

## Impact of the SCF signaling pathway on leukemia stem cell-mediated ATL initiation and progression in an HBZ transgenic mouse model

### Supplementary Material

**Supplemental Table 1: Antibodies, probes, and cytokines used in this study**

Product name	Clone/CatNo	Application	manufacture
anti-CD4	RM4.5	FCM	eBioscience
anti-CD8a	53-6.7	FCM	eBioscience
anti-CD16/32	93	FCM	eBioscience
anti-CD38	90	FCM	eBioscience
anti-CD71	R17217	FCM	eBioscience
anti-CD117	2B8	FCM	BioLegend
anti-CD3	Ab5690	IHC	Abcam
anti-B220	RA3-6B2	IHC	eBioscience
anti-Ter119	TER-119	IHC	eBioscience
Donkey anti-Rat Alexa 488	17A2	IHC	Jackson Immuno Laboratories
Donkey anti-Rabbit - Cy3	RM4.5	IHC	Jackson Immuno Laboratories
<i>HTLV-1 Tax</i>	VF6-19283	ISH/FCM	Affymetrix
<i>HTLV-1 Hbz</i>	VF6-19284	ISH/FCM	Affymetrix
<i>Hlf</i>	VB1-17786	ISH/FCM	Affymetrix
<i>Junb</i>	VB1-17776	ISH/FCM	Affymetrix
<i>Jund</i>	VB1-17779	ISH/FCM	Affymetrix
<i>Fosb</i>	VB1-12485	ISH/FCM	Affymetrix
<i>Tie1</i>	VB1-17781	ISH/FCM	Affymetrix
<i>mActb</i>	VB1-10350	ISH/FCM	Affymetrix
<i>c-kit</i>	VB1-11509	ISH/FCM	Affymetrix
rMurine IL-2	212-12	CC	Peprotech
rMurine IL-3	212-13	CC	Peprotech
rMurine IL-6	216-16	CC	Peprotech
rMurine SCF	250-03	CC	Peprotech
rMurine Flt3-Ligand	250-31L	CC	Peprotech
rMurine TPO	315-14	CC	Peprotech

**FCM:** Flowcytometry, **IHC:** Immunohistochemistry, **ISH:** *in situ* Hybridization, **CC:** Cell Culture, **JIL:** Jackson Immuno Laboratories

