

Smad4 suppresses the tumorigenesis and aggressiveness of neuroblastoma through repressing the expression of heparanase

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Supplementary Figure Legends

Supplementary Figure S1 Mining the transcription factor crucial for HPSE expression. **a**, over-lapping analysis revealing Smad4, LEF1, and PPARG as potential transcription factors of HPSE in NB. **b**, real-time quantitative RT-PCR and dual-luciferase assays showing the *PPARG* levels and *HPSE* promoter activity in NB cells. **c**, the correlation between *Smad4*, *LEF1*, and *HPSE* levels in NB or neuroblastic tumor tissues derived from R2 microarray analysis and visualization platform (<http://r2.amc.nl>).

Supplementary Figure S2 Effects of TGF-β on HPSE expression in NB cells. **a**, nuclear run-on assay indicating the nascent transcript levels of *HPSE* in NB cells stably transfected with

mock, *Smad4*, sh-Scb, or sh-Smad4. **b**, dual-luciferase assay indicating the activity of Smad4 response element reporter (pSBE4-luc) and *HPSE* promoter reporter in NB cells stably transfected with sh-Scb or sh-Smad4, and those treated with TGF- β . * $P<0.01$ vs. mock or sh-Scb.

Supplementary Figure S3 Effects of BMP-2 on HPSE expression in NB cells. **a** and **b**, immunofluorescence and western blot assays indicating the localization of Smad4 and expression of HPSE in IMR32 and BE(2)-C cells stably transfected with mock or *Smad4*, and those treated with TGF- β or LY364947. **c**, Co-IP and western blot assays showing the interaction of Smad4 with p-Smad1 or p-Smad2 in IMR32 cells stably transfected with mock, *Smad4*, *Smad4 (D351H)*, or *Smad4 (R361H)*. **d**, **e**, and **f**, western blot, real-time quantitative RT-PCR, and dual-luciferase assays indicating the expression levels and promoter activity of *HPSE* in NB cells stably transfected with mock, *Smad4*, *Smad4 (D351H)*, or *Smad4 (R361H)*. **g**, **h**, and **i**, dual-luciferase and real-time quantitative RT-PCR assays showing the activity of Smad4 response element reporter (pSBE4-luc) or BMP/Smad transcriptional reporter (pGL3-BRE-Luc), and *HPSE* transcript levels in NB cells transfected with activin A-neutralizing antibody, BMP-2, or LDN-193189, and those transfected with mock or *Smad4*. * $P<0.01$ vs. mock; $^{\Delta}P<0.01$ vs. control untreated with BMP-2 or LDN-193189.

Supplementary Figure S4 Effects of Smad4 and LEF1 on the HPSE activity, growth, invasion, and angiogenesis of NB cells. Dual-luciferase (**a**), colorimetric (**b**), MTT colorimetric (**c**), soft agar (**d**), matrigel invasion (**e**), tube formation (**f**) assays showing the cellular *HPSE* promoter activity, HPSE enzymatic activity, viability, growth, invasion, and angiogenesis capability of NB cells stably transfected with empty vector (mock) or *Smad4*, and those

co-transfected with *LEF1*. * $P<0.01$ vs. mock.

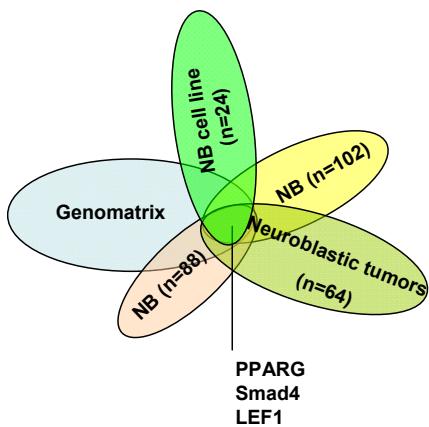
Supplementary Figure S5 Expression of Smad4, LEF1, and HPSE in public databases. **a** and **b**, the datasets derived from Oncogenomics (<https://pob.abcc.ncifcrf.gov/cgi-bin/JK>) and TARGET (<https://target.nci.nih.gov/dataMatrix/>) indicating the copy number and mutation frequency of *Smad4* and *HPSE* in NB. **c** and **d**, the expression of *Smad4*, *LEF1*, and *HPSE* in NB tissues derived from R2 microarray analysis and visualization platform (<http://r2.amc.nl>).

Supplementary Figure S6 Patients' survival analysis. Kaplan–Meier survival plots of NB cases derived from R2 microarray analysis and visualization platform (<http://r2.amc.nl>) showing the survival probability of patients with high or low expression of Smad4 or HPSE.

Supplementary Figure S7 Interaction between Smad4 and LEF1 in NB cells. Co-IP and western blot assays showing the interaction between Smad4 and LEF1 in IMR32 cells transfected with different Myc-tagged *Smad4* truncates and FLAG-tagged *LEF1* construct.

