

## Erratum

### **Demarcation of ilarviruses based on the phylogeny of RNA2-encoded RdRp and a generic ramped annealing RT-PCR**

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The original version of this paper unfortunately contained a mistake on page 1690, second column, first paragraph, primer concentrations were given incorrectly.

The correct sentence (starting in first column last paragraph) is:

“The first RT-PCR was carried out with 1 µl of RNA in a final volume of 25 µl containing 10 mM Tris–HCl [pH 8.8], 50 mM KCl, 1.5 mM MgCl<sub>2</sub>, 0.1% Triton X-100, 0.25 mM of each deoxyribonucleoside triphosphate (dNTP), 5.0 mM DTT, 5% dimethyl sulphoxide (DMSO), 12 units of ribonuclease inhibitor (HT Biotechnology, England), 0.7 units of Superscript<sup>TM</sup> II Rnase H-Reverse Transcriptase (Invitrogen, The Netherlands), 0.7 units of Avian Myeloblastosis Virus Reverse Transcriptase (Finnzymes, Finland), 1.5 units of *Taq* Platinum DNA Polymerase (Invitrogen, The Netherlands) **and 1 µM each of the primers** “Ilapol up2” and “Ilapol do4”. Nested PCR reaction (20 µl) was performed using 1 µl of the first RT-PCR product. The reaction mixture contained 10 mM Tris-HCl [pH 8.8], 50 mM KCl, 1.5 mM MgCl<sub>2</sub>, 0.1% Triton X-100, 0.2 mM of each dNTP, 1 unit of *Taq* Platinum DNA Polymerase **and 1 µM each of** “Ilapolnest up3” and “Ilapolnest d5” primers.”

Authors apologize for the error.