

## DDR2 controls the epithelial-mesenchymal-transition-related gene expression via c-Myb acetylation upon matrix stiffening

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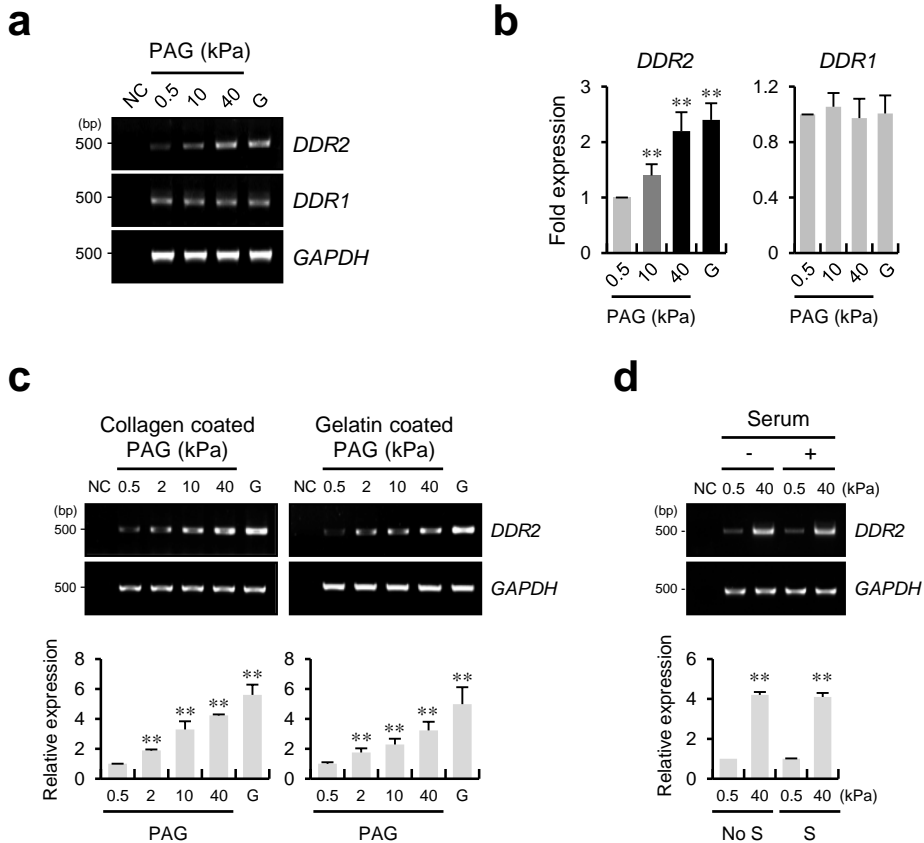
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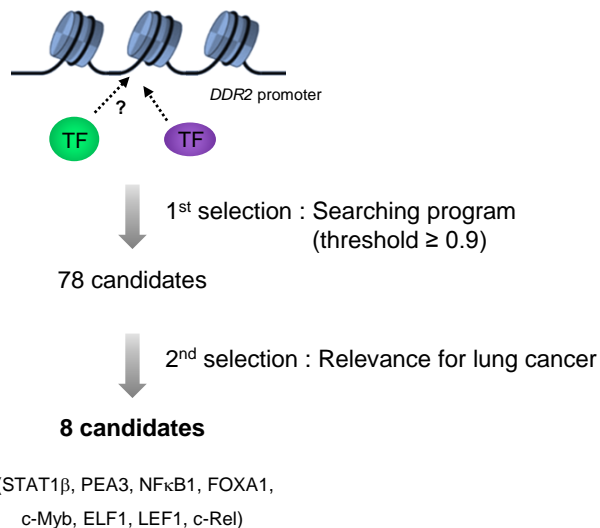
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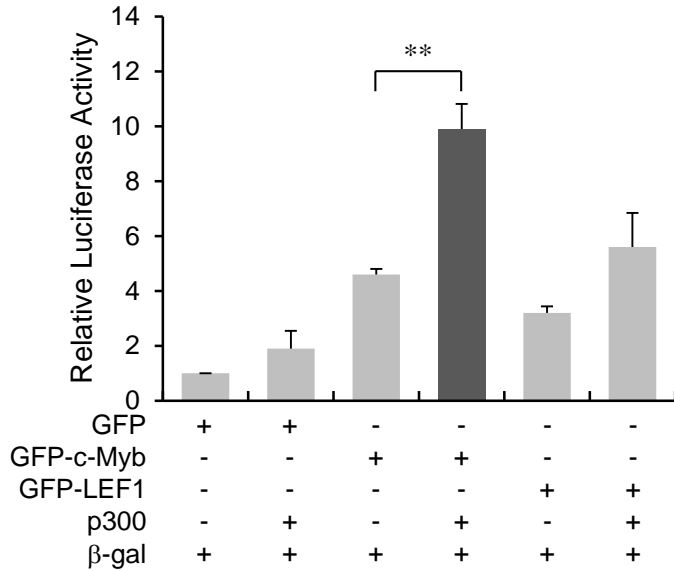
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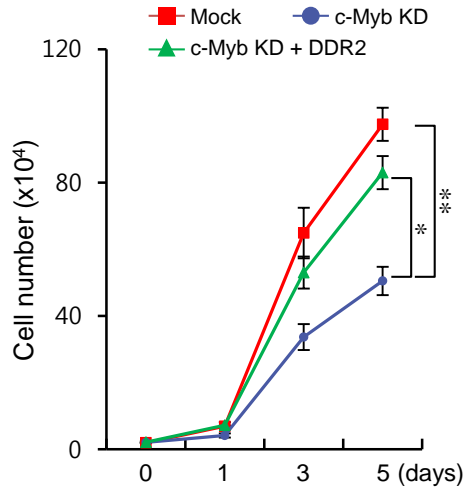
**Supplementary Figure 1.** Expression of *DDR2* transcripts is dependent on matrix rigidity. (a,b) RT-PCR (a) and qRT-PCR (b) analyses of expression levels of *DDR1*, *DDR2*, and *GAPDH* in H1299 cells incubated on a PAG of various stiffness or on glass (G). (c) RT-PCR analysis of the expression levels of *DDR2* and *GAPDH* in H1299 cells incubated on a PAG of various stiffness or glass coated with collagen or gelatine. (d) RT-PCR analysis of expression levels of *DDR2* and *GAPDH* in H1299 cells incubated on a PAG in the absence or presence of serum. Band densities were measured and normalised to *GAPDH* expression levels under the same conditions. NC, negative control. Data represent the mean of three independent experiments  $\pm$  SEM. \*\*  $P < 0.01$ .



