

The Effect of Bezuclastinib on the Pathobiology of Advanced Systemic Mastocytosis: Results From the Pivotal Apex Trial

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INTRODUCTION

Bone Marrow Pathobiology Underpins the Clinical and Biological Complexity of AdvSM

- AdvSM is a life-threatening disease characterized by abnormal mast cell accumulation, driven by the gain-of-function KIT p.D816V mutation in most patients^{1,4}
 - Subtypes: aggressive SM (ASM), SM with associated hematologic neoplasm (SM-AHN), mast cell leukemia (MCL)
- Pathological and molecular markers of disease help determine AdvSM subtype and disease severity, monitor treatment response, and predict prognosis and survival⁵⁻¹⁰
 - Per 2022 WHO criteria, SM diagnosis requires 1 major plus 1 minor OR ≥3 minor criteria⁵

Major Criterion	Minor Criteria
Multifocal dense mast cell infiltrates (≥15 in aggregates) in BM or other extracutaneous organ(s)	Aberrant CD25, CD30, and/or CD2 expression on mast cells in BM, blood, or other extracutaneous organ(s) >25% atypical or spindle-shaped mast cells in infiltrates in BM or other extracutaneous organ(s) Baseline serum tryptase >20 ng/mL in the absence of AHN, levels should be adjusted in the case of Hct† KIT p.D816V or other activating <i>KIT</i> mutation detected in BM or other extracutaneous organ(s)

Bezuclastinib is an Investigational, Oral, Highly Selective Tyrosine Kinase Inhibitor with Potent Activity Against KIT D816V

- Results from the Apex study of bezuclastinib demonstrated favorable efficacy and safety in AdvSM¹¹
- Herein we report the effect of bezuclastinib on pathological markers of SM from the pivotal portion of Apex

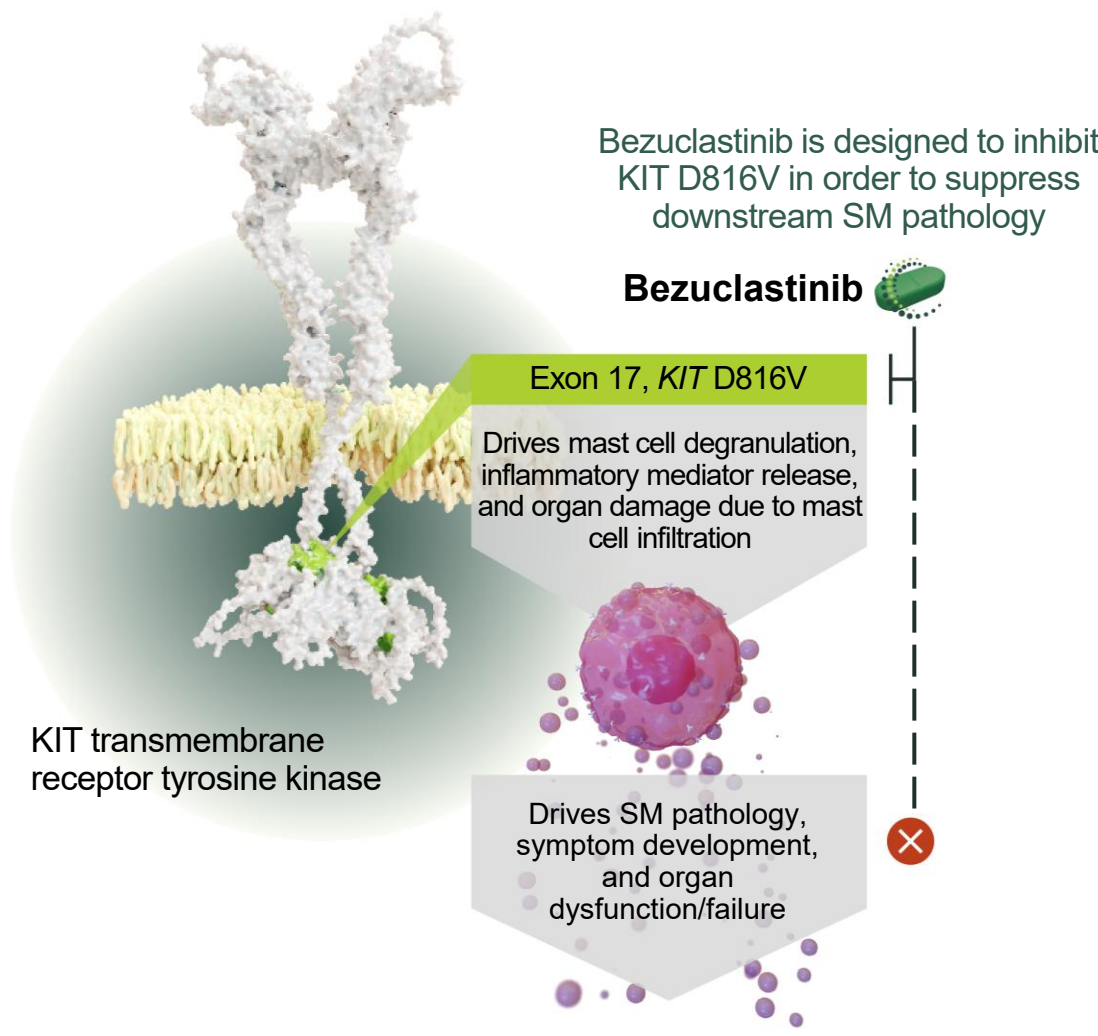
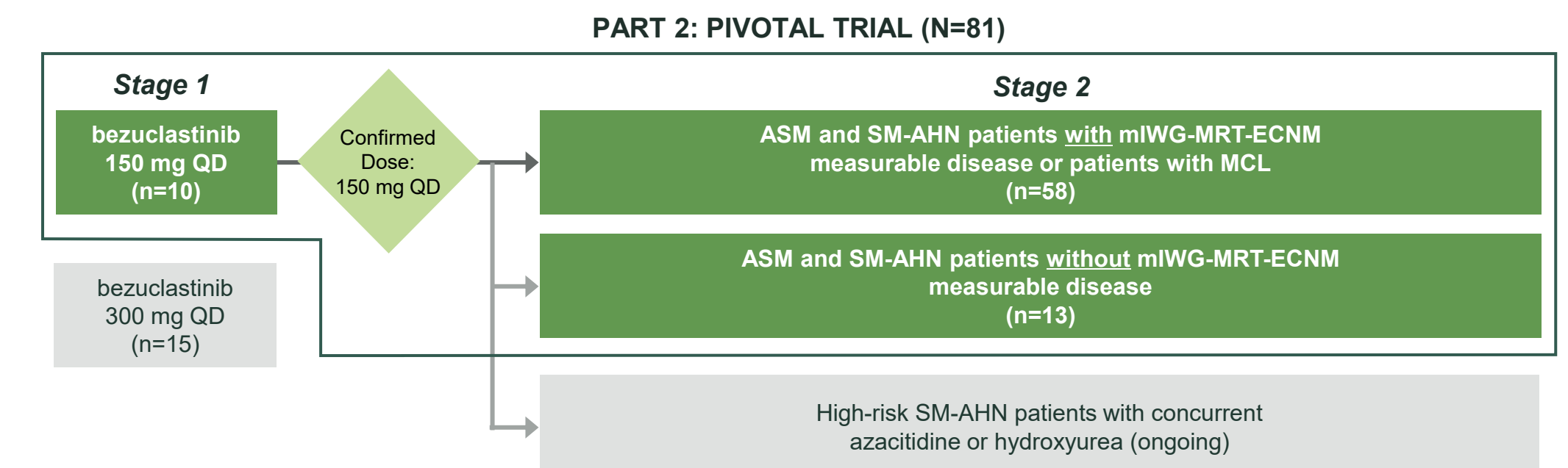


