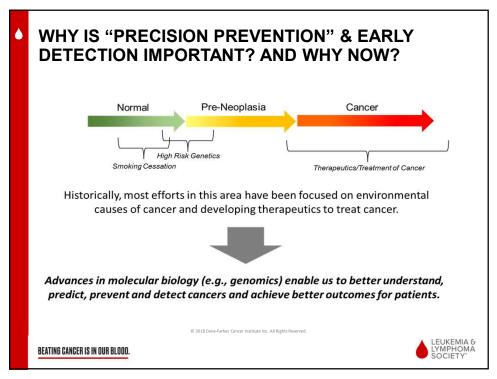
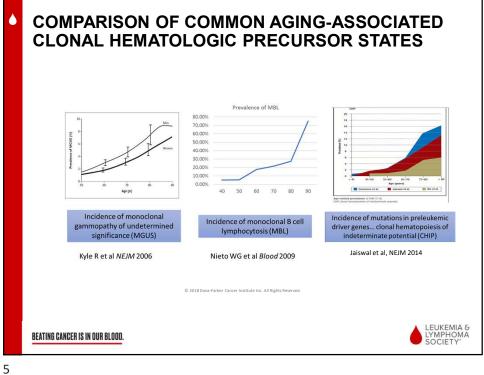
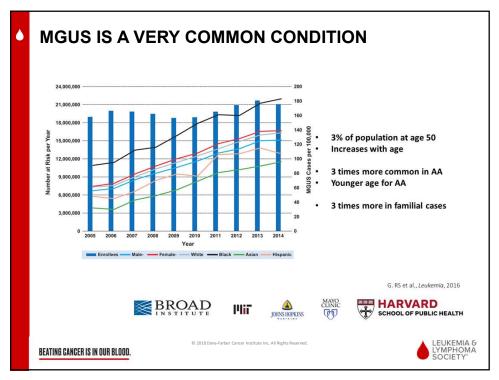
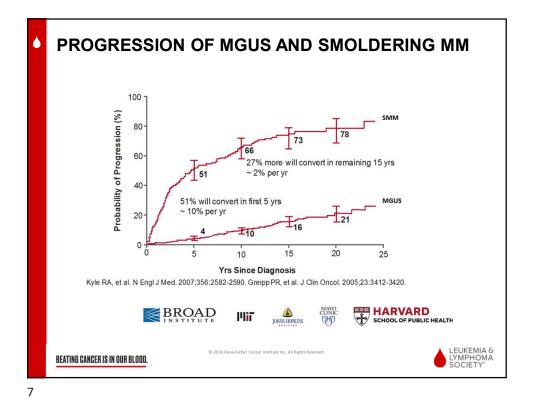


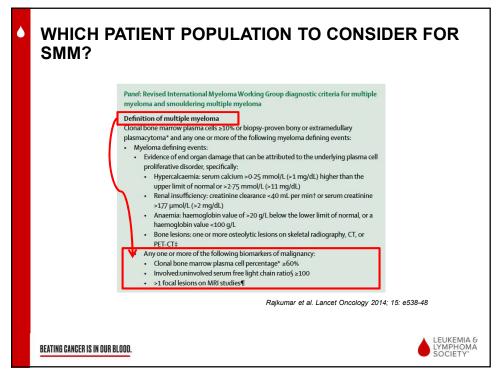
			MILD	MODERATE	SEVERE	CI	s
olon	NORMAL 5-20	INITIATED	adenoma	PRE-C	ANCER 5-15 years		CANCE
lead and neck	tobacco	r	dysplastic			years	
cervix	CIN 1		9–13 years	akid	CIN 3/CI	s	10-20 years
ung (smokers)				20-40 pack-year	5		
preast	atypical hyperplasia					DCIS	6-10 years
prostate		20 years	PIN	≥10 years	latent cancer		3-15 years

