

ERCC8 Rabbit mAb
Catalog # AP76489**Specification****ERCC8 Rabbit mAb - Product Information**

Application	WB, IP
Primary Accession	Q13216
Reactivity	Human
Host	Rabbit
Clonality	Monoclonal Antibody
Calculated MW	44055

ERCC8 Rabbit mAb - Additional Information**Gene ID** 1161**Other Names**

ERCC8

Dilution

WB~~1/500-1/1000

IP~~N/A

Format

Liquid

ERCC8 Rabbit mAb - Protein Information**Name** ERCC8 {ECO:0000303|PubMed:19894250, ECO:0000312|HGNC:HGNC:3439}**Function**

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: [12732143](http://www.uniprot.org/citations/12732143) target="_blank">12732143, PubMed: [16751180](http://www.uniprot.org/citations/16751180) target="_blank">16751180, PubMed: [16964240](http://www.uniprot.org/citations/16964240) target="_blank">16964240, PubMed: [32142649](http://www.uniprot.org/citations/32142649) target="_blank">32142649, PubMed: [34526721](http://www.uniprot.org/citations/34526721) target="_blank">34526721, PubMed: [38316879](http://www.uniprot.org/citations/38316879) target="_blank">38316879, PubMed: [38600235](http://www.uniprot.org/citations/38600235) target="_blank">38600235, PubMed: [38600236](http://www.uniprot.org/citations/38600236) target="_blank">38600236). 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: [12732143](http://www.uniprot.org/citations/12732143) target="_blank">12732143, PubMed: [16751180](http://www.uniprot.org/citations/16751180) target="_blank">16751180, PubMed: [16964240](http://www.uniprot.org/citations/16964240) target="_blank">16964240).

target="_blank">16964240, PubMed:32142649, PubMed:32355176, PubMed:34526721, PubMed:38316879, PubMed:38600235, PubMed:38600236). 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).

ERCC8 Rabbit mAb - Protocols

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

- [Western Blot](#)
- [Blocking Peptides](#)
- [Dot Blot](#)
- [Immunohistochemistry](#)
- [Immunofluorescence](#)
- [Immunoprecipitation](#)
- [Flow Cytometry](#)
- [Cell Culture](#)

ERCC8 Rabbit mAb - Images

