

# A case study investigation into the role of CD4<sup>+</sup> tumor-infiltrating lymphocytes in a metastatic melanoma patient with a complete response to adoptive cell therapy

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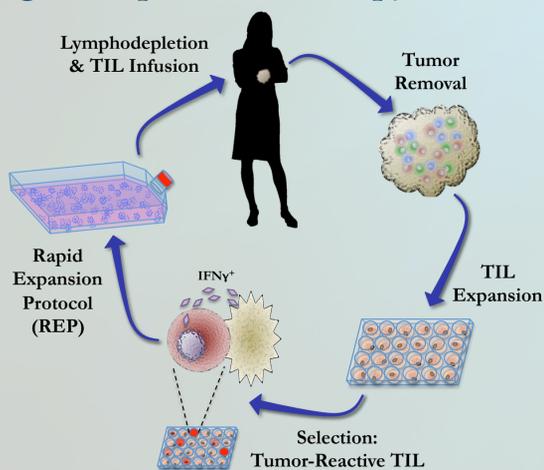
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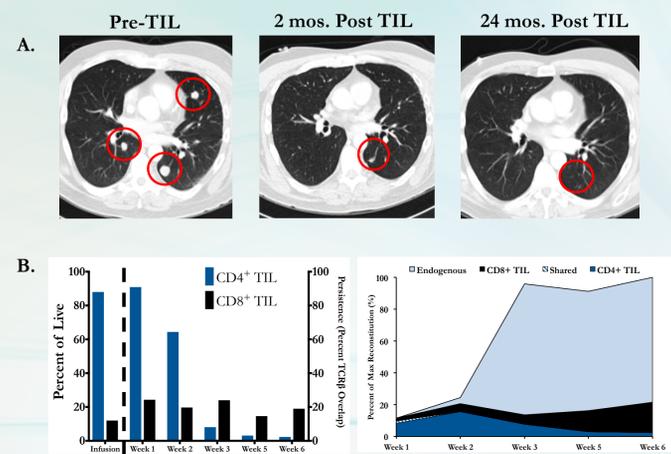
## ABSTRACT

Immunotherapy for cancer has long been focused on the generation of CD8<sup>+</sup> cytotoxic T lymphocyte responses, independent of their dynamic CD4<sup>+</sup> T cell counterpart. One promising approach, adoptive cell transfer (ACT) of tumor-infiltrating lymphocytes (TIL), has yielded response rates ranging from 28-55%. Although lasting and complete responses have been achieved, there is substantial opportunity for improvement. Investigation into the role of CD4<sup>+</sup> TIL in this setting remains critically underexplored. CD4<sup>+</sup> T cells recognize tumor antigen presented on MHC Class II either directly on tumor cells or indirectly through antigen presenting cells (APCs) and are able to elicit potent anti-tumor responses under the appropriate conditions. Here, we present a case study of a metastatic melanoma patient who received adoptive transfer of a predominantly (88%) CD4<sup>+</sup> TIL product. This patient demonstrated a complete response (CR) to therapy despite a lack of detection of IFN $\gamma$  in the infusion product *in vitro* when these TIL were co-cultured with autologous tumor prior to ACT. Tumor recognition was also absent when CD8<sup>+</sup> TIL were isolated and stimulated directly with HLA-matched tumor lines, indicating a lack of recognition of shared melanoma antigens presented on MHC Class I. Longitudinal analysis of the peripheral blood of this patient confirmed that the infused CD4<sup>+</sup> TIL persisted after therapy for at least six weeks. Whole exome sequencing (WES) performed on the TIL surgical specimen discovered 88 non-synonymous single nucleotide variants (SNVs) as candidate neoantigens. Predicted binding of the resulting mutant peptides to autologous HLA molecules generated a predominantly MHC Class II restricted profile, with 81.8% of variants capable of MHC Class II presentation and greater than half exclusive to MHC Class II only. CD4<sup>+</sup> TIL were screened for tumor antigen recognition by upregulation of OX40 and 41BB after stimulation with autologous APCs loaded with mutant peptides. Nearly half (49.2%) of CD4<sup>+</sup> TIL responded to tumor-derived peptides. These CD4<sup>+</sup> TIL were then sorted into tumor-reactive and non-reactive subsets for further clonal analysis of phenotype and transcriptional profile (scRNASeq) of these T cells in order to characterize the nature of the CD4<sup>+</sup> TIL response to tumor antigen. Overall, thorough interrogation of this patient's case study demonstrated evidence of CD4<sup>+</sup> TIL involvement in a complete clinical response after ACT. Ongoing studies will define the precise role of tumor-reactive CD4<sup>+</sup> T cells in the anti-tumor immune response and provide the framework for future investigation into their function and therapeutic efficacy.

