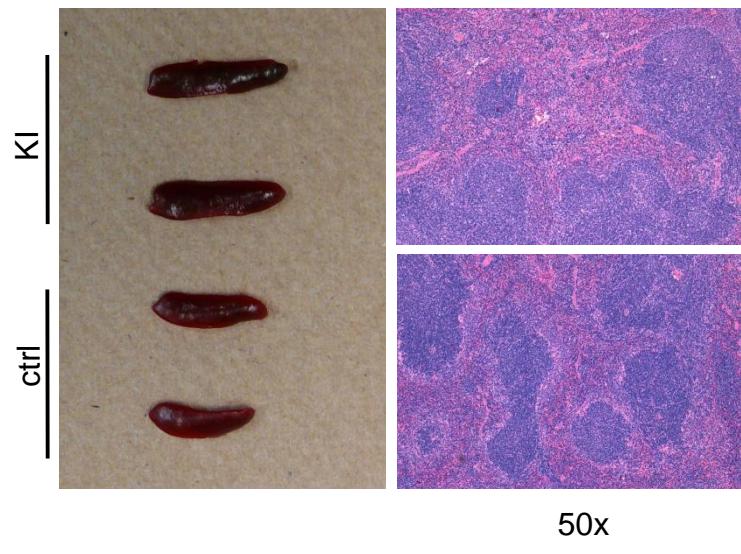
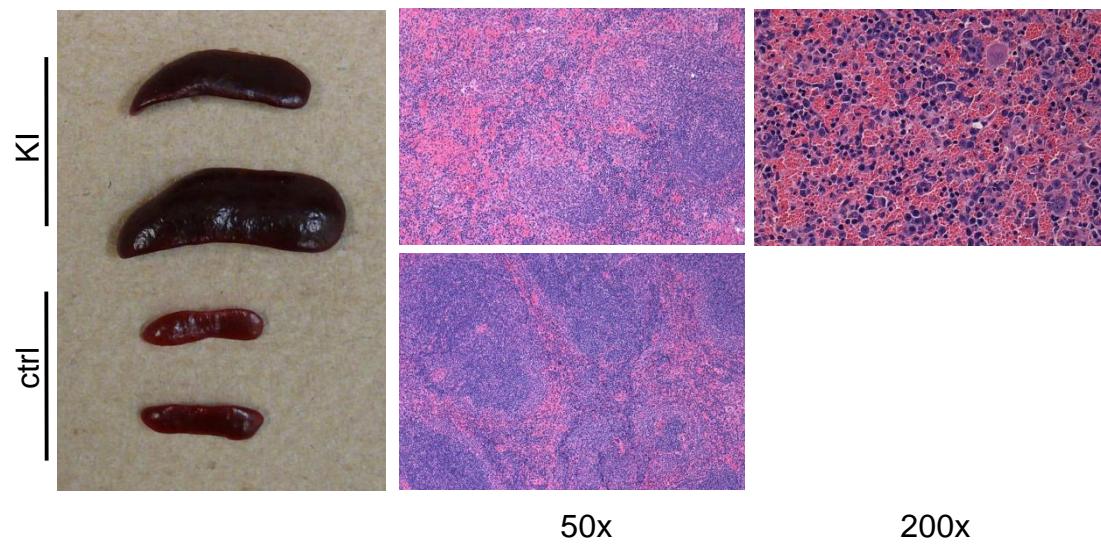


