

# **Establishment of active chromatin structure at enhancer elements by mixed-lineage leukemia 1 to initiate estrogen-dependent gene expression**

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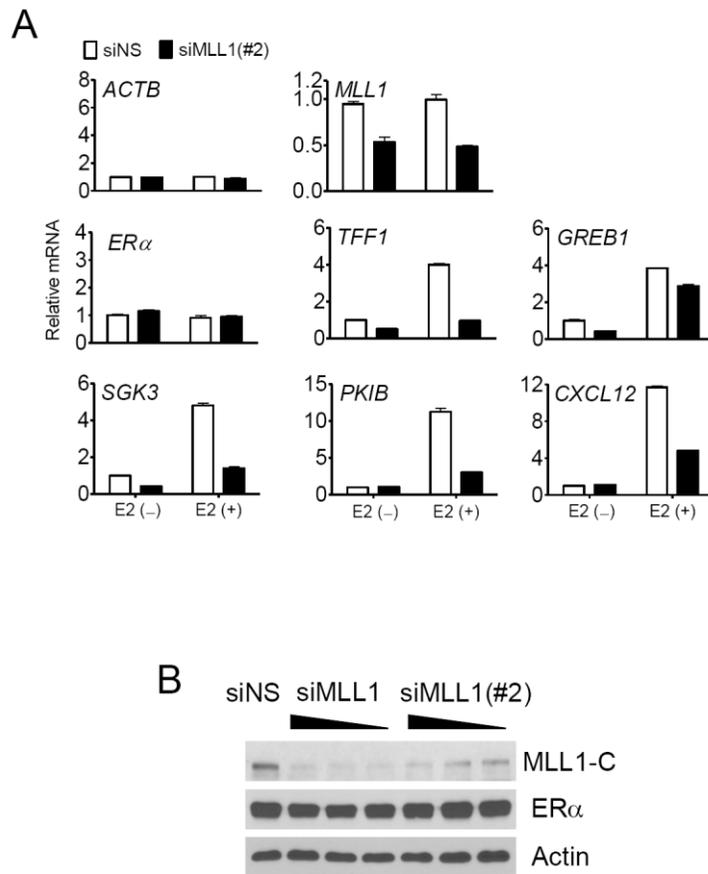
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## **Supplementary data**

### Inventory of Supplementary data

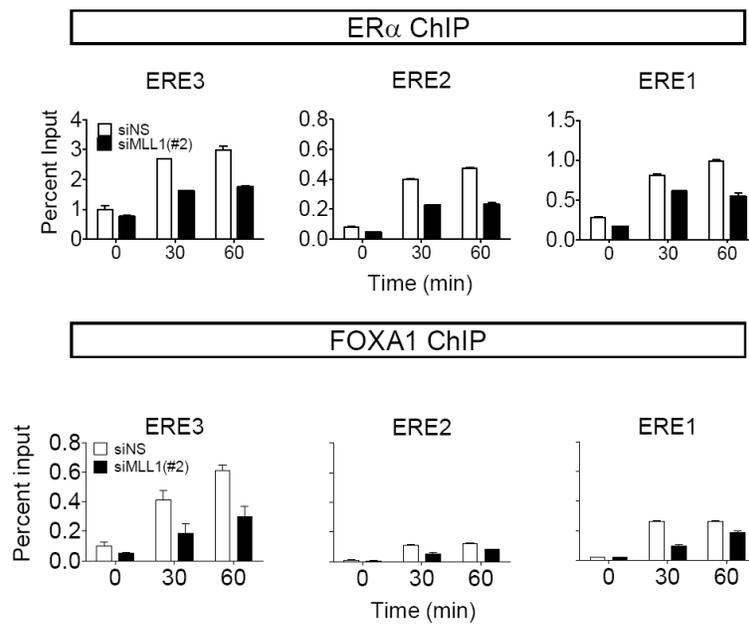
1. Supplementary Fig. 1, Requirement of MLL1 for the expression of endogenous ER $\alpha$  target genes.
2. Supplementary Fig. 2, Effect of MLL1 depletion on the recruitment of ER $\alpha$  and FOXA1 to TFF1 EREs
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5. Supplementary Fig. 5, ER $\alpha$  binding to *TFF1*, *GREB1*, *CTSD*, and *MYC* is dependent on MLL1.