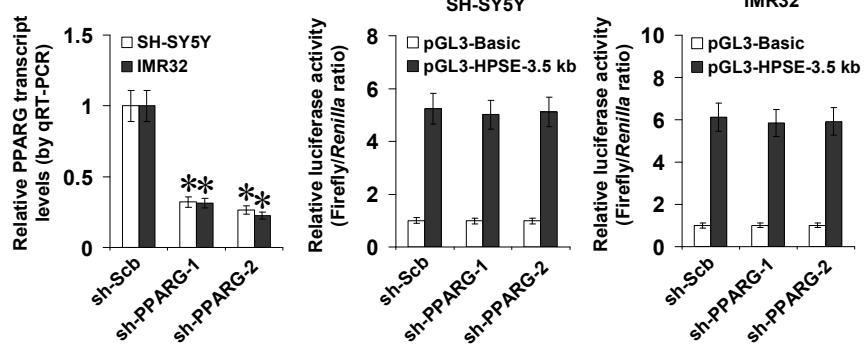
Supplementary Figure S1

a

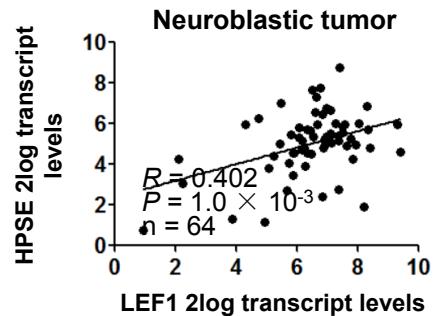
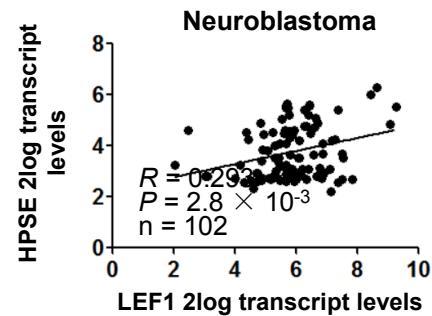
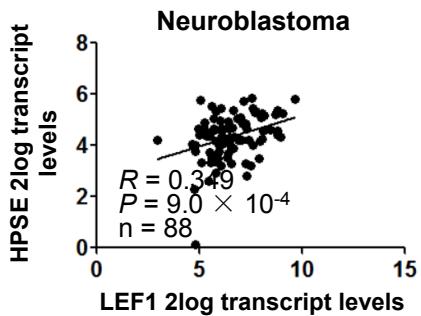
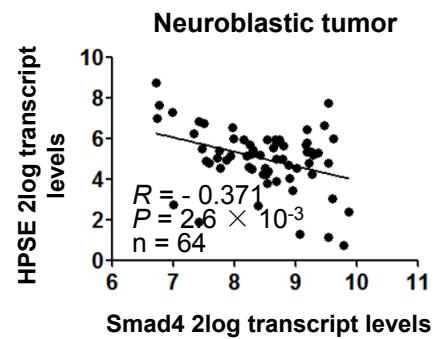
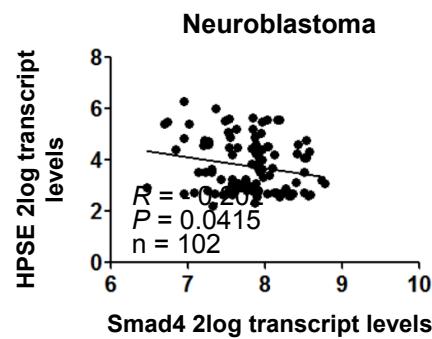
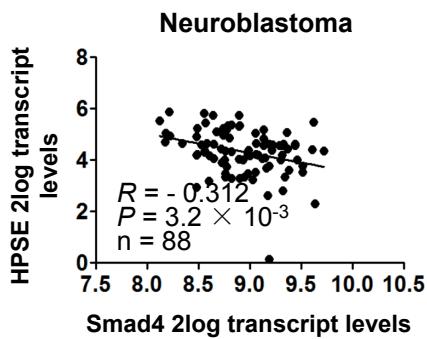


