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Society for Immunotherapy of Cancer

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Fueling immunotherapy through modulation of the metabolic axis



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Presenter Disclosure Information

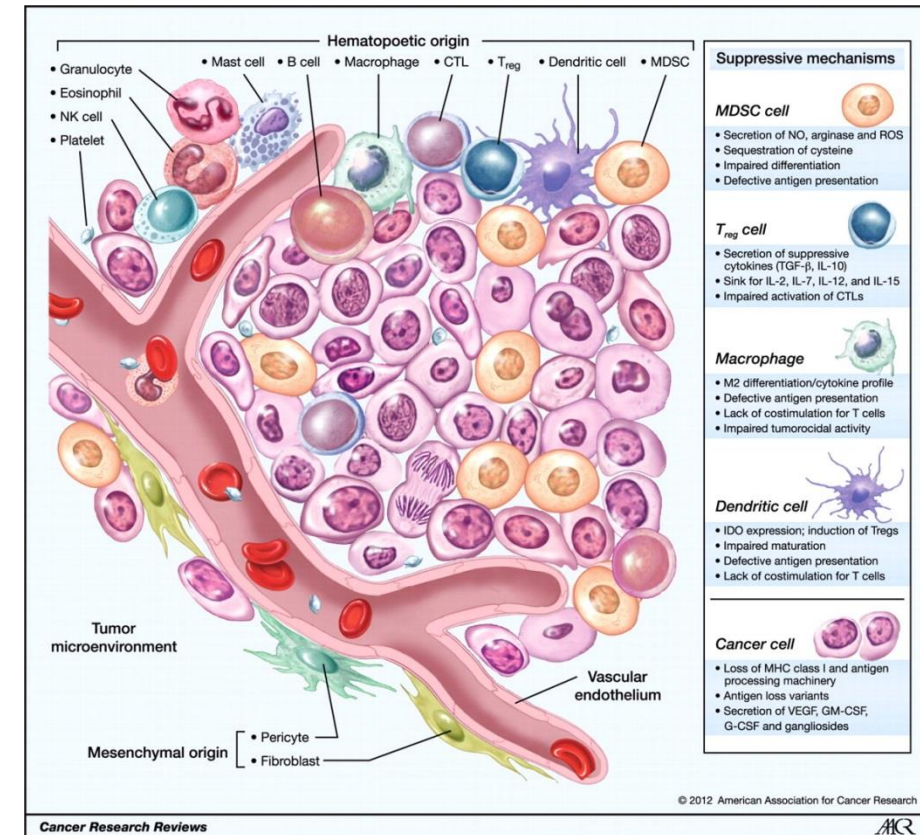
Greg M. Delgoffe

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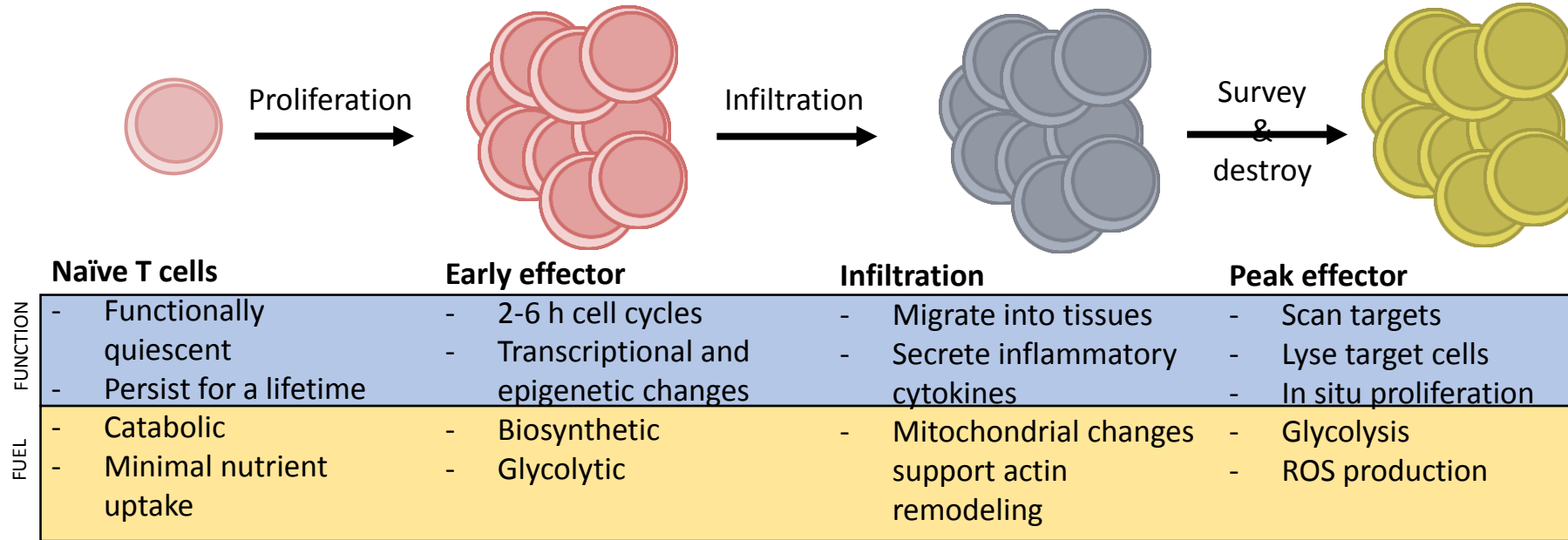
Merck & Co., Received, Consultant

Tumor microenvironment metabolism as a patient-specific resistance mechanism to immunotherapy

- Progressing tumors develop a potently immunosuppressive **tumor microenvironment**, which fosters tumor adaptation and progression
- Characterized by many different differentiation stages of tumor cell, altered fibroblasts, and a constellation of immunosuppressive cell types
 - Myeloid derived suppressor cells
 - Tumor associated macrophages
 - Regulatory T cells
- The tumor microenvironment also generates a **distinct metabolic landscape**
 - Hypoxia
 - Acidosis
 - Hypoglycemia
 - Saturated fats and ketone bodies
 - Depleted essential amino acids