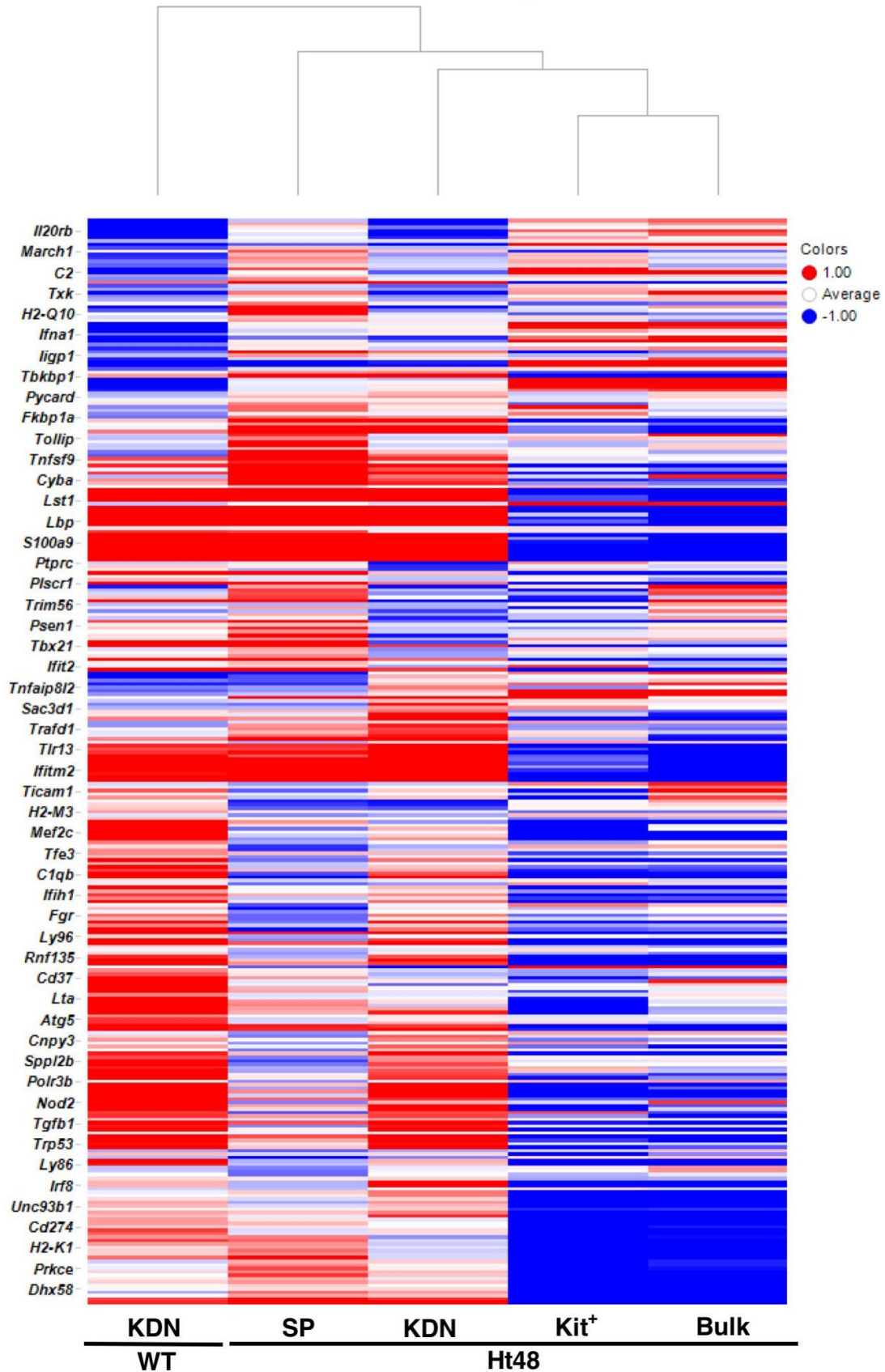
**Supplemental Table 2: Primers pairs used in this study.**

<b>Gene Symbol</b>	<b>forward</b>	<b>Reverse</b>
<i>HBZ</i>	ctgtgcttgacggtttgcta	cctctttctcccgtctttt
<i>Hlf</i>	gctgggcaaatacaagaaca	gacaggaaacaagctgtcca
<i>Junb</i>	gcctccgggacagtactttt	gggctttgacaaaaccgtcc
<i>Jund</i>	tcttgggctgctcaaactcg	tcgggtagaggaactgcgta
<i>Tie1</i>	agcagagtttgaagtgccca	tgacattgaccttgaagcgc
<i>Fosb</i>	tgcatcgaaactgggcagt	atggccgagtgggaatgagat
<i>Klra4</i>	acttacagcacacaggcaga	tggaattctggcagctctgt
<i>Robo4</i>	agaaaggctcgtggatgcagt	tagcgcattggtccctgtt
<i>Klra7</i>	attgccacgataactgcagc	agaagatcattgcctggcct
<i>Cxcl16</i>	aattggctggatgctggcta	agcagcgccaacaagaaaag
<i>Ndr2</i>	atagctacaacaaccgccga	acattccaccacggcatctt
<i>mouseβ-actin</i>	cagccttcctcttgggatgg	ctgtgtggcatagaggtctttacg