Fig. S1



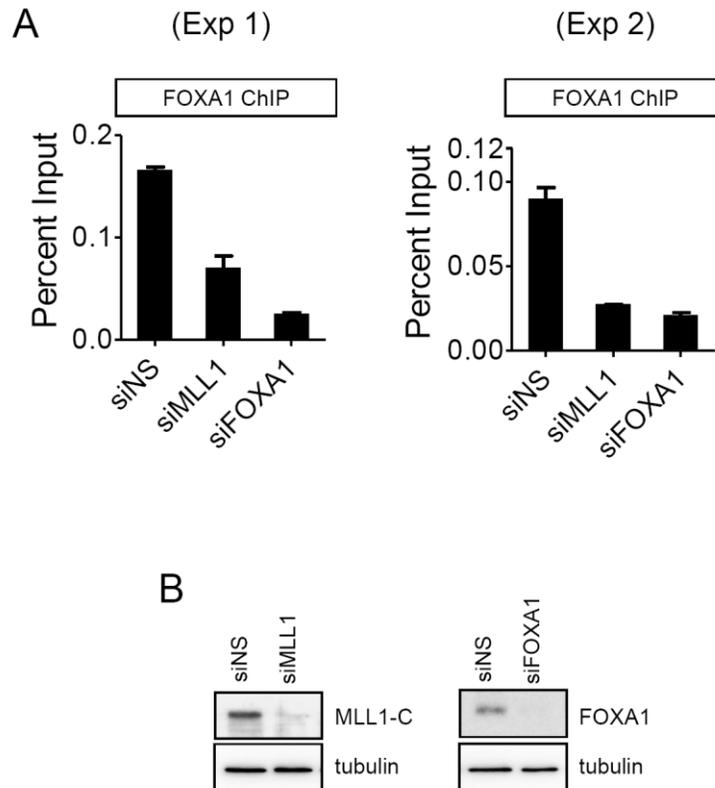
**Supplementary Figure 1.** Requirement of MLL1 for the expression of endogenous ER $\alpha$  target genes. (A) Effect of reduced MLL1 on the expression of estrogen-responsive genes. MCF-7 cells were transfected with siMLL1(#2) (which has a different target sequence from siMLL1) or non-specific siRNA (siNS), and treated with E2 (10 nM) for 16 hours before harvest. Total RNA was analyzed by qRT-PCR. Levels of all mRNAs were normalized to that of GAPDH mRNA. (B) Depletion of MLL1 protein by siMLL1 or siMLL1(#2) transfection. MCF-7 cells were transfected with siMLL1, siMLL1(#2) or siNS as in (A) and grown in hormone free-media for 72 h. Levels of ER $\alpha$ ,  $\beta$ -actin and C-terminal cleavage product of MLL1 (MLL1-C) were assessed by immunoblotting.

Fig. S2



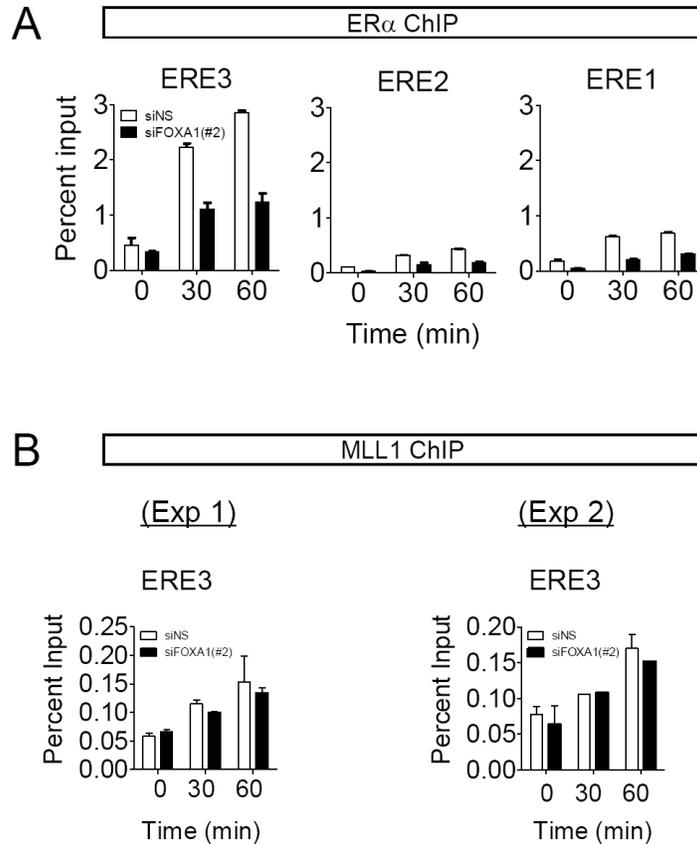
**Supplementary Figure 2.** Effect of MLL1 depletion on the recruitment of ER $\alpha$  and FOXA1 to *TFF1* EREs. ChIP assays were performed as detailed in Figure 2 after transfection with siNS or siMLL1(#2). The MCF-7 cells were treated with E2 (100 nM) or ethanol for 60 min. The amount of the indicated region of the *TFF1* gene precipitated by ER $\alpha$  or FOXA1 antibodies was determined by qPCR.