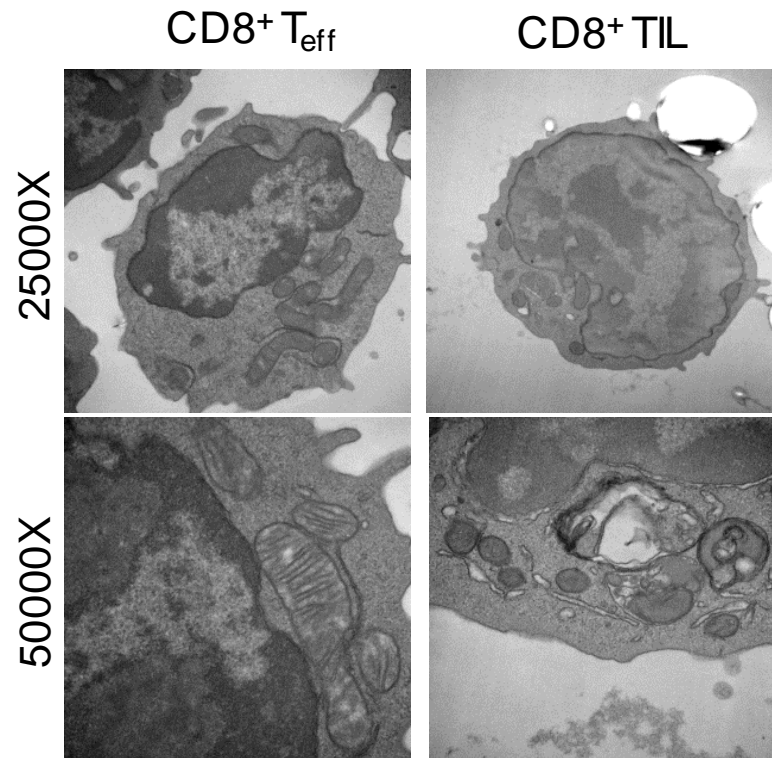


T cell activation is metabolically demanding

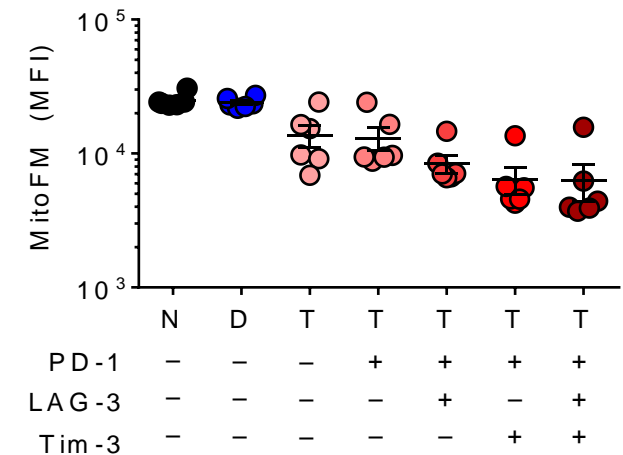
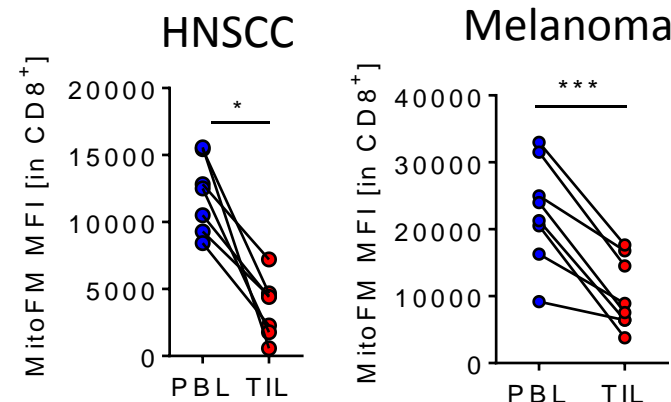


Thus, access to nutrients and the ability to process them represents a mechanism by which T cells can be regulated

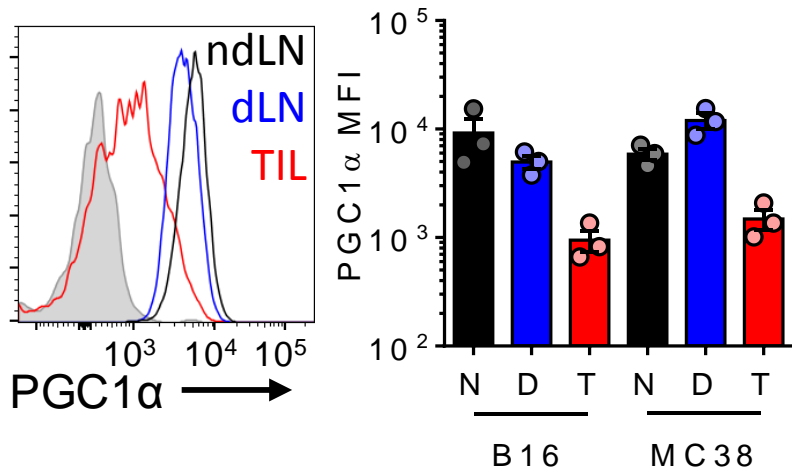
What is the metabolic phenotype of tumor infiltrating T cells?



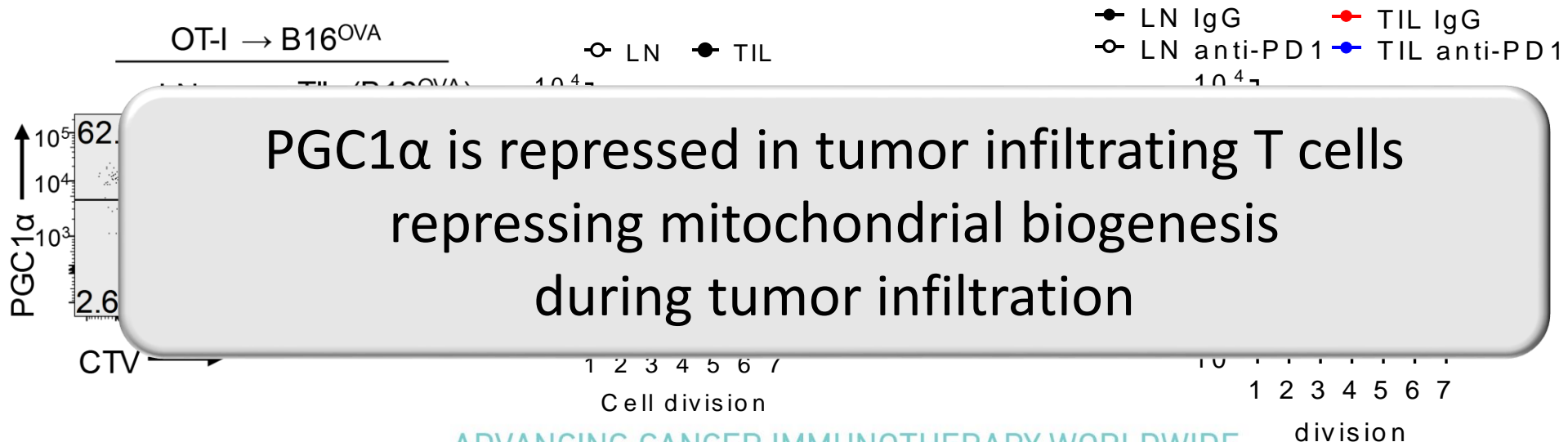
- Tumor infiltrating T cells in murine and human cancer show defects in glucose uptake and dysfunctional mitochondria
- Loss of T cell metabolic sufficiency correlates with upregulation of co-inhibitory molecules, but occurs even in the presence of PD-1 blockade!



What is the molecular mechanism of mitochondrial loss in intratumoral T cells?