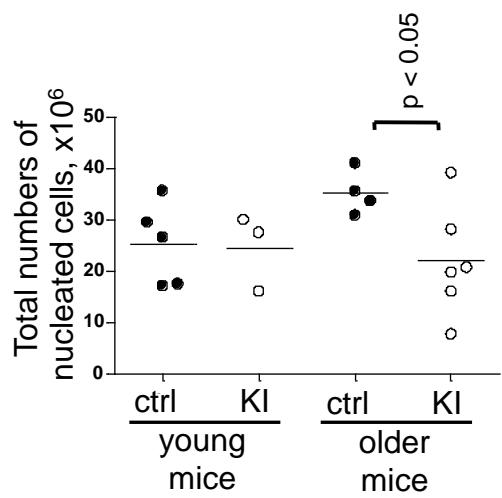
a young mice



b older mice



c bone marrow



d bone marrow

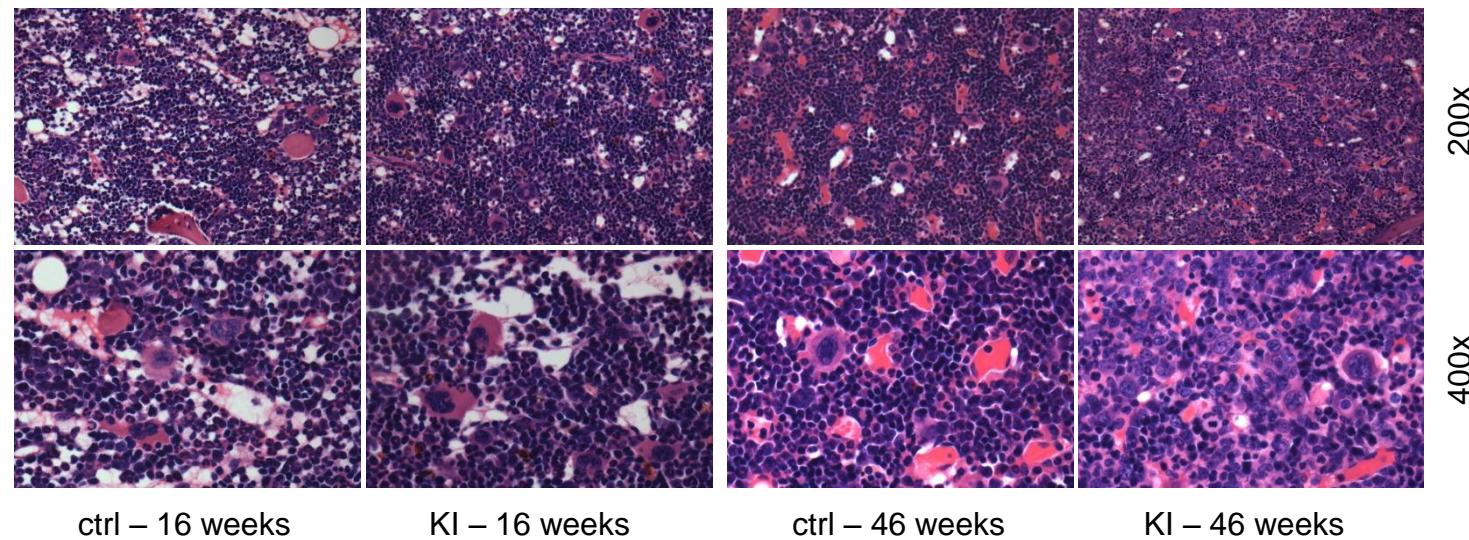
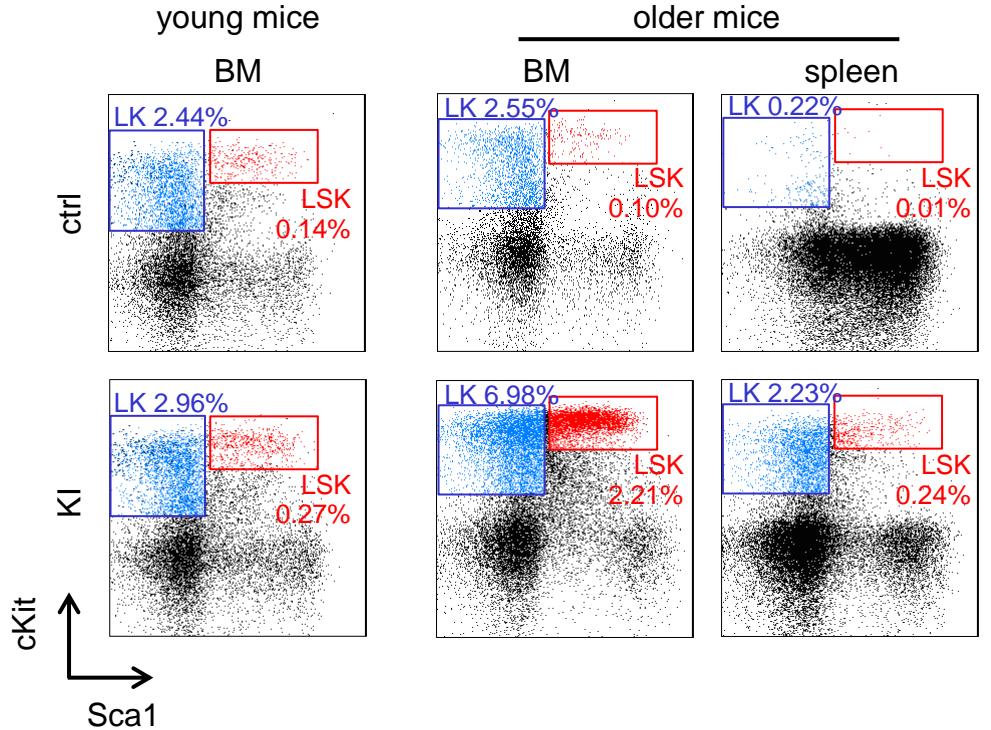
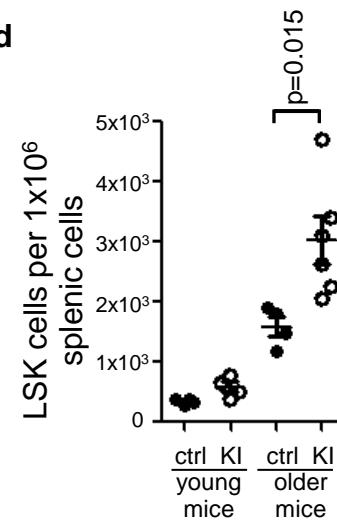