Fig. S3



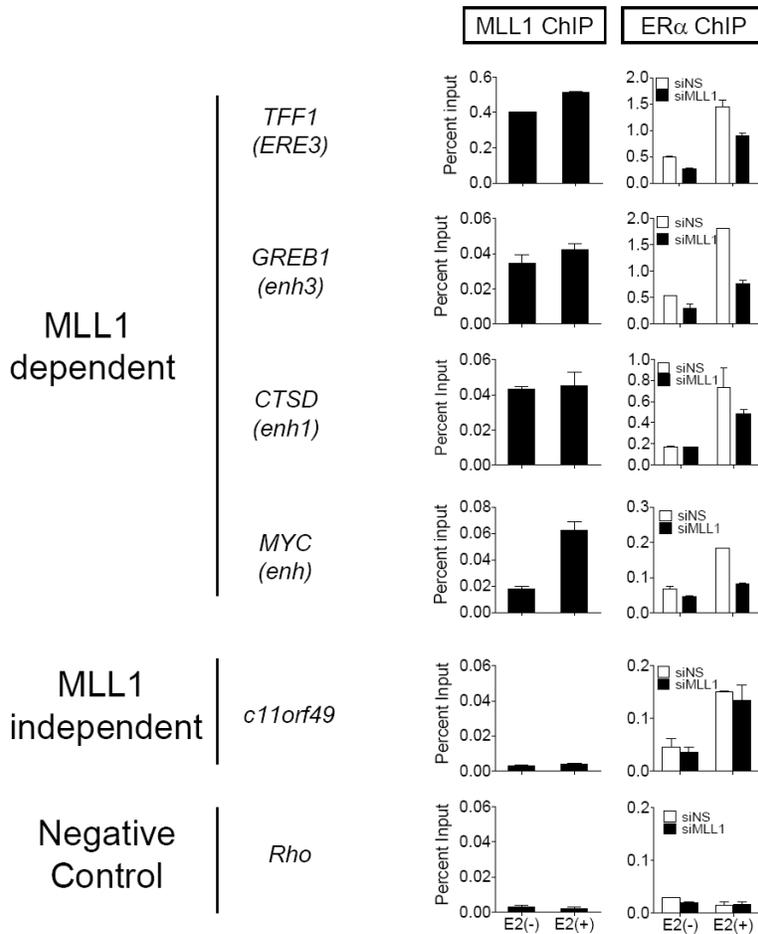
**Supplementary Figure 3.** Effect of MLL1 depletion on the recruitment of FOXA1 in the absence of E2. (A) ChIP assays were performed as detailed in Figure 2 after transfection with siNS, siMLL1 or siFOXA1. The MCF-7 cells were treated with ethanol for 60 min. The amount of the ERE3 region of the TFF1 gene precipitated by FOXA1 antibodies was determined by qPCR. (B) Depletion of MLL1 or FOXA1 protein by siMLL1 or siFOXA1 transfection. MCF-7 cells were transfected with siNS, siMLL1 or siFOXA1 as in (A) and grown in hormone free-media for 72 h. Levels of tubulin, FOXA1, and C-terminal cleavage of MLL1 (MLL1-C) were assessed by immunoblotting.

Fig. S4



**Supplementary Figure 4.** Effect of FOXA1 depletion on the recruitment of ER $\alpha$  or MLL1. (A-B) ChIP assays were performed as detailed in Figure 2 after transfection with siNS or siFOXA1(#2). The MCF-7 cells were treated with E2 (100 nM) for the indicated time periods. The amount of the indicated region of the *TFF1* gene precipitated by ER $\alpha$  or MLL1 antibodies was determined by qPCR. (B) Data shown from two independent experiments

Fig. S5



**Supplementary Figure 5.** ER $\alpha$  binding to *TFF1*, *GREB1*, *CTSD*, and *MYC* is dependent on MLL1. ChIP assays were performed as detailed in Figure 2 after transfection with siNS or siMLL1. The MCF-7 cells were treated with E2 (100 nM) or ethanol for 60 min. The amount of the indicated region of the *TFF1*, *GREB1*, *CTSD*, *MYC*, *c11orf49* and *Rho* gene precipitated by ER $\alpha$  or MLL1 antibodies was determined by qPCR.

