

Supplementary File for

LSD1-ERR α complex requires NRF1 to positively regulate transcription and cell invasion

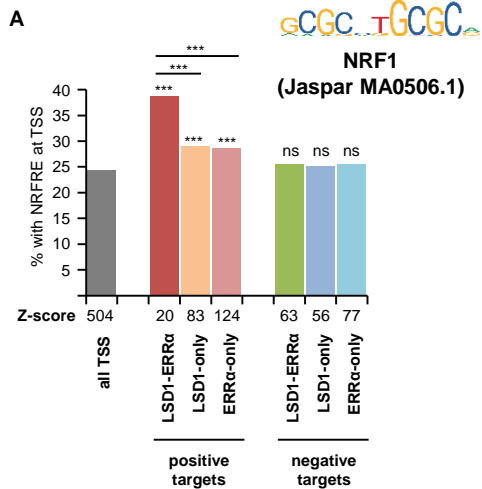
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Contains Supplementary Figures S1, S2, S3 and Supplementary Table S1.

Supplementary Figure S1. Additional bio-informatics data.

a. Same as Fig.1b using the displaying the percent of sequences displaying at least 85% similarity to the indicated NRF1 matrix. Except where indicated, significance is calculated relative to "all TSS" with ***: $p < 0.005$, ns: nonsignificant.

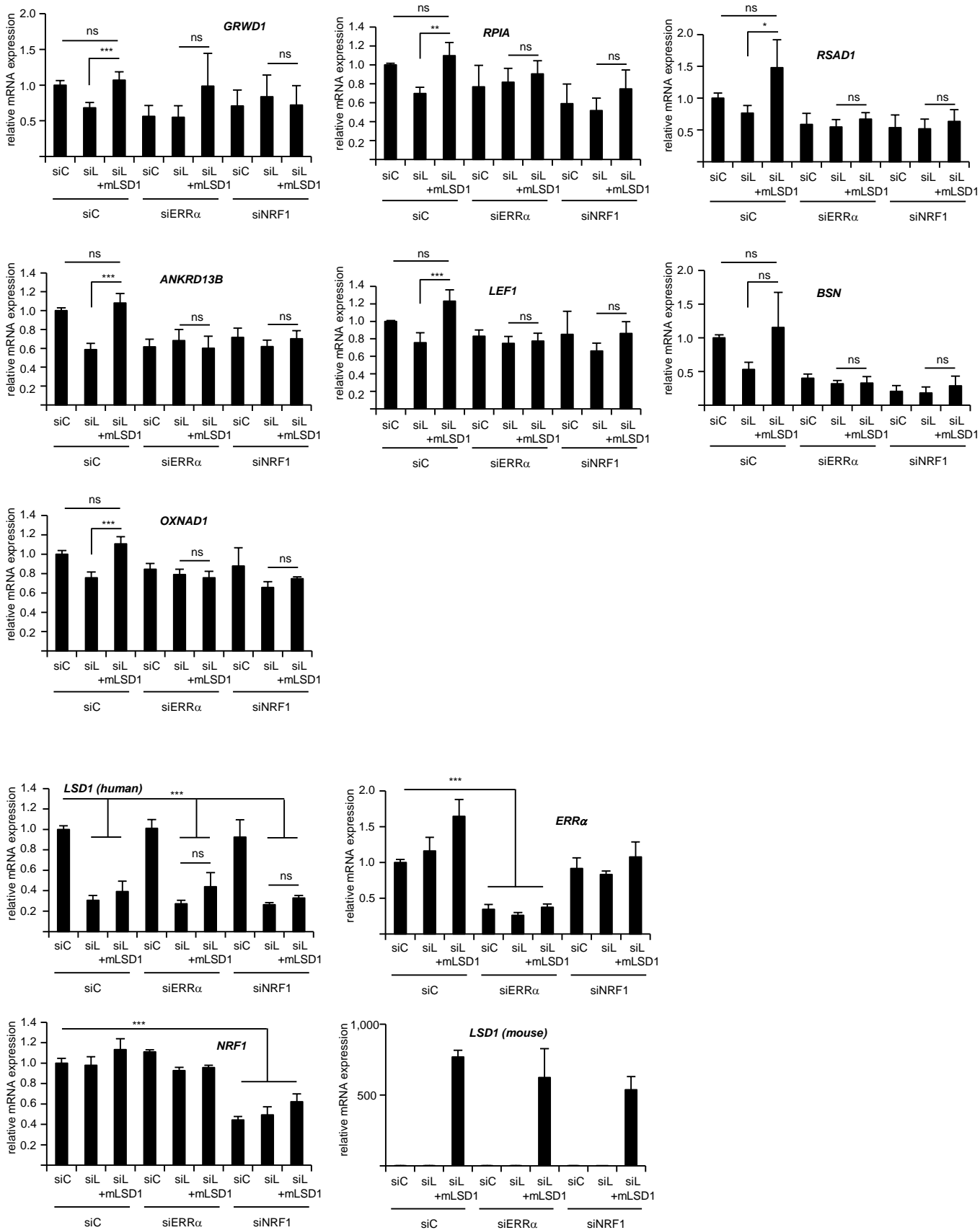
b. Identification of the best (according to % score) putative NRF1 response elements (NRFRE) in the promoters (-150; +50 relative to TSS) of the ten genes of interest used in this study. Sequence, position on chromosome and relative to TSS is indicated.