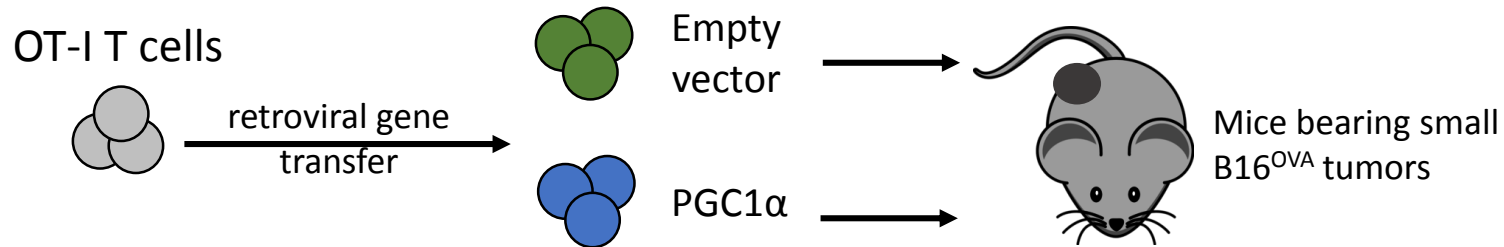


- Mitochondrial biogenesis, the process by which new mitochondria are generated, is programmed in part by the transcriptional co-activator PGC1α
- PGC1α protein is dramatically reduced in TIL CD8⁺ T cells
- PGC1α loss occurs progressively as T cells become activated in tumors
- PGC1α downregulation occurs independently of PD-1 signaling

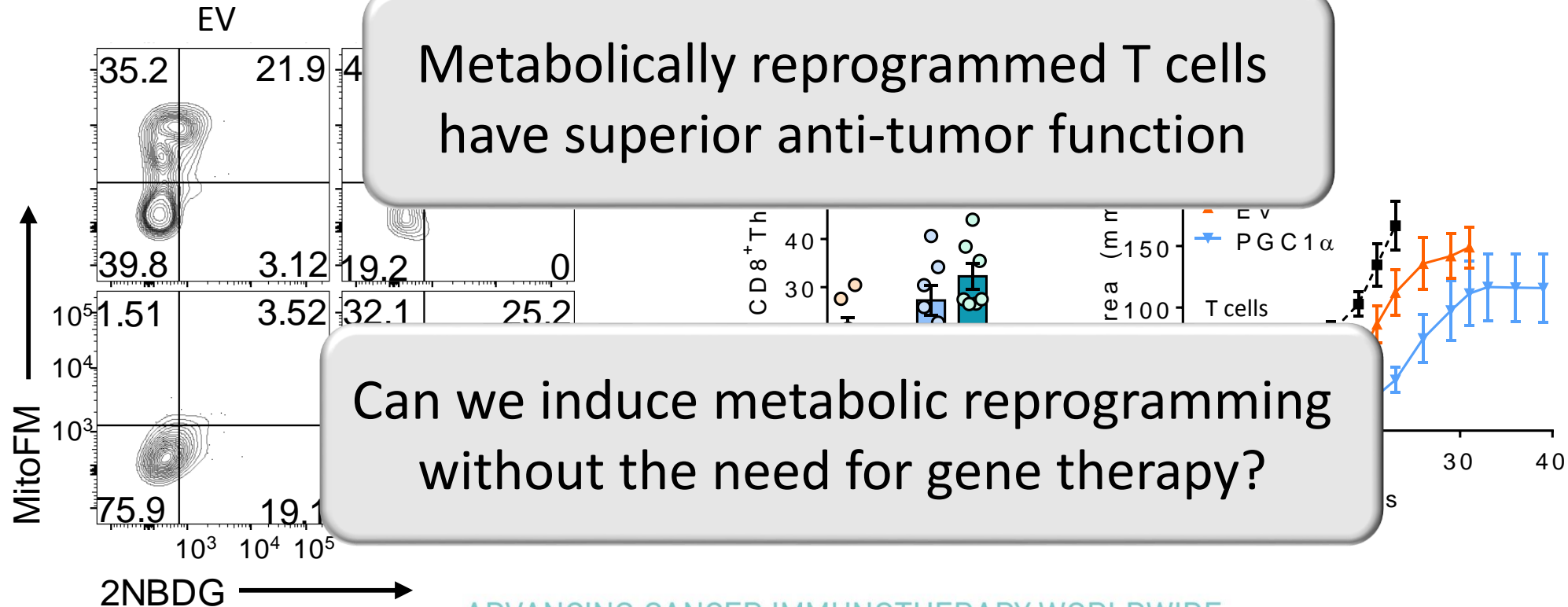


PGC1α is repressed in tumor infiltrating T cells
repressing mitochondrial biogenesis
during tumor infiltration

Can T cells be reprogrammed metabolically to bolster function in the tumor?



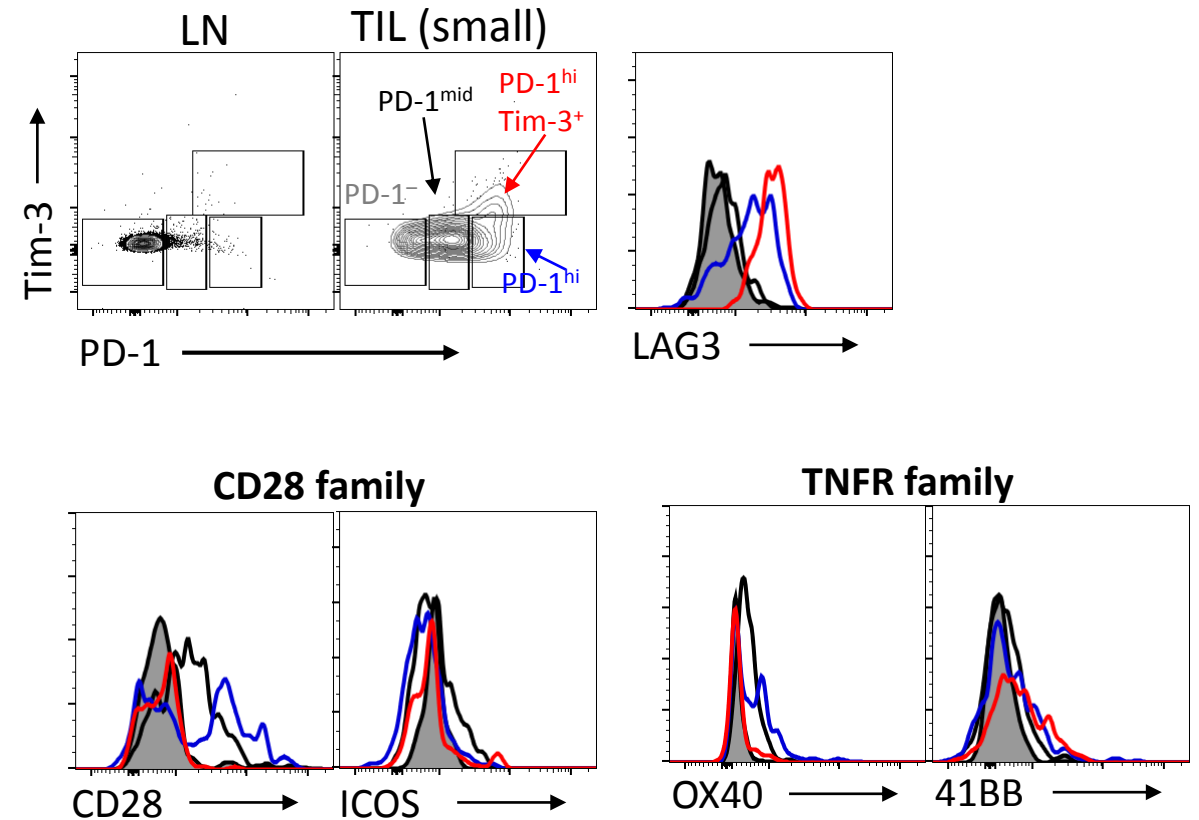
Metabolically reprogrammed T cells have superior anti-tumor function



Can we induce metabolic reprogramming without the need for gene therapy?

