



Multiple Myeloma and disparities

Irene Ghobrial, MD

Lavine Family Chair of Preventative Cancer Therapy

Professor of Medicine

Harvard Medical School

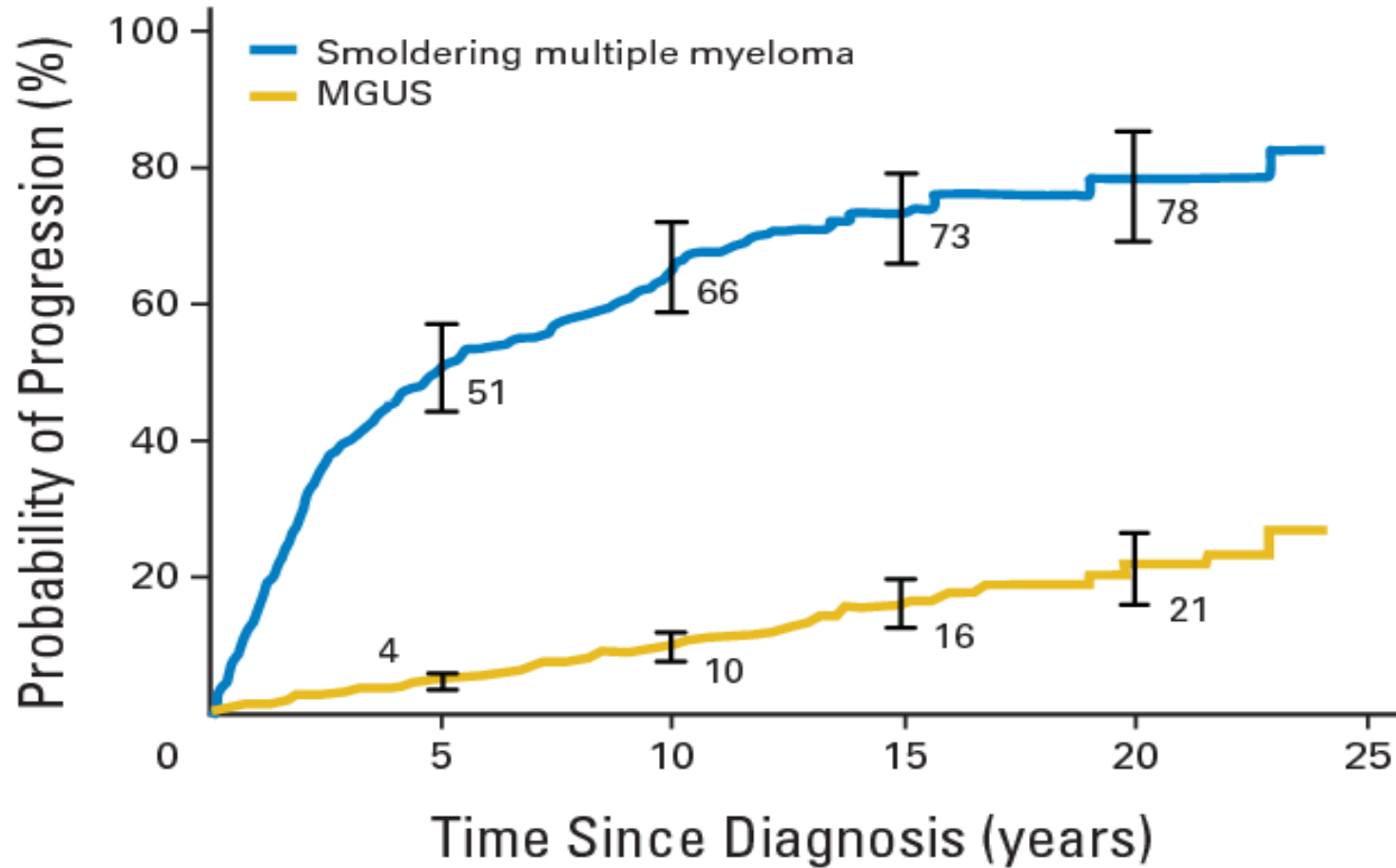
Dana-Farber Cancer Institute

Boston, MA



Dana-Farber
Cancer Institute

Multiple myeloma is always preceded by MGUS and SMM



Myeloma is the most common blood cancer in African Americans

And the incidence is growing....

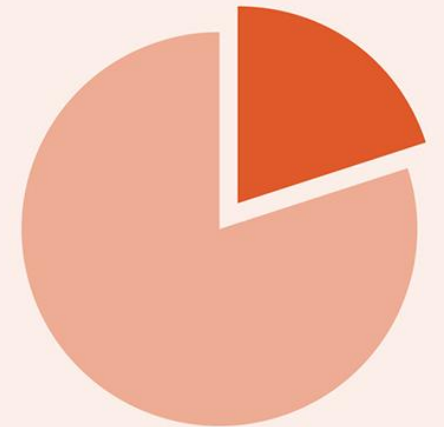
African Americans have **>2x the incidence rate** of MM compared to white Americans¹

By 2034 it is estimated that African Americans will make up roughly **24% of the newly diagnosed** MM population¹

African Americans currently represent about

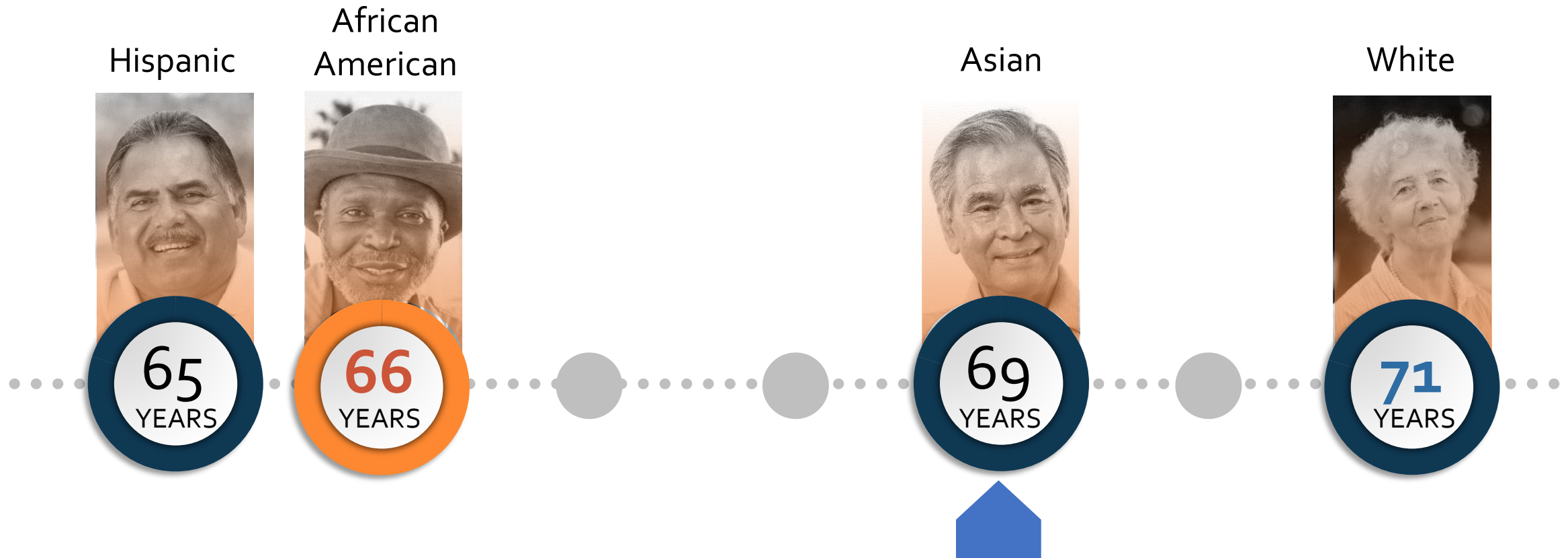
20% or **1 out of 5**

patients living with multiple myeloma



¹American Cancer Society. *Cancer Facts and Figures for African Americans 2019-2021*.

African Americans are younger at diagnosis by about 5 years



There is a **LONGER** time from symptom onset to diagnosis in African Americans

The average myeloma patient sees their primary care doctor **THREE** times with symptoms and signs consistent with MM.

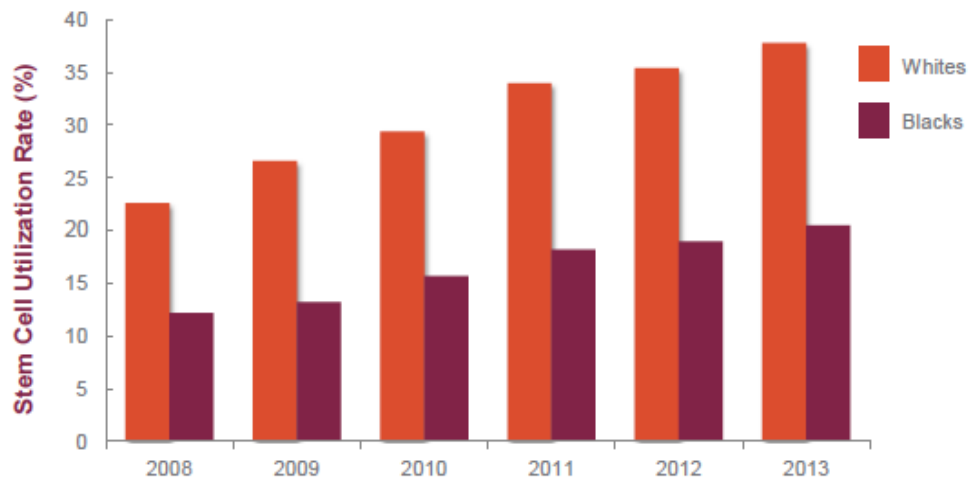
- Confounding diagnoses (like diabetes), Access to diagnostics and care, Awareness in primary care providers, Timely referral to specialists...

African Americans have only HALF the survival of White Americans



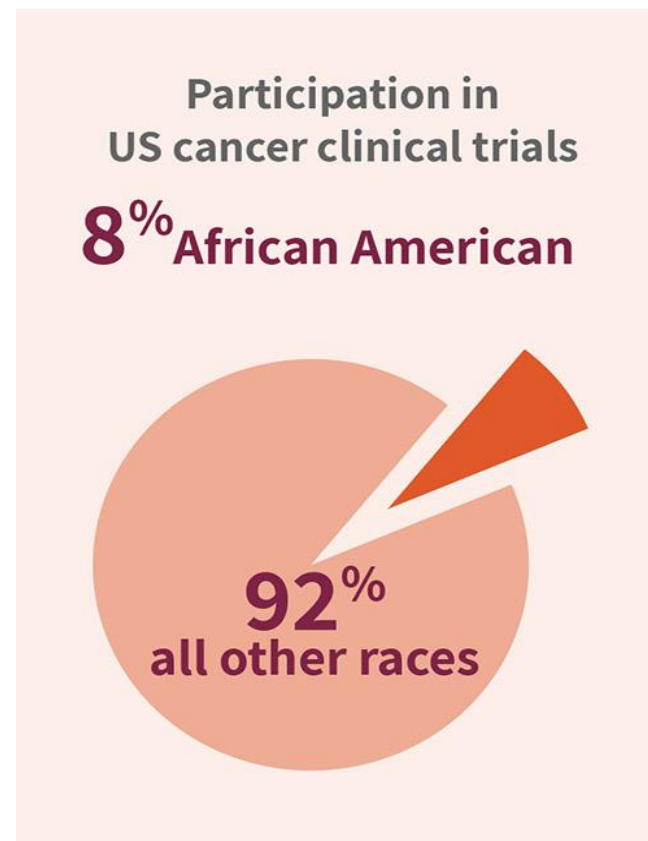
Huge progress in survival in MM but this has not been realized to the same extent in African Americans

An analysis from the Center for International Blood and Marrow Transplant Research Database (CIBMTR, N=28,450) showed increased utilization differed by race



Less likely to receive the critical treatments for MM – The 3 Ts: Triplets, Transplants, and CAR T cell therapy

Less likely to participate in clinical trials (the 4th T)



