

# PMS2

Rabbit Monoclonal antibody(Mab) Catalog # AD80243

### **Product Information**

| Application       | IHC-P         |
|-------------------|---------------|
| Primary Accession | <u>P54278</u> |
| Reactivity        | Human         |
| Host              | Rabbit        |
| Clonality         | Monoclonal    |
| Clone Names       | 157G1F5       |
| Calculated MW     | 95797         |

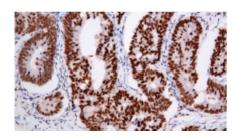
## **Additional Information**

| Gene ID<br>Gene Name<br>Other Names | 5395<br>PMS2 ( <u>HGNC:9122</u> )<br>Mismatch repair endonuclease PMS2, 3.1, DNA mismatch repair protein<br>PMS2, PMS1 protein homolog 2, PMS2 ( <u>HGNC:9122</u> ) |
|-------------------------------------|---|
| Dilution                            | IHC-P~~Ready-to-use   |
| Storage                             | Maintain refrigerated at 2-8°C.   |
| Precautions                         | PMS2 Antibody is for research use only and not for use in diagnostic or therapeutic procedures.   |

### **Protein Information**

| Name     | PMS2 ( <u>HGNC:9122</u> )   |
|----------|---|
| Function | Component of the post-replicative DNA mismatch repair system (MMR)<br>(PubMed: <u>30653781</u> , PubMed: <u>35189042</u> ). Heterodimerizes with MLH1 to form<br>MutL alpha. DNA repair is initiated by MutS alpha (MSH2-MSH6) or MutS beta<br>(MSH2-MSH3) binding to a dsDNA mismatch, then MutL alpha is recruited to<br>the heteroduplex. Assembly of the MutL-MutS-heteroduplex ternary complex<br>in presence of RFC and PCNA is sufficient to activate endonuclease activity of<br>PMS2. It introduces single-strand breaks near the mismatch and thus<br>generates new entry points for the exonuclease EXO1 to degrade the strand<br>containing the mismatch. DNA methylation would prevent cleavage and<br>therefore assure that only the newly mutated DNA strand is going to be<br>corrected. MutL alpha (MLH1-PMS2) interacts physically with the clamp loader<br>subunits of DNA polymerase III, suggesting that it may play a role to recruit<br>the DNA polymerase III to the site of the MMR. Also implicated in DNA<br>damage signaling, a process which induces cell cycle arrest and can lead to<br>apoptosis in case of major DNA damages. Possesses an ATPase activity, but in<br>the absence of gross structural changes, ATP hydrolysis may not be necessary |

## Images



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