

40% 20% 0% 10,000 2,000 400 100 Granulocytes CD4T CD8T other B cells







104

10³

Sample

10⁶ Total Reads 10⁴ 10⁵ Water Sample









Figure S1. White Blood Cell Population Size at Various HSC Transplantation Doses. *Related to Figure 1*.

(A–C) White blood cell population sizes were measured (A) one, (B) two, and (C) six months after transplantation by flow cytometry. Bar segments represent group means. Error bars represent standard error of the mean. 7-8 mice were used per transplantation dose.

Figure S2. Replicate Analysis of barcode quantification. *Related to Figure 1*.

Blood samples were lysed and split into replicate samples. Replicates were separately extracted for genomic DNA, amplified by PCR, and quantified using high throughput sequencing. Each data point represents a single barcode. The x and y axes represent the abundance of a barcode in each replicate sample.

Figure S3. Removal of Sequencing Background. Related to Figure 1.

Molecular grade water was processed via PCR and sent for sequencing. False positive barcodes detected in the water were quantified. Barcodes from actual samples were only considered valid if they exceeded 99% of the signals from water samples. Each graph shows the barcodes of a mouse sample compared to the water sample. Each blue dot represents a barcode. The red line represents the 99% threshold.

Figure S4. Chimerism over Time at Different Transplantation Doses. *Related to Figure 1*.

(A–D) Donor, helper, and host chimerisms in (A) granulocytes, (B) B cells, (C) CD4T cells, and (D) CD8T cells were measured monthly by flow cytometry. Data points represent group means. Error bars depict standard error of the mean. 7-8 mice were used per transplantation dose.

Figure S5. Early Granulocyte Production Changes with HSC Dose. Related to Figure 5.

Pie charts show the percent of granulocyte production by each category of clones one and two months after transplantation. Values are the mean percentage for each HSC dose group. See also Tables S6-S7. 7-8 mice were used per transplantation dose.

Figure S6. Alternate Methods of Computing Clonal Dominance Produce Similar Results. *Related to Figure 6*.

The number of clones that produced more than 0.1% of all granulocytes at each HSC transplantation dose. Each diamond represents one mouse; horizontal black lines represent group means.

Figure S7. Lineage Bias Analysis of Granulocytes vs. T cells. Related to Figure 7.

Granulocyte production was compared to T-cell production for each HSC clone. A clone was considered to be "specialized" to a cell type if it only produced that cell type, "biased" to a cell type if it favored the production of that cell type, and "balanced" if its relative production of the two cell types was similar. The mean percent is displayed for each HSC dose. Error bars represent the standard error of the mean. 7-8 mice were used per transplantation dose.

	1 mc	onth			2 mor	nths			6 months			
	10000	2000	400		10000	2000	400		10000	2000	400	
2000	***			2000	***			2000	***			
400	***	NS		400	***	**		400	***	*		
100	***	**	NS	100	***	***	NS	100	***	**	NS	

NS, not significant

Table \$	S2
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	Early-diffe	erentiatin	g		9	Early	only
	10000	2000	400		400	10000	2000
2000	***			2000		***	
400	***	*		400		***	*
100	***	***	NS	100	NS	***	**
		ltunon				ato ond o	
			400		400		
2000	10000 ***	2000	400	2000	400	*	2000
400	***	NS		400		***	NS
100	***	*	NS	100	NS	***	NS
	arly and a	ill cell typ	es		es	Speci	alized
	10000	2000	400		400	10000	2000
2000	***			2000		* * *	
400	***	*		400		* * *	NS
100	***	**	NS	100	NS	***	**
E	Early and s	specialize	ed		ed	Late and s	
	10000	2000	400		400	10000	2000
2000	NS	2000	.00	2000		**	2000
400	**	NS		400		*	NS
100	***	*	NS	100	NS	***	NS