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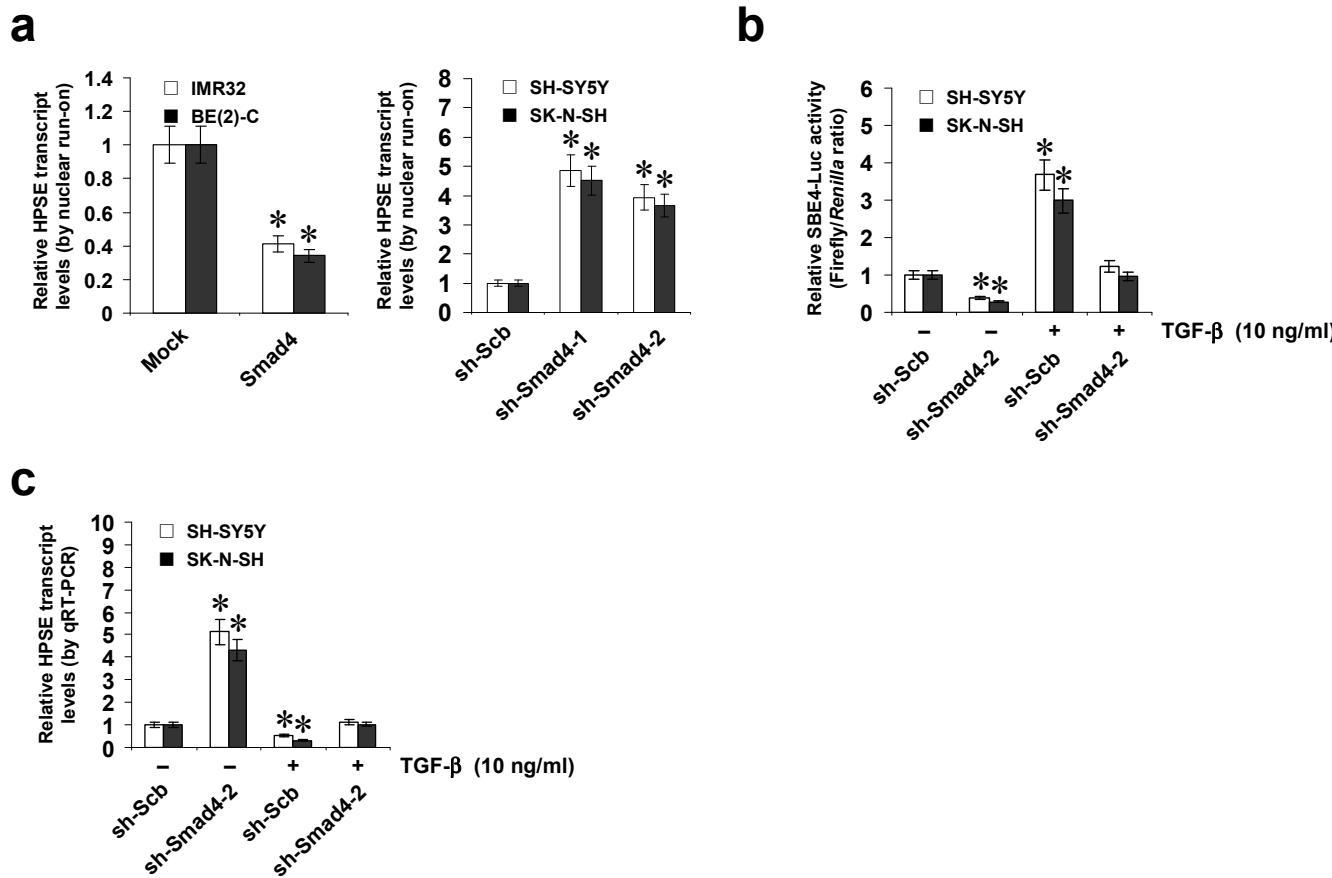
Correlation between PPARG and HPSE expression		
R2 datasets	R -value	P -value
NB (n=102)	0.314	0.001326016
NB (n=88)	0.249	0.019311638
Neuroblastic tumor (n=64)	0.411	0.000752421
NB cell lines (n=24)	0.487	0.015850924



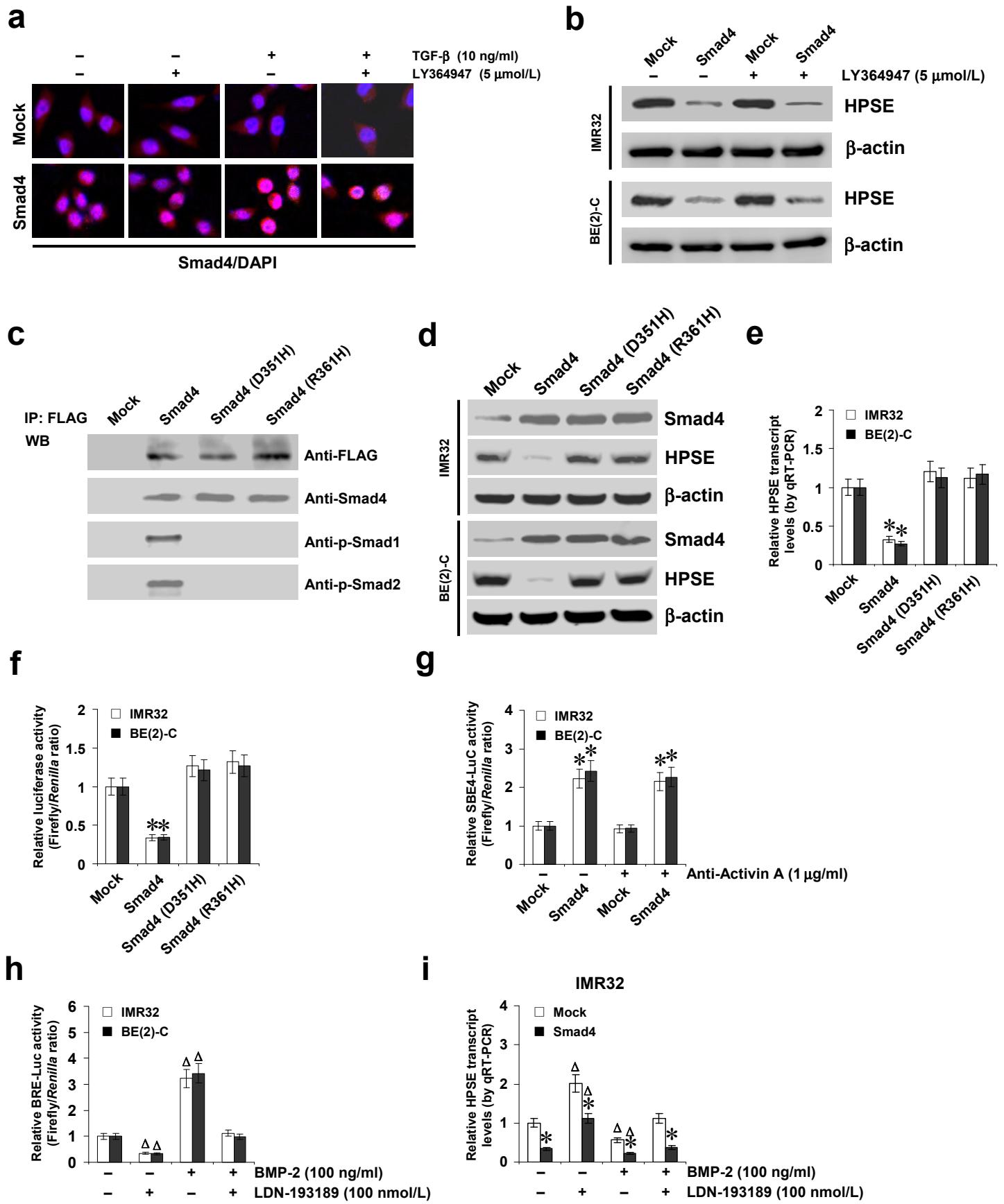
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Supplementary Figure S2