Costimulatory molecules in the TME

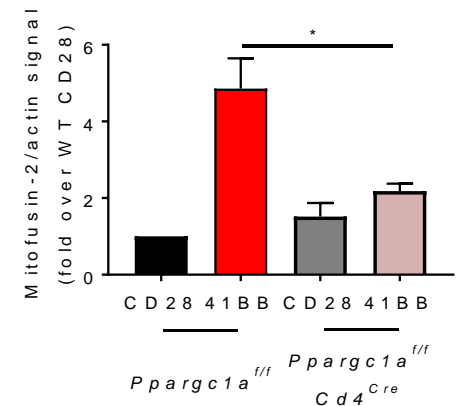
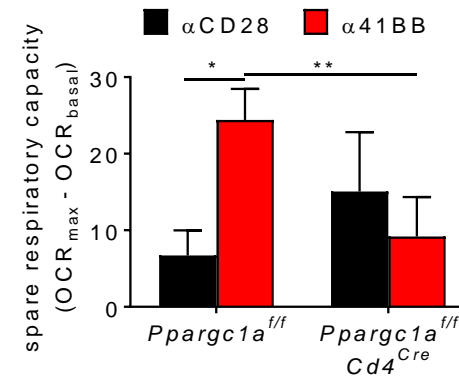
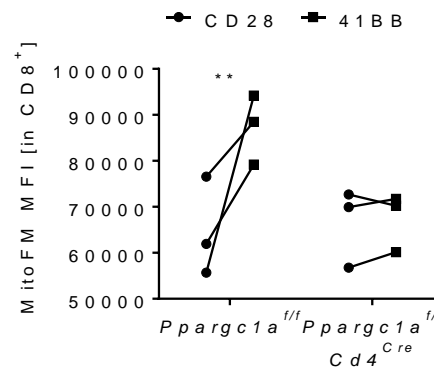
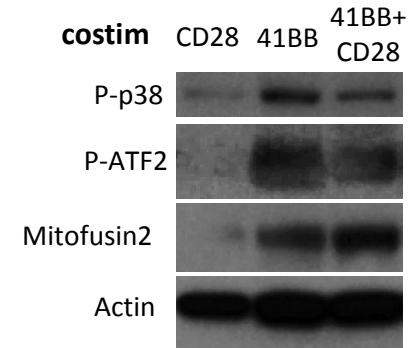
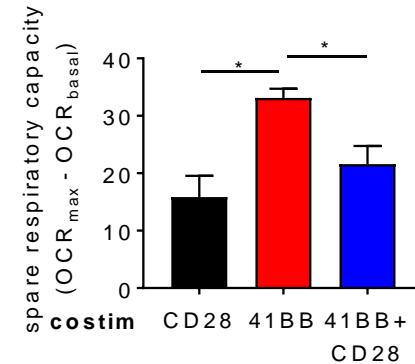
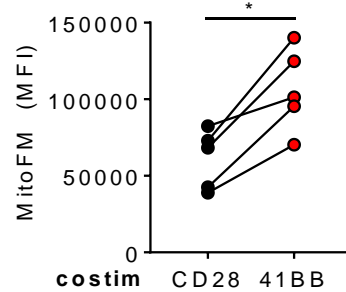
- Previous work has highlighted the importance of costimulation in metabolic reprogramming, but typically in a nontumor setting
- The costimulatory/inhibitory molecules expressed by functional/dysfunctional states of TIL may provide insight into how to properly rejuvenate them
- We assayed which molecules were expressed on T cells in various functional states as defined by PD-1 and Tim-3 in small melanomas
- LAG3, as expected, progressively rose as T cells upregulated PD-1 and began making Tim-3
- CD28 family members were markedly repressed as T cells became more and more phenotypically 'exhausted'
- OX40 was repressed in terminally exhausted T cells
- 41BB, however, was expressed on the most exhausted T cells



Can 41BB ligation provide metabolic support to the most exhausted TIL?

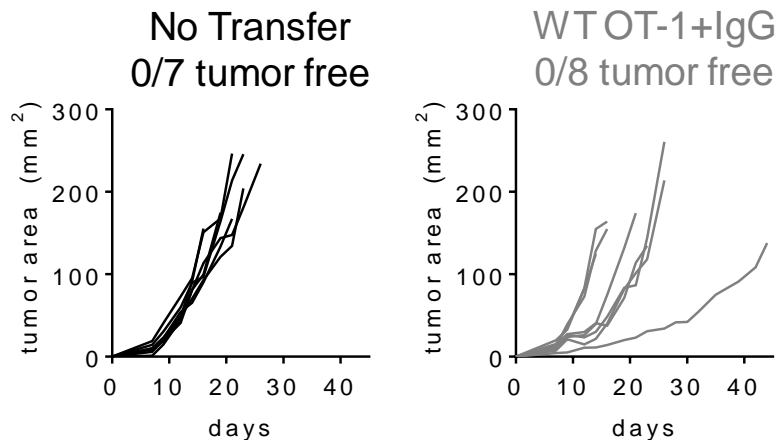
Can 41BB support mitochondrial metabolism?