Gormley et al, Blood Cancer Discovery 2021

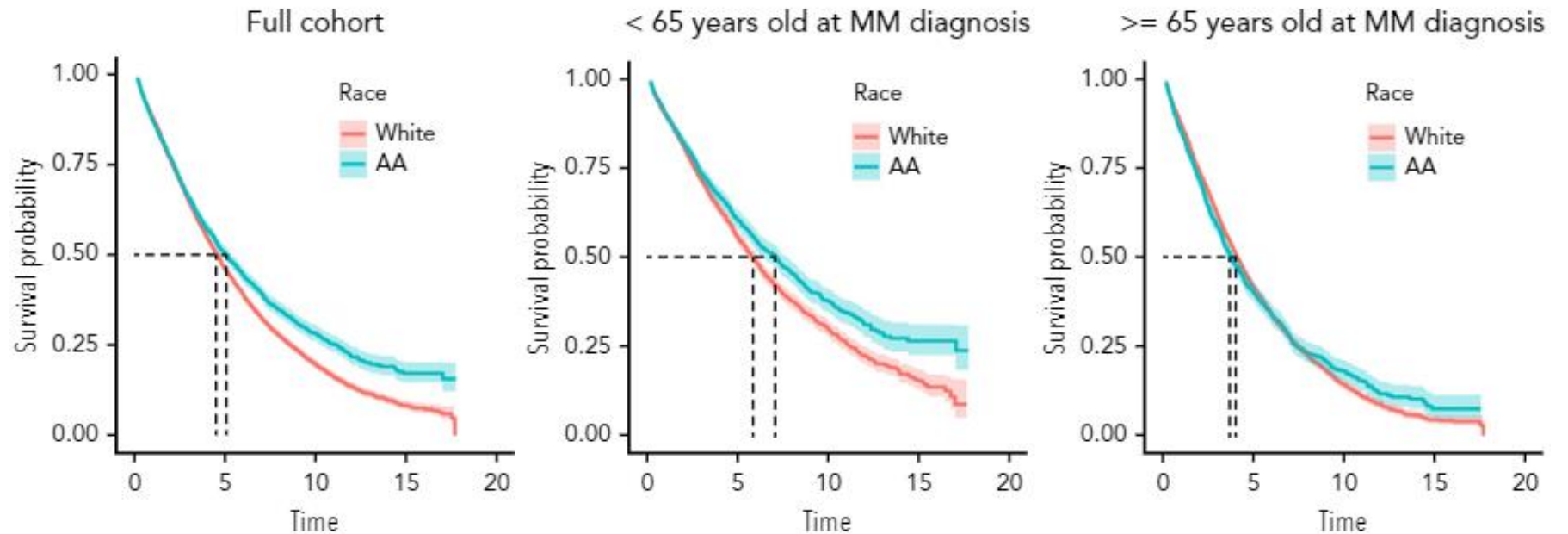
When African Americans receive equal access to care, their survival outcomes are equal, and at times, better than White patients

SCIENCE IN SOCIETY

Recommendations on Eliminating Racial Disparities in Multiple Myeloma Therapies: A Step toward Achieving Equity in Healthcare



Nicole Gormley¹, Lola Fashoyin-Aje¹, Trevan Locke², Joseph M. Unger³, Richard F. Little⁴, Ajay Nooka⁵, Khalid Mezzi⁶, Mihaela Popa-McKiver⁷, Rachel Kobos⁸, Yelak Biru⁹, Tiffany H. Williams¹⁰, and Kenneth C. Anderson¹¹



African ancestry associated with less aggressive disease

Higher prevalence of ^[a]:

t(11;14)

t(14;16)

t(14;20)

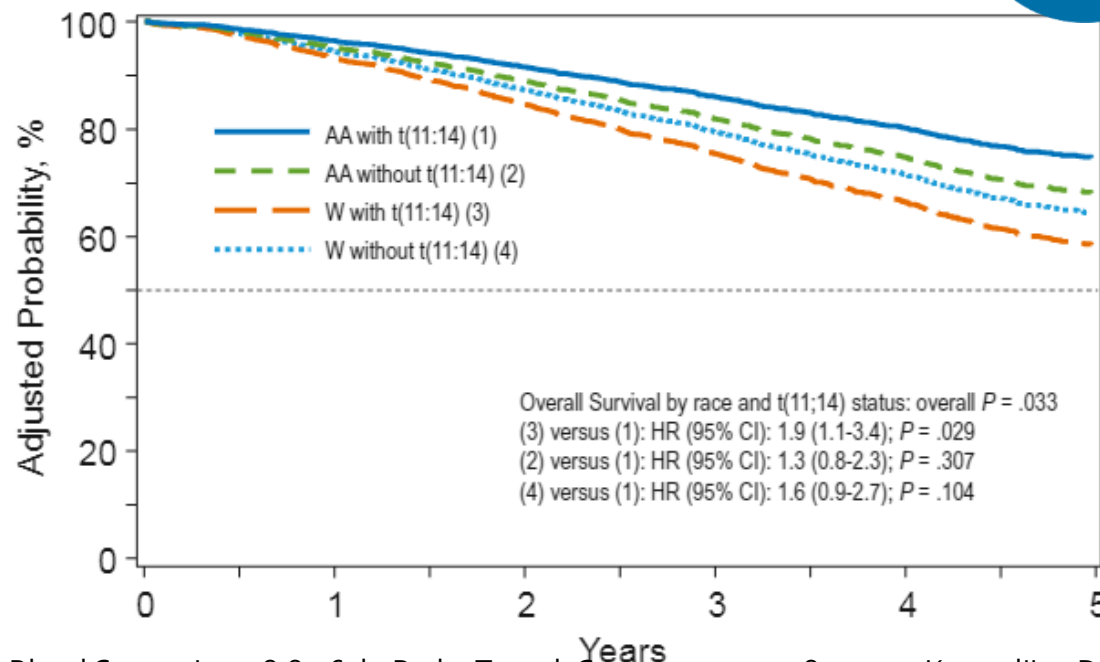


Lower prevalence^[c]:

13q deletion

17p deletion

- Absence of 17p deletion associated with better survival among younger African Americans vs White counterparts^[d]



There are biologic differences in African Americans of lower risk disease

MM patients with the **highest levels of African ancestry** demonstrate

- a **higher** prevalence of:
 - t(11:14)
 - t(14;16)
 - t(14;20)
- a **lower** prevalence of:
 - 13q deletion
 - **17p deletion**
 - However, if present in AA patients under the age of 65, median survival rate is less compared to Whites

Cytogenetic differences (RVD 1000) - Emory

| Cytogenetic abnormality | Caucasians (N=619) | African-American (N=352) | P-value |
|-------------------------|--------------------|--------------------------|---------|
| 1q gains | 111 (18.8%) | 37 (10.8%) | 0.001 |
| T(11;14) | 66 (11.5%) | 55 (16.1%) | 0.043 |
| T(4;14) | 25 (4.3%) | 18 (5.3%) | 0.512 |
| T(14;16) | 16 (2.8%) | 10 (2.9%) | 0.888 |
| del17p | 70 (12.1%) | 23 (6.7%) | 0.009 |
| del13 | 168 (29.2%) | 70 (20.5%) | 0.004 |

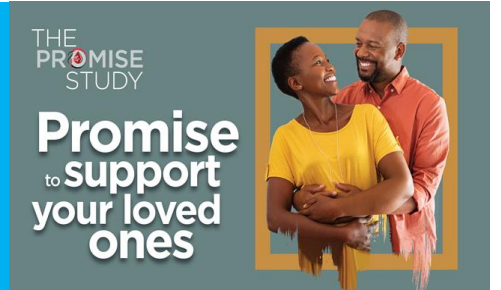
Ailawadhi et al. (2018); Baker et al. (2013); Baughn et al. (2018); Cirstea et al. (2019); Kazandjian et al. (2019); Munjuluri (2019)

Early detection and interception initiatives:

1

Screening Early

- Cancer screening saves lives
- A blood sample is easier than colonoscopy!
- High risk individuals have a risk of about 13% or more



2

Risk Stratification

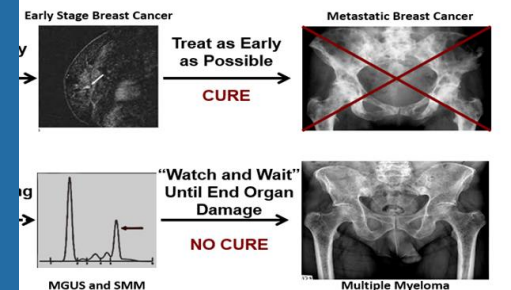
- Who is truly at risk of progression?
- Adding genomic and immune biomarkers for more precise risk assessment
- Blood instead of bone marrow



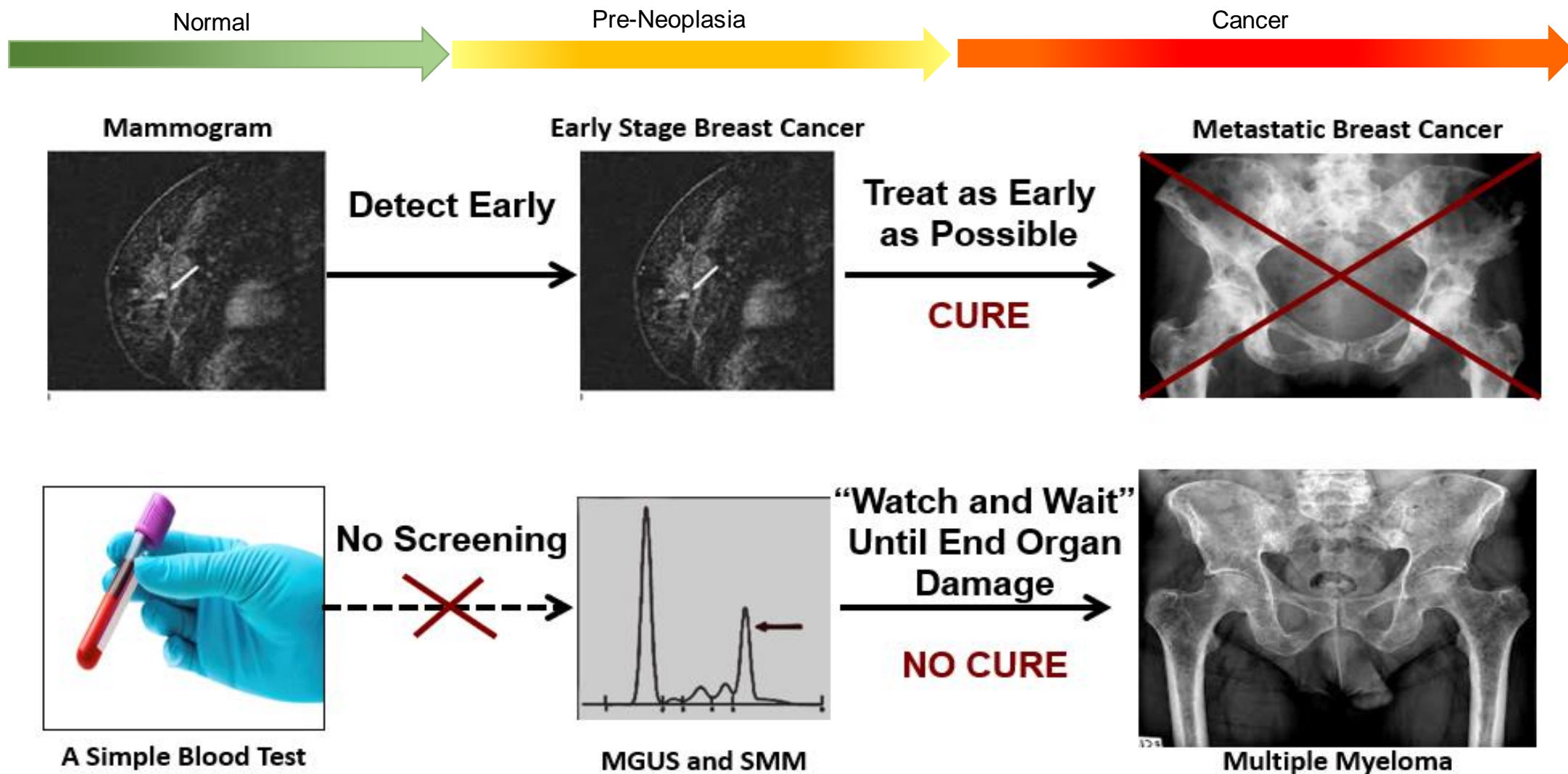
3

Early Interception

- More efficacious therapies that can cure early
- No clinical benefit of waiting until you have end organ damage
- Early use of late-stage therapies



We urgently need early detection in precursor stages of Multiple Myeloma



PROMISE

Predicting Progression of Developing Myeloma in a High-Risk Screened Population



How do I join?