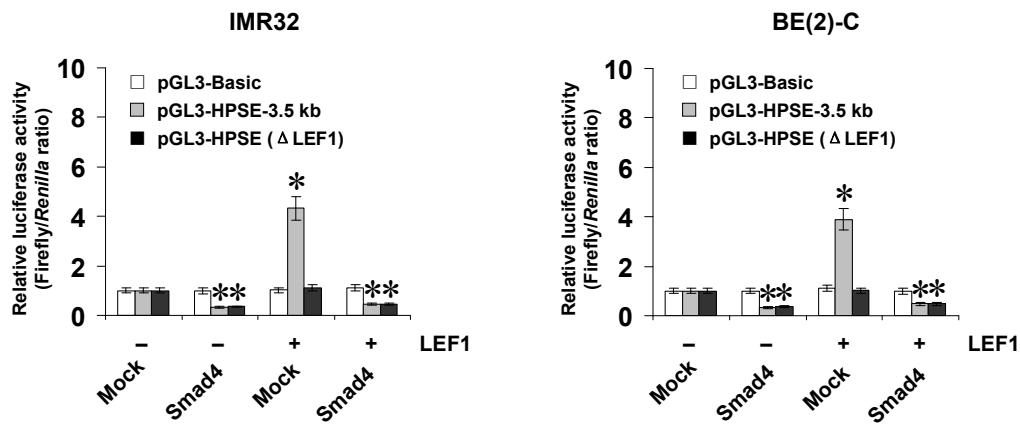


Supplementary Figure S3

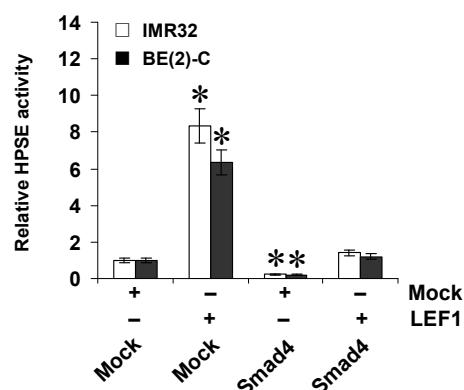


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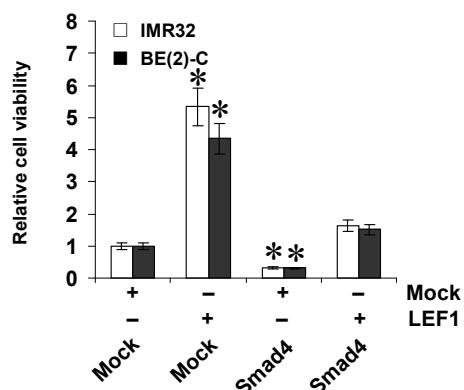
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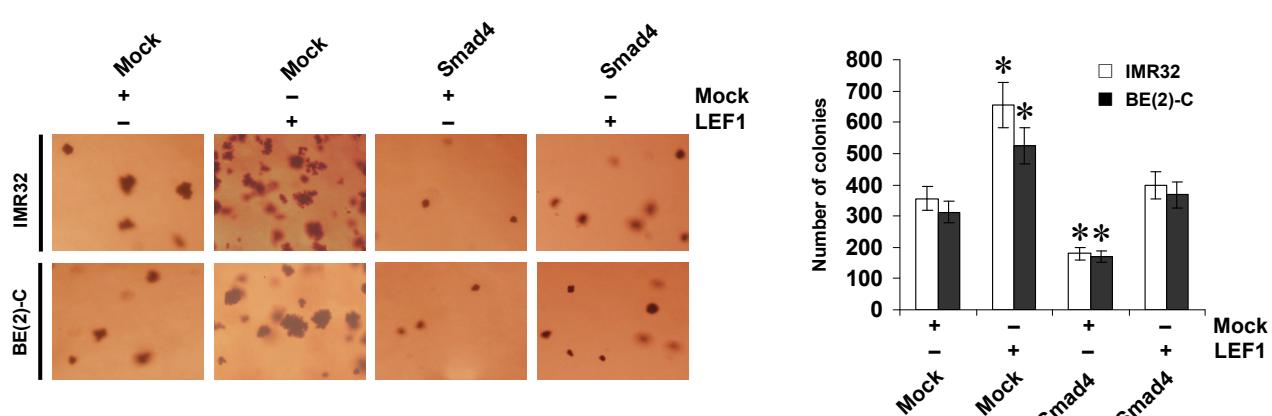
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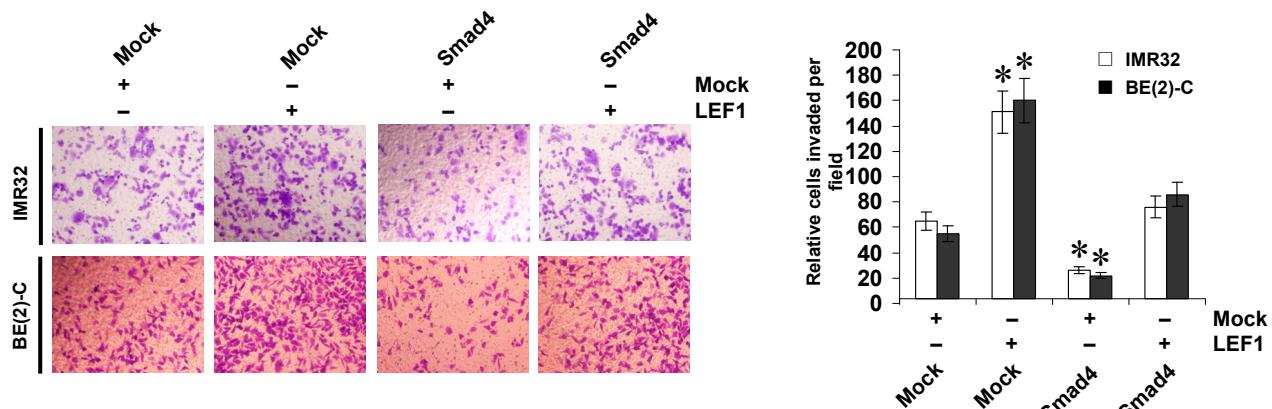
