



Erratum: Genomic analysis of ERVWE2 locus in patients with Multiple sclerosis: absence of genetic association but potential role of Human Endogenous retrovirus type W elements in molecular mimicry with myelin antigen

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A commentary on

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by do Olival, G. S., Faria, T. S., Nali, L. H. S., de Oliveira, A. C. P., Casseb, J., Vidal, J. E., Cavenaghi, V. B., Tilbery, C. P., Moraes, L., Fink, M. C. S., Sumita, L. M., Perron, H., and Romano, C.

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The correct sequences of the primers described in this work are HW_MSChX_F-TGGGTGAAGTAAGTCCAACAG and HW_MSChX_R-TGAAGAACGTATCCAG CCTACA. The amplified fragment corresponds to nucleotides 21227–22122 of the human chromosome Xq22.3 (GenBank ID AL390039.10).

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