

SITC 2019

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& Convention Center

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Society for Immunotherapy of Cancer



Characteristics of the microbiome of complete responders to Anti-PD1 and healthy individuals: Implications for donor selection and clinical trial design

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Society for Immunotherapy of Cancer

#SITC2019

Disclosure

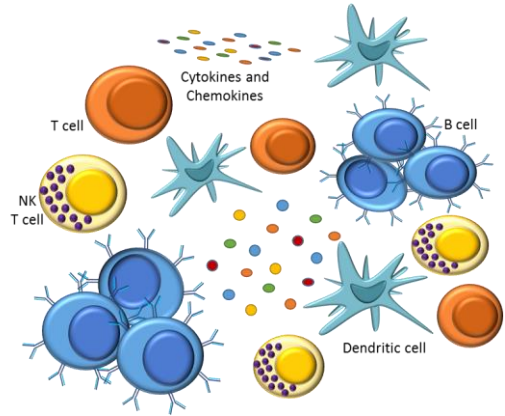
- I have no potential disclosure with this presentation
- My PI (Jennifer Wargo) is a co-inventor on patent submitted by The University of Texas MD Anderson Cancer Center to the US Patent and Trademark Office on modulating gut microbes to improve responses to immune checkpoint blockade (Patent # PCT/US1/53717)

Responses are dependent on factors shaping tumor growth and immunity

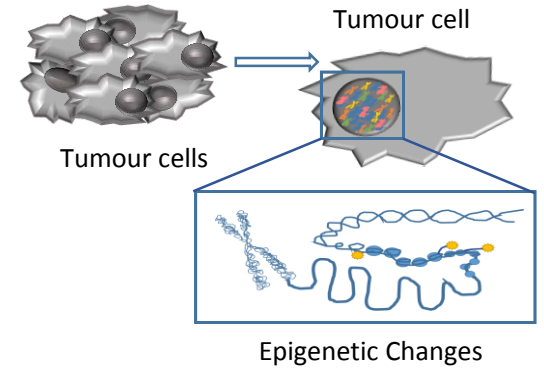
Cogdill, Andrews, Wargo - *British Journal of Cancer* May 2017

Systemic Immunity

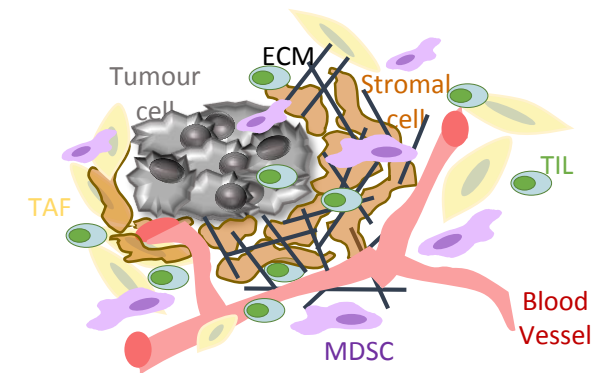
Innate and Adaptive



Tumour Genome and Epigenome



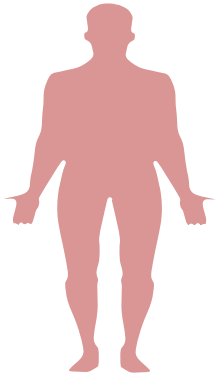
Tumour Microenvironment



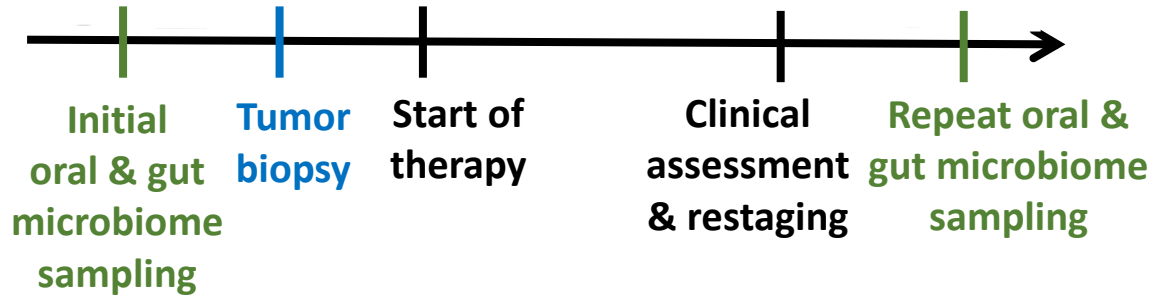
There is a significant microbial contribution to the total makeup of our cellular composition as well as our DNA that dramatically influences our physiology