To sign up, visit:

www.PromiseStudy.org

Or you can use
your phone to
scan this QR
code



What's involved?
Participants will:

- Complete a brief survey
- Sign a consent form
- Share a small blood sample

Sign up online. No travel to Dana-Farber is required. Visit your local QUEST Diagnostics and return your kit by mail.

As a token of our appreciation,
participants can request a \$50 gift card.



www.enroll.promisestudy.org

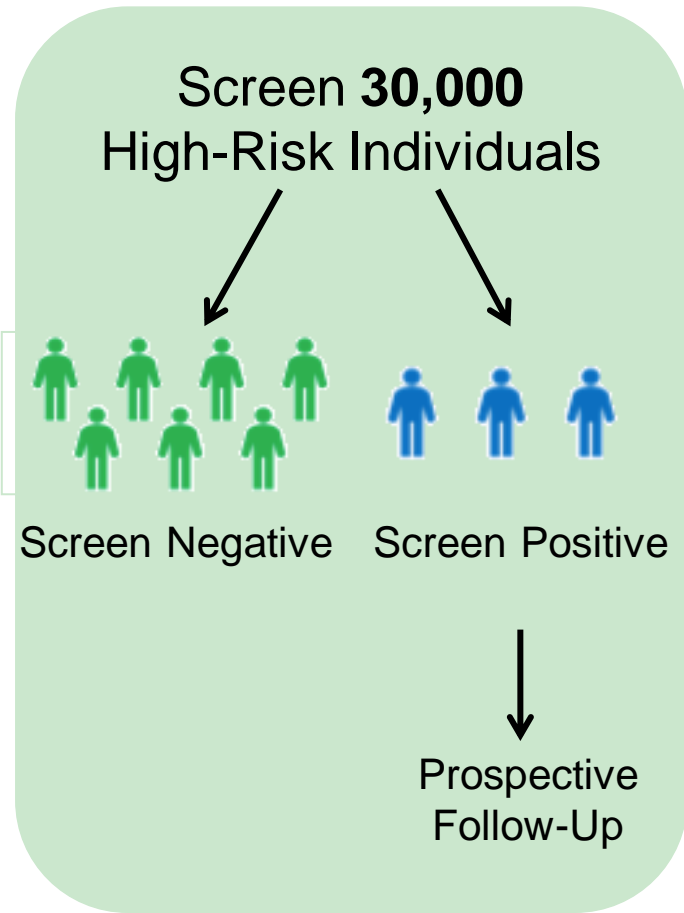
ClinicalTrials.gov Identifier: NCT03689595

Current Status: Actively Recruiting

Phase: Cohort Study

www.PromiseStudy.org

PROMISE



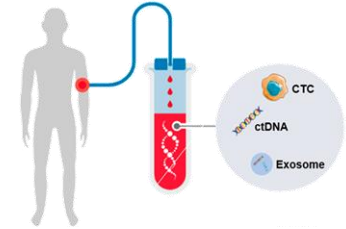
Genetics and Genomics

Viktor Adalsteinsson

Gad Getz

Irene Ghobrial

Develop novel **biomarkers for diagnosis**



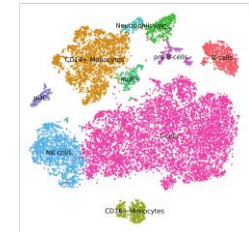
Epidemiology

Tim Rebbeck

Lorelei Mucci

Catherine Marinac

Establish new **risk stratification tools**



Bone Marrow Niche

Ivan Borrello

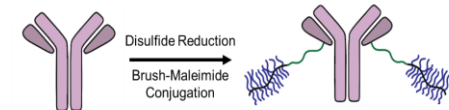
Irene Ghobrial

Imaging and Therapeutics

Jeremiah Johnson

Irene Ghobrial

Generate new tools to **prevent disease progression**



Inclusion Criteria



Adults \geq 30 years old who are:

African American (self-identified)

*Risks are 2-3 times higher
for this group*



African Americans

First-degree relative of a patient with a blood cancer

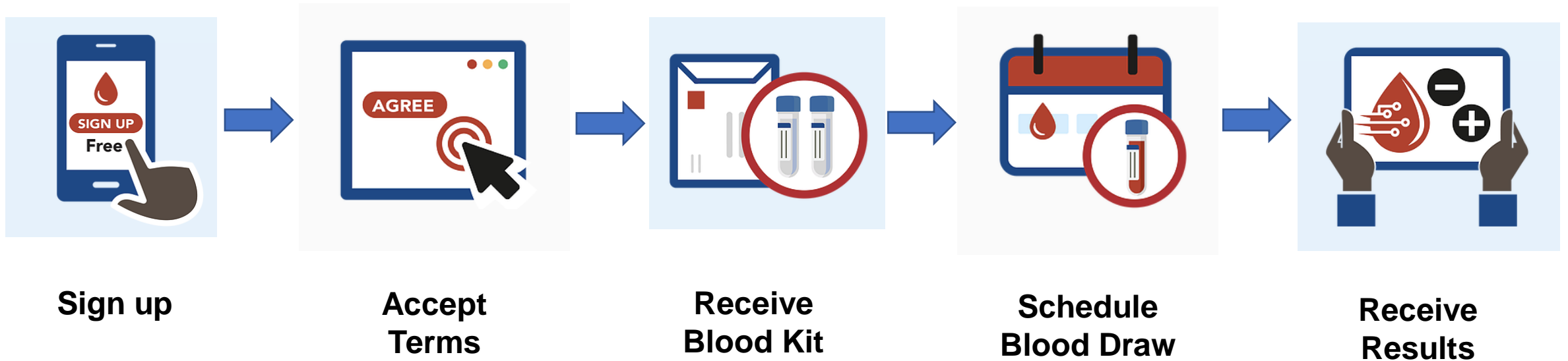
*Risks are higher when a parent, sibling or child has a
blood cancer or myeloma precursor condition*



Close Family Relatives
of a person with myeloma
or its precursor conditions

Expanded inclusion for those who are 18+ with a strong family history (2 or more 1st and/or 2nd degree relatives)

How Does the PROMISE Screening Process Work?



The PROMISE Study: Test Results

After the Mayo Clinic has tested the samples, the results are interpreted by our clinical team.

Positive Cohort

- * The research nurse reaches out by phone to notify the participant and answer questions
- * The participant signs the Positive Consent Form and completes their Baseline Questionnaire
- * The participant finds a hematologist/oncologist for follow-up and samples are banked every few months

Test Negative

Negative Cohort

- * Participants are notified via email of their negative result.
- * Participants are asked to finish the remaining forms on their PROMISE Dashboard, notably the Baseline Questionnaire

Test Positive

THE PROMISE STUDY

On-The-Ground



**Rahway, NJ Agape
Church Health Fair**



Los Angeles, CA



Ayanna Pressley, MA Representative

Minority Recruitment



2022 Indiana Black & Minority Health Fair



177 participants enrolled over 4 days

- 1 Light chain myeloma
- 21 MGUS
- 30 MGIP

COVID-era zoom educational sessions

ASK THE EXPERTS:

Discussing precursor conditions, clinical trials and the future with COVID-19

Dr. Irene Ghobrial is joined by Dr. Gormley and Dr. Bindu Kanapuram at the FDA Oncology Center of Excellence. **Tuesday May 26 from 4:00 - 5:00 PM WEBINAR.**

Submit your questions to precursor@partners.org or leave a comment below!

To join, visit the **Promise Study YouTube channel** or use the link below: https://www.youtube.com/watch?v=eK2lyNVIQ_Y

Do you have questions about multiple myeloma? Do you want more information on how COVID19 affects your precursor condition?

ASK OUR TOP EXPERTS!



Dr. Karen Winkfield
Wake Forest
Baptist Health



Dr. Irene Ghobrial
Dana-Farber Cancer
Institute



Dr. Craig Cole
Michigan State
University

JOIN US FOR A WEBINAR

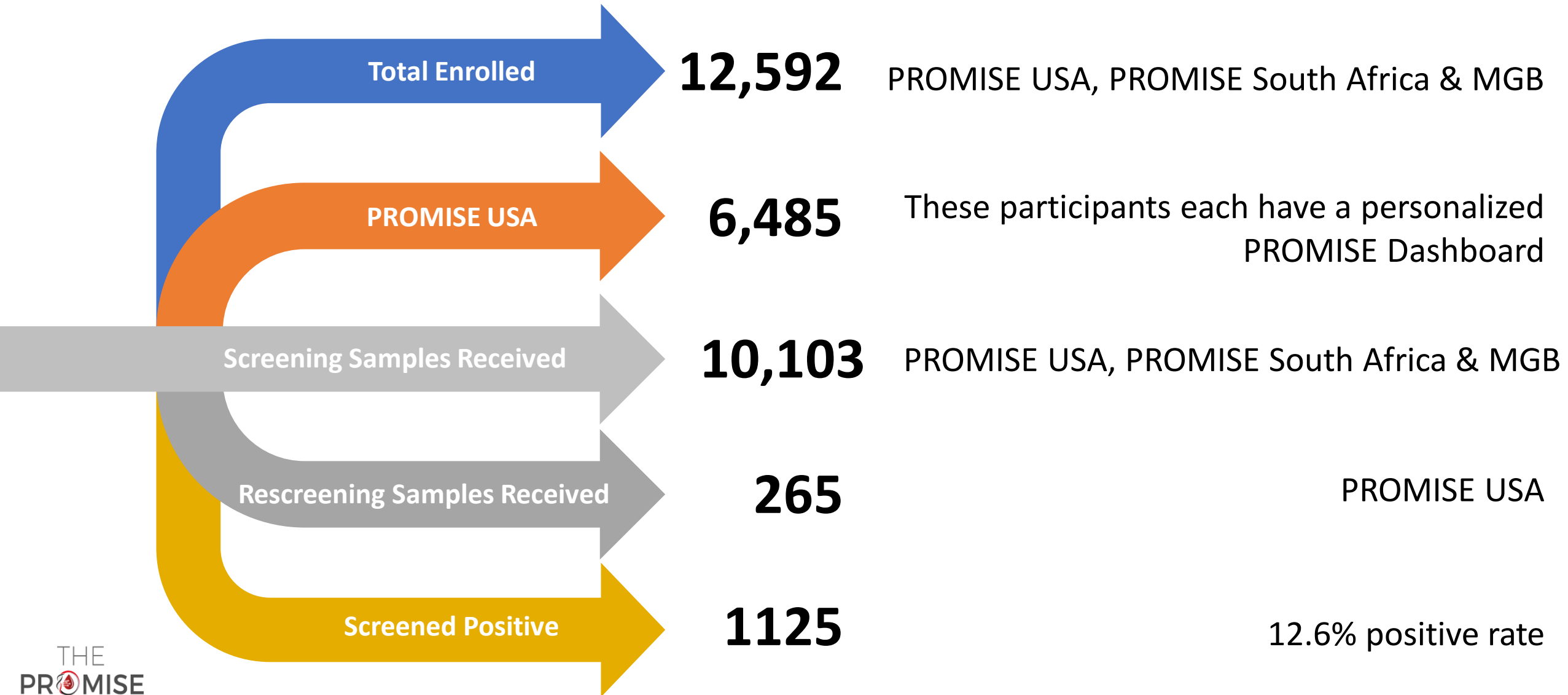
Tuesday, April 21 4:30-5:15 PM EST

Hosted on the Promise Study YouTube Channel

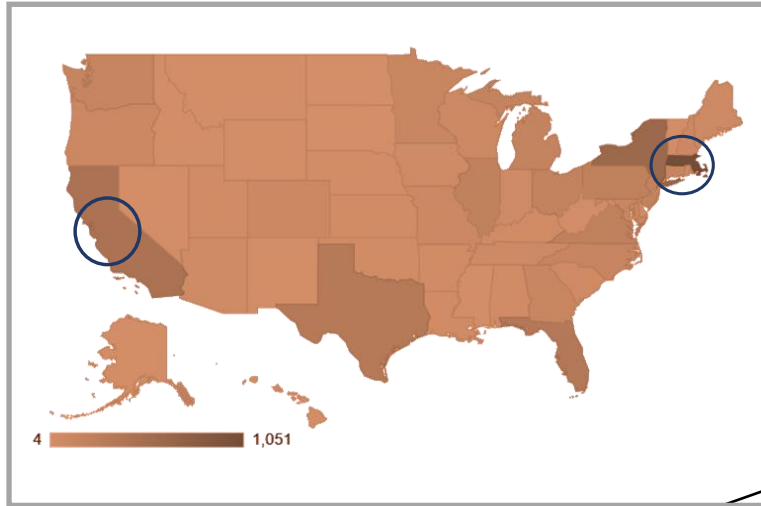
Submit questions to precursor@partners.org

or leave a comment below!

