

# Phenotypic characterisation of virus-specific T cells in treated HIV-1 infection: Profiling total and multimer-specific CD8 T cells

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## Introduction

Maturation and activation of T cells is critical for effective immune control of viruses including HIV-1. In patients with established HIV-1 infection, a skewed maturation/differentiation profile has been linked to viral pathogenesis and is not restored by effective combination (c)ART (1, 2). Furthermore, chronic immune activation is one of the strongest predictors of HIV-1 disease progression (3). In HIV-1 negative individuals, asymptomatic CMV is associated with higher T-cell activation (4). cART reduces T-cell activation, however little is known about its effect on CD8 T-cell maturation. We evaluated the magnitude and characteristics of both HIV-1- and CMV-specific T-cell responses in order to further elucidate the effect of cART on virus-specific T-cell differentiation and activation.

## Methods

• Activation (HLA-DR/CD38) and maturation (CD45RA/CCR7) profiles of total CD8<sup>+</sup> T cells from 31 cART-treated HIV-1<sup>+</sup> CMV<sup>+</sup> co-infected patients (median 420 CD4 T cells/ $\mu$ l blood) and 5 HIV-1 negative healthy controls for comparison, were examined ex-vivo. This involved surface staining of peripheral blood mononuclear cells (PBMC) with monoclonal antibodies: CD3, CD8, CD45RA, CCR7, CD38 and HLA-DR for flow cytometric analysis.

• PBMC of 7 HIV-1<sup>+</sup> CMV<sup>+</sup> patients were also stained with HA9-B\*3501 pentamer, TM10-B\*0702 pentamer (all ProlImmune Ltd, Oxford, UK) and TM10-B\*0702 dextramer (Immudex, Copenhagen, Denmark) in order to identify antigen-specific CD8<sup>+</sup> T cells.

• Non-parametric intergroup analysis was performed using Mann-Whitney U test, and paired data was analysed by Wilcoxon signed-rank test, with significance defined as  $p < 0.05$ .

## References

1. Champagne et al. (2001) *Nature*, 410, 106-111
2. Migueles et al. (2009) *J Virol*, 83, 11876-11889
3. Hazenberg et al. (2003) *AIDS*, 17, 1881-1888
4. Hunt et al. (2011) *J Inf Dis*, 203, 1474-1483
5. Champagne et al. (2001) *Nature*, 410, 106-111
6. Carrington and Walker (2012) *Annu Rev Med*, 63, 131-145

### Gating strategy

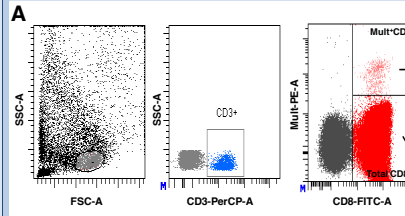
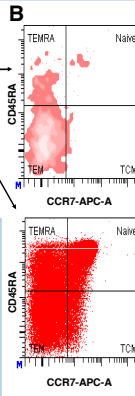
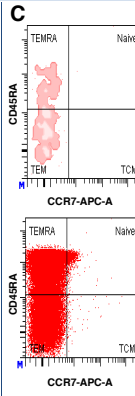


Figure 1. Representative example of staining (A). Total and multimer-specific CD8<sup>+</sup> T cells are shown, expressing CCR7 and CD45RA (B and C) and CD38 and HLA-DR (D).

### Differentiation CMV-specific



### Differentiation HIV-1-specific



### Activation CMV-specific

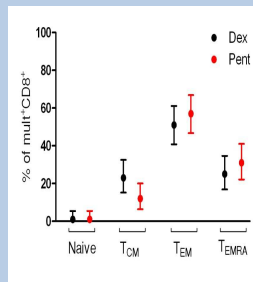
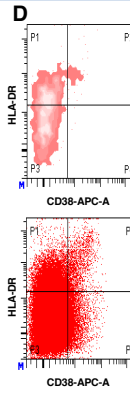


Figure 2. Comparison of the proportion of multimer+CD8<sup>+</sup> memory subsets observed using dextramer and pentamer technologies in parallel. The two technologies were comparable. Symbols represent percentage distribution for each technology and error bars indicate 95% confidence intervals.

