

Poster presentation

Open Access

PI7-24. Interfering overlapping epitopes contribute to the subdominance of an HLA-A2-restricted HIV Gag-specific epitope

J Kan-Mitchell*¹, AM Nyakeriga¹, JR Salkowitz¹, MC Costanzo¹, J Sidney², A Sette² and V Ayyavoo³

Address: ¹Biological Sciences, University of Texas at El Paso, El Paso, TX, USA, ²La Jolla Institute of Allergy and Infectious Disease, La Jolla, CA, USA and ³University of Pittsburgh School of Public Health, Pittsburgh, PA, USA

* Corresponding author

from AIDS Vaccine 2009
Paris, France. 19–22 October 2009

Published: 22 October 2009

Retrovirology 2009, 6(Suppl 3):P306 doi:10.1186/1742-4690-6-S3-P306

This abstract is available from: <http://www.retrovirology.com/content/6/S3/P306>

© 2009 Kan-Mitchell et al; licensee BioMed Central Ltd.

Background

Recapitulating immunodominant A2-restricted HIV-specific CTLs that do not suppress virus *in vivo* is unlikely to be an effective vaccine strategy.

Methods

To detect novel subdominant determinants in HIV-1 Gag, we primed CD8+ T cells from eight seronegative donors with autologous dendritic cells transduced to express Gag. T cells were re-stimulated weekly with monocytes pulsed with 123 15-mer overlapping peptides (OLPs) spanning Gag. Responses were identified with OLP matrix pools followed by interrogation at the single OLP level in responsive pools using IFN- γ ELISPOT assays.

Results

One OLP (Gag145-159) was recognized by all donors. Of note, this reactivity predominated in five of eight T cell cultures. Fine mapping with progressively truncated peptides revealed three overlapping epitopes: RTLNAWVKV (RV9), RTLNAWVKVV (RV10) and TLNAWVKVV (TV9). TV9 is a known epitope that is rarely recognized *in vivo*. In contrast, RV9 and RV10 are novel. Although the latter bound with lower affinities to HLA-A2 than TV9, RV9- and RV10-cultures were readily generated. RV9- and RV10-T cells were cytotoxic, secreting cytokines and suppressive of HIV replication *in vitro*.

Conclusion

In sum, we report two novel naturally processed and presented epitopes in HIV p24 that are recognized by pre-infection T cell repertoires. Further studies are needed to explain why these reactivities are rare during infections and more importantly, whether this conserved region of the HIV proteome has value as a prophylactic vaccine for A2 individuals.