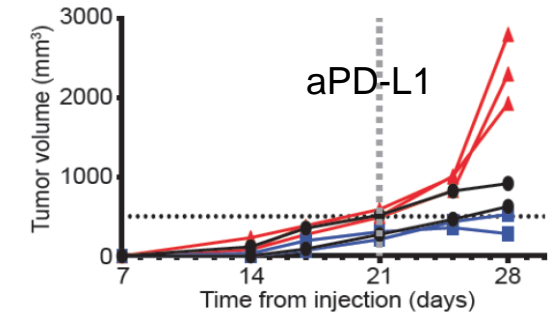
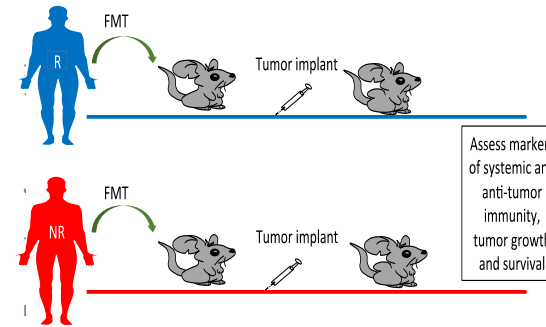
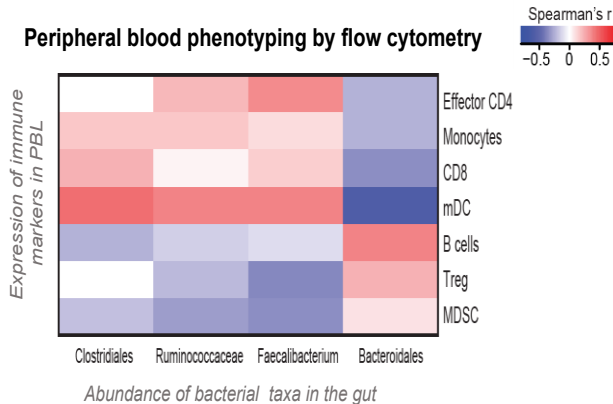
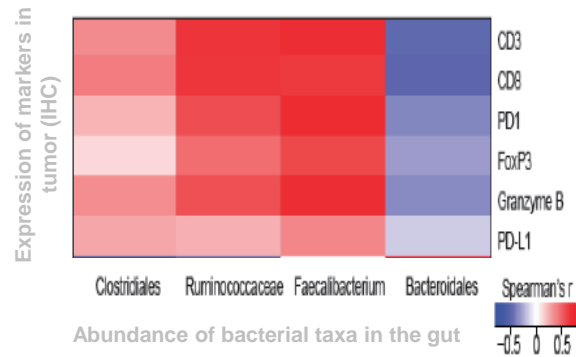
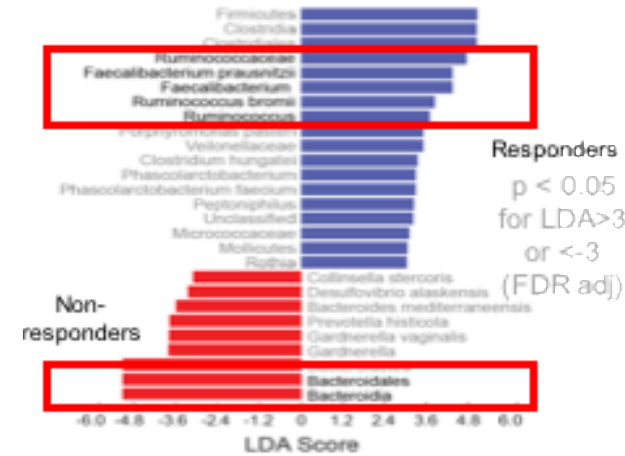
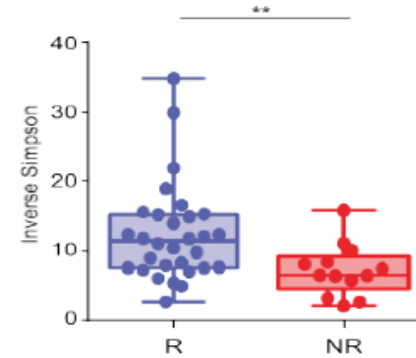
We studied oral and gut (fecal) microbiome signatures in patients with melanoma



n = 233 patients



Responders to anti-PD-1 had a higher diversity and differential composition of gut bacteria



Importantly, these “favorable” signatures in the gut microbiome were associated with better immune profiles in the tumor and in the blood of patients, with validation in murine models



