Contrasting evolution of diversity at two disease-associated chicken genes

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Unfortunately, an incorrect version of Fig. 2a has been published. Please find the correct version below.

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Fig. 2a Median-joining networks of chicken population haplotypes for IL1B. Populations are denoted in the legend. Branch lengths are proportional to the number of sequence differences between haplotypes. The outgroup samples are represented by the colourless nodes.

V represents the green JF sequences; *F* the grey francolin; *B* the bamboo partridge; *G* the grey JF; *C* the Ceylon JF; *R* the red JF and *R/JF* the genome sequence. There are no differences between the genome sequence and red JF sample