

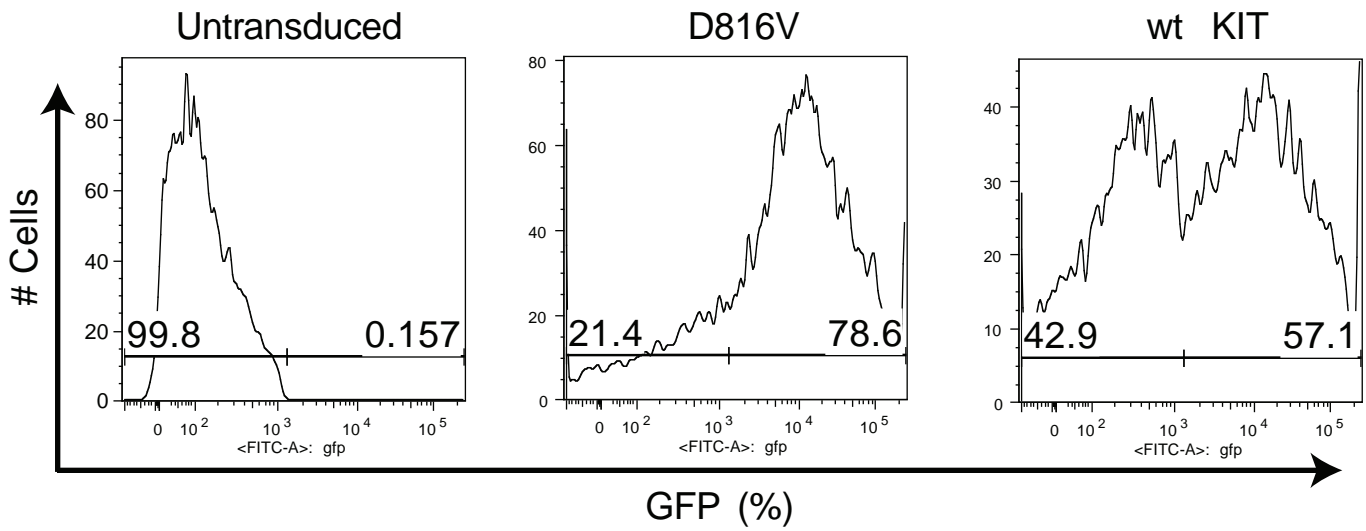
Table S1

Transduction efficiency (%) of BM cells using stable viral producing lines (co-culture)

BM cells transduced	816V (%)	816Y(%)	Wt (%)	GFP (%)
MxCre-CM	12-79 (N = 7)	10-78 (N =9)	8-57 (N =10)	52-77 (N = 3)
wt	12-47 (N = 6)	9-69 (N = 6)	15-43 (N = 6)	34-56 (N = 2)