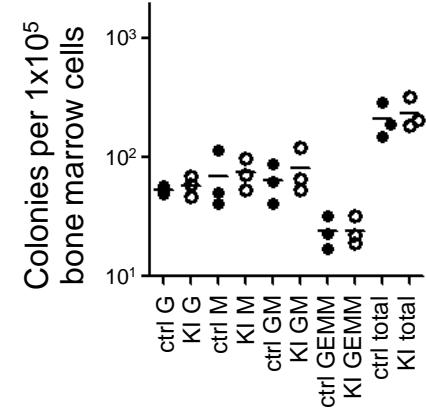
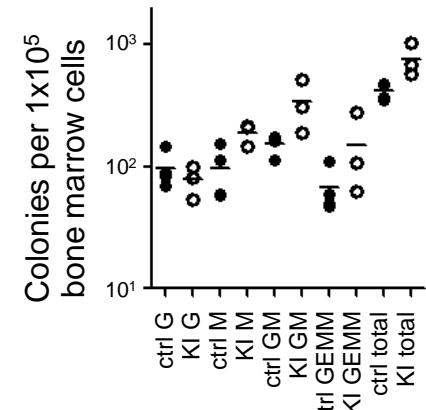
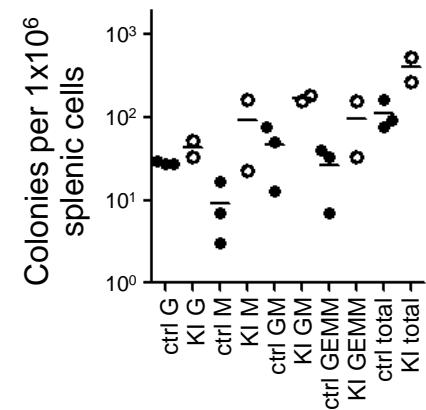
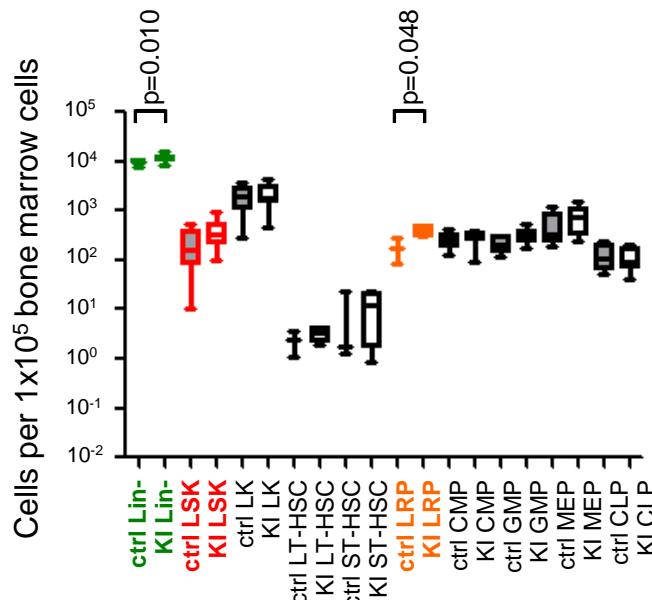
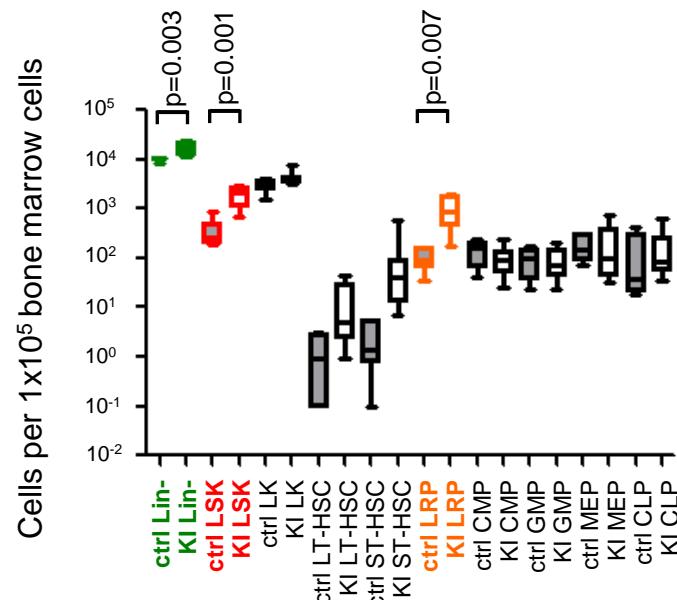


