## Retrovirology



Poster presentation

**Open Access** 

## Development of a Human Bone Marrow Progenitor Cell Line to Examine HIV-I Susceptibility and LTR Activity

Aikaterini Alexaki\*<sup>‡</sup>, Michael Nonnemacher and Brian Wigdahl

Address: Department of Microbiology and Immunology and Institute for Molecular Medicine and Infectious Disease, Drexel University College of Medicine, Philadelphia, PA, USA

from 2005 International Meeting of The Institute of Human Virology Baltimore, USA, 29 August – 2 September 2005

Published: 8 December 2005

Retrovirology 2005, 2(Suppl 1):P8 doi:10.1186/1742-4690-2-S1-P8

Previous studies have suggested that the bone marrow compartment may play an integral role in the genesis of HIV-1 dementia (HIVD). Interestingly, CD34+/CD38pluripotent stem cells within the bone marrow are refractile to HIV-1 infection. The CD34+/CD38+ TF-1 cell line has been selected as a model to study HIV-1 infection during the differentiation process of hematopoietic progenitor cells. A number of cytokines such as GM-CSF, M-CSF, IL-1 $\beta$ , TNF- $\alpha$ , and IL-4 were used to induce differentiation and activation of TF-1 cells and their surface marker expression was monitored by flow cytometry. Interestingly, IL-1β treatment, alone or in combination with TNFα, lead to up-regulation of CXCR4 and CCR5 surface presentation, and preservation of CD4 expression possibly providing an optimal cellular phenotype for HIV-1 infection of this cell population. The surface marker expression after this treatment also correlated with a more differentiated phenotype. To begin exploring the potential of these cells to support productive HIV-1 replication, a series of stably transfected cell lines were developed. To this end, macrophage-, T cell- and dual-tropic long terminal repeats (LTRs) were coupled to the gene encoding green fluorescent protein. These cell lines were utilized to explore the functional properties of specific cis-acting regulatory elements in LTR function within the bone marrow precursor cell population.