PROMISE by the numbers



PROMISE by the numbers



**12,592
Participants
Enrolled**



**PROMISE
US**

**PROMISE
South Africa**

MGB Biobank

**Family history
8,653**

**African Descent
3,866**

A look at some of PROMISE participants

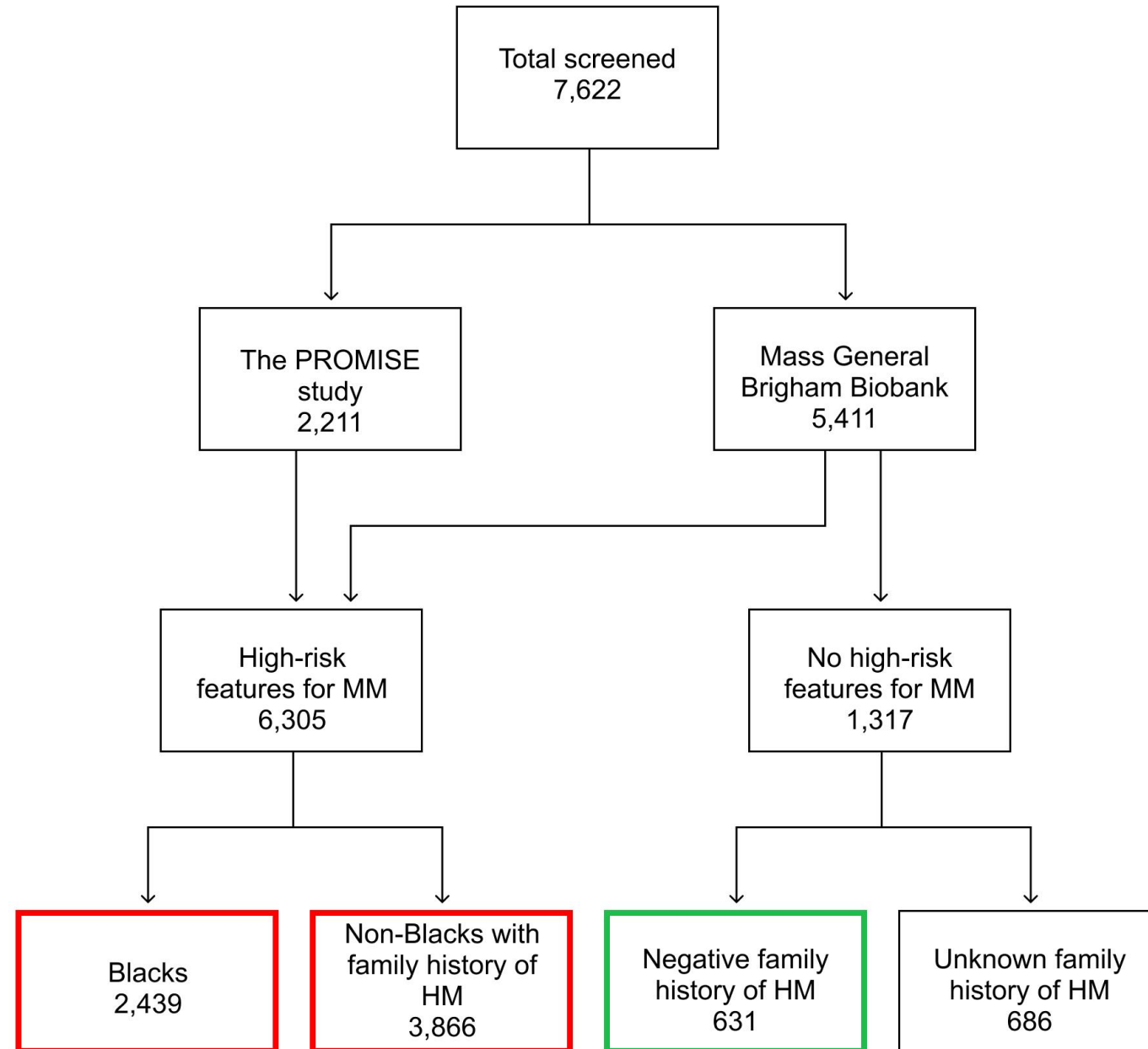
A family of 9 siblings

Numerous sets of twins

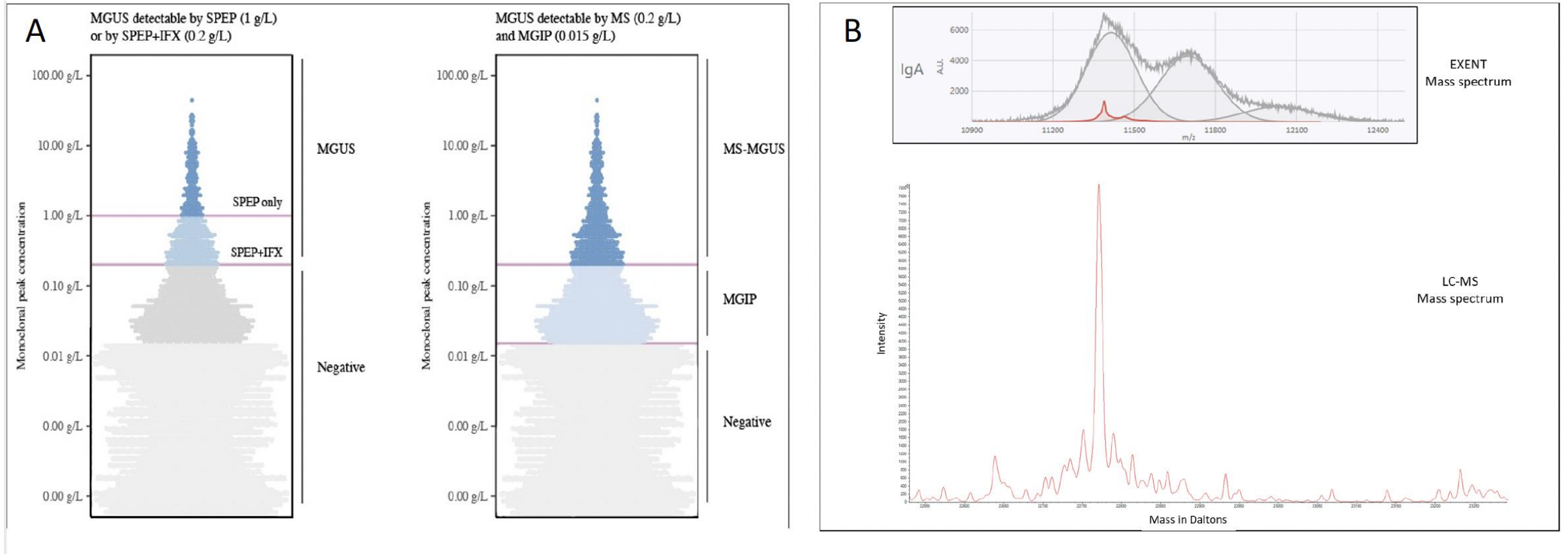
Located on the northern-most
inhabited island off the coast of
Alaska

A team of janitors at a public
high school

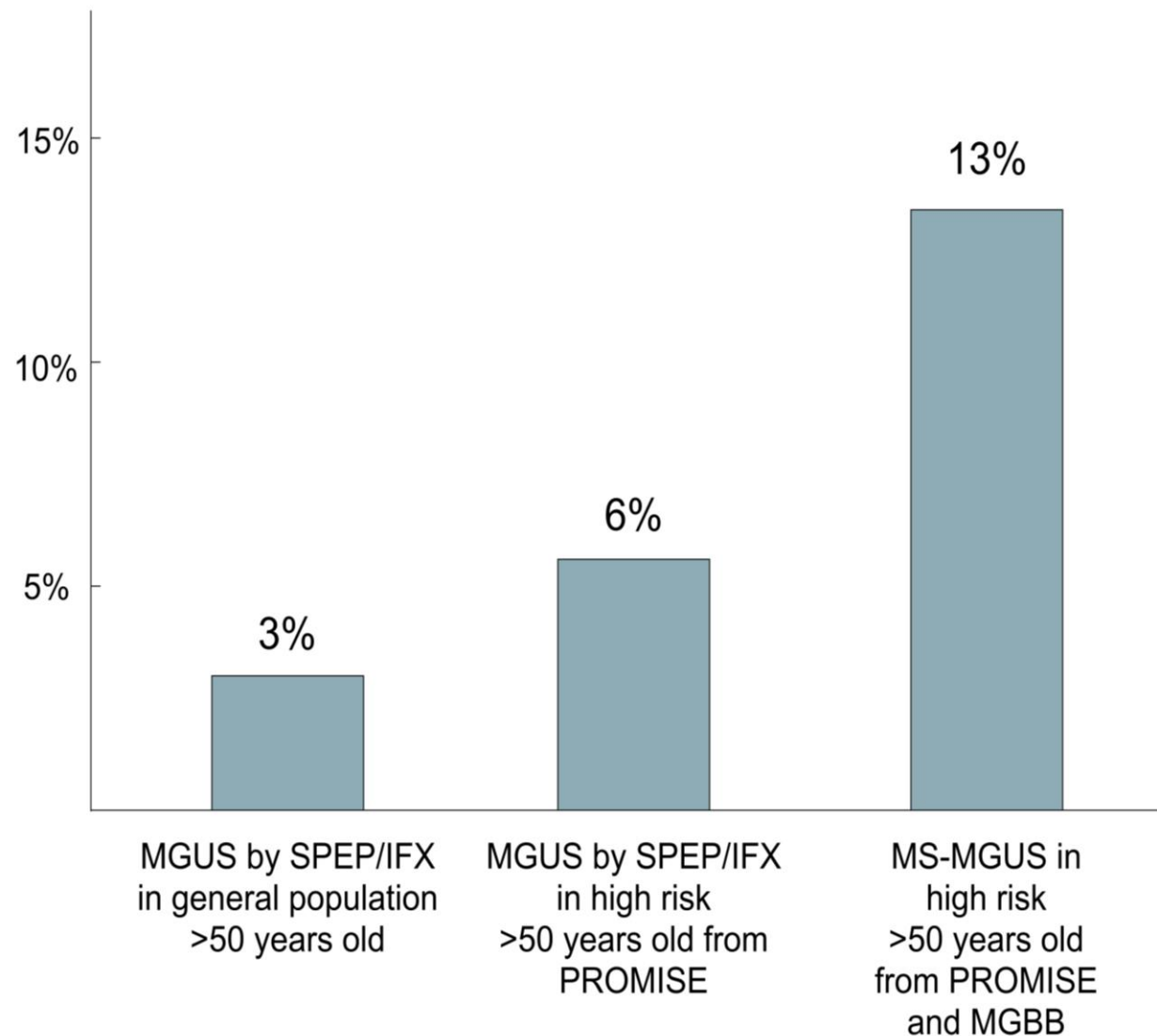
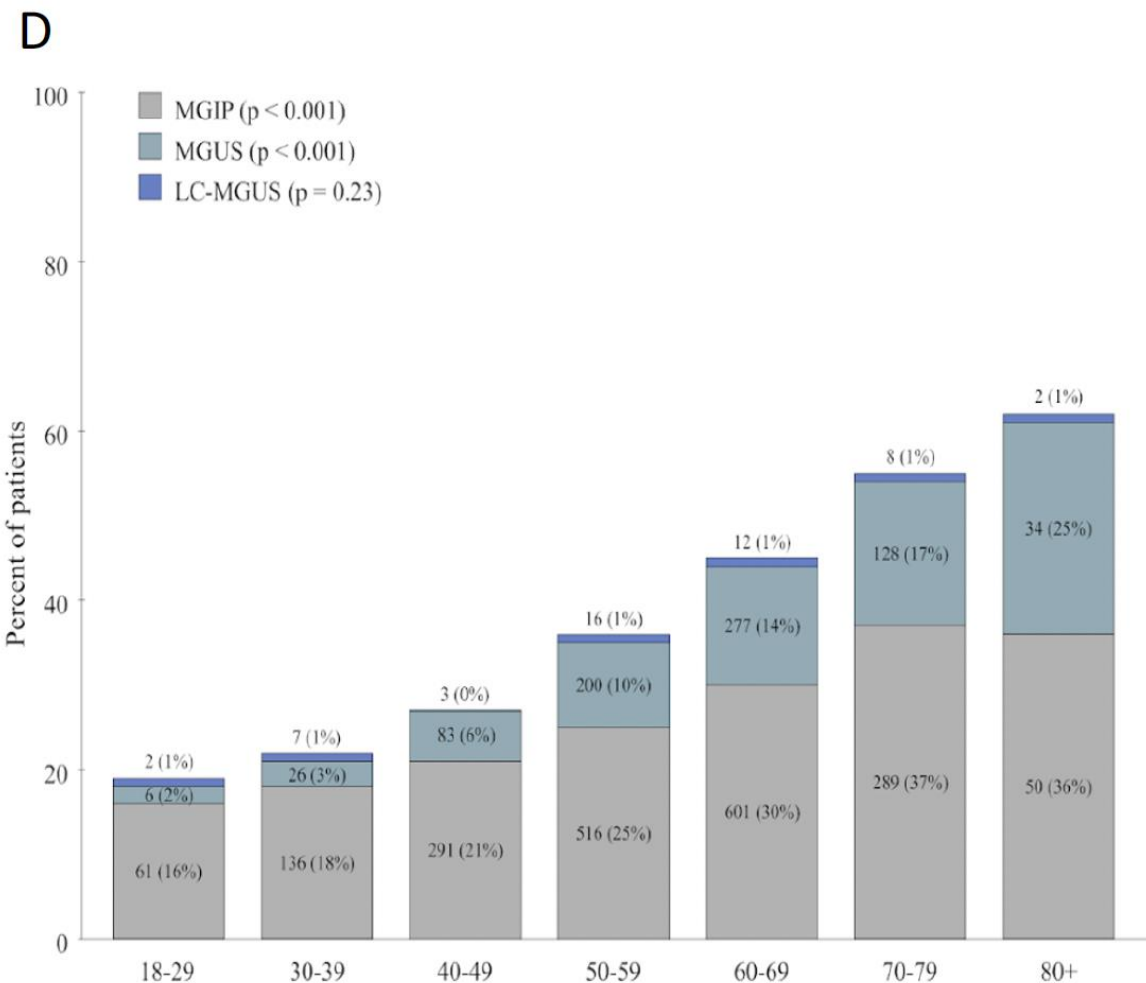
PROMISE early results: Prevalence of MGUS in a high-risk population