Figure 1

a

Gated on live lineage-negative cells

**d****e****f****g****b** young mice**c** older mice**Figure 2**

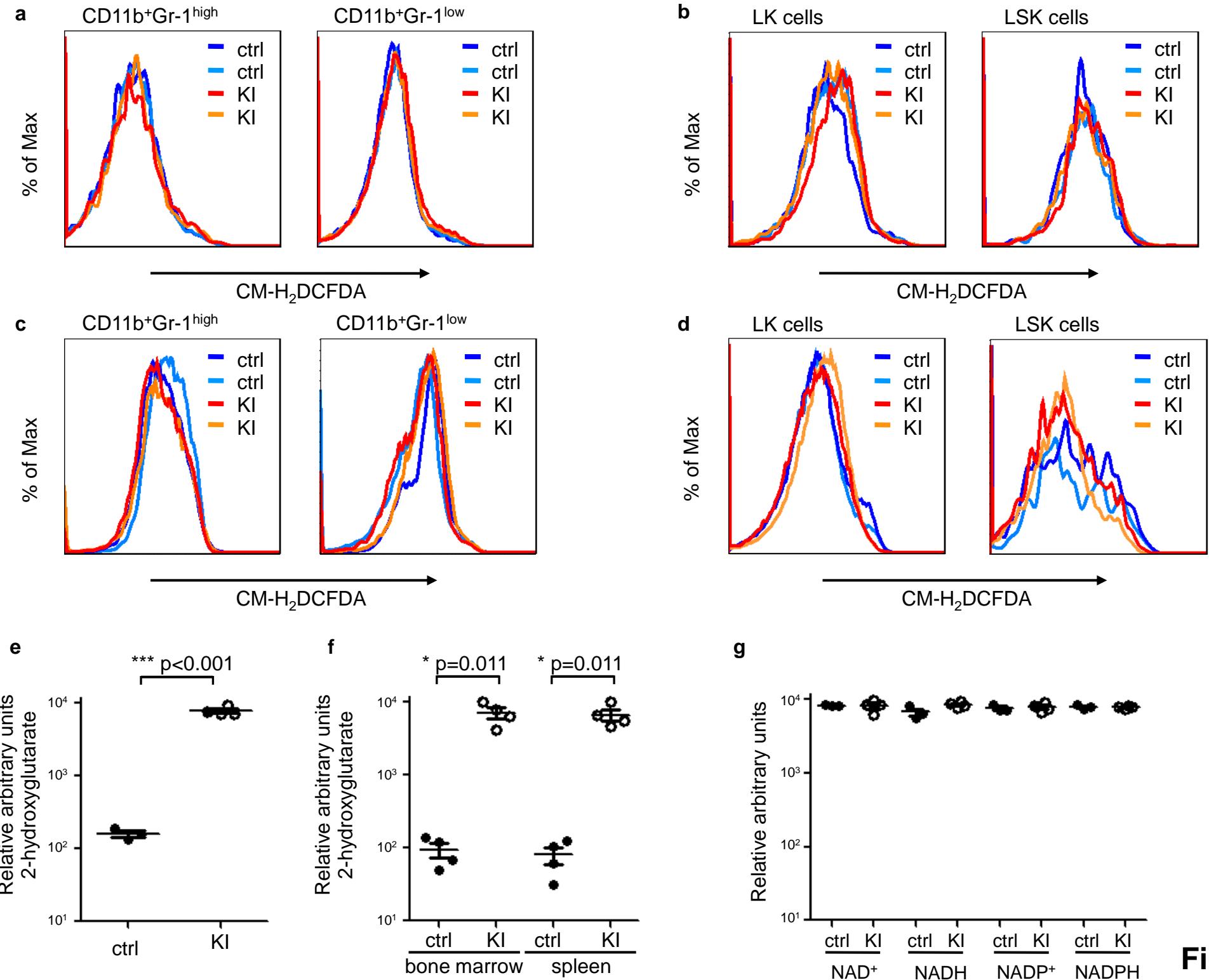
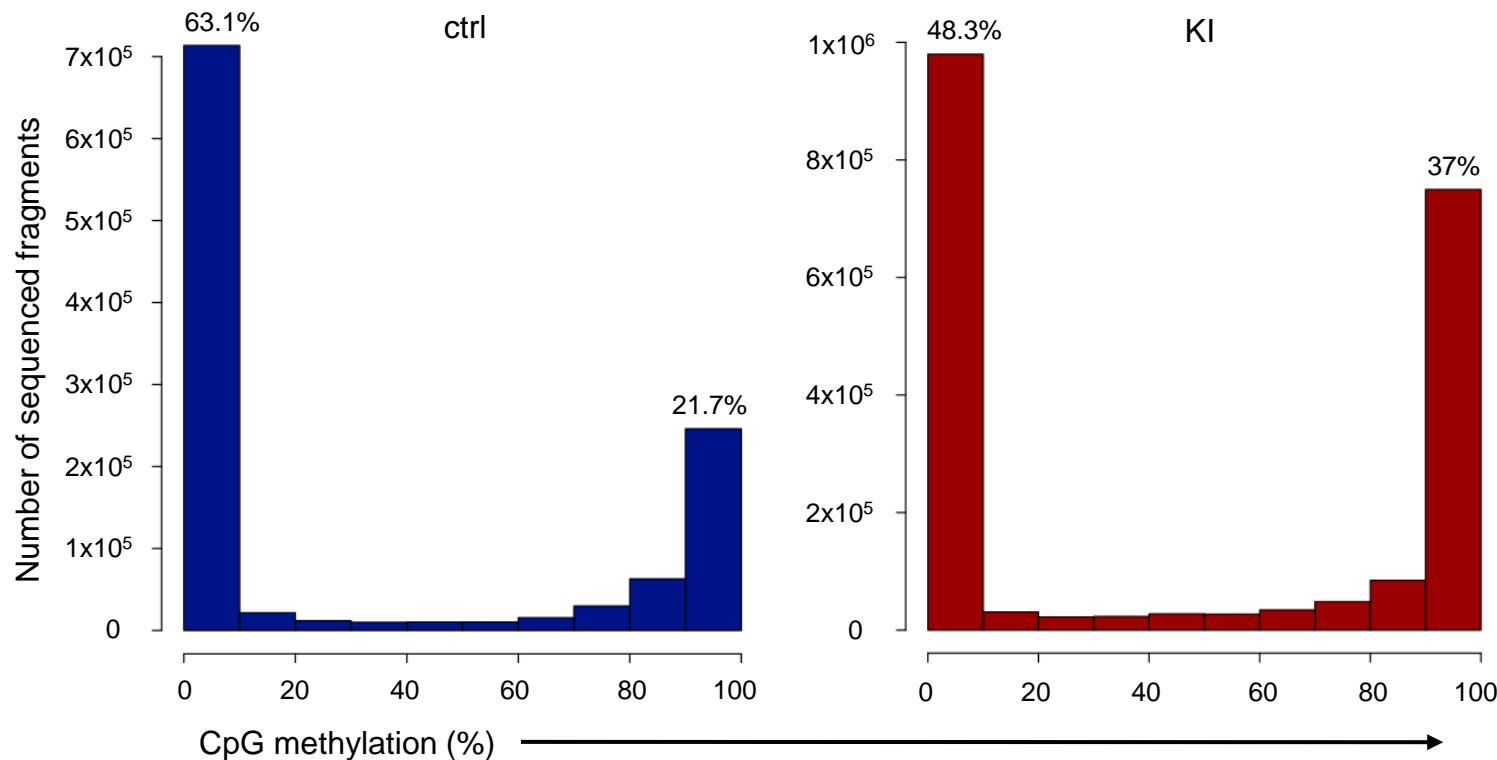
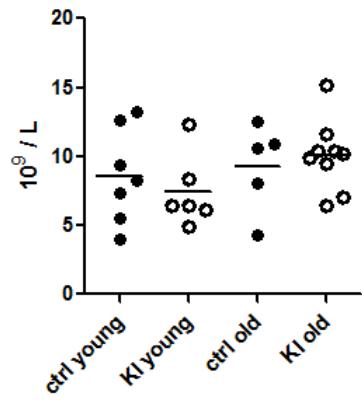