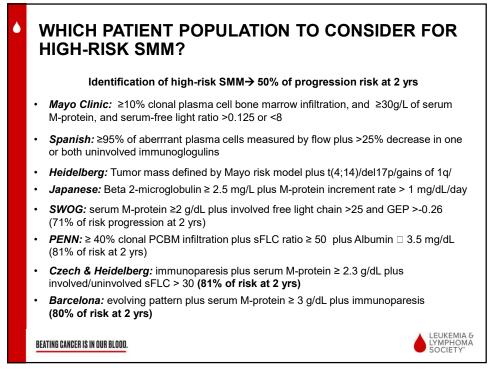


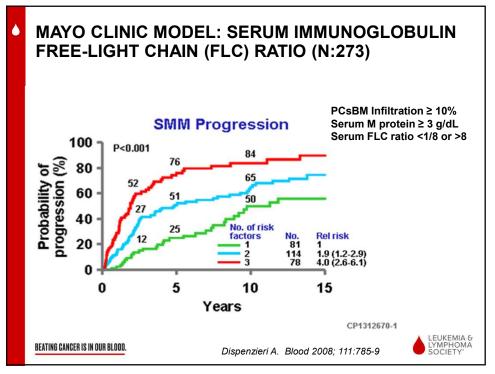


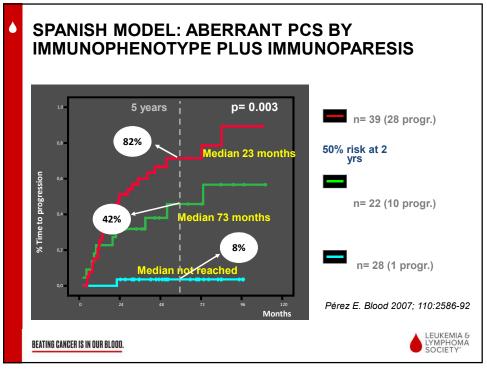












DEL(17P), T(4;14), AND +1Q21 PREDICT PROGRESSION FROM SMOLDERING TO SYMPTOMATIC MM (N=248)

del(17p13), t(4;14), +1q21 showed significant impact on TTP

	TTP	P
All pts	4.9 years	
+1q21 versus no gain of 1q21	3.7 years 5.3 years	0.013
del(17p13) versus no del(17p13)	2.7 versus 4.9 years	0.019
t(4;14) versus no t(4;14)	2.9 versus 5.2 years	0.021
HD versus NHD	3.9 versus 5.7 years	0.036

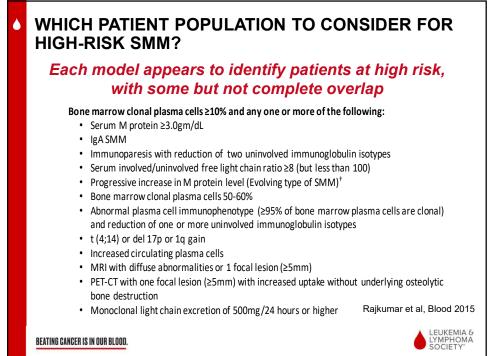
Multivariate analysis: t(4;14), +1q21, HD, reduction of uninvolved immunoglobulins and risk score defined by Kyle et al. as independent factors for adverse outcome

Conclusion: specific chromosomal aberrations drive transition from asymptomatic to symptomatic disease

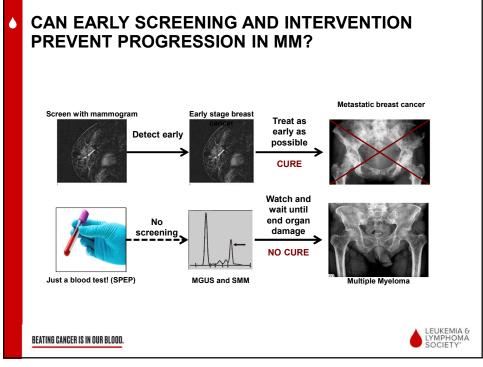
Neben et al. JCO 2013

LEUKEMIA & LYMPHOMA SOCIETY*

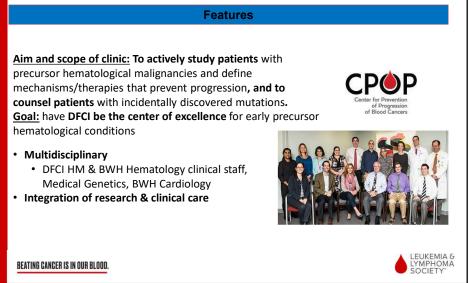
BEATING CANCER IS IN OUR BLOOD.