## Fig 1. Adoptive Cell Therapy with TIL

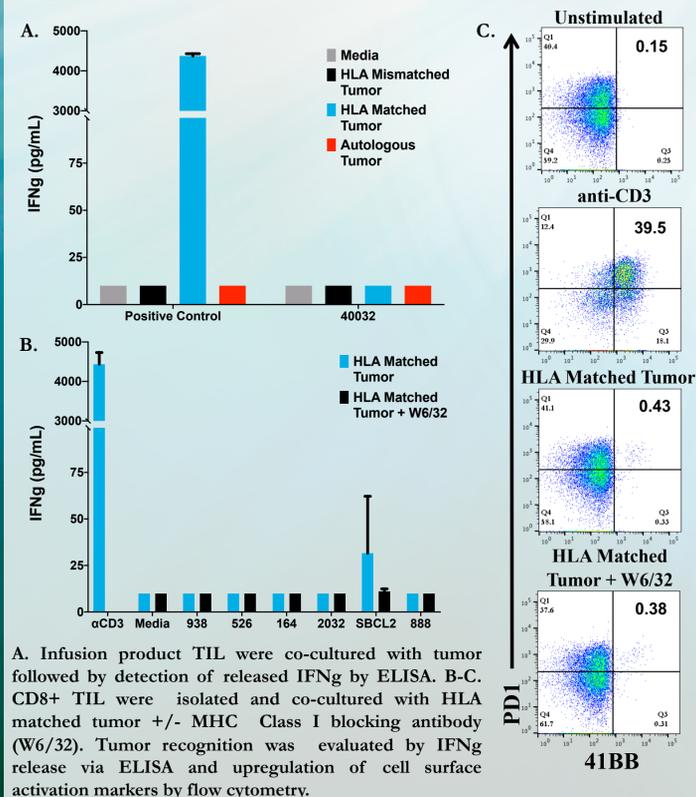


## Fig 2. CD4<sup>+</sup> TIL persist *in vivo*



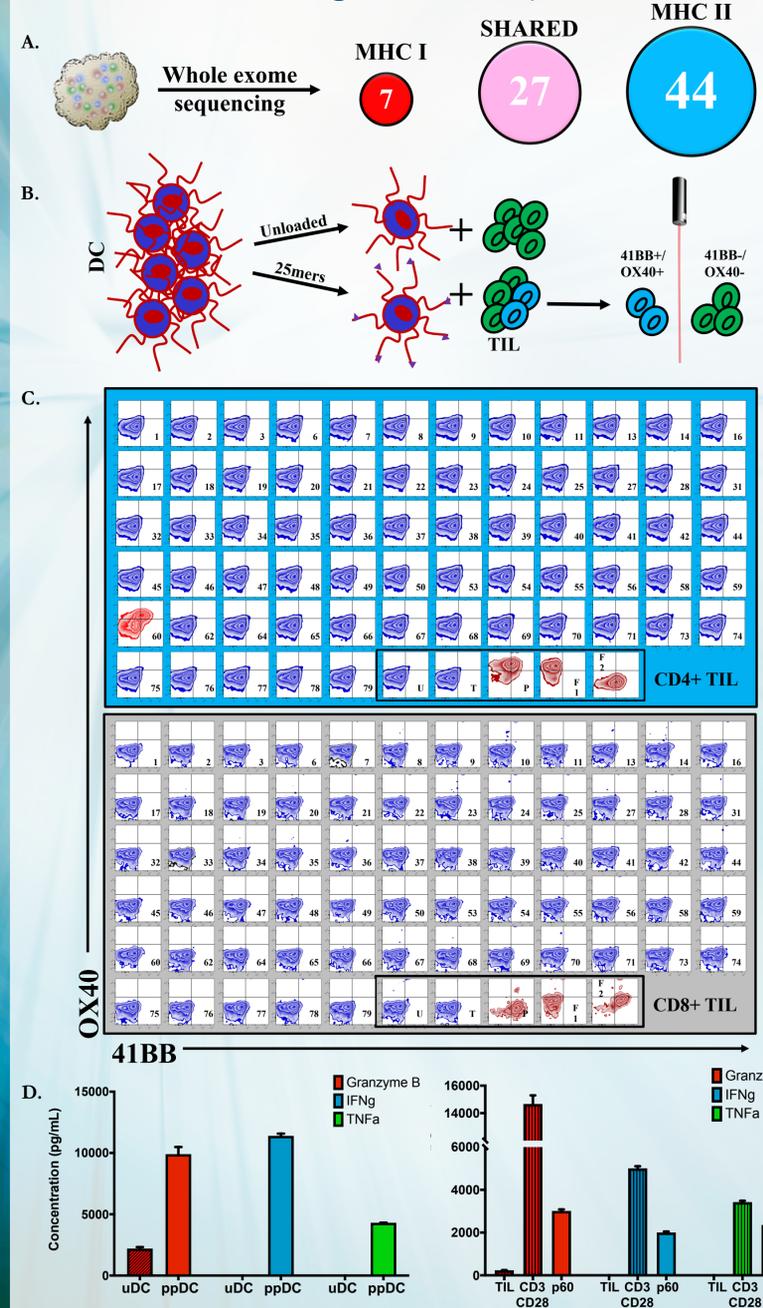
A. CT scans of lung metastases from complete responder (CR) infused with predominantly CD4<sup>+</sup> TIL [1]. B. Persistence measured by TCR $\beta$  overlap between CD4<sup>+</sup> and CD8<sup>+</sup> TIL and weekly PBMC repertoire.