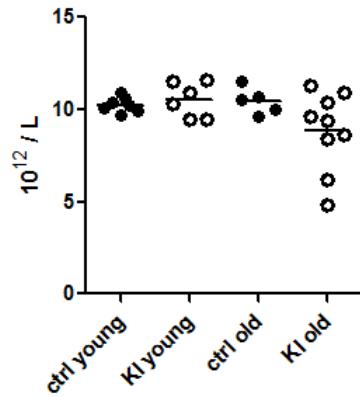
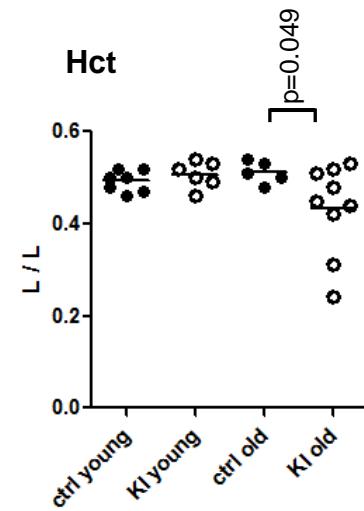
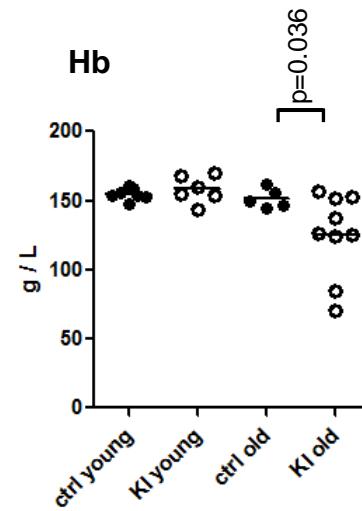
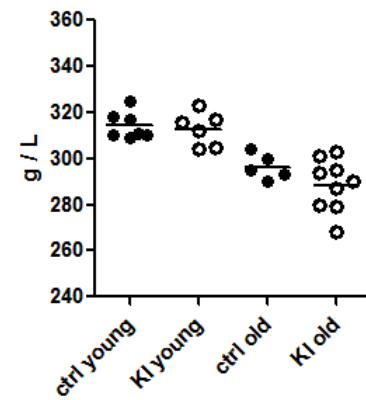
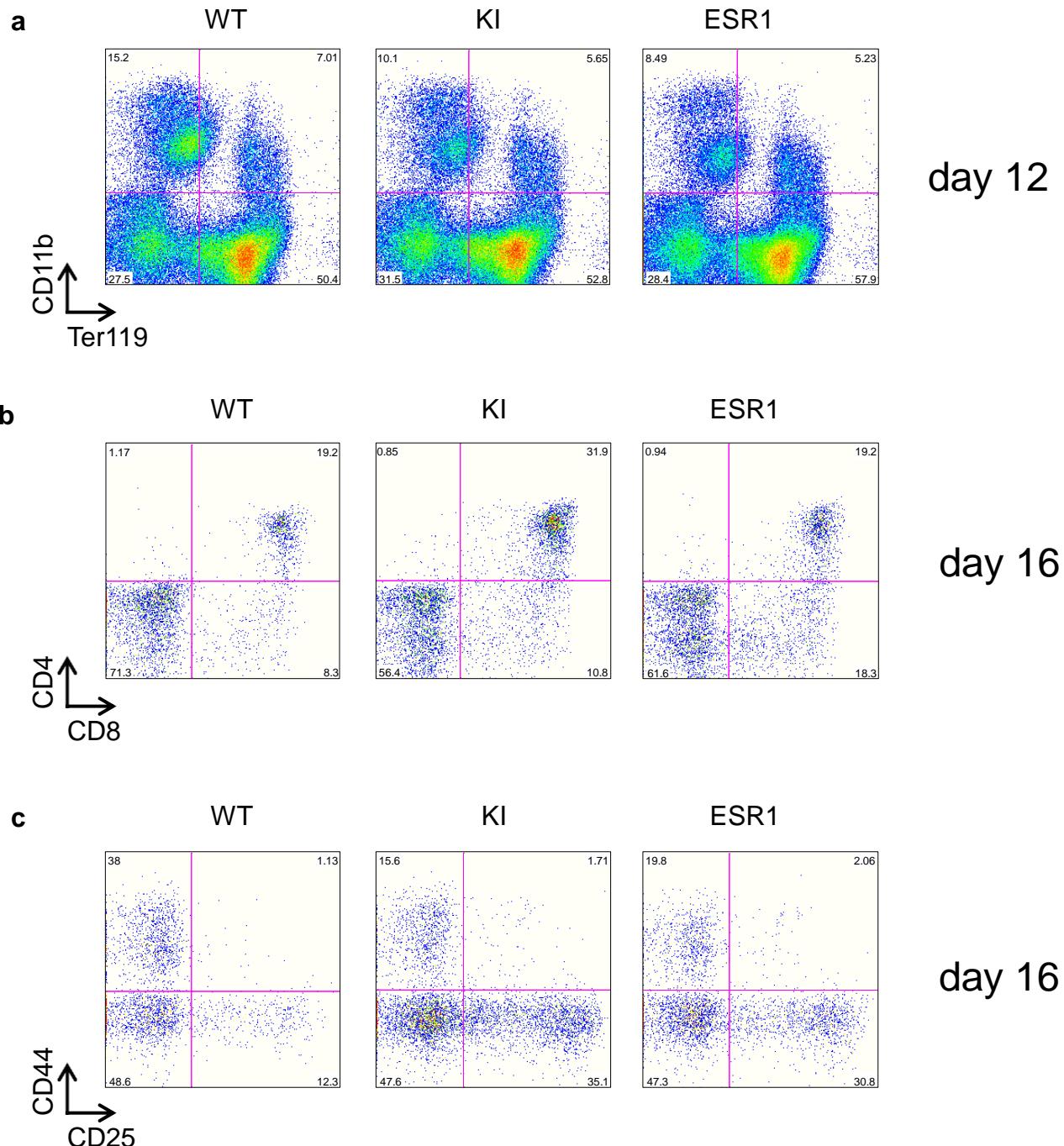


