known about such mechanisms, the model proposed here is simple and potentially useful in analysing the complicated forces concerning extranuclear genomic evolution.

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ERRATUM

In Takahata & Slatkin (*Genetical Research* 1983, **42**, 257–265), two sentences should be read as:

but, in contrast, the configuration of the genome and the mode of transmission both make a large difference (on page 257, lines 11–12 up)

Contribution no. 1528 from the National Institute of Genetics, Mishima, Shizuoka-ken 411, Japan (on page 264, above REFERENCES).

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