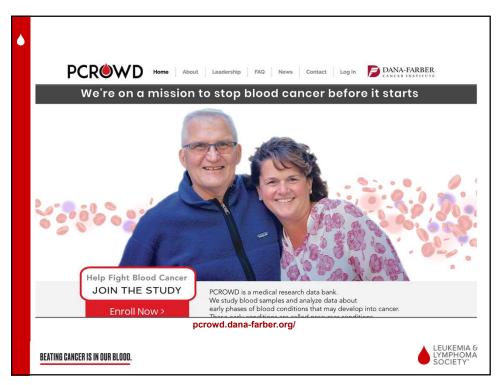


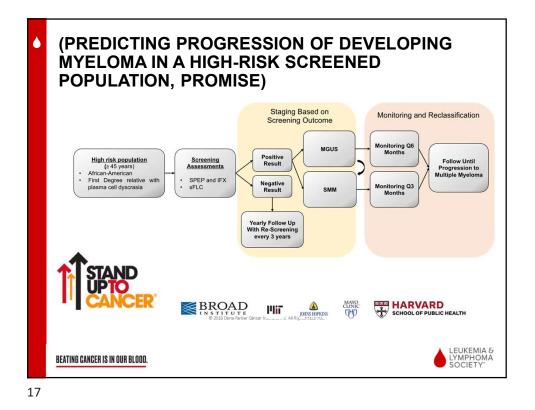




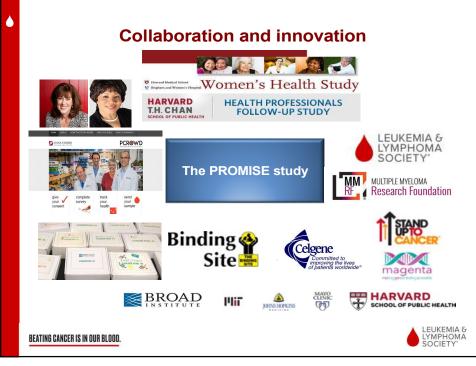
NEW CPOP CLINIC (CENTER FOR PREVENTION OF PROGRESSION OF BLOOD CANCERS)





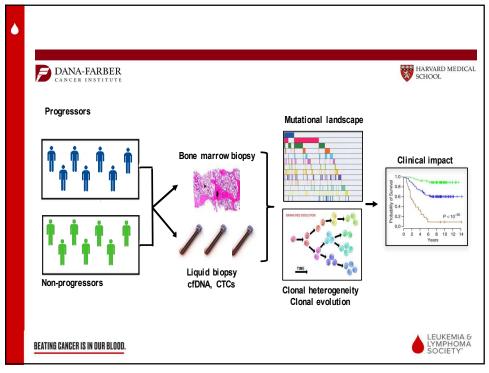


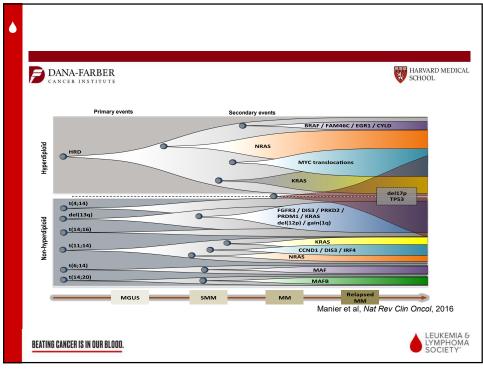
PROJECT SUMMARY ۵ Aim 1. PROMISE screen Aim 2. Genomic Characteristics **Develop novel biomarkers** N = 50,000 Gaddy Getz (Principal) Benjamin Ebert 000 Irene Ghobrial David Liu Π Negative Cohort N = 47.0 Viktor Adalsteinsson Jihye Park 10 Aim 3. Race/Obesity Tim Rebbeck David Liu 0.51 Positive Cohort N = 3,000 Joseph Mikhael Lorelei Mucci Catherine Marinac Т Ί Develop new therapies to Aim 4. Microenvironment Irene Ghobrial (Leader) Ivan Borrello (Co-Leader) Joseph Mikhael (Principal) Lorelei Mucci (Principal) Tim Rebbeck (Principal) prevent/delay progression Ivan Borrello Irene Ghobrial Jihye Park 0.8 Tim Rebuces, Support by: Advocates: Jennifer Ahlstrom and Cheryl Boyce • Broad Institute: DDP • MMRF/MMRC, LLS, Celgene, SU2C advertisement team 0.6 Aim 5. Imaging / Therapeutic 0.4 -Jeremiah Johnson (Principal) 0.2 P < 10-50 Irene Ghobrial Alexandre Detappe 0.0 4 6 8 10 12 1 © 2018 Dana-Farber Cancer Institute Inc. All Rights R HARVARD SCHOOL OF PUBLIC JOHNS HOPKINS MAYO BROAD 14117 GP SCHOOL OF PUBLIC HEALTH LEUKEMIA & LYMPHOMA SOCIETY* BEATING CANCER IS IN OUR BLOOD.

