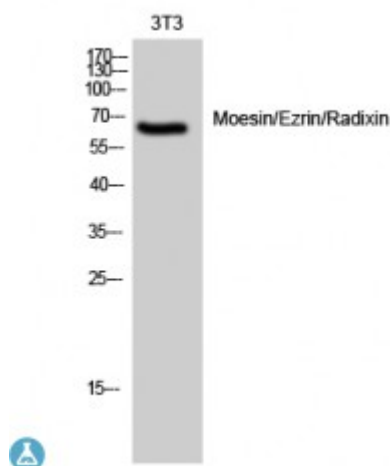


## Anti-Moesin/Ezrin/Radixin antibody



<b>Description</b>	Rabbit polyclonal to Moesin/Ezrin/Radixin.
<b>Model</b>	STJ94178
<b>Host</b>	Rabbit
<b>Reactivity</b>	Human, Mouse, Rat
<b>Applications</b>	ELISA, IHC, WB
<b>Immunogen</b>	Synthesized peptide derived from human Moesin/Ezrin/Radixin around the non-phosphorylation site of T558.
<b>Immunogen Region</b>	500-580 aa
<b>Gene ID</b>	<a href="#">4478</a>
<b>Gene Symbol</b>	<a href="#">MSN</a>
<b>Dilution range</b>	WB 1:500-1:2000IHC 1:100-1:300ELISA 1:20000
<b>Specificity</b>	Moesin/Ezrin/Radixin Polyclonal Antibody detects endogenous levels of Moesin/Ezrin/Radixin protein.
<b>Tissue Specificity</b>	In all tissues and cultured cells studied.
<b>Purification</b>	The antibody was affinity-purified from rabbit antiserum by affinity-chromatography using epitope-specific immunogen.
<b>Note</b>	For Research Use Only (RUO).
<b>Protein Name</b>	Moesin Membrane-organizing extension spike protein
<b>Molecular Weight</b>	67 kDa
<b>Clonality</b>	Polyclonal

<b>Conjugation</b>	Unconjugated
<b>Isotype</b>	IgG
<b>Formulation</b>	Liquid in PBS containing 50% glycerol, 0.5% BSA and 0.02% sodium azide.
<b>Concentration</b>	1 mg/ml
<b>Storage Instruction</b>	Store at -20°C, and avoid repeat freeze-thaw cycles.
<b>Database Links</b>	<a href="https://www.ncbi.nlm.nih.gov/Protein/300988">HGNC:7373OMIM:300988</a>
<b>Alternative Names</b>	Moesin Membrane-organizing extension spike protein
<b>Function</b>	Probably involved in connections of major cytoskeletal structures to the plasma membrane. May inhibit herpes simplex virus 1 infection at an early stage. Plays a role in regulating the proliferation, migration, and adhesion of human lymphoid cells and participates in immunologic synapse formation .
<b>Sequence and Domain Family</b>	The [IL]-x-C-x-x-[DE] motif is a proposed target motif for cysteine S-nitrosylation mediated by the iNOS-S100A8/A9 transnitrosylase complex.
<b>Cellular Localization</b>	Cell membrane Cytoplasm, cytoskeleton Apical cell membrane Cell projection, microvillus membrane. Phosphorylated form is enriched in microvilli-like structures at apical membrane . Increased cell membrane localization of both phosphorylated and non-phosphorylated forms seen after thrombin treatment.
<b>Post-translational Modifications</b>	Phosphorylation on Thr-558 is crucial for the formation of microvilli-like structures. Phosphorylation by ROCK2 suppresses the head-to-tail association of the N-terminal and C-terminal halves resulting in an opened conformation which is capable of actin and membrane-binding . Phosphorylation on Thr-558 by STK10 negatively regulates lymphocyte migration and polarization. S-nitrosylation of Cys-117 is induced by interferon-gamma and oxidatively-modified low-density lipoprotein (LDL(ox)) implicating the iNOS-S100A8/9 transnitrosylase complex.