

**Supplemental Data Table S6.** Comparison of the frequencies of mtDNA haplogroups in healthy controls and pediatric AML patients

Haplogroup	Frequency (N, %)						<i>P</i>	Haplogroup	Frequency (N, %)						<i>P</i>
	Healthy controls				Pediatric AML patients				Healthy controls				Pediatric AML patients		
	Korean <sup>†</sup>		Korean <sup>‡</sup>						Korean <sup>†</sup>		Korean <sup>‡</sup>				
A	4	5.7	3	5.3			F	3	4.3						
A4					1	1.8	F1a	1	1.4	2	3.5				
A5	2	2.9					F1ac					2	3.6		
A5a					2	3.6	F1b	2	2.9	1	1.8				
A5b					1	1.8	F1b1					1	1.8		
Subtotal	6	8.6	3	5.3	4	7.3	0.964	F1c			1	1.8			
B	4	5.7					F3a	1	1.4						
B4	2	2.9	3	5.3	2	3.6	0.923	F4a				1	1.8		
B4a	4	5.7			2	3.6	Subtotal	7	10.0	4	7.0	4	7.3	0.754	
B4b1			1	1.8	1	1.8	G1a	4	5.7						
B4c1a					2	3.6	G1a1					1	1.8		
B4c1c					1	1.8	G2a	2	2.9	2	3.5	1	1.8	0.614	
B4d					1	1.8	G3	1	1.4						
B5a	1	1.4	2	3.5			G4a			2	3.5				
B5a2					1	1.8	G4b			1	1.8				
B5b	1	1.4	2	3.5	2	3.6	0.629	Subtotal	7	10.0	5	8.8	2	3.6	0.177
Subtotal	12	17.1	8	14.0	12	21.8	0.323	M	1	1.4			1	1.8	
D	2	2.9					M7a1			2	3.5	2	3.6		
D4*	11	15.7	16	28.1	3	5.5	0.008	M7b2	1	1.4	3	5.3	2	3.6	0.866
D4a	2	2.9			3	5.5		M7c1	1	1.4	3	5.3	1	1.8	0.614
D4b1	1	1.4					M8a			3	5.3	1	1.8		
D4b2b					1	1.8	M8a2					1	1.8		
D4e1					1	1.8	M9a			1	1.8				
D4g1					1	1.8	M10	4	5.7	5	8.8				
D4j1					1	1.8	Subtotal	7	10.0	17	29.8	8	14.5	0.479	
D4n					1	1.8	N9a	3	4.3	1	1.8				
D5	4	5.7	2	3.5	2	3.6	0.742	N9a1	1	1.4			3	5.5	
D5a	2	2.9					N9a2	3	4.3						
D5a2					1	1.8	Subtotal	7	10.0	1	1.8	3	5.5	0.826	
D5b	1	1.4			2	3.6	Y			1	1.8				
D5c					2	3.6	Y1	1	1.4						
Subtotal	23	32.9	18	31.6	18	32.7	0.953	Y1b					3	5.5	
							Subtotal	1	1.4	1	1.8	3	5.5	0.141	
							Z					1	1.8		
							Subtotal					1	1.8		
							Total	70	100.0	57	100.0	55	100.0		

\*Odds ratio for risk of pediatric AML development=0.213675 (95% CI; 0.061898-0.73762).  
Healthy controls of reference 22<sup>†</sup> and reference 23<sup>‡</sup>.