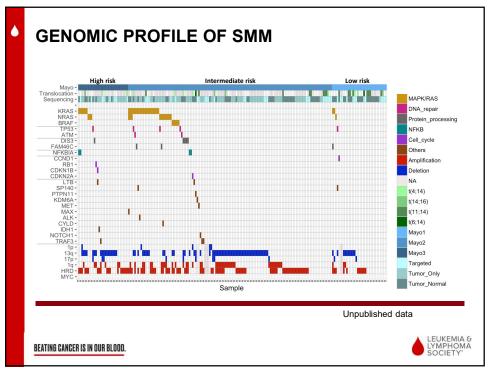


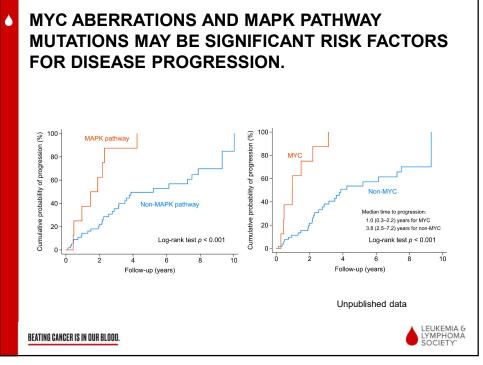


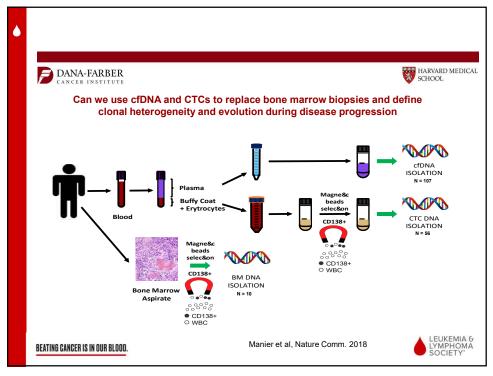


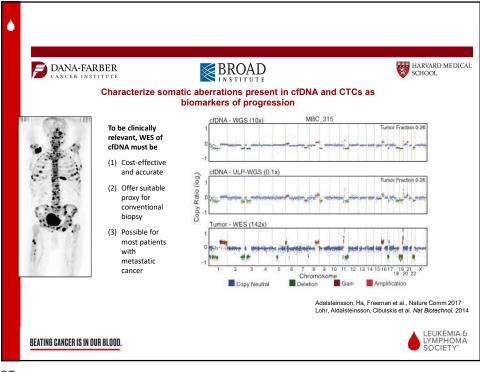


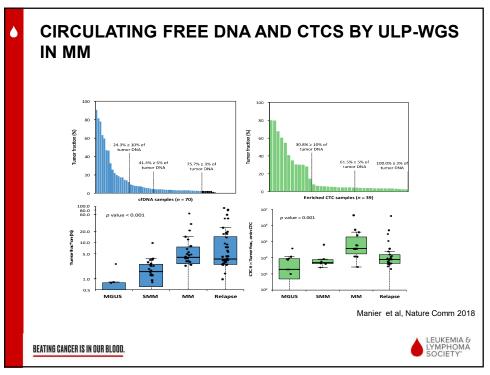


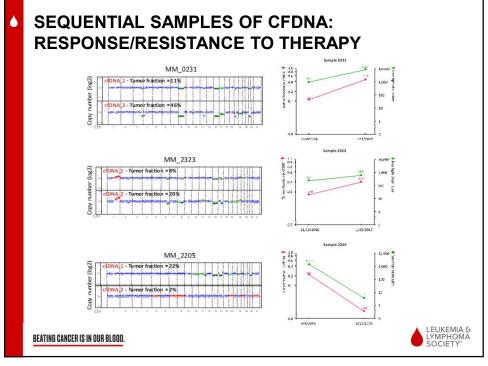


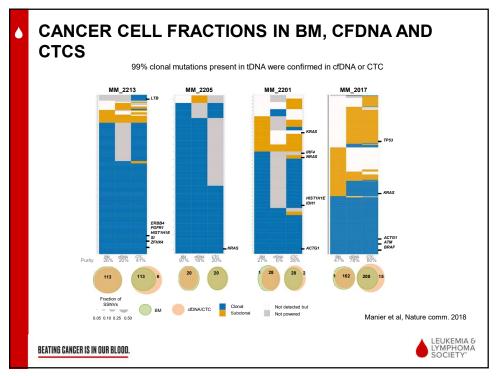


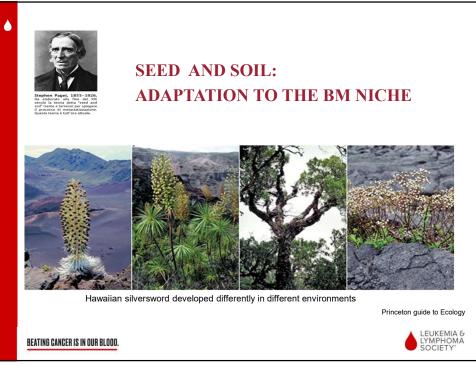