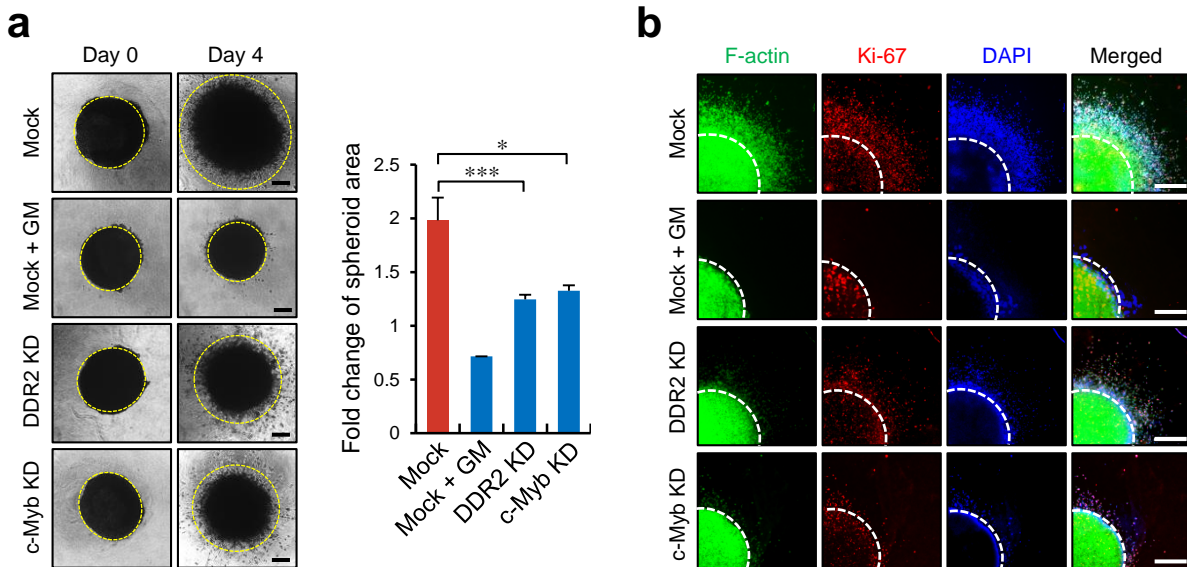
**Supplementary Figure 2.** The experimental scheme employed to analyse transcription factors that regulate *DDR2* promoter activity in lung cancer. Seventy-eight candidates are listed in Supplementary Table 1, and eight candidates (STAT1 $\beta$ , PEA3, NF $\kappa$ B1, FOXA1, c-Myb, ELF1, LEF1, and c-Rel) were eventually selected.



