

# **ERCC8 Rabbit mAb**

Catalog # AP78498

#### **Product Information**

Application WB, IP
Primary Accession Q13216
Reactivity Human
Host Rabbit

**Clonality** Monoclonal Antibody

Isotype IgG

**Conjugate** Unconjugated

**Immunogen** A synthesized peptide derived from human ERCC8

**Purification** Affinity Purified

Calculated MW 44055

### **Additional Information**

**Gene ID** 1161

Other Names ERCC8

**Dilution** WB~~1/500-1/1000 IP~~N/A

Format Liquid in 10mM PBS, pH 7.4, 150mM sodium chloride, 0.05% BSA, 0.02%

sodium azide and 50% glycerol.

**Storage** Store at 4°C short term. Aliquot and store at -20°C long term. Avoid

freeze/thaw cycles.

### **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, PubMed:16751180,

PubMed: 16964240, PubMed: 32142649, PubMed: 34526721,

PubMed:38316879, PubMed:38600235, PubMed: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, PubMed:16751180,

D. LAA. 14.0004040 D. LAA. 100440040 D. LAA. 100055476

PubMed:<u>16964240</u>, PubMed:<u>32142649</u>, PubMed:<u>32355176</u>, PubMed:<u>34526721</u>, PubMed:<u>38316879</u>, PubMed:<u>38600235</u>,

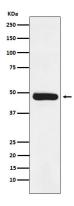
PubMed: 38600236). The CSA complex also promotes the ubiquitination and

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).

## **Images**



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