

**Phospho-EP300(S1834) Blocking Peptide**  
**Synthetic peptide**  
**Catalog # BP3842a****Specification****Phospho-EP300(S1834) Blocking Peptide -  
Product Information**

Primary Accession [Q09472](#)  
Other Accession [NP\\_001420.2](#)

**Phospho-EP300(S1834) Blocking Peptide -  
Additional Information**

**Gene ID** 2033

**Other Names**

Histone acetyltransferase p300, p300 HAT,  
E1A-associated protein p300, EP300, P300

**Target/Specificity**

The synthetic peptide sequence is selected  
from aa 1827-1841 of HUMAN EP300

**Format**

Peptides are lyophilized in a solid powder  
format. Peptides can be reconstituted in  
solution using the appropriate buffer as  
needed.

**Storage**

Maintain refrigerated at 2-8°C for up to 6  
months. For long term storage store at  
-20°C.

**Precautions**

This product is for research use only. Not  
for use in diagnostic or therapeutic  
procedures.

**Phospho-EP300(S1834) Blocking Peptide -  
Protein Information**

**Name** EP300

**Synonyms** P300

**Function**

Functions as histone acetyltransferase and  
regulates transcription via chromatin  
remodeling (PubMed:<a href="http://www.u

**Phospho-EP300(S1834) Blocking Peptide -  
Background**

This gene encodes the adenovirus  
E1A-associated cellular  
p300 transcriptional co-activator protein. It  
functions as histone  
acetyltransferase that regulates transcription  
via chromatin  
remodeling and is important in the processes  
of cell proliferation  
and differentiation. It mediates cAMP-gene  
regulation by binding  
specifically to phosphorylated CREB protein.  
This gene has also  
been identified as a co-activator of HIF1A  
(hypoxia-inducible  
factor 1 alpha), and thus plays a role in the  
stimulation of  
hypoxia-induced genes such as VEGF. Defects  
in this gene are a  
cause of Rubinstein-Taybi syndrome and may  
also play a role in  
epithelial cancer.

**Phospho-EP300(S1834) Blocking Peptide -  
References**

Zhang, M., et al. J. Immunol.  
185(7):3960-3969(2010)  
Bailey, S.D., et al. Diabetes Care  
33(10):2250-2253(2010)  
Vempati, R.K., et al. J. Biol. Chem.  
285(37):28553-28564(2010)  
Reynoird, N., et al. EMBO J.  
29(17):2943-2952(2010)  
Jang, E.R., et al. Biochem. Biophys. Res.  
Commun. 397(4):637-643(2010)

niprot.org/citations/23415232"  
target="\_blank">23415232</a>,  
PubMed:<a href="http://www.uniprot.org/citations/23934153"  
target="\_blank">23934153</a>,  
PubMed:<a href="http://www.uniprot.org/citations/8945521"  
target="\_blank">8945521</a>). Acetylates  
all four core histones in nucleosomes.  
Histone acetylation gives an epigenetic tag  
for transcriptional activation (PubMed:<a href="http://www.uniprot.org/citations/23415232" target="\_blank">23415232</a>,  
PubMed:<a href="http://www.uniprot.org/citations/23934153"  
target="\_blank">23934153</a>,  
PubMed:<a href="http://www.uniprot.org/citations/8945521"  
target="\_blank">8945521</a>). Mediates  
cAMP-gene regulation by binding  
specifically to phosphorylated CREB protein.  
Mediates acetylation of histone H3 at  
'Lys-122' (H3K122ac), a modification that  
localizes at the surface of the histone  
octamer and stimulates transcription,  
possibly by promoting nucleosome  
instability. Mediates acetylation of histone  
H3 at 'Lys-27' (H3K27ac) (PubMed:<a href="http://www.uniprot.org/citations/23911289" target="\_blank">23911289</a>). Also  
functions as acetyltransferase for  
non-histone targets, such as ALX1, HDAC1,  
PRMT1 or SIRT2 (PubMed:<a href="http://www.uniprot.org/citations/12929931" target="\_blank">12929931</a>,  
PubMed:<a href="http://www.uniprot.org/citations/16762839" target="\_blank">16762839</a>,  
PubMed:<a href="http://www.uniprot.org/citations/18722353" target="\_blank">18722353</a>).  
Acetylates 'Lys- 131' of ALX1 and acts as its  
coactivator (PubMed:<a href="http://www.uniprot.org/citations/12929931" target="\_blank">12929931</a>).  
Acetylates SIRT2 and is proposed to  
indirectly increase the transcriptional  
activity of TP53 through acetylation and  
subsequent attenuation of SIRT2  
deacetylase function (PubMed:<a href="http://www.uniprot.org/citations/18722353" target="\_blank">18722353</a>).  
Acetylates HDAC1 leading to its inactivation  
and modulation of transcription (PubMed:<a href="http://www.uniprot.org/citations/16762839" target="\_blank">16762839</a>).  
Acetylates 'Lys-247' of EGR2 (By similarity).