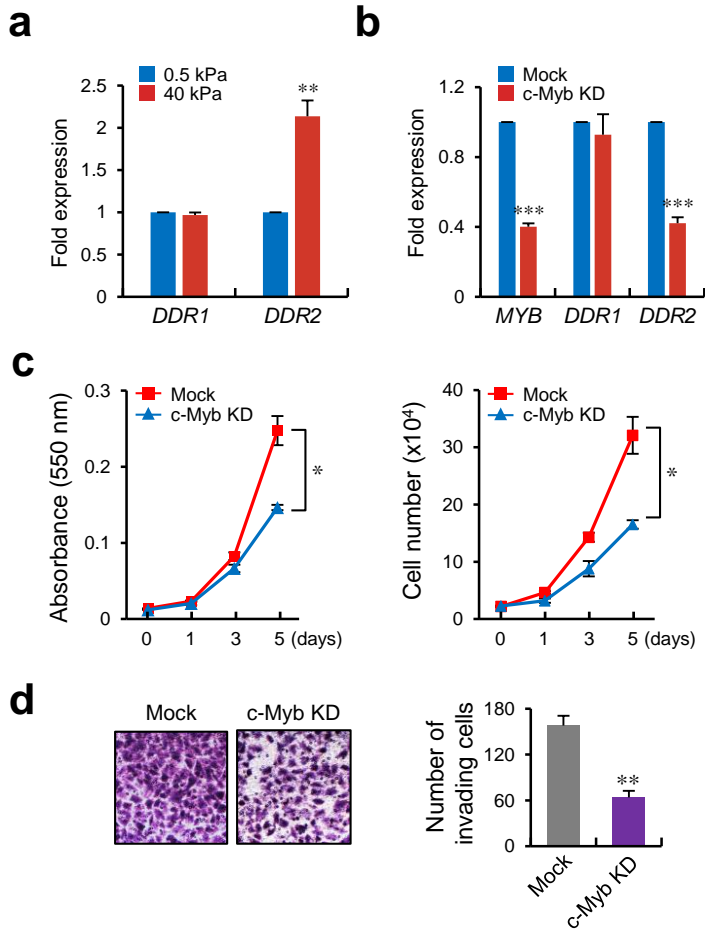
**Supplementary Figure 3.** Coexpression of c-Myb with p300 induces *DDR2* promoter activity. Relative expression of *DDR2* WT promoter-driven luciferase reporters in c-Myb-, LEF1-, or p300-overexpressing H1299 cells. Data represent the mean of three independent experiments  $\pm$  SEM. \*\* $P < 0.01$ .



