

POSTER PRESENTATION

Open Access

Detection and quantification of STLV-1 and SFV proviral load in blood and saliva of naturally infected non-human primates

Sandrine Alais^{1,2,3,4,5}, Amandine Pasquier^{1,2,3,4,5,6}, Jocelyn Turpin^{1,2,3,4,5}, Rejane Rua⁷, Antoine Gessain⁷, Romain Lacoste⁸, Renaud Mahieux^{1,2,3,4,5*}

From 17th International Conference on Human Retroviruses: HTLV and Related Viruses Trois Ilets, Martinique. 18-21 June 2015

Simian T Lymphotropic Virus type 1 (STLV-1) and Simian Foamy Virus (SFV) retroviruses infect Old World non-human primates (NHP) and humans. Interhuman transmission has been described for HTLV-1 but not for SFV. SFV infection is asymptomatic in its hosts, while STLV-1 and its human counterpart HTLV-1 are the etiologic agents of Adult T-cell Leukemia/Lymphoma. Both STLV-1 and SFV can be zoonotically transmitted from NHP to humans through severe bites, thus involving contact between virus-containing saliva in the donor and blood in the recipient. Surprisingly, while the presence of both SFV RNA and DNA has been characterized into the saliva of NHP, neither STLV-1 DNA, nor STLV-1 RNA was quantified. Thus, the goal of our study was to search for STLV-1 provirus in the cells present in the saliva of NHP and then to quantify the proviral load of both viruses. We took advantages of a cohort of 45 papio anubis, naturally infected by STLV-1. We first assessed SFV infection and then potential SFV/STLV-1 co-infections. To this end, we designed semi-nested PCR and qPCR protocols (1) to diagnose infection and (2) to quantify STLV-1 and/or SFV proviral load in peripheral blood cells and in saliva. First, STLV-1 provirus was detected by semi-nested PCR in 8/10 blood samples tested, but only in the saliva of 1/10 NHP who had a high STLV-1 proviral load in peripheral blood cells. SFV DNA was detected by nested-PCR in blood samples from 10/10 baboons and in the saliva of 8/10 animals. A second study performed with 20 animals will be presented. We will show whether a correlation exists between blood and saliva STLV-1/SFV

proviral load and whether infection with one retrovirus impacts proviral load of the other. Altogether, our current results suggest that SFV is more frequently present in saliva than STLV-1. This should impact the ability of both viruses to be zoonotically transmitted through bites.

Authors' details

¹Equipe Oncogenèse Rétrovirale, International Center for Research in Infectiology, INSERM U1111 – CNRS UMR5308, France. ²Equipe labellisée "Ligue Nationale Contre le Cancer", International Center for Research in Infectiology, INSERM U1111 – CNRS UMR5308, France. ³International Center for Research in Infectiology, INSERM U1111 – CNRS UMR5308, France. ⁴Ecole Normale Supérieure de Lyon, Lyon, France. ⁵Université Lyon 1, Lyon, 69364 Cedex 07, France. ⁶Ecole Pratique des Hautes Etudes, Pasteur Institute, Paris, 75724 Cedex 15, France. ⁷Epidémiologie et Physiopathologie des Virus Oncogènes, CNRS UMR 3569, Pasteur Institute, Paris, 75724 Cedex 15, France. ⁸Station de Primatologie-UPS846-CNRS, Rousset sur Arc, 13790, France.

Published: 28 August 2015

doi:10.1186/1742-4690-12-S1-P4

Cite this article as: Alais *et al.*: Detection and quantification of STLV-1 and SFV proviral load in blood and saliva of naturally infected non-human primates. *Retrovirology* 2015 **12**(Suppl 1):P4.

¹Equipe Oncogenèse Rétrovirale, International Center for Research in Infectiology, INSERM U1111 – CNRS UMR5308, France Full list of author information is available at the end of the article



^{*} Correspondence: renaud.mahieux@ens-lyon.fr