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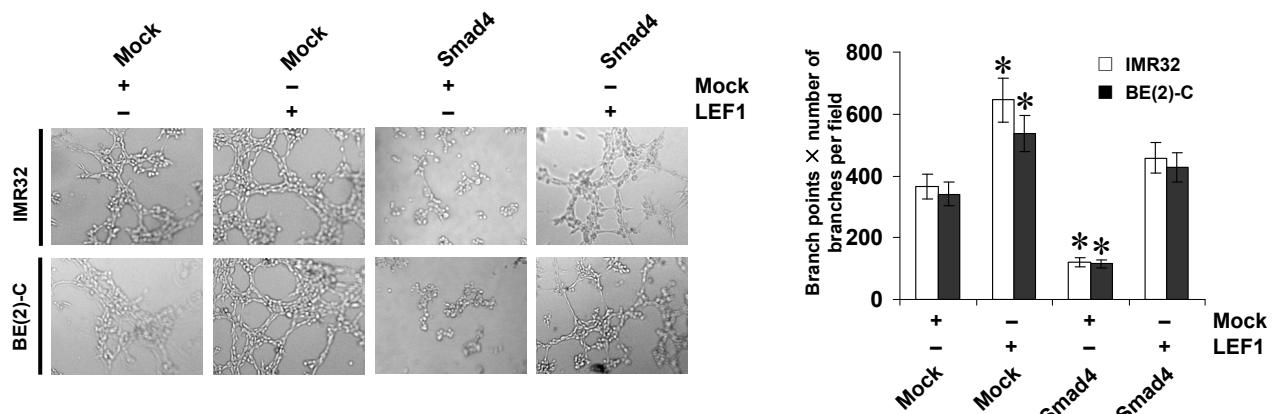
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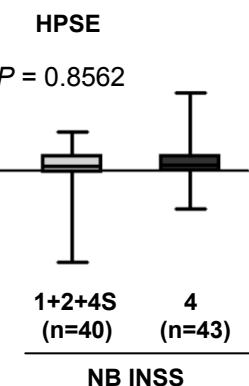
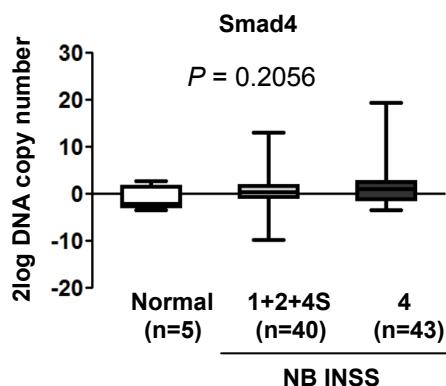


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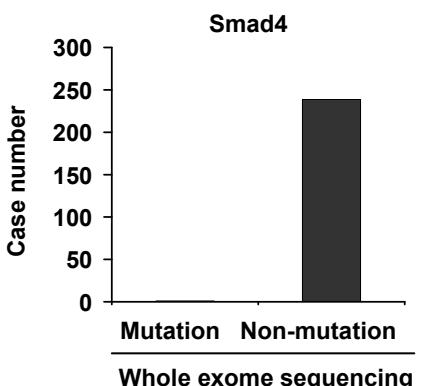