Figure 3

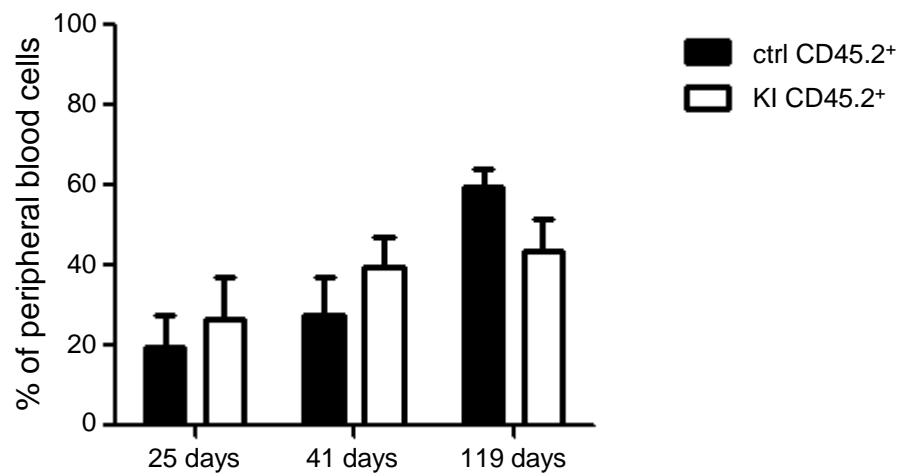
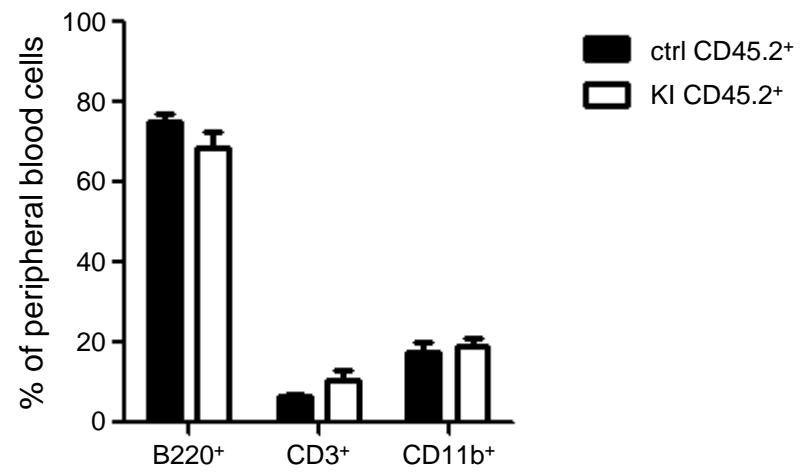
a**b****Figure 4**

