

## FANCM Rabbit pAb

CatalogNo: YN0712

### Key Features

#### Host Species

- Rabbit

#### Reactivity

- Human, Mouse

#### Applications

- WB, ELISA

#### MW

- 225kD (Observed)

#### Isotype

- IgG

### Recommended Dilution Ratios

WB 1:500-2000

ELISA 1:5000-20000

### Storage

**Storage\*** -15°C to -25°C/1 year (Do not lower than -25°C)

**Formulation** Liquid in PBS containing 50% glycerol, and 0.02% sodium azide.

### Basic Information

**Clonality** Polyclonal

### Immunogen Information

**Immunogen** Synthesized peptide derived from part region of human protein. AA 440-490

**Specificity** FANCM Polyclonal Antibody detects endogenous levels of protein.

### Target Information

**Gene name** FANCM KIAA1596

<b>Protein Name</b>	Fanconi anemia group M protein (Protein FANCM) (ATP-dependent RNA helicase FANCM) (Fanconi anemia-associated polypeptide of 250 kDa) (FAAP250) (Protein Hef ortholog)		
	<b>Organism</b>	<b>Gene ID</b>	<b>UniProt ID</b>
	Human	<a href="#">57697</a> ;	<a href="#">Q8IYD8</a> ;
	Mouse		<a href="#">Q8BGE5</a> ;
<b>Cellular Localization</b>	Nucleus .		
<b>Tissue specificity</b>	Expressed in germ cells of fetal and adult ovaries. In fetal ovaries, it is present in oogonia but expression is stronger in pachytene stage oocytes. Expressed in oocytes arrested at the diplotene stage of prophase I during the last trimester of pregnancy and in adults (PubMed:29231814). Expressed in the testis (PubMed:30075111).		
<b>Function</b>	Disease:Defects in FANCM are a cause of Fanconi anemia (FA) [MIM:227650]. FA is a genetically heterogeneous, autosomal recessive disorder characterized by progressive pancytopenia, a diverse assortment of congenital malformations, and a predisposition to the development of malignancies. At the cellular level it is associated with hypersensitivity to DNA-damaging agents, chromosomal instability (increased chromosome breakage), and defective DNA repair.,Function:ATPase required for FANCD2 ubiquitination, a key reaction in DNA repair. Binds to ssDNA but not to dsDNA.,PTM:Phosphorylated; hyperphosphorylated in response to genotoxic stress.,sequence Caution:Intron retention.,similarity:Belongs to the DEAD box helicase family. DEAH subfamily.,similarity:Contains 1 helicase ATP-binding domain.,similarity:Contains 1 helicase C-terminal domain.,subunit:Belongs to the multisubunit FA complex composed of FANCA, FANCB, FANCC, FANCE, FANCF, FANCG, FANCL/PHF9, FANCM and FAAP24. The complex is not found in FA patients. Interacts with FAAP24. Interacts with EME1.,		

## | Validation Data

## | Contact information

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