NS, not significant

	Early-diffe	erentiatin	g
	10000	2000	400
2000	NS		
400	NS	NS	
100	**	**	NS
	A II II		
	All Cel	types	
	10000	2000	400
2000	NS		
400	NS	NS	
100	NS	NS	NS
C			400
	10000	2000	400
2000	NS		
400	NS	NS	
100	NS	NS	NS
E	Early and s	specialize	ed
	10000	2000	400
2000	NS		
400	NS	NS	
100	NS	NS	NS

NS, not significant

p<0.01 *p<0.001

Table S3

Gr 10,000				Gr 2,000			Gr 400					Gr 100			
	L all	L Sp	E all		L all	L Sp	E all		L all	L Sp	E all		L all	L Sp	E all
L Sp	NS	-		L Sp	NS	-		L Sp	NS	-		L Sp	NS	-	
E all	***	***		Eall	**	**		Eall	**	*		Eall	***	***	
E Sp	NS	NS	NS	E Sp	NS	NS	**	E Sp	NS	NS	NS	E Sp	NS	NS	***
	B 10,000B 2,000														
		1,000													
	Lall	L Sp	Eall		Lai	L Sp	Eall		Lall	L Sp	Eall		Lan	L Sp	E all
L Sp	NS			L Sp	NS			L Sp	×			L Sp	NS		
E all	***	***		E all	***	***		E all	*	NS		E all	**	***	
E Sp	**	**	***	E Sp	*	*	***	E Sp	NS	NS	NS	E Sp	NS	NS	***
	47.4	0.000			47.0	000			47.	100			47	100	
	41 10	0,000			<u>412</u>	,000			414	+00			41	100	
	L all	L Sp	E all		L all	L Sp	E all		L all	L Sp	E all		L all	L Sp	E all
L Sp	NS			L Sp	NS			L Sp	NS			L Sp	NS		
E all	***	***		E all	***	***		E all	***	***		E all	***	***	
E Sp	**	**	***	E Sp	**	**	***	E Sp	*	*	*	E Sp	NS	NS	***
	8T 1	0 0 0 0			8T 2	000			8T 4	100			87	100	
	<u>ו וט</u>	l Sn	Eall		<u>ייי</u> ו או	,000 I Sn	E all			1 Sn	E all				
		LOP				LOP				LOP				LOP	
L SP	112			L Sp	112			L Sp	IN2			L SD	IN S		
E all	***	***		E all	***	***		E all	***	***		⊢ all	***	***	
E Sp	***	***	***	E Sp	**	**	***	E Sp	**	*	NS	E Sp	NS	NS	***

L all, Late and all cell types; L Sp, Late and specialized; E all, Early and all cell types; E Sp, Early and specialized; NS, not significant

Gr	- Late	te and all Gr - Late and S					5p	Gr -	- Early	and	all	Gr – Early and Sp				
	10000	2000	400		10000	2000	400		10000	2000	400		10000	2000	400	
2000	NS			2000	NS			2000	NS			2000	NS			
400	*	NS		400	NS	NS		400	NS	NS		400	NS	NS		
100	*	NS	NS	100	NS	NS	NS	100	NS	NS	NS	100	NS	NS	NS	
B ce	ell - La	te and	d all	B ce	ell - Lat	e and	Sp	B cel	I - Ear	ly and	d all	B ce	ll - Ear	ly and	Sp	
	10000	2000	400		10000	2000	400		10000	2000	400		10000	2000	400	
2000	NS			2000	NS			2000	NS			2000	NS			
400	**	NS		400	*	**		400	NS	*		400	NS	NS		
100	**	NS	NS	100	NS	NS	NS	100	NS	NS	*	100	**	NS	NS	
AT cell - Late and all AT cell - Late and Sn					d Sp	4T ce	II - Ea	rlv an	d all	4T ce	II – Ea	rlv an	d Sp			
														,	1-	
	10000	2000	400		10000	2000	400		10000	2000	400		10000	2000	400	
2000	**			2000	NS			2000	NS			2000	NS			
400	***	NS		400	NS	NS		400	NS	NS		400	NS	NS		
100	***	NS	NS	100	NS	NS	NS	100	NS	NS	NS	100	NS	NS	NS	
8T ce	- - e	ate an	d all	8T c	ell - La	te and	l Sn	8T ce	8T cell Early and all			8T ce	II – Fa	rlv an	d Sn	
	40000															
	10000	2000	400		10000	2000	400		10000	2000	400		10000	2000	400	
2000	NS			2000	NS			2000	NS			2000	NS			
400	***	NS		400	NS	NS		400	NS	NS		400	NS	NS		
100	**	NS	NS	100	*	NS	NS	100	NS	NS	NS	100	NS	NS	NS	

