

Table 1: **Experimental evidence for regulatory connections in the T-cell gene regulatory network.** Confidence levels for regulatory interaction are given as follows: 1 – Evidence in non-T cells only; 2 – Evidence from gene perturbation in T-cell progenitors alone, or from cis-regulatory analysis in T-cell progenitors alone; 3 – Evidence from gene perturbation in T-cell progenitors and cis-regulatory analysis. P – post-transcriptional regulatory interaction. FL – fetal liver, BM – bone marrow. Cited references are given below.

Source	Sign	Target	Level	Experiment	Cell Type	Reference
Bcl11b	+	CD3e	2	Bcl11b over-expression + <i>in vitro</i> T-cell culture + mRNA analysis	Bcl11b ^{-/-} FL Lin-Sca1+Kit ⁺ (LSK)	[1]
Bcl11b	-	CEBPa	2	Bcl11b over-expression + <i>in vitro</i> T-cell culture + mRNA analysis	Bcl11b ^{-/-} FL Lin-Sca1+Kit ⁺ (LSK)	[1]
Bcl11b	-	Id2	1	mRNA expression profiling Chip-seq using Bcl11b antibody	Bcl11b ^{-/-} double positive thymocytes	[2]
Bcl11b	-	Kit	2	Bcl11b over-expression + <i>in vitro</i> T-cell culture + surface expression analysis using flow cytometry	Bcl11b ^{-/-} FL Lin-Sca1+Kit ⁺ (LSK)	[1]
Bcl11b	+	pTa	2	Bcl11b over-expression + <i>in vitro</i> T-cell culture + mRNA analysis	FL Lin-Sca1+Kit ⁺ (LSK) cells	[1]
Bcl11b	-	PU.1	2	Bcl11b over-expression + <i>in vitro</i> T-cell culture + mRNA analysis	FL Lin-Sca1+Kit ⁺ (LSK) cells	[1]
Deltex	-	Notch signaling	P	Deltex over-expression + Notch reporter measurements	Jurkat T-cell line	[3]
E protein activity	-	Gata3	2	<i>In vitro</i> T-cell culture + mRNA expression analysis	HEB ^{-/-} Rag1 ^{-/-} DN3 cells	[4]
				mRNA expression analysis	E47 ^{bm/bm} thymocytes (bm – DNA-binding site mutation)	[5]
E protein activity	+	Gata3	1	Analysis of Gata3 cis-regulatory element (WT/mutated E-protein binding site)	Jurkat T-cell line	[6]
E protein activity	+	Gfi1	2	E47 over-expression + mRNA expression analysis	1F9 E2A ^{-/-} T-cell lymphoma line	[7]
E protein activity	+	Gfi1b	2	E47 over-expression + mRNA expression analysis / ChIP analysis using E2A antibody	E2A ^{-/-} T-cell lymphoma	[8]
				E47 over-expression + mRNA expression analysis	E2A ^{-/-} Lin- BM cells (cultured)	[9]
E protein activity	+	HEBAIt	2	Heb over-expression + mRNA expression analysis	P2C2 immature T-cell line	[10]
E protein activity	+	Hes1	2	E47 over-expression + mRNA expression analysis	E2A ^{-/-} Lin- BM cells (cultured)	[9]
E protein activity	+	IL-7R α	2	mRNA expression profiling	E2A ^{-/-} Lin-Sca1+Kit+Flt3-hi (LMPP)	[11]
				E47 over-expression + mRNA expression analysis	E2A ^{-/-} Lin- BM cells (cultured)	[9]
E protein activity	+	Lat	2	E47 over-expression + mRNA expression analysis	E2A ^{-/-} Lin- BM cells (cultured)	[9]
E protein activity	+	Notch1	3	mRNA expression profiling	E2A ^{-/-} Lin-Sca1+Kit+Flt3-hi (LMPP)	[11]
				ChIP analysis using E2A antibody E47 over-expression + Notch1 promoter activity measurements	Rag2 ^{-/-} DN3 thymocytes NIH-3T3 cell line	[12]
E protein activity	+	Notch3	2	E47 over-expression + mRNA expression analysis	E2A ^{-/-} Lin- BM cells (cultured)	[9]
E protein	+	pTa	3	E47 over-expression + mRNA	E2A ^{-/-} Lin- BM cells	[9]

activity				expression analysis	(cultured)	
E protein activity	+	Rag-1	2	mRNA expression profiling	E2A ^{-/-} Lin-Sca1+Kit+Flt3-hi (LMPP)	[11]
Scl/Lyl1/E	+	Id3	1	Lyl1 over-expression + mRNA expression analysis	human AML cells	[13]
Scl/Lyl1/E	+	Hhex	1	Analysis of Hhex enhancer containing Scl sites Chip-Seq using Scl antibody on putative enhancer	416B progenitor cell line HPC-7 progenitor cell line	[14] [15]
Scl/Lyl1/E	+	Kit	1	mRNA expression analysis	immature B-cells (B220 ⁺) from wildtype/SCL transgenic mice	[16]
