

# Prognostic impact of *SF3B1* mutation and multilineage dysplasia in myelodysplastic syndromes with ring sideroblasts: a Mayo Clinic study of 170 informative cases

Faiqa Farrukh,<sup>1</sup> Maymona Abdelmagid,<sup>1</sup> Abhishek Mangaonkar,<sup>1</sup> Mrinal Patnaik,<sup>1</sup> Aref Al-Kali,<sup>1</sup> Michelle A. Elliott,<sup>1</sup> Kebede H. Begna,<sup>1</sup> Christopher C. Hook,<sup>1</sup> William J. Hogan,<sup>1</sup> Animesh Pardanani,<sup>1</sup> Mark R. Litzow,<sup>1</sup> Rhett P. Ketterling,<sup>2</sup> Naseema Gangat,<sup>1</sup> Daniel A. Arber,<sup>3</sup> Attilio Orazi,<sup>4</sup> Rong He,<sup>2</sup> Kaaren Reichard<sup>2</sup> and Ayalew Tefferi<sup>1</sup>

<sup>1</sup>Division of Hematology, Department of Medicine and Laboratory Medicine, Mayo Clinic, Rochester, MN; <sup>2</sup>Division of Hematopathology, Department of Medicine and Laboratory Medicine, Mayo Clinic, Rochester, MN; <sup>3</sup>University of Chicago, Chicago, IL and <sup>4</sup>Department of Pathology, Texas Tech University Health Sciences Center, El Paso, TX, USA

**Correspondence:** A. Tefferi  
[tefferi.ayalew@mayo.edu](mailto:tefferi.ayalew@mayo.edu)

**Received:** November 21, 2023.  
**Accepted:** February 29, 2024.  
**Early view:** March 7, 2024.

<https://doi.org/10.3324/haematol.2023.284719>

©2024 Ferrata Storti Foundation  
Published under a CC BY-NC license 

**Supplemental table 1:** Phenotypic and genotypic characteristics of 170 patients with myelodysplastic syndromes with ring sideroblasts (MDS-RS), stratified by presence or absence of multilineage dysplasia

Variables	All patients n=170	MDS-RS-MLD n=87 (51%)	MDS-RS-SLD n=83 (49%)	P value
Age in years; median (range)	73 (41-94)	74 (47-94)	72 (41-89)	0.81
Males; n (%)	114 (67)	62 (71)	52 (63)	0.23
Hemoglobin (g/dl) median (range)*	9.6 (6.4-13.7)	9.3 (6.4-13.7)	9.9 (6.5-12.2)	0.19
Transfusion dependence; n (%)	38 (22)	23 (26)	15 (18)	0.19
Leukocytes $\times 10^9/l$ ; median (range)	N=169  5.1 (0.7-13.1)	N=87  4.4 (0.7-13.1)	N=82  5.6 (1.4-12.7)	<b>&lt;0.01</b>
ANC $\times 10^9/l$ ; median (range)	N=167  2.8 (0.2-9.4)	N=87  2.6 (0.2-9.4)	N=80  3 (0.2-8.4)	0.06
ANC $<1 \times 10^9/l$ ; n (%)	11 (7)	9 (10)	2 (2)	<b>0.02</b>
ALC $\times 10^9/l$ ; median (range)	N=166  1.3 (0.04-3.6)	N=86  1.2 (0.2-3.1)	N=80  1.6 (0.04-3.6)	<b>0.001</b>
ALC $<1.2 \times 10^9/l$ ; n (%)	65 (39)	40 (47)	25 (31)	<b>0.04</b>
AMC $\times 10^9/l$ ; median (range)	N=165  0.4 (0.02-1.6)	N=85  0.4 (0.02-1.6)	N=80  0.5 (0.06-1.1)	0.15
Platelets $\times 10^9/l$ ; median (range)	N=169  229 (20-599)	N=87  202 (20-599)	N=82  275 (46-585)	<b>&lt;0.01</b>

<b>Platelets &lt;100 × 10<sup>9</sup>/l; n (%)</b>	19 (11)	16 (18)	3 (4)	<b>&lt;0.01</b>
<b>Bone marrow blast %; median (range)</b>	1 (0-4)	1 (0-4)	1 (0-4)	0.14
<b>Peripheral blood blast %; median (range)</b>	0 (0-1)	0 (0-1)	0 (0-0)	0.37
<b>Karyotype (n evaluable = 166)</b>				<b>&lt;0.01</b>
<i>Normal; n (%)</i>	121 (73)	53 (63)	68 (83)	
<i>Sole loss of "Y"; n (%)</i>	9 (5)	5 (6)	4 (5)	
<i>Sole trisomy 8; n (%)</i>	8 (5)	3 (2)	5 (3)	
<i>Sole 20q-; n (%)</i>	7 (4)	4 (5)	3 (4)	
<i>Complex; n (%)</i>	5 (3)	5 (6)	0	
<i>-7/7q-; n (%)</i>	3 (2)	3 (4)	0	
<i>Other; n (%)</i>	13 (8)	11 (13)	2 (2)	
<b>Mutations** (n evaluable = 145)</b>				
<i>SF3B1; n (%)</i>	126 (87)	59 (82)	67 (92)	0.08
<i>TET2; n (%)</i>	37 (25)	15 (21)	22 (30)	0.2
<i>DNMT3A; n (%)</i>	28 (19)	13 (18)	15 (21)	0.7
<i>ASXL1; n (%)</i>	16 (11)	7 (10)	9 (12)	0.61
<i>SRSF2; n (%)</i>	7 (5)	2 (3)	5 (7)	0.25
<i>TP53; n (%)</i>	7 (5)	3 (4)	4 (5)	0.73
<i>IDH1; n (%)</i>	3 (2)	3 (4)	0 (0)	0.08
<i>CSF3R; n (%)</i>	3 (2)	0 (0)	3 (4)	0.08
<b>Erythropoietin IU/l; median (range)</b>	N=122 69 (7-5461)	N=59 73 (13-5461)	N=63 61 (7-1648)	0.36
<b>LDH U/l; median (range)</b>	N=83 194 (112-538)	N=46 185 (112-413)	N=37 198 (112-538)	0.47
<b>Bone marrow ring sideroblasts %; median (range)</b>	40 (5-80)	35 (10-80)	45 (5-80)	0.11

<b>Median follow-up for patients alive.</b> <b>median years (range)</b>	5.2 (0.13-12.6)	5.4 (0.12-9)	5.0 (1.0-12.6)	0.93
<b>Deaths; n (%)</b>	104 (61)	63 (72)	41 (49)	<b>&lt;0.01</b>
<b>Leukemic transformation; n (%)</b>	8 (5)	6 (7)	2 (2)	0.16
<b>Allogeneic stem cell transplant; n (%)</b>	7 (4)	5 (6)	2 (2)	0.27

\*Hemoglobin for patients with transfusion independent anemia at time of diagnosis.

\*\*mutations present in at least 3 patients.

Abbreviations: ALC, absolute lymphocyte count; ANC, absolute neutrophil count; AMC, absolute monocyte count; IPSS-R, Revised International Prognostic Scoring System; MDS-RS, MDS with ring sideroblasts; MDS-RS-MLD, MDS with ring sideroblasts and multilineage dysplasia; MDS-RS-SLD, MDS with ring sideroblasts and single lineage dysplasia

**Supplemental table 2:** Phenotypic and genotypic characteristics of 145 patients with myelodysplastic syndromes with ring sideroblasts (MDS-RS), stratified by presence or absence of *SF3B1* mutation

Variables	All patients N=145	<i>SF3B1</i> mutated n=126	<i>SF3B1</i> wild-type n=19	P value
Age in years; median (range)	72 (41-94)	72 (41-94)	72 (55-83)	0.6
Males; n (%)	98 (68)	82 (65)	16 (84)	0.08
Hemoglobin (mg/dl) median (range)*	9.6 (6.5-13.7)	9.5 (6.5-13.5)	10.2 (6.6-13.7)	0.2
Transfusion dependence; n (%)	36 (25)	33 (26)	3 (16)	0.31
Leukocytes $\times 10^9/l$ ; median (range)	5.2 (0.7-13.1)	5.5 (1.5-13.1)	3.7 (0.7-9.4)	<0.01
ANC $\times 10^9/l$ ; median (range)	3 (0.2-9.8)	3 (0.4-9.8)	1.9 (0.2-5.6)	<0.01
ANC $< 1 \times 10^9/l$ ; n (%)	9 (6)	5 (4)	4 (21)	0.02
ALC $\times 10^9/l$ ; median (range)	1.4 (0.04-3.6)	1.4 (0.2-3.6)	1 (0.04-3.1)	0.03
ALC $< 1.2 \times 10^9/l$ ; n (%) ( <i>n evaluable</i> = 143)	54 (38)	41 (33)	13 (68)	<0.01
AMC $\times 10^9/l$ ; median (range)	0.5 (0.04-1.6)	0.5 (0.06-1.6)	0.2 (0.04-0.9)	0.02
Platelets $\times 10^9/l$ ; median (range)	226 (20-599)	247 (25-599)	85 (20-406)	<0.01
Platelets $< 100 \times 10^9/l$ ; n (%)	15 (10)	4 (3)	11 (58)	<0.01
Karyotype ( <i>n evaluable</i> = 143)				

<i>Normal; n (%)</i>	109 (76)	98 (79)	11 (58)	0.08
<i>Sole loss of "Y"; n (%)</i>	7 (5)	5 (4)	2 (11)	
<i>Sole trisomy 8; n (%)</i>	5 (4)	5 (4)	0	
<i>Sole 20q-; n (%)</i>	7 (5)	5 (4)	2 (11)	
<i>Complex; n (%)</i>	5 (4)	2 (2)	3 (16)	
<i>-7/7q-; n (%)</i>	2 (1)	2 (2)	0	
<i>Other; n (%)</i>	8 (6)	7 (6)	1 (5)	
<b>Mutations**</b>				
<i>DNMT3A; n (%)</i>	28 (19)	27 (21)	1 (5)	0.06
<i>TET2; n (%)</i>	36 (25)	28 (22)	8 (42)	0.08
<i>ASXL1; n (%)</i>	16 (11)	15 (12)	1 (5)	0.34
<i>SRSF2; n (%)</i>	7 (5)	2 (2)	5 (26)	<b>&lt;0.01</b>
<i>TP53; n (%)</i>	7 (5)	4 (3)	3 (16)	<b>0.04</b>
<i>RUNX1; n (%)</i>	5 (3)	3 (2)	2 (10)	<b>&lt;0.01</b>
<i>IDH1; n (%)</i>	3 (2)	0	3 (16)	<b>&lt;0.01</b>
<i>IDH2; n (%)</i>	3 (2)	3 (2)	0	0.35
<i>CSF3R; n (%)</i>	3 (2)	3 (2)	0	0.35
<i>CBL; n (%)</i>	3 (2)	3 (2)	0	0.35
<i>ZRSR2; n (%)</i>	3 (2)	2 (2)	1 (5)	0.95
<i>U2AF1; n (%)</i>	2 (1)	0	2 (10)	<b>0.01</b>
<i>EZH2; n (%)</i>	2 (1)	2 (2)	0	0.45
<b>Erythropoietin IU/l; median (range) (n evaluable = 106)</b>	68 (10-5461)	67 (10-5461)	81 (20-380)	0.54
<b>LDH U/l; median (range) (n evaluable = 65)</b>	198 (112-538)	181 (112-441)	225 (128-538)	0.1
<b>Bone marrow ring sideroblasts %; median (range)</b>	35 (5-80)	40 (5-80)	30 (15-70)	0.09
<b>MDS-RS subclassification.</b>				
<b>MDS-RS-MLD; n (%)</b>	72 (50)	59 (47)	13 (68)	
<b>MDS-RS-SLD; n (%)</b>	73 (50)	67 (53)	6 (32)	

<b>Median follow-up for living patients; median years (range)</b>	5.1 (0.12-12.6)	5.2 (0.12-12.6)	4.6 (1.9-8.3)	0.09
<b>Deaths; n (%)</b>	88 (61)	79 (63)	9 (47)	0.2
<b>Leukemic transformation; n (%)</b>	6 (4)	4 (3)	2 (10)	0.2
<b>Allogeneic hematopoietic stem cell transplant; n (%)</b>	5 (4)	4 (3)	1 (5)	0.7

\*Hemoglobin for patients with transfusion independent anemia at time of diagnosis.

\*\*mutations present in at least 2 patients.

Abbreviations: ALC, absolute lymphocyte count; ANC, absolute neutrophil count; AMC, absolute monocyte count; IPSS-R, Revised International Prognostic Scoring System; MDS-RS-MLD, Myelodysplastic syndromes with ring sideroblasts and multilineage dysplasia; MDS-RS-SLD, Myelodysplastic syndromes with ring sideroblasts and single lineage dysplasia