Late and all, Late and all cell types; Late and Sp, Late and specialized; Early and all, Early and all cell types; Early and Sp, Early and specialized; NS, not significant

1 n	nonth all	cell typ	bes		1 mont	h Sp	1	1 month early only				
	10000	2000	400		10000	2000	400		10000	2000	400	
2000	NS			2000	NS			2000	NS			
400	NS	NS		400	NS	NS		400	*	NS		
100	**	*	NS	100	NS	NS	NS	100	***	*	*	
2 n	nonth all	cell typ	bes		2 mont	h Sp		2	2 month early only			
	10000	2000	400		10000	2000	400		10000	2000	400	
2000	NS			2000	NS			2000	NS			
400	NS	NS		400	NS	NS		400	*	NS		
100	**	NS	NS	100	NS	NS	NS	100	**	NS	NS	

Sp, Specialized; NS, not significant

1 month 10,000			1 month 2,000			1 n	nonth 4	400	1 n	1 month 100			
	all	Sp		all	Sp		all	Sp		all	Sp		
Sp	***		Sp	***		Sp	*		Sp	NS			
e.o.	***	NS	e.o.	*	*	e.o.	NS	NS	e.o.	NS	NS		
2 mo	nths 1	10,000	2 months 2,000			2 months 400			2 m	2 months 100			
	all	Sp		all	Sp		all	Sp		all	Sp		
Sp	**		Sp	NS		Sp	NS		Sp	**			
e.o.	NS	***	e.o.	**	*	e.o.	*	NS	e.o.	***	NS		

all, all cell types; Sp, specialized; e.o., Early only; NS, not significant

Table S1. Significance Values for the Line Graph Shown in Figure 2A. Related to Figure 2.

Significance of the comparisons between HSC dose groups for the number of granulocyte-producing clones detected at each time point. NS, not significant.

Table S2. Significance Values for the Number of Clones in Each Category of the Venn Diagrams Shown in Figure 4. Related to Figure 4.

Significance of the comparisons between HSC dose groups. NS, not significant.

Table S3. Significance Values for the Proportion of Clones in Each Category of the Venn DiagramsShown in Figure 4. *Related to Figure 4*.

Significance of the comparisons between HSC dose groups. NS, not significant.

Table S4. Significance Values for the Comparisons of Blood Production by Each Category of the PieCharts Shown in Figure 5. *Related to Figure 5*.

Significance of the comparisons for each cell type and HSC dose. L all, late and all cell types; L Sp, late and specialized; E all, early and all cell types; E Sp, early and specialized; NS, not significant.

Table S5. Significance Values for the Comparisons of Blood Production at Each Transplantation Dose of the Pie Charts Shown in Figure 5. *Related to Figure 5*.

Significance of the comparisons for each cell type and category of clonal blood production. Late and All, late and all cell types; Late and Sp, late and specialized; Early and All, early and all cell types; Early and Sp, early and specialized; NS, not significant.

Table S6. Significance Values for the Comparisons of Granulocyte Production at Each HSCTransplantation Dose Group of the Pie Charts Shown in Figure S5. *Related to Figure 5*.

Significance of the comparisons for each category of clone and time point. Sp, Early specialized; NS, not significant.

Table S7. Significance Values for the Comparisons of Granulocyte Production by Each Category of the Pie Charts Shown in Figure S5. *Related to Figure 5*.

Significance of the comparisons for each time point and HSC transplantation dose. Sp, Early specialized; all, Early all cell types; e.o., early only; NS, not significant.