TCT2021 LBA5

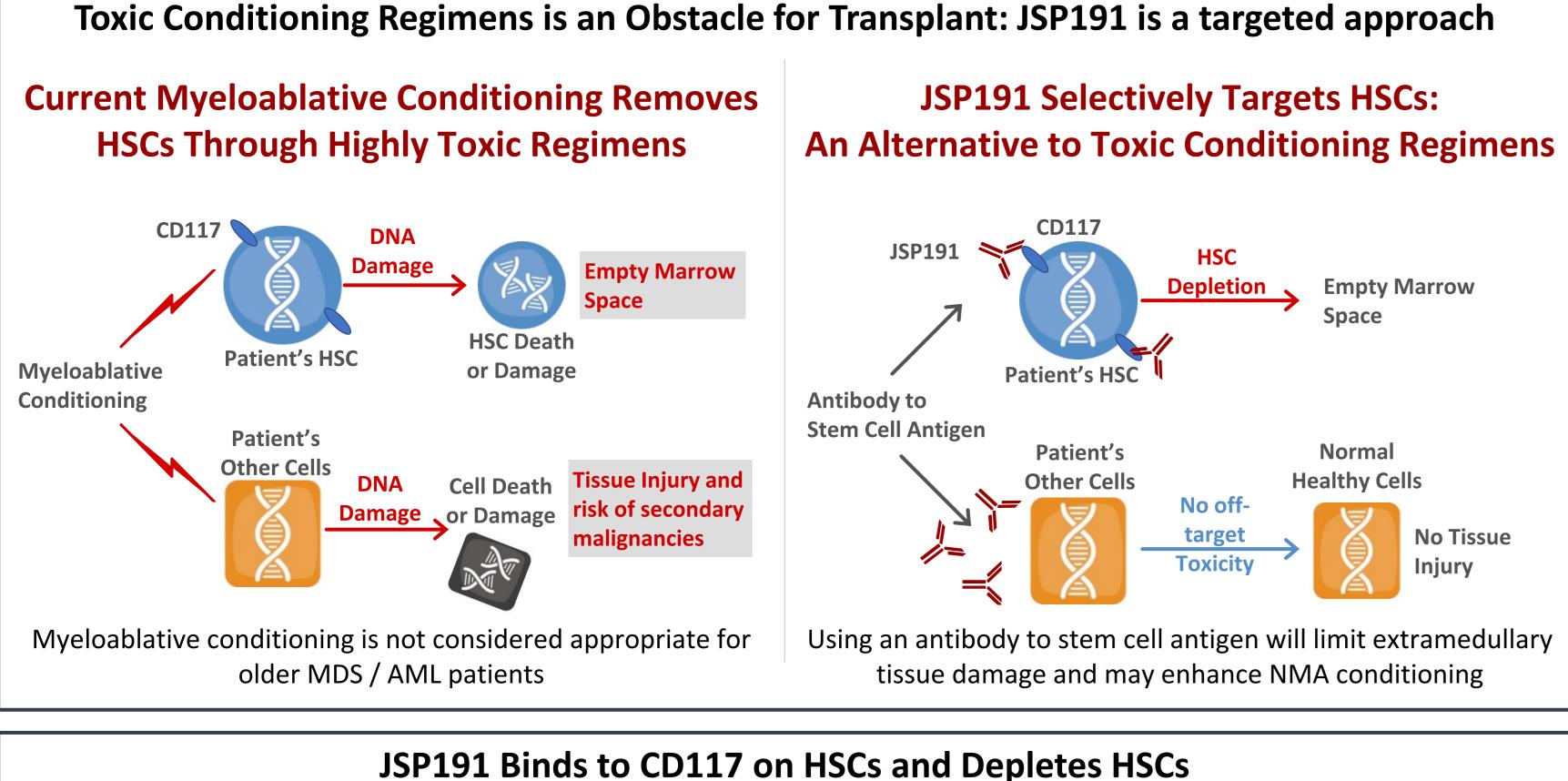
Phase 1 Study of JSP191, an Anti-CD117 Monoclonal Antibody, with Low Dose Irradiation and Fludarabine in Older Adults with MRD-Positive AML/MDS Undergoing Allogeneic HCT