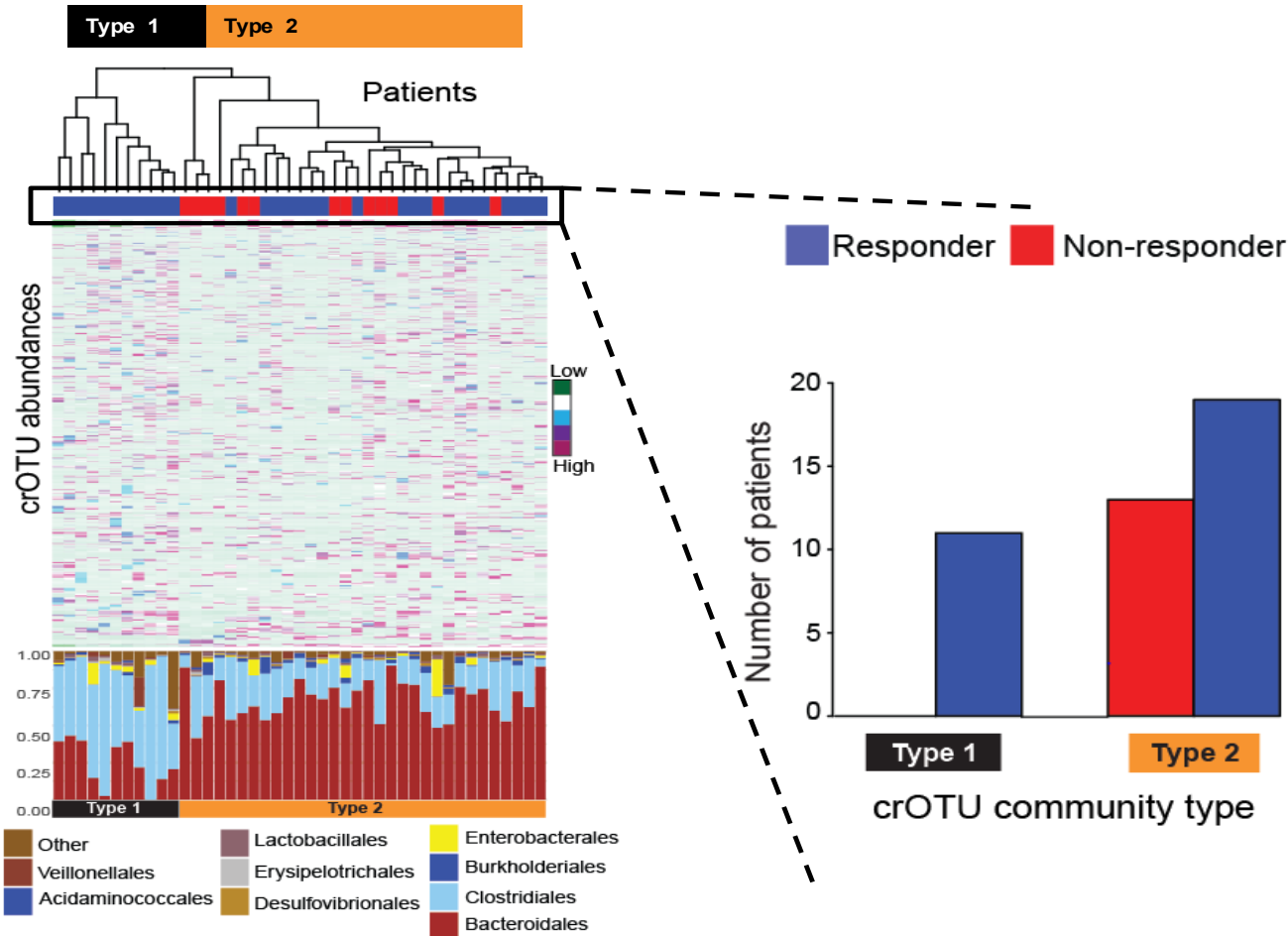
Deepak Gopalakrishnan PhD

Gopalakrishnan, Spencer et al, Science 2018

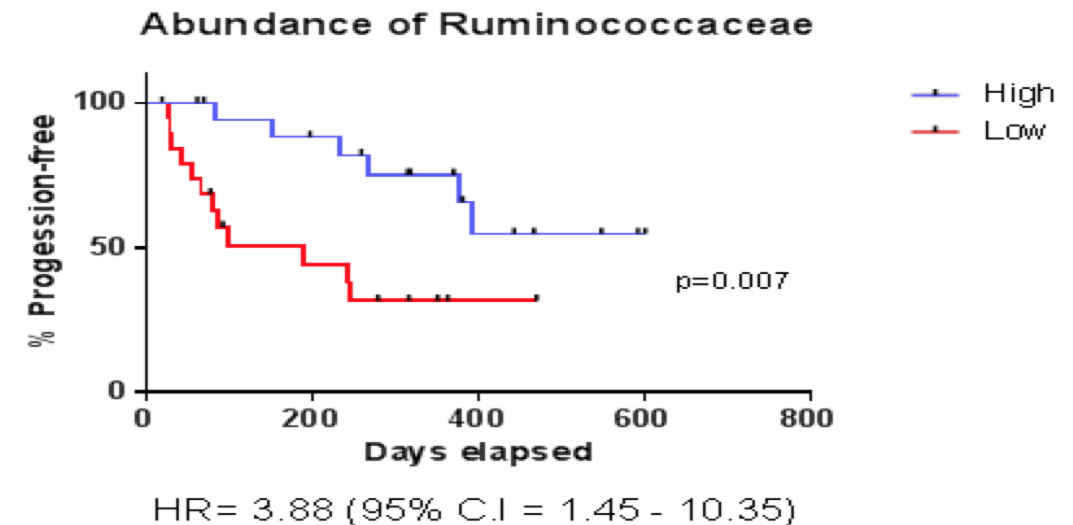
Christine Spencer PhD



In our cohort, we identified a gut microbiome “signature” with a high likelihood of response to anti-PD-1 (type I), with subsequent validation in a larger cohort

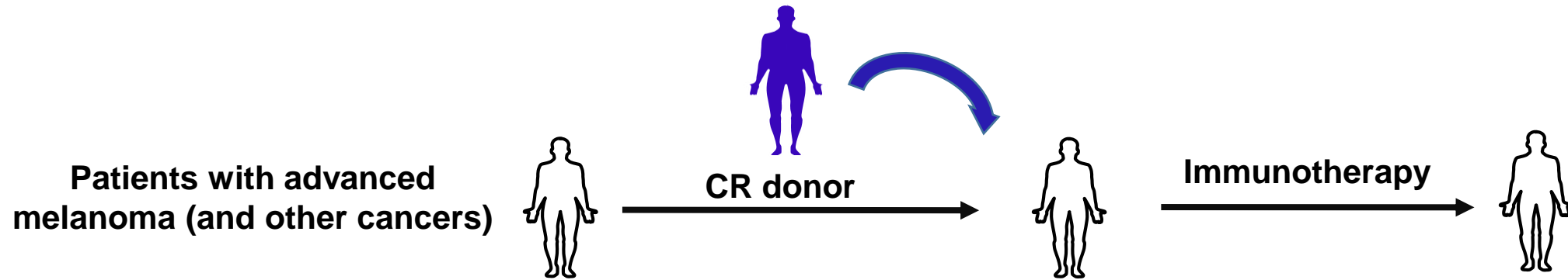


Notably, the relative abundance of Ruminococcaceae was a good surrogate marker for a type I signature, with patients having a high abundance doing better on therapy