## Primer sequences for ChIP experiment

TFF1 (ERE1)

5'-GGCAGGCTCTGTTTGCTTAAAGAGCG-3' (forward)

5'-GGCCATCTCTCACTATGAATCACTTCTGC-3' (reverse)

TFF1 (ERE2)

5'-CCTCCCCAGCTCACGTTGT-3' (forward)

5'-GGGTTGCATTTAAGGGACCTT-3' (reverse)

TFF1 (ERE3)

5'-GTCGTTGCCAGCGTTTCC-3' (forward)

5'-CTTCTCCACGCCCTGTAAATTT-3' (reverse)

TFF1 (-4.8 kb)

5'-CGGTGGCCAGAGAGCATAA-3' (forward)

5'-CCCTTCGGGTGCCACTCAA-3' (reverse)

## Primer sequences for qRT-PCR

ACTB

5'-ACCCCATCGAGCACGGCATCG-3' (forward)

5'-GTCACCGGAGTCCATCACGATG-3' (reverse)

TFF1

5'-GAACAAGGTGATCTGCG-3' (forward)

5'-TGGTATTAGGATAGAAGCACCA-3' (reverse)

GREB1

5'-CAAAGAATAACCTGTTGGCCCTGC-3' (forward)

5'-GACATGCCTGCGCTCTCATACTTA-3' (reverse)

PgR

5'-GTGCCTATCCTGCCTCTCAATC-3' (forward)

5'-CCCGCCGTCGTAACCTTCG-3' (reverse)

SGK3

5'-TGAGGCCAGGAGTGAGTCTT-3' (forward)

5'-TATCATCTGGTCCAGCAACA-3' (reverse)

PKIB

5'-CCAATTTTGCATCTTCAGCA-3' (forward)

5'-GGCTTTTCCAATTGGTCTTG-3' (reverse)

MYC

5'-CTCTCAACGACAGCAGCTCG-3' (forward)  
5'-CAACATCGATTTCTTCCTCATCTTC-3' (reverse)

#### CCND1

5'-AAGCTCAAGTGGAACCT-3' (forward)  
5'-AGGAAGTTGTTGGGGC-3' (reverse)

#### CXCL12

5'-TCAGCCTGAGCTACAGATGC-3' (forward)  
5'-CTTTAGCTTCGGGTCAATGC-3' (reverse)

#### TFF1 pre-mRNA

5'-ACCATGGAGAACAAGGTGATC-3' (forward)  
5'-TAAAACAGTGGCTCCTGGCG-3' (reverse)

#### GREB1 pre-mRNA

5'-GATAAAAGCAACGTGCGTCTC-3' (forward)  
5'-TCTTGCACAATTCCATCGAG-3' (reverse)

#### CCND1 pre-mRNA

5'-CCTACTTCAAATGTGTGCAGAAG-3' (forward)  
5'-CAACAAGTTGCAGGGAAGTC-3' (reverse)

#### PgR pre-mRNA

5'-CAGGTCTACCCGCCCTATCT-3' (forward)  
5'-TATATTCTGCGCCACCTTC-3' (reverse)

#### GAPDH

5'-TCTGGTAAAGTGGATATTGTTG-3' (forward)  
5'-GATGGTGATGGGATTTCC-3' (reverse)

### Sequences of siRNAs

#### siNS

5'-UUCUCCGAACGUGUCACGUdTdT-3' (sense)  
5'-ACGUGACACGUUCGGAGAAdTdT-3' (anti-sense)

#### siMLL1

5'-GAUUCGAACACCCAGUUAUdTdT-3' (sense)  
5'-AUAACUGGGUGUUCGAAUCdTdT-3' (anti-sense)

#### siMLL1(#2)

5'-GCACUGUUAACAUAUCCACdTdT-3' (sense)  
5'-GUGGAAUGUUUAACAGUGCdTdT-3' (anti-sense)

siFOXA1

5'-GAGAGAAAAAAUCAACAGCdTdT-3' (sense)

5'-GCUGUUGAUUUUUUCUCUCdTdT-3' (anti-sense)

siFOXA1(#2)

5'-GCGAAGUUUAAUGAUCCACdTdT-3' (sense)

5'-GUGGAUCAUUAACUUCGCdTdT-3' (anti-sense)