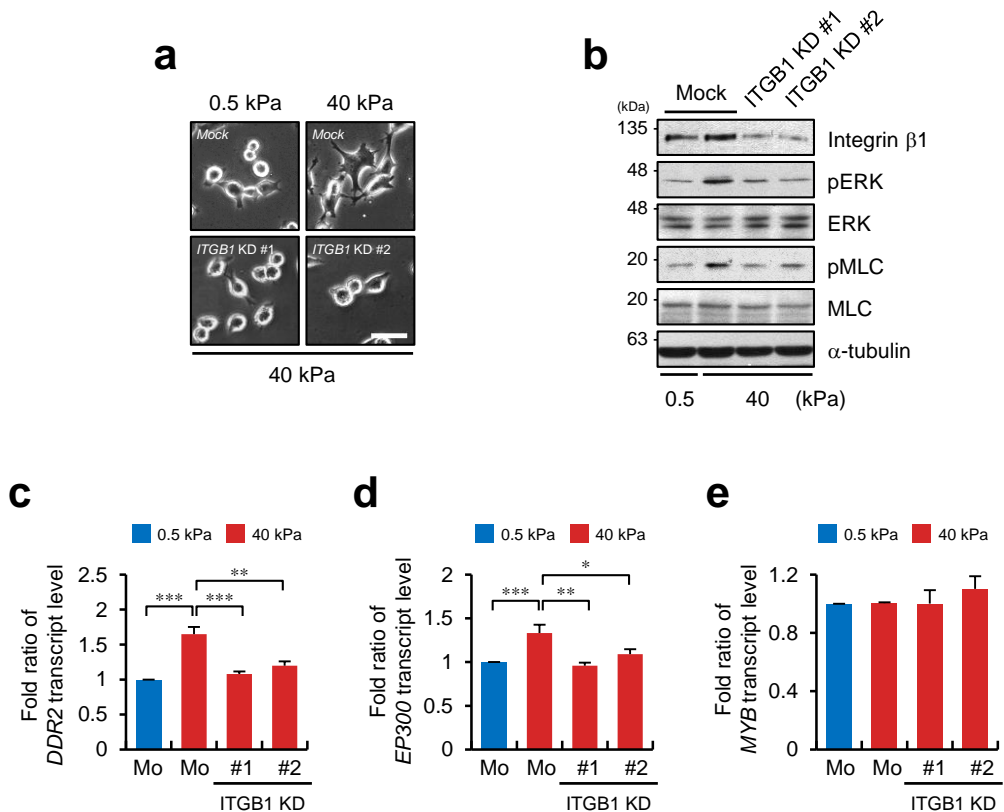
**Supplementary Figure 4.** Inhibition of proliferation in c-Myb knockdown H1299 cells is reversed by DDR2 overexpression. Quantification of proliferation of H1299 cells incubated on a stiff matrix for the indicated periods. The cells were trypsinized and counted. Data represent the mean of three independent experiments  $\pm$  SEM. \* $P < 0.05$ , \*\* $P < 0.01$ .



**Supplementary Figure 5.** A knockdown of c-Myb and DDR2 inhibits cancer invasion in a 3D environment. H1299 cells expressing mock (with or without 5  $\mu$ M GM6001), anti-DDR2, or anti-c-Myb shRNA were incubated for 2 days to form spheroids as described in the *Methods* section. Collagen-embedded spheroids were further incubated for 4 days, fixed, and stained with phalloidin, an anti-Ki-67 antibody, and DAPI. The spheroids were examined by phase contrast microscopy (**a**) and fluorescence microscopy (**b**). GM: GM6001. Scale bars: 200  $\mu$ m (a) and 500  $\mu$ m (b). Data represent the mean of three independent experiments  $\pm$  SEM. \* $P$  < 0.05, \*\*\* $P$  < 0.001.