Gopalakrishnan et al, Science 2018 + confidential unpublished data * DO NOT POST *

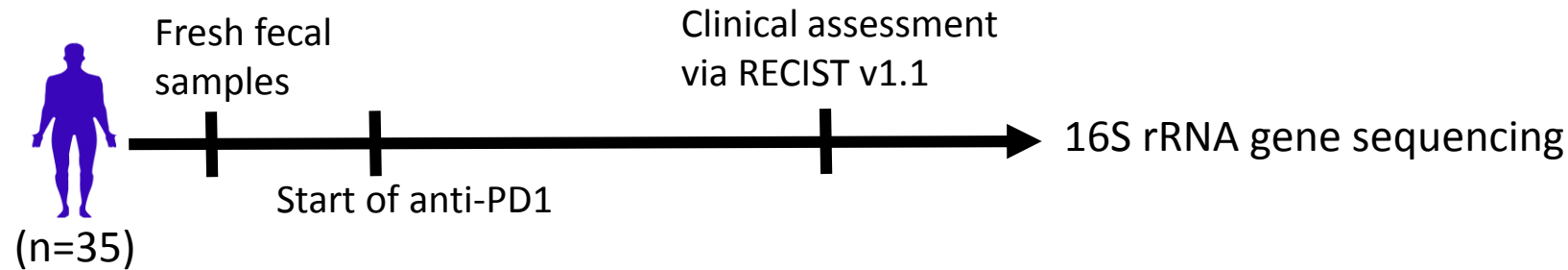
Based on these findings, studies are underway aiming to modulate the gut microbiome using fecal microbiota transplant (FMT) and other strategies
(with some success as recently reported by 2 groups at AACR)



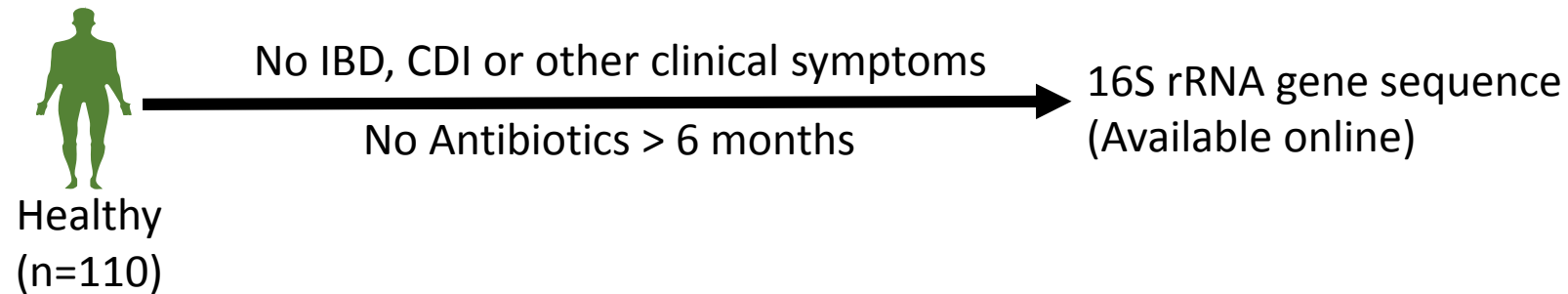
*However not all complete responders have what we would consider to be a “favorable” gut microbiome,
And there is the potential to explore the use of FMT from healthy individuals in these types of studies*

