

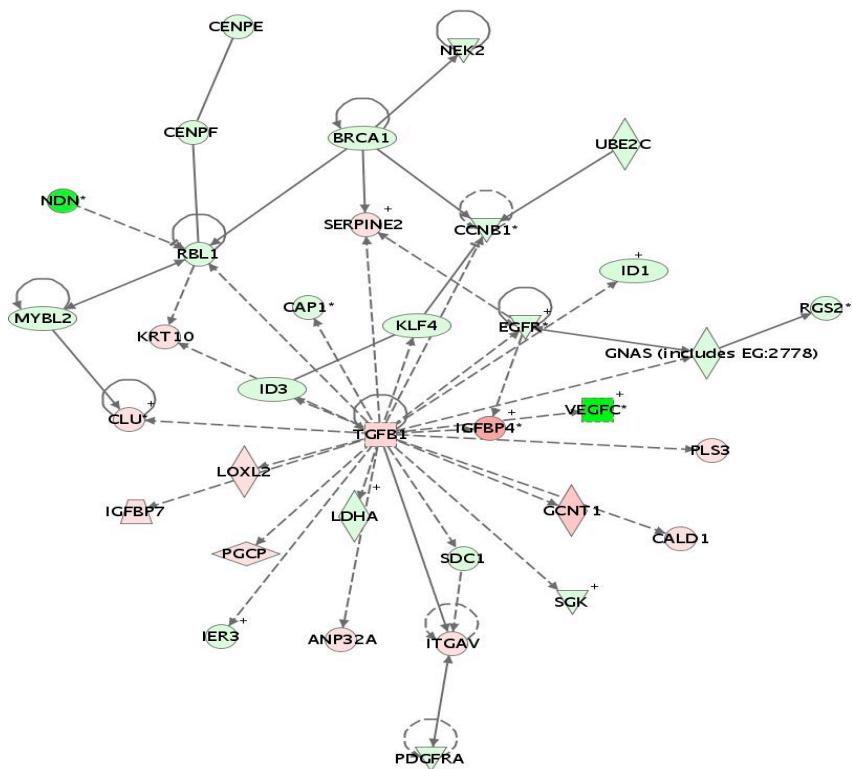
## Supplemental Data 1

*Pathway analysis between HNF-1 $\alpha$  (+/-)/RIP-Tag beta-cell line and HNF-1 $\alpha$  (+/+)/RIP-Tag beta-cell line or HNF-1 $\alpha$  (-/-)/RIP-Tag beta-cell line and HNF-1 $\alpha$  (+/+)/RIP-Tag beta-cell line.*

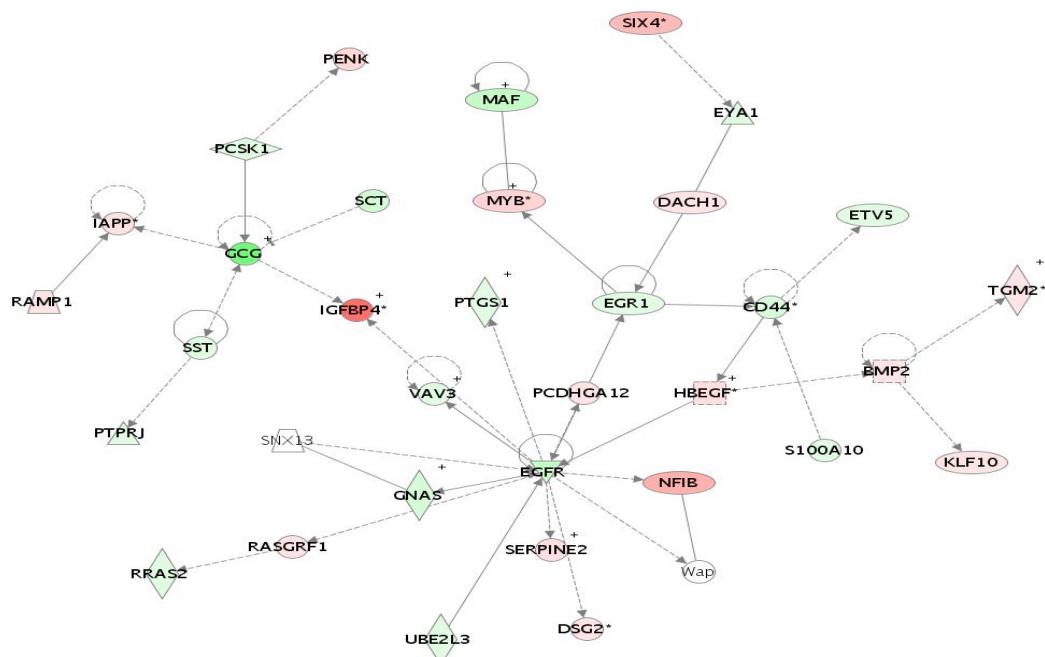
Network	Score	Focus genes	Top functional categories
<b>H1A (+/-)/RIP-Tag versus H1A (+/+)/RIP-Tag beta-cell line</b>			
1	60	35	Cellular Growth and Proliferation, Cell Death, Cancer
2	25	20	Cellular Assembly and Organization, Cell Cycle, DNA Replication, Recombination, and Repair
3	19	17	Cellular Movement, Cellular Assembly and Organization, Nervous System Development and Function
4	18	16	Gene Expression, Cancer, Cell Cycle
5	18	16	Cancer, Cellular Growth and Proliferation, Immunological Disease
6	18	16	Cancer, Cellular Growth and Proliferation, DNA Replication, Recombination, and Repair
7	18	16	Cellular Growth and Proliferation, Renal and Urological Disease, Drug Metabolism
<b>H1A (-/-)/RIP-Tag versus H1A (+/+)/RIP-Tag beta-cell line</b>			
1	49	33	Cellular Growth and Proliferation, Cell Signaling, Nucleic Acid Metabolism
2	20	19	Cell Death, Dermatological Diseases and Conditions, Connective Tissue Disorders
3	17	17	Cell Cycle, Cellular Movement, Cellular Development
4	15	16	Molecular Transport, RNA Trafficking, Cell-To-Cell Signaling and Interaction
5	14	15	Carbohydrate Metabolism, Small Molecule Biochemistry, Cardiovascular System Development and Function
6	14	15	Cellular Movement, Cancer, Reproductive System Disease
7	14	15	Hematological System Development and Function, Cardiovascular Disease, Hematological Disease
8	14	15	Cell Signaling, Cellular Assembly and Organization, Cellular Compromise