- 41BB costimulation of murine T cells results in cells with increased mitochondrial mass
- Costimulation of 41BB does not induce increased mitochondrial oxidation, but rather improves mitochondrial capacity
- 41BB ligation promotes increases in PGC1 α expression and a program of mitochondrial fusion and biogenesis
- PGC1 α is required for 41BB-mediated metabolic reprogramming



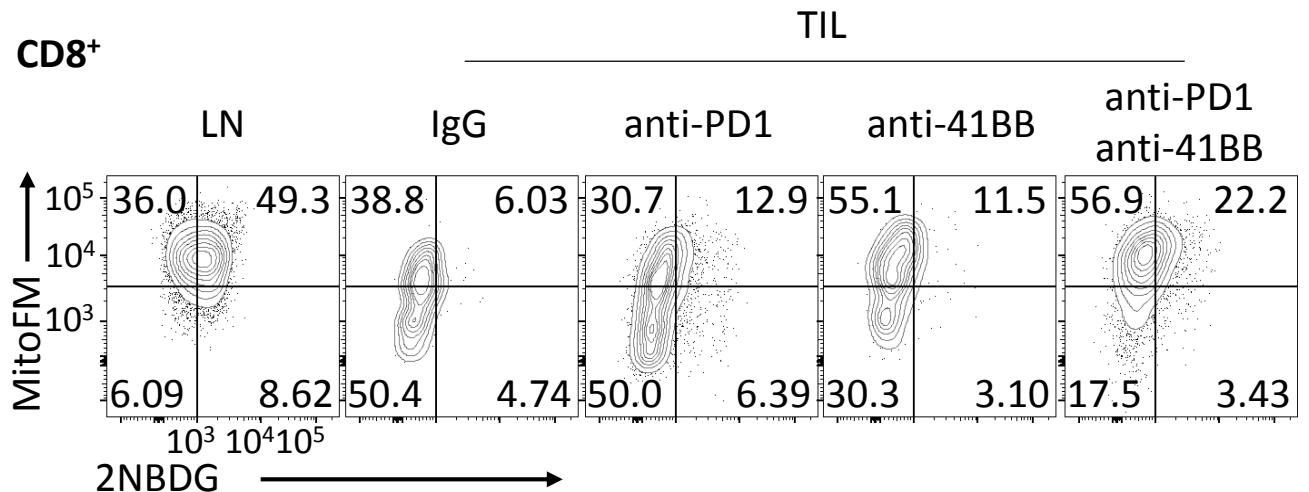
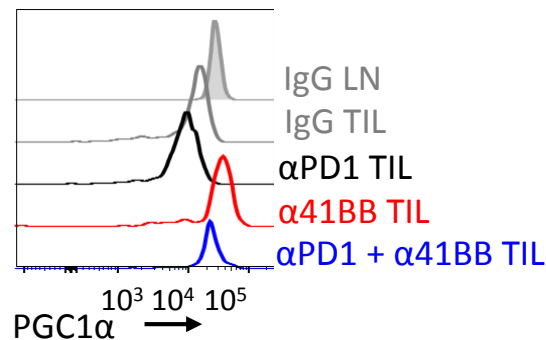
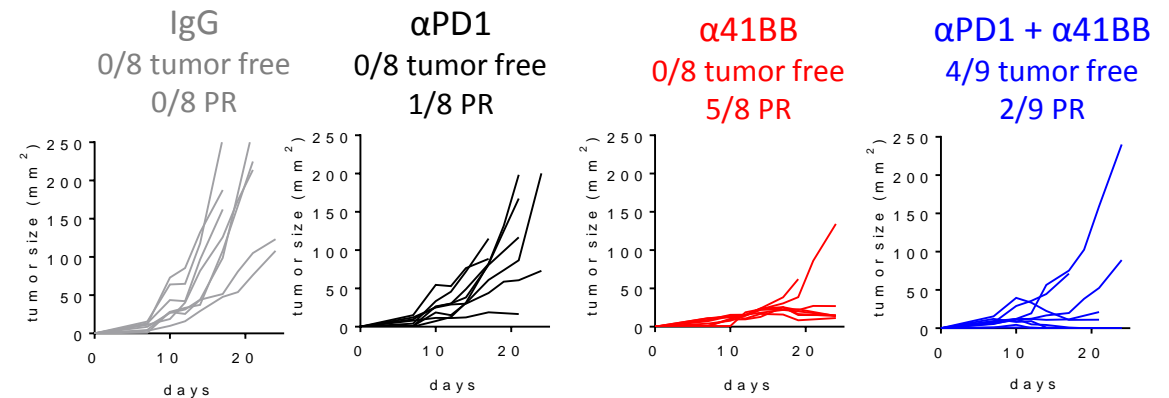
How does metabolic support by 41BB impact immunotherapy?

- Adoptive cell therapy using activated OT-I T cells transferred into B16^{OVA} bearing mice results in a very limited response
- ACT into mice receiving 41BB agonists results in potent synergy resulting in a majority of mice experiencing complete regression
- This 41BB effect requires the expression of PGC1 α by the therapeutic T cell



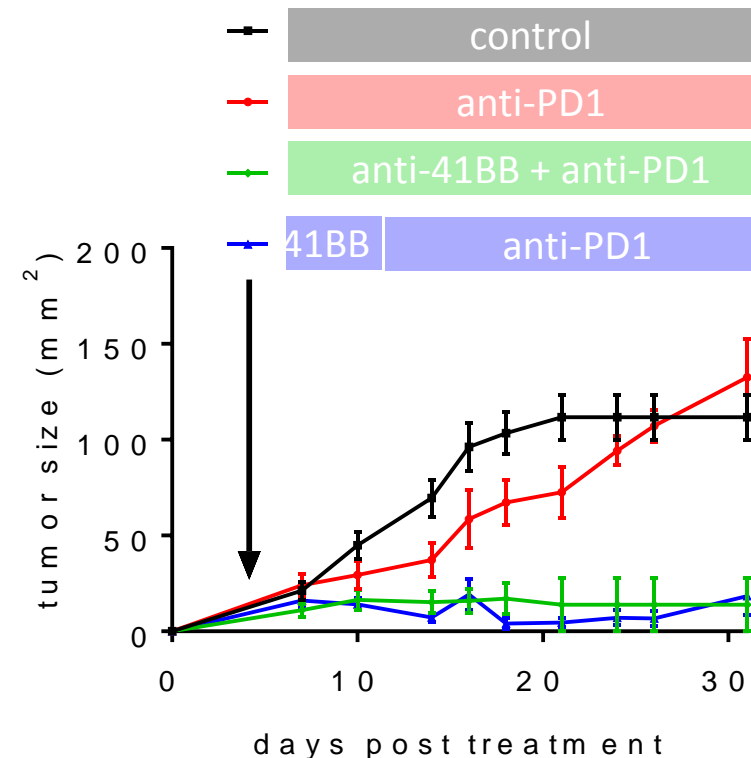
41BB also impacts checkpoint blockade immunotherapy

- As previously reported, 41BB combinatorial therapy promotes tumor clearance
- PGC1 α is significantly upregulated in TIL of 41BB treated animals
- 41BB agonism reverses the mitochondrial insufficiency observed in TIL