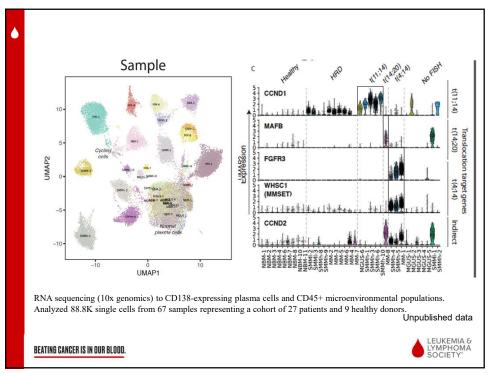


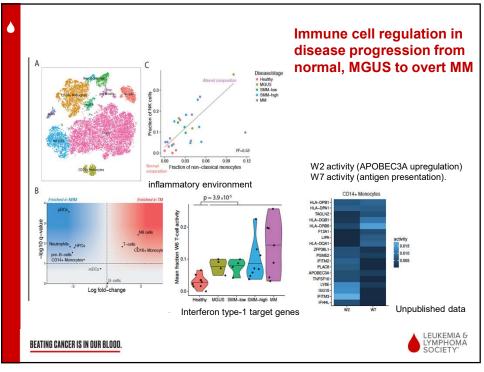




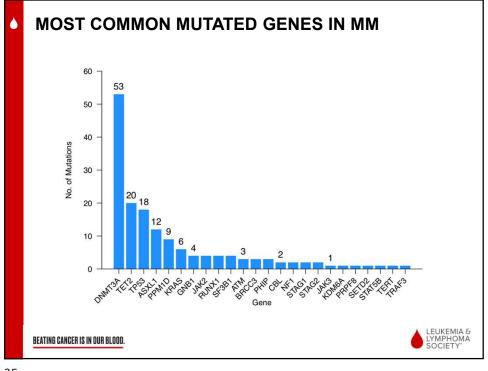


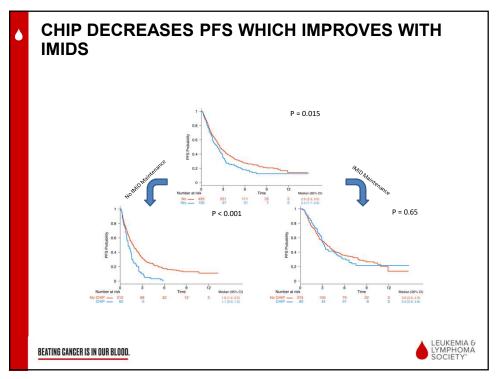


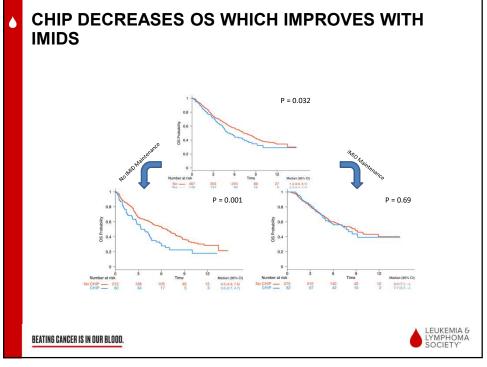


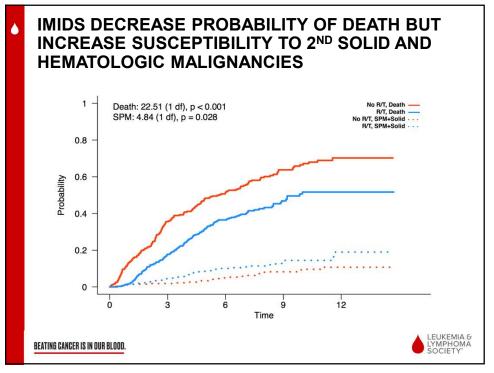


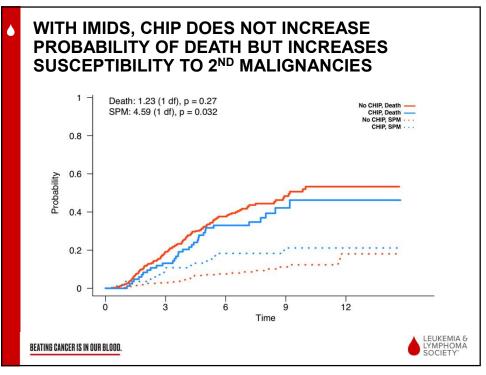


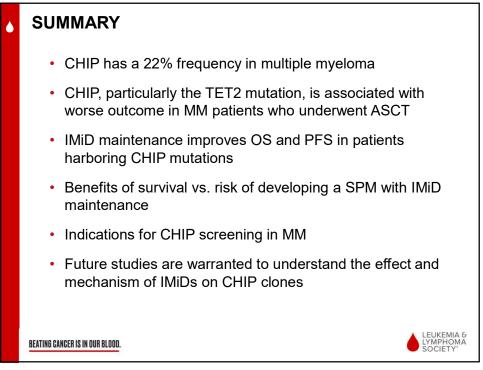