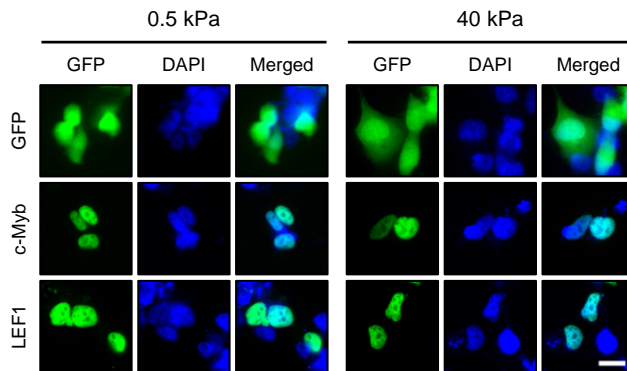


**Supplementary Figure 6.** The knockdown of c-Myb inhibits proliferation and invasiveness of A549 cells. **(a)** qRT-PCR analysis of *DDR1* and *DDR2* on 0.5- and 40-kPa PAGs in A549 cells. **(b)** qRT-PCR of *MYB*, *DDR1* and *DDR2* on 40 kPa PAGs in mock shRNA and c-Myb KD A549 cells. **(c)** Quantification of cell proliferation in mock or anti-c-Myb shRNA-expressing A549 cells. Graphs show absorbance of crystal violet (left) and the number of counted cells (right). **(d)** Quantification of invasion. Data represent the mean of three independent experiments  $\pm$  SEM. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.005$ .



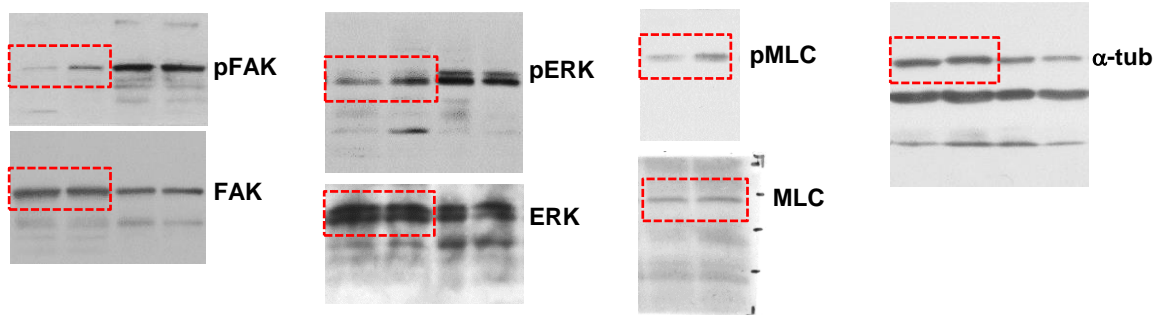
**Supplementary Figure 7.** Cellular contractility on a stiff matrix regulates the expression of *DDR2* and *EP300* via integrin  $\beta$ 1. **(a)** A phase contrast image of H1299 cells transfected with mock siRNA or anti-integrin  $\beta$ 1 siRNA (#1 and #2) on 0.5- and 40-kPa PAGs. Scale bar, 50  $\mu$ m. **(b)** Western blot analysis of mock or anti-integrin  $\beta$ 1 siRNA-transfected H1299 cells. **(c,d,e)** qRT-PCR analysis of expression of *DDR2* (c), *EP300* (d), and *MYB* (e) in mock and integrin  $\beta$ 1 knockdown cells. qRT-PCR values were normalised to *GAPDH*. Mo: Mock. Data represent the mean of three independent experiments  $\pm$  SEM. \* $P$ < 0.05, \*\* $P$ < 0.01, \*\*\* $P$ < 0.005.



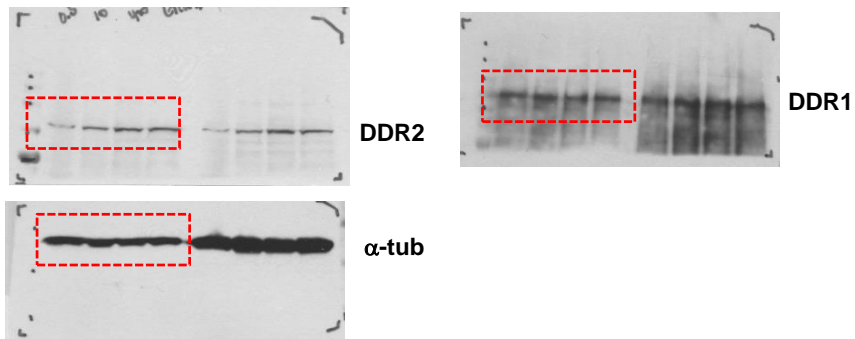


**Supplementary Figure 8.** Cellular localisation of c-Myb and LEF1 does not depend on matrix stiffness. Fluorescent images of H1299 cells transfected with plasmids encoding GFP, GFP-c-Myb, or GFP-LEF1 and incubated on a PAG. Nuclei were stained with DAPI. Scale bar, 10  $\mu$ m.