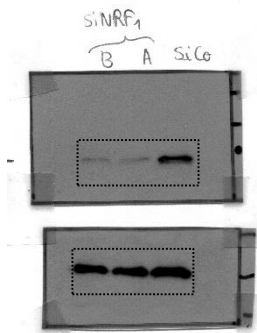


B

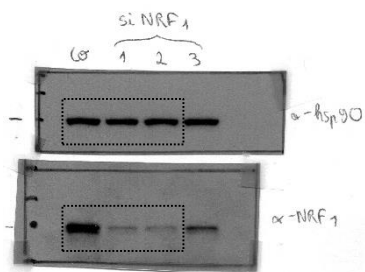
Gene ID	Chr.	NRFRE first nucleotide position		strand	% score	NRFRE sequence
		on chromosome	rel. to TSS			
ONECUT2	18	55102819	-98	+	91.8	CCGCACGCGCA
ZNF768	16	30535412	-42	+	84.6	GCCCAGGCCCC
GRWD1	19	48949286	37	-	88.5	ACACGTGCGCC
RPIA	2	88991018	-144	+	87.2	GCGATTGCGCA
RSAD1	17	48556174	13	+	92.2	CCGGCTGCGCA
ANKRD13B	17	27916641	-146	+	90	CCCCCTGCGCG
LEF1	4	109002670	-70	+	87.1	GCAGAGGCGCA
TMEM198	2	220408794	32	-	87.8	GAGCCCGCCCG
BSN	3	49591895	-17	+	99.6	GCGCATGCGCA
OXNAD1	3	16306683	-13	-	87.8	GCGCGGGCGAG

Supplementary Figure S2. LSD1 requires both NRF1 and $ERR\alpha$ to activate common target genes. Same as Fig. 4b, examining the expression of the indicated target genes.

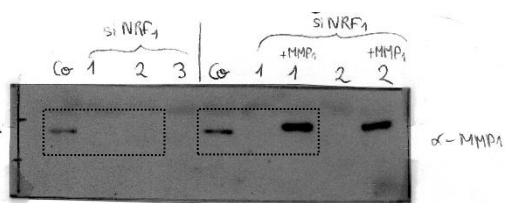




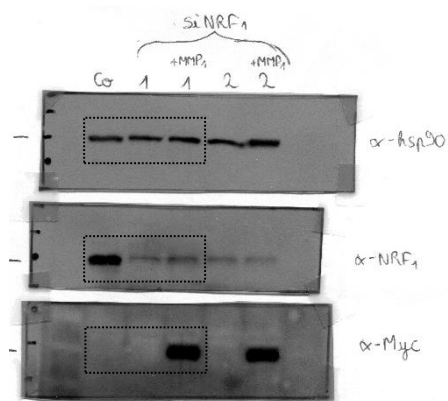
For Fig. 2A.
Note that the blots were left-right inverted in Fig. 2A.



