

PMS2-FC recombinant protein

YD3031 Catalog No:

Reactivity: Human;

>90% as determined by SDS-PAGE **Purity:**

Gene Name: PMS2

Mismatch repair endonuclease PMS2 (EC 3.1.-.-) (DNA mismatch repair protein **Protein Name:**

PMS2) (PMS1 protein homolog 2)

Amino acid:391-440, with FC tag. Sequence:

Human Gene Id: 5395

Human Swiss Prot

No:

P54278

Formulation: Phosphate-buffered solution

Source: Mammalian cells

-15°C to -25°C/1 year(Avoid freeze / thaw cycles) **Storage Stability:**

Function: Component of the post-replicative DNA mismatch repair system (MMR)

> (PubMed:30653781, PubMed:35189042). Heterodimerizes with MLH1 to form MutL alpha. DNA repair is initiated by MutS alpha (MSH2-MSH6) or MutS beta (MSH2-MSH3) binding to a dsDNA mismatch, then MutL alpha is recruited to the heteroduplex. Assembly of the MutL-MutS-heteroduplex ternary complex in presence of RFC and PCNA is sufficient to activate endonuclease activity of PMS2. It introduces single-strand breaks near the mismatch and thus generates new entry points for the exonuclease EXO1 to degrade the strand containing the mismatch. DNA methylation would prevent cleavage and therefore assure that only the newly mutated DNA strand is going to be corrected. MutL alpha

> (MLH1-PMS2) interacts physically with the clamp loader subunits of DNA polymerase III, suggesting that it may play a role to recruit the DNA polymerase III

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Subcellular Location:

Nucleus.



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