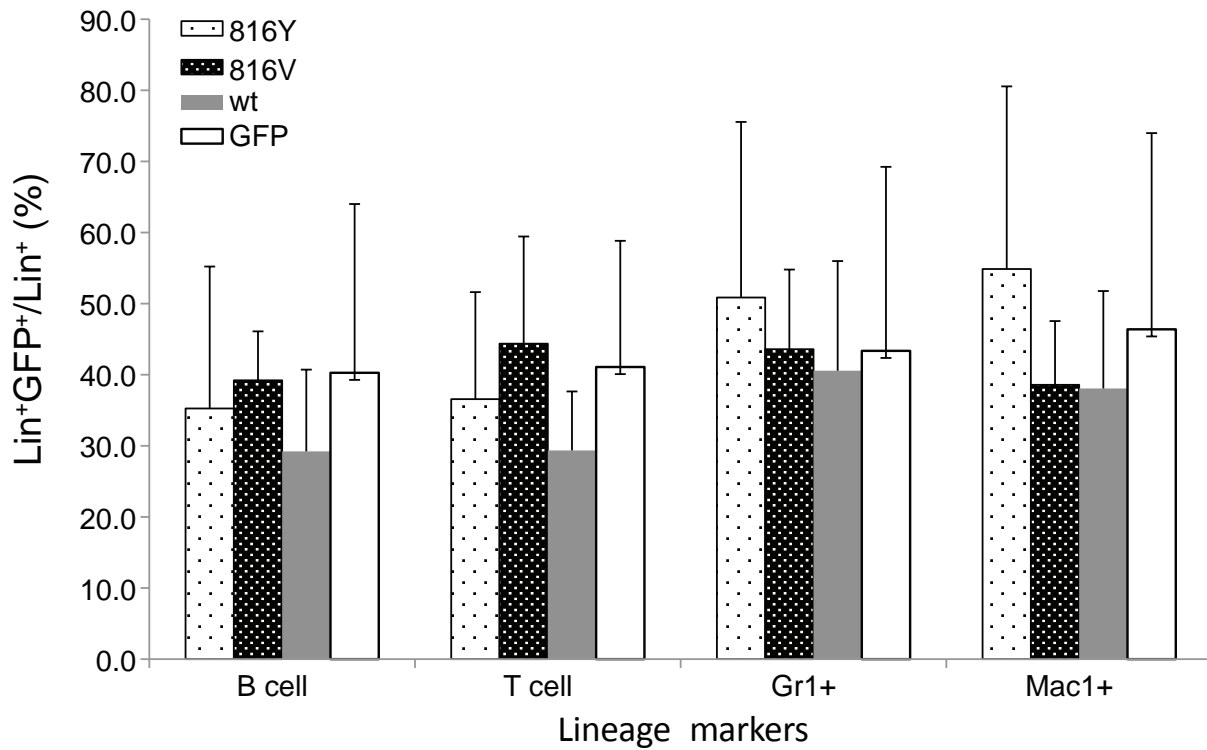
N: number of experiments

Figure S1



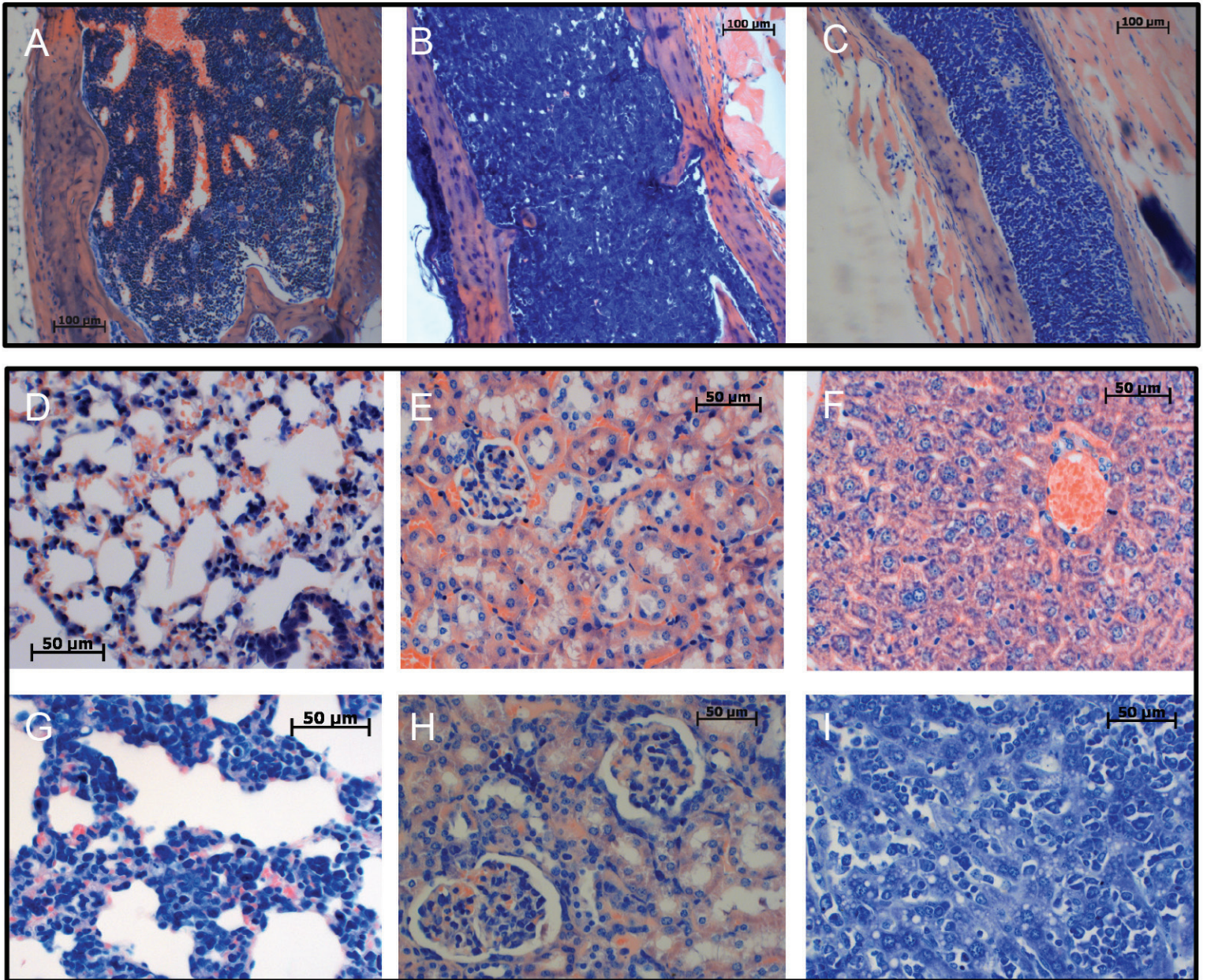
**Transduction efficiency of *Cbfb*<sup>+/-56m</sup>; *Tg*(Mx1-Cre) BM cells co-cultured with stable retroviral producing lines for D816V KIT or wt KIT. The Y-axis is the number of cells and the X-axis is GFP.**

Figure S2



**Contribution of KIT or GFP vector transduced *Cbfb*<sup>+56m</sup>; *Tg(Mx1-Cre)* BM cells to different lineages in the peripheral blood, 12 weeks after transplantation.** White dotted bars, D816Y KIT (n = 12); black dotted bars: D816V KIT (n = 9); grey solid bars: wt KIT (n = 6); white solid bars: GFP vector (n = 7).

Figure S3



**Sections of leukemia infiltrated tissues by Wright-Giemsa staining. A:** WT BM; **B:** BM from Cbfb<sup>+/56m</sup>; Tg(Mx1-Cre)/KIT<sup>D816Y</sup> leukemic mouse; **C:** BM from secondary transplantation of Cbfb<sup>+/56m</sup>; Tg(Mx1-Cre)/KIT<sup>D816Y</sup> leukemic cells; **D:** WT lung; **E:** WT kidney; **F:** WT liver; **G:** lung from a Cbfb<sup>+/56m</sup>; Tg(Mx1-Cre)/KIT<sup>D816Y</sup> leukemic mouse; **H:** kidney from a Cbfb<sup>+/56m</sup>; Tg(Mx1-Cre)/KIT<sup>D816Y</sup> leukemic mouse; **I:** liver from a Cbfb<sup>+/56m</sup>; Tg(Mx1-Cre)/KIT<sup>D816Y</sup> leukemic mouse.