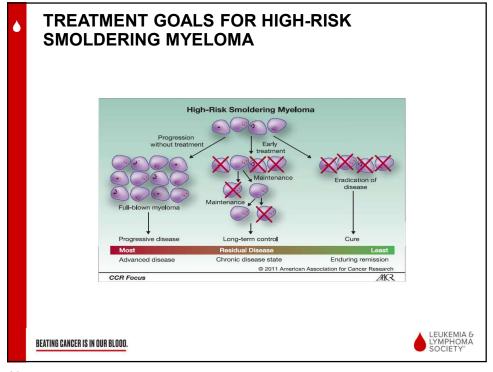


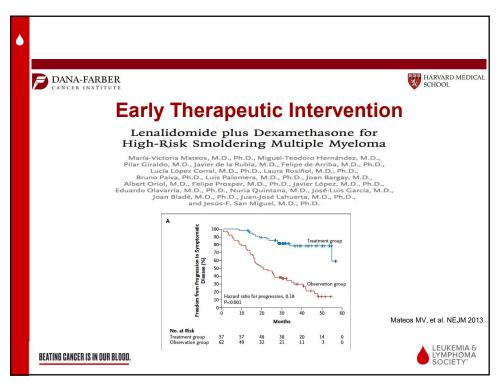


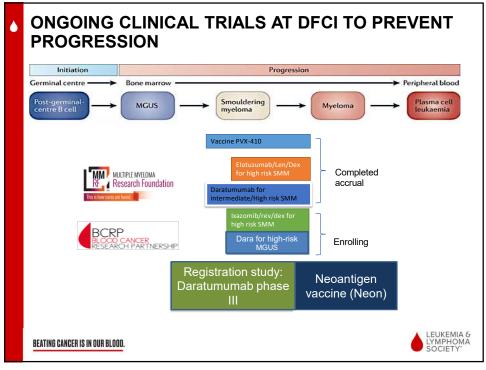


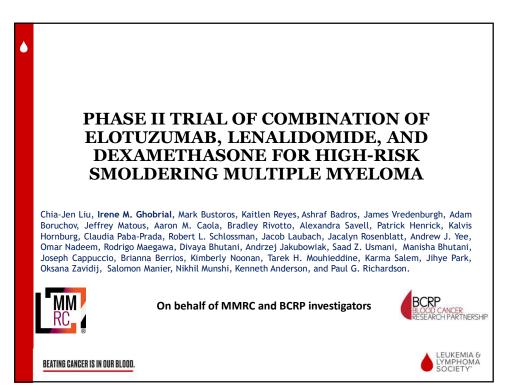


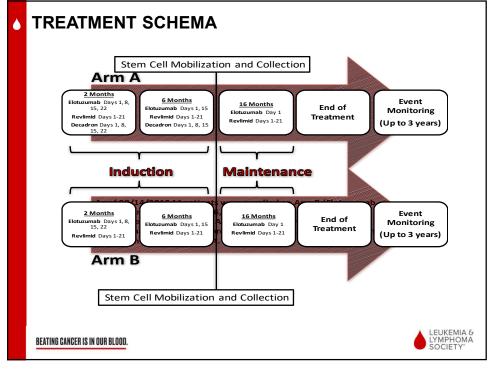




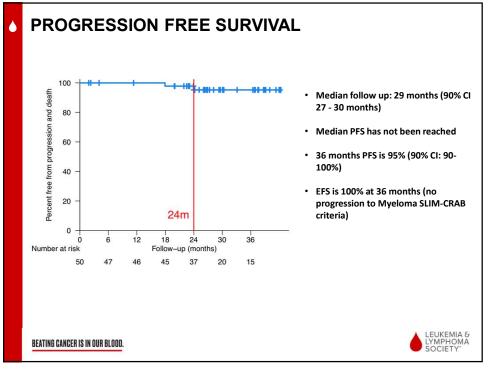


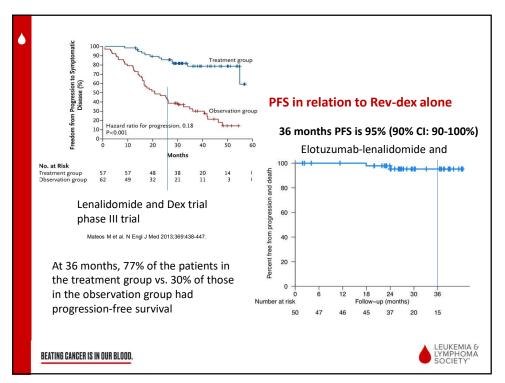


