

SUPPLEMENTAL MATERIAL

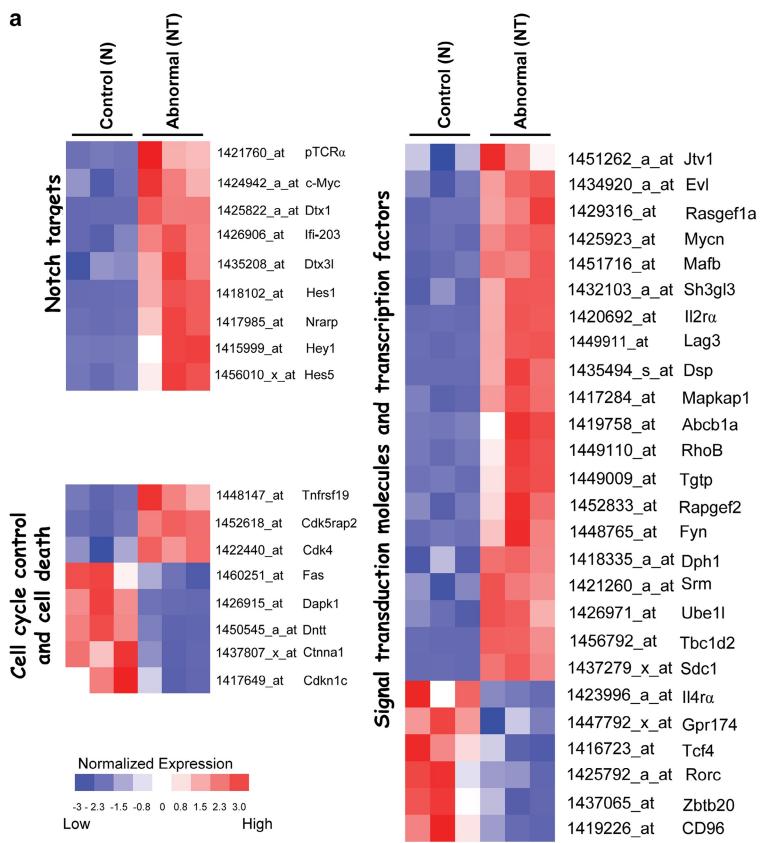
Li et al., <http://www.jem.org/cgi/content/full/jem.20081561/DC1>**a**

Figure S1. Global gene expression. Gene expression profiling was performed in control CD4 $^{+}$ 8 $^{+}$ versus abnormal nonmalignant ICN1-overexpressing CD4 $^{+}$ 8 $^{+}$ cells (a) and abnormal nonmalignant versus tumor CD4 $^{+}$ 8 $^{+}$ cells (b).

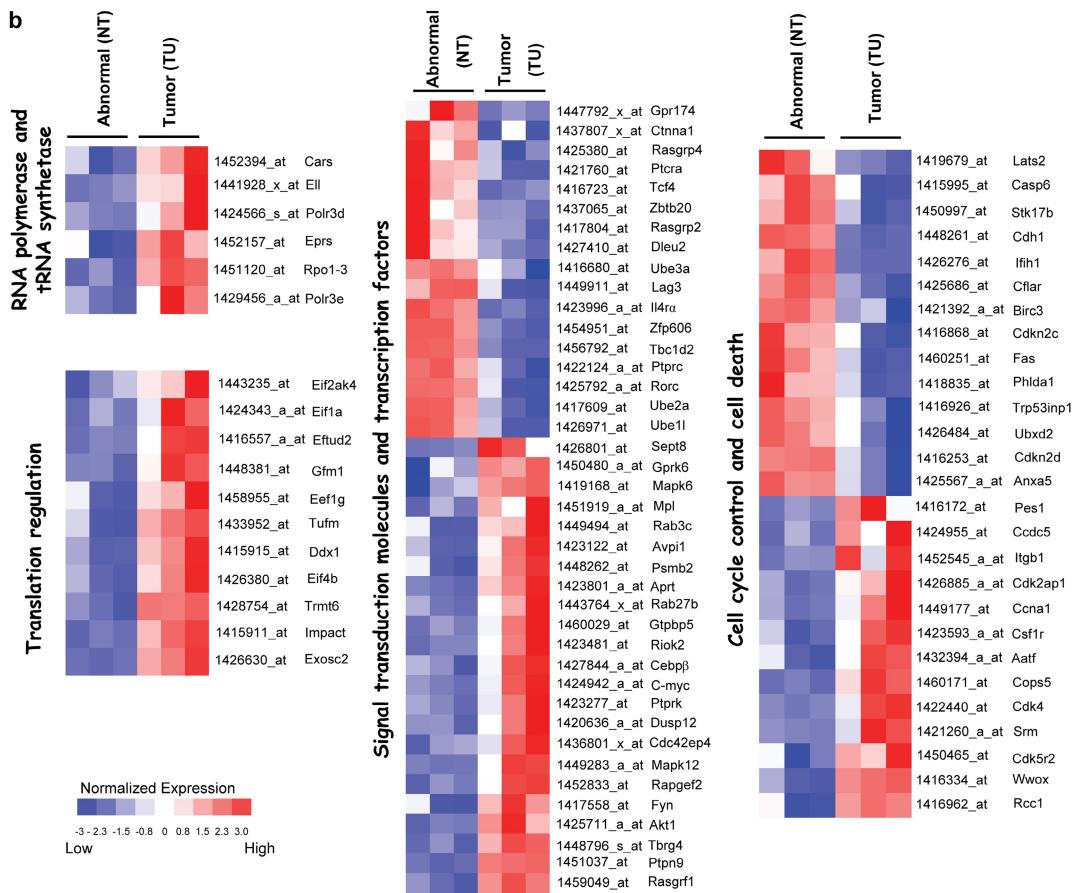
b

Table S1. Tumor development in nude mice

Time after ICN1 BMT	Sorted Cells (2x10 ⁶)	Post injection in nude recipient mice	Tumor Development
2 weeks	GFP ⁺ /DP	2wk	—
		2mo	—
		6mo	—
6 weeks	GFP ⁺ /DP	2wk	+
		3.5wk	+
	GFP ⁺ /CD4 ⁺ CD8 ⁺	2wk	+
		3.5wk	+
		3.5wk	+
8 weeks	GFP ⁺ /DP	2wk	+
		3.5wk	+
	GFP ⁺ /CD4 ⁺ CD8 ⁺	2wk	+
		3.5wk	+

Tumorigenicity assays in nu/nu homozygous mice. To determine which populations of GFP⁺ cells are tumorigenic, transplantation assays were performed in nu/nu homozygous mice. 10⁶ GFP⁺/CD4⁺8⁺ (GFP⁺/DP) splenocytes at 2 wk after BMT, and 10⁶ GFP⁺/DP or GFP⁺/CD4⁺8⁺ splenocytes at 6 and 8 wk after BMT were washed, pelleted, resuspended in 200 µl of PBS, and intraperitoneally injected into nude recipient mice. Blood samples of nude mice were analyzed by FACS, and mice were killed when displaying obvious clinical symptoms of malignancy.