Supplementary Figure S5

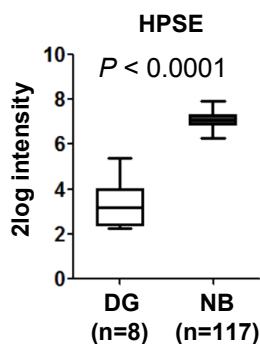
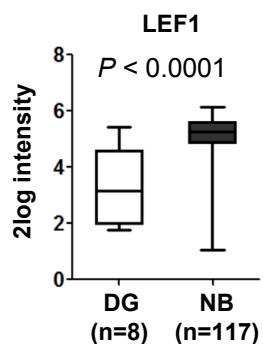
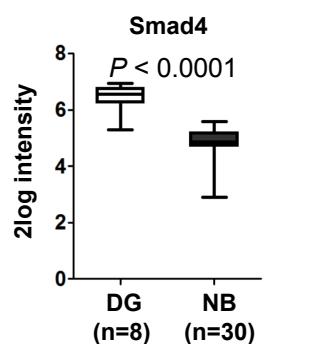
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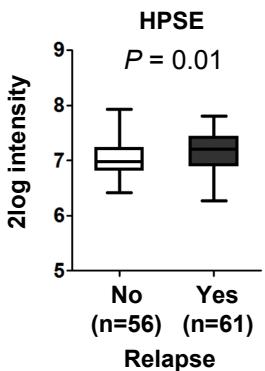
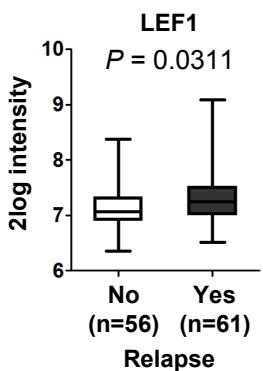
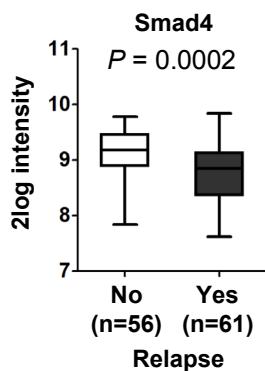
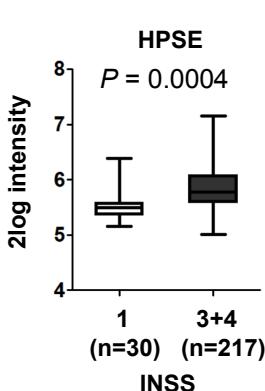
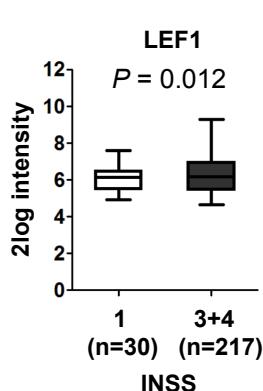
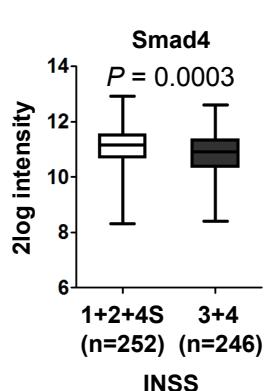
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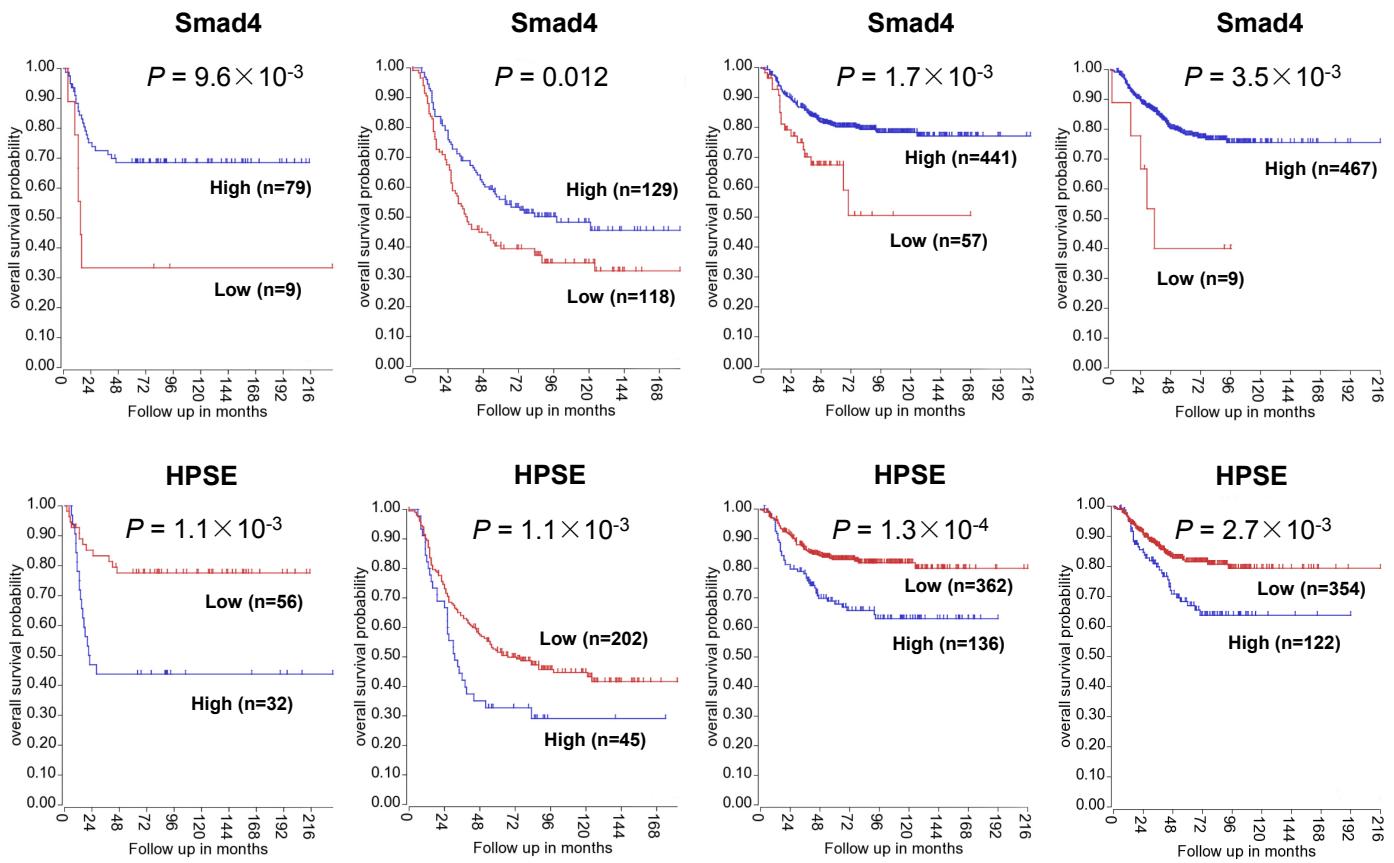
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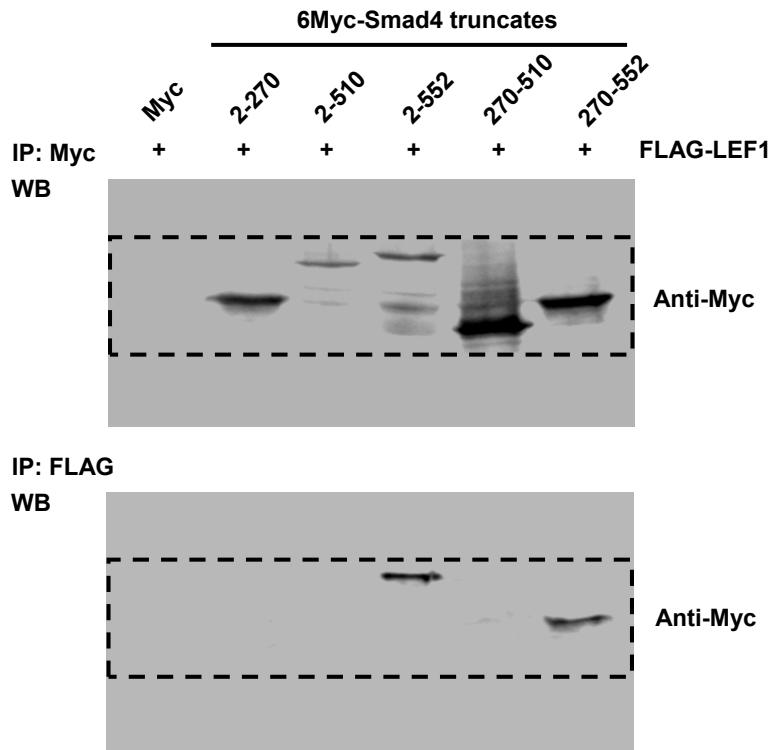
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Supplementary Figure S6