## Fig 3. CD8<sup>+</sup> TIL fail to recognize tumor



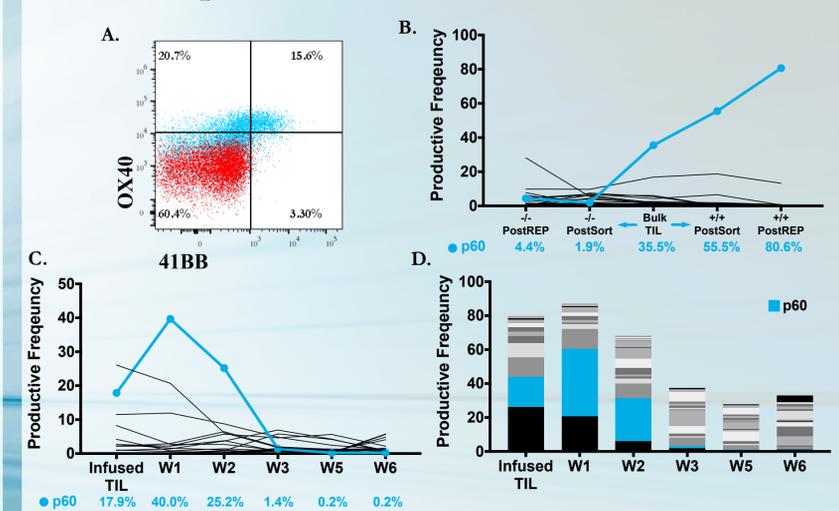
A. Infusion product TIL were co-cultured with tumor followed by detection of released IFN $\gamma$  by ELISA. B-C. CD8<sup>+</sup> TIL were isolated and co-cultured with HLA mismatched tumor +/- MHC Class I blocking antibody (W6/32). Tumor recognition was evaluated by IFN $\gamma$  release via ELISA and upregulation of cell surface activation markers by flow cytometry.

## Fig 4. Polyfunctional neoantigen-specific CD4<sup>+</sup> TIL detected via immunogenomic analysis



A. Mutant 25mers were predicted for MHC binding from whole exome sequencing. B-D. TIL were stimulated with peptides loaded on dendritic cells (DC) and assessed for increased cell surface expression of OX40 and 41BB by flow cytometry and effector molecule production by ELISA.

## Fig 5. Clonal tracing demonstrates *in vitro* enrichment and *in vivo* persistence



A. Neoantigen-specific TIL were sorted on OX40/41BB expression (+/+ vs. -/-) in response to peptide. B-D. Sorted and expanded (REP) TIL populations were sequenced at the TCR $\beta$  locus and tracked from infusion product across weekly patient peripheral blood samples for clonal persistence.

## CONCLUSIONS & FUTURE DIRECTIONS

- Patient achieved a complete response (CR) after ACT with predominantly CD4<sup>+</sup> TIL
- CD4<sup>+</sup> TIL were highly functional in response to autologous tumor peptides
- Neoantigen-specific CD4<sup>+</sup> TIL were enriched by OX40/41BB selection, expanded and demonstrated strong persistence within the patient
- Identify TCR $\alpha\beta$  and mutant peptide cognate pair
- Determine the transcriptional profile of these neoantigen CD4<sup>+</sup> TIL
- Further elucidate the function of these CD4<sup>+</sup> TIL *in vitro* and *in vivo*
- Expand analysis to additional patient samples to understand importance of CD4<sup>+</sup> TIL in ACT

## ACKNOWLEDGEMENTS

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Moffitt Cancer Center has licensed Intellectual Property related to the proliferation and expansion of bulk tumor-infiltrating lymphocytes (TIL) to Iovance Biotherapeutics. MH, AAS, and SPT are inventors on such Intellectual Property.

## REFERENCES

1. Pilon-Thomas S, et al. *J Immunother*. 2012; 35: 615-20.