Scl/Lyl1/E	-	pTa	2	mRNA expression analysis Scl over-expression + pT α enhancer activity analysis	Scl-Lmo1 transgenic mice AD10.1 immature T-cell line	[17]
Scl/Lyl1/E	+	Runx1	1	Scl over-expression + mRNA expression profiling ChIP analysis using Scl antibody	SCL ^{-/-} yolk sac cell line yolk sac / FL	[18]
Scl/Lyl1/E	+	Runx3	1	Scl over-expression + mRNA expression profiling ChIP analysis using Scl antibody	SCL ^{-/-} yolk sac cell line yolk sac / FL	[18]
Ets1	+	TCR β	2	Ets1 over-expression + analysis of TCR β enhancer activity	p19 cell line	[19]
Gata3	-	CEBPa	2	Gata3 over-expression + mRNA expression analysis	Bcl2-transgenic thymocytes	[20]
Gata3	+	Cpa3	3	Gata3 over-expression + mRNA expression analysis	Bcl2-transgenic thymocytes	[20]
Gata3	+	Deltex	1	Gata3 over-expression + mRNA expression analysis	adult DN1 thymocytes	[21]
Gata3	+	Gfi1b	2	Gata3 over-expression + mRNA expression analysis	Bcl2-transgenic thymocytes	[20]
Gata3	-	IL-7R α	2	Gata3 over-expression + mRNA expression analysis	Bcl2-transgenic thymocytes	[20]
Gata3	+	Kit	2	Gata3 over-expression + mRNA expression analysis	Bcl2-transgenic thymocytes	[20]
Gata3	-	Lef1	2	Gata3 over-expression + mRNA expression analysis	Bcl2-transgenic thymocytes	[20]
Gata3	+	Notch3	2	Gata3 over-expression + mRNA expression analysis	Bcl2-transgenic thymocytes	[20]
Gata3	-	pTa	2	Gata3 over-expression + mRNA expression analysis	Bcl2-transgenic thymocytes	[20]
Gata3	-	PU.1	2	Gata3 over-expression + mRNA expression analysis	Bcl2-transgenic thymocytes	[20]
Gata3	-	Rag-1	2	Gata3 over-expression + mRNA expression analysis	Bcl2-transgenic thymocytes	[20]
Gata3	+	Scl/Tal1	2	Gata3 over-expression + mRNA expression analysis	Bcl2-transgenic thymocytes	[20]
Gata3	-	TCF-1	2	Gata3 over-expression + mRNA expression analysis	Bcl2-transgenic thymocytes	[20]
Gata3	+	TCR β	2	Measurement of TCR β enhancer activity (wildtype/mutated GATA binding site)	p5424 T-cell line	[22]
Gfi1	+	Lmo2	1	Gfi1 knockdown + <i>in situ</i> RNA hybridization	Zebrafish embryos	[23]
Gfi1	-	PU.1	3	mRNA expression analysis; ChIP analysis using Gfi1 antibody Gfi1 over-expression/knockdown + <i>in</i>	Gfi1 ^{-/-} Lin-Sca1+Kit+ (LSK) cells Zebrafish embryos	[24] [23]

				<i>situ</i> RNA hybridization		
Gfi1	+	Rag-1	1	Gfi1 knockdown + <i>in situ</i> RNA hybridization	Zebrafish embryos	[23]
Gfi1	+	Scf/Tal1	1	Gfi1 knockdown + <i>in situ</i> RNA hybridization	Zebrafish embryos	[23]
Gfi1b	+	Bcl11b	1	Gfi1b knockdown + mRNA expression profiling	K562 myeloid leukemic line	[25]
Gfi1b	-	Gata3	2	Gfi1b over-expression + mRNA expression analysis	E2A ^{-/-} T-cell lymphoma	[8]
Gfi1b	-	Gfi1	2	Gfi1b over-expression + mRNA expression analysis mRNA expression analysis	E2A ^{-/-} T-cell lymphoma Gfi1b transgenic thymocytes	[8] [26]
Ikaros	-	Gfi1	2	Analysis of Gfi1-GFP reporter expression by flow cytometry	Ikaros ^{-/-} Lin-Sca1+Kit ⁺ (LSK) cells with Gfi1-GFP reporter	[24]
Ikaros	-	PU.1	2	mRNA expression analysis	Ikaros ^{-/-} Lin-Sca1+Kit ⁺ (LSK) cells	[24]
Ikaros	-	Runx1	2	mRNA expression analysis	Ikaros ^{-/-} adult DN thymocytes	[27]
IL-7R/Stat	-	Bcl11b	2	<i>In vitro</i> T-cell culture + IL7 drop + mRNA expression analysis	FL Lin-Sca1+Kit ⁺ (LSK) cells	[1]
IL-7R/Stat	+	CEBPa	2	<i>In vitro</i> T-cell culture + IL7 drop + mRNA expression analysis	FL Lin-Sca1+Kit ⁺ (LSK) cells	[1]
IL-7R/Stat	+	Ebf1	1	constitutive IL7 signaling activation + mRNA expression analysis	IL-7R α ^{-/-} pre-pro B-cells	[28]
IL-7R/Stat	+	GATA3	1	Flow cytometry with intracellular staining using GATA3 antibody ChIP with Stat5 antibody	Stat5 ^{-/-} Th2 cells (n.b. cells were stimulated with IL-2 and IL-33)	[29]
