

Outcomes and treatment patterns in primary and secondary plasma cell leukemia: insights from a large US cohort study

Shagun Singh,^{1*} Supriya Peshin,^{2*} Betsy C. Wertheim,³ Ashley Larsen,⁴ Iloabueke Chineke,⁵ Douglas W. Sborov,⁶ Damian Green,⁷ Michaela Liedtke,⁸ Marie Okoniewski,⁹ Mohammed Wazir,¹⁰ Omar Nadeem,¹¹ Levanto G. Schachter,¹² Dominique DeGraaff,¹³ Victoria A. Vardell,¹⁴ David G. Coffey¹⁵ and Krisstina Gowin¹⁶

¹Internal Medicine, Banner University Medical Center, Tucson, AZ; ²Norton Community Hospital, Ballad Health, Norton, VA; ³University of Arizona Cancer Center, Tucson, AZ; ⁴Department of Medicine, University of Arizona, Tucson, AZ; ⁵Rocky Mountain Oncology, Casper, WY; ⁶Huntsman Cancer Institute at the University of Utah, Salt Lake City, UT; ⁷University of Miami, Miller School of Medicine, Miami, FL; ⁸Stanford Medicine, Department of Hematology, Palo Alto, CA; ⁹University of Kansas Cancer Center, Kansas City, KS; ¹⁰UMass Memorial Medical Center, Division of Hematology/Oncology, Worcester, MA; ¹¹Harvard Medical School, Dana-Faber Cancer Institute, Boston, MA; ¹²UCLA Health, Hematology Oncology, San Luis Obispo, CA; ¹³University of Washington, Department of Oncology and Hematology, Seattle, WA; ¹⁴University of Utah, Huntsman Cancer Institute, Salt Lake City, UT; ¹⁵Divison of Myeloma, Sylvester Comprehensive Cancer Center, University of Miami, Miami, FL and ¹⁶Department of Supportive Care Medicine, Hematology/HCT, City of Hope, CA, USA

**SS and SP contributed equally as first authors.*

Correspondence: K. Gowin
kgowin@coh.org

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Supplement Table 1. Patient characteristics, mean \pm SD or n (%)

Characteristic	Total n = 153	Primary PCL n = 96	Secondary PCL n = 57	P
Type of transplant(s) at induction				< 0.001
Tandem auto/auto	1 (0.8)	1 (1.1)	0 (0.0)	
Tandem auto/allo	11 (8.3)	10 (11.1)	1 (2.4)	
Single auto	51 (38.6)	45 (50.0)	6 (14.3)	
Single allo	5 (3.8)	4 (4.4)	1 (2.4)	
No transplant at induction	64 (48.5)	30 (33.3)	34 (81.0)	
Type of transplant(s) at relapse				0.613
Tandem auto/auto	1 (0.8)	1 (1.1)	0 (0.0)	
Tandem auto/allo	1 (0.8)	1 (1.1)	0 (0.0)	
Single auto	4 (3.1)	2 (2.2)	2 (5.0)	
Single allo	4 (3.1)	4 (4.4)	0 (0.0)	
No transplant at relapse	120 (92.3)	82 (91.1)	38 (95.0)	
CCND1/IgH t(11;14) at dx				0.323
Absent at diagnosis	39 (49.4)	26 (45.6)	13 (59.1)	
Present at diagnosis	40 (50.6)	31 (54.4)	9 (40.9)	
<i>n missing</i>	74	39	35	
CCND1/IgH t(11;14) ever				1.000
Never present	50 (50.0)	35 (49.3)	15 (51.7)	
Ever present	50 (50.0)	36 (50.7)	14 (48.3)	
<i>n missing</i>	53	25	28	

Missing data for each characteristic (total): Type of transplant(s) at induction (21), Type of transplant(s) at relapse (23), t(11;14) at dx (74), t(11;14) ever (53)

Missing data for each characteristic (primary PCL): Type of transplant(s) at induction (6), Type of transplant(s) at relapse (6), t(11;14) at dx (39), t(11;14) ever (25)

Missing data for each characteristic (secondary PCL): Type of transplant(s) at induction (15), Type of transplant(s) at relapse (17), t(11;14) at dx (35), t(11;14) ever (28)

Supplement Table 2. Associations with survival (Total, n = 153)

Characteristic	Age-adjusted HR (95% CI)	Multivariate ¹ HR (95% CI)	p-value
Secondary PCL	4.54 (3.01–6.83)	2.37 (1.47–3.84)	< 0.001
Any transplant	0.12 (0.06–0.21)	0.16 (0.08–0.29)	< 0.001
High-risk genetic feature	1.28 (0.76–2.14)	1.10 (0.63–1.92)	0.749
DEL17p	1.56 (0.90–2.71)	1.31 (0.72–2.40)	0.376
t(4;14)	1.21 (0.64–2.30)	0.92 (0.45–1.88)	0.818
t(14;16)	0.63 (0.28–1.39)	0.89 (0.38–2.08)	0.790
Extramedullary disease	1.57 (0.99–2.48)	0.97 (0.57–1.66)	0.919
CCND1/IgH t(11;14) at dx	0.63 (0.35–1.11)	0.65 (0.34–1.26)	0.203
CCND1/IgH t(11;14) ever	0.80 (0.49–1.32)	0.79 (0.46–1.36)	0.393

¹ Further adjusted for PCL type and any transplant

Supplement Table 3. Patient characteristics, by transplant status, mean ± SD or n (%)

Characteristic	Never transplant n = 96	Ever transplant n = 57	P
Age at diagnosis	64.8 ± 10.0	56.4 ± 9.0	< 0.001
Lab measures at diagnosis			
LDH (U/L)	501.5 ± 675.6	347.4 ± 238.9	0.509
Albumin (g/dL)	3.2 ± 0.7	3.4 ± 0.8	0.224
Creatinine (mg/dL)	1.8 ± 1.7	5.1 ± 19.8	0.733
Beta-2 microglobulin (?)	11.5 ± 14.8	10.1 ± 10.2	0.837
Secondary PCL			< 0.001
Primary	28 (46.7)	65 (86.7)	
Secondary	32 (53.3)	10 (13.3)	

Supplement Table 3. Patient characteristics, by transplant status, mean ± SD or n (%)

Characteristic	Never transplant n = 96	Ever transplant n = 57	P
High-risk genetic feature			0.500
No	12 (36.4)	24 (46.2)	
Yes	21 (63.6)	28 (53.8)	
Extramedullary disease			0.505
No	32 (71.1)	54 (78.3)	
Yes	13 (28.9)	15 (21.7)	
First line of chemotherapy			0.111
Proteasome inhibitor triplet	16 (45.7)	31 (70.5)	
VTD PACE-like regimen	10 (28.6)	8 (18.2)	
Daratumamab quadruplet	1 (2.9)	1 (2.3)	
Other	8 (22.9)	4 (9.1)	
Best response to first line of chemo			0.041
Less than CR	19 (61.3)	10 (32.3)	
CR or better	12 (38.7)	21 (67.7)	
MRD Negative (10^{-6}) (ever)			< 0.001
Yes	1 (2.5)	9 (16.1)	
No	18 (45.0)	3 (5.4)	
Unknown	21 (52.5)	44 (78.6)	
CCND1/IgH t(11;14) at dx			1.000
Absent at diagnosis	13 (50.0)	22 (48.9)	
Present at diagnosis	13 (50.0)	23 (51.1)	
CCND1/IgH t(11;14) ever			1.000
Never present	16 (51.6)	30 (51.7)	
Ever present	15 (48.4)	28 (48.3)	

Note: P-value for comparing never versus ever transplant from Wilcoxon rank-sum test (continuous variables) or Fisher's exact test (categorical variables).

High-risk genetic features, as defined by the revised 2016 IMWG consensus, include the presence of t(4;14), t(14;16), Del17p13, a non-hyper diploid karyotype and gain (1q)

Missing data for never transplant: LDH (26), Albumin (15), Creatinine (18), beta-2 microglobulin (19), High-risk genetic feature (27), Extra medullary disease (15), First line chemotherapy (25), Best response to first line of chemo (29), MRD response (20), t(11;14) at dx (34), t(11;14) ever (29)

Missing data for ever transplant: LDH (43), Albumin (32), Creatinine (36), beta-2 microglobulin (34), High-risk genetic feature (23), Extra medullary disease (06), First line chemotherapy (31), Best response to first line of chemo (44), MRD response (19), t(11;14) at dx (30), t(11;14) ever (17)