Figure developed by Cogent Biosciences, Inc.

METHODOLOGY

Apex (NCT04996875): Phase 2, Multipart, Multicenter, Open-Label Study Evaluating Efficacy and Safety of Bezuclastinib in Patients With AdvSM

Eligibility*
- ASM, SM-AHN, or MCL per WHO 2022 Classification - No restrictions on prior therapy - Serum tryptase ≥20 ng/mL - Platelet count ≥50×10⁹/L



Part 2 Endpoints
Primary: ORR (CR/CRh + PR + CI) per miWG-MRT-ECNM criteria
Key Secondary: ORR (CR/CRh + PR) per Pure Pathologic Response (PPR) criteria
Other Secondary & Exploratory*: Changes in objective measures of disease (MC & BM characteristics)

- BM biopsy and aspirate, serum tryptase, and KIT p.D816V variant allele frequency (VAF) collections were tested centrally at baseline, every three 28-day cycles for the first 12 months, and then every six cycles
- Morphology, mast cell burden, aggregation, and immunophenotype (tryptase, CD25, CD30) were assessed
- Immunohistochemical (tryptase, CD25, CD30, CD34, CD117) and special (reticulin, trichrome) stains were performed on BM sections

*Additional criteria and endpoints are defined in the protocol.

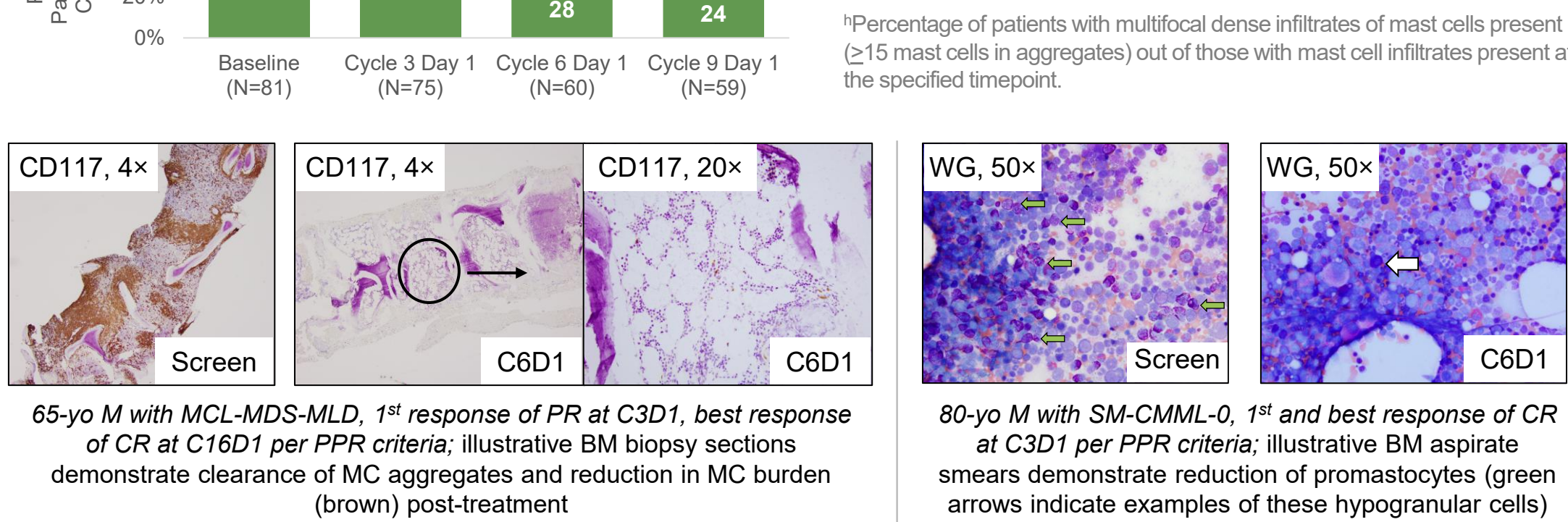
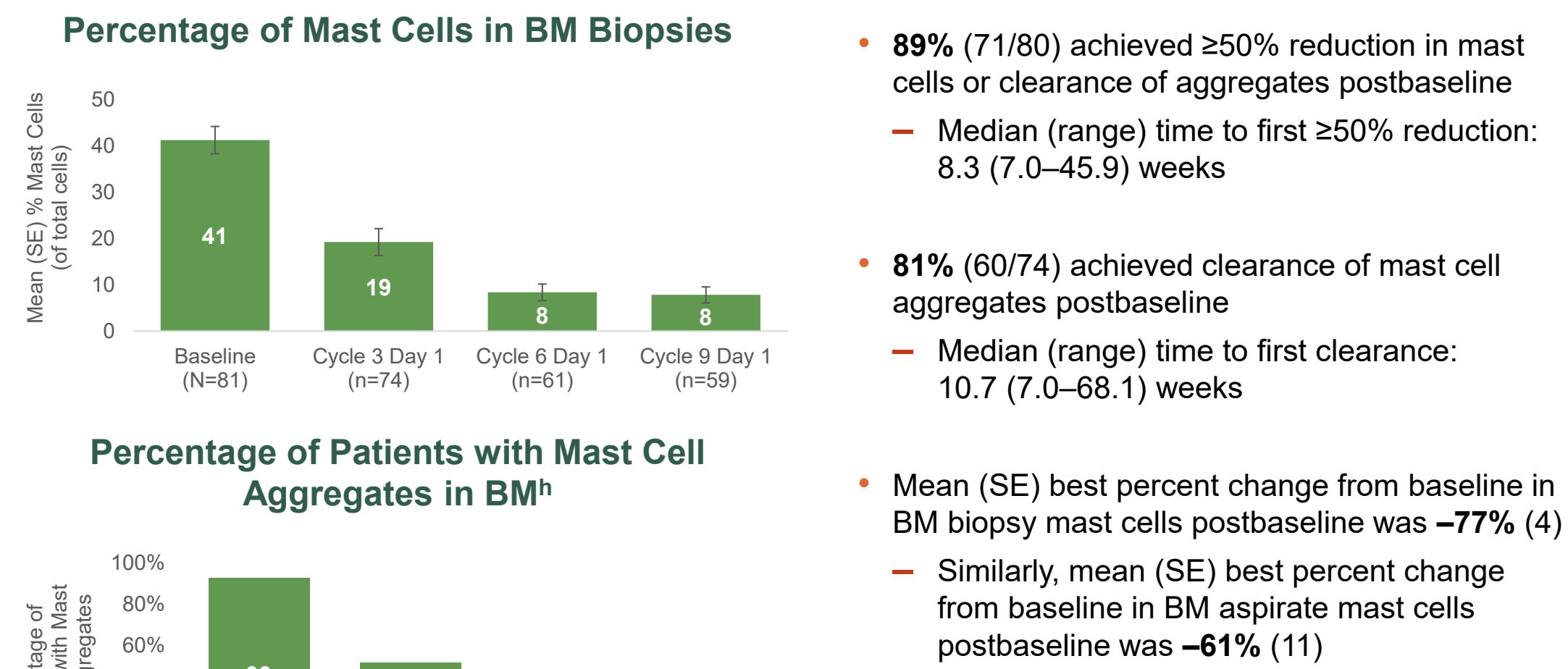
Baseline Characteristics Reflected a Broad AdvSM Population

Characteristic	Bezuclastinib 150 mg QD (N=81)	Characteristic	Bezuclastinib 150 mg QD (N=81)
Female, n (%)	23 (28)	Median <i>KIT</i> p.D816V in peripheral blood, % (range)	6.7 (0-91.4)
Median age, years (range)	70 (43-87)	<i>KIT</i> p.D816V VAF detected*, n (%)	71 (90) [†]
ECOG Performance Status, n (%)	0: 22 (27) ≥2: 40 (49)	Median BM mast cell burden, % (range)	35 (3-95)
AdvSM subtype per central eligibility review, n (%)	ASM: 11 (14) SM-AHN ^b : 57 (70) MCL ^c : 13 (16)	Median serum tryptase, ng/mL (range)	191 (23-1987)
<i>SRSF2/ASXL1/RUNX1</i> mutation detected in peripheral blood, n (%)	48 (59)	Myelofibrosis Grade, n (%)^d	MF-0: 7 (9) MF-1: 38 (48) MF-2: 26 (33) MF-3: 9 (11)
Prior TKI therapy for AdvSM, n (%)	28 ^e (35)	ORR-Evaluable Populations, n (%)	
Prior midostaurin	24 (30)	Patients evaluable per miWG-MRT-ECNM criteria by Eligibility Committee	68 (84)
Prior avapritinib	6 (7)	Patients evaluable per PPR criteria	81 (100)
Prior imatinib	3 (4)		

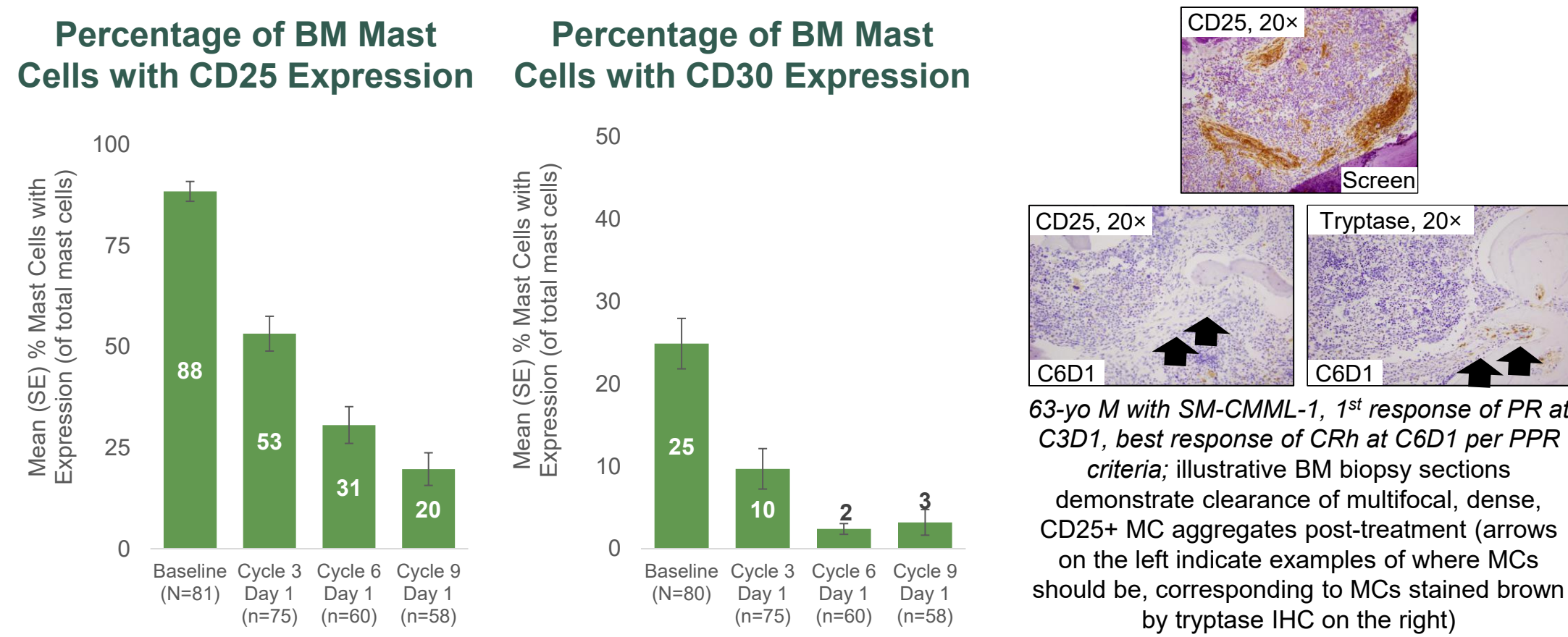
- Median (range) duration of treatment at data cutoff: 63.0 (1.0-135.3) weeks

^aAHN subtypes: CMML (n=37), MDS (n=6), MPN (n=4), MDS/MPN-UJ (n=10). ^bIncludes 2 with MCL-AHN (MDS-MLD). ^c5 received 2 prior TKIs. ^dDroplet digital PCR limit of detection=0.03%. ^eOf 79 with baseline assessment. ^fOf 80 with baseline assessment.

Bezuclastinib Reduced Mast Cell Burden and Cleared Pathologic Mast Cell Aggregates, the Major Criterion for SM Diagnosis, in Bone Marrow

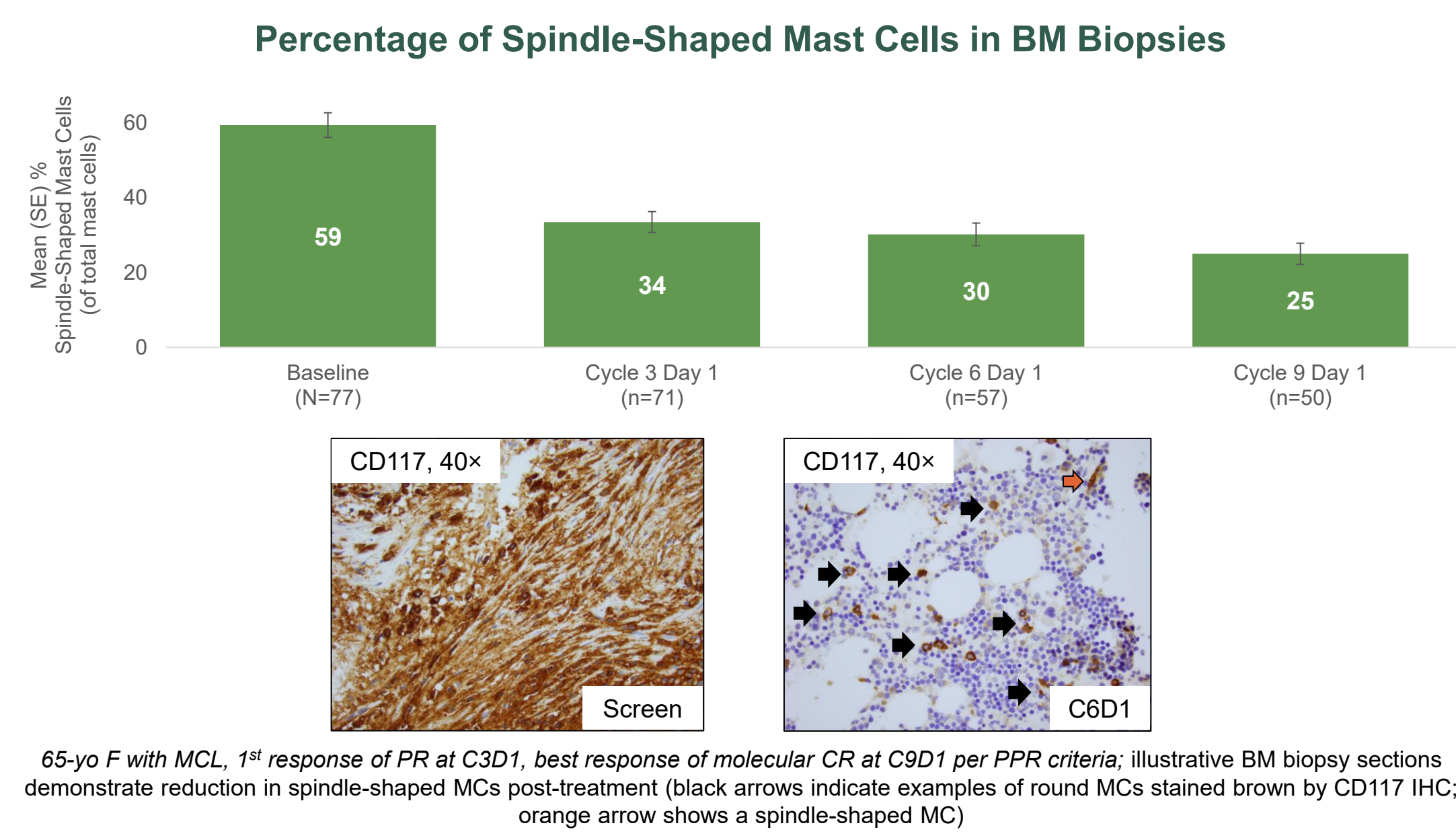


Bezuclastinib Reduced Aberrant CD25 and CD30 Expression on Mast Cells in Bone Marrow, Supporting Targeted Mast Cell Modulation

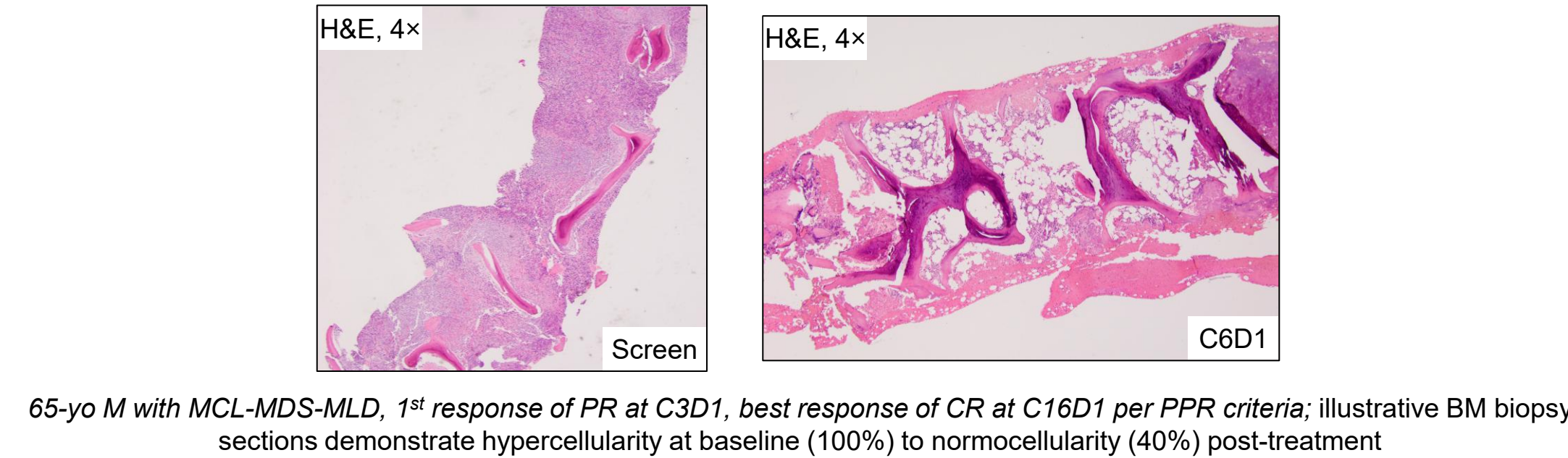
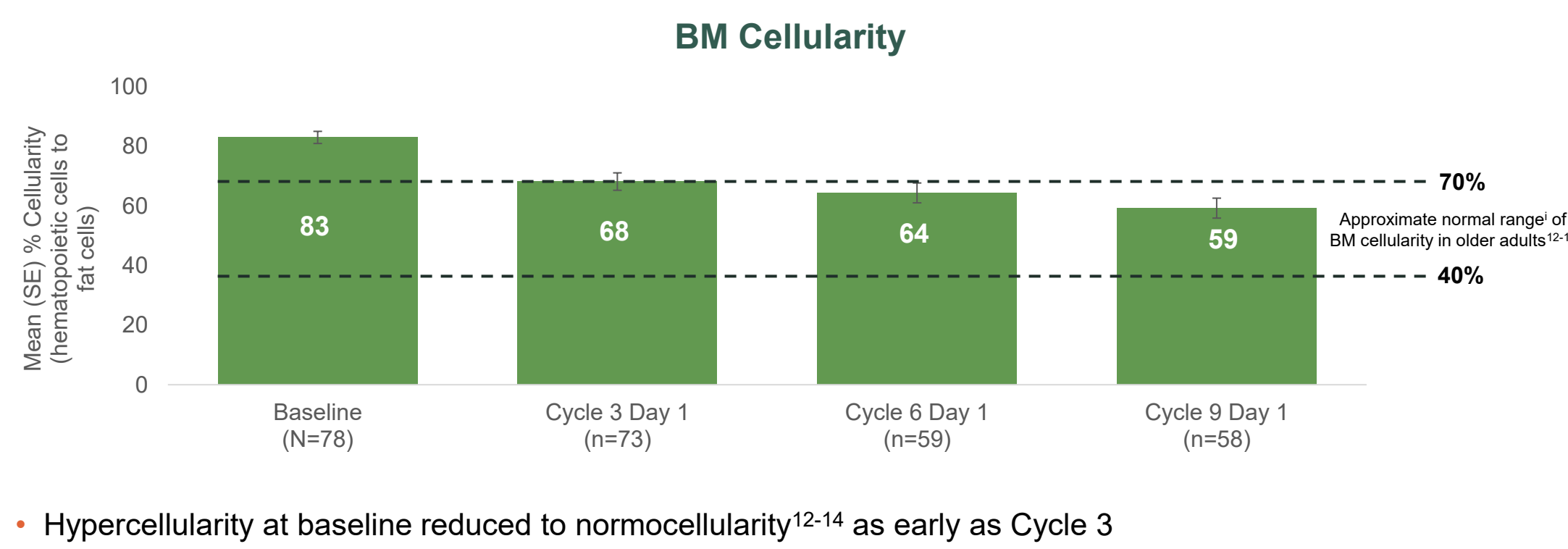


RESULTS

Bezuclastinib Reduced Atypical Mast Cell Morphology in Bone Marrow

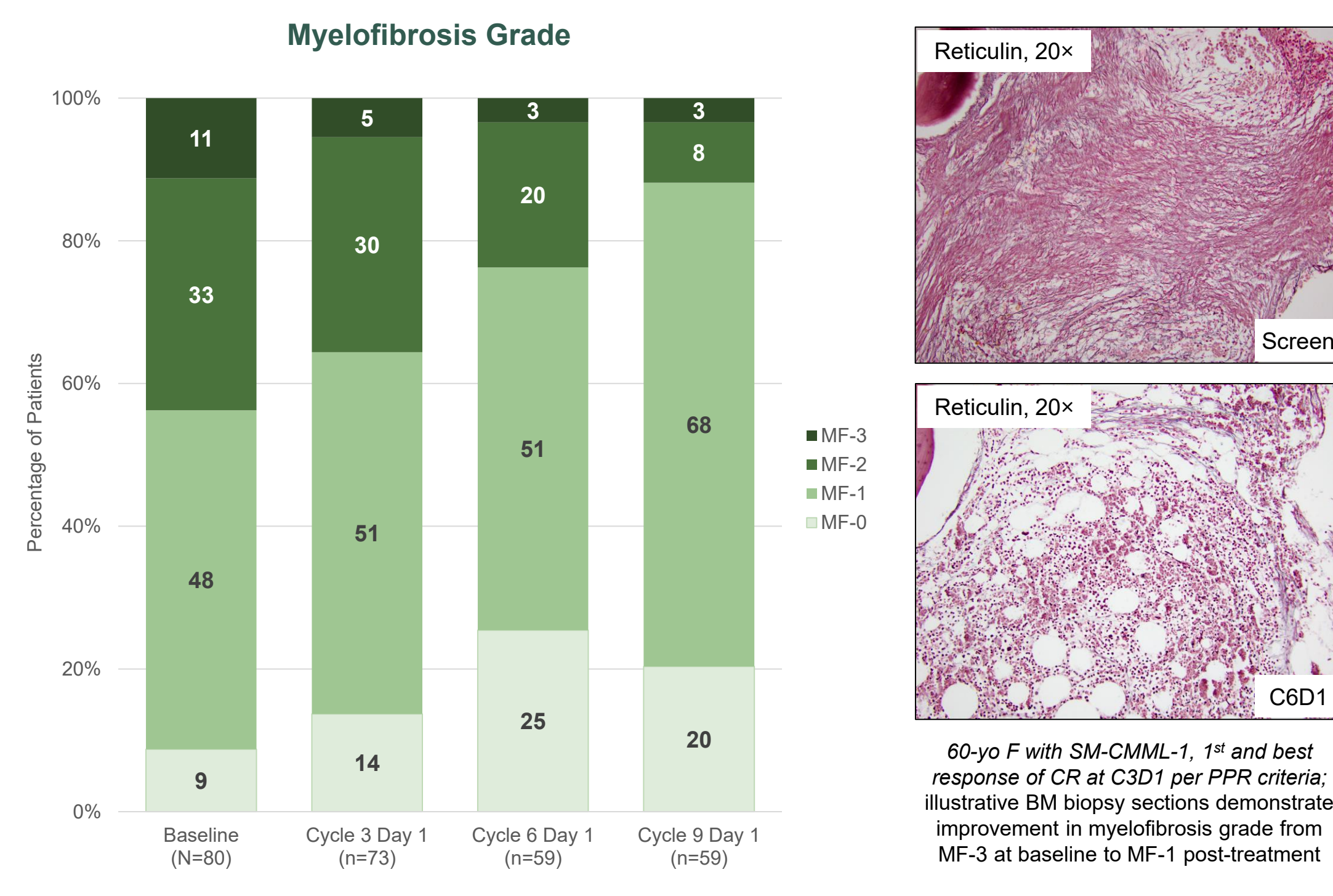


Bezuclastinib Reduced Bone Marrow Cellularity

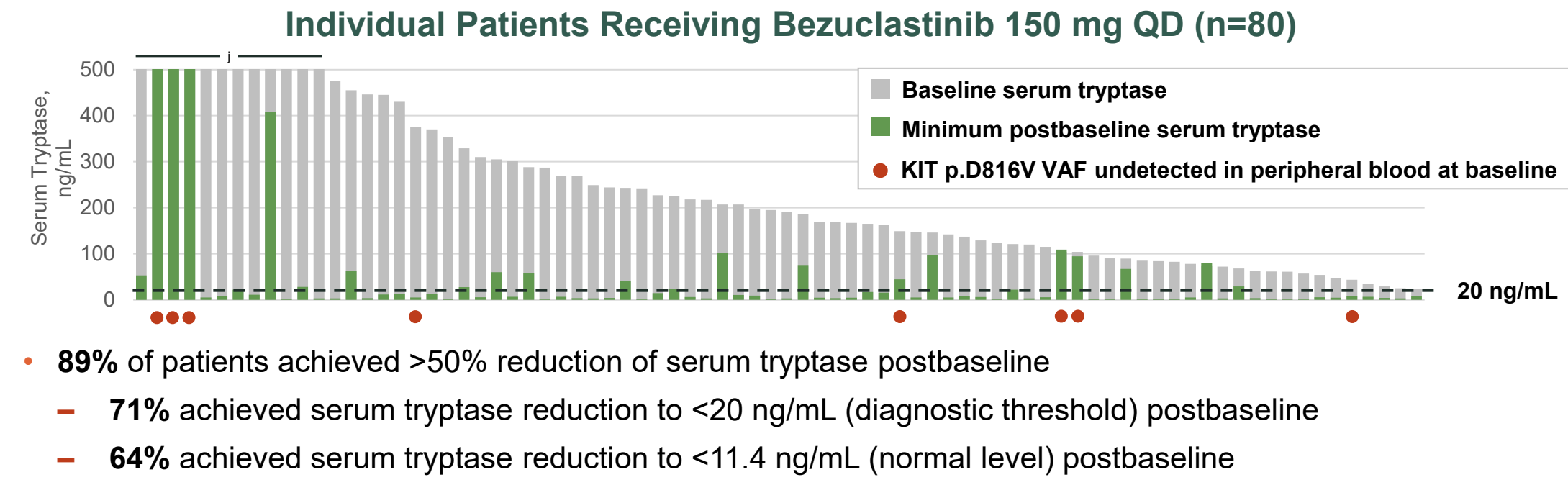


Normal BM cellularity range is age-dependent.

Bezuclastinib Treatment Led to Improvement in Myelofibrosis Grade

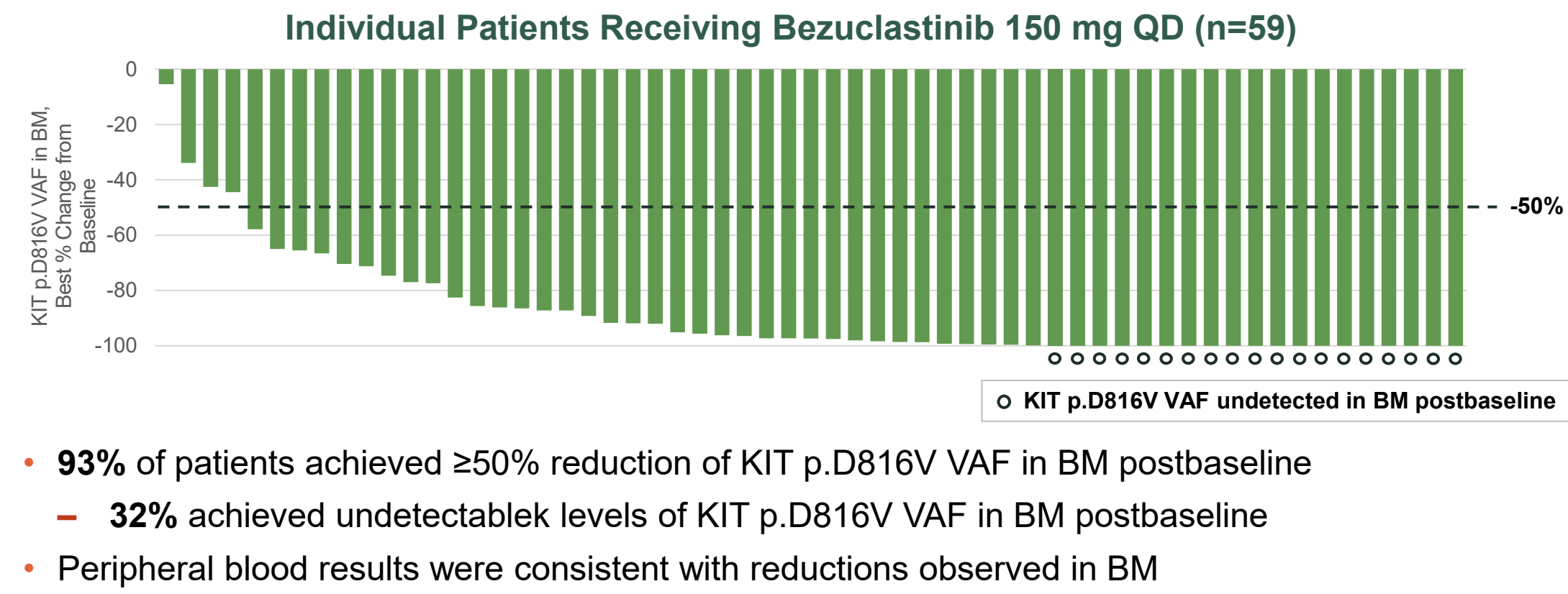


Bezuclastinib Reduced Serum Tryptase to Below Diagnostic Threshold Levels



^aPatients with serum tryptase levels >500 ng/mL, axis is truncated for visual purposes.

Bezuclastinib Reduced KIT p.D816V Variant Allele Frequency (VAF) in Bone Marrow and Blood



^aDroplet digital PCR limit of detection=0.03%.

Bezuclastinib Demonstrated High Response Rates in AdvSM per Pure Pathologic Response (PPR) Criteria

Best Confirmed Response		All Evaluable Patients (N=81)
Overall Response Rate, n (%)	CR+CRh+PR	66 (81)
Best Overall Response, n (%)	Complete Remission (CR+CRh)	50 (62)
	Molecular CR/CRh	20 (25)
	Partial Remission (PR)	16 (20)
	Stable Disease	14 (17)
	Progressive Disease	0
	Not Evaluable	1 (1)

CONCLUSIONS

Pathobiology Results from the Apex Study Support the Potential of Bezuclastinib to Treat Underlying Disease Biology in AdvSM

- Bezuclastinib demonstrated robust changes in disease pathology, with effects observed as early as the first assessment at 8 weeks
 - High pure pathologic response (PPR) rates
 - Improvement (including normalization) in bone marrow mast cell distribution (aggregates and burden), atypical morphology, and aberrant immunophenotype
 - Improvements in broader bone marrow characteristics (cellularity, myelofibrosis)
 - Majority achieved normalization of serum tryptase
 - Approximately one-third achieved undetectable levels of *KIT* p.D816V VAF in bone marrow
- Marked reductions in objective measures of disease in most patients signifies impact on KIT-driven pathology and modification of underlying AdvSM disease with bezuclastinib treatment

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Abbreviations: AdvSM, advanced SM; AHN, associated hematologic neoplasm; ASM, aggressive SM; BM, bone marrow; CI, clinical improvement; CMML, chronic myelomonocytic leukemia; CR, complete remission; CRh, CR with partial hematologic recovery; CXDX, Cycle X Day X; ECOG, Eastern Cooperative Oncology Group; F, female; Hct, hereditary alpha-tryptasemia; H&E, hematoxylin and eosin stain; IHC, immunohistochemistry; M, male; MC, mast cell; MCL, mast cell leukemia; MCL-AHN, MCL with AHN; MDS, myelodysplastic syndrome; MDS/MPN-U, MDS/MPN undifferentiated; MF, myelofibrosis; miWG-MRT-ECNM, modified International Working Group–Myeloproliferative Neoplasms Research and Treatment–European Competence Network on Mastocytosis; MLD, multilineage dysplasia; MPN, myeloproliferative neoplasm; ORR, overall response rate; PCR, polymerase chain reaction; PPR, Pure Pathologic Response; PR, partial remission; QD, once daily; SE, standard error; SM, systemic mastocytosis; SM-AHN, SM with AHN; TKI, tyrosine kinase inhibitor; VAF, variant allele frequency; WG, Wright-Giemsa stain; WHO, World Health Organization; yo, year-old.

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