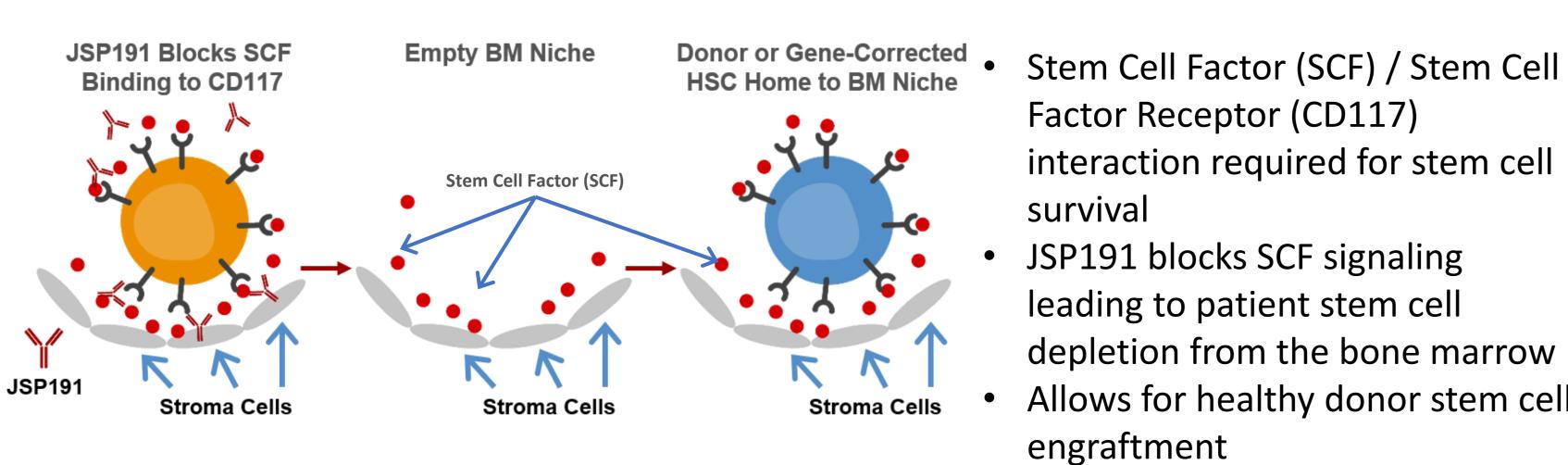
Lori Muffly, MD, MS¹, Hye-Sook Kwon, PhD², Michelle Chin, BS¹, Cara Lieber², Steve Smith³, Judith A. Shizuru, MD, PhD¹, Wendy W. Pang, MD, PhD² and Andrew S. Artz, MD⁴ ¹Department of Medicine, Division of Blood and Marrow Transplantation, Stanford University School of Medicine, Stanford, CA, ²Jasper Therapeutics, Inc., Redwood City, CA, ³Independent, San Jose, CA, ⁴Hematology/Hematopoietic Cell Transplant, City of Hope National Medical Center, Duarte, CA