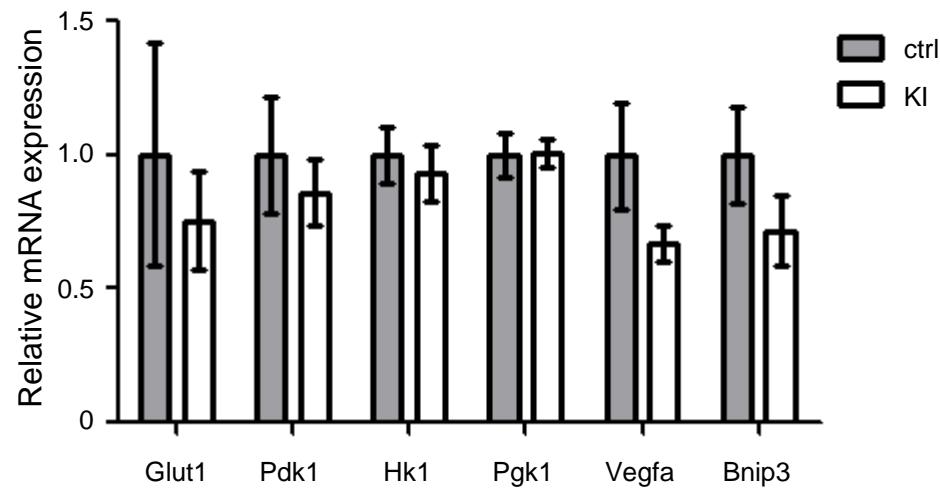
a WBC**b RBC****c Hct****d Hb****e MCHC****f**

	Young mice (7-16 weeks)					Older mice (42-46 weeks)				
	WT Mean \pm SEM	N =	KI Mean \pm SEM	N =	p-value	WT Mean \pm SEM	N =	KI Mean \pm SEM	N =	p-value
WBC	8.600 ± 1.295	7	7.400 ± 1.076	6	0.4924	9.260 ± 1.435	5	10.07 ± 0.851	9	0.6458
RBC	10.26 ± 0.156	7	10.55 ± 0.383	6	0.5050	10.46 ± 0.323	5	8.844 ± 0.718	9	0.0674
Hct	0.493 ± 0.009	7	0.507 ± 0.012	6	0.3803	0.512 ± 0.011	5	0.433 ± 0.033	9	0.0491*
Hb	154.9 ± 1.710	7	158.3 ± 4.072	6	0.4612	151.6 ± 3.311	5	125.3 ± 10.13	9	0.0359*
MCHC	314.3 ± 2.244	7	312.8 ± 3.005	6	0.7075	296.4 ± 2.502	5	286.3 ± 4.308	9	0.1112