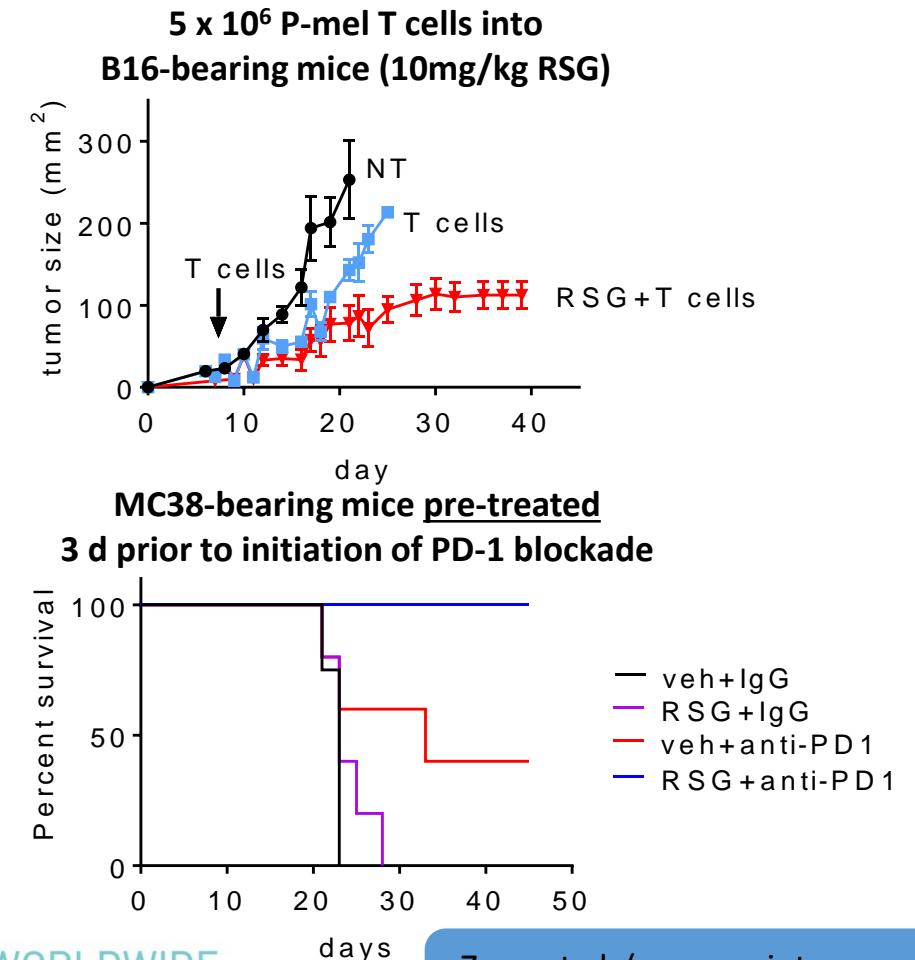
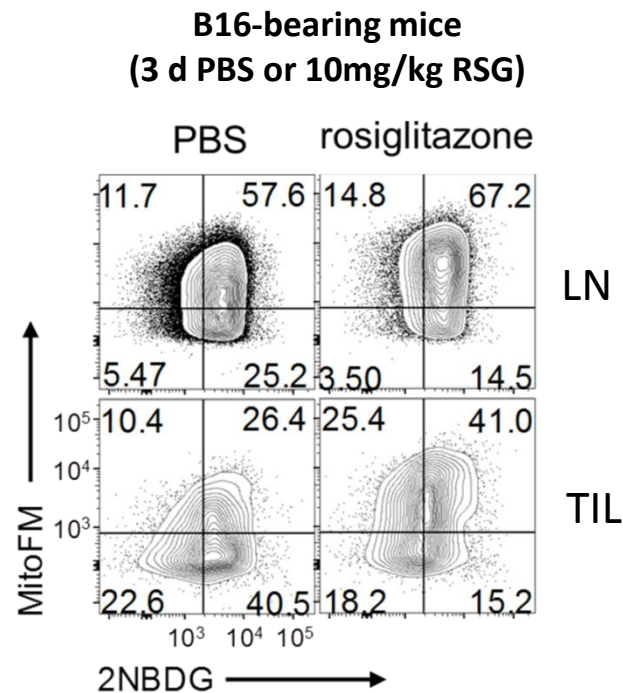
Does metabolic support need to be continuous?

- 41BB has been studied clinically as an immunotherapeutic target, but with significant toxicities
- Our data suggested though that the metabolic support afforded by 41BB ligation occurred quickly and persisted after cessation of treatment
- We asked in our mouse model whether a short-course of 41BB treatment could provide similar therapeutic benefits
- Treatment of mice with 41BB for just three days sensitized mice to PD-1 immunotherapy, with results indistinguishable from the continuous co-treatment



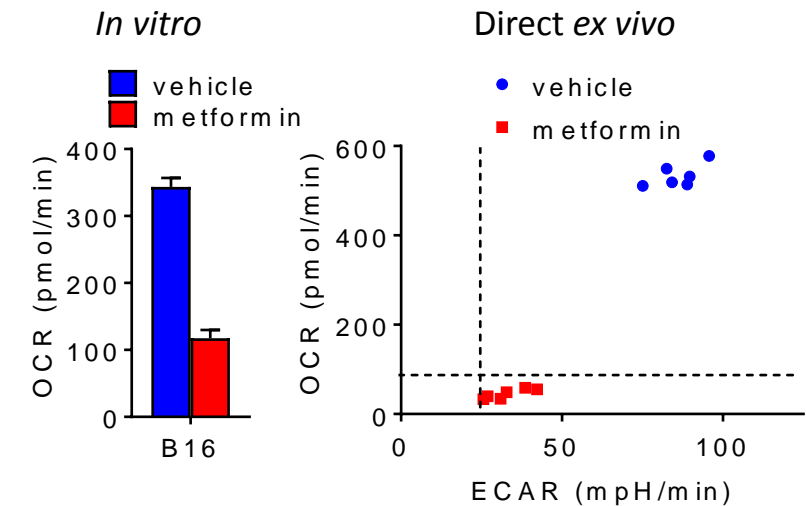
Does metabolic support need to be so heroic?

- Rosiglitazone, a synthetic PPAR agonist, can stimulate mitochondrial biogenesis in tumor infiltrating and peripheral T cells
- Glitazones can improve adoptive T cell therapy
- Even three days of pre-treatment with RSG can prime T cells for response to checkpoint blockade



The elephant in the room... the tumor cell

- Our data so far has focused on providing a T cell with a metabolic advantage
- However, it is competition with the **tumor cell** that drives the need for this energetic support
- Analysis of tumor cell metabolism and response to immunotherapy suggested it was the oxidative axis that was especially detrimental to T cell function
- We identified the widely prescribed type II diabetes drug **metformin** as a tumor microenvironment remodeler
- Metformin acts in part as a (weak) complex I inhibitor, inhibiting mitochondrial oxygen consumption
- Epidemiological studies have revealed that individuals taking metformin have a reduced risk of cancer and do better on various therapies
- Preclinical data suggest that the anti-tumor metformin effect requires T cells