**Supplemental Table 3: List of up-regulated genes in Ht48 SP cells.**

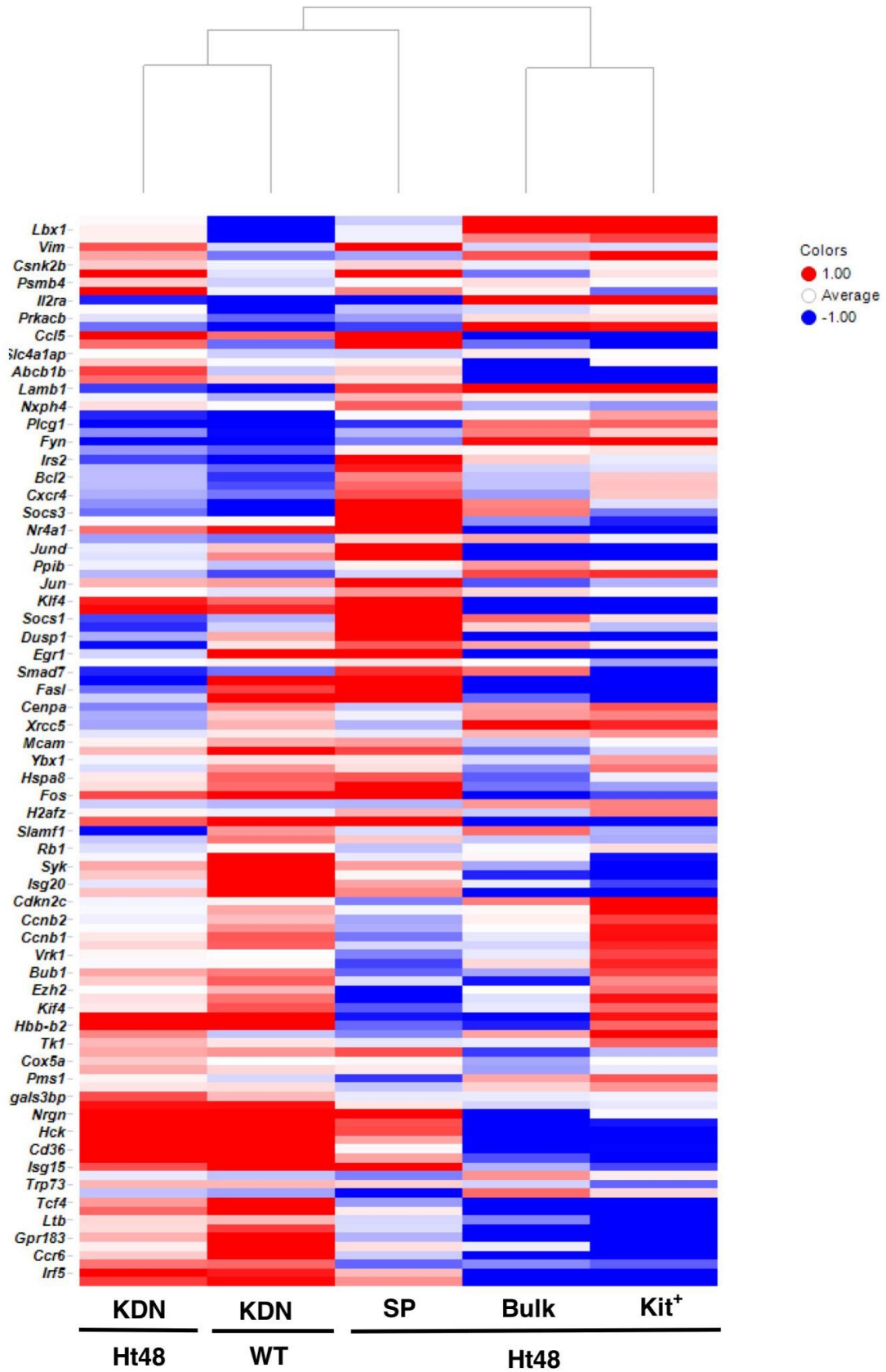
For Table S5, please see the attached Excel file

# Immune regulation gene set



**Supplemental Figure 1. Global gene expression profile of Ht48 cells.** Ht48 cells were subdivided into c-kit<sup>+</sup> cells, c-kit<sup>+</sup>CD4/8DN (**KDN**) cells, SP cells, and normal WT counterpart c-kit<sup>+</sup>CD4/8DN (**WT-KDN**) cells. Unsupervised hierarchical clustering of 632 immune regulatory gene expression profiles obtained from each fraction using an Agilent DNA microarray system. Immune regulatory genes were selected by the GO category.

# ATL related gene set



**Supplemental Figure 2 Global gene expression profile of Ht48 cells.** Ht48 cells were subdivided into c-kit<sup>+</sup> cells, c-kit<sup>+</sup>CD4/8DN (**KDN**) cells, SP cells, and normal WT counterpart c-kit<sup>+</sup>CD4/8DN (**WT-KDN**) cells. Unsupervised hierarchical clustering of 250 HTLV-1 infection- and ATL-associated gene expression profiles obtained from each fraction using an Agilent DNA microarray system.