Supplementary figure 2 |

a**b**



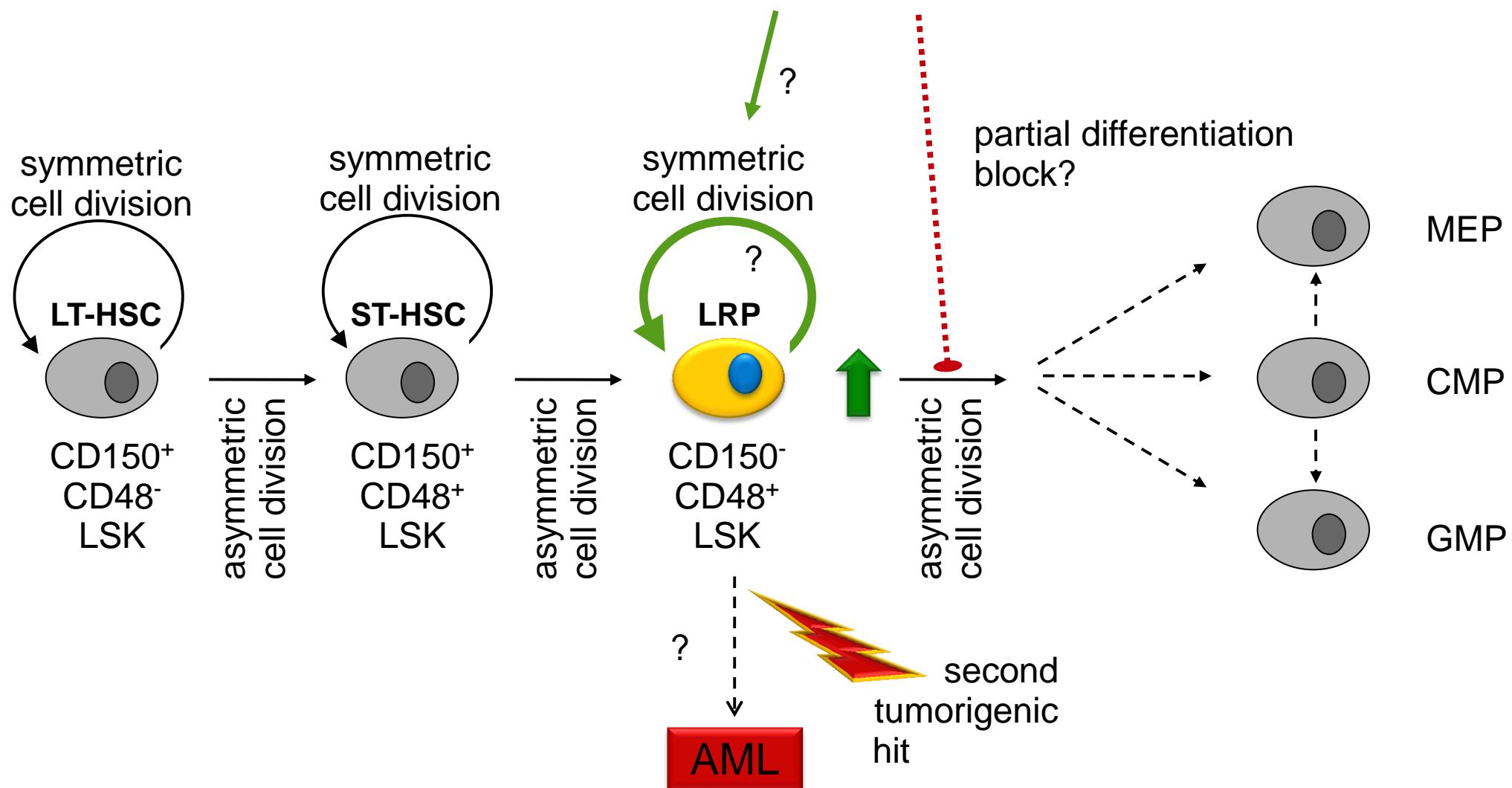
Supplementary figure 4 |



LysMCre-
R132H-IDH1-KI



- Elevated 2HG levels
 - Changes in histone methylation
 - Changes in DNA methylation



Supplementary figure 5 |

	genotype				
	WT-	WT+	HET-	HET+ *	total
all	64 (23.0%)	77 (27.8%)	69 (24.8%)	68 (24.4%)	278
female	35	38	31	28	132
male	29	39	38	40	146

	Young mice (7-16 weeks)					Older mice (42-46 weeks)				
	WT Mean \pm SEM	N=	KI Mean \pm SEM	N=	p-value	WT Mean \pm SEM	N=	KI Mean \pm SEM	N=	p-value
Lin ⁻	9.85 \pm 0.23	14	11.26 \pm 0.44	14	0.0100*	10.12 \pm 0.437	6	17.19 \pm 1.612	8	0.0029**
LSK (Lin ⁻ Sca1 ^{+c} Kit ⁺)	0.223 \pm 0.061	9	0.401 \pm 0.082	10	0.1003	0.360 \pm 0.102	6	1.946 \pm 0.311	8	0.0013**
LK (Lin ⁻ Sca1 ^{+c} Kit ⁺)	1.987 \pm 0.366	9	2.167 \pm 0.364	10	0.7313	3.065 \pm 0.3535	6	4.329 \pm 0.529	5	0.0724
LT-HSC (CD150 ^{+CD48⁻LSK)}	0.0023 \pm 0.0007	3	0.0032 \pm 0.0005	4	0.3694	0.0013 \pm 0.0006	6	0.0133 \pm 0.0061	8	0.0889
ST-HSC (CD150 ^{+CD48^{+LSK)}}	0.0090 \pm 0.0075	3	0.0117 \pm 0.0052	4	0.7911	0.0024 \pm 0.0009	6	0.1104 \pm 0.0719	8	0.1770
LRP (CD150 ^{-CD48^{+LSK)}}	0.1840 \pm 0.0594	3	0.3939 \pm 0.0450	4	0.0480*	0.1022 \pm 0.0194	6	0.9992 \pm 0.2358	8	0.0068**
CMP (CD34 ^{+CD16/32^{med}LK)}	0.2640 \pm 0.0480	5	0.2967 \pm 0.0429	6	0.6255	0.1417 \pm 0.0300	6	0.1009 \pm 0.0234	8	0.3090
GMP (CD34 ^{+CD16/32^{high}LK)}	0.2220 \pm 0.0381	5	0.3283 \pm 0.0488	6	0.1244	0.0973 \pm 0.0240	6	0.0967 \pm 0.0236	8	0.9864
MEP (CD34 ^{-CD16/32^{low}LK)}	0.5200 \pm 0.1820	5	0.7667 \pm 0.1880	6	0.3734	0.1790 \pm 0.0429	6	0.2104 \pm 0.0885	8	0.7566
CLP (Lin ⁻ Il7Ra ^{+Sca1^{med}c} Kit ^{med})	0.1283 \pm 0.0308	6	0.1129 \pm 0.0238	7	0.7002	0.1263 \pm 0.0942	4	0.1695 \pm 0.0904	6	0.7504

	Forward oligonucleotide (5'→3')	Reverse oligonucleotide (5'→3')	Fragment length
Glut1	gtcaccatctggagctgttc	cacctgcctctcgaaagatg	139bp
Vegfa	catgcggatcaaaccctcacca	ctggcttggctgtctttcttg	101bp
Pgk1	ccttcctaagggtgctcaaca	gtcaacaggcaaggtaatcttc	128bp
Hk1	ctgtctggacgacatcagaac	caggatgttacggacgatctc	127bp
Bnip3	cacagctactctcagcatgagaa	cagacgcctccaatgttagatc	145bp
Pdk1	ggaagatcgacagactctcaac	gaggcgtatatggcaatc	136bp
Hprt	ggtggagatgatctctaactt	ccagcaagctgcaaccctaac	142bp