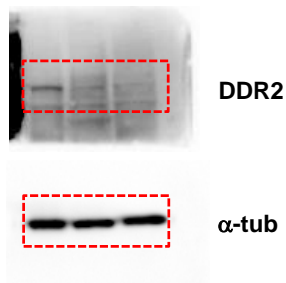
**Figure 1c**



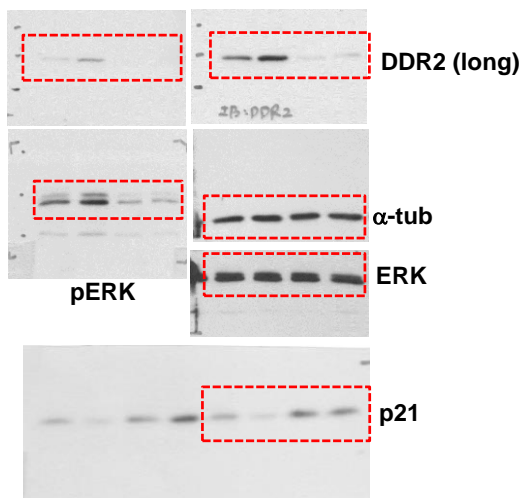
**Figure 2a**



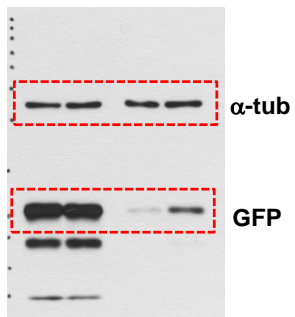
**Figure 2b**



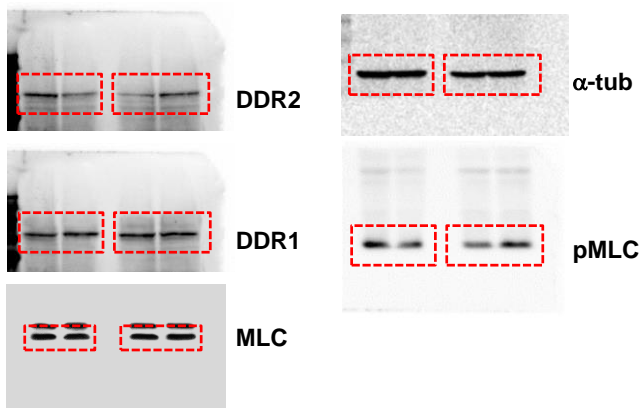
**Figure 2d**



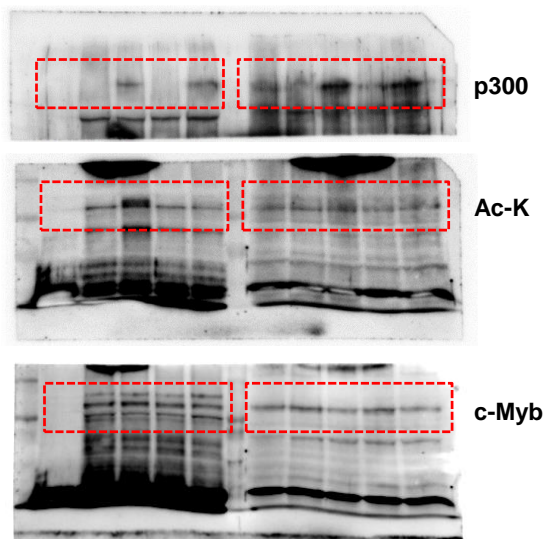
**Figure 3b**



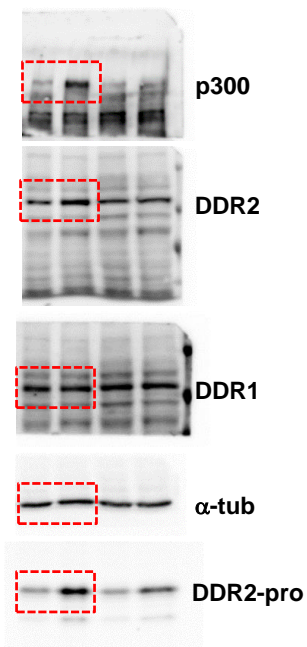
**Figure 4e and 4g**



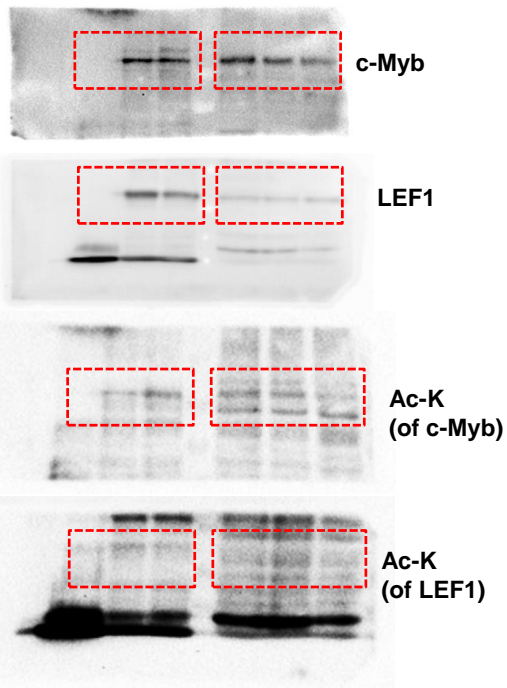
**Figure 5d**



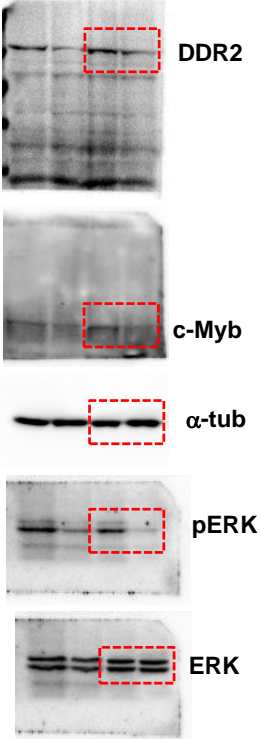
**Figure 5f**



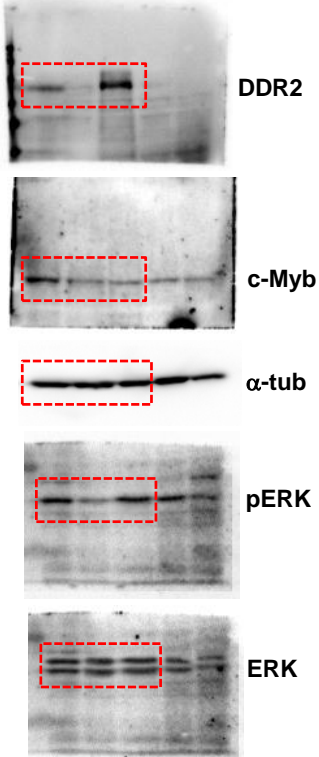
**Figure 5a**



**Figure 6c**



**Figure 6e**



**Supplementary Figure 9.** Full length blots. Red dotted lines show the cropping locations.

**Supplementary Table 1.** Seventy-eight transcription factors that are predicted to bind to the *DDR2* promoter region.

No	Name	Description
1	FOXP3	Forkhead box P3
2	RXR $\alpha$	Retinoid X receptor alpha
3	NR3C1	Nuclear receptor subfamily 3, group C, member 1 (glucocorticoid receptor)
4	RAR $\beta$	Retinoic acid receptor beta
5	GR $\alpha$	Glucocorticoid receptor alpha
6	C/EBP $\alpha$	CCAAT/enhancer binding protein alpha
7	C/EBP $\beta$	CCAAT/enhancer binding protein beta
8	POU2F2	POU class 2 homeobox 2
9	c-Myb	Transcription factor c-Myb
10	FOXA1	Forkhead box A1
11	FOXA2	Forkhead box A2
12	XBP-1	X-box binding protein 1
13	TFIID	Transcription factor II D
14	c-Ets-1	V-ets erythroblastosis virus E26 oncogene homolog 1
15	c-Ets-2	V-ets erythroblastosis virus E26 oncogene homolog 2
16	IRF-1	Interferon regulatory factor 1