Supplementary Figure S7



Supplementary Table S1 Smad4 expression in human NB tissues

Group	Total number	Smad4 expression				Positive rates (%)	P-Value
		-	+	++	+++		
Age							
<1 year	20	12	3	3	2	40.0	0.778
≥1 year	22	14	5	2	1	36.4	
Differentiation							
Well differentiated	8	0	2	3	3	100.0	
Poorly differentiated	28	21	5	2	0	25.0	<0.001
Undifferentiated	6	5	1	0	0	16.7	
MKI							
<200	17	6	3	5	3	64.7	0.002
>200	25	20	5	0	0	20.0	
INSS stages							
Stage 1-2	14	4	3	4	3	71.4	
Stage 3-4	20	16	4	0	0	20.0	0.016
Stage 4S	8	6	1	1	0	25.0	

Smad4, SMAD family member 4; MKI, mitosis karyorrhexis index; INSS, international neuroblastoma staging system.

Supplementary Table S2 Correlation among the expression of Smad4, LEF1, and HPSE

	Smad4 expression				LEF1 expression			
	Low	High	R-value	P-value	Low	High	R-value	P-value
HPSE expression								
Low	4	6	- 0.583	<0.001	8	2	0.553	< 0.001
High	30	2			6	26		

Smad4, SMAD family member 4; LEF1, lymphoid enhancer-binding factor 1; HPSE, heparanase; Pearson's correlation coefficient was applied to determine the expression correlation.

