



POSTER PRESENTATION

Open Access

Evaluation of pathways and new host proteins involved in CD4 down-modulation during HIV-1 infection

Alessia Landi^{1*}, Mostafa Bentahir^{1,2}, Jolien Vermeire¹, Hanne Vanderstraeten¹, Veronica Ianucci¹, Bruno Verhasselt¹

From Frontiers of Retrovirology 2011
Amsterdam, The Netherlands. 3-5 October 2011

Down-modulation of the CD4 receptor expressed on T cells is one of the hallmarks of Human Immunodeficiency Virus (HIV) infection and it is believed to confer to the virus a selective replicative advantage *in vivo*. HIV has evolved redundant mechanisms to remove the receptor from the cell surface and accelerate its degradation. This process is mainly mediated by three viral proteins: Vpu, Env and Nef. Up to date, the mechanisms that lead to CD4 depletion from the surface of CD4+ T lymphocytes, the natural targets of HIV, are still poorly understood and only partially characterized. We are interested in the discovery of pathways and human proteins involved in the process, in order to eventually find potential new drug targets. To pursue our aim, we first performed a functional screening on HeLa CD4+ cells expressing Nef, using a shRNA lentiviral interference delivery system targeting the whole human genome compatible with an Affymetrix GeneChip Microarray. The read out was the rescue of the CD4 high phenotype, despite Nef expression. The results were analyzed with Affymetrix Expression Console 1.1 software and the web-based gene ontology resource DAVID. After four different screens we obtained a final list of 75 genes enriched in the cells sorted for high surface CD4. These genes appear to code for proteins involved in endocytic trafficking, trans-Golgi trafficking and lysosomal degradation pathways. We decided to validate our hits in primary blood CD4+ T cells, the most relevant for this kind of studies, and we set up a protocol for the transduction of these cells with lentiviral and retroviral vectors, necessary respectively for the expression of the shRNAs and the viral proteins Vpu, Nef and Env. The

actual down-regulation of the gene expression will be validated via quantitative Real Time PCR and Western Blot and the experiments will be eventually repeated in the context of HIV-1 infection.

Author details

¹Department of Clinical Chemistry, Microbiology, and Immunology, Ghent University, Belgium. ²Present address: Centre de Technologies Moléculaires Appliquées, Ecole de Santé Publique, Brussels, Belgium.

Published: 3 October 2011

doi:10.1186/1742-4690-8-S2-P43

Cite this article as: Landi et al.: Evaluation of pathways and new host proteins involved in CD4 down-modulation during HIV-1 infection. *Retrovirology* 2011 **8**(Suppl 2):P43.

Submit your next manuscript to BioMed Central and take full advantage of:

- Convenient online submission
- Thorough peer review
- No space constraints or color figure charges
- Immediate publication on acceptance
- Inclusion in PubMed, CAS, Scopus and Google Scholar
- Research which is freely available for redistribution

Submit your manuscript at
www.biomedcentral.com/submit



¹Department of Clinical Chemistry, Microbiology, and Immunology, Ghent University, Belgium
Full list of author information is available at the end of the article