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## corrigendum

# Sequence and comparative analysis of the chicken genome provide unique perspectives on vertebrate evolution

**International Chicken Genome Sequencing Consortium**

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In Table 5 of this Article, the last four values listed in the ‘Copy number’ column were incorrect. These should be: LTR elements, 30,000; DNA transposons, 20,000; simple repeats, 140,000; and satellites, 4,000. These errors do not affect any of the conclusions in our paper. □