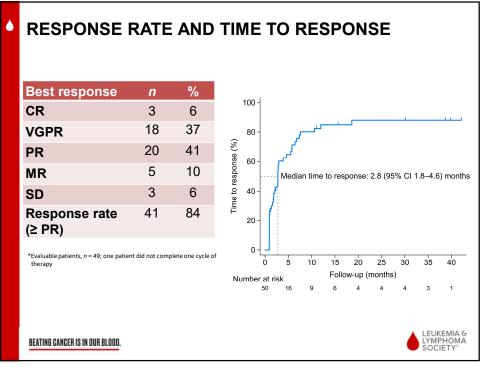


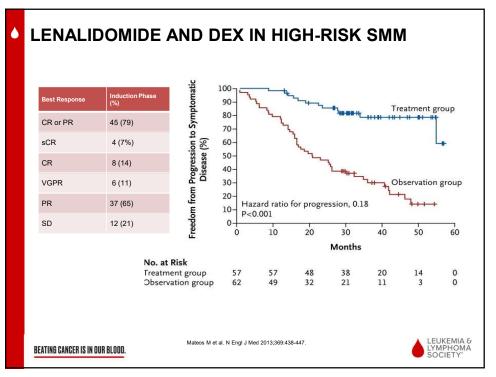


	Total (<i>n</i> = 50)			
Characteristics	n	%		
Median age, years (range)	62 (29–79)			
Male sex	18	36.0		
Race				
White	41	82.0		
Black	7	14.0		
Heavy-chain type				
lgG	33	66.0		
IgA	15	30.0		
BM plasma (%)	20.0 (10.0–60.0)			
β2-microglobulin, mg/dL	2.1 (0.8–5.9)			

