CORRIGENDUM

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Corrigendum: Elephant shark genome provides unique insights into gnathostome evolution

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The ranges of accession numbers JW861113–JW881738 and KA353634–KA353668 (20,661 in total) cited in this Article include 3,630 sequences that were unrelated and hence removed from the NCBI GenBank database before the analysis of the data and submission of the manuscript to *Nature*. None of those sequences were used or are cited anywhere in the Article. The sequences that were used in the analysis and cited in the Article are available under BioProject ID PRJNA168475.