## Supplemental Data 2

A



B



### Supplemental Data 3

*Real-time PCR analysis for SERPINE2, IGFBP4, TGFB1 and IGFBP7 in HNF-1 $\alpha$*

	H1A (+/+)–Tag	H1A (+/-)–Tag	H1A (−/−)–Tag
n	4	4	4
SERPINE2	1.0±0.1	13.4±2.5*	13±0.8*
IGFBP4	1.0±0.1	4.9±0.6*	16.8±0.6*#
TGFB1	1.0±0.1	150.6±18.*	28.0±3.9*#
IGFBP7	1.0±0.1	4.1±0.1*	2.3±0.1*#

(+/+)/RIP-Tag, HNF-1 $\alpha$  (+/-)/RIP-Tag and HNF-1 $\alpha$  (−/−)/RIP-Tag beta-cell lines.

\* $P < 0.05$  compared with H1A (+/+)–Tag

# $P < 0.05$  compared with H1A (+/-)–Tag

## Supplemental Data 4

*Changes of beta-cell related genes in HNF-1 $\alpha$  (+/+)/RIP-Tag, HNF-1 $\alpha$  (+/-)/RIP-Tag and HNF-1 $\alpha$  (-/-)/RIP-Tag beta-cell lines.*

Accession	gene	fold change			
		+/-	+/+	-/-	+/-
<b>hormone and growth factors</b>					
NM_008386	insulin I		1.26		1.23
NM_008387	insulin II		1.07		1.01
AF276754	glucagon	0.25	↓	0.07	↓
<b>transcription factors</b>					
AK020261	Pdx-1		0.87		0.99
NM_010446	forkhead box A2 (Foxa-2)		1.97	↑	1.63 ↑
AF357883	Nkx-6.1		1.18		1.25
NM_010919	Nkx-2.2		1.60	↑	1.39
AF031150	Pax-4		1.67	↑	1.12
BC011272	Pax-6		1.12		1.21
AK008017	Ngn-3		1.00		1.15
BQ176915	isl-1	0.64	↓	0.92	
BM116592	NeuroD-1		1.29		0.87
<b>glucose metabolism</b>					
NM_013631	liver pyruvate kinase (L-Pk)		0.99		0.87
NM_031197	glucose transporter type 2 (Glut2)		1.13		0.93
BC024112	aldolase B		0.87		0.87
BC011139	glucokinase		1.58	↑	1.56 ↑
<b>others</b>					
NM_010512	insulin-like growth factor 1 (IGF-1)		1.09		1.06
NM_009864	E-cadherin		1.60	↑	1.02
AI314694	Tmem27		1.17		0.65 ↓

Abbreviations: +/+, H1A(+/+)/RIP-Tag; +/-, H1A(+/-)/RIP-Tag; -/-, H1A(-/-)/RIP-Tag

A single arrow and a double arrow show changes of 1.5~ 2-fold or > 2-fold, respectively.