17	NFATC1	Nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 1
18	NFATC2	Nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 2
19	AP-2 $\alpha$ A	Transcription factor AP-2-alpha A
20	p53	Tumor protein p53
21	AR	Androgen receptor
22	GATA-1	GATA binding protein 1 (globin transcription factor 1)
23	GATA-2	GATA binding protein 2 (globin transcription factor 2)
24	STAT1 $\beta$	Signal transducer and activator of transcription 1beta
25	STAT4	Signal transducer and activator of transcription 4
26	STAT5A	Signal transducer and activator of transcription 5a
27	ATF3	Activating transcription factor 3
28	NFI/CTF	Nuclear factor I-CCAAT-binding transcription factor
29	NF-1	Neurofibromin 1
30	SRY	Sex-determining region Y
31	TCF-4E	Transcription factor 7 like 2, T cell specific, HMG box
32	VDR	Vitamin D (1,25- dihydroxyvitamin D3) receptor

33	PXR-1	Pregnane X receptor
34	c-Jun	Jun proto-oncogene
35	c-Fos	FBJ murine osteosarcoma viral oncogene homolog
36	HNF-1A	HNF1 homeobox A
37	HNF-1B	HNF1 homeobox B
38	YY1	YY1 transcription factor
39	POU2F1	POU class 2 homeobox 1
40	TEAD2	TEA domain family member 2
41	RBP-Jκ	Recombination signal binding protein for immunoglobulin kappa J region
42	PEA3	Phosphatidylinositol-4-phosphate 5-kinase and related FYVE finger-containing proteins signal transduction mechanisms
43	COUP-TF1	Nuclear receptor subfamily 2, group F, member 1
44	LEF-1	Lymphoid enhancer-binding factor 1
45	NF-κB1	Nuclear factor of kappa light polypeptide gene enhancer in B-cells
