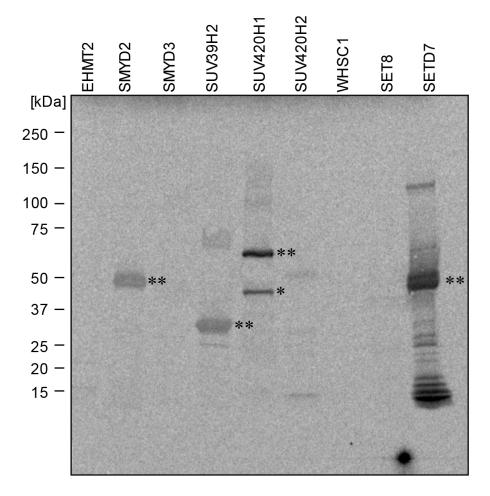
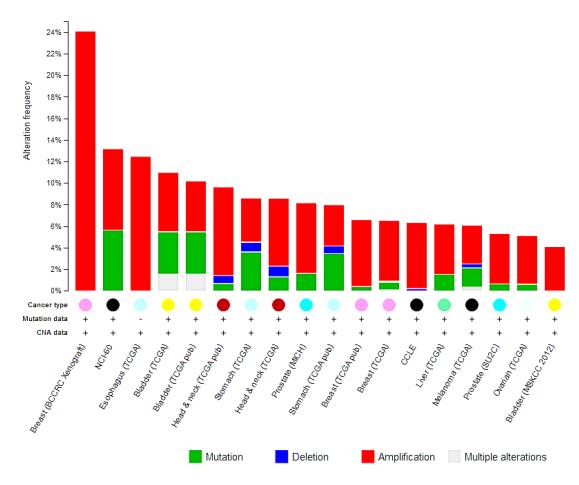
# **SUV420H1** enhances the phosphorylation and transcription of ERK1 in cancer cells

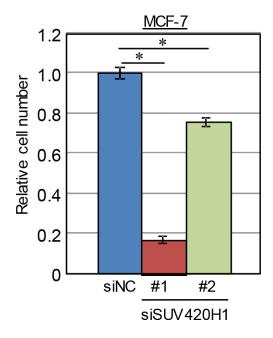
## **Supplementary Materials**



**Supplementary Figure S1: SUV420H1 methylates ERK1** *in vitro*. Recombinant ERK1 protein was incubated with a variety of methyltransferases, and methylation signal was detected by autoradiography. \*ERK1 methylation band. \*\*automethylation band.



**Supplementary Figure S2: Cross-cancer alteration summary for** *SUV420H1***.** The data were obtained from the TCGA database (http://www.cbioportal.org/public-portal/).



Supplementary Figure S3: Involvement of SUV420H1 in the growth of the breast cancer cell MCF-7. Relative cell number shows the value normalized to siNC-treated cells. Mean  $\pm$  SD of four independent experiments. P values were calculated using Student's t-test (\*P < 0.05).

#### **Supplementary Table S1: Information of certificated cell lines**

Name	Origin	Certification institution	Tested method	DNA profile or characteristics
MCF7	human breast cancer	ATCC	STR	Amelogenin: X/X CSF1PO: 10/10 D13S317: 11/11 D16S539: 11/12 D5S818: 12/12 D7S820: 8/9 THO1: 6/6 TPOX: 9/12 vWA: 14/15 D3S1358: 16/16 D21S11: 30/30 D18S51: 14/14 Penta E: 7/12 Penta D: 12/12 D8S1179: 10/14 FGA: 23/25 CLS 2011.
293T	human embryonic kidney fibroblast	ATCC	STR	Amelogenin: X CSF1PO: 11, 12 D13S317: 12, 14 D16S539: 9, 13 D5S818: 8, 9 D7S820: 11 THO1: 7, 9.3 TPOX: 11 vWA: 16, 18, 19
HeLa	human cervix carcinoma	ATCC	STR	Amelogenin: X, Y CSF1PO: 11, 12 D13S317: 11, 14 D16S539: 9, 11 D5S818: 11, 12 D7S820: 10, 11 THO1: 8 TPOX: 8 vWA: 15

ATCC; American Type Culture Collection

## **Supplementary Table S2: Characteristics of SCCHN cell lines**

Cell name	TNM stage	Specimen site	Gender	HPV status
HN-SCC-151	T3N0M0	Oral tongue	_	HPV16-negative
FaDu	-	Hypopharynx	Male	HPV16-negative

## Supplementary Table S3: Primer sequences for quantitative RT-PCR

Gene name	Primer sequence
GAPDH (housekeeping gene)-f	5' GCAAATTCCATGGCACCGTC 3'
GAPDH (housekeeping gene)-r	5' TCGCCCCACTTGATTTTGG 3'
SDH (housekeeping gene)-f	5' TGGGAACAAGAGGGCATCTG 3'
SDH (housekeeping gene)-r	5' CCACCACTGCATCAAATTCATG 3'
SUV420H1-f	5' AGAAATCATTGCAAGCGGCTGGAG 3'
SUV420H1-r	5' CTGGCTCCTTATCTTTTTAATGG 3'

#### **Supplementary Table S4: siRNA sequences**

siRNA name	Sequence		
	Target#1	Sense: 5' AUCCGCGCGAUAGUACGUA 3'	
siNegative control		Antisense: 5' UACGUACUAUCGCGCGGAU 3'	
(Cocktail)	Target#2	Sense: 5' UUACGCGUAGCGUAAUACG 3'	
		Antisense: 5' CGUAUUACGCUACGCGUAA 3'	
	Target#3	Sense: 5' UAUUCGCGCGUAUAGCGGU 3'	
		Antisense: 5' ACCGCUAUACGCGCGAAUA 3'	
siSUV420H1#1		Sense: 5' CUGCGUUUAUAAACCAUGA 3'	
		Antisense: 5' UCAUGGUUUAUAAACGCAG 3'	
siSUV420H1#2		Sense: 5' GUGUCAACUGGUCGAGAUA 3'	
		Antisense: 5' UAUCUCGACCAGUUGACAC 3'	