We studied gut microbiome signatures in complete responders vs. healthy individuals

Complete Responder (CR) cohort: Metastatic melanoma patients on Anti-PD1

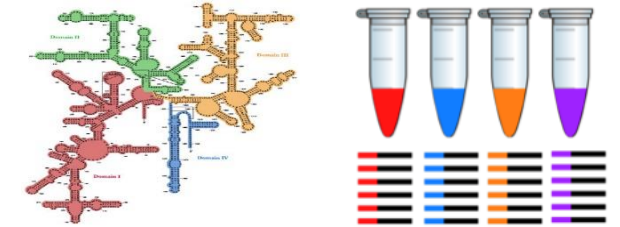


Healthy cohort: sequence data from American Gut Project (McDonald et al. 2018)



16S rRNA: Microbial Taxonomy

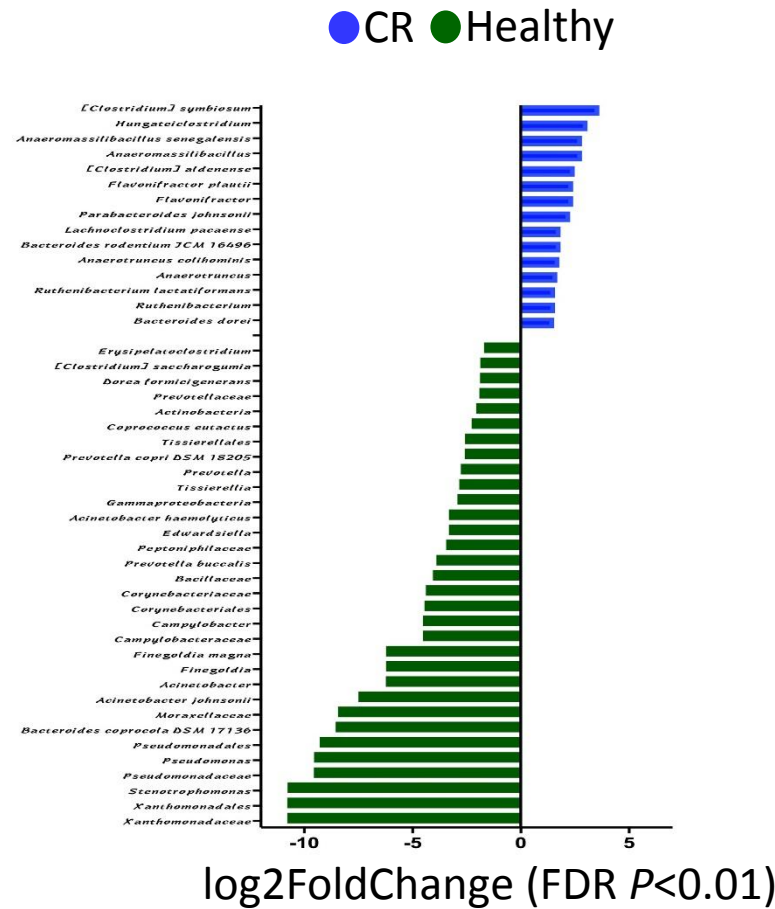
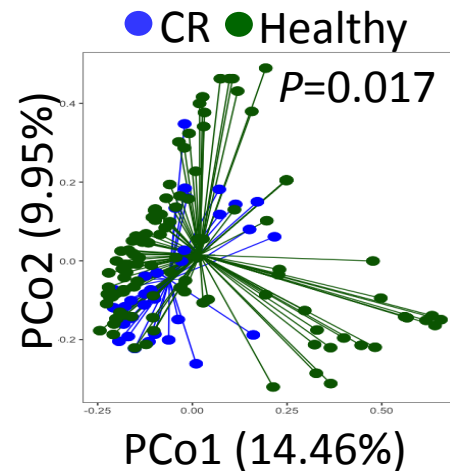
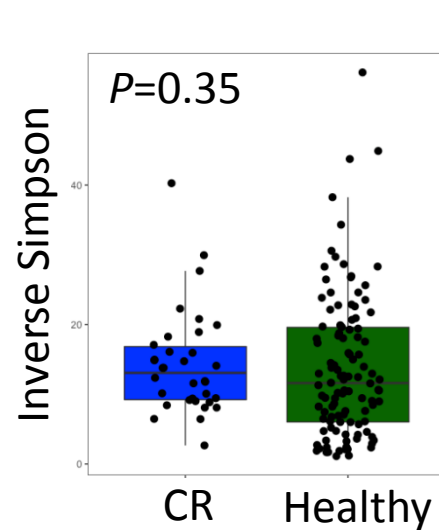
Illumina MiSeq