IL-7R/Stat	+	Kit	2	<i>In vitro</i> T-cell culture + IL7 drop + mRNA expression analysis	FL Lin-Sca1+Kit ⁺ (LSK) cells	[1]
IL-7R/Stat	-	Lck	2	<i>In vitro</i> T-cell culture + IL7 drop + analysis of Lck reporter	FL Lin-Sca1+Kit ⁺ (LSK) cells	[1]
IL-7R/Stat	-	Lef1	2	IL-7 stimulation + mRNA expression analysis	IL-7R α transgenic immature thymocytes	[30]
IL-7R/Stat	-	pTa	2	<i>In vitro</i> T-cell culture + IL7 drop + mRNA expression analysis	FL Lin-Sca1+Kit ⁺ (LSK) cells	[1]
IL-7R/Stat	+	PU.1	2	<i>In vitro</i> T-cell culture + IL7 drop + mRNA expression analysis	FL Lin-Sca1+Kit ⁺ (LSK) cells	[1]
IL-7R/Stat	-	TCF-1	2	<i>In vitro</i> T-cell culture + IL7 drop + mRNA expression analysis IL-7 stimulation + mRNA expression analysis	FL Lin-Sca1+Kit ⁺ (LSK) cells IL-7R α transgenic immature thymocytes	[1] [30]
Lmo2	+	E2A/Scf	P	EMSA assay using Lmo2 and Tal1 antibody	Lmo transgenic immature T-cell line	[31]
Lmo2	+	Hhex	2	mRNA expression analysis	Lmo transgenic DN3 leukemic cells	[32]
Lmo2	+	Kit	2	mRNA expression analysis	Lmo transgenic DN3 leukemic cells	[32]
Lmo2	+	Lyl1	2	mRNA expression analysis	Lmo transgenic DN3 leukemic cells	[32]
Myb	+	Gata3	1	dominant negative Myb over-expression + mRNA expression analysis	E16 cell line	[33]
Notch signaling	+	Bcl11b	3	<i>In vitro</i> culture (+/-DL1) + mRNA expression analysis ChIP analysis using CSL antibody	Thy1 ⁺ fetal thymocytes adult thymocytes	[34] [35]

Notch signaling	+	CD25	2	<i>In vitro</i> culture (+/-DL1) + surface expression analysis using flow cytometry surface expression analysis using flow cytometry	FL Lin-Kit+ cells DN3 thymocytes with inhibitor of Notch signaling (DNMAML)	[36] [37]
Notch signaling	+	CD3e	2	<i>In vitro</i> culture (+/-DL1) + mRNA expression analysis	Bcl2-transgenic thymocytes	[20]
Notch signaling	+	CD3g	2	<i>In vitro</i> culture (+/-DL1) + mRNA expression analysis	Bcl2-transgenic thymocytes	[20]
Notch signaling	-	CEBPa		<i>In vitro</i> culture (+/-DL1) + mRNA expression analysis	Thy1+ fetal thymocytes	[34]
Notch signaling	+	Deltex	3	<i>In vitro</i> culture (+/-DL1) + mRNA expression analysis <i>In vitro</i> culture (+/-DL1) + mRNA expression analysis <i>In vitro</i> culture (+/-DL1) + mRNA expression analysis	FL Lin-Kit+ cells Thy1+fetal thymocytes Bcl2-transgenic thymocytes	[36] [34] [20] many earlier references
Notch signaling	+	E2A (weak)	2	<i>In vitro</i> culture (+/-DL1) + mRNA expression analysis	Bcl2-transgenic thymocytes	[20]
Notch signaling	-	Ebf1	2	<i>In vitro</i> culture (+/-DL1) + mRNA expression analysis	FL Lin-Kit+ cells	[36]
Notch signaling	+	Gata3	2	<i>In vitro</i> culture (+/-DL1) + mRNA expression analysis (connection absent in humans)	FL Lin-Kit+ cells	[36] [38, 39]
Notch signaling	+	Gfi1b	2	<i>In vitro</i> culture (+/-DL1) + mRNA expression analysis	Bcl2-transgenic thymocytes	[20]
Notch signaling	+	HEBAIt	2	<i>In vitro</i> culture (+/-DL1) + mRNA expression analysis	Thy1+ fetal thymocytes	[34]
Notch signaling	+	Hes1	2	<i>In vitro</i> culture (+/-DL1) + mRNA expression analysis	FL Lin-Kit+ cells	[36] many earlier references
Notch signaling	-	Id3	2	<i>In vitro</i> culture (+/-DL1) + mRNA expression analysis <i>In vitro</i> culture (+/-DL1) + mRNA expression analysis	Thy1+ fetal thymocytes Bcl2-transgenic thymocytes	[34] [20]
Notch signaling	+	IL-7R α	3	Notch ICN over-expression + <i>in vitro</i> culture + surface expression analysis by flow Notch ICN over-expression + analysis of IL-7R α promoter activity	DN1 thymocytes 293T and Jurkat cell lines	[40]
Notch signaling	+	Lat	2	<i>In vitro</i> culture (+/-DL1) + mRNA expression analysis	Bcl2-transgenic thymocytes	[20]