TANFORD UNIVERSITY

Background

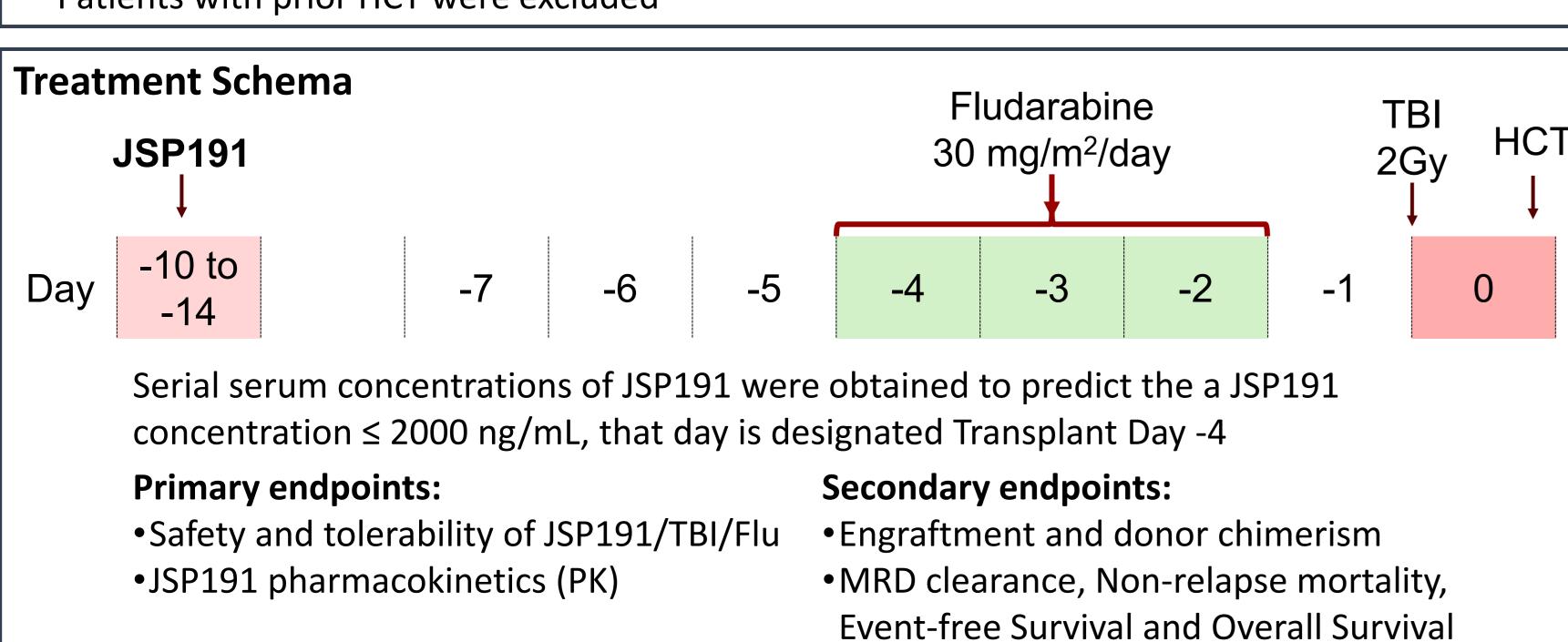
Myelodysplastic syndrome (MDS) and acute myeloid leukemia (AML) are hematologic malignancies primarily affecting older adults. Allogeneic hematopoietic cell transplantation (HCT) is potentially curative for MDS/AML, but intensive conditioning limits its application in older or frail patients. Non-myeloablative (NMA) or reduced intensity conditioning (RIC) achieves tolerability at the expense of heightened disease relapse; thus, innovative strategies to reduce relapse while maintaining low toxicity are needed. We are developing a first-in-class monoclonal antibody (mAb), JSP191, which targets and depletes normal and MDS/AML disease-initiating hematopoietic stem cells (HSC). JSP191 acts by inhibiting stem cell factor (SCF) binding to CD117 (c-Kit) present on HSC. We and others showed in pre-clinical models that HSC depletion and donor cell engraftment can be enhanced by combining anti-CD117 mAb with low dose total body radiation (TBI). Based on these data, we hypothesized that the addition of JSP191 prior to NMA HCT conditioning of 200 cGy TBI and fludarabine (Flu) would result in clearance of disease, lower toxicity, and reduced relapse in older patients with MDS/AML and measurable residual disease (MRD). This Phase 1 trial evaluates this clinical hypothesis (NCT#04429191).





Key Inclusion Criteria

- Patients with AML or MDS
- \geq 60 years or with HCT-Cl \geq 3
- Minimal Identifiable Disease (MID) or Measurable Residual Disease (MRD) detected by cytogenetics (cyto), difference from normal flow cytometry (flow), or next-generation sequencing (NGS)
- HLA matched related or unrelated donor
- Patients with prior HCT were excluded

