			% of Peripheral Blood			
Primary	% CD45.2+	% CD45.2+	B Cells	T Cells	Myeloid	
Transplant		Granulocytes				
Total	20.97±5.16	35.24±7.96	9.74±3.09	3.34±1.90	6.96±1.55	
GFP ^{Hi}	30.25±6.04	64.60±6.27	14.01±3.50	2.66±1.47	12.50±1.77	
GFP ^{Lo}	10.07±3.16	22.56±6.57	5.85±2.38	0.91±0.38	2.96±0.95	
Secondary						
Transplant						
Total	14.55±2.87	20.54±7.98	5.97±1.33	4.06±1.29	4.28±1.65	
GFP ^{Hi}	12.03±3.95	29.70±7.58	11.61±5.79	1.13±0.46	4.04±1.11	
GFP ^{Lo}	1.30±0.65	1.40±0.49	0.80±0.41	0.26±0.21	0.20±0.06	

Table S1 – Related to Figure 1. Donor cell contribution to peripheral blood. White blood cell and granulocyte contributions, as well as donor derived lineage contributions to total peripheral blood at the end of primary (22 weeks) or secondary (24 weeks) transplantation. Data are represented as mean \pm SEM of 8-14 mice per group from two independent experiments.

	Mice Reconstituted	Repopulating Cell Frequency (1/x)	Repopulating Cell Range (95% CI)	Cell #/ Mouse	Cell # Range (95% CI)	CD45.2 ⁺ Chimerism (% ± SEM)
GFP ^L ° CD41 ⁺	6/12	21.6	9.57 <x<49.0< th=""><th>0.69</th><th>1.57>x>0.31</th><th>1.12±0.67%</th></x<49.0<>	0.69	1.57>x>0.31	1.12±0.67%
GFP ^{Lo} CD41 ⁻	7/11	14.8	6.85 <x<32.1< th=""><th>1.01</th><th>2.19>x>0.47</th><th>13.09±11.73%</th></x<32.1<>	1.01	2.19>x>0.47	13.09±11.73%
GFP ^{Hi} CD41 ⁺	4/6	13.7	4.87 <x<38.2< th=""><th>1.09</th><th>3.08>x>0.39</th><th>37.59±19.55%</th></x<38.2<>	1.09	3.08>x>0.39	37.59±19.55%
GFP ^{Hi} CD41 ⁻	7/9	9.97	4.42 <x<22.5< th=""><th>1.50</th><th>3.39>x>0.67</th><th>27.70±6.62%</th></x<22.5<>	1.50	3.39>x>0.67	27.70±6.62%

Table S2 – Related to Figure 3. Limiting dilution analysis of aging HSC populations. CD45.2* chimerism was determined at 24 weeks in primary transplantation, and averaged between all mice considered repopulated regardless of retrospectively identified repopulating cell type.

Table S3: Antibody Panel Related to STAR Methods

Peripheral Blood	CLPs 17A2 RA3-6B2 M1/70 (After Transplantation) (Biotin) (Biotin) (Biotin)	Myeloid Progenitors 17A2 RA3-6B2 M1/70 (After Transplantation) (Biotin) (Biotin) (Biotin)	HSPC Analysis 17A2 RA3-6B2 M1/70 (After Transplantation) (Biotin) (Biotin) (Biotin)	HSC panel 17A2 RA3-6B2 M1/70 (Cell Cycle) (Biotin) (Biotin) (Biotin)	HSC panel 17A2 RA3-6B2 M1/70 (LRC Analysis) (Biotin) (Biotin) (Biotin)	CD3s B220 CD11b	Lineage Cocktai
	RB6-8C5 TER (Biotin) (Bi	RB6-8C5 TER (Biotin) (Bi	RB6-8C5 TER (Biotin) (Bi	RB6-8C5 TER (Biotin) (Bi	RB6-8C5 TER (Biotin) (Bi	Gr1 Te	Cocktail
	TER-119 APC/Cy7	TER-119 APC/Cy7	TER-119 APC/Cy7	TER-119 PE/Cy5 or (Biotin) APC	TER-119 PE/Cy5 or (Biotin) eFluor450	Ter119 Streptavidin	
	7 2B8 (APC)	7 2B8 (APC)	7 2B8 (PE)		or 2B8 50 (PE or APC)	din c-Kit	
	D7 (PacificBlue)	D7 (PacificBlue)	D7 HM48-1 (PerCP/Cy5.5	2B8 (FITC or D7 (APC/Cy7 PerCP/Cy5.5) or PacificBlue)	D7 (APC/Cy7 or PacificBlue)	Sca-1	
			HM48-1 (PerCP/Cy5.5)	HM48-1 (Biotin)	HM48-1 (Biotin)	CD48	
	A2F10 (PE)			A2F10 (Biotin)	A2F10 (Biotin)	FK2	Stem/F
			TC15-12F12.2 (PE/Cy7)	TC15-12F12.2 (PE/Cy7)	TC15-12F12.2 (PE/Cy7)	CD150	Prog
			eBioMWReg31 (APC)	eBioMWReg31 (APC)	eBioMWReg31 (APC)	CD41	
	A7R34 (PE/Cy7)					IL-7Rα	
		93 (PE/Cy7)				FcyRIII	
		RAM34 (AF700)				CD34	
				SolA15 (PE)		Ki67	Cel
	0		0	DAPI		DNA Content	Cell Cycle
104 (PE) (F	104 (AF700)	104 (PE)	104 (AF700)			CD45.2	
GK1.5 (PE/Cy5)						CD4	
53-6.7 (PE/Cy5)						CD8a	Periphe
RA3-6B2 (PE/Cy7)						B220	Peripheral Blood
53-6.7 RA3-6B2 M1/70 RB6-8C5 (PE/Cy5) (PE/Cy7) (APC/Cy7) (APC/Cy7)						CD11b	
RB6-8C5 (APC/Cy7)						Gr1	
DAPI	DAPI	DAPI	DAPI		DAPI or PI	Viability	

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