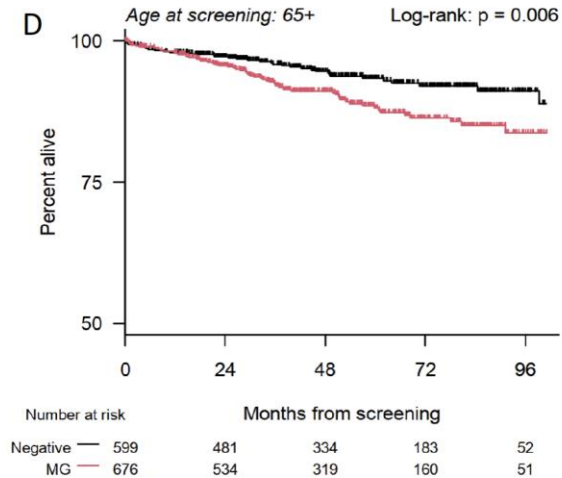
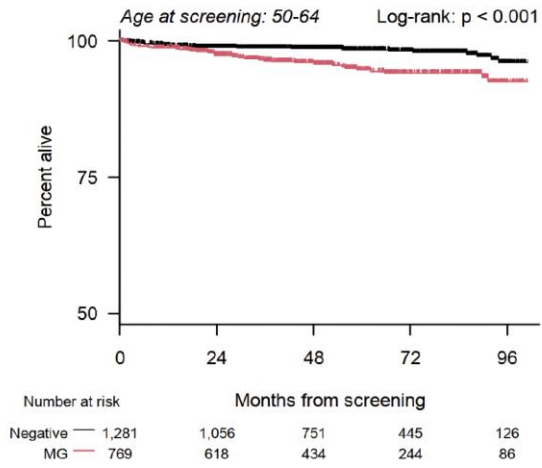
Prevalence of MGUS in a high-risk population



Prevalence of MGUS in a high-risk population



Worse overall survival and association with all cause mortality

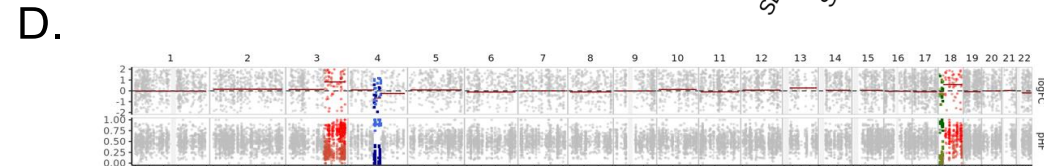
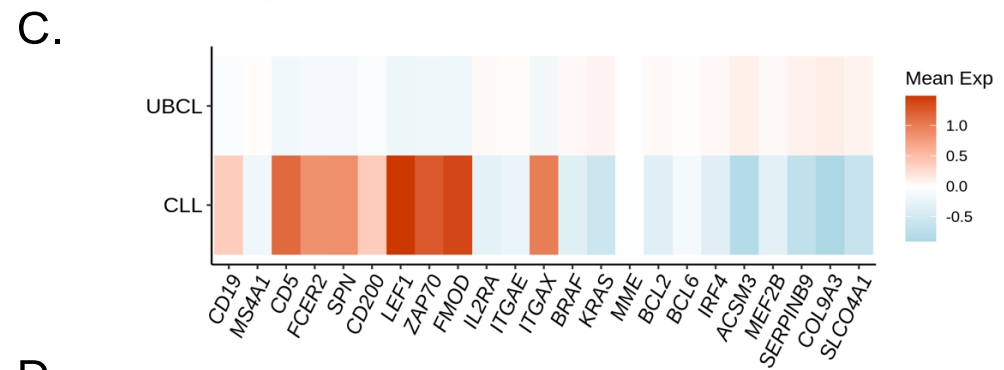
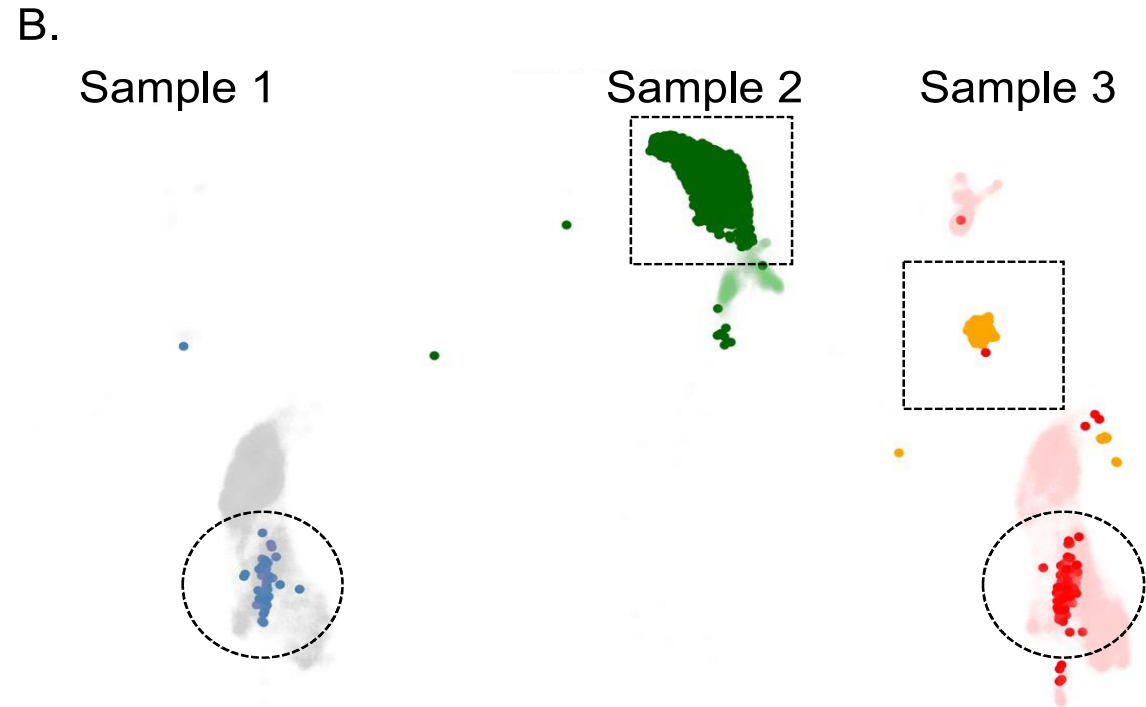
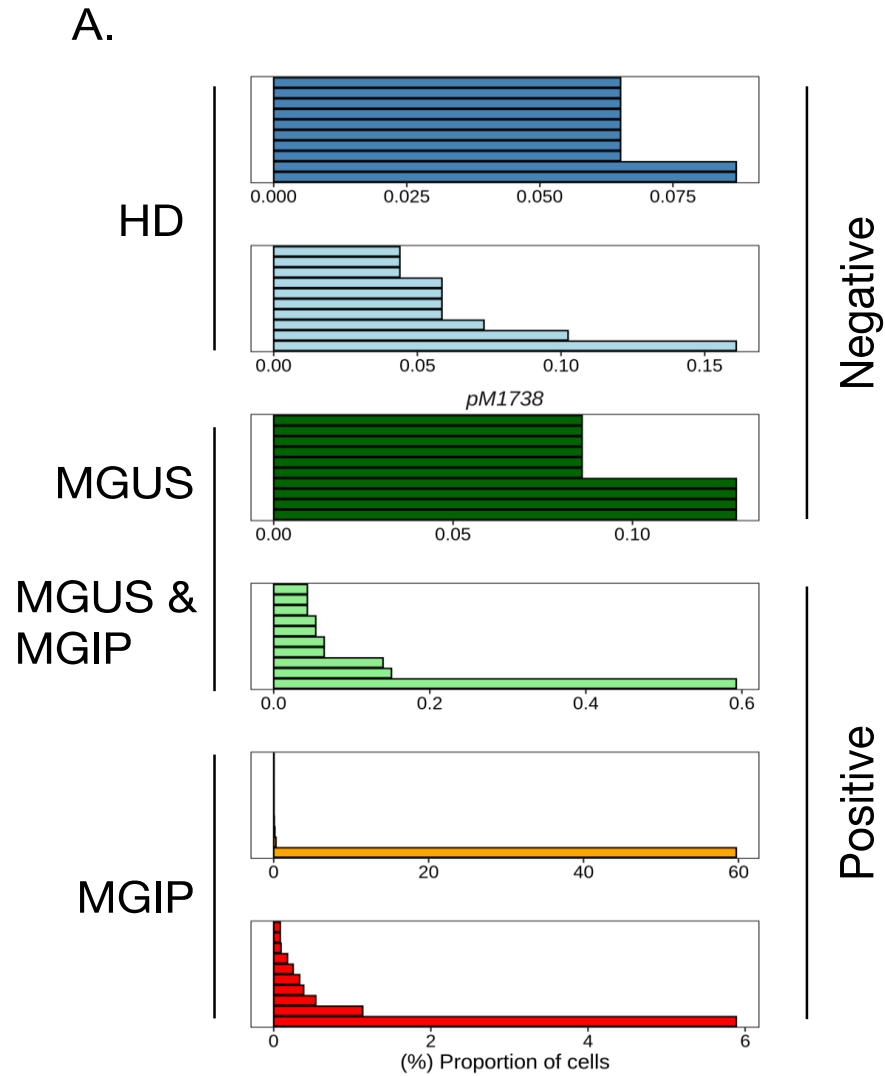


| Term | N (%) | HR (95% CI) | p-value |
|--|-------------|--------------------|---------|
| MS status | | | |
| Negative | 3,455 (64) | Reference | |
| Positive | 1,939 (36) | 1.55 (1.16, 2.08) | 0.003 |
| Age | | | |
| 10-year increase | 5,394 (100) | 1.43 (1.25, 1.65) | < 0.001 |
| Gender | | | |
| Female | 3,410 (63) | Reference | |
| Male | 1,984 (37) | 1.55 (1.17, 2.07) | 0.003 |
| Race/family history risk classification | | | |
| Non-Black, no family history | 631 (12) | Reference | |
| Black | 2,249 (42) | 4.72 (1.73, 12.86) | 0.002 |
| Non-Black, family history | 1,829 (34) | 3.14 (1.13, 8.69) | 0.028 |
| Unknown | 685 (13) | 5.85 (2.07, 16.51) | < 0.001 |
| Charlson comorbidity index | | | |
| 1-unit increase | 5,394 (100) | 1.57 (1.37, 1.81) | < 0.001 |