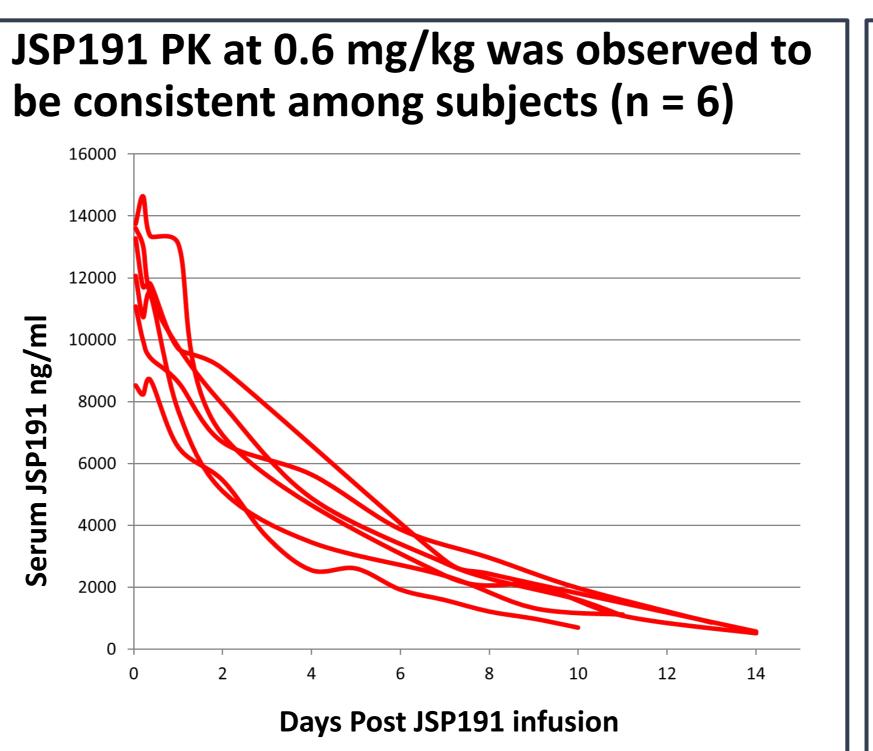




Subject	ect Age Pierresia Prior Therapy				
Number	Sex	Diagnosis	for MDS or AML	Donor	
003	74F	AML	Azacitidine/ Venetoclax	Matched unrelated	
004	70M	MDS	Erythropoietin	Matched related	
005	68M	MDS	Azacitidine	Matched unrelated	
009	74M	MDS	None	Matched unrelated	
010	65M	AML	Cytarabine/Idarubicin (7+3) +Midostaurin Azacitidine/ Venetoclax	Matched unrelated	
011	69M	AML	Cytarabine/ Daunorubicin (7+3) Cytarabine/ Daunorubicin (5+2)	Matched related	

JSP191 when added to TBI/Flu appears to be a safe and tolerable

- No infusion reactions
- No treatment related toxicities Protocol allows for outpatient conditioning All subjects are still on study



JSP191 Conditioning Leads to Successful Transplant and Conversion to MRD-Negative/MRD Reduction in First Five Evaluable Subjects

			-			
Subject Number	MRD at Screening	MRD at TD+28	MRD at TD+56	MRD at TD+90		
Subject Number	NGS, Flow, or Cyto	NGS, Flow, or Cyto	NGS, Flow, or Cyto	NGS, Flow, or Cyto		
	DNMT3A (VAF: 4.7%)	DNMT3A (VAF: 0.3%)	DNMT3A (VAF: 0.4%)	Subject still on study – assessments TBD		
003	RUNX1 (VAF: 1.7%)	RUNX1 (VAF: 0.3%)	RUNX1 (VAF: 0.3%)			
	PTPN11 (VAF: 0.7%)	NEG	NEG			
	ASXL1 (VAF: 0.3%)	NEG	ND	NEG		
004	PTPN11 (VAF: 0.4%)	NEG	ND	NEG		
	Del(20q)	NEG	ND	NEG		
	DNMT3A (VAF: 25.2%)	NEG	ND	Cubicat still an atualu		
005	SRSF2 (VAF: 0.3%)	NEG	ND	Subject still on study – assessments TBD		
	Flow 3.1%	NEG	ND			
009	Complex Cytogenetics	QNS	Subject still on study	y – assessments TBD		
009	Flow 0.7%	NEG	Subject still off study			
010	ASXL1 (VAF: 1.5%)	NEG	Subject still on study	accoccmonts TPD		
010	KMT2A duplication	KMT2A duplication	Subject still off study	- assessments TBD		
011	SRSF2 (VAF: 14.6%) Subject still on study – assessments TBD					
VAF: Variable allele frequency						

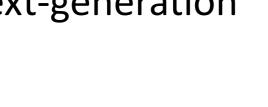
VAF: Variable allele frequency QNS: unable to obtain sufficient sample, will be repeated at TD+56 ND: MRD assays obtained on TD+56 only if TD+28 is positive

Mutation clearance after transplantation for MDS is associated with an improved Progression-Free Survival (Duncavage et al, NEJM 2018; 379:1028-41).

