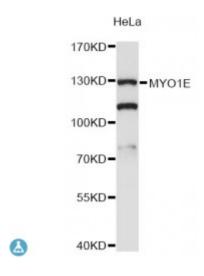


## **Anti-MYO1E Antibody**



**Description** This gene encodes a member of the nonmuscle class I myosins which are a

subgroup of the unconventional myosin protein family. The  $\,$ 

unconventional myosin proteins function as actin-based molecular motors. Class I myosins are characterized by a head (motor) domain, a regulatory domain and a either a short or long tail domain. Among the class I myosins, this protein is distinguished by a long tail domain that is involved

in crosslinking actin filaments. This protein localizes to the cytoplasm and may be involved in intracellular movement and membrane trafficking.

Mutations in this gene are the cause of focal segmental

glomerulosclerosis-6. This gene has been referred to as myosin IC in the literature but is distinct from the myosin IC gene located on chromosome

17.

Model STJ114999

**Host** Rabbit

**Reactivity** Human

**Applications** WB

Immunogen Recombinant fusion protein containing a sequence corresponding to amino

acids 809-1108 of human MYO1E (NP\_004989.2).

**Gene ID** 4643

Gene Symbol MYO1E

**Dilution range** WB 1:1000 - 1:2000

**Tissue Specificity** Expressed in the immune system, In the kidney, predominantly expressed in

the glomerulus, including podocytes

**Purification** Affinity purification

**Note** For Research Use Only (RUO).

Protein Name Unconventional myosin-Ie Myosin-Ic Unconventional myosin 1E

Molecular Weight 127.062 kDa

**Clonality** Polyclonal

**Conjugation** Unconjugated

**Isotype** IgG

**Formulation** PBS with 0.02% sodium azide, 50% glycerol, pH7.3.

**Storage Instruction** Store at -20C. Avoid freeze / thaw cycles.

Database Links <u>HGNC:7599OMIM:601479</u>

Alternative Names Unconventional myosin-Ie Myosin-Ic Unconventional myosin 1E

**Function** Myosins are actin-based motor molecules with ATPase activity,

Unconventional myosins serve in intracellular movements, Their highly divergent tails bind to membranous compartments, which are then moved

relative to actin filaments, Binds to membranes containing anionic

phospholipids via its tail domain, Required for normal morphology of the glomerular basement membrane, normal development of foot processes by kidney podocytes and normal kidney function, In dendritic cells, may control the movement of class II-containing cytoplasmic vesicles along the actin cytoskeleton by connecting them with the actin network via ARL14EP and

ARL14,

Cellular Localization Cytoplasm, Cytoplasm, cytoskeleton, Cytoplasmic vesicle, Cytoplasmic

vesicle, clathrin-coated vesicle, Cell junction,

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