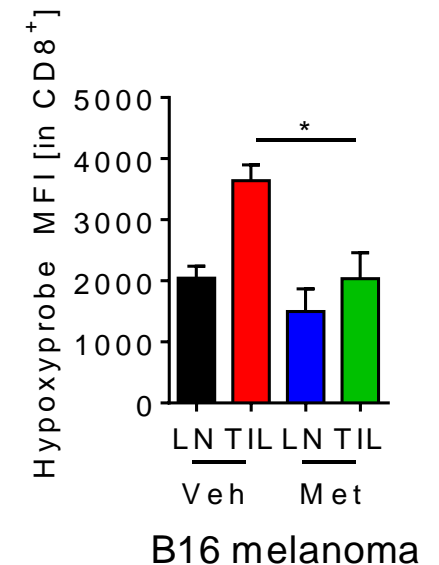
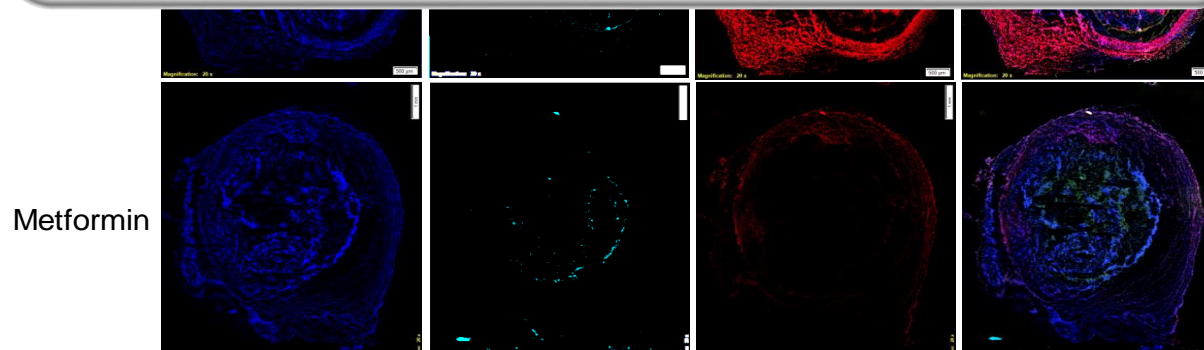
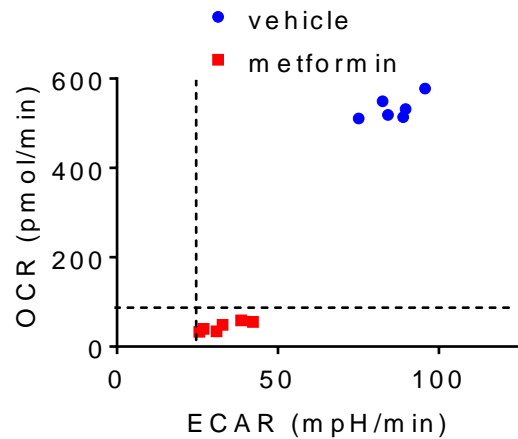
- Thus, we hypothesized that metformin might act by inhibiting tumor oxygen consumption, generating a less hypoxic environment for T cells

Does metformin inhibit tumor cell oxygen consumption?

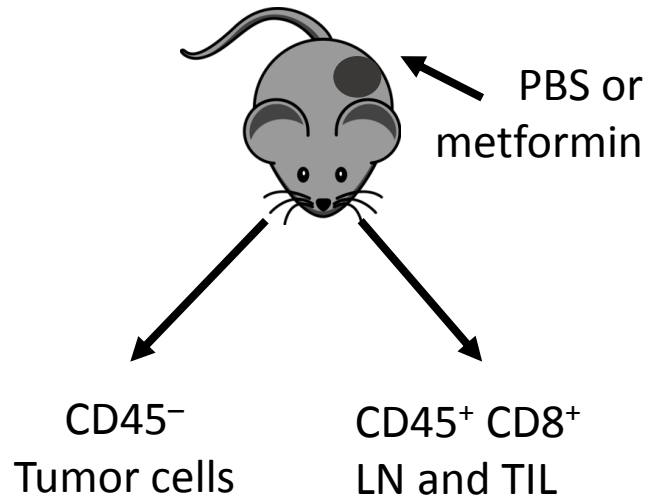
- Metformin treatment of tumor bearing mice remodels TME metabolism
- Loss of oxidative metabolism in tumor cells reduces tumor hypoxia
- Metformin treatment reduces tumor hypoxia experienced by T cells *in vivo*

B16-bearing mice treated 4 days with metformin

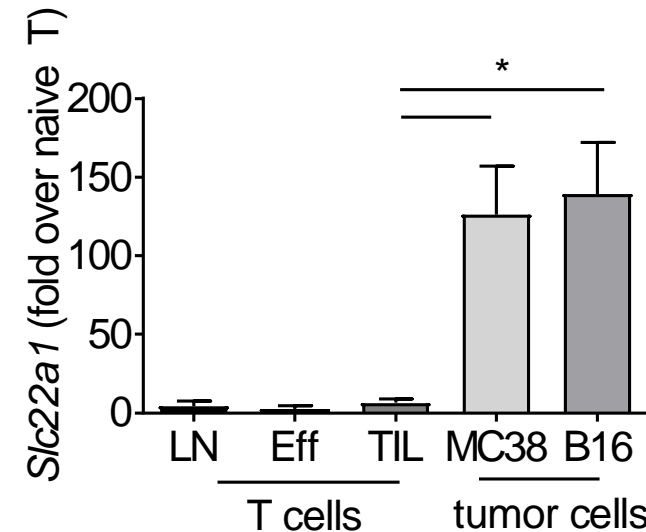
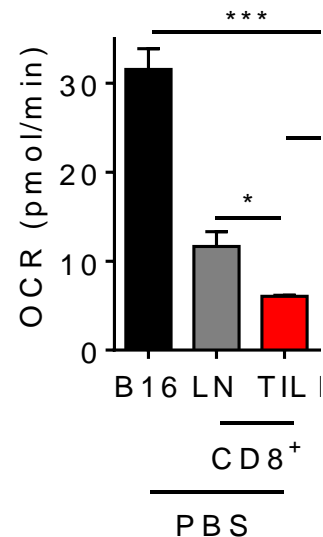
Metformin-mediated inhibition of tumor cell metabolism reduces tumor microenvironment hypoxia



How does metformin affect T cell metabolism?

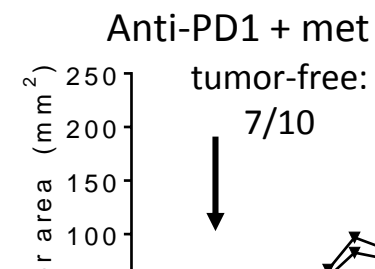
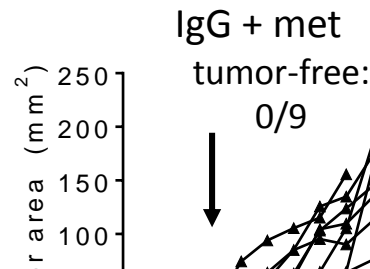
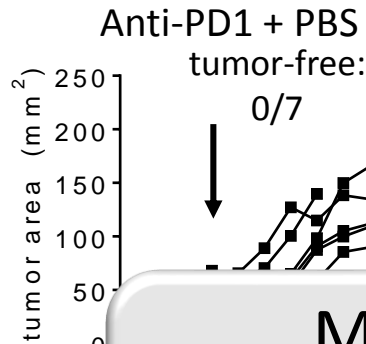
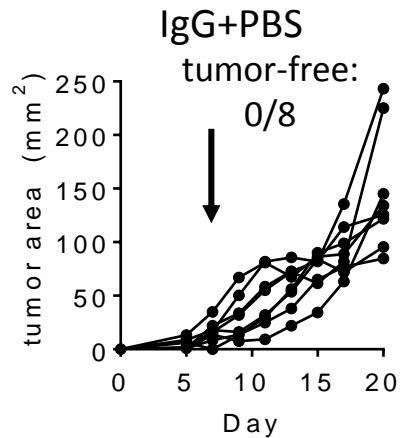