Acts as a TFAP2A-mediated transcriptional coactivator in presence of CITED2 (PubMed:<a href="http://www.uniprot.org/citations/12586840" target="\_blank">12586840</a>). Plays a role as a coactivator of NEUROD1-dependent transcription of the secretin and p21 genes and controls terminal differentiation of cells in the intestinal epithelium. Promotes cardiac myocyte enlargement. Can also mediate transcriptional repression. Acetylates FOXO1 and enhances its transcriptional activity (PubMed:<a href="http://www.uniprot.org/citations/15890677" target="\_blank">15890677</a>). Acetylates BCL6 which disrupts its ability to recruit histone deacetylases and hinders its transcriptional repressor activity (PubMed:<a href="http://www.uniprot.org/citations/12402037" target="\_blank">12402037</a>). Participates in CLOCK or NPAS2-regulated rhythmic gene transcription; exhibits a circadian association with CLOCK or NPAS2, correlating with increase in PER1/2 mRNA and histone H3 acetylation on the PER1/2 promoter (PubMed:<a href="http://www.uniprot.org/citations/14645221" target="\_blank">14645221</a>). Acetylates MTA1 at 'Lys-626' which is essential for its transcriptional coactivator activity (PubMed:<a href="http://www.uniprot.org/citations/16617102" target="\_blank">16617102</a>). Acetylates XBP1 isoform 2; acetylation increases protein stability of XBP1 isoform 2 and enhances its transcriptional activity (PubMed:<a href="http://www.uniprot.org/citations/20955178" target="\_blank">20955178</a>). Acetylates PCNA; acetylation promotes removal of chromatin-bound PCNA and its degradation during nucleotide excision repair (NER) (PubMed:<a href="http://www.uniprot.org/citations/24939902" target="\_blank">24939902</a>). Acetylates MEF2D (PubMed:<a href="http://www.uniprot.org/citations/21030595" target="\_blank">21030595</a>). Acetylates and stabilizes ZBTB7B protein by antagonizing ubiquitin conjugation and degradation, this mechanism may be involved in CD4/CD8 lineage differentiation (PubMed:<a href="http://www.uniprot.org/citations/20810990" target="\_blank">20810990</a>).

Acetylates GABPB1, impairing GABPB1 heterotetramerization and activity (By similarity). In addition to protein acetyltransferase, can use different acyl-CoA substrates, such as (2E)-butenoyl-CoA (crotonyl-CoA), butanoyl-CoA (butyryl-CoA), 2-hydroxyisobutanoyl-CoA (2-hydroxyisobutyryl-CoA), lactoyl-CoA or propanoyl-CoA (propionyl-CoA), and is able to mediate protein crotonylation, butyrylation, 2-hydroxyisobutyrylation, lactylation or propionylation, respectively (PubMed:<a href="http://www.uniprot.org/citations/17267393" target="\_blank">17267393</a>, PubMed:<a href="http://www.uniprot.org/citations/25818647" target="\_blank">25818647</a>, PubMed:<a href="http://www.uniprot.org/citations/29775581" target="\_blank">29775581</a>, PubMed:<a href="http://www.uniprot.org/citations/31645732" target="\_blank">31645732</a>). Acts as a histone crotonyltransferase; crotonylation marks active promoters and enhancers and confers resistance to transcriptional repressors (PubMed:<a href="http://www.uniprot.org/citations/25818647" target="\_blank">25818647</a>). Histone crotonyltransferase activity is dependent on the concentration of (2E)-butenoyl-CoA (crotonyl-CoA) substrate and such activity is weak when (2E)-butenoyl-CoA (crotonyl-CoA) concentration is low (PubMed:<a href="http://www.uniprot.org/citations/25818647" target="\_blank">25818647</a>). Also acts as a histone butyryltransferase; butyrylation marks active promoters (PubMed:<a href="http://www.uniprot.org/citations/17267393" target="\_blank">17267393</a>). Catalyzes histone lactylation in macrophages by using lactoyl-CoA directly derived from endogenous or exogenous lactate, leading to stimulates gene transcription (PubMed:<a href="http://www.uniprot.org/citations/31645732" target="\_blank">31645732</a>). Acts as a protein-lysine 2-hydroxyisobutyryltransferase; regulates glycolysis by mediating 2-hydroxyisobutyrylation of glycolytic enzymes (PubMed:<a href="http://www.uniprot.org/citations/29775581" target="\_blank">29775581</a>).

target="\_blank">29775581</a>).  
Functions as a transcriptional coactivator  
for SMAD4 in the TGF-beta signaling  
pathway (PubMed:<a href="http://www.uni  
prot.org/citations/25514493"  
target="\_blank">25514493</a>).  
Acetylates PCK1 and promotes PCK1  
anaplerotic activity (PubMed:<a href="http:  
//www.uniprot.org/citations/30193097"  
target="\_blank">30193097</a>).  
Acetylates RXRA and RXRG (PubMed:<a href="http://www.uniprot.org/citations/17761950" target="\_blank">17761950</a>).

#### **Cellular Location**

Cytoplasm. Nucleus. Chromosome.  
Note=Localizes to active chromatin:  
Colocalizes with histone H3 acetylated  
and/or crotonylated at 'Lys-18' (H3K18ac  
and H3K18cr, respectively)  
(PubMed:25818647). In the presence of  
ALX1 relocates from the cytoplasm to the  
nucleus. Colocalizes with ROCK2 in the  
nucleus (PubMed:12929931).

#### **Phospho-EP300(S1834) Blocking Peptide - Protocols**

Provided below are standard protocols that you  
may find useful for product applications.

- [Blocking Peptides](#)