Factor Receptor (CD117) interaction required for stem cell

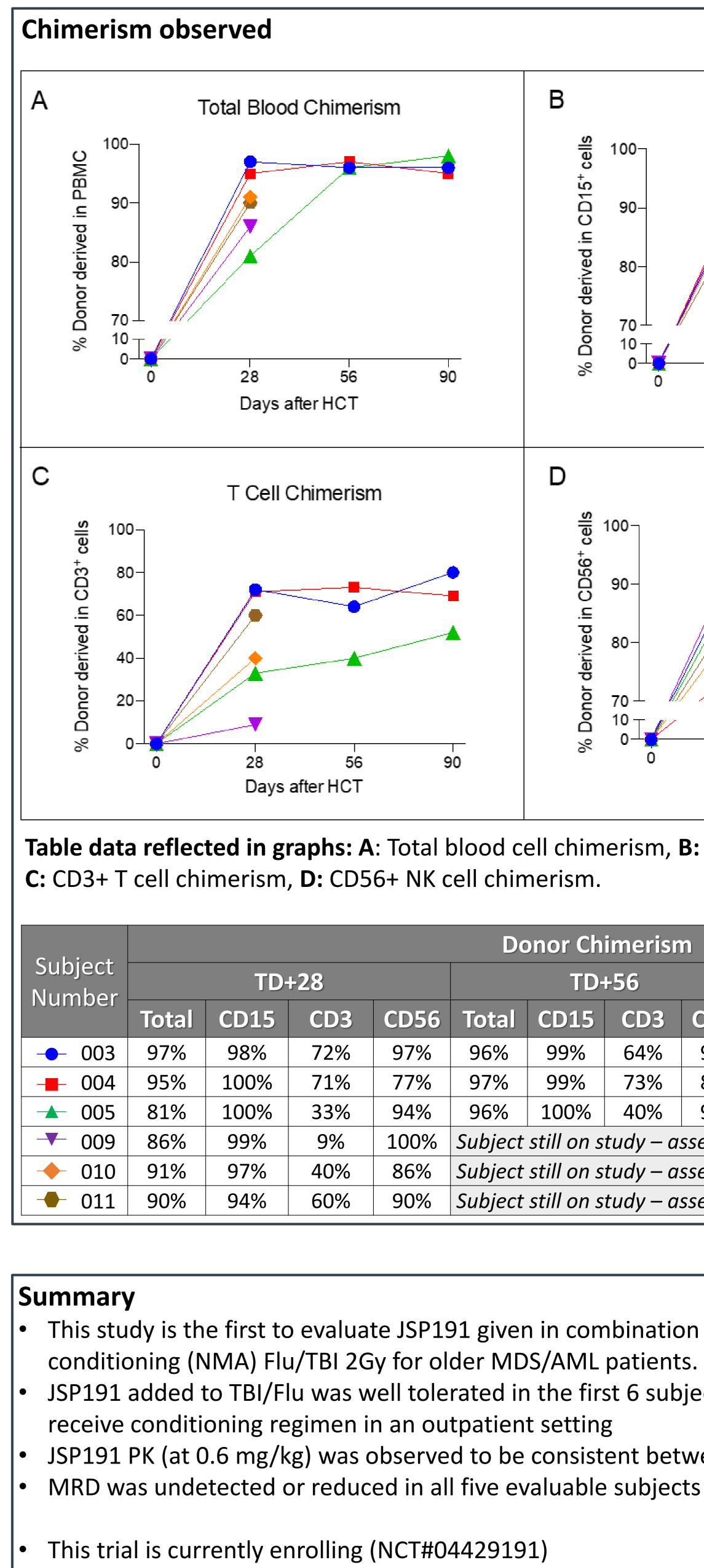
JSP191 blocks SCF signaling leading to patient stem cell depletion from the bone marrow Allows for healthy donor stem cell





Engraftment observed

Subject Number	Days from HCT to ANC >500
003	23
004	22
005	26
009	23
010	22
011	19



