

# ERCC8 Antibody

Rabbit mAb

Catalog # AP92619

## Product Information

<b>Application</b>	WB, IP
<b>Primary Accession</b>	<a href="#">Q13216</a>
<b>Reactivity</b>	Human
<b>Clonality</b>	Monoclonal
<b>Other Names</b>	CKN1; CSA; ERCC8;
<b>Isotype</b>	Rabbit IgG
<b>Host</b>	Rabbit
<b>Calculated MW</b>	44055

## Additional Information

<b>Dilution</b>	WB 1:500~1:2000 IP 1:50
<b>Purification</b>	Affinity-chromatography
<b>Immunogen</b>	A synthesized peptide derived from human ERCC8
<b>Description</b>	Substrate-recognition component of the CSA complex, a DCX (DDB1-CUL4-X-box) E3 ubiquitin-protein ligase complex, involved in transcription-coupled nucleotide excision repair.
<b>Storage Condition and Buffer</b>	Rabbit IgG in phosphate buffered saline , pH 7.4, 150mM NaCl, 0.02% sodium azide and 50% glycerol. Store at +4°C short term. Store at -20°C long term. Avoid freeze / thaw cycle.

## Protein Information

<b>Name</b>	ERCC8 {ECO:0000303   PubMed:19894250, ECO:0000312   HGNC:HGNC:3439}
<b>Function</b>	Substrate-recognition component of the CSA complex, a DCX (DDB1-CUL4-X-box) E3 ubiquitin-protein ligase complex, involved in transcription-coupled nucleotide excision repair (TC-NER), a process during which RNA polymerase II-blocking lesions are rapidly removed from the transcribed strand of active genes (PubMed: <a href="#">12732143</a> , PubMed: <a href="#">16751180</a> , PubMed: <a href="#">16964240</a> , PubMed: <a href="#">32142649</a> , PubMed: <a href="#">34526721</a> , PubMed: <a href="#">38316879</a> , PubMed: <a href="#">38600235</a> , PubMed: <a href="#">38600236</a> ). Following recruitment to lesion-stalled RNA polymerase II (Pol II), the CSA complex mediates ubiquitination of Pol II subunit POLR2A/RPB1 at 'Lys- 1268', a critical TC-NER checkpoint, governing RNA Pol II stability and initiating DNA damage excision by TFIIH recruitment (PubMed: <a href="#">12732143</a> , PubMed: <a href="#">16751180</a> , PubMed: <a href="#">16964240</a> , PubMed: <a href="#">32142649</a> , PubMed: <a href="#">32355176</a> , PubMed: <a href="#">34526721</a> , PubMed: <a href="#">38316879</a> , PubMed: <a href="#">38600235</a> , PubMed: <a href="#">38600236</a> ). The CSA complex also promotes the ubiquitination and subsequent proteasomal degradation of ERCC6/CSB in a UV-dependent manner; ERCC6 degradation is essential for the recovery of RNA synthesis

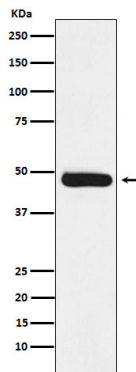
after transcription-coupled repair (PubMed:[16751180](#)). Also plays a role in DNA double-strand breaks (DSSBs) repair by non-homologous end joining (NHEJ) (PubMed:[29545921](#)).

### Cellular Location

Nucleus. Chromosome Nucleus matrix. Note=Recruited to lesion- stalled RNA polymerase II (Pol II) sites by ERCC6/CSB (PubMed:32355176). UV-induced translocation to the nuclear matrix is dependent on ERCC6/CSB (PubMed:26620705).

### Images

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Western blot analysis of ERCC8 expression in Molt4 cell lysate.

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