0 2 4 6 8 10
Hazard ratio

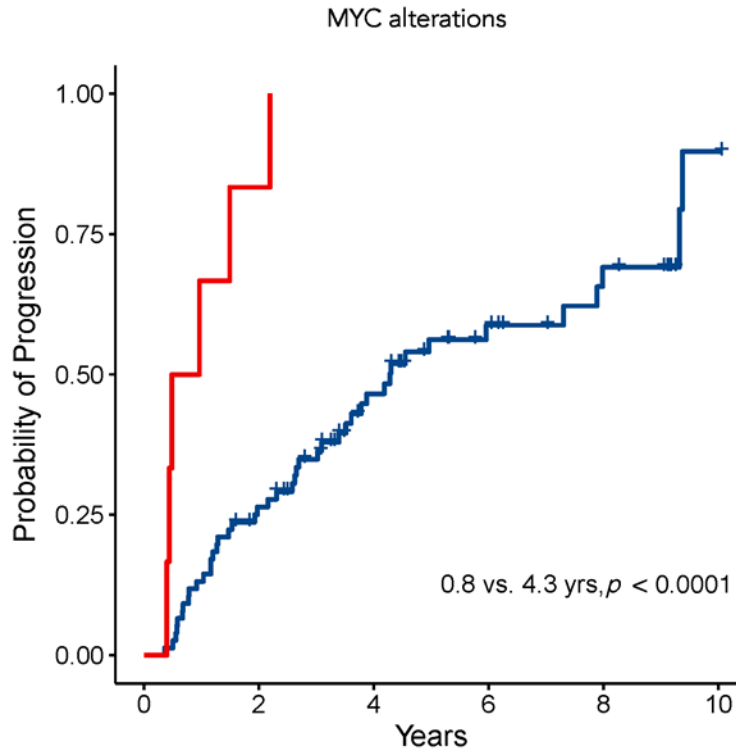
Multi-variable Cox model in all patients, n=5,394

MGIP identifies a lymphoid clone in the peripheral blood



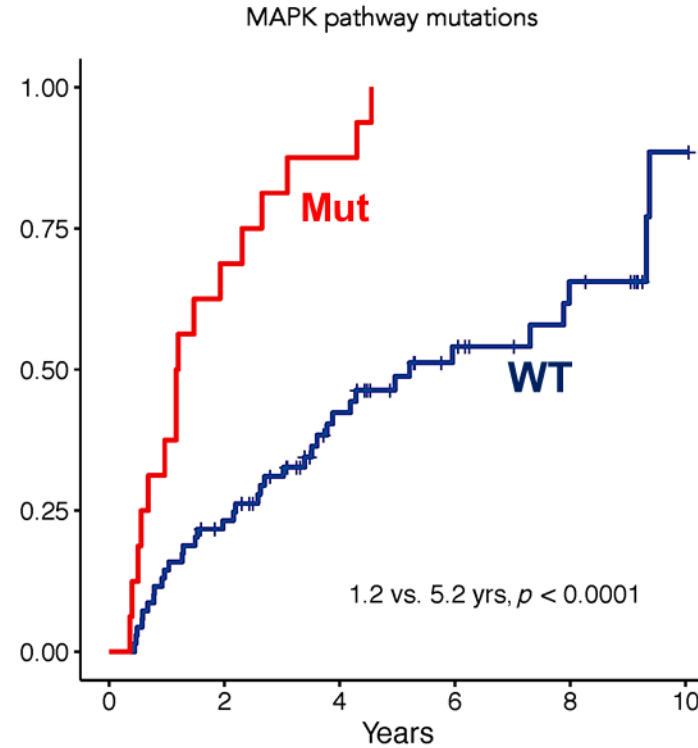
Preliminary unpublished data

Risk stratification of MGUS and SMM to predict progression to MM



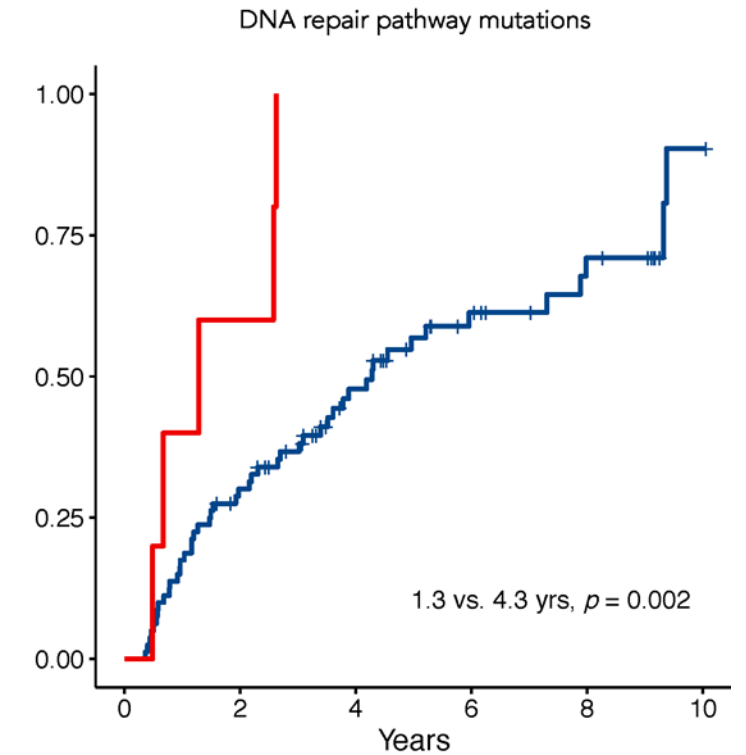
Number at risk

| | | | | | | |
|---|----|----|----|----|---|---|
| 0 | 76 | 54 | 30 | 16 | 9 | 1 |
| 1 | 6 | 1 | 0 | 0 | 0 | 0 |



Number at risk

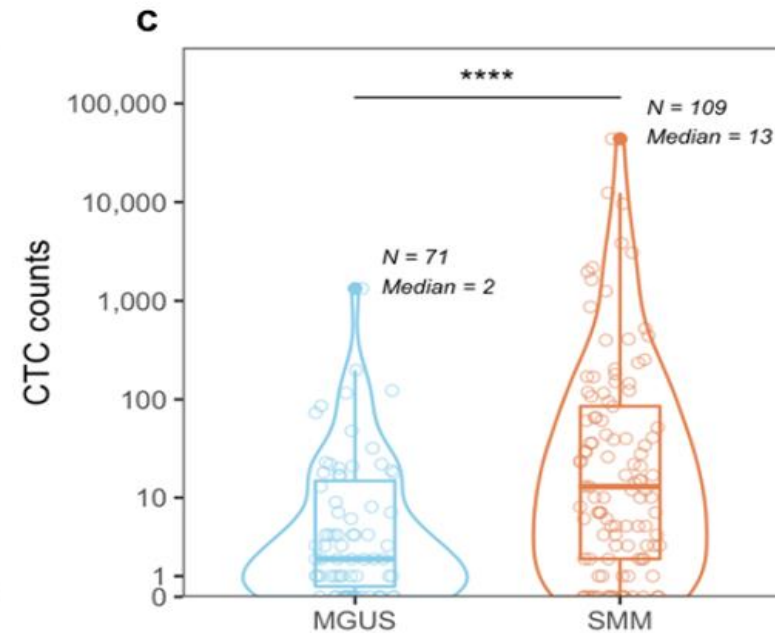
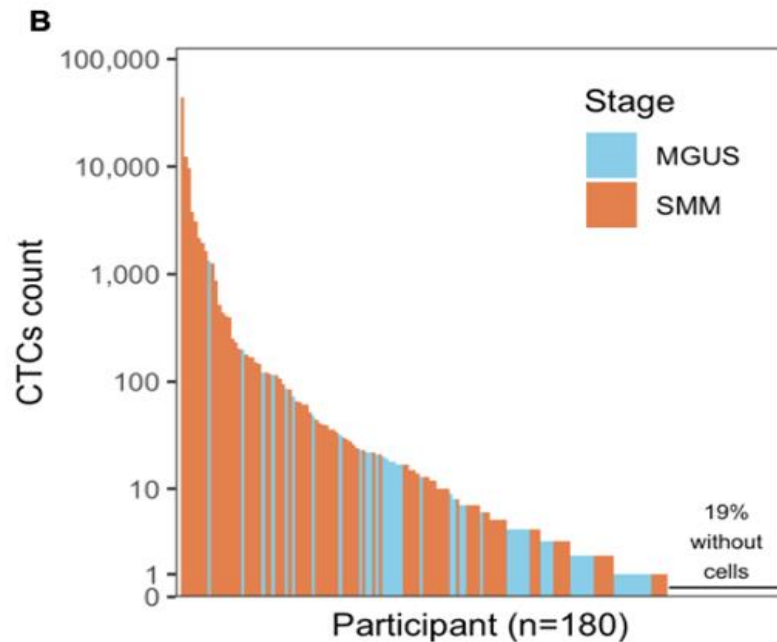
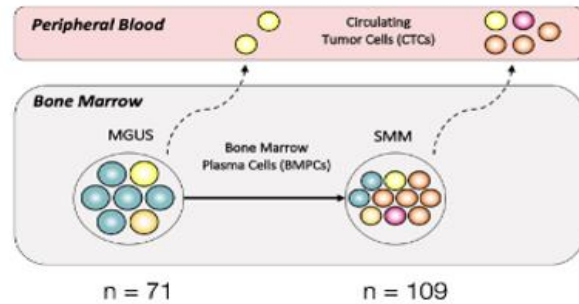
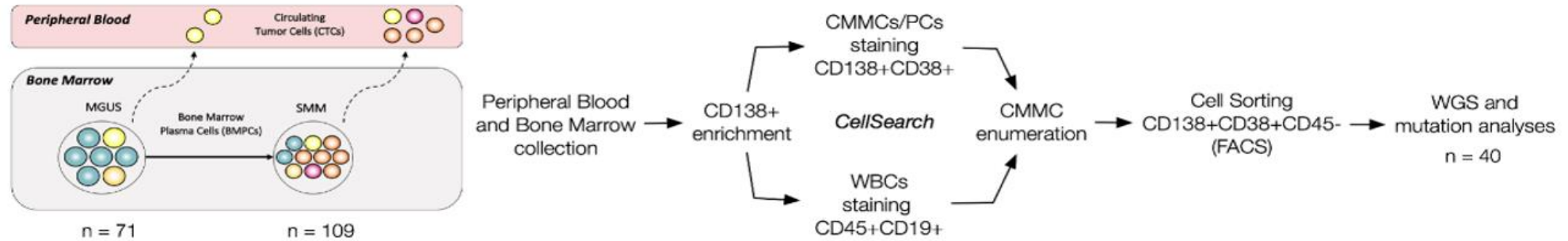
| | | | | | | |
|-----|----|----|----|----|---|---|
| WT | 69 | 51 | 29 | 16 | 9 | 1 |
| Mut | 16 | 5 | 2 | 0 | 0 | 0 |



