



## **Anti-MIF polyclonal antibody (DPABY-723)**

This product is for research use only and is not intended for diagnostic use.

## PRODUCT INFORMATION

| Antigen Description | This gene encodes a lymphokine involved in cell-mediated immunity, immunoregulation, and inflammation. It plays a role in the regulation of macrophage function in host defense through the suppression of anti-inflammatory effects of glucocorticoids. This lymphokine and the JAB1 protein form a complex in the cytosol near the peripheral plasma membrane, which may indicate an additional role in integrin signaling pathways. [provided by RefSeq] |
|---------------------|---|
| Immunogen           | Recombinant fragment corresponding to a region within amino acids 1 and 115 of MIF (Uniprot ID#P14174)  |
| Isotype             | IgG   |
| Source/Host         | Rabbit  |
| Species Reactivity  | Human   |
| Purification        | Purified by antigen-affinity chromatography.  |
| Conjugate           | Unconjugated  |
| Applications        | ELISA Pr*, ICC/IF, IHC-P, WB  |
| Molecular Weight    | 12 kDa  |
| Positive Control    | 293T, A431, H1299, HeLaS3, HepG2, Molt-4, Raji, NIH-3T3   |
| Format              | Liquid  |
| Concentration       | 1 mg/ml   |
| Size                | 25 μΙ   |
| Buffer              | 0.1M Tris, 0.1M Glycine, 10% Glycerol (pH7). 0.01% Thimerosal was added as a preservative.  |

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| Preservative | None  |
|--------------|---|
| Storage      | Keep as concentrated solution. Aliquot and store at -31°C or below. Avoid multiple freezethaw cycles. |

## **GENE INFORMATION**

| Gene Name           | MIF macrophage migration inhibitory factor (glycosylation-inhibiting factor) [ Homo sapiens (human) ]   |
|---------------------|---|
| Official Symbol     | MIF   |
| Synonyms            | MIF; macrophage migration inhibitory factor (glycosylation-inhibiting factor); GIF; GLIF; MMIF; macrophage migration inhibitory factor; L-dopachrome isomerase; L-dopachrome tautomerase; phenylpyruvate tautomerase; |
| Entrez Gene ID      | 4282  |
| Protein Refseq      | NP 002406   |
| UniProt ID          | <u>I4AY87</u>   |
| Chromosome Location | 22q11.23  |
| Pathway             | Adipogenesis; Phenylalanine metabolism; Spinal Cord Injury; Tyrosine metabolism;  |
| Function            | chemoattractant activity; cytokine activity; cytokine receptor binding; dopachrome isomerase activity; phenylpyruvate tautomerase activity; protein binding; receptor binding;  |