

AUTHOR'S CORRECTION

Several Groups among Human Herpesvirus 6 Strains Can Be Distinguished by Southern Blotting and Polymerase Chain Reaction

JEAN-THIERRY AUBIN, HELENE COLLANDRE, DANIEL CANDOTTI, DIDIER INGRAND,
CHRISTINE ROUZIOUX, MARIANNE BURGARD, STEPHANE RICHARD,
JEAN-MARIE HURAU, AND HENRI AGUT

*Laboratoire de Bactériologie-Virologie, C.E.R.V.I., and Laboratoire de Neurohistologie, Hôpital de la Pitié-Salpêtrière,
75013 Paris, and Unité d'Oncologie Virale, Institut Pasteur, and Laboratoire de Bactériologie-Virologie,
Hôpital Necker, 75015 Paris, France*

Volume 29, no. 2, p. 367-372 and Fig. 5: Position 154 in the nucleotide sequences of strains TAN, HST, and MBE is a C. Note that the *Hind*III and *Taq*I restriction sites depicted for group II human herpesvirus 6 strains are not present in this area (close to nucleotide 154).