For Fig. 5B.
NRF1 and hsp90 parts



For Fig. 5B and 5E
MMP1 parts



For Fig. 5E
Myc, NRF1 and Hsp90 parts

Supplementary Table S1: Oligonucleotides and siRNAs used in this study

Oligonucleotides for expression studies

36b4	GTCACTGTGCCAGCCAGAA	TCAATGGTGCCCCTGGAGAT
KDM1A (LSD1) human	ACCACAACAGACCCAGAAGG	CTCGGTGGACAAGCACAGTA
ESRRA	CAAGCGCCTCTGCCTGGTCT	ACTCGATGCTCCCCTGGATG
NRF1	GCACCTTTGGAGAATGTGGT	AATTCCGTCGATGGTGAGAG
ONECUT2	AAATCTGGCAGGGAGACCTT	GGTTCCTTGCTCTTTGCGTTT
ZNF768	GGGTACCTCAGAGGCAACAT	GGGTTCAAACCTCTGGGCTTT
GRWD1	TGGTCACCGACTGAGAACAC	GCTCCAGCTGATGACATTGA
RPIA	TAGTCGCTTCATCGTGATCG	GATTCCCTTGTGCCACTGAT
RSAD1	CGCAGCTGAGATGTACCAGA	AACGCCAAGGTAAGTACCAC
ANKRD13B	GGCAAGGTCAAAGGCTGTAA	GTGATCAGGGTCCCATTTTG
LEF1	TGGAAAACGAAGCTCATTCC	GGGTTGGCAGTGATTGTCTT
TMEM198	GTGCTGTTTGTGGAGTCG	TCTCGGTAGCAGAGGAGGAA
BSN	CCAGCCAAACTTCAACACCT	AGCCCTCTGCATCTGACAGT
OXNAD1	CAGCAGCTAAGGTGTGTGGA	AACCCACCAACCACAGAGAC
ABCF1	CAGTGCCAACCAGTGATGAG	GGCAGGCTTAGGAGGATGTT
SNX12	CCGAGGAGATGAAGGGATCT	TGTAGGCAGCGTTCATTCTG
ELK3	CCAAAGGCTTGGAAATCTCA	CGGAGTCAGAAGCAATCCAT
DARS	CCTGAGGCAGAAGGAGAAGA	GACTGGAGACGGAAGACTGC
CFDP1	GTTCCTTCAGCTCTGCCATC	TGAAGCTCTCCCAGTCCAGT
ASB13	GGAGCATCTGGACTGTGTCA	GTGGTGAAGGGCAGTCTCAT
TM9SF4	CCAGAACGATCCCCTAGAAA	TCTCAGCACCTCTCCCAGAT
NDUFB2	ACTGCTGGAGATGGTGGAGT	CACATGAGTCCGCTGAAGAA
MYC	GCCACGCTCCACACATCAG	TCTTGGCAGCAGGATAGTCCTT
MMP1	AGGTCTCTGAGGGTCAAGCA	CTGGTTGAAAAGCATGAGCA
ACLY	TGCCGACTACATCTGCAAAG	GGTTCAGCAAGGTCAGCTTC
NEK7	TGGATGAGCAATCACAAGGA	TAGCCCATATCCGGTCGTAA
UHMK1	TGGCCTGCAGAGTGATACAG	AGCAGAACTGTTTGCCTTCC
Kdm1a mouse	CGAATGACCTCTCAGGAAGC	CCAGCCATAACTGCAATGTG

Oligonucleotides for ChIP on TSS

ONECUT2	GCCCTGATGGACTGAATGAA	GTGCCAGACTTTCCATTGT
ZNF768	TTCTTCTCAGGCTTTGGAG	CAGCAACAGCAAAGCACTTC
GRWD1	GATTAGCGGTCCCAGGAGTT	CTGCTGACACCCGGTAAGAG
RPIA	CGGAGAGCATTATGGGATTG	GGGTCTTGCTGGGAAATGTA
RSAD1	AGTTCTGAGCCAGGCACATC	CCTCCTGTCGGTCTTTCC
ANKRD13B	TCCTCTCTCCCCATCATCC	GGTCCCCACAGCTGCATA
LEF1	ACACACCCCAAAACCAAGAC	GAAGGAGGTGGTGATTGAGG
TMEM198	GGGTTGGCAGTGATTGTCTT	TCTGCTCCTGACGTCATCTC
BSN	CAGAGGCCGCTTCTAGTGG	CGCGGTTGTATCATGCTG
OXNAD1	TGGAGATGGCTGAACTAGGG	GCGAGCCCCACATTCTATT
ABCF1	AGGCTTCCCAGACAGTCGTA	GAGCTTCGCCGTCATCTC
SNX12	TGTGCTGCTAGGCAGGGTAT	GAGTGGAGGCCATATTGAGG
ELK3	GGCGGAAAAGCCTGTTTA	GAGCAGGAAGTTGGAGCAGT
MMP1	AGGAATCCATAAGGGGAGGA	CGGTGTCACCAGTGCTATCT

siRNAs

ERR α #1	GGCAGAAACCUAUCUCAGGUU	CCUGAGAUAGGUUUCUGCCUC
ERR α #2	GAAUGCACUGGUGUCACAUCUGCUG	CAGCAGAUAGACACCAGUGCUUC
LSD1#1	ACUUUGUAACUGUCGAGCUGC	GCAGCUCGACAGUUACAAAGU
LSD1#2	CCACGGAGCGACAGAGCGAGC	GCUCGCUCUGUCGUCCGUGG
NRF1#1	CAGUCACUAUGGCGCUUAA	UUAAGCGCCAUAUGUGACUG
NRF1#2	CACGUUUGCUUCGAAACU	AGUUUCCGAAGCAAACGUG