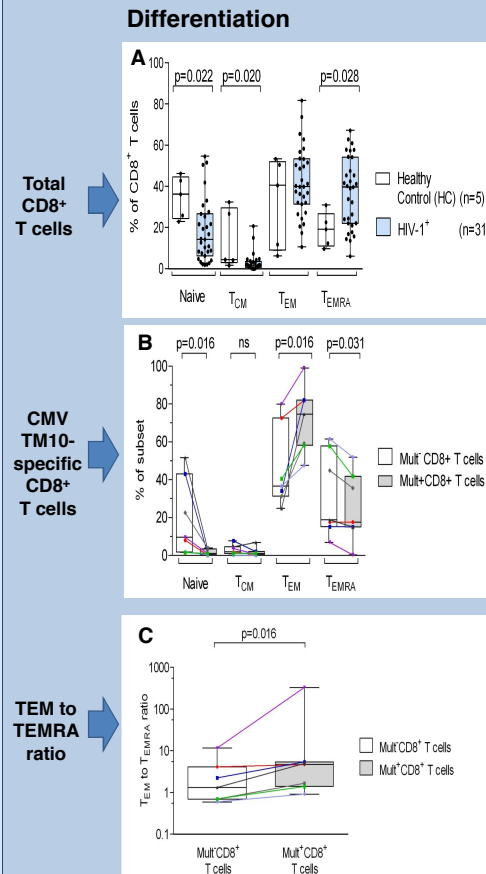


Figure 3. Differentiation profile of total CD8<sup>+</sup> T cells in HIV-1<sup>+</sup> patients (A; blue bars) and healthy controls (A; white bars) and mult+CD8<sup>+</sup> T cells (white bars) and mult+CD8<sup>+</sup> T cells (grey bars) in HIV-1<sup>+</sup> patients (data from B and C). Each coloured line represents one individual.

### Activation

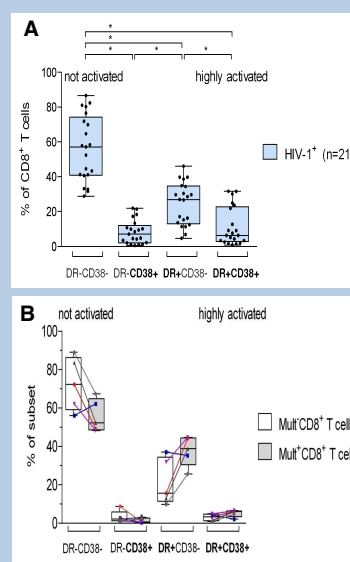


Figure 4. Activation levels of total CD8<sup>+</sup> T cells (A), and mult+CD8<sup>+</sup> T cells and mult+CD8<sup>+</sup> T cells (B). Data from HIV-1<sup>+</sup> patients. Each coloured line represents one individual.

## Results

Analysis of multimer-specific CD8<sup>+</sup> T cells and their differentiation profile (Figure 1 A, B and C) showed that the two multimer technologies were comparable (Figure 2). HIV-1<sup>+</sup> patients showed a significant reduction in naïve and central memory (T<sub>CM</sub>) compartments ( $p=0.022$  and  $0.020$  respectively), and a significant increase in terminally differentiated (T<sub>EMRA</sub>) subset within total CD8<sup>+</sup> T cells ( $p=0.028$ ), compared to healthy controls (Figure 3 A). CMV TM10-specific CD8<sup>+</sup> T cells had a significantly higher T<sub>EM</sub>:T<sub>EMRA</sub> ratio (median= 4.65), compared to the rest of the CD8 T-cell pool (median= 1.32;  $p=0.016$ ; Figure 3 B and C). The majority of HIV-1<sup>+</sup> individuals had a low proportion of CD8<sup>+</sup> T cells that were highly activated (Figure 4 A). There was no significant difference in the activation levels between CMV TM10-specific CD8<sup>+</sup> T cells and total CD8<sup>+</sup> T cells (Figure 4 B).

## Conclusions

- Maturation profiles of total and multimer-specific CD8<sup>+</sup> T cells indicate a shift towards the T<sub>EM</sub> and T<sub>EMRA</sub> subset.
- Although CMV-specific CD8<sup>+</sup> T cells have been shown to be predominantly of the T<sub>EMRA</sub> subset in untreated HIV-1<sup>+</sup> individuals (5), here we show that in cART-treated HIV-1<sup>+</sup> patients CMV-specific CD8 T cells are primarily at the effector memory (T<sub>EM</sub>) stage of differentiation.
- There is no difference in the activation level of CMV TM10-specific CD8<sup>+</sup> T cells compared to total CD8<sup>+</sup> T cells, even in the context of cART.

## Future Studies

Further studies aim to look at the role of different HLA alleles on the specificity and avidity of T-cell immune responses and how this affects HIV-1 disease progression (6).

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