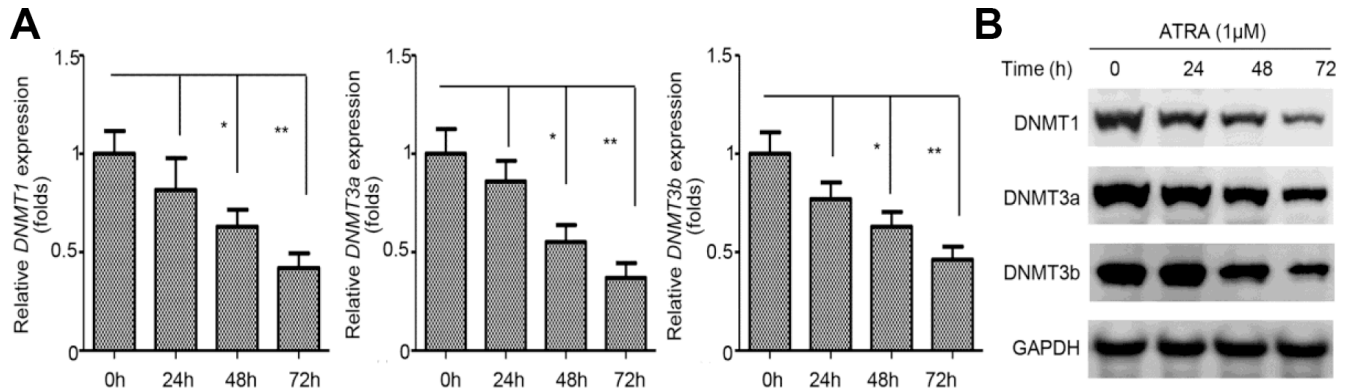
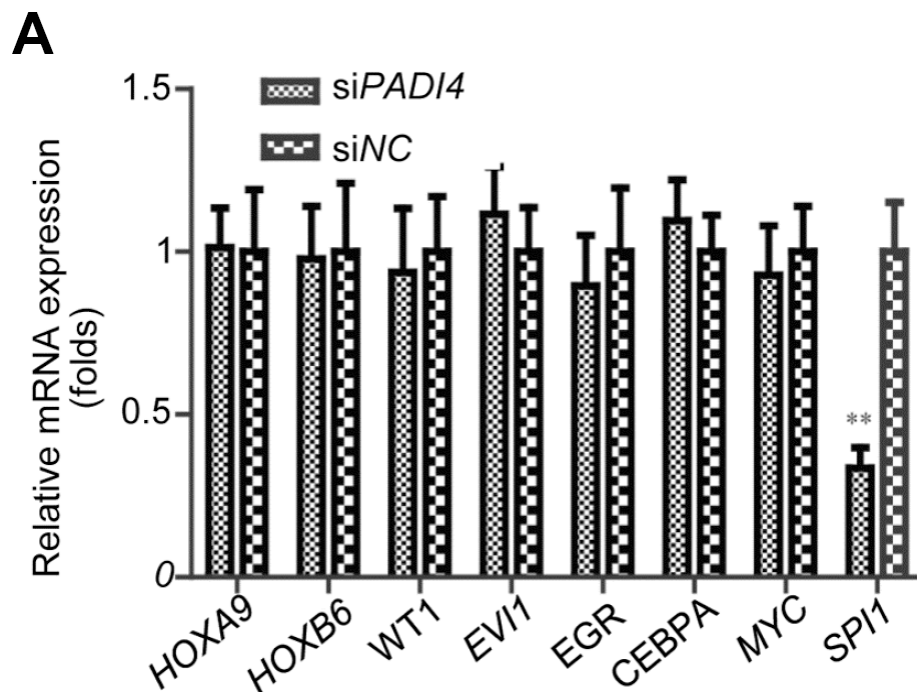


A novel PAD4/SOX4/PU.1 signaling pathway is involved in the committed differentiation of acute promyelocytic leukemia cells into granulocytic cells