Notch signaling	+	Lck	2	<i>In vitro</i> culture (+/-DL1) + mRNA expression analysis	Bcl2-transgenic thymocytes	[20]
Notch signaling	+	Lef1	2	<i>In vitro</i> culture (+/-DL1) + mRNA expression analysis	Bcl2-transgenic thymocytes	[20]
Notch signaling	+	Notch1	2	<i>In vitro</i> culture (+/-DL1) + mRNA expression analysis ChIP using CSL antibody Analysis of Notch 1 reporter construct	Bcl2-transgenic thymocytes Rag2-/- thymocytes NIH-3T3 cell line	[20] [12]
Notch signaling	+	Notch3	2	<i>In vitro</i> culture (+/-DL1) + mRNA expression analysis	Bcl2-transgenic thymocytes	[20]
Notch signaling	+	Nrarp	2	Notch IC over-expression + mRNA expression analysis Notch IC over-expression + analysis of Nrarp promoter activity	AKR1010 T-cell line 293T cells	[41] [42]
Notch signaling	+	pTa	3	Notch ICD over-expression + analysis of pT α enhancer activity <i>In vitro</i> culture (+/-DL1) + mRNA	293 cell line Thy1+ fetal thymocytes	[43] [34]

				expression analysis <i>In vitro</i> culture (+/-DL1) + mRNA expression analysis	Bcl2-transgenic thymocytes	[20]
Notch signaling	+	Rag-1	2	<i>In vitro</i> culture (+/-DL1) + mRNA expression analysis	Bcl2-transgenic thymocytes	[20]
Notch signaling	+	Runx1	3	<i>In vitro</i> culture (+/-DL1) + mRNA expression analysis <i>In vitro</i> culture (+/-DL1) + mRNA expression analysis	Thy1+ fetal thymocytes	[34]
					Bcl2-transgenic thymocytes NIH-3T3 cell line	[20] [44]
Notch signaling	+	TCF-1	2	<i>In vitro</i> culture (+/-DL1) + mRNA expression analysis	FL Lin-Kit+CD27+ progenitors	[45]
					Lin-cKit+Sca1+ (LSK) cells	[46]
Nrarp	-	Notch signaling	P	Nrarp over-expression + mRNA expression analysis of Notch responsive genes	AKR1010 T-cell line	[47]
PU.1	+	Bcl11a	2	PU.1 over-expression + mRNA expression analysis	Thy1+ fetal thymocytes	[34]
PU.1	-	Bcl11b	2	PU.1 over-expression + mRNA expression analysis	Thy1+ fetal thymocytes	[34]
PU.1	-	CD3e	2	PU.1 over-expression + mRNA expression analysis	Thy1+ fetal thymocytes	[34]
PU.1	-	CD3g	2	PU.1 over-expression + mRNA expression analysis	Thy1+ fetal thymocytes	[34]
PU.1	-	E2A	2	PU.1 over-expression + mRNA expression analysis	Thy1+ fetal thymocytes	[34]
PU.1	-	Ets1	2	PU.1 over-expression + mRNA expression analysis	Thy1+ fetal thymocytes	[34]
PU.1	-	Gata3 activity	P	ChIP using Gata3 antibody	PU.1-/- CD4+ T-cells	[48]
PU.1	-	Gfi1	2	PU.1 over-expression + mRNA expression analysis	Thy1+ fetal thymocytes	[34]
PU.1	-	HEB	2	PU.1 over-expression + mRNA expression analysis	Thy1+ fetal thymocytes	[34]
PU.1	-	HEBAIt	2	PU.1 over-expression + mRNA expression analysis	Thy1+ fetal thymocytes	[34]
PU.1	-	Hes1	2	PU.1 over-expression + mRNA expression analysis	Thy1+ fetal thymocytes	[34]
PU.1	+	Id2	2	PU.1 over-expression + mRNA expression analysis	Thy1+ fetal thymocytes	[34]
PU.1	-	Id3	2	PU.1 over-expression + mRNA expression analysis	Thy1+ fetal thymocytes	[34]
PU.1	-	Ikaros	2	PU.1 over-expression + mRNA expression analysis	Thy1+ fetal thymocytes	[34]
PU.1	+	IL-7R α	2	mRNA expression analysis ChIP using PU.1 antibody	FL PU.1-/- FL cells FL-derived pro-B cells	[49]
PU.1	-	IL-7R α	2	PU.1 over-expression + mRNA expression analysis	Thy1+ fetal thymocytes	[34]
PU.1	-	Kit	2	PU.1 over-expression + surface expression analysis by flow cytometry	Thy1+ fetal thymocytes	[34]
PU.1	-	Lat	2	PU.1 over-expression + mRNA expression analysis	Thy1+ fetal thymocytes	[34]
PU.1	-	Lck	2	PU.1 over-expression + mRNA expression analysis	Thy1+ fetal thymocytes	[34]
PU.1	+	Lmo2	1	PU.1 over-expression + Lmo2 promoter activity measurements ChIP using PU.1 antibody	293T cell line 416B myeloid cell line	[50]
PU.1	+	Lyl1	1	ChIP using PU.1 antibody Lyl1 promoter activity measurements	416B myeloid cell line	[51]

