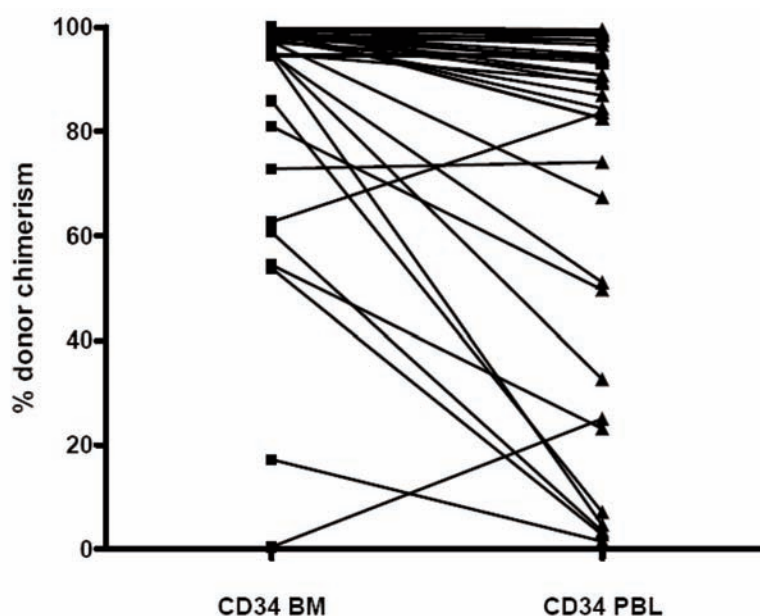


Monitoring of donor chimerism in sorted CD34⁺ peripheral blood cells allows the sensitive detection of imminent relapse after allogeneic stem cell transplantation

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Online Supplementary Figure 1. Paired analysis of donor chimerism in CD34⁺ peripheral blood and CD34⁺ BM cells. The graph shows results of donor chimerism analysis in CD34 sorted peripheral blood (PBL, filled triangles) vs. bone marrow (BM, filled squares) samples taken from the same patient on the same day. Both PBL and BM values of each patient are connected by a line.

Online Supplementary Table 1. Patient characteristics.

Characteristic		N (%)
Disease		
-AML		67 (74)
-ALL		16 (18)
-MDS		7 (8)
Cytogenetics/ Molecular marker		
- Normal karyotype		37 (41)
- Numeric aberration/Deletion		18 (20)
- Translocation/Inversion		22 (24)
- Inversion 16		6 (7)
- Bcr/abl (ALL)		5 (6)
- unknown		13 (15)
Median age (range)		
Sex		
-Male		49 (54)
-Female		41 (46)
Donor		
-Sibling		35 (39)
-Unrelated		55 (61)
HLA match		
-Matched*		76 (85)
-Mismatch		12 (13)
-Haploidentical		2 (2)
Risk category		
-Low risk+		42 (47)
-High Risk#		48 (53)
Graft source		
-Peripheral blood		82 (91)
-Bone marrow		8 (9)
Conditioning therapy		
-Intensive		50 (56)
-Reduced intensity		40 (44)

AML: acute myeloid leukemia; ALL: acute lymphoblastic leukemia; MDS: myelodysplastic syndrome; *, A, B, C, DRB1 and DQB1 allele matched; +, AML/ALL in 1st or 2nd complete remission and MDS with \leq 5% blasts; # AML/ALL > 2nd CR or not in remission and MDS > 5% blasts.

Online Supplementary Table 2. Cox regression models.

Outcome	Risk factor	Hazard ratio	95% CI	p-value
Probability of relapse	Risk group (High vs. low)	2.4	1.1-5.3	0.03
	CD34+ chimerism < 80%	3.5	1.5-7.8	0.003
Overall survival	Risk group (High vs. low)	2.4	1.2-4.6	0.01
	CD34+ chimerism < 80%	1.9	1.0-3.5	0.05
Disease-free survival	Risk group (High vs. low)	2.4	1.3-4.5	0.06
	CD34+ Chimerism < 80%	2.4	1.3-4.6	0.08

AML: acute myeloid leukemia; ALL: acute lymphoblastic leukemia; MDS: myelodysplastic syndrome; *, A, B, C, DRB1 and DQB1 allele matched; +, AML/ALL in 1st or 2nd complete remission and MDS with \leq 5% blasts; # AML/ALL > 2nd CR or not in remission and MDS > 5% blasts.