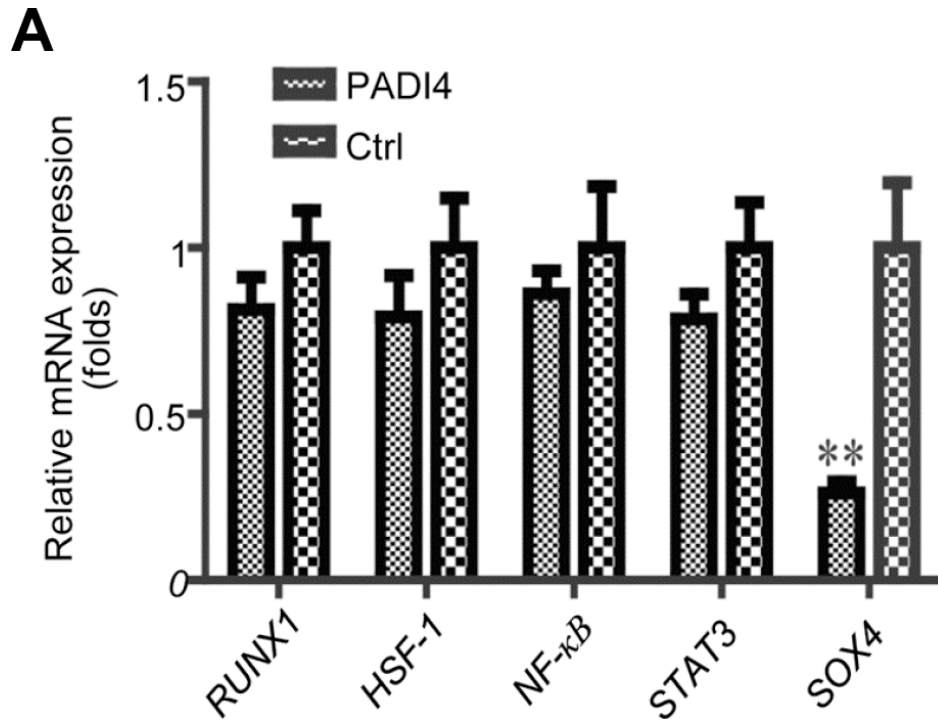
Supplementary Materials



Supplementary Figure S1: ATRA treatment suppresses the expression of DNMTs. (A) and (B) Expression of three DNMTs was detected by RT-PCR (A) and Western blot analysis (B) after treatment with ATRA (1 μ M) at indicated timepoints.



Supplementary Figure 2: PU.1 was regulated by PAD4. (A) Expression of *SPI1*, *HOXA9*, *HOXB6*, *WT1*, *EVI1*, and *EGR* after silencing PAD4 in ATRA-induced HL-60 cells. $**P < 0.01$. Data are means of biological triplicates (\pm standard error) and representative of triplicate experiments.



Supplementary Figure 3: SOX4 was regulated by PAD4. (A) Expression of *RUNX1*, *HSF-1*, *NF- κ B*, *STAT3*, and *SOX4* after silencing PAD4 in ATRA-induced HL-60 cells. ** $P < 0.01$. Data are means of biological triplicates (\pm standard error) and representative of triplicate experiments.

Supplementary Table S1: Primers used in this study

Gene	Forward	Reverse
qRT-PCR		
<i>PADI4</i>	GTTTAGGGTCAGACAGTCTGG	AGATGTGAGTAGTGGCACATGC
<i>SOX4</i>	ACACTGGTGGCAGGTTAAGG	CGCTGTTTGGATTTCTGAT
<i>PU.1</i>	CTGTTGGACCTGCTCCGCA	TGCGGAGCAGGTCCAACAG
<i>GAPDH</i>	AATGGGCAGCCGTTAGGAAA	GCGCCAATACGACCAAATC
<i>RUNX1</i>	GGACGCCAGAAGGAAGTCAA	TCGCAGCCAGGAAAGAAGTT
<i>HSF-1</i>	CCGTGGACACCCTCTTGTC	GAGCTCATTCTTGTCAGGCA
<i>STAT3</i>	CTGCCCATACCTGAAGACC	TCCTCACATGGGGGAGGTAG
<i>NF-κB</i>	CCAACAGATGGCCCATACCT	AACCTTTGCTGGTCCCACAT
ChIP		
PU.1-1	GATGGATGGGTAGATGAGTG	CAGCCATCCACAAATCCACC
PU.1-2	TGGATGGCTGGATGAATGAG	ATGCAGTATCTTTTTGGTAT
PU.1-3	ACTGCATTTGTATGTTTATC	CCAATTCCATCCATGTTGC
PU.1-4	GATACTGCATTTGTATGTTT	GATAATGTCCTCCAATTCC
PU.1-5	AGGACATTATCATATGCGAA	CATTCATTCATCCATCCATC
PU.1-6	TGAATGAATGAAGGGGTAGG	CAGTCCATCAGTCCACAA
PU.1-7	GTGGACTGATGGACTGATAG	CCCATGCATCTATTTGTC
PU.1-8	GACAGACAAATAGATGCATG	CCTAGGGCTCTGTTTCCAGCC
PU.1-9	ATAGATGCATGGGAAAACAG	TGGGTCTTATACCCTCC
PU.1-10	AGCAGCACTATGCTGAAGAC	GCTCTAACCCAACAAATGC
PU.1-11	TAGAGCAAAAAGCCTCCCAGT	AACCCGTTTTGCATAAATCTCT
PU.1-12	ACGGGTTGGGGCGGTGATGT	AATGCAGAGCCCCTCAGGATG
SOX4-I	TTAAAGAGCGTGCAAGAA	AGCCAAGACTGTGAAAGG
SOX4-II	CAGTATTCACCCTTACCC	GATTTCTGCTGCCTCTTT
SOX4-III	TACGGAGCACTACCTAAT	GTAAATCCCTTCAGAACC
SOX4-IV	CCCTTCGATTCAAGTAAC	TAGGCAGATTTCCAGAGT
SOX4-V	ACTTTATAGAGGGTTGTTGT	GTCTGCCTGTTATTACTT
Plasmid		
pGL3- SOX4 (P1)	CGAGCTCAGGTGCCTGTGTGTTT	GGAAGATCTTCCGCCTCGCGCCTCTT
pGL3- SOX4 (P2)	CGAGCTCCAATGGAATGGCAGGGT	GGAAGATCTTCCGCCTCGCGCCTCTT
pGL3- SOX4 (P3)	CGAGCTACAATACATCAGGTGC	GGAAGATCTTCCGCCTCGCGCCTCTT
pGL3- SOX4 (P4)	CGAGCTCGAGCACTACCTAATGTG	GGAAGATCTTCCGCCTCGCGCCTCTT
pGL3- SOX4 (P5)	CGAGCTCTGCACCAGAGGCTGATT	GGAAGATCTTCCGCCTCGCGCCTCTT
pGL3- SOX4 (P6)	CGAGCTCAGGTGCCTGTGTGTTT	GGAAGATCTAATCAGCCTCTGGTGCA