				(WT/mutated PU.1 site)		
PU.1	-	Myb	2	PU.1 over-expression + mRNA expression analysis	Thy1+ fetal thymocytes	[34]
PU.1	+	PU.1	1	transfection with PU.1 regulatory element construct (+/- mutation to PU.1 binding site) + analysis of reporter expression / ChIP using PU.1 antibody	416B myeloid cell line	[52]
PU.1	-	Rag-1	2	PU.1 over-expression + mRNA expression analysis	Thy1+ fetal thymocytes	[34]
PU.1	-	Runx3	1	PU.1 over-expression + mRNA expression analysis	Thy1+ fetal thymocytes	[34]
PU.1	-	TCF-1	2	PU.1 over-expression + mRNA expression analysis	Thy1+ fetal thymocytes	[34]
PU.1	-	Zap70	2	PU.1 over-expression + mRNA expression analysis	Thy1+ fetal thymocytes	[34]
Runx1	-	PU.1	3	mRNA expression analysis	Runx1 ^{-/-} DN2, DN3 thymocytes	[53]
				Runx1/Runx1 dominant negative over-expression + analysis of PU.1 cis-regulatory element	P2C2 immature T-cell line + Raw264 myeloid cell line	[54]
Runx1	-	Rag-1	2	Analysis of Rag1-GFP reporter expression by flow cytometry (wildtype/mutated Runx1 site)	Rag1 GFP-reporter transgenic DN thymocytes	[55]
Runx1	+	TCR β	1	Runx1 over-expression + analysis of TCR β enhancer activity	p19 cell line	[19]
TCF-1	-	PU.1	1	Analysis of PU.1 cis-regulatory element (WT/mutated Tcf site)	EL4 T-cell line	[56]
TCF-1	+	Lef1	2	Wnt pathway activation, ChiP using TCF antibody	DLD1 cancer cell line	[57]
				TCF-1 over-expression + mRNA expression analysis	Lin-cKit+Sca1+ (LSK) cells	[46]
TCF-1	+	TCF-1	3	TCF-1 over-expression + mRNA expression analysis	Lin-cKit+Sca1+ (LSK) cells	[46]
				ChIP analysis using TCF-1 antibody		
TCF-1	+	Bcl11b	3	TCF-1 over-expression + mRNA expression analysis	Lin-cKit+Sca1+ (LSK) cells	[46]
				ChIP analysis using TCF-1 antibody		
TCF-1	+	GATA3	3	TCF-1 over-expression + mRNA expression analysis	Lin-cKit+Sca1+ (LSK) cells	[46]
				ChIP analysis using TCF-1 antibody	TCF ^{-/-} Th2 cells	[58]
TCF-1	+	CD25	3	TCF-1 over-expression + mRNA expression analysis	Lin-cKit+Sca1+ (LSK) cells	[46]
				ChIP analysis using TCF-1 antibody		

References

1. Ikawa, T, Hirose, S, Masuda, K, Kakugawa, K, Satoh, R, Shibano-Satoh, A, Kominami, R, Katsura, Y, Kawamoto, H. *An essential developmental checkpoint for production of the T cell lineage*. Science 2010. 329(5987):93-6.
2. Kastner, P, Chan, S, Vogel, WK, Zhang, LJ, Topark-Ngarm, A, Golonzhka, O, Jost, B, Le Gras, S, Gross, MK, Leid, M. *Bcl11b represses a mature T-cell gene expression program in immature CD4(+)CD8(+) thymocytes*. Eur J Immunol 2010. 40(8):2143-54.
3. Izon, DJ, Aster, JC, He, Y, Weng, A, Karnell, FG, Patriub, V, Xu, L, Bakkour, S, Rodriguez, C, Allman, D, Pear, WS. *Deltex1 redirects lymphoid progenitors to the B cell lineage by antagonizing Notch1*. Immunity 2002. 16(2):231-43.
4. Braunstein, M, Anderson, MK. *HEB-deficient T-cell precursors lose T-cell potential and adopt an alternative pathway of differentiation*. Mol Cell Biol 2011. 31(5):971-82.
5. Jia, J, Dai, M, Zhuang, Y. *E proteins are required to activate germline transcription of the TCR Vbeta8.2 gene*. Eur J Immunol 2008. 38(10):2806-20.
6. Gregoire, JM, Romeo, PH. *T-cell expression of the human GATA-3 gene is regulated by a non-lineage-specific silencer*. J Biol Chem 1999. 274(10):6567-78.
7. Schwartz, R, Engel, I, Fallahi-Sichani, M, Petrie, HT, Murre, C. *Gene expression patterns define novel roles for E47 in cell cycle progression, cytokine-mediated signaling, and T lineage development*. Proc Natl Acad Sci U S A 2006. 103(26):9976-81.
8. Xu, W, Kee, BL. *Growth factor independent 1B (Gfi1b) is an E2A target gene that modulates Gata3 in T-cell lymphomas*. Blood 2007. 109(10):4406-14.