Acknowledgements

- this clinical trial



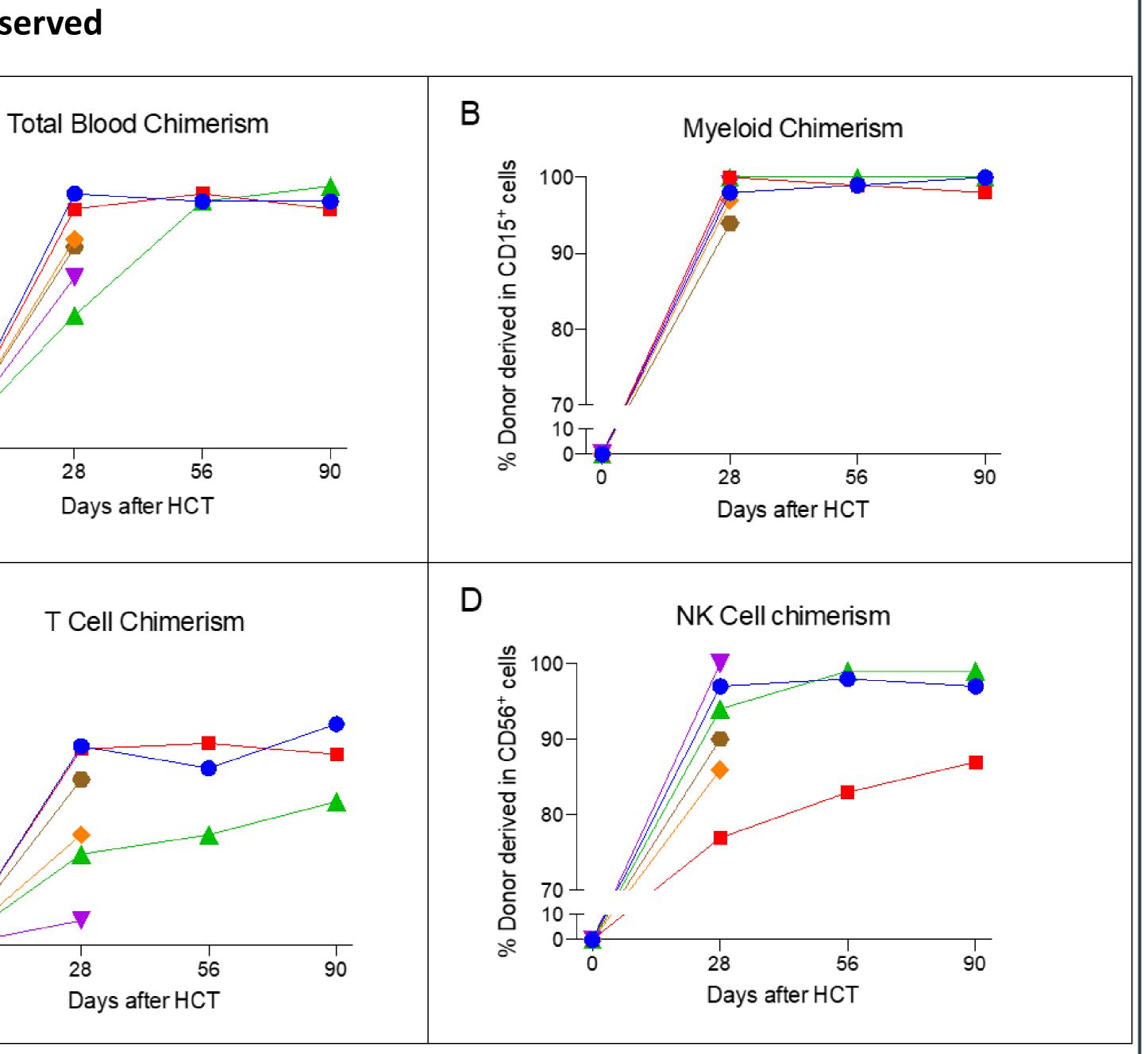


Table data reflected in graphs: A: Total blood cell chimerism, B: CD15+ Myeloid cell chimerism, **C:** CD3+ T cell chimerism, **D:** CD56+ NK cell chimerism.

	Donor Chimerism									
D+28			TD+56				TD+90			
	CD3	CD56	Total	CD15	CD3	CD56	Total	CD15	CD3	CD56
	72%	97%	96%	99%	64%	98%	96%	100%	80%	97%
	71%	77%	97%	99%	73%	83%	95%	98%	69%	87%
	33%	94%	96%	100%	40%	99%	98%	100%	52%	99%
	9%	100%	Subject still on study – assessments TBD							
	40%	86%	Subject still on study – assessments TBD							
	60%	90%	Subject still on study – assessments TBD							

This study is the first to evaluate JSP191 given in combination with non-myeloablative

- JSP191 added to TBI/Flu was well tolerated in the first 6 subjects; protocol allows for subjects to receive conditioning regimen in an outpatient setting
- JSP191 PK (at 0.6 mg/kg) was observed to be consistent between subjects
- MRD was undetected or reduced in all five evaluable subjects at TD+28 and are all still on study

This trial is currently enrolling (NCT#04429191)

The 2021 TCT Meetings Digital Experience

February 8-12, 2021

We wish to thank Kevin N. Heller, Janet Hurt, Joe Laver, Susan Prohaska, Beverly Smith, and Bin Yao for their tireless effort to generate data and support this clinical trial

Jasper Therapeutics and the Investigators thank the patients and families for participating in