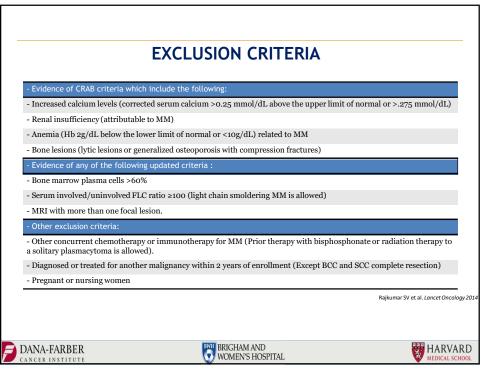




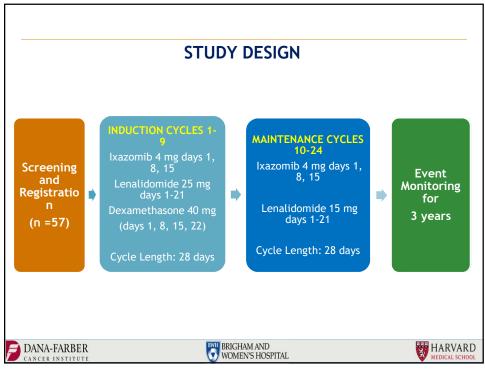








INCLUSION CRITERIA				
- Age ≥ 18 years - Bone marrow clonal plasma c	ells ≥10% and any one or more of the followi	ng:		
- Serum M protein ≥3.0g/dL	ř	0		
- IgA SMM				
- Immunoparesis with reduction of	two uninvolved immunoglobulin isotypes			
- Serum involved/uninvolved free light	ght chain ratio ≥8 (but less than 100)			
- Progressive increase in M protein	evel (Evolving type of SMM)†			
- Bone marrow clonal plasma cells 5	0-60%			
- Abnormal plasma cell immunophe more uninvolved immunoglobulin i	enotype (≥95% of bone marrow plasma cells are⊂cl sotypes	onal) and reduction of one or		
- t (4;14) or del 17p or 1q gain				
- MRI with diffuse abnormalities or	1 focal lesion			
- PET-CT with focal lesion with incr	eased uptake without underlying osteolytic bone des	struction		
- Monoclonal light chain excretion of	f 500mg/24 hours or higher*			
† Increase in serum monoclonal protein by ≥1 * Monoclonal Light Chain Smoldering	0% on two successive evaluations within a 6 month period.	Rajkumar SV, Mateos MV, Landgren O. B		
ANA FADRED	BYE BRIGHAM AND	H A RV		



	Total (n = 2	.9)			
Characteristics	n	%			
Median age, years (range)	61 (41-73)				
Male sex	15	51.7		Total (n = 29)
Heavy-chain type			High-risk criteria	n	%
IgG	19	65.5			
IgA	9	31.0	Bone marrow infiltration > 50%		27.5
No heavy chain	1	3.5	M-protein ≥ 3.0 g/dL	8	27.5
Light chain type			FLC ratio > 8		
Kappa	20	69.0	FLC ratio > 8		65.5
Lambda	9	31.0	Immunoparesis		75.8
Median (range)			IgA isotype SMM		31.0
Plasma cell percentage	20 (10-55)		0 51		
M-protein	1.7 (0.4-4.1)		Evolving M-protein	16	55.1
Absolute FLC ratio	11.6 (1.2-153.5)		High-risk cytogenetics*		54.2
Calcium	9.6 (8.7–10.4)				
Creatinine	0.9 (0.6–1.3)		High-risk cytogenetics are defined by the p	esence of t(4;14), t(14;16),
Hemoglobin, g/dL	13.0 (10.5-15.6)		deletion, and 1q gain.		
Median treatment cycles	14 (3–22)				
Median follow up time	15 months (4.2-22)				