9. Ikawa, T, Kawamoto, H, Goldrath, AW, Murre, C. *E proteins and Notch signaling cooperate to promote T cell lineage specification and commitment*. J Exp Med 2006. 203(5):1329-42.
10. Wang, D, Claus, CL, Vaccarelli, G, Braunstein, M, Schmitt, TM, Zuniga-Pflucker, JC, Rothenberg, EV, Anderson, MK. *The basic helix-loop-helix transcription factor HEBAIt is expressed in pro-T cells and enhances the generation of T cell precursors*. J Immunol 2006. 177(1):109-19.
11. Dias, S, Mansson, R, Gurbuxani, S, Sigvardsson, M, Kee, BL. *E2A proteins promote development of lymphoid-primed multipotent progenitors*. Immunity 2008. 29(2):217-27.
12. Yashiro-Ohtani, Y, He, Y, Ohtani, T, Jones, ME, Shestova, O, Xu, L, Fang, TC, Chiang, MY, Intlekofer, AM, Blacklow, SC, Zhuang, Y, Pear, WS. *Pre-TCR signaling inactivates Notch1 transcription by antagonizing E2A*. Genes Dev 2009. 23(14):1665-76.
13. San-Marina, S, Han, Y, Suarez Saiz, F, Trus, MR, Minden, MD. *Lyl1 interacts with CREB1 and alters expression of CREB1 target genes*. Biochim Biophys Acta 2008. 1783(3):503-17.
14. Donaldson, IJ, Chapman, M, Kinston, S, Landry, JR, Knezevic, K, Piltz, S, Buckley, N, Green, AR, Gottgens, B. *Genome-wide identification of cis-regulatory sequences controlling blood and endothelial development*. Hum Mol Genet 2005. 14(5):595-601.
15. Wilson, NK, Miranda-Saavedra, D, Kinston, S, Bonadies, N, Foster, SD, Calero-Nieto, F, Dawson, MA, Donaldson, IJ, Dumon, S, Frampton, J, Janky, R, Sun, XH, Teichmann, SA, Bannister, AJ, Gottgens, B. *The transcriptional program controlled by the stem cell leukemia gene Scf/Tal1 during early embryonic hematopoietic development*. Blood 2009. 113(22):5456-65.
16. Lecuyer, E, Herblot, S, Saint-Denis, M, Martin, R, Begley, CG, Porcher, C, Orkin, SH, Hoang, T. *The SCL complex regulates c-kit expression in hematopoietic cells through functional interaction with Sp1*. Blood 2002. 100(7):2430-40.
17. Herblot, S, Steff, AM, Hugo, P, Aplan, PD, Hoang, T. *SCL and LMO1 alter thymocyte differentiation: inhibition of E2A-HEB function and pre-T alpha chain expression*. Nat Immunol 2000. 1(2):138-44.
18. Landry, JR, Kinston, S, Knezevic, K, de Bruijn, MF, Wilson, N, Nottingham, WT, Peitz, M, Edenhofer, F, Pimanda, JE, Ottersbach, K, Gottgens, B. *Runx genes are direct targets of Scf/Tal1 in the yolk sac and fetal liver*. Blood 2008. 111(6):3005-14.
19. Kim, WY, Sieweke, M, Ogawa, E, Wee, HJ, Englmeier, U, Graf, T, Ito, Y. *Mutual activation of Ets-1 and AML1 DNA binding by direct interaction of their autoinhibitory domains*. EMBO J 1999. 18(6):1609-20.
20. Taghon, T, Yui, MA, Rothenberg, EV. *Mast cell lineage diversion of T lineage precursors by the essential T cell transcription factor GATA-3*. Nat Immunol 2007. 8(8):845-55.
21. Wang, HC, Perry, SS, Sun, XH. *Id1 attenuates Notch signaling and impairs T-cell commitment by elevating Deltex1 expression*. Mol Cell Biol 2009. 29(17):4640-52.

22. Yang, XO, Doty, RT, Hicks, JS, Willerford, DM. *Regulation of T-cell receptor D beta 1 promoter by KLF5 through reiterated GC-rich motifs.* Blood 2003. 101(11):4492-9.
23. Wei, W, Wen, L, Huang, P, Zhang, Z, Chen, Y, Xiao, A, Huang, H, Zhu, Z, Zhang, B, Lin, S. *Gfi1.1 regulates hematopoietic lineage differentiation during zebrafish embryogenesis.* Cell Res 2008. 18(7358):677-85.
24. Spooner, CJ, Cheng, JX, Pujadas, E, Laslo, P, Singh, H. *A recurrent network involving the transcription factors PU.1 and Gfi1 orchestrates innate and adaptive immune cell fates.* Immunity 2009. 31(4):576-86.
25. Koldehoff, M, Zakrzewski, JL, Klein-Hitpass, L, Beelen, DW, Elmaagacli, AH. *Gene profiling of growth factor independence 1B gene (Gfi-1B) in leukemic cells.* Int J Hematol 2008. 87(1):39-47.