- Tumor cell metabolism dwarfs that of TIL metabolism, consistent with many previous reports
- Metformin treatment inhibits tumor cell OCR, resulting in increased TIL OCR direct *ex vivo*
- Tumor cells express much higher levels of the metformin transporter OCT1 (encoded by *Slc22a1*), suggesting the effect of metformin on TIL T cells is *indirect*

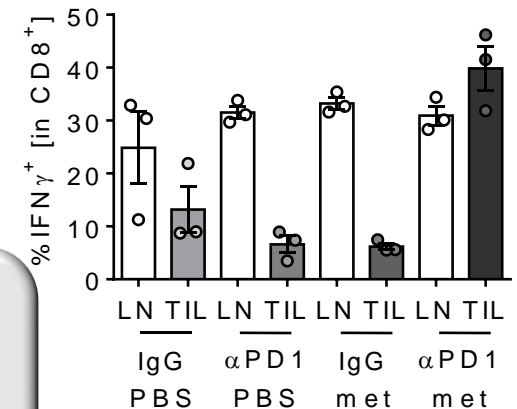


Does metformin treatment synergize with immunostimulatory therapies?

B16



Metformin acts to metabolically remodel the tumor microenvironment to be more permissive to immunotherapy

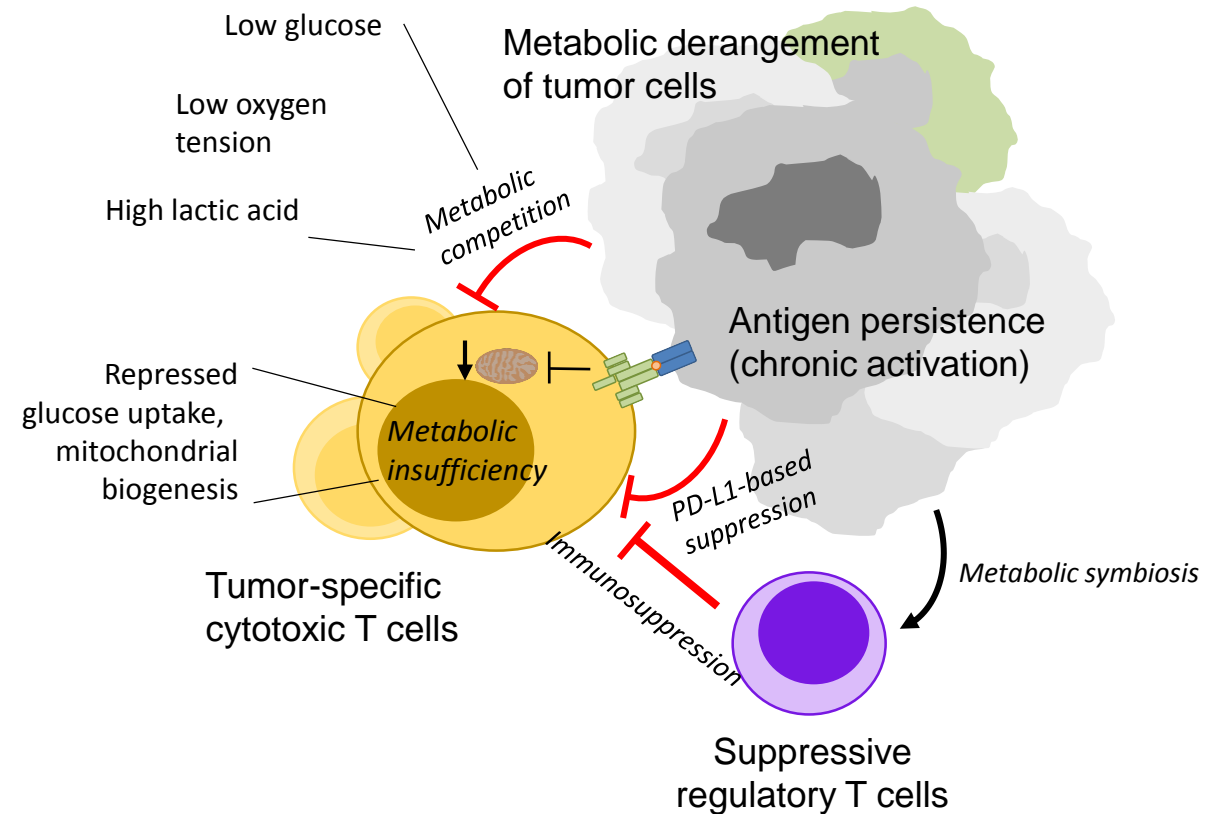


These exciting preclinical data have led to **funded/in development clinical trials combining metformin and PD-1 blockade in several human cancers**

- Metastatic melanoma
- Locally advanced/recurrent-metastatic head and neck cancer
- Other PD-1 refractory indications

Metabolic barriers to antitumor immunity

- T cells must be reactivated immunologically and bolstered metabolically to fully support their effector function in the tumor microenvironment
- Genetic engineering, costimulatory immunotherapy, or pharmacologic rejuvenation may help counteract these pathways and create superior therapeutic T cells
- Metabolic intervention need not be continuous, although sequencing is likely important
- Targeting oxidative or glycolytic tumor cell metabolism can remodel the microenvironment to be permissive to immunotherapy (and may have differential effects on effector versus regulatory populations)



• Delgoffe Lab

- Nicole Scharping
- Ashley Menk
- *Lucy Zeng*
- Dayana Rivadeneira
- McLane Watson
- Paolo Vignali
- *Becca Moreci*

- Robert Ferris
- Yana Najjar
- Dario A. A. Vignali
- Mark Shlomchik

• Abigail Overacre-Delgoffe

• Jonathan Powell

Interested in a postdoc in the
Delgoffe lab? Contact me!

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Immunometabolism in full effect here at SITC

P156 - Nicole (metabolic underpinnings of T cell exhaustion)

P152 - Ashley (41BB-mediated reprogramming)

P484 - Dayana (oncolytic virus reprogramming)

Coming up soon in this session... Mac (T_{reg} cell metabolism)

ADVANCING CANCER IMMUNOTHERAPY WORLDWIDE

- Center for Biologic Imaging
- UPMC Flow Cytometry/Animal Facilities
- Tumor Microenvironment Center
- Department of Immunology
- **Patients and their families**

• Funding sources:

- Sidney Kimmel Foundation for Cancer Research
- Stand Up 2 Cancer Innovative Research Grant
- Melanoma and Skin Cancer SPORE
- Head and Neck Cancer SPORE
- Alliance for Cancer Gene Therapy
- NIH Director's New Innovator Award (DP2AI136598)

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