CORRIGENDUM

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Clinical characteristics and the identification of novel mutations of *COL1A1* and *COL1A2* in 61 Chinese patients with osteogenesis imperfecta

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Following the publication of this article, an interested reader drew to our attention that, in Table IV, we describe a c.1081C>T, p.Arg361X mutation in collagen type I, alpha 2 (*COL1A2*). Codon 361 is a glycine residue, not an arginine, according to the transcript, Z74616.1. We re-examined the original data, and identified that the mutation c.1081C>T, p.Arg361X was of the collagen type I, alpha 1 *COL1A1* gene, and furthermore, it was not novel in the mutation spectrum of *COL1A1*. Therefore, in the paper, the c.1081C>T, p.Arg361X mutation should have appeared in Table III (not Table IV), which described the clinical and genetic characteristics of probands with mutations in *COL1A1*, with an indication in the final column of the Table that it was not novel in the mutation spectrum of *COL1A1*.

This error did not affect the major conclusions drawn in this study. We sincerely apologize for this mistake, and thank the reader of our article who drew this matter to our attention.