46	E2F-1	E2F transcription factor 1
47	SP1	Sp1 transcription factor
48	MAZ	MYC-associated zinc finger protein (purine-binding transcription factor)
49	TBP	TATA box binding protein
50	MEF-2A	Myocyte enhancer factor 2A
51	HOXD9	Homeobox D9
52	HOXD10	Homeobox D910
53	NF-Y	Nfy Histone-like transcription factor NFY protein family (IC)
54	Elk-1	Member of ETS oncogene family
55	EBF	Transcription factor COE1-like
56	Ik-1	IKAROS family zinc finger 1 (Ikaros)
57	ELF-1	E74-like factor 1 (ets domain transcription factor)
58	PU.1	SPI1 (PU 1) spleen focus forming virus (SFFV) proviral integration oncogene
59	WT1	Wilms tumor 1
60	Egr-3	Early growth response 3
61	AhR:Arnt	Aryl hydrocarbon receptor nuclear translocator
62	PPARα	Peroxisome proliferator-activated receptor alpha
63	TCF-4	Transcription factor 4
64	IRF-2	Interferon regulatory factor 2
65	USF2	Upstream transcription factor 2, c-fos interacting
66	Pax2	Paired box 2

67	Nkx-2.1	Homeobox protein Nkx-2.1
68	Sox-5	sex determining region Y-box 5
69	HSF1	Heat shock transcription factor 1
70	HSF2	Heat shock transcription factor 2
71	CP2	Transcription factor CP2
72	NF-E2	Nuclear factor, erythroid 2
73	MZF1	Myeloid zinc finger 1
74	Evi-1	Ecotropic viral integration site 1
75	c-Rel	V-rel avian reticuloendotheliosis viral oncogene homolog
76	NRF-2	Nf-E2 related factor 2
77	Lyf-1	Lymphoid transcription factor LyF-1
78	Pbx-1	Pre-B-cell leukemia homeobox 1

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