26. Doan, LL, Porter, SD, Duan, Z, Flubacher, MM, Montoya, D, Tschlis, PN, Horwitz, M, Gilks, CB, Grimes, HL. *Targeted transcriptional repression of Gfi1 by GFI1 and GFI1B in lymphoid cells.* Nucleic Acids Res 2004. 32(8):2508-19.
27. Chari, S, Winandy, S. *Ikaros regulates Notch target gene expression in developing thymocytes.* J Immunol 2008. 181(9):6265-74.
28. Kikuchi, K, Lai, AY, Hsu, CL, Kondo, M. *IL-7 receptor signaling is necessary for stage transition in adult B cell development through up-regulation of EBF.* J Exp Med 2005. 201(8):1197-203.
29. Guo, L, Wei, G, Zhu, J, Liao, W, Leonard, WJ, Zhao, K, Paul, W. *IL-1 family members and STAT activators induce cytokine production by Th2, Th17, and Th1 cells.* Proc Natl Acad Sci U S A 2009. 106(32):13463-8.
30. Yu, Q, Erman, B, Park, JH, Feigenbaum, L, Singer, A. *IL-7 receptor signals inhibit expression of transcription factors TCF-1, LEF-1, and RORgamma: impact on thymocyte development.* J Exp Med 2004. 200(6):797-803.
31. Grutz, GG, Bucher, K, Lavenir, I, Larson, T, Larson, R, Rabbitts, TH. *The oncogenic T cell LIM-protein Lmo2 forms part of a DNA-binding complex specifically in immature T cells.* EMBO J 1998. 17(16):4594-605.
32. McCormack, MP, Young, LF, Vasudevan, S, de Graaf, CA, Codrington, R, Rabbitts, TH, Jane, SM, Curtis, DJ. *The Lmo2 oncogene initiates leukemia in mice by inducing thymocyte self-renewal.* Science 2010. 327(5967):879-83.
33. Maurice, D, Hooper, J, Lang, G, Weston, K. *c-Myb regulates lineage choice in developing thymocytes via its target gene Gata3.* EMBO J 2007. 26(15):3629-40.
34. Franco, CB, Scripture-Adams, DD, Proekt, I, Taghon, T, Weiss, AH, Yui, MA, Adams, SL, Diamond, RA, Rothenberg, EV. *Notch/Delta signaling constrains reengineering of pro-T cells by PU.1.* Proc Natl Acad Sci U S A 2006. 103(32):11993-8.
35. Li, P, Burke, S, Wang, J, Chen, X, Ortiz, M, Lee, SC, Lu, D, Campos, L, Goulding, D, Ng, BL, Dougan, G, Huntly, B, Gottgens, B, Jenkins, NA, Copeland, NG, Colucci, F, Liu, P. *Reprogramming of T cells to natural killer-like cells upon Bcl11b deletion.* Science 2010. 329(5987):85-9.
36. Taghon, TN, David, ES, Zuniga-Pflucker, JC, Rothenberg, EV. *Delayed, asynchronous, and reversible T-lineage specification induced by Notch/Delta signaling.* Genes Dev 2005. 19(8):965-78.
37. Maillard, I, Tu, L, Sambandam, A, Yashiro-Ohtani, Y, Millholland, J, Keeshan, K, Shestova, O, Xu, L, Bhandoola, A, Pear, WS. *The requirement for Notch signaling at the beta-selection checkpoint in vivo is absolute and independent of the pre-T cell receptor.* J Exp Med 2006. 203(10):2239-45.
38. Van de Walle, I, De Smet, G, De Smedt, M, Vandekerckhove, B, Leclercq, G, Plum, J, Taghon, T. *An early decrease in Notch activation is required for human TCR-alpha/beta lineage differentiation at the expense of TCR-gammadelta T cells.* Blood 2009. 113(13):2988-98.
39. Weerkamp, F, Luis, TC, Naber, BA, Koster, EE, Jeannotte, L, van Dongen, JJ, Staal, FJ. *Identification of Notch target genes in uncommitted T-cell progenitors: No direct induction of a T-cell specific gene program.* Leukemia 2006. 20(11):1967-77.
40. Gonzalez-Garcia, S, Garcia-Peydro, M, Martin-Gayo, E, Ballestar, E, Esteller, M, Bornstein, R, de la Pompa, JL, Ferrando, AA, Toribio, ML. *CSL-MAML-dependent Notch1 signaling controls T lineage-specific IL-7R{alpha} gene expression in early human thymopoiesis and leukemia.* J Exp Med 2009. 206(4):779-91.
41. Krebs, LT, Deftos, ML, Bevan, MJ, Gridley, T. *The Nrarp gene encodes an ankyrin-repeat protein that is transcriptionally regulated by the notch signaling pathway.* Dev Biol 2001. 238(1):110-9.
42. Piro, P, van Grunsven, LA, Marine, JC, Huylebroeck, D, Bellefroid, EJ. *Direct regulation of the Nrarp gene promoter by the Notch signaling pathway.* Biochem Biophys Res Commun 2004. 322(2):526-34.
43. Reizis, B, Leder, P. *Direct induction of T lymphocyte-specific gene expression by the mammalian Notch signaling pathway.* Genes Dev 2002. 16(3):295-300.