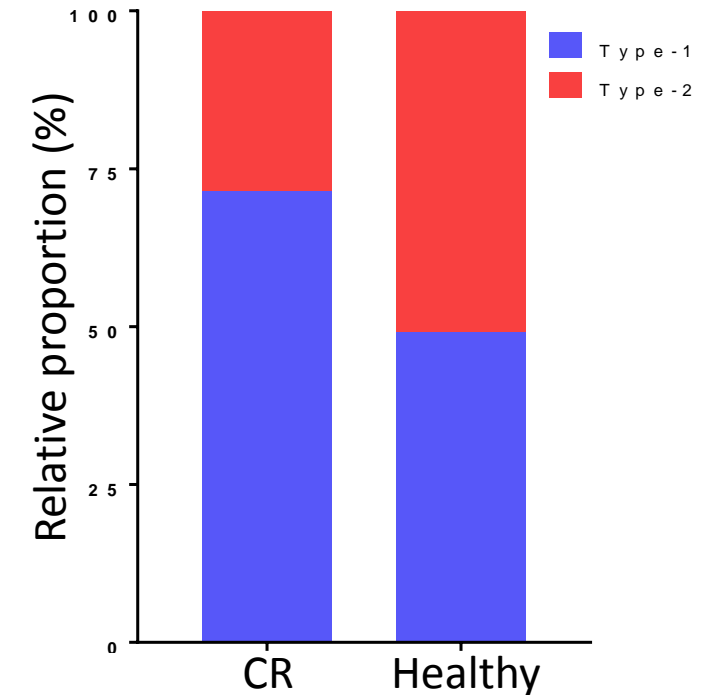
1. V4 region of 16S rRNA gene: Taxonomy by NCBI. Analyzed by QIIME, USEARCH, vsearch, R
2. PICRUST2 (Douglas et al. 2019): No. of reference genomes 20,000. Functional genes: KEGG. Metabolic pathways: MetaCyc

Murine Melanoma Model

In these studies, CRs and healthy donors had diverse microbiomes overall, with differential composition and proportions of a type I signature between the groups