Supplementary Table S3 Oligonucleotide sets used for constructs, inhibitors and short hairpin RNAs

Oligo Set	Sequences
pcDNA3.1-Smad4	5'-CGCGGATCCATGGACAATATGTCTATTACG-3' (sense); 5'-CCGCTCGAGTCAGTCTAAAGGTTGGGTC-3' (antisense)
pEF-FLAG-Smad4 (D351H)	5'-TGTTACTGTTCATGGATACTGGACCCCTCTGGAGGAGATCG-3' (sense); 5'-CACGTATCCATGAACAGTAACAATAGGGCAGCTGAAGGAAC-3' (antisense)
pEF-FLAG-Smad4 (R361H)	5'-GGAGGAGATCACTTTGTTGGTCAACTCTCCAATGTCCAC-3' (sense); 5'-CCAAACAAAAGTGATCTCCTCCAGAAGGGTCCACGTATCCAT-3' (antisense)
pcDNA3.1-HPSE	5'-CGGGTACCATGCTGCTGCGCTCGAACGCCTG-3' (sense); 5'-CCGCTCGAGTCAGATGCAAGCAGCAACTTGG-3' (antisense)
pCMV-3Tag-1A-LEF1	5'-CGCGGATCCATGCCCAACTCTCCGGAGGAGGT-3' (sense); 5'-CCCAAGCTTCAGATGTAGGCAGCTGTCATTCT-3' (antisense)
pGL3-HPSE mut (△ Smad4)	5'-TGTGAGGCCGCTGAGGGCAGATCGCGAGGTCAAGGAGATTGAGACC-3' (sense); 5'-GCGATCTGCCCTCAGCGGCCCTCACAAAATGCTAGGATTGCAGGCA-3' (antisense)
pGL3-HPSE mut (△ LEF1)	5'-TTTGGGTGGCGACTCTCTTCCAGCTGCAGTTAGCGTATGCTG-3' (sense); 5'-CTGGAAAGAGAGTCGCCACCCAAAATTCTTATCAACTTCTCATT-3' (antisense)
sh-Scb	5'-CCGGCGAACGATCGAGTAAACGGACTCGAGTCCGTTACTCGATCGTCGCTTT T-3' (sense); 5'-AATTCAAAAAGCGAACGATCGAGTAAACGGACTCGAGTCCGTTACTCGATCGTC GC-3' (antisense)
sh-Smad4-1	5'-CCGGGCAGACAGAAACTGGATTAAACTCGAGTTAACAGTTCTGTCTGCTTT T-3' (sense); 5'-AATTCAAAAAGCAGACAGAAACTGGATTAAACTCGAGTTAACAGTTCTGTCTG C-3' (antisense)
sh-Smad4-2	5'-CCGGCCTGAGTATTGGTGTCCATTCTCGAGAACATGGAACACCAAAACTCAGGTTT T-3' (sense); 5'-AATTCAAAAACCTGAGTATTGGTGTCCATTCTCGAGAACATGGAACACCAAAACTCAG G-3' (antisense)
sh-PPARG-1	5'-CCGGCGCCTGCATCTCACCTTATCTCGAGATAAGGTGGAGATGCAGGCGCTT TT-3' (sense); 5'-AATTCAAAAAGCGCCTGCATCTCACCTTATCTCGAGATAAGGTGGAGATGCAGGC GC-3' (antisense)
sh-PPARG-2	5'-CCGGCGAACGCTTATCTATGACAGACTCGAGTCTGTCATAGATAAGCTCGCTTT T-3' (sense); 5'-AATTCAAAAAGCGAACGCTTATCTATGACAGACTCGAGTCTGTCATAGATAAGCTCG C-3' (antisense)

Smad4, SMAD family member 4; HPSE, heparanase; LEF1, lymphoid enhancer-binding factor 1; sh-Scb, scramble short hairpin RNA; PPARG, peroxisome proliferator-activated receptor gamma.

Supplementary Table S4 Primer sets used for qRT-PCR, nuclear run-on, and ChIP

Primer set	Primers	Sequence	Product size (bp)	Application
HPSE	Forward	5'-GAATGGACGGACTGCTAC-3'	261	qRT-PCR, nuclear run-on
	Reverse	5'-CCAAAGAATACTTGCCTCA-3'		
Smad4	Forward	5'-TGCCTCACCAACCAAAACGG-3'	255	qRT-PCR
	Reverse	5'-CCAAACAAAAGCGATCTCCTCC-3'		
LEF1	Forward	5'-AAATAAAGTGCCGTGGTGC-3'	280	qRT-PCR
	Reverse	5'-TGAGGATGGTAGGGTTGCC-3'		
PPARG	Forward	5'-GAACGACCAAGTAACCTCCCTC-3'	166	qRT-PCR
	Reverse	5'-GGGCTCCATAAAGTCACCAAAA-3'		
β -actin	Forward	5'-ATCTACGAGGGTATGCC-3'	227	qRT-PCR
	Reverse	5'-TAGCTCTTCCAGGGAG-3'		
HPSE Set 1 (-2347/-2148)	Forward	5'-AAATTGGTATGACTGGGCATGG-3'	200	ChIP
	Reverse	5'-TCAGCCTCCCGAGTAGCTGGACTA-3'		
HPSE Set 2 (-1505/-1268)	Forward	5'-ACCTTGGCCTCCAAAATGCTGG-3'	238	ChIP
	Reverse	5'-GGCATGTGATCATGTTGGTGCTTA-3'		
HPSE Set 3 (-673/-476)	Forward	5'-ACACTTGTATGGGGCTTCAG-3'	198	ChIP
	Reverse	5'-CTCTGGCAAGGCAGTTCTCAG-3'		

HPSE, heparanase; Smad4, SMAD family member 4; LEF1, lymphoid enhancer-binding factor 1; PPARG, peroxisome proliferator-activated receptor gamma.