



## HIV-1 and SUMOylation

Guillaume Beauclair, Ali Saïb, Alessia Zamborlini

► **To cite this version:**

Guillaume Beauclair, Ali Saïb, Alessia Zamborlini. HIV-1 and SUMOylation. *Retrovirology*, BioMed Central, 2013, 10 (Suppl 1), pp.P106. <inserm-00868804>

**HAL Id: inserm-00868804**

**<http://www.hal.inserm.fr/inserm-00868804>**

Submitted on 2 Oct 2013

**HAL** is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers.

L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.



POSTER PRESENTATION

Open Access

# HIV-1 and SUMOylation

Guillaume Beauclair<sup>1</sup>, Ali Saïb<sup>1,2</sup>, Alessia Zamborlini<sup>1,2\*</sup>

From *Frontiers of Retrovirology: Complex retroviruses, retroelements and their hosts*  
Cambridge, UK. 16-18 September 2013

## Background

The HIV virus hijacks cellular machineries to replicate. Post-translational modifications, like acetylation, phosphorylation or ubiquitination, are no exception. We are interested in studying the interplay between HIV and the SUMOylation pathway.

SUMOylation occurs via an enzymatic cascade requiring an E1 activating enzyme (SAE1/SAE2 heterodimer) and an E2 conjugating enzyme (Ubc9). This modification is reversible thanks to SUMO isopeptidases (SENPs). Despite Ubc9 alone being sufficient to transfer SUMO moieties to acceptor lysine residues *in vitro*, E3 SUMO ligases promote specificity and/or efficiency of SUMOylation *in vivo*. So far, several E3 SUMO ligases have been characterized, such as nuclear pore complex component Ran-binding protein 2 (RanBP2) and protein-inhibitor of activated STAT (PIAS) family proteins.

We have recently shown that HIV integrase, the viral enzyme that is responsible for viral genome integration into host cellular chromosome, is SUMOylated, and that virions harboring a SUMOylation-defective integrase are less infectious than WT viruses. Our data suggest that this modification is important for efficient viral replication.

## Materials and methods

To detect protein SUMOylation, cells expressing His-tagged SUMO protein together with the protein of interest (HIV integrase) were lysed in denaturing conditions followed by enrichment on NiNTA column. Bound proteins were eluted by boiling in Laemmli sample buffer supplemented with imidazole. Intracellular localization was studied by Duo LINK approach. Protein-protein interactions were analyzed by immunoprecipitation and GST pull-down assays.

## Results and conclusions

To better characterize the involvement of SUMOylation during HIV replication, we are currently studying the role of E3 SUMO ligases and SENPs in the conjugation of SUMO to integrase. We established that HIV integrase interacts with PIAS family member proteins and we showed that integrase and PIAS colocalise. Moreover, we demonstrate that PIAS1 and PIAS4, but not other PIAS family members, increase whereas SENP1 decreases integrase SUMOylation. We also analyzed the effect of PIAS over-expression in virus-producing or target cells and do not find a significant variation of viral infectivity. Interestingly, during HIV infection we show a decrease of the expression levels of one of the members of PIAS family. Experiments are ongoing to elucidate the mechanisms underlying this interplay between HIV and the SUMOylation pathway.

## Authors' details

<sup>1</sup>CNRS/UP7 UMR7212, INSERM U944, Institut Universitaire d'Hématologie-Université Paris7 Diderot, Paris, 75010, France. <sup>2</sup>Conservatoire des Arts et Métiers, Paris, 75003, France.

Published: 19 September 2013

doi:10.1186/1742-4690-10-S1-P106

Cite this article as: Beauclair *et al.*: HIV-1 and SUMOylation. *Retrovirology* 2013 **10**(Suppl 1):P106.

<sup>1</sup>CNRS/UP7 UMR7212, INSERM U944, Institut Universitaire d'Hématologie-Université Paris7 Diderot, Paris, 75010, France

Full list of author information is available at the end of the article