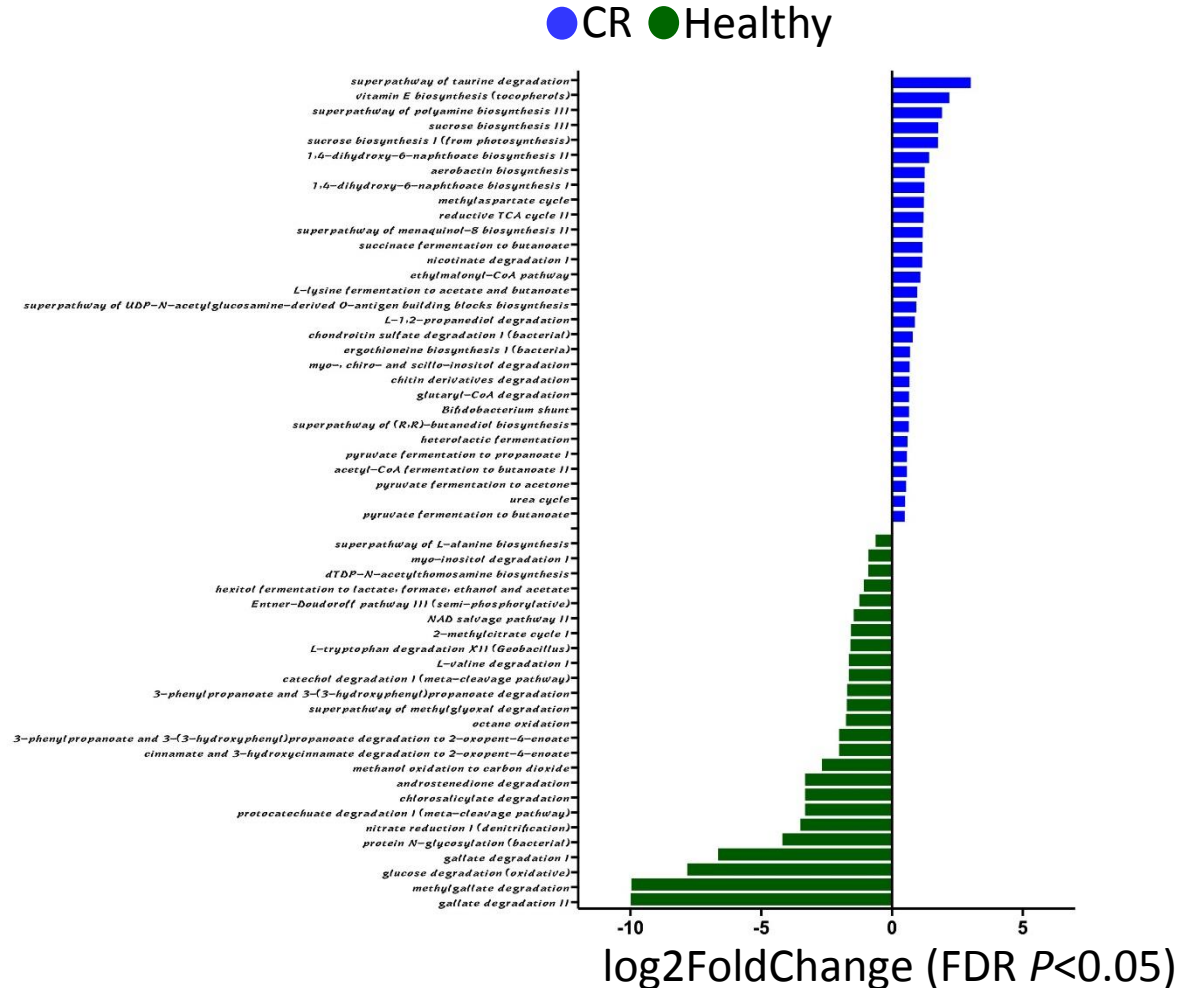
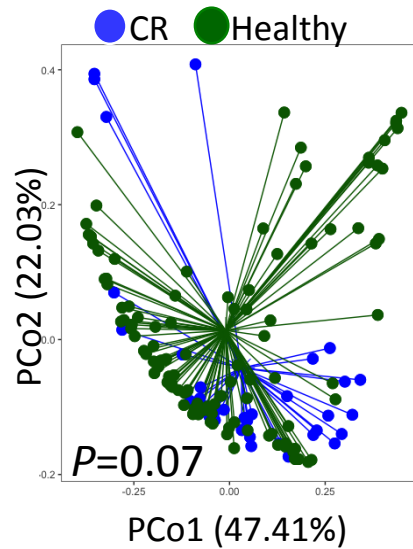
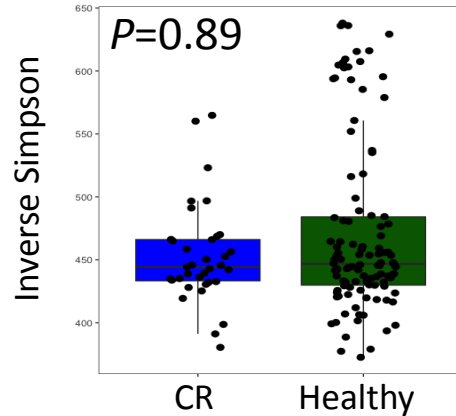
Not all CRs have Type-1 signature