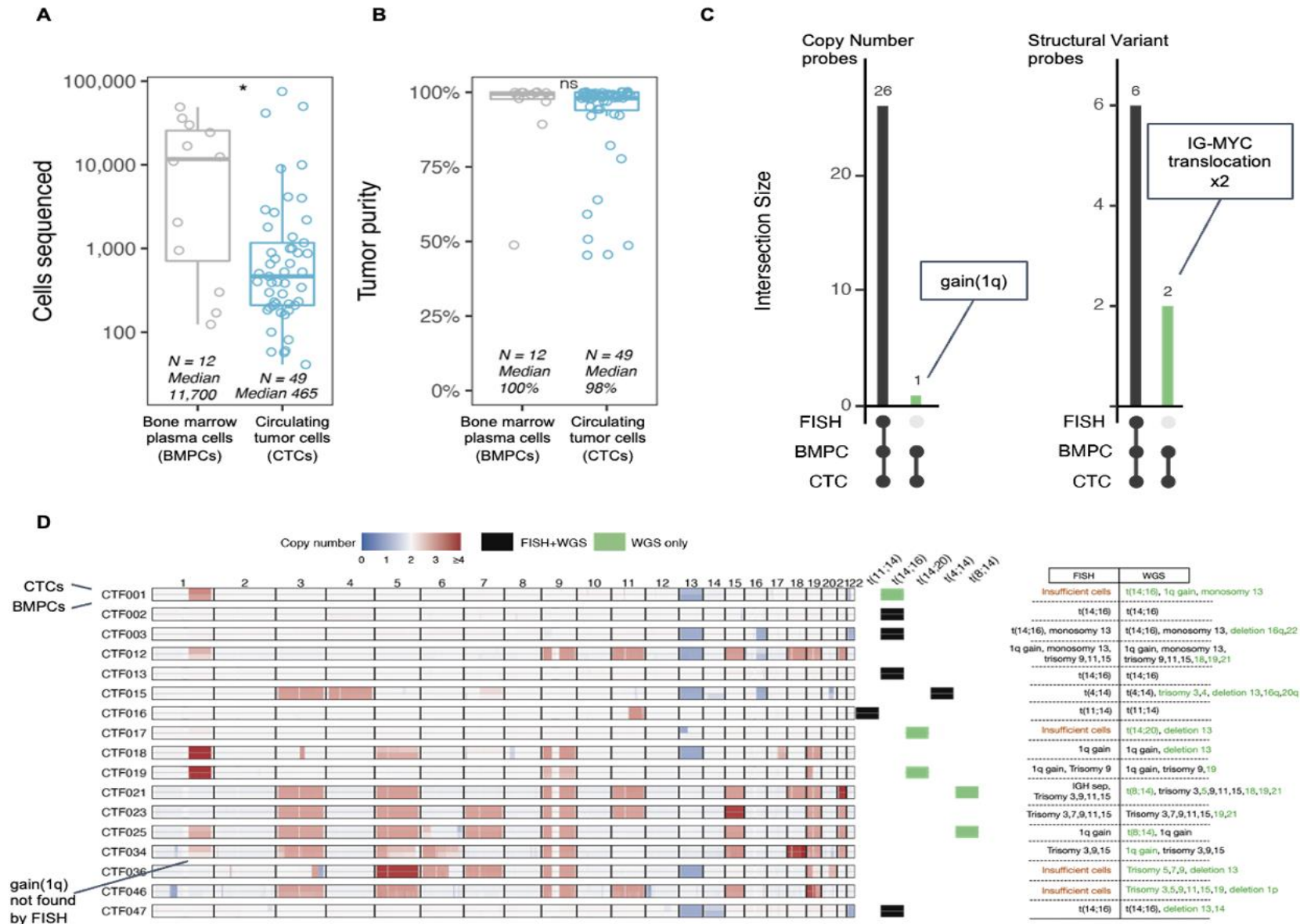
Number at risk

| | | | | | | |
|---|----|----|----|----|---|---|
| 0 | 80 | 54 | 31 | 16 | 9 | 1 |
| 1 | 5 | 2 | 0 | 0 | 0 | 0 |

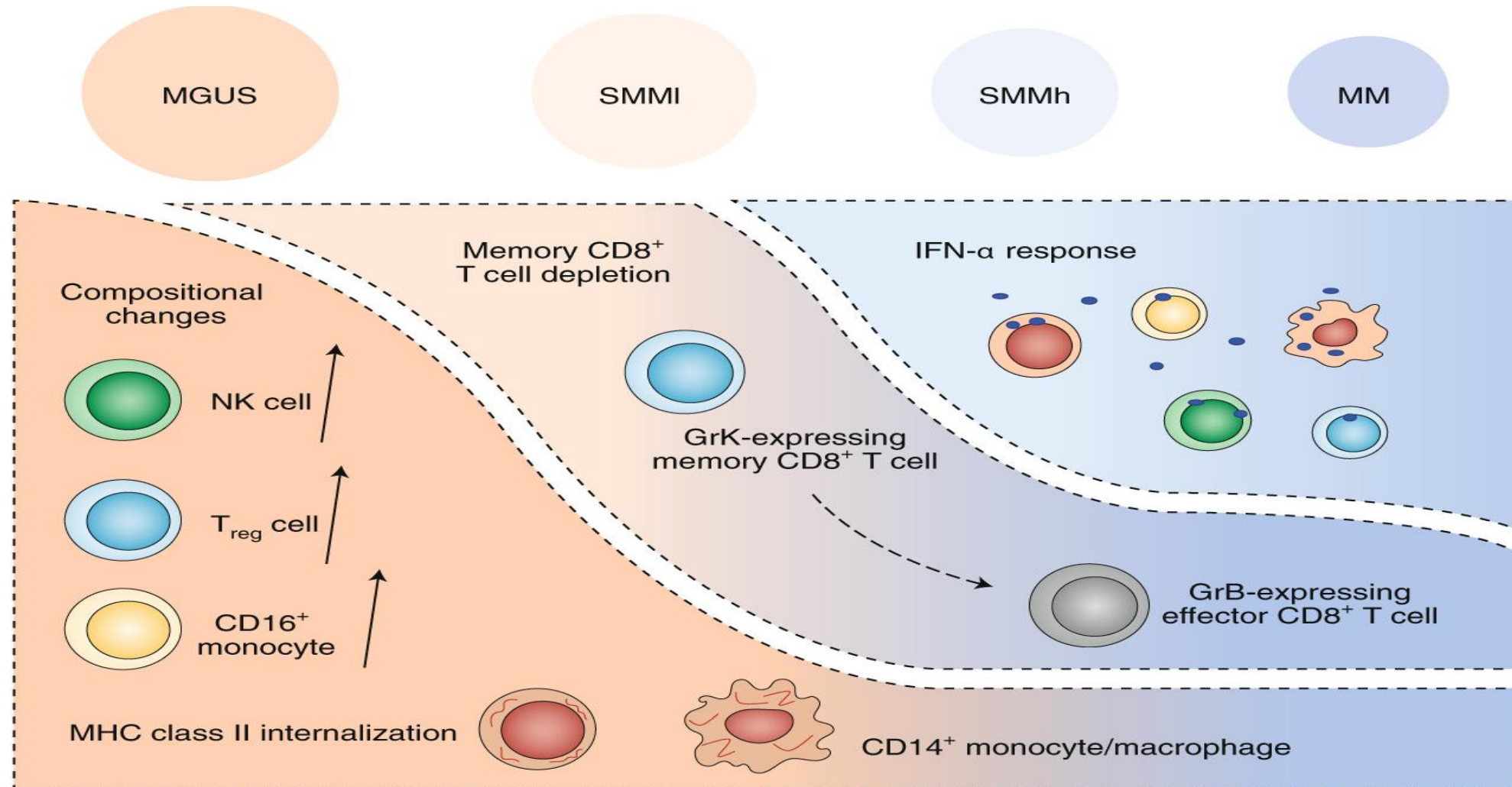
MinimuMM-seq: WGS of CTCs for minimally invasive molecular characterization of clonal evolution



Whole genome sequencing can replace FISH



Early alterations in the immune microenvironment



Bailur et al. JCI Insight 2019

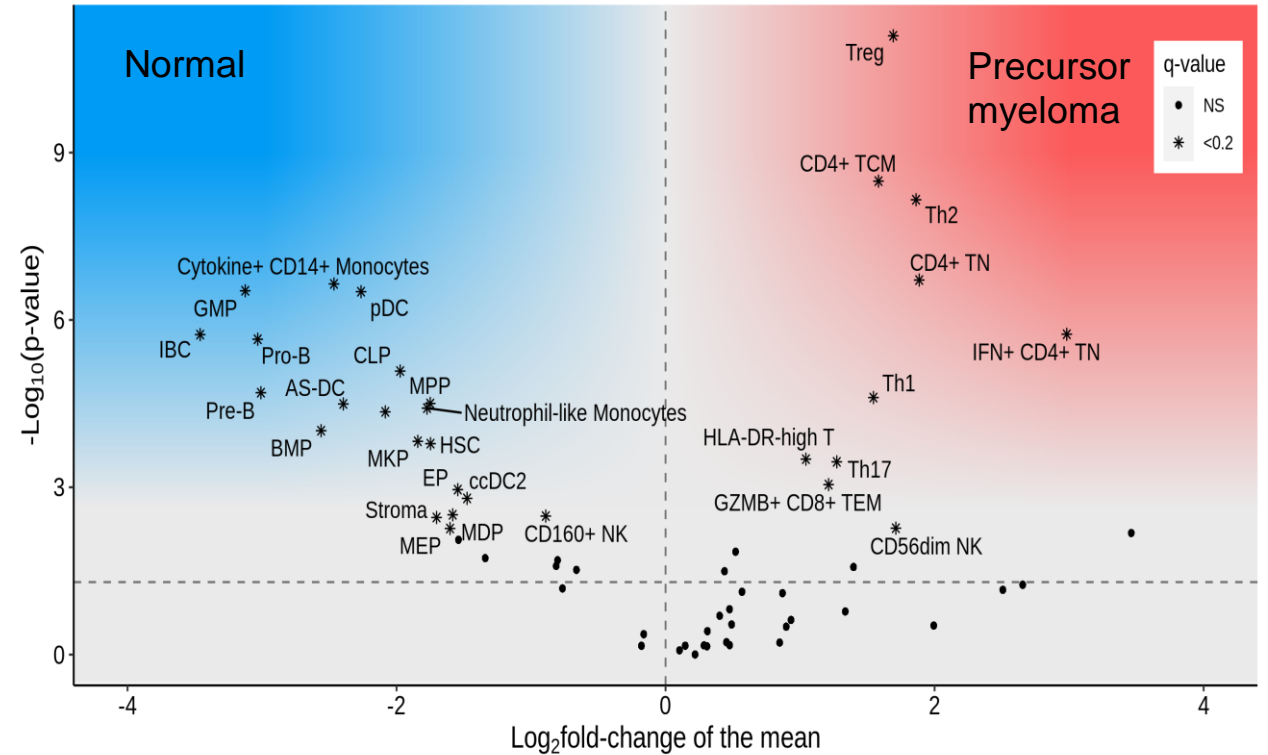
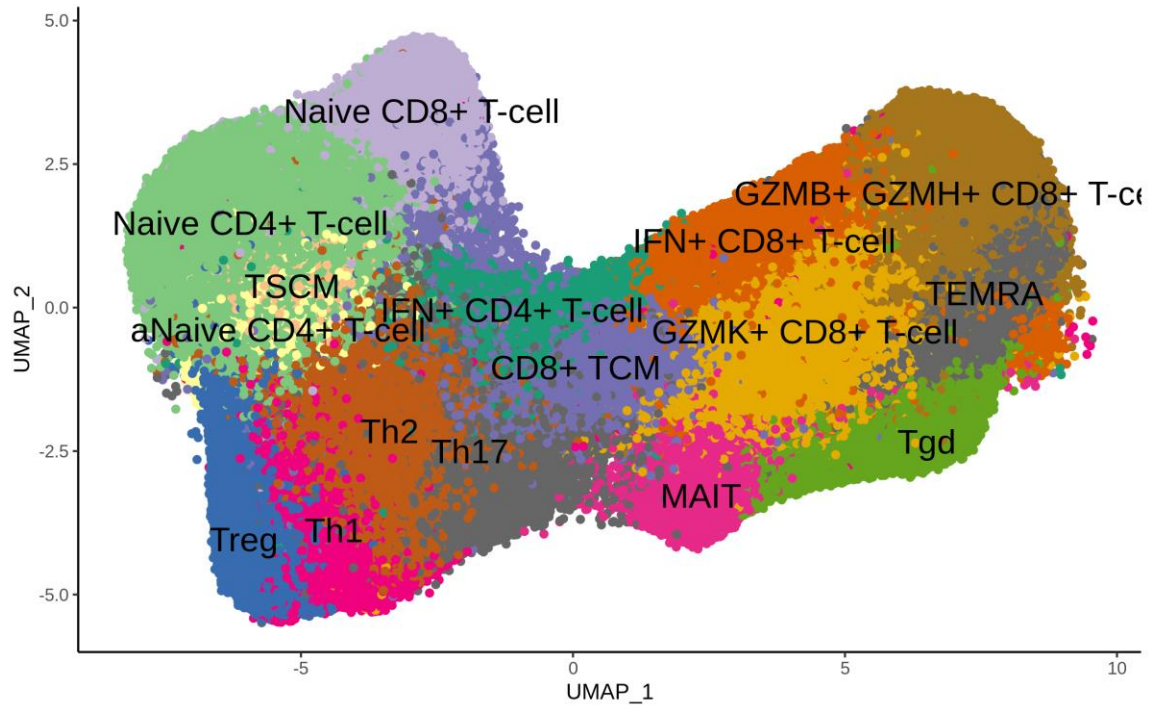
Kourelis et al. Blood Cancer J 2019

Zavidij et al. Nat Cancer 2020

Bashin et al. ASH 2020

Liu et al. Nat Comm 2021

Can we identify healthy from precursor MM by immune cell sequencing



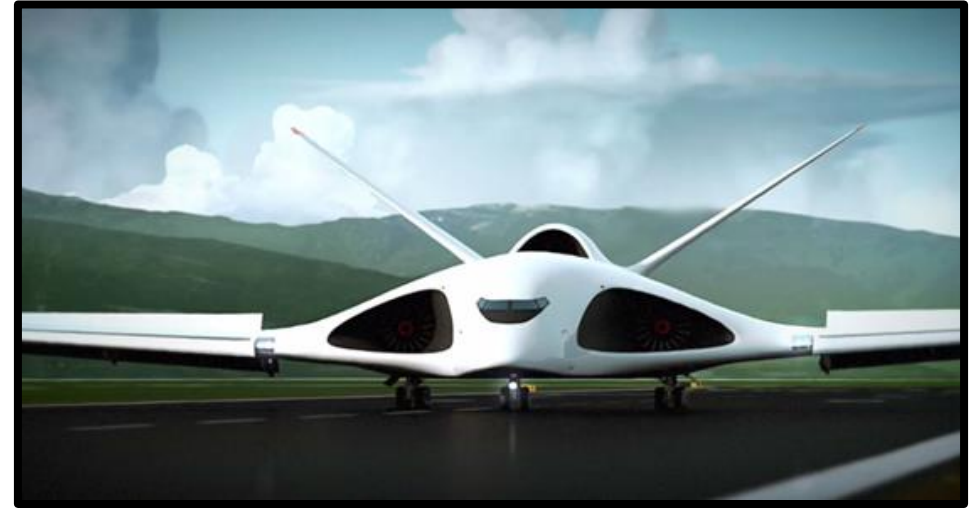
Single-cell RNA-sequencing (n=190), Healthy, MGUS and smoldering myeloma. Bone marrow and peripheral blood

Our First Attempts of therapy in SMM



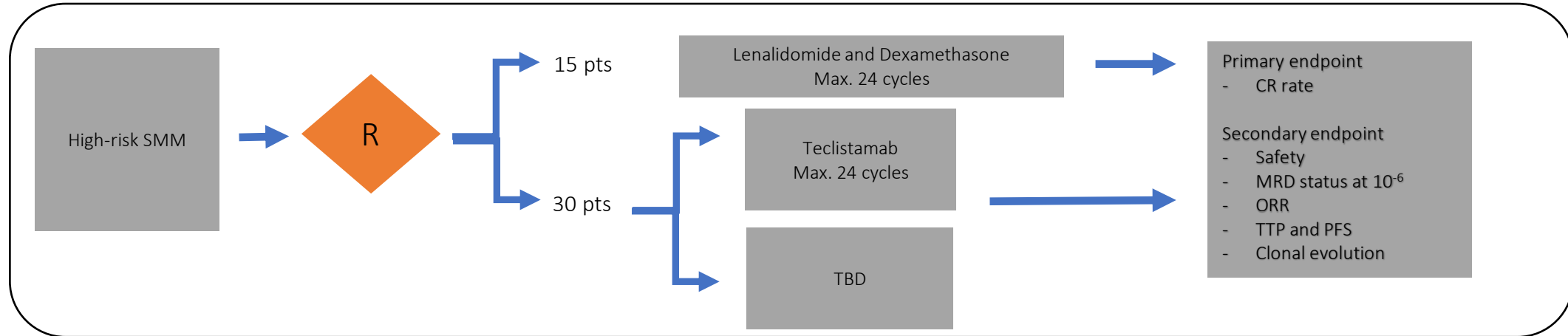
- Lenalidomide was the first proof of principle that early therapeutic intervention works in high risk SMM
- Possible immune regulation
- No overall survival benefit yet
- Cannot truly predict who had benefit and who had clonal selection and tumor resistance

Where we are heading



- Develop precision interception based on genomic/immune profile
- t11:14- venetoclax
- Vaccine therapy for MGUS
- Immunotherapy early to control the clone without the need of traditional myeloma therapy
- Identify markers of response or resistance

Immuno-PRISM (Precision Intervention Smoltering Myeloma): A Randomized Phase II Platform Study of Select Immunotherapies for High-Risk Smoldering Myeloma



**If a participant randomized to the control arm experiences confirmed IMWG disease progression at any time during the treatment period, they may choose to receive the investigational agent for up to 24 cycles.*

Inclusion Criteria:

High risk SMM defined as having 1 of the following 2 criteria:

1. High risk per "20-2-20" Criteria defined as presence of any two of the following:

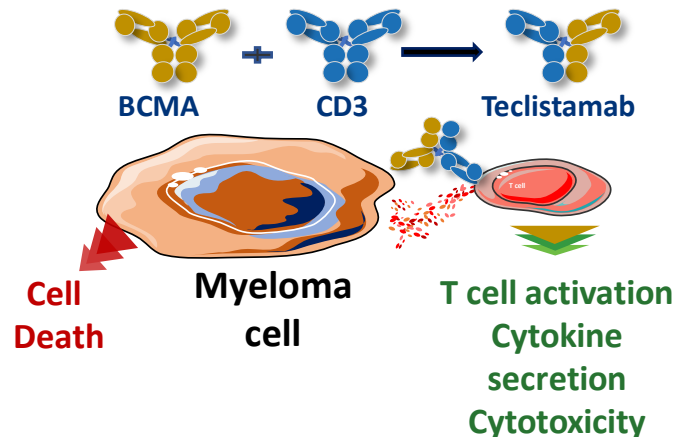
- Serum M spike ≥ 2 gm/dL
- Involved to uninvolved free light chain (FLC) ratio ≥ 20
- Bone marrow PC% $\geq 20\%$

OR **total score** of 9 using the following scoring system:

- FLC Ratio: $>10-25 = 2$, $>25-40 = 3$, $>40 = 5$
- Serum M Protein (g/dL): $>1.5-3 = 3$, $>3 = 4$
- BMPC%: $>15-20 = 2$, $>20-30 = 3$, $>30-40 = 5$, $>40 = 6$
- FISH abnormality (t(4,14), t(14,16), 1q gain, or del13q = 2)

2. Presence of $\geq 10\%$ BMPC and at least one of the following:

- **Evolving pattern**
- **Abnormal PC immunophenotype** ($\geq 95\%$ of BMPCs are clonal) and reduction of ≥ 1 uninvolved immunoglobulin isotype. (Only IgG; IgA and IgM will be considered)
- **High risk cytogenetics** defined as presence of t(4;14), t(14;16), t(14;20), 17p deletion, TP53 mutation, 1q21 gain



Teclistamab Dosing:

Cycle 1

- Step-up dose: days 1 and 3
- Treatment Dose: days 8, 15, 22

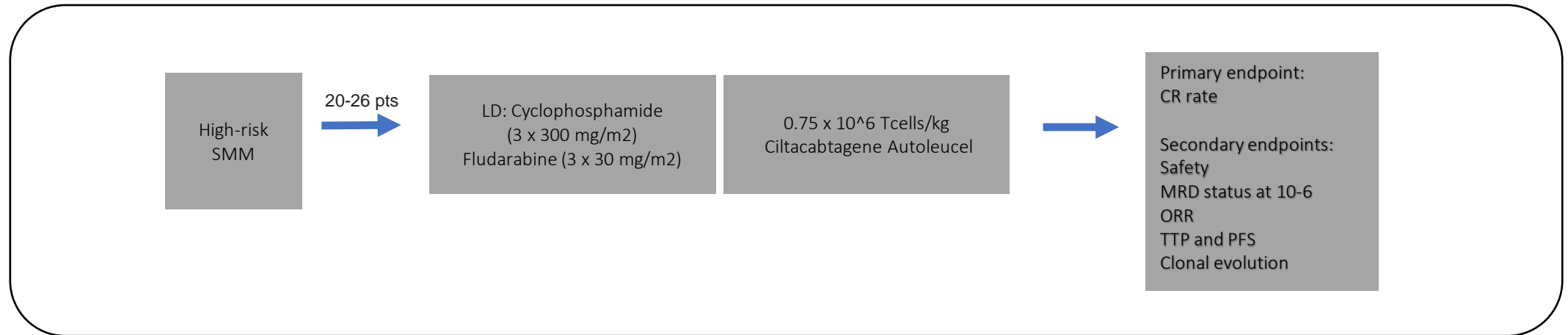
Cycle 2:

- Teclistamab (subcutaneous): Days 1, 8, 15 and 22

Cycle 3-24

- Teclistamab (subcutaneous): Days 1 and 15

CAR-PRISM (Precision Intervention Smoldering Myeloma): Ciltacabtagene Autoleucel in High-Risk Smoldering Myeloma



Inclusion Criteria:

High risk SMM defined as having 1 of the following 2 criteria:

1. High risk per "20-2-20" Criteria defined as presence of any two of the following:

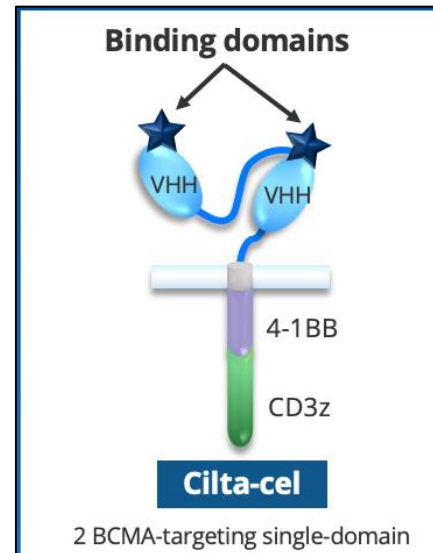
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Cilta-Cel Dosing:

- First 3 patients at 0.5×10^6 /kg cells
- Subsequent patients at 0.75×10^6 /kg cells
- Staggered enrollment for first 3 patients
- Safety criteria



Dana-Farber
Cancer Institute

Lab members:

Mike Agius
Michelle Aranha
Jean Baptiste Alberg
Luca Bertamini
Cody Boehner
Amanda Cao
CJ Curry
Julia Colchie
Ankit Dutta
Habib El-Khoury
Daniel Mallor
Camille Gladieux
Laura Hevenor
Anna Justis
Yoshinobu Konishi
David Lee

Clinical members:

Omar Nadeem- CPOP director
Brian Sheehan
Meredith Bertoni
Annie Cowan
Maya Davis
Erika Howowitz
Elisabeth Kitzenberg
Elizabeth Murphy
Vidhi Patel
Jacqueline Perry
Hira Shretha
Kelsey Tague
Rebekah Medina

Elizabeth Lightbody
Oliver Lomas
Mahshid Rahmat
Romanos Pistofidis
Andrea Poletti
Cinnie Seokojo
Michael Timonian
Katherine Tiwle
Shirley Wang



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Research Foundation



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LYMPHOMA
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NATIONAL
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