter J. De Jong, Aleksandar Milosavljevic. Long-range massively parallel mate pair sequencing detects distinct mutations and similar patterns of structural mutability in two breast cancer cell lines. *Cancer Genetics* 2011;204:447–457.

The article was published omitting two authors: Cristian Coarfa, *Department of Molecular and Human Genetics, Baylor College of Medicine, Houston, TX*; and Caroline Schoenherr, *Lester and Sue Smith Breast Center, Baylor College of Medicine, Houston, TX*.

We apologize for any confusion or inconvenience this may have caused.

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