Confidential unpublished data * DO NOT POST *

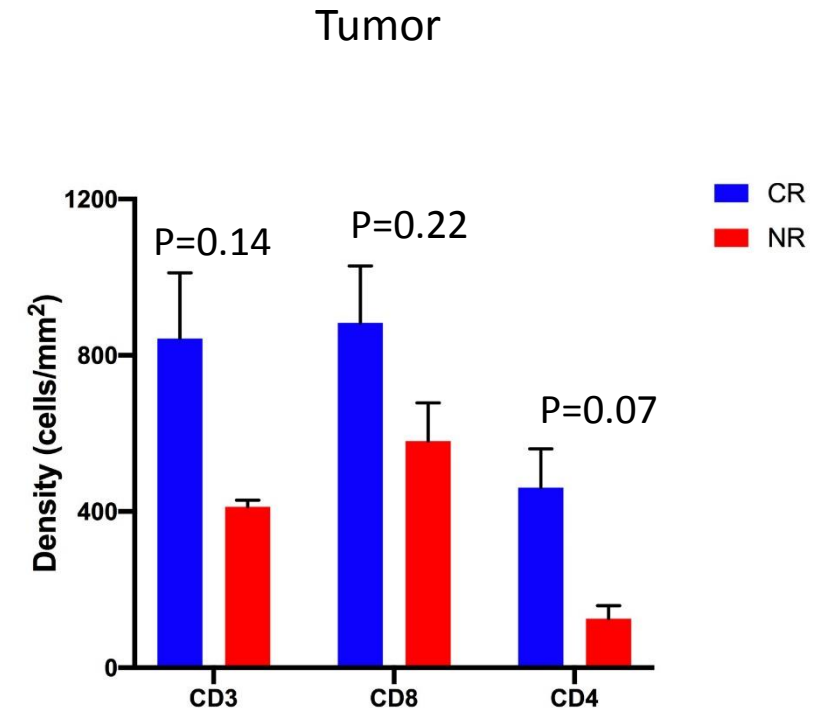
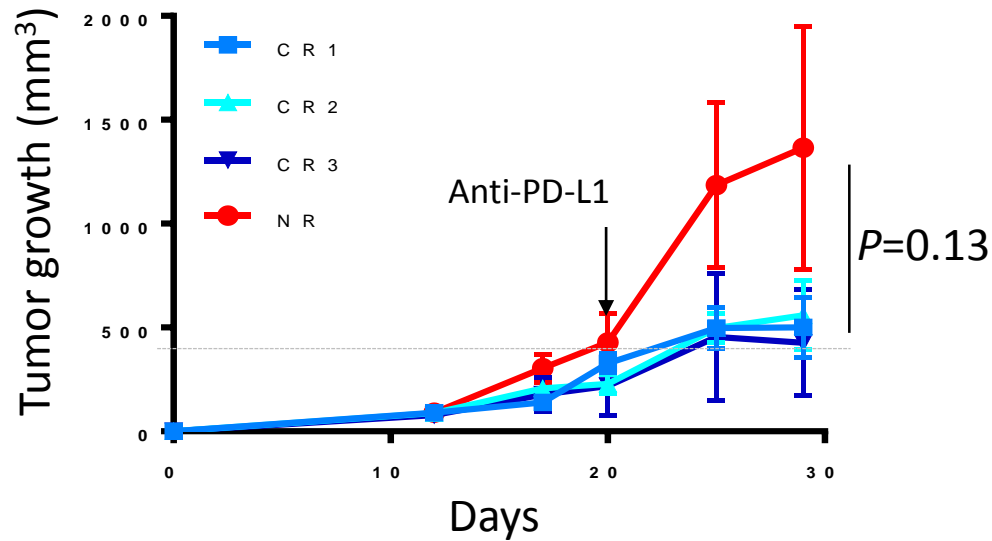
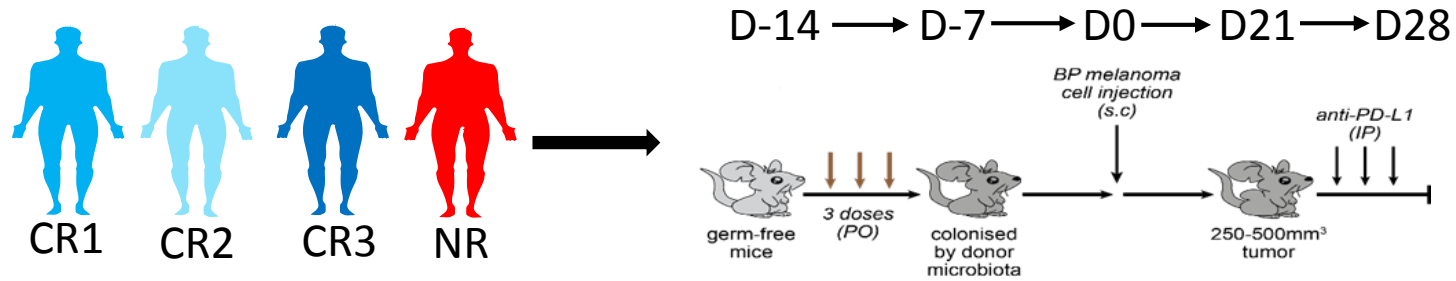
Functional differences between CRs and healthy individuals were also noted, though these need to be interrogated with dedicated metabolomic profiling

Functional genes diversity



Confidential unpublished data * DO NOT POST *

We performed pilot studies in germ-free mouse models, demonstrating good tumor control in mice receiving CR donor FMT (compared to non-responder FMT)



Confidential unpublished data * DO NOT POST *

Take Home Messages and Next Steps

- There is now strong evidence for the impact of gut microbes in immunotherapy response, though optimal strategies to modulate gut microbes remain incompletely understood
- Treatment with FMT has been associated with responses, however optimal donors remain unknown (and there are some risks associated with FMT)
- Further studies to better understand determinants of enhanced response are needed, and are currently underway (including deeper studies in larger cohorts, and studies using highly selected FMT donors)
- There is still a great deal to learn, and this is best accomplished through collaboration (*and we owe this to our patients!*)

Acknowledgements



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Patients & their families!!

Biospecimen collection team

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Sakara Waller

Elizabeth Sirmans

Eliza Posada

Baylor CMMR-sequencing team

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Julie Gardner



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for Cancer Research

Extra slide