44. Nakagawa, M, Ichikawa, M, Kumano, K, Goyama, S, Kawazu, M, Asai, T, Ogawa, S, Kurokawa, M, Chiba, S. *AML1/Runx1 rescues Notch1-null mutation-induced deficiency of para-aortic splanchnopleural hematopoiesis.* Blood 2006. 108(10):3329-34.

45. Tydell, CC, David-Fung, ES, Moore, JE, Rowen, L, Taghon, T, Rothenberg, EV. *Molecular dissection of prethymic progenitor entry into the T lymphocyte developmental pathway.* J Immunol 2007. 179(1):421-38.
46. Weber, BN, Chi, AW, Chavez, A, Y., Y-O, Yang, Q, Shestova, O, Bhandoola, A. *A critical role for TCF-1 in T-lineage specification and differentiation.* Nature 2011. 476(7358):63-68.
47. Yun, TJ, Bevan, MJ. *Notch-regulated ankyrin-repeat protein inhibits Notch1 signaling: multiple Notch1 signaling pathways involved in T cell development.* J Immunol 2003. 170(12):5834-41.
48. Chang, HC, Han, L, Jabeen, R, Carotta, S, Nutt, SL, Kaplan, MH. *PU.1 regulates TCR expression by modulating GATA-3 activity.* J Immunol 2009. 183(8):4887-94.
49. DeKoter, RP, Lee, HJ, Singh, H. *PU.1 regulates expression of the interleukin-7 receptor in lymphoid progenitors.* Immunity 2002. 16(2):297-309.
50. Landry, JR, Bonadies, N, Kinston, S, Knezevic, K, Wilson, NK, Oram, SH, Janes, M, Piltz, S, Hammett, M, Carter, J, Hamilton, T, Donaldson, IJ, Lacaud, G, Frampton, J, Follows, G, Kouskoff, V, Gottgens, B. *Expression of the leukemia oncogene Lmo2 is controlled by an array of tissue-specific elements dispersed over 100 kb and bound by Tal1/Lmo2, Ets, and Gata factors.* Blood 2009. 113(23):5783-92.
51. Chan, WY, Follows, GA, Lacaud, G, Pimanda, JE, Landry, JR, Kinston, S, Knezevic, K, Piltz, S, Donaldson, IJ, Gambardella, L, Sablitzky, F, Green, AR, Kouskoff, V, Gottgens, B. *The paralogous hematopoietic regulators Lyl1 and Scl are coregulated by Ets and GATA factors, but Lyl1 cannot rescue the early Scl^{-/-} phenotype.* Blood 2007. 109(5):1908-16.
52. Okuno, Y, Huang, G, Rosenbauer, F, Evans, EK, Radomska, HS, Iwasaki, H, Akashi, K, Moreau-Gachelin, F, Li, Y, Zhang, P, Gottgens, B, Tenen, DG. *Potential autoregulation of transcription factor PU.1 by an upstream regulatory element.* Mol Cell Biol 2005. 25(7):2832-45.
53. Huang, G, Zhang, P, Hirai, H, Elf, S, Yan, X, Chen, Z, Koschmieder, S, Okuno, Y, Dayaram, T, Growney, JD, Shivdasani, RA, Gilliland, DG, Speck, NA, Nimer, SD, Tenen, DG. *PU.1 is a major downstream target of AML1 (RUNX1) in adult mouse hematopoiesis.* Nat Genet 2008. 40(1):51-60.
54. Zarnegar, MA, Chen, J, Rothenberg, EV. *Cell-type-specific activation and repression of PU.1 by a complex of discrete, functionally specialized cis-regulatory elements.* Mol Cell Biol 2010. 30(20):4922-39.
55. Yannoutsos, N, Barreto, V, Misulovin, Z, Gazumyan, A, Yu, W, Rajewsky, N, Peixoto, BR, Eisenreich, T, Nussenzweig, MC. *A cis element in the recombination activating gene locus regulates gene expression by counteracting a distant silencer.* Nat Immunol 2004. 5(4):443-50.
56. Rosenbauer, F, Owens, BM, Yu, L, Tumang, JR, Steidl, U, Kutok, JL, Clayton, LK, Wagner, K, Scheller, M, Iwasaki, H, Liu, C, Hackanson, B, Akashi, K, Leutz, A, Rothstein, TL, Plass, C, Tenen, DG. *Lymphoid cell growth and transformation are suppressed by a key regulatory element of the gene encoding PU.1.* Nat Genet 2006. 38(1):27-37.
57. Li, TW, Ting, JH, Yokoyama, NN, Bernstein, A, van de Wetering, M, Waterman, ML. *Wnt activation and alternative promoter repression of LEF1 in colon cancer.* Mol Cell Biol 2006. 26(14):5284-99.
58. Yu, Q, Sharma, A, Oh, SY, Moon, HG, Hossain, MZ, Salay, TM, Leeds, KE, Du, H, Wu, B, Waterman, ML, Zhu, Z, Sen, JM. *T cell factor 1 initiates the T helper type 2 fate by inducing the transcription factor GATA-3 and